# Package 'miesmuschel'

July 9, 2024

Title Mixed Integer Evolution Strategies

Description Evolutionary black box optimization algorithms building on the 'bbotk' package. 'miesmuschel' offers both ready-to-use optimization algorithms, as well as their fundamental building blocks that can be used to manually construct specialized optimization loops. The Mixed Integer Evolution Strategies as described by Li et al. (2013) <doi:10.1162/EVCO\_a\_00059> can be implemented, as well as the multi-objective optimization algorithms NSGA-II by Deb, Pratap, Agarwal, and Meyarivan (2002) <doi:10.1109/4235.996017>.

URL https://github.com/mlr-org/miesmuschel

BugReports https://github.com/mlr-org/miesmuschel/issues

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Collate 'dictionaries.R' 'MiesOperator.R' 'Filtor.R' 'FiltorMaybe.R'

'FiltorNull.R' 'FiltorProxy.R' 'FiltorSurrogate.R'

'FiltorSurrogateProgressive.R' 'FiltorSurrogateTournament.R'

'Mutator.R' 'MutatorCmpMaybe.R' 'MutatorDiscreteUniform.R'

'MutatorErase.R' 'MutatorGauss.R' 'MutatorMaybe.R'

'MutatorNull.R' 'MutatorProxy.R' 'MutatorSequential.R'

'ParamSetShadow.R' 'OperatorCombination.R' 'Recombinator.R'

'Selector.R' 'mies\_methods.R' 'OptimizerMies.R'

'RecombinatorCmpMaybe.R' 'RecombinatorConvex.R'

'RecombinatorConvexPair.R' 'RecombinatorCrossoverNary.R'

'RecombinatorCrossoverUniform.R' 'RecombinatorMaybe.R'

'RecombinatorNull.R' 'RecombinatorProxy.R'

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'TerminatorGenerationStagnation.R' 'TerminatorGenerations.R'
'TunerMies.R' 'bibentries.R' 'repr.R' 'utils.R' 'utils_mo.R'
'zzz.R'
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miesmuschel-package miesmuschel: Mixed Integer Evolution Strategies

# **Description**

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miesmuschel offers both an Optimizer and a Tuner for general MIES-optimization, as well as all the building blocks for building a custom optimization algorithm that is more flexible and can be used for research into novel evolution strategies.

The call-graph of the default algorithm in OptimizerMies / TunerMies is as follows, and is shown here as an overview over the mies\_\* functions, and how they are usually connected. (Note that only the exported mies\_\* functions are shown.) See the help information of these functions for more info.

```
OptimizerMies$.optimize(inst)
|- mies_prime_operators()  # prime operators on instance's search_space
|- mies_init_population()  # sample and evaluate first generation
| `- mies_evaluate_offspring()  # evaluate sampled individuals
| `- inst$eval_batch()  # The OptimInst's evaluation method
`- repeat # Repeat the following until terminated
```

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#### See Also

Useful links:

- https://github.com/mlr-org/miesmuschel
- Report bugs at https://github.com/mlr-org/miesmuschel/issues

crate\_env

Set a Function's Environment

#### **Description**

Useful to represent functions efficiently within repr().

## Usage

```
crate_env(fun, namespace = "R_GlobalEnv", ..., selfref = NULL)
```

# **Arguments**

fun (function)

Function of which the environment should be set.

namespace (character(1))

Name of the namespace, as given by <a href="mailto:environmentName">environmentName</a>(), to be used as the (parent of the) environment of fun. Special values "R\_GlobalEnv" (global environment), "R\_BaseEnv" (base environment; note this one is non-standard within R), "R\_EmptyEnv" (empty environment). The content of the namespace is not be a life to the life of the

modified. Default "R\_GlobalEnv".

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```
Content of environments within which to place fun.

selfref (character(1) | named integer | NULL)

If character(1): The name of the entry of the first element in . . . that refers to the function itself. If a named integer, then the values indicate the lists in . . . where the reference to the function should be placed see examples. Default NULL: No reference to the function itself is present.
```

#### Value

function: The given fun with changed environment.

```
identity2 = crate_env(function(x) x, "base")
identical(identity, identity2) # TRUE
y = 1
f1 = mlr3misc::crate(function(x) x + y, y, .parent = .GlobalEnv)
f2 = crate_{env}(function(x) x + y, "R_GlobalEnv", list(y = 1))
# Note identical() does not apply because both contain (equal, but not
# identical) 'y = 1'-environments
all.equal(f1, f2) # TRUE
f1(10) # 10 + 1 == 11
factorial1 = mlr3misc::crate(
  function(x) if (x > 0) x * factorial1(x - 1) else 1,
  y, .parent = .GlobalEnv
)
environment(factorial1)$factorial1 = factorial1
factorial2 = crate_env(
  function(x) if (x > 0) x * factorial1(x - 1) else 1,
  "R_GlobalEnv", list(y = 1), selfref = "factorial1")
# putting 'factorial1' into the list (or repeating function(x) ....)
# would *not* work, since we want:
identical(environment(factorial2)$factorial1, factorial2) # TRUE
all.equal(factorial1, factorial2) # TRUE
g = crate_env(function(x) x + y + z, "miesmuschel",
  list(y = 1), list(z = 2), selfref = c(X = 1, Y = 2, Z = 2))
g(0) # 0 + 1 + 2 == 3
identical(environment(g)$X, g)
identical(parent.env(environment(g))$Y, g)
identical(parent.env(environment(g))$Z, g)
identical(
  parent.env(parent.env(environment(g))),
  loadNamespace("miesmuschel")
)
```

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 ${\sf dict\_filtors}$ 

Dictionary of Filtors

#### **Description**

Dictionary of Filtors

# Usage

dict\_filtors

#### **Format**

An object of class DictionaryFiltor (inherits from DictionaryEx, Dictionary, R6) of length 15.

#### Methods

Methods inherited from Dictionary, as well as:

```
    help(key, help_type)
        (character(1), character(1))
        Displays help for the dictionary entry key. help_type is one of "text", "html", "pdf" and given as the help_type argument of R's help().
```

## See Also

Other dictionaries: dict\_mutators, dict\_recombinators, dict\_scalors, dict\_selectors, mut()

dict\_filtors\_maybe

Filtor-Combination that Filters According to Two Filtors

# **Description**

Filtor that wraps two other Filtors given during construction and chooses which operation to perform. Each of the resulting n\_filter individuals is chosen either from \$filtor, or from \$filtor\_not.

This makes it possible to implement filter methods such as random interleaving, where only a fraction of p individuals were filtered and the others were not.

Letting the number of individuals chosen by \$filtor be n\_filter\_f, then n\_filter\_f is either fixed set to round(n\_filter \* p), (when random\_choise is FALSE) or to rbinom(1, n\_filter, p) (when random\_choice is TRUE).

When random\_choice is FALSE, then \$needed\_input() is calculated directly from \$needed\_input() of \$filtor and \$filtor\_not, as well as n\_filter\_f and n\_filter - n\_filter\_f.

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When random\_choice is TRUE, then \$needed\_input() is considers the "worst case" from \$filtor and \$filtor\_not, and assumes that \$needed\_input() is monotonically increasing in its input argument.

To make the worst case less extreme, the number of individuals chosen with random\_choice set to TRUE is limited to  $qbinom(-20, n_filter, p, log.p = TRUE)$  (with lower.tail FALSE and TRUE for  $filter_not$ , respectively), which distorts the binomial distribution with probability 1 - exp(-20) or about 1 - 0.5e-9.

# **Configuration Parameters**

This operator has the configuration parameters of the Filtors that it wraps: The configuration parameters of the operator given to the filtor construction argument are prefixed with "maybe.", the configuration parameters of the operator given to the filtor\_not construction argument are prefixed with "maybe\_not.".

Additional configuration parameters:

- p::numeric(1)
  Probability per individual (when random\_choise is TRUE), or fraction of individuals (when random\_choice is FALSE), that are chosen from \$filtor instead of \$filtor\_not. Must be set by the user.
- random\_choice :: logical(1)
  Whether to sample the number of individuals chosen by \$filtor according to rbinom(1, n\_filter, p), or to use a fixed fraction. Initialized to FALSE.

# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of filtor and filtor\_not.

# **Dictionary**

This Filtor can be created with the short access form ftr() (ftrs() to get a list), or through the the dictionary dict\_filtors in the following way:

```
# preferred:
ftr("maybe", <filtor> [, <filtor_not>])
ftrs("maybe", <filtor> [, <filtor_not>]) # takes vector IDs, returns list of Filtors
# long form:
dict_filtors$get("maybe", <filtor> [, <filtor_not>])
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Filtor -> FiltorMaybe
```

# **Active bindings**

```
filtor (Filtor)
```

Filtor being wrapped. This operator gets run with probability / proportion p (configuration parameter).

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```
filtor_not (Filtor)
```

Alternative Filtor being wrapped. This operator gets run with probability / proportion 1 - p (configuration parameter).

# Methods

#### **Public methods:**

- FiltorMaybe\$new()
- FiltorMaybe\$prime()
- FiltorMaybe\$clone()

Method new(): Initialize the FiltorMaybe object.

```
Usage:
```

```
FiltorMaybe$new(filtor, filtor_not = FiltorNull$new())
```

#### Arguments:

```
filtor (Filtor)
```

Filtor to wrap. This operator gets run with probability p (Configuration parameter).

The constructed object gets a *clone* of this argument. The \$filtor field will reflect this value.

```
filtor_not (Filtor)
```

Another Filtor to wrap. This operator runs when filtor is not chosen. By default, this is FiltorNull, i.e. no filtering. With this default, the FiltorMaybe object applies the filtor operation with probability / proportion p, and no operation at all otherwise.

The constructed object gets a *clone* of this argument. The \$filtor\_not field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to filtor and filtor\_not during construction.

```
Usage:
```

```
FiltorMaybe$prime(param_set)
```

Arguments:

```
param_set (ParamSet)
```

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage

```
FiltorMaybe$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

#### See Also

```
Other filtors: Filtor, FiltorSurrogate, dict_filtors_null, dict_filtors_proxy, dict_filtors_surprog, dict_filtors_surtour
```

Other filtor wrappers: dict\_filtors\_proxy

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# **Examples**

```
library("mlr3")
library("mlr3learners")

fm = ftr("maybe", ftr("surprog", lrn("regr.lm"), filter.pool_factor = 2), p = 0.5)
p = ps(x = p_dbl(-5, 5))
known_data = data.frame(x = 1:5)
fitnesses = 1:5
new_data = data.frame(x = c(0.5, 1.5, 2.5, 3.5, 4.5))

fm$prime(p)

fm$needed_input(2)

fm$operate(new_data, known_data, fitnesses, 2)

fm$param_set$values$p = 0.33

fm$needed_input(3)

fm$operate(new_data, known_data, fitnesses, 3)
```

dict\_filtors\_null

Null-Filtor

# Description

Null-filtor that does not perform filtering. Its needed\_input() is always the output\_size, and operate() selects the first n\_filter values from its input.

Useful in particular with operator-wrappers such as FiltorProxy, and to make filtering optional.

# **Configuration Parameters**

This operator has no configuration parameters.

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Filtor can be created with the short access form ftr() (ftrs() to get a list), or through the the dictionary dict\_filtors in the following way:

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```
# preferred:
ftr("null")
ftrs("null") # takes vector IDs, returns list of Filtors
# long form:
dict_filtors$get("null")
```

# Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Filtor-> FiltorNull
```

# Methods

# **Public methods:**

- FiltorNull\$new()
- FiltorNull\$clone()

**Method** new(): Initialize the FiltorNull object.

Usage:

FiltorNull\$new()

Method clone(): The objects of this class are cloneable with this method.

Usage

FiltorNull\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

# See Also

```
\label{lem:optimizer} Other\ filtors: FiltorSurrogate, dict\_filtors\_maybe, dict\_filtors\_proxy, dict\_filtors\_surprog, dict\_filtors\_surtour
```

```
fn = ftr("null")

p = ps(x = p_dbl(-5, 5))
known_data = data.frame(x = 1:5)
fitnesses = 1:5

new_data = data.frame(x = c(2.5, 4.5))

fn$prime(p)

fn$needed_input(1)

fn$operate(new_data, known_data, fitnesses, 1)
```

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dict\_filtors\_proxy

Proxy-Filtor that Filters According to its Configuration Parameter

# **Description**

Filtor that performs the operation in its operation configuration parameter. This can be used to make filtor operations fully parametrizable.

# **Configuration Parameters**

• operation :: Filtor

Operation to perform. Must be set by the user. This is primed when \$prime() of SelectorProxy is called, and also when \$perate() is called, to make changing the operation as part of self-adaption possible. However, if the same operation gets used inside multiple SelectorProxy objects, then it is recommended to \$clone(deep = TRUE) the object before assigning them to operation to avoid frequent re-priming.

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("proxy")
sels("proxy") # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("proxy")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Filtor -> FiltorProxy
```

# Methods

# **Public methods:**

- FiltorProxy\$new()
- FiltorProxy\$prime()
- FiltorProxy\$clone()

**Method** new(): Initialize the FiltorProxy object.

Usage:

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```
FiltorProxy$new()
```

Usage:

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the operator given to the operation configuration parameter. Note that this modifies the \$param\_set\$values\$operation object.

```
FiltorProxy$prime(param_set)

Arguments:

param_set (ParamSet)
    Passed to MiesOperator$prime().

Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.

Usage:
FiltorProxy$clone(deep = FALSE)

Arguments:
deep Whether to make a deep clone.
```

#### See Also

```
Other filtors: FiltorSurrogate, dict_filtors_maybe, dict_filtors_null, dict_filtors_surprog, dict_filtors_surtour

Other filtor wrappers: dict_filtors_maybe
```

```
library("mlr3")
library("mlr3learners")
fp = ftr("proxy")
p = ps(x = p_dbl(-5, 5))
known_data = data.frame(x = 1:5)
fitnesses = 1:5
new_data = data.frame(x = c(2.5, 4.5))

fp$param_set$values$operation = ftr("null")
fp$prime(p)
fp$operate(new_data, known_data, fitnesses, 1)

fp$param_set$values$operation = ftr("surprog", lrn("regr.lm"), filter.pool_factor = 2)
fp$operate(new_data, known_data, fitnesses, 1)
```

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# **Description**

Performs progressive surrogate model filtering. A surrogate model is used, as described in the parent class FiltorSurrogate. The filtering is "progressive" in that successive values are filtered more agressively.

# Algorithm

Given the number n\_filter of of individuals to sample, and the desired pool size at round i pool\_size(i), progressive surrogate model filtering proceeds as follows:

- 1. Train the surrogate\_learner LearnerRegr on the known\_values and their fitnesses.
- 2. Take pool\_size(1) configurations, predict their expected performance using the surrogate model, and put them into a pool P of configurations to consider.
- 3. Initialize i to 1.
- 4. Take the individual that is optimal according to predicted performance, remove it from P and add it to solution set S.
- 5. If the number of solutions in S equals n\_filter, quit.
- 6. If pool\_size(i + 1) is larger than pool\_size(i), take the next pool\_size(i + 1) pool\_size(i) configurations, predict their expected performance using the surrogate model, and add them to P. Otherwise, remove pool\_size(i) pool\_size(i + 1) random individuals from the pool. The size of P ends up being pool\_size(i + 1) i, as i individuals have also been removed and added to S.
- 7. Increment i, jump to 4.

(The algorithm presented here is optimized for clarity; the actual implementation does all the surrogate model prediction in one go, but is functionally equivalent).

pool\_size(i) is calculated as round(n\_filter \* pool\_factor \* (pool\_factor\_last / pool\_factor)
^ (i / n\_filter)), i.e. a log-linear interpolation from pool\_factor \* n\_filter to pool\_factor\_last
\* n\_filter.

The pool\_factor and pool\_factor\_last configuration parameters of this algorithm determine how agressively the surrogate model is used to filter out sampled configurations. If the filtering is agressive (large values), then more "exploitation" at the cost of "exploration" is performed. When pool\_factor is small but pool\_factor\_last is large (or vice-versa), then different individuals are filtered with different agressiveness, potentially leading to a tradeoff between "exploration" and "exploitation".

When pool\_factor\_last is set, it defaults to pool\_factor, with no new individuals added and no individuals removed from the filter pool during filtering. It is equivalent to taking the top n\_filter individuals out of a sample of n\_filter \* pool\_factor.

dict\_filtors\_surprog

#### **Configuration Parameters**

FiltorSurrogateProgressive's configuration parameters are the hyperparameters of the FiltorSurrogate base class, as well as:

- filter.pool\_factor :: numeric(1) pool\_factor parameter of the progressive surrogate model filtering algorithm, see the corresponding section. Initialized to 1. Together with the default of filter.pool\_factor\_last, this is equivalent to random sampling new individuals.
- filter.pool\_factor\_last :: numeric(1) pool\_factor\_last parameter of the progressive surrogate model filtering algorithm, see the corresponding section. When not given, it defaults to filter.pool\_factor, equivalent to taking the top n\_filter from n\_filter \* pool\_factor individuals.

# **Supported Operand Types**

See FiltorSurrogate about supported operand types.

# **Dictionary**

This Filtor can be created with the short access form ftr() (ftrs() to get a list), or through the the dictionary dict\_filtors in the following way:

```
# preferred:
ftr("surprog", <surrogate_learner> [, <surrogate_selector>])
ftrs("surprog", <surrogate_learner> [, <surrogate_selector>]) # takes vector IDs, returns list of Filt
# long form:
dict_filtors$get("surprog", <surrogate_learner> [, <surrogate_selector>])
```

# **Super classes**

```
miesmuschel::MiesOperator -> miesmuschel::Filtor -> miesmuschel::FiltorSurrogate -
> FiltorSurrogateProgressive
```

#### Methods

# **Public methods:**

- FiltorSurrogateProgressive\$new()
- FiltorSurrogateProgressive\$clone()

Method new(): Initialize the FiltorSurrogateProgressive.

```
Usage:
FiltorSurrogateProgressive$new(
   surrogate_learner,
   surrogate_selector = SelectorBest$new()
)
Arguments:
```

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```
surrogate_learner (mlr3::LearnerRegr)
   Regression learner for the surrogate model filtering algorithm.
   The $surrogate_learner field will reflect this value.
surrogate_learner (mlr3::LearnerRegr)
   Regression learner for the surrogate model filtering algorithm.
   The $surrogate_learner field will reflect this value.
surrogate_selector (Selector) Selector for the surrogate model filtering algorithm.
   The $surrogate_selector field will reflect this value.
surrogate_selector (Selector) Selector for the surrogate model filtering algorithm.
   The $surrogate_selector field will reflect this value.

Method clone(): The objects of this class are cloneable with this method.

Usage:
FiltorSurrogateProgressive$clone(deep = FALSE)

Arguments:
deep Whether to make a deep clone.
```

# See Also

```
Other filtors: Filtor, FiltorSurrogate, dict_filtors_maybe, dict_filtors_null, dict_filtors_proxy, dict_filtors_surtour
```

# **Examples**

```
library("mlr3")
library("mlr3learners")
fp = ftr("surprog", lrn("regr.lm"), filter.pool_factor = 2)

p = ps(x = p_dbl(-5, 5))
known_data = data.frame(x = 1:5)
fitnesses = 1:5
new_data = data.frame(x = c(2.5, 4.5))

fp$prime(p)

fp$needed_input(1)

fp$operate(new_data, known_data, fitnesses, 1)
```

### **Description**

Performs tournament surrogate model filtering. A surrogate model is used, as described in the parent class FiltorSurrogate.

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### Algorithm

Selects individuals from a tournament by taking the top per\_tournament individuals, according to surrogate\_selector and as predicted by surrogate\_learner, from a sample of tournament\_size(i), where tournament\_size(1) is given by tournament\_size, tournament\_size(ceiling(n\_filter / per\_tournament)) is given by tournament\_size\_last, and tournament\_size(i) for i between these values is linearly interpolated on a log scale.

# **Configuration Parameters**

FiltorSurrogateProgressive's configuration parameters are the hyperparameters of the FiltorSurrogate base class, as well as:

- filter.per\_tournament :: integer(1)

  Number of individuals to select from each tournament. If per\_tournament is not a divider of n\_filter, then the last tournament selects a random subset of n\_filter %% per\_tournament individuals out of the top per\_tournament individuals. Initialized to 1.
- filter.tournament\_size :: numeric(1)

  Tournament size used for filtering. If tournament\_size\_last is not given, all n\_filter individuals are selected in batches of per\_tournament from tournaments of this size. If it is given, then the actual tournament size is interpolated between tournament\_size and tournament\_size\_last on a logarithmic scale. Tournaments with tournament size below per\_tournament select per\_tournament individuals without tournament, i.e. no filtering. Initialized to 1.
- filter.tournament\_size\_last :: numeric(1)

  Tournament size used for the last tournament, see description of tournament\_size. Defaults to tournament\_size when not given, i.e. all tournaments have the same size.

#### **Supported Operand Types**

See FiltorSurrogate about supported operand types.

# **Dictionary**

This Filtor can be created with the short access form ftr() (ftrs() to get a list), or through the the dictionary dict\_filtors in the following way:

```
# preferred:
ftr("surtour", <surrogate_learner> [, <surrogate_selector>])
ftrs("surtour", <surrogate_learner> [, <surrogate_selector>])  # takes vector IDs, returns list of Filt
# long form:
dict_filtors$get("surtour", <surrogate_learner> [, <surrogate_selector>])
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Filtor -> miesmuschel::FiltorSurrogate -
> FiltorSurrogateTournament
```

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### Methods

#### **Public methods:**

```
• FiltorSurrogateTournament$new()
```

```
• FiltorSurrogateTournament$clone()
```

```
Method new(): Initialize the FiltorSurrogateTournament.
```

```
Usage:
FiltorSurrogateTournament$new(
  surrogate_learner.
  surrogate_selector = SelectorBest$new()
)
Arguments:
surrogate_learner (mlr3::LearnerRegr)
    Regression learner for the surrogate model filtering algorithm.
    The $surrogate_learner field will reflect this value.
surrogate_learner (mlr3::LearnerRegr)
    Regression learner for the surrogate model filtering algorithm.
    The $surrogate_learner field will reflect this value.
surrogate_selector (Selector) Selector for the surrogate model filtering algorithm.
    The $surrogate_selector field will reflect this value.
surrogate_selector (Selector) Selector for the surrogate model filtering algorithm.
    The $surrogate_selector field will reflect this value.
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

```
FiltorSurrogateTournament$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

#### See Also

```
Other filtors: Filtor, FiltorSurrogate, dict_filtors_maybe, dict_filtors_null, dict_filtors_proxy, dict_filtors_surprog
```

```
library("mlr3")
library("mlr3learners")
fp = ftr("surtour", lrn("regr.lm"), filter.tournament_size = 2)

p = ps(x = p_dbl(-5, 5))
known_data = data.frame(x = 1:5)
fitnesses = 1:5
new_data = data.frame(x = c(2.5, 4.5))

fp$prime(p)
```

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```
fp$needed_input(1)
fp$operate(new_data, known_data, fitnesses, 1)
```

dict\_mutators

Dictionary of Mutators

# **Description**

Dictionary of Mutators

# Usage

dict\_mutators

#### **Format**

An object of class DictionaryMutator (inherits from DictionaryEx, Dictionary, R6) of length 15.

#### Methods

Methods inherited from Dictionary, as well as:

help(key, help\_type)
 (character(1), character(1))
 Displays help for the dictionary entry key. help\_type is one of "text", "html", "pdf" and given as the help\_type argument of R's help().

# See Also

Other dictionaries: dict\_filtors, dict\_recombinators, dict\_scalors, dict\_selectors, mut()

```
dict_mutators_cmpmaybe
```

Mutator Choosing Action Component-Wise Independently

# Description

Mutator that chooses which operation to perform probabilistically. The Mutator wraps two other Mutators given during construction, and both of these operators are run. The ultimate result is sampled from the results of these operations independently for each individuum and component: with probability p (configuration parameter), the result from the Mutator given to the mutator construction argument is used, and with probability p-1 the one given to mutator\_not is used.

# **Configuration Parameters**

This operator has the configuration parameters of the Mutators that it wraps: The configuration parameters of the operator given to the mutator construction argument are prefixed with "cmpmaybe.", the configuration parameters of the operator given to the mutator\_not construction argument are prefixed with "cmpmaybe\_not.".

Additional configuration parameters:

• p:: numeric(1)
Probability per component with which to apply the operator given to the mutator construction argument. Must be set by the user.

# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of mutator and mutator\_not.

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("cmpmaybe", <mutator> [, <mutator_not>])
muts("cmpmaybe", <mutator> [, <mutator_not>]) # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("cmpmaybe", <mutator> [, <mutator_not>])
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Mutator -> MutatorCmpMaybe
```

# **Active bindings**

```
mutator (Mutator)
    Mutator being wrapped. This operator gets run with probability p (configuration parameter).
mutator_not (Mutator)
    Alternative Mutator being wrapped. This operator gets run with probability 1 - p (configuration parameter).
```

#### Methods

# **Public methods:**

- MutatorCmpMaybe\$new()
- MutatorCmpMaybe\$prime()
- MutatorCmpMaybe\$clone()

**Method** new(): Initialize the MutatorCmpMaybe object.

Usage:

```
MutatorCmpMaybe$new(mutator, mutator_not = MutatorNull$new())
       Arguments:
       mutator (Mutator)
           Mutator to wrap. Component-wise results of this operator are used with probability p
           (Configuration parameter).
           The constructed object gets a clone of this argument. The $mutator field will reflect this
           value.
       mutator_not (Mutator)
           Another Mutator to wrap. Results from this operator are used when mutator is not chosen.
           By default, this is MutatorNull, i.e. no operation.
           With this default, the MutatorCmpMaybe object applies the mutator operation with proba-
           bility p, and no operation at all otherwise.
           The constructed object gets a clone of this argument. The $mutator_not field will reflect
           this value.
     Method prime(): See MiesOperator method. Primes both this operator, as well as the wrapped
     operators given to mutator and mutator_not during construction.
       Usage:
       MutatorCmpMaybe$prime(param_set)
       Arguments:
       param_set (ParamSet)
           Passed to MiesOperator$prime().
       Returns: invisible self.
     Method clone(): The objects of this class are cloneable with this method.
       Usage:
       MutatorCmpMaybe$clone(deep = FALSE)
       Arguments:
       deep Whether to make a deep clone.
See Also
    Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict_mutators_erase,
    dict_mutators_gauss, dict_mutators_maybe, dict_mutators_null, dict_mutators_proxy,
    dict_mutators_sequential, dict_mutators_unif
    Other mutator wrappers: OperatorCombination, dict_mutators_maybe, dict_mutators_proxy,
    dict_mutators_sequential
```

```
set.seed(1)
mcm = mut("cmpmaybe", mut("gauss", sdev = 5), p = 0.5)
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5))
data = data.frame(x = rep(0, 5), y = rep(0, 5))
mcm$prime(p)
```

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```
mcm$operate(data)

mcm$param_set$values$p = 0.2

mcm$operate(data)

mcm2 = mut("cmpmaybe",
    mutator = mut("gauss", sdev = 0.01),
    mutator_not = mut("gauss", sdev = 10),
    p = 0.5
)

mcm2$prime(p)
mcm2$operate(data)
```

 ${\sf dict\_mutators\_erase}$ 

Uniform Sample Mutator

# Description

"Mutates" individuals by forgetting the current value and sampling new individuals from scratch. Since the information loss is very high, this should in most cases be combined with MutatorCmpMaybe.

# **Configuration Parameters**

• initializer :: function
Function that generates the initial population as a Design object, with arguments param\_set
and n, functioning like paradox::generate\_design\_random or paradox::generate\_design\_lhs.
This is equivalent to the initializer parameter of mies\_init\_population(), see there for
more information. Initialized to generate\_design\_random().

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

#### **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("erase")
muts("erase") # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("erase")
```

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# Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Mutator-> MutatorErase
```

#### Methods

#### **Public methods:**

- MutatorErase\$new()
- MutatorErase\$clone()

**Method** new(): Initialize the MutatorErase object.

Usage:

MutatorErase\$new()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

MutatorErase\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

```
Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict_mutators_cmpmaybe, dict_mutators_gauss, dict_mutators_maybe, dict_mutators_null, dict_mutators_proxy, dict_mutators_sequential, dict_mutators_unif
```

# Examples

```
set.seed(1)
mer = mut("erase")
p = ps(x = p_lgl(), y = p_fct(c("a", "b", "c")), z = p_dbl(0, 1))
data = data.frame(x = rep(TRUE, 5), y = rep("a", 5),
    z = seq(0, 1, length.out = 5),
    stringsAsFactors = FALSE) # necessary for R <= 3.6

mer$prime(p)
mer$operate(data)</pre>
```

dict\_mutators\_gauss

Gaussian Distribution Mutator

# **Description**

Individuals are mutated with an independent normal random variable on each component.

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# **Configuration Parameters**

• sdev :: numeric

Standard deviation of normal distribuion. This is absolute if sdev\_is\_relative is FALSE, and multiplied with each individual component's range (upper - lower) if sdev\_is\_relative is TRUE. This may either be a scalar, in which case it is applied to all input components, or a vector, in which case it must have the length of the input and applies to components in order in which they appear in the priming ParamSet. Must be set by the user.

- sdev\_is\_relative :: logical(1)
  Whether sdev is absolute (FALSE) or relative to component range (TRUE). Initialized to FALSE.
- truncated\_normal :: logical(1)
  Whether to draw individuals from a normal distribution that is truncated at the bounds of each component (TRUE), or to draw from a normal distribution and then restrict to bounds afterwards (FALSE). The former (TRUE) will lead to very few to no samples landing on the exact bounds (analytically it would be none almost surely, but this is subject to machine precision), the latter (FALSE) can lead to a substantial number of samples landing on the exact bounds. Initialized to FALSE.

# **Supported Operand Types**

```
Supported Domain classes are: p_int ('ParamInt'), p_dbl ('ParamDbl')
```

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("gauss")
muts("gauss") # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("gauss")
```

# Super classes

```
miesmuschel::Mutator -> miesmuschel::Mutator -> miesmuschel::MutatorNumeric -
> MutatorGauss
```

## Methods

### **Public methods:**

- MutatorGauss\$new()
- MutatorGauss\$clone()

**Method** new(): Initialize the MutatorGauss object.

```
Usage:
MutatorGauss$new()
```

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**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
MutatorGauss$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_erase, dict\_mutators\_maybe, dict\_mutators\_null, dict\_mutators\_proxy, dict\_mutators\_sequential, dict\_mutators\_unif

### **Examples**

```
set.seed(1)
mg = mut("gauss", sdev = 0.1)
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5))
data = data.frame(x = rep(0, 5), y = rep(0, 5))
mg$prime(p)
mg$operate(data)
mg$param_set$values$sdev = 100
mg$operate(data)
```

dict\_mutators\_maybe

Mutator Choosing Action Probabilistically

# Description

Mutator that chooses which operation to perform probabilistically. The Mutator wraps two other Mutators given during construction, and for each individuum, the operation to perform is sampled: with probability p (configuration parameter), the Mutator given to the mutator construction argument is applied, and with probability p-1 the one given to mutator\_not is applied.

# **Configuration Parameters**

This operator has the configuration parameters of the Mutators that it wraps: The configuration parameters of the operator given to the mutator construction argument are prefixed with "maybe.", the configuration parameters of the operator given to the mutator\_not construction argument are prefixed with "maybe\_not.".

Additional configuration parameters:

• p:: numeric(1)
Probability per individual with which to apply the operator given to the mutator construction argument. Must be set by the user.

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# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of mutator and mutator\_not.

#### **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("maybe", <mutator> [, <mutator_not>])
muts("maybe", <mutator> [, <mutator_not>]) # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("maybe", <mutator> [, <mutator_not>])
```

#### Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Mutator -> MutatorMaybe
```

### **Active bindings**

```
mutator (Mutator)
    Mutator being wrapped. This operator gets run with probability p (configuration parameter).
mutator_not (Mutator)
    Alternative Mutator being wrapped. This operator gets run with probability 1 - p (configuration parameter).
```

# Methods

### **Public methods:**

- MutatorMaybe\$new()
- MutatorMaybe\$prime()
- MutatorMaybe\$clone()

**Method** new(): Initialize the MutatorMaybe object.

```
Usage:
MutatorMaybe$new(mutator, mutator_not = MutatorNull$new())
Arguments:
mutator (Mutator)
```

Mutator to wrap. This operator gets run with probability p (configuration parameter).

The constructed object gets a *clone* of this argument.

The \$mutator field will reflect this value.

```
mutator_not (Mutator)
```

Another Mutator to wrap. This operator runs when mutator is not chosen. By default, this is MutatorNull, i.e. no operation. With this default, the MutatorMaybe object applies the mutator operation with probability p, and no operation at all otherwise.

The constructed object gets a *clone* of this argument. The \$mutator\_not field will reflect this value.

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**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to mutator and mutator\_not during construction.

```
Usage:
MutatorMaybe$prime(param_set)
Arguments:
param_set (ParamSet)
    Passed to MiesOperator$prime().
Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.
Usage:
MutatorMaybe$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

```
Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict_mutators_cmpmaybe, dict_mutators_erase, dict_mutators_gauss, dict_mutators_null, dict_mutators_proxy, dict_mutators_sequential, dict_mutators_unif
```

Other mutator wrappers: OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_proxy, dict\_mutators\_sequential

```
set.seed(1)
mm = mut("maybe", mut("gauss", sdev = 5), p = 0.5)
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5))
data = data.frame(x = rep(0, 5), y = rep(0, 5))
mm$prime(p)
mm$operate(data)
mm$param_set$values$p = 0.3
mm$operate(data)

mm2 = mut("maybe",
    mutator = mut("gauss", sdev = 0.01),
    mutator_not = mut("gauss", sdev = 10),
    p = 0.5
)

mm2$prime(p)
mm2$prime(p)
mm2$operate(data)
```

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```
dict_mutators_null Null Mutator
```

# Description

Null-mutator that does not perform any operation on its input. Useful in particular with operator-wrappers such as MutatorMaybe or MutatorCombination.

# **Configuration Parameters**

This operator has no configuration parameters.

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("null")
muts("null") # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("null")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Mutator -> MutatorNull
```

# Methods

### **Public methods:**

```
• MutatorNull$new()
```

• MutatorNull\$clone()

**Method** new(): Initialize the MutatorNull object.

```
Usage:
```

MutatorNull\$new()

Method clone(): The objects of this class are cloneable with this method.

```
Usage:
```

```
MutatorNull$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

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#### See Also

Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_erase, dict\_mutators\_gauss, dict\_mutators\_maybe, dict\_mutators\_proxy, dict\_mutators\_sequential, dict\_mutators\_unif

# **Examples**

```
mn = mut("null")
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5), z = p_lgl())
data = data.frame(x = rep(0, 5), y = rep(0, 5), z = rep(TRUE, 5))
mn$prime(p)
mn$operate(data)
```

dict\_mutators\_proxy

Proxy-Mutator that Mutates According to its Configuration parameter

# **Description**

Mutator that performs the operation in its operation configuration parameter. This is useful, e.g., to make OptimizerMies's mutation operation fully parametrizable.

# **Configuration Parameters**

• operation :: Mutator

Operation to perform. Must be set by the user. This is primed when \$prime() of MutatorProxy is called, and also when \$operate() is called, to make changing the operation as part of self-adaption possible. However, if the same operation gets used inside multiple MutatorProxy objects, then it is recommended to \$clone(deep = TRUE) the object before assigning them to operation to avoid frequent re-priming.

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("proxy")
muts("proxy") # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("proxy")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Mutator -> MutatorProxy
```

#### Methods

#### **Public methods:**

- MutatorProxy\$new()
- MutatorProxy\$prime()
- MutatorProxy\$clone()

**Method** new(): Initialize the MutatorProxy object.

```
Usage:
```

MutatorProxy\$new()

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the operator given to the operation configuration parameter. Note that this modifies the \$param\_set\$values\$operation object.

```
Usage:
```

```
MutatorProxy$prime(param_set)
```

Arguments:

```
param_set (ParamSet)
```

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage.

MutatorProxy\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

```
\label{lem:continuity} Other \ mutators: \ Mutator, \ Mutator Discrete, \ Mutator Numeric, \ Operator Combination, \ dict_mutators\_cmpmaybe, \ dict_mutators\_erase, \ dict_mutators\_gauss, \ dict_mutators\_maybe, \ dict_mutators\_null, \ dict_mutators\_sequential, \ dict_mutators\_unif
```

Other mutator wrappers: OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_maybe, dict\_mutators\_sequential

```
set.seed(1)
mp = mut("proxy", operation = mut("gauss", sdev = 0.1))
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5))
data = data.frame(x = rep(0, 5), y = rep(0, 5))
mp$prime(p)
```

```
mp$operate(data)
mp$param_set$values$operation = mut("null")
mp$operate(data)
```

dict\_mutators\_sequential

Run Multiple Mutator Operations in Sequence

# **Description**

Mutator that wraps multiple other Mutators given during construction and uses them for mutation in sequence.

# **Configuration Parameters**

This operator has the configuration parameters of the Mutators that it wraps: The configuration parameters of the operator given to the mutators construction argument are prefixed with "mutator\_1", "mutator\_2", ... up to "mutator\_#", where # is length(mutators).

# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of the Mutators given in mutators.

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("sequential", <mutators>)
muts("sequential", <mutators>) # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("sequential", <mutators>)
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Mutator -> MutatorSequential
```

# **Active bindings**

```
mutators (list of Mutator)
```

Mutators being wrapped. These operators get run sequentially in order.

# Methods

#### **Public methods:**

- MutatorSequential\$new()
- MutatorSequential\$prime()
- MutatorSequential\$clone()

**Method** new(): Initialize the MutatorSequential object.

```
Usage:
```

MutatorSequential\$new(mutators)

Arguments:

```
mutators (list of Mutator)
```

Mutators to wrap. The operations are run in order given to mutators. The constructed object gets a *clone* of this argument. The \$mutators field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to mutator and mutator\_not during construction.

```
Usage:
```

MutatorSequential\$prime(param\_set)

Arguments:

param\_set (ParamSet)

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

MutatorSequential\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

# See Also

```
Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict_mutators_cmpmaybe, dict_mutators_erase, dict_mutators_gauss, dict_mutators_maybe, dict_mutators_null, dict_mutators_proxy, dict_mutators_unif
```

Other mutator wrappers: OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_maybe, dict\_mutators\_proxy

```
set.seed(1)
# dataset:
# - x1 is mutated around +- 10
# - x2 influences sdev of mutation of x1
ds = data.frame(x1 = 0, x2 = c(.01, 0.1, 1))
```

```
p = ps(x1 = p_dbl(-10, 10), x2 = p_dbl(0, 10))
\# operator that only mutates x1, with sdev given by x2
gauss_x1 = mut("combine",
  operators = list(
   x1 = mut("gauss", sdev_is_relative = FALSE),
   x2 = mut("null")
  ),
  adaptions = list(x1.sdev = function(x) x$x2)
)
gauss_x1$prime(p)
gauss_x1$operate(ds) # see how x1[1] changes little, x1[3] changes a lot
# operator that mutates x1 with sdev given by x2, as well as x2. However,
# the value that x2 takes after mutation does not influence the value that
# the mutator of x1 "sees" -- although x2 is mutated to extreme values,
\# mutation of x1 happens as in `gauss_x1`.
gauss_x1_x2 = mut("combine",
  operators = list(
   x1 = mut("gauss", sdev_is_relative = FALSE),
   x2 = mut("gauss", sdev = 100)
  ),
  adaptions = list(x1.sdev = function(x) x$x2)
)
gauss_x1_x2$prime(p)
gauss_x1_x2$operate(ds) # see how x1 changes in similar ways to above
# operator that mutates sequentially: first x2, and then x1 with sdev given
# by x2. The value that x2 takes after mutation *does* influence the value
# that the mutator of x1 "sees": x1 is mutated either to a large degree,
# or not at all.
gauss_x2_then_x1 = mut("sequential", list(
   mut("combine",
      operators = list(
       x1 = mut("null"),
        x2 = mut("gauss", sdev = 100)
      )
   ),
    mut("combine",
      operators = list(
       x1 = mut("gauss", sdev_is_relative = FALSE),
       x2 = mut("null")
     ),
      adaptions = list(x1.sdev = function(x) x$x2)
))
gauss_x2_then_x1$prime(p)
gauss_x2_then_x1$operate(ds)
```

34 dict\_mutators\_unif

dict\_mutators\_unif Uniform Discrete Mutator

# **Description**

Discrete components are mutated by sampling from a uniform distribution, either from all possible values of each component, or from all values except the original value.

Since the information loss is very high, this should in most cases be combined with MutatorCmpMaybe.

# **Configuration Parameters**

can\_mutate\_to\_same :: logical(1)
 Whether to sample from entire range of each parameter (TRUE) or from all values except the current value (FALSE). Initialized to TRUE.

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_fct ('ParamFct')
```

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("unif")
muts("unif") # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("unif")
```

# Super classes

```
miesmuschel::MutatorDiscrete
-> MutatorDiscreteUniform
```

# Methods

#### **Public methods:**

- MutatorDiscreteUniform\$new()
- MutatorDiscreteUniform\$clone()

**Method** new(): Initialize the MutatorDiscreteUniform object.

Usage:
MutatorDiscreteUniform\$new()

**Method** clone(): The objects of this class are cloneable with this method.

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```
Usage:
MutatorDiscreteUniform$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

# See Also

Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_erase, dict\_mutators\_gauss, dict\_mutators\_maybe, dict\_mutators\_null, dict\_mutators\_proxy, dict\_mutators\_sequential

# **Examples**

```
set.seed(1)
mdu = mut("unif")
p = ps(x = p_lgl(), y = p_fct(c("a", "b", "c")))
data = data.frame(x = rep(TRUE, 5), y = rep("a", 5),
    stringsAsFactors = FALSE) # necessary for R <= 3.6
mdu$prime(p)
mdu$operate(data)
mdu$param_set$values$can_mutate_to_same = FALSE
mdu$operate(data)</pre>
```

dict\_recombinators

Dictionary of Recombinators

# Description

Dictionary of Recombinators

# Usage

dict\_recombinators

### **Format**

An object of class DictionaryRecombinator (inherits from DictionaryEx, Dictionary, R6) of length 15.

# Methods

Methods inherited from Dictionary, as well as:

help(key, help\_type)
 (character(1), character(1))
 Displays help for the dictionary entry key. help\_type is one of "text", "html", "pdf" and given as the help\_type argument of R's help().

#### See Also

Other dictionaries: dict\_filtors, dict\_mutators, dict\_scalors, dict\_selectors, mut()

dict\_recombinators\_cmpmaybe

Recombinator Choosing Action Component-Wise Independently

# **Description**

Recombinator that chooses which operation to perform probabilistically and independently for each component. The Recombinator wraps two other Recombinators given during construction, and both of these operators are run. The ultimate result is sampled from the results of these operations independently for each individuum and component: with probability p (configuration parameter), the result from the Recombinator given to the recombinator construction argument is used, and with probability p - 1 the one given to recombinator\_not is used.

The values of \$n\_indivs\_in and \$n\_indivs\_out is set to the corresponding values of the wrapped Recombinators. Both recombinator and recombinator\_not must currently have the same respective \$n\_indivs\_in and \$n\_indivs\_out values.

# **Configuration Parameters**

This operator has the configuration parameters of the Recombinators that it wraps: The configuration parameters of the operator given to the recombinator construction argument are prefixed with "cmpmaybe.", the configuration parameters of the operator given to the recombinator\_not construction argument are prefixed with "cmpmaybe\_not.".

Additional configuration parameters:

• p:: numeric(1)
Probability per component with which to use the result of applying the operator given to the recombinator construction argument. Must be set by the user.

# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of recombinator and recombinator\_not.

# **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("cmpmaybe", <recombinator> [, <recombinator_not>])
recs("cmpmaybe", <recombinator> [, <recombinator_not>]) # takes vector IDs, returns list of Recombinat
# long form:
dict_recombinators$get("cmpmaybe", <recombinator> [, <recombinator_not>])
```

#### Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Recombinator -> RecombinatorCmpMaybe
```

#### **Active bindings**

```
recombinator (Recombinator)
```

Recombinator being wrapped. This operator gets run with probability p (configuration parameter).

```
recombinator_not (Recombinator)
```

Alternative Recombinator being wrapped. This operator gets run with probability 1 - p (configuration parameter).

#### Methods

### **Public methods:**

- RecombinatorCmpMaybe\$new()
- RecombinatorCmpMaybe\$prime()
- RecombinatorCmpMaybe\$clone()

**Method** new(): Initialize the RecombinatorCmpMaybe object.

Usage:

RecombinatorCmpMaybe\$new(recombinator, recombinator\_not = NULL)

Arguments:

```
recombinator (Recombinator)
```

Recombinator to wrap. Component-wise results of this operator are used with probability p (Configuration parameter).

The constructed object gets a *clone* of this argument. The \$recombinator field will reflect this value.

```
recombinator_not (Recombinator)
```

Another Recombinator to wrap. Results from this operator are used when recombinator is not chosen. By default, this is RecombinatorNull, i.e. no operation, with both n\_indivs\_in and n\_indivs\_out set to match recombinator. This does not work when recombinator has n\_indivs\_in < n\_indivs\_out, in which case this argument must be set explicitly.

With this default, the RecombinatorCmpMaybe object applies the recombinator operation with probability p, and no operation at all otherwise.

The constructed object gets a *clone* of this argument. The \$recombinator\_not field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to recombinator and recombinator\_not during construction.

```
Usage:
```

```
RecombinatorCmpMaybe$prime(param_set)
```

Arguments:

```
param_set (ParamSet)
```

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

RecombinatorCmpMaybe\$clone(deep = FALSE)

Arguments.

deep Whether to make a deep clone.

#### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_null, dict\_recombinators\_prox\_dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xond dict\_recombinators\_xounif

Other recombinator wrappers: OperatorCombination, dict\_recombinators\_maybe, dict\_recombinators\_proxy, dict\_recombinators\_sequential

## **Examples**

```
set.seed(1)
rm = rec("cmpmaybe", rec("swap"), p = 0.5)
p = ps(x = p_int(1, 8), y = p_dbl(1, 8), z = p_lgl())
data = data.frame(x = 1:8, y = 1:8, z = rep(TRUE, 4))
rm$prime(p)
rm$operate(data)
rm$param_set$values$p = 0.3
rm$operate(data)
# equivalent to rec("cmpmaybe", rec("swap", keep_complement = FALSE), p = 0.7)
rm2 = rec("cmpmaybe",
  recombinator = rec("null", 2, 1),
  recombinator_not = rec("swap", keep_complement = FALSE),
  p = 0.3
)
rm2$prime(p)
rm2$operate(data)
```

dict\_recombinators\_convex

Convex Combination Recombinator

## **Description**

Numeric Values between various individuals are recombined via component-wise convex combination (or weighted mean). The number of individuals over which the convex combination is taken must be determined during construction as n\_indivs\_in.

The number of output individuals is always 1, i.e. n\_indivs\_in are used to create one output value. When using this recombinator in a typical EA setting, e.g. with mies\_generate\_offspring, it is therefore recommended to use a parent-selector where the expected quality of selected parents does not depend on the number of parents selected when n\_indivs\_in is large: sel("tournament") is preferred to sel("best").

## **Configuration Parameters**

• lambda:: numeric | matrix Combination weights; these are normalized to sum to 1 internally. Must either be a vector of length n\_indivs\_in, or a matrix with n\_indivs\_in rows and as many columns as there are components in the values being operated on. Must be non-negative, at least one value per column must be greater than zero, but it is not necessary that they sum to 1.

Initialized to rep(1, n\_indivs\_in), i.e. equal weights to all individuals being operated on.

## **Supported Operand Types**

```
Supported Domain classes are: p_db1 ('ParamDbl')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("convex")
recs("convex") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("convex")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Recombinator -> RecombinatorConvex
```

#### Methods

#### **Public methods:**

- RecombinatorConvex\$new()
- RecombinatorConvex\$clone()

**Method** new(): Initialize the RecombinatorConvex object.

```
Usage:
```

```
RecombinatorConvex$new(n_indivs_in = 2)
```

```
Arguments:
n_indivs_in (integer(1))
```

Number of individuals to consider at the same time. When operating, the number of input individuals must be divisible by this number. Default 2.

The \$n\_indivs\_in field will reflect this value.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

RecombinatorConvex\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_null, dict\_recombinators\_proxidict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xonadict\_recombinators\_xounif

## **Examples**

```
set.seed(1)
rcvx = rec("convex", n_indivs_in = 3)
p = ps(x = p_dbl(-5, 5), y = p_dbl(-5, 5), z = p_dbl(-5, 5))
data = data.frame(x = 0:5, y = 0:5, z = 0:5)
rcvx$prime(p)
rcvx$operate(data)  # mean of groups of 3

rcvx = rec("convex", 3, lambda = c(0, 1, 2))$prime(p)
rcvx$operate(data)  # for groups of 3, take 1/3 of 2nd and 2/3 of 3rd row
lambda = matrix(c(0, 1, 2, 1, 1, 1, 0, 0), ncol = 3)
lambda
rcvx = rec("convex", 3, lambda = lambda)$prime(p)
rcvx$operate(data)  # componentwise different operation
```

dict\_recombinators\_cvxpair

Convex Combination Recombinator for Pairs

## Description

Numeric Values between various individuals are recombined via component-wise convex combination (or weighted mean). Exactly two individuals are being recombined, and the lambda configuration parameter determines the relative weight of the first individual in each pair for the first result, and the relative weight of the second individual for the complement, if initialized with keep\_complement set to TRUE.

## **Configuration Parameters**

• lambda:: numeric
Combination weight. If keep\_complement is TRUE, then two individuals are returned for each
pair of input individuals: one corresponding to lambda \* <1st individual> + (1-lambda) \* <2nd individual>,
and one corresponding to (1-lambda) \* <1st individual> + lambda \* <2nd individual>
(i.e. the complement). Otherwise, only the first of these two is generated. Must either be a
scalar, or a vector with length equal to the number of components in the values being operated
on. Must be between 0 and 1.
Initialized to 0.5.

## **Supported Operand Types**

```
Supported Domain classes are: p_dbl ('ParamDbl')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("convex")
recs("convex") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("convex")
```

## Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Recombinator-> miesmuschel::RecombinatorPair
-> RecombinatorConvexPair
```

#### Methods

### **Public methods:**

- RecombinatorConvexPair\$new()
- RecombinatorConvexPair\$clone()

Method new(): Initialize the RecombinatorConvexPair object.

```
Usage:
```

RecombinatorConvexPair\$new(keep\_complement = TRUE)

Arguments:

```
keep_complement (logical(1))
```

Whether the operation should keep both resulting individuals (TRUE), or only the first and discard the complement (FALSE). Default TRUE. The \$keep\_complement field will reflect this value.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
RecombinatorConvexPair$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

## See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_maybe, dict\_recombinators\_null, dict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xonadict\_recombinators\_xounif

## **Examples**

```
set.seed(1)
rcvx = rec("cvxpair")
p = ps(x = p_dbl(-5, 5), y = p_dbl(-5, 5), z = p_dbl(-5, 5))
data = data.frame(x = 0:5, y = 0:5, z = 0:5)
rcvx$prime(p)
rcvx$operate(data) # mean of groups of 2
# with the default value of lambda = 0.5, the default of
# keep_complement = TRUE means that pairs of equal values are generated;
# consider setting keep_complement = FALSE int that case.
rcvx$param_set$values$lambda = 0.1
rcvx$operate(data)
```

dict\_recombinators\_maybe

Recombinator Choosing Action Probabilistically

## **Description**

Recombinator that chooses which operation to perform probabilistically. The Recombinator wraps two other Recombinators given during construction, and for each group of  $n_i$  individuals, the operation to perform is sampled: with probability p (configuration parameter), the Recombinator given to the recombinator construction argument is applied, and with probability p - 1 the one given to recombinator\_not is applied.

The values of \$n\_indivs\_in and \$n\_indivs\_out is set to the corresponding values of the wrapped Recombinators. Both recombinator and recombinator\_not must currently have the same respective \$n\_indivs\_in and \$n\_indivs\_out values.

#### **Configuration Parameters**

This operator has the configuration parameters of the Recombinators that it wraps: The configuration parameters of the operator given to the recombinator construction argument are prefixed with "maybe.", the configuration parameters of the operator given to the recombinator\_not construction argument are prefixed with "maybe\_not.".

Additional configuration parameters:

 p::numeric(1)
 Probability per group of n\_indivs\_in individuals with which to apply the operator given to the recombinator construction argument. Must be set by the user.

## **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of recombinator and recombinator\_not.

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("maybe", <recombinator> [, <recombinator_not>])
recs("maybe", <recombinator> [, <recombinator_not>]) # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("maybe", <recombinator> [, <recombinator_not>])
```

#### Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Recombinator -> RecombinatorMaybe
```

## **Active bindings**

```
recombinator (Recombinator)
    Recombinator being wrapped. This operator gets run with probability p (configuration parameter).
recombinator_not (Recombinator)
    Alternative Recombinator being wrapped. This operator gets run with probability 1 - p (configuration parameter).
```

#### Methods

#### **Public methods:**

- RecombinatorMaybe\$new()
- RecombinatorMaybe\$prime()
- RecombinatorMaybe\$clone()

**Method** new(): Initialize the RecombinatorMaybe object.

```
Usage:
```

RecombinatorMaybe\$new(recombinator, recombinator\_not = NULL)

Arguments:

```
recombinator (Recombinator)
```

Recombinator to wrap. This operator gets run with probability p (Configuration parameter).

The constructed object gets a *clone* of this argument. The \$recombinator field will reflect this value.

```
recombinator_not (Recombinator)
```

Another Recombinator to wrap. This operator runs when recombinator is not chosen. By default, this is RecombinatorNull, i.e. no operation, with both n\_indivs\_in and n\_indivs\_out set to match recombinator. This does not work when recombinator has n\_indivs\_in < n\_indivs\_out, in which case this argument must be set explicitly.

With this default, the RecombinatorMaybe object applies the recombinator operation with probability p, and no operation at all otherwise.

The constructed object gets a *clone* of this argument. The \$recombinator\_not field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to recombinator and recombinator\_not during construction.

```
- Usuge.
```

RecombinatorMaybe\$prime(param\_set)

Arguments:

param\_set (ParamSet)

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

RecombinatorMaybe\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_null, dict\_recombinators\_product\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xonadict\_recombinators\_xounif

Other recombinator wrappers: OperatorCombination, dict\_recombinators\_cmpmaybe, dict\_recombinators\_proxy, dict\_recombinators\_sequential

## **Examples**

```
set.seed(1)
rm = rec("maybe", rec("xounif", p = 1), p = 0.5)
p = ps(x = p_int(1, 8), y = p_dbl(1, 8), z = p_lgl())
data = data.frame(x = 1:8, y = 1:8, z = rep(TRUE, 4))

rm$prime(p)
rm$operate(data)

rm$param_set$values$p = 0.3
rm$operate(data)

rm2 = rec("maybe",
    recombinator = rec("xounif", p = 1),
    recombinator_not = rec("xounif", p = 0.5),
    p = 0.5
)

rm2$prime(p)
rm2$operate(data)
```

dict\_recombinators\_null

Null-Recombinator

## **Description**

Null-recombinator that does not perform any operation on its input. Useful in particular with operator-wrappers such as RecombinatorMaybe or RecombinatorCombination.

n\_indivs\_in and n\_indivs\_out can be set during construction, where n\_indivs\_out must be less or equal n\_indivs\_in. If it is strictly less, then the operation returns only the first n\_indivs\_out individuals out of each n\_indivs\_in sized group.

## **Configuration Parameters**

This operator has no configuration parameters.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("null")
recs("null") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("null")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Recombinator -> RecombinatorNull
```

## Methods

#### **Public methods:**

- RecombinatorNull\$new()
- RecombinatorNull\$clone()

**Method** new(): Initialize base class components of the Recombinator.

```
Usage:
```

```
RecombinatorNull$new(n_indivs_in = 1, n_indivs_out = n_indivs_in)
```

Arguments:

```
n_indivs_in (integer(1))
```

Number of individuals to consider at the same time. When operating, the number of input individuals must be divisible by this number. Setting this number to a number unequal 1 is mostly useful when incorporating this operator in wrappers such as RecombinatorMaybe or RecombinatorCombination. Default 1.

The \$n\_indivs\_in field will reflect this value.

```
n_indivs_out (integer(1))
```

Number of individuals that result for each n\_indivs\_in lines of input. Must be at most n\_indivs\_in. If this is less than n\_indivs\_in, then only the first n\_indivs\_out individuals out of each n\_indivs\_in sized group are returned by an operation. Setting this number to a number unequal 1 is mostly useful when incorporating this operator in wrappers such as RecombinatorMaybe or RecombinatorCombination. Default equal to n\_indivs\_in. The \$n\_indivs\_out field will reflect this value.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

RecombinatorNull\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_product\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xonoict\_recombinators\_xounif

## **Examples**

```
rn = rec("null")
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5), z = p_lgl())
data = data.frame(x = 1:4, y = 0:3, z = rep(TRUE, 4))
rn$prime(p)
rn$operate(data)
rn_half = rec("null", n_indivs_in = 2, n_indivs_out = 1)
rn_half$prime(p)
rn_half$operate(data)
```

dict\_recombinators\_proxy

Proxy-Recombinator that Recombines According to its Configuration parameter

## Description

Recombinator that performs the operation in its operation configuration parameter. This is useful, e.g., to make OptimizerMies's recombination operation fully parametrizable.

## **Configuration Parameters**

• operation :: Recombinator

Operation to perform. Must be set by the user. This is primed when \$prime() of RecombinatorProxy is called, and also when \$perate() is called, to make changing the operation as part of self-adaption possible. However, if the same operation gets used inside multiple RecombinatorProxy objects, then it is recommended to \$clone(deep = TRUE) the object before assigning them to operation to avoid frequent re-priming.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("proxy")
recs("proxy") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("proxy")
```

## Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Recombinator-> RecombinatorProxy
```

#### Methods

#### **Public methods:**

- RecombinatorProxy\$new()
- RecombinatorProxy\$prime()
- RecombinatorProxy\$clone()

**Method** new(): Initialize the RecombinatorProxy object.

```
Usage:
```

```
RecombinatorProxy$new(n_indivs_in = 2, n_indivs_out = n_indivs_in)
```

Arguments:

```
n_indivs_in (integer(1))
```

Number of individuals to consider at the same time. When operating, the number of input individuals must be divisible by this number. Furthermore, the Recombinator assigned to the operation configuration parameter must have an n\_indivs\_in that is a divisor of this number. Default 2.

The \$n\_indivs\_in field will reflect this value.

```
n_indivs_out (integer(1))
```

Number of individuals that result for each n\_indivs\_in lines of input. Must be at most n\_indivs\_in. The ratio of \$n\_indivs\_in to \$n\_indivs\_out of the Recombinator assigned to the operation configuration parameter must be the same as n\_indivs\_in to n\_indivs\_out of this object. Default equal to n\_indivs\_in.

The \$n\_indivs\_out field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the operator given to the operation configuration parameter. Note that this modifies the \$param\_set\$values\$operation object.

```
Usage:
```

```
RecombinatorProxy$prime(param_set)
```

Arguments:

```
param_set (ParamSet)
```

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage.

```
RecombinatorProxy$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xond dict\_recombinators\_xounif

Other recombinator wrappers: OperatorCombination, dict\_recombinators\_cmpmaybe, dict\_recombinators\_maybe, dict\_recombinators\_sequential

## **Examples**

```
set.seed(1)
rp = rec("proxy", operation = rec("xounif"))
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5), z = p_lgl())
data = data.frame(x = 1:4, y = 0:3, z = rep(TRUE, 4))

rp$prime(p)
rp$operate(data)  # default operation: null

rp$param_set$values$operation = rec("xounif", p = 0.5)
rp$operate(data)
```

dict\_recombinators\_sbx

Simulated Binary Crossover Recombinator

## **Description**

Numeric Values between two individuals are recombined via component-wise independent simulated binary crossover. See Deb (1995) for more details.

This operator is applied to all components; It is common to apply the operator to only some randomly chosen components, in which case the rec("cmpmaybe") operator should be used; see examples.

## **Configuration Parameters**

• n:: numeric

Non-negative distribution index of the polynomial distribution for each component. Generally spoken, the higher n, the higher the probability of creating near parent values. This may either be a scalar in which case it is applied to all input components, or a vector, in which case it must have the length of the input components and applies to components in order in which they appear in the priming ParamSet. Initialized to 1.

## **Supported Operand Types**

```
Supported Domain classes are: p_int ('ParamInt'), p_dbl ('ParamDbl')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("sbx")
recs("sbx") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("sbx")
```

### Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Recombinator-> miesmuschel::RecombinatorPair
-> RecombinatorSimulatedBinaryCrossover
```

#### Methods

#### **Public methods:**

- RecombinatorSimulatedBinaryCrossover\$new()
- RecombinatorSimulatedBinaryCrossover\$clone()

Method new(): Initialize the RecombinatorSimulatedBinaryCrossover object.

Usage:

RecombinatorSimulatedBinaryCrossover\$new(keep\_complement = TRUE)

Arguments:

```
keep_complement (logical(1))
```

Whether the operation should keep both resulting individuals (TRUE), or only the first and discard the complement (FALSE). Default TRUE. The \$keep\_complement field will reflect this value.

Method clone(): The objects of this class are cloneable with this method.

Usage:

RecombinatorSimulatedBinaryCrossover\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

## References

Deb, Kalyanmoy, Agrawal, Bhushan R, others (1995). "Simulated binary crossover for continuous search space." *Complex systems*, **9**(2), 115–148.

## See Also

```
Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict_recombinators_cmpmaybe, dict_recombinators_convex, dict_recombinators_cvxpair, dict_recombinators_maybe, dict_recombinators_nudict_recombinators_proxy, dict_recombinators_sequential, dict_recombinators_swap, dict_recombinators_xounif
```

### **Examples**

```
set.seed(1)
rsbx = rec("cmpmaybe", rec("sbx"), p = 0.5)
p = ps(x = p_dbl(-5, 5), y = p_dbl(-5, 5), z = p_dbl(-5, 5))
data = data.frame(x = 0:5, y = 0:5, z = 0:5)
rsbx$prime(p)
rsbx$operate(data)

rsbx = rec("sbx", n = c(0.5, 1, 10))
rsbx$prime(p)
rsbx$prime(p)
rsbx$operate(data)
```

dict\_recombinators\_sequential

Run Multiple Recombinator Operations in Sequence

## **Description**

Recombinator that wraps multiple other Recombinators given during construction and uses them for mutation in sequence.

When subsequent Recombinators have mismatching n\_indivs\_out/n\_indivs\_in, then RecombinatorSequential tries to match them by running them multiple times. If e.g. recombinators[[1]]\$n\_indivs\_out is 2 and recombinators[[2]]\$n\_indivs\_in is 1, then recombinators[[2]] is run twice, once for each output of recombinators[[1]].

When the allow\_lcm\_packing argument is FALSE, then an error is given if neither n\_indivs\_out of a Recombinator divides n\_indivs\_in of the following Recombinator, nor n\_indivs\_in of the latter divides n\_indivs\_out of the former even when considering that the former is run multiple times. If allow\_lcm\_packing is TRUE, then both recombinators are run multiple times, according to the lowest common multiple ("lcm") of the two.

However, allow\_lcm\_packing can lead to very large values of n\_indivs\_in/n\_indivs\_out, so it may instead be preferred to add RecombinatorNull objects with fitting n\_indivs\_in/n\_indivs\_out values to match subsequent recombinators.

### **Configuration Parameters**

This operator has the configuration parameters of the Recombinators that it wraps: The configuration parameters of the operator given to the recombinators construction argument are prefixed with "recombinator\_1", "recombinator\_2", ... up to "recombinator\_#", where # is length(recombinators).

Additional configuration parameters:

• shuffle\_between :: logical(1)
Whether to reorder values between invocations of recombinators. Initialized to TRUE.

### **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of the Recombinators given in recombinators.

### **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("sequential", <recombinators>)
recs("sequential", <recombinators>) # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("sequential", <recombinators>)
```

## Super classes

```
miesmuschel::MiesOperator->miesmuschel::Recombinator->RecombinatorSequential
```

## **Active bindings**

```
recombinators (list of Recombinator)
Recombinators being wrapped. These operators get run sequentially in order.

allow_lcm_packing (logical(1))
Whether to allow lowest common multiple packing.
```

#### Methods

#### **Public methods:**

- RecombinatorSequential\$new()
- RecombinatorSequential\$prime()
- RecombinatorSequential\$clone()

Method new(): Initialize the RecombinatorSequential object.

```
Usage:
```

```
RecombinatorSequential$new(recombinators, allow_lcm_packing = FALSE)
```

Arguments:

```
recombinators (list of Recombinator)
```

Recombinators to wrap. The operations are run in order given to recombinators. The constructed object gets a *clone* of this argument. The \$recombinators field will reflect this value.

```
allow_lcm_packing (logical(1))
```

Whether to allow lowest common multiple packing. Default FALSE. The \$allow\_lcm\_packing field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to recombinator and recombinator\_not during construction.

```
Usage:
RecombinatorSequential$prime(param_set)
Arguments:
param_set (ParamSet)
    Passed to MiesOperator$prime().
Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.
Usage:
RecombinatorSequential$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

## See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_swap, dict\_recombinators\_xonary, dict\_recombinators\_xounif

Other recombinator wrappers: OperatorCombination, dict\_recombinators\_cmpmaybe, dict\_recombinators\_maybe, dict\_recombinators\_proxy

## **Examples**

```
set.seed(1)

ds = data.frame(a = c(0, 1), b = c(0, 1))
p = ps(a = p_dbl(0, 1), b = p_dbl(0, 1))

convex = rec("cvxpair", lambda = 0.7)
swap = rec("swap")

convex_then_swap = rec("sequential", list(convex, swap))

ds

convex$prime(p)$operate(ds)

swap$prime(p)$operate(ds)

convex_then_swap$prime(p)$operate(ds)
```

## **Description**

Values between two individuals are exchanged. This is relatively useless as an operator by itself, but is used in combination with RecombinatorCmpMaybe to get a recombinator that is crossing over individuals uniformly at random. Because this is such a frequently-used operation, the RecombinatorCrossoverUniform pseudo-class exists as a shortcut.

## **Configuration Parameters**

None.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("swap")
recs("swap") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("swap")
```

## Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Recombinator-> miesmuschel::RecombinatorPair
-> RecombinatorSwap
```

## Methods

### **Public methods:**

- RecombinatorSwap\$new()
- RecombinatorSwap\$clone()

**Method** new(): Initialize the RecombinatorCrossoverSwap object.

```
Usage:
RecombinatorSwap$new(keep_complement = TRUE)
```

```
Arguments:
```

```
keep_complement (logical(1))
```

Whether the operation should keep both resulting individuals (TRUE), or only the first and discard the complement (FALSE). Default TRUE. The \$keep\_complement field will reflect this value.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

```
RecombinatorSwap$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

#### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_xoudict\_recombinators\_xounif

## **Examples**

```
set.seed(1)
rs = rec("swap")
p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5), z = p_dbl(-5, 5))
data = data.frame(x = 0:5, y = 0:5, z = 0:5)

rs$prime(p)
rs$operate(data)

rx = rec("cmpmaybe", rec("swap"), p = 0.5)  # the same as 'rec("xounif")'
rx$prime(p)
rx$operate(data)
```

dict\_recombinators\_xonary

N-ary Crossover Recombinator

## Description

Values are chosen componentwise independently at random from multiple individuals. The number of individuals must be determined during construction as n\_indivs\_in.

The number of output individuals is always 1, i.e. n\_indivs\_in are used to create one output value. When using this recombinator in a typical EA setting, e.g. with mies\_generate\_offspring, it is therefore recommended to use a parent-selector where the expected quality of selected parents does not depend on the number of parents selected when n\_indivs\_in is large: sel("tournament") is preferred to sel("best").

## **Configuration Parameters**

• p:: numeric | matrix

Sampling weights these are normalized to sum to 1 internally. Must either be a vector of length n\_indivs\_in, or a matrix with n\_indivs\_in rows and as many columns as there are components in the values being operated on. Must be non-negative, at least one value per column must be greater than zero, but it is not necessary that they sum to 1.

Initialized to rep(1, n\_indivs\_in), i.e. uniform sampling from all individuals being operated on.

## **Supported Operand Types**

```
Supported Domain classes are: p_dbl ('ParamDbl')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("convex")
recs("convex") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("convex")
```

#### Super classes

```
miesmuschel::Recombinator
```

## Methods

#### Public methods:

- RecombinatorCrossoverNary\$new()
- RecombinatorCrossoverNary\$clone()

Method new(): Initialize the RecombinatorConvex object.

```
Usage:
```

```
RecombinatorCrossoverNary$new(n_indivs_in = 2)
```

Arguments:

```
n_indivs_in (integer(1))
```

Number of individuals to consider at the same time. When operating, the number of input individuals must be divisible by this number. Default 2.

The \$n\_indivs\_in field will reflect this value.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

```
RecombinatorCrossoverNary$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

#### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swadict\_recombinators\_xounif

## **Examples**

```
set.seed(1)
rxon = rec("xonary", n_indivs_in = 3)
p = ps(x = p_dbl(-5, 5), y = p_dbl(-5, 5), z = p_dbl(-5, 5))
data = data.frame(x = 0:5, y = 0:5, z = 0:5)

rxon$prime(p)
rxon$operate(data)  # uniform sampling from groups of 3

rxon = rec("xonary", 3, p = c(0, 1, 2))$prime(p)
# for groups of 3, take with probability 1/3 from 2nd and with probability 2/3 from 3rd row rxon$operate(data)

pmat = matrix(c(0, 1, 2, 1, 1, 1, 1, 0, 0), ncol = 3)
pmat

rxon = rec("xonary", 3, p = pmat)$prime(p)
rxon$operate(data)  # componentwise different operation
```

dict\_recombinators\_xounif

Crossover Recombinator

## **Description**

Values between two individuals are exchanged with component-wise independent probability.

This is a pseudo-class: It does not create a single R6-object of a class; instead, it creates the object rec("cmpmaybe", rec("swap"), p = 0.5), making use of the RecombinatorCmpMaybe and RecombinatorSwap operators.

### Usage

RecombinatorCrossoverUniform(keep\_complement = TRUE)

### **Arguments**

```
keep_complement
```

(logical(1))

Whether the operation should keep both resulting individuals (TRUE), or only the first and discard the complement (FALSE). Default TRUE. The \$keep\_complement field will reflect this value.

## Value

```
an object of class Recombinator: rec("cmpmaybe", rec("swap")).
```

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the the dictionary dict\_recombinators in the following way:

```
# preferred:
rec("xounif")
recs("xounif") # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("xounif")
```

### See Also

Other recombinators: OperatorCombination, Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swadict\_recombinators\_xonary

## **Examples**

```
set.seed(1)
rx = rec("xounif")
print(rx)

p = ps(x = p_int(-5, 5), y = p_dbl(-5, 5), z = p_dbl(-5, 5))
data = data.frame(x = 0:5, y = 0:5, z = 0:5)

rx$prime(p)
rx$operate(data)

rx$param_set$values$p = 0.3
rx$operate(data)
```

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dict\_scalors

Dictionary of Scalors

## **Description**

Dictionary of Scalors

## Usage

dict\_scalors

#### **Format**

An object of class DictionaryScalor (inherits from DictionaryEx, Dictionary, R6) of length 15.

#### Methods

Methods inherited from Dictionary, as well as:

```
    help(key, help_type)
        (character(1), character(1))
        Displays help for the dictionary entry key. help_type is one of "text", "html", "pdf" and given as the help_type argument of R's help().
```

## See Also

Other dictionaries: dict\_filtors, dict\_mutators, dict\_recombinators, dict\_selectors, mut()

```
dict_scalors_aggregate
```

Scalor giving Weighted Sum of Multiple Scalors

## **Description**

Scalor that applies multiple other Scalors and calculates their weighted sum.

## **Configuration Parameters**

This operation has the configuration parameters of the Scalors that it wraps: The configuration parameters of the operator given to the scalors construction argument are prefixed with "scalor\_1", "scalor\_2", ... up to "scalor\_#", where # is length(scalors).

Additional configuration parameters:

- weight\_1, weight\_2, ...: numeric(1) Weight factors of scalors[[1]], scalors[[2]], etc. Depending on scaling, the outputs of scalors is multiplied with this (when scaling is "linear" or "rank"), or ties between ranks are broken with it (when scaling is "tiebreak"). Initialized to 1.
- scaling:: character(1)
  How to calculate output values, one of "linear", "rank" or "tiebreak". When scaling is "linear", then the output is calculated as the weighted sum of the outputs of scalors, weighted by weight\_1, weight\_2 etc. When scaling is "rank", then the output is calculated as the weighted sum of the rank() of scalors, weighted by weight\_1, weight\_2 etc., with ties broken by average. When scaling is "tiebreak", then the output is calculated as the averaged rank() of the scalors with the highest weight\_#, with ties broken by the average rank() of the second highest weight\_#, with remaining ties broken by scalors with third highest weight\_# etc. Initialized to "linear".
- scale\_output :: logical(1)
  Whether to scale the output to lie between 0 and 1. Initialized to FALSE.

### **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("aggregate", <scalors>)
scls("aggregate", <scalors>) # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("aggregate", <scalors>)
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorAggregate
```

## **Active bindings**

```
scalors (list of Scalor)
Scalors being wrapped. These operators are run and their outputs weighted.
```

#### Methods

#### **Public methods:**

- ScalorAggregate\$new()
- ScalorAggregate\$prime()
- ScalorAggregate\$clone()

**Method** new(): Initialize the ScalorAggregate object.

```
Usage:
ScalorAggregate$new(scalors)
Arguments:
```

```
scalors (list of Scalor)
```

Scalors to wrap. The operations are run and weighted by weight\_# configuration parameters, depending on the scaling configuration parameter. The constructed object gets a *clone* of this argument. The \$scalors field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to scalors during construction.

```
Usage:
    ScalorAggregate$prime(param_set)
    Arguments:
    param_set (ParamSet)
        Passed to MiesOperator$prime().
    Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.
    Usage:
    ScalorAggregate$clone(deep = FALSE)
    Arguments:
    deep Whether to make a deep clone.
```

#### See Also

Other scalors: Scalor, dict\_scalors\_domcount, dict\_scalors\_fixedprojection, dict\_scalors\_hypervolume, dict\_scalors\_nondom, dict\_scalors\_one, dict\_scalors\_proxy, dict\_scalors\_single

Other scalor wrappers: dict\_scalors\_fixedprojection, dict\_scalors\_proxy

## **Examples**

```
# only 2 * second objective
sa$param_set$values[c("weight_1", "weight_2")] = c(0, 2)
sa$operate(data, fitnesses)
```

dict\_scalors\_domcount Scalor Counting Dominating Individuals

## **Description**

Scalor that returns a the number of (weakly, epsilon-) dominated or dominating individuals for each individuum.

### **Configuration Parameters**

- output :: character(1)
  What to count: individuals that are being dominated by the point under consideration("count\_dominated"), or individuals that do not dominate the point under consideration ("count\_not\_dominating").
  In both cases, a larger output means the individual is "better", in some way, according to the fitness values. Initialized with "count\_not\_dominating".
- epsilon:: numeric Epsilon-value for non-dominance, as used by rank\_nondominated. Initialized to 0.
- jitter :: logical(1)
  Whether to add random jitter to points, with magnitude sqrt(.Machine\$double.eps) relative to fitness values. This is used to effectively break ties.
- scale\_output :: logical(1)
  Whether to scale output by the total number of individuals, giving output between 0 and 1 (inclusive) when TRUE or integer outputs ranging from 0 and nrow(fitnesses) (inclusive) when FALSE. Initialized to TRUE.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("domcount")
scls("domcount") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("domcount")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorDomcount
```

#### Methods

#### **Public methods:**

- ScalorDomcount\$new()
- ScalorDomcount\$clone()

**Method** new(): Initialize the ScalorNondom object.

```
Usage:
```

ScalorDomcount\$new()

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage.
```

ScalorDomcount\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

Other scalors: Scalor, dict\_scalors\_aggregate, dict\_scalors\_fixedprojection, dict\_scalors\_hypervolume, dict\_scalors\_nondom, dict\_scalors\_one, dict\_scalors\_proxy, dict\_scalors\_single

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## **Examples**

```
p = ps(x = p_dbl(-5, 5))
data = data.frame(x = rep(0, 5))

sd = scl("domcount")
sd$prime(p)

(fitnesses = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2))

# to see the fitness matrix, use:
## plot(fitnesses, pch = as.character(1:5))

# note that for both 2 and 4, all points do not dominate them
# their value is therefore 1
sd$operate(data, fitnesses)

sd$param_set$values$scale_output = FALSE
sd$operate(data, fitnesses)

sd$param_set$values$output = "count_dominated"
# point 4 dominates three other points, point 2 only one other point.
sd$operate(data, fitnesses)
```

```
dict_scalors_fixedprojection
```

Multi-Objective Fixed Projection Scalor

## Description

Scalor that returns the maximum of a set of projections.

Priming PS must contain a "scalarization\_weights" tagged p\_uty that contains weight matrices (Nobjectives x Nweights) or vectors (if Nweights is 1).

## **Configuration Parameters**

• scalarization :: function

Function taking a fitness-matrix fitnesses (Nindivs x Nobjectives, with higher values indicating higher desirability) and a list of weight matrices weights (Nindivs elements of Nobjectives x Nweights matrices; positive weights should indicate a positive contribution to scale) and returns a matrix of scalarizations (Nindivs x Nweights, with higher values indicating greater desirability).

While custom functions can be used, it is recommended to use a Scalarizer, such as scalarizer\_linear(), or scalarizer\_chebyshev().

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("fixedprojection")
scls("fixedprojection") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("fixedprojection")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorFixedProjection
```

### **Active bindings**

```
weights_component_id (numeric(1)) search space component identifying the weights by which to scalarize.
```

#### Methods

#### **Public methods:**

- ScalorFixedProjection\$new()
- ScalorFixedProjection\$prime()
- ScalorFixedProjection\$clone()

Method new(): Initialize the ScalorFixedProjection object.

```
Usage:
```

ScalorFixedProjection\$new(weights\_component\_id = "scalarization\_weights")

Arguments:

```
weights_component_id (character(1))
```

Id of the search space component identifying the weights by which to scalarize. Default "scalarization\_weights".

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the operator given to the operation configuration parameter. Note that this modifies the \$param\_set\$values\$operation object.

```
Usage:
```

ScalorFixedProjection\$prime(param\_set)

Arguments:

```
param_set (ParamSet)
```

Passed to MiesOperator\$prime().

Returns: invisible self.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

ScalorFixedProjection\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

## See Also

 $Other scalors: Scalor, dict\_scalors\_aggregate, dict\_scalors\_domcount, dict\_scalors\_hypervolume, dict\_scalors\_nondom, dict\_scalors\_one, dict\_scalors\_proxy, dict\_scalors\_single$ 

Other scalor wrappers: dict\_scalors\_aggregate, dict\_scalors\_proxy

## **Examples**

```
set.seed(1)
```

```
dict_scalors_hypervolume
```

Hypervolume Scalor

## **Description**

Scalor that returns the hypervolume of each individual, relative to nadir and as a contribution over baseline. The returned scalar value is the measure of all points that have fitnesses that are

- greater than the respective value in nadir in all dimensions, and
- smaller than the respective value in the given point in all dimensions, and
- greater than all points in baseline in at least one dimension.

baseline should probably be a paradox::ContextPV and generate fitness values from the Archive in the context using mies\_get\_fitnesses.

## **Configuration Parameters**

- scale\_output :: logical(1)
  Whether to scale output to lie between 0 and 1.
- nadir:: numeric
   Nadir of fitness values relative to which hypervolume ution is calculated.
- baseline:: matrix
   Fitness-matrix with one column per objective, giving a population over which the hypervolume improvement should be calculated.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("hypervolume")
scls("hypervolume") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("hypervolume")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorHypervolume
```

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#### Methods

#### **Public methods:**

```
• ScalorHypervolume$new()
```

• ScalorHypervolume\$clone()

```
Method new(): Initialize the ScalorHypervolume object.
```

```
Usage:
```

```
ScalorHypervolume$new()
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

```
ScalorHypervolume$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

## See Also

Other scalors: Scalor, dict\_scalors\_aggregate, dict\_scalors\_domcount, dict\_scalors\_fixedprojection, dict\_scalors\_nondom, dict\_scalors\_one, dict\_scalors\_proxy, dict\_scalors\_single

## **Examples**

```
sv = scl("hypervolume")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that ScalorHV does not depend on data content
data = data.frame(x = rep(0, 5))
fitnesses = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2)
sv$param_set$values$baseline = matrix(c(1, 1), ncol = 2)
sv$param_set$values$nadir = c(0, -1)
sv$prime(p)
sv$operate(data, fitnesses)
```

dict\_scalors\_nondom

Nondominated Sorting Scalor

## **Description**

Scalor that returns a the rank of the pareto-front in nondominated sorting as scale. Higher ranks indocate higher fitnesses and therefore "better" individuals.

dict\_scalors\_nondom

## **Configuration Parameters**

- epsilon
- nadir
- jitter
- scale\_output
- tiebreak

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("nondom")
scls("nondom") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("nondom")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorNondom
```

#### Methods

## **Public methods:**

- ScalorNondom\$new()
- ScalorNondom\$clone()

**Method** new(): Initialize the ScalorNondom object.

```
Usage:
ScalorNondom$new()
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
ScalorNondom$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

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## See Also

Other scalors: Scalor, dict\_scalors\_aggregate, dict\_scalors\_domcount, dict\_scalors\_fixedprojection, dict\_scalors\_hypervolume, dict\_scalors\_one, dict\_scalors\_proxy, dict\_scalors\_single

### **Examples**

## **Description**

Scalor that returns a the fitness value of a single objective dimension as scale.

## **Configuration Parameters**

• objective :: integer(1) objective to return as scale, ranges from 1 (the default, first objective) to the number of objectives of the function being optimized.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("one")
scls("one") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("one")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorOne
```

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## Methods

#### **Public methods:**

```
• ScalorOne$new()
```

```
• ScalorOne$clone()
```

```
Method new(): Initialize the ScalorOne object.
```

```
Usage:
ScalorOne$new()
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
ScalorOne$clone(deep = FALSE)
Arguments:
```

deep Whether to make a deep clone.

## See Also

Other scalors: Scalor, dict\_scalors\_aggregate, dict\_scalors\_domcount, dict\_scalors\_fixedprojection, dict\_scalors\_hypervolume, dict\_scalors\_nondom, dict\_scalors\_proxy, dict\_scalors\_single

## **Examples**

```
so = scl("one")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that ScalorOne does not depend on data content
data = data.frame(x = rep(0, 5))
fitnesses = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2)
so$prime(p)
so$operate(data, fitnesses)
so$param_set$values$objective = 2
so$operate(data, fitnesses)
```

dict\_scalors\_proxy

Proxy-Scalor that Scales According to its Configuration parameter

## Description

Scalor that performs the operation in its operation configuration parameter. This is useful, e.g., to make SelectorBest's operation fully parametrizable.

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## **Configuration Parameters**

• operation :: Scalor

Operation to perform. Initialized to ScalorSingleObjective. This is primed when \$prime() of ScalorProxy is called, and also when \$perate() is called, to make changing the operation as part of self-adaption possible. However, if the same operation gets used inside multiple ScalorProxy objects, then it is recommended to \$clone(deep = TRUE) the object before assigning them to operation to avoid frequent re-priming.

## **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

## **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("proxy")
scls("proxy") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("proxy")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorProxy
```

## Methods

#### **Public methods:**

- ScalorProxy\$new()
- ScalorProxy\$prime()
- ScalorProxy\$clone()

**Method** new(): Initialize the ScalorProxy object.

```
Usage:
ScalorProxy$new()
```

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the operator given to the operation configuration parameter. Note that this modifies the \$param\_set\$values\$operation object.

```
Usage:
ScalorProxy$prime(param_set)
Arguments:
```

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```
param_set (ParamSet)
    Passed to MiesOperator$prime().

Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.

Usage:
ScalorProxy$clone(deep = FALSE)

Arguments:
deep Whether to make a deep clone.
```

## See Also

Other scalors: Scalor, dict\_scalors\_aggregate, dict\_scalors\_domcount, dict\_scalors\_fixedprojection, dict\_scalors\_hypervolume, dict\_scalors\_nondom, dict\_scalors\_one, dict\_scalors\_single

Other scalor wrappers: dict\_scalors\_aggregate, dict\_scalors\_fixedprojection

# Examples

```
set.seed(1)
sp = scl("proxy")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that ScalorOne does not depend on data content
data = data.frame(x = rep(0, 5))
fitnesses = c(1, 5, 2, 3, 0)

sp$param_set$values$operation = scl("one")
sp$prime(p)
sp$operate(data, fitnesses)
```

## **Description**

Scalor that uses a single given objective, throwing an error in case it is used in a multi-objective problem.

In contrast to ScalorOne, this Scalor throws an error when more than one objective is present. When this Scalor gets used as the default value, e.g. for a Selector, then it forces the user to make an explicit decision about what Scalor to use in a multi-objective setting.

## **Configuration Parameters**

No configuration parameters.

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# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

### **Dictionary**

This Scalor can be created with the short access form scl() (scls() to get a list), or through the the dictionary dict\_scalors in the following way:

```
# preferred:
scl("single")
scls("single") # takes vector IDs, returns list of Scalors
# long form:
dict_scalors$get("single")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Scalor -> ScalorSingleObjective
```

#### Methods

### **Public methods:**

- ScalorSingleObjective\$new()
- ScalorSingleObjective\$clone()

**Method** new(): Initialize the ScalorSingleObjective object.

Usage:

ScalorSingleObjective\$new()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

ScalorSingleObjective\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

```
Other scalors: Scalor, dict_scalors_aggregate, dict_scalors_domcount, dict_scalors_fixedprojection, dict_scalors_hypervolume, dict_scalors_nondom, dict_scalors_one, dict_scalors_proxy
```

```
ss = scl("single")

p = ps(x = p_dbl(-5, 5))

# dummy data; note that ScalorOne does not depend on data content

data = data.frame(x = rep(0, 5))

fitnesses_so = c(1, 5, 2, 3, 0)
```

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```
fitnesses_mo = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2)
ss$prime(p)
ss$operate(data, fitnesses_so)
try(ss$operate(data, fitnesses_mo))
```

dict\_selectors

Dictionary of Selectors

# Description

Dictionary of Selectors

## Usage

```
dict_selectors
```

#### **Format**

An object of class DictionarySelector (inherits from DictionaryEx, Dictionary, R6) of length 15.

# Methods

Methods inherited from Dictionary, as well as:

```
    help(key, help_type)
        (character(1), character(1))
        Displays help for the dictionary entry key. help_type is one of "text", "html", "pdf" and given as the help_type argument of R's help().
```

# See Also

Other dictionaries: dict\_filtors, dict\_mutators, dict\_recombinators, dict\_scalors, mut()

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## **Description**

Selector that selects the top n\_select individuals based on the fitness value, breaking ties randomly. When n\_select is larger than the number of individuals, the selection wraps around: All nrow(values) individuals are selected at least floor(nrow(values) / n\_select) times, with the top nrow(values) %% n\_select individuals being selected one more time.

### **Configuration Parameters**

• shuffle\_selection :: logical(1)
Whether to shuffle the selected output. When this is TRUE, selected individuals are returned in random order, so when this operator is e.g. used in mies\_generate\_offspring(), then subsequent recombination operators effectively operate on pairs (or larger groups) of random individuals. Otherwise they are returned in order, and recombination operates on the first batch of n\_indivs\_in returned individuals first, then the second batch etc. in order. Initialized to TRUE (recommended).

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

#### **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("best")
sels("best") # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("best")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> miesmuschel::SelectorScalar
-> SelectorBest
```

#### Methods

#### **Public methods:**

- SelectorBest\$new()
- SelectorBest\$clone()

```
Method new(): Initialize the SelectorBest object.
    Usage:
    SelectorBest$new(scalor = ScalorSingleObjective$new())
    Arguments:
    scalor (Scalor)
        Scalor to use to generate scalar values from multiple objectives, if multi-objective optimization is performed. Initialized to ScalorSingleObjective: Doing single-objective optimization normally, throwing an error if used in multi-objective setting: In that case, a Scalor needs to be explicitly chosen.
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
SelectorBest$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

```
Other selectors: Selector, SelectorScalar, dict_selectors_maybe, dict_selectors_null, dict_selectors_proxy, dict_selectors_random, dict_selectors_sequential, dict_selectors_tournament
```

# **Examples**

```
sb = sel("best")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that SelectorBest does not depend on data content
data = data.frame(x = rep(0, 5))
fitnesses = c(1, 5, 2, 3, 0)
sb$prime(p)
sb$operate(data, fitnesses, 2)
sb$param_set$values$shuffle_selection = FALSE
sb$operate(data, fitnesses, 4)
```

dict\_selectors\_maybe Selector-Combination that Selects According to Two Selectors

#### **Description**

Selector that wraps two other Selectors given during construction and uses both for selection proportionally. Each of the resulting n\_select individuals is chosen either from \$selector, or from \$selector\_not.

This makes it possible to implement selection methods such as random interleaving, where only a fraction of p individuals were selected by a criterion, while the others are taken randomly.

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#### Algorithm

To perform selection, n\_selector\_in rows of values are given to \$selector, and the remaining nrow(values) - n\_selector\_in rows are given to \$selector\_not. Both selectors are used to generate a subset of selected individuals: \$selector generates n\_selector\_out individuals, and \$selector\_not generates n\_select - n\_selector\_out individuals.

n\_selector\_in is either set to round(nrow(values) \* p\_in) when proportion\_in is "exact", or to rbinom(1, nrow(values), p\_in) when proportion\_in is "random".

n\_selector\_out is set to round(n\_select \* p\_out) when proportion\_out is "exact", or to rbinom(1, n\_select, p\_out) when proportion\_out is "random".

When odds\_correction is TRUE, then p\_out is adjusted depending on the used n\_selector\_in value before being applied. Let odds(p) = p/(1-p). Then the effective p\_out is set such that odds(effective p\_out) = odds(p\_out) \* n\_selector\_in / (nrow(values) - n\_selector\_in) / odds(p\_in). This corrects for the discrepancy between the chosen p\_in and the effective proportion of n\_selector\_in / nrow(values) caused either by rounding errors or when proportion\_in is "random".

When p\_in is exactly 1 or exactly 0, and p\_out is not equal to p\_in, then an error is given.

If nrow(values) is 1, then this individuum is returned and \$selector / \$selector\_not are not called.

If try\_unique is TRUE, then n\_selector\_out is set to at most n\_selector\_in and at least n\_select - nrow(values) + n\_selector\_in, and an error is generated when nrow(values) is less than n\_select.

If try\_unique is FALSE and odds\_correction is TRUE and n\_selector\_in is either 0 or nrow(values), then \$p\_out is set to either 0 or 1, respectively.

If try\_unique is FALSE and odds\_correction is FALSE and n\_selector\_in is either 0 or nrow(values), and n\_selector\_out is not equal to 0 or n\_select, respectively, then n\_selector\_in is increased / decreased by 1 to give \$selector\_not / \$selector at least one individuum to choose from. While this behaviour may seem pathological, it is to ensure continuity with sampled values of n\_selector\_in that are close to 0 or n\_select.

If n\_selector\_out is n\_select or 0, or if n\_selector\_in is nrows(values) - 1 or 1, then only \$selector\_not is executed, respectively; possibly with a subset of values if n\_selector\_in differs from nrow(values) / 0.

### **Configuration Parameters**

This operator has the configuration parameters of the Selectors that it wraps: The configuration parameters of the operator given to the selector construction argument are prefixed with "maybe.", the configuration parameters of the operator given to the selector\_not construction argument are prefixed with "maybe\_not.".

Additional configuration parameters:

p\_in :: numeric(1)
 Probability per individu

Probability per individual (when random\_choise is TRUE), or fraction of individuals (when random\_choice is FALSE), that are given to \$selector instead of \$selector\_not. This may be overriden when try\_unique is TRUE, in which case at least as many rows are given to \$selector and \$selector\_not as they are generating output values respectively. When this is exactly 1 or exactly 0, then p\_out must be equal to p\_in. Must be set by the user.

- p\_out :: numeric(1)
  - Probability per output value (when random\_choise is TRUE), or fraction of output values (when random\_choice is FALSE), that are generated by \$selector instead of \$selector\_not. When this values is not given, it defaults to p\_in.
- shuffle\_input :: logical(1)
  - Whether to distribute input values randomly to \$selector / \$selector\_not. If FALSE, then the first part of values is given to \$selector. This only randomizes *which* lines of values are given to \$selector / \$selector\_not, but it does not necessarily reorder the lines of values given to each. In particular, if p\_out is 0 or 1, then no shuffling takes place. Initialized to TRUE.
- proportion\_in :: character(1)
  When set to "random", sample the number of individuals given to \$selector according to rbinom(1, nrow(values), p\_in). When set to "exact", give \$selector round(nrow(values) \* p\_in) individuals. Initialized to "exact".
- proportion\_out :: character(1)
  When set to "random", sample the number of individuals generated by \$selector according to rbinom(1, n\_select, p\_out). When set to "exact", have \$selector generate round(n\_select \*p\_out) individuals.
- odds\_correction :: logical(1)
   When set, the effectively used value of p\_out is set to 1 / (1 + ((nrow(values) n\_selector\_in) \* p\_in \* (1 p\_out)) / (n\_selector\_in \* p\_out \* (1 p\_in))), see the Algorithm section. Initialized to FALSE.
- try\_unique :: logical(1)
  Whether to give at least as many rows of values to each of \$selector and \$selector\_not as they are generating output values. This should be set to TRUE whenever SelectorMaybe is used to select unique values, and can be set to FALSE when selecting values multiple times is acceptable. When this is TRUE, then having n\_select > nrow(values) generates an error. Initialized to TRUE.

# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of selector and selector\_not.

### **Dictionary**

This Filtor can be created with the short access form ftr() (ftrs() to get a list), or through the the dictionary dict\_filtors in the following way:

```
# preferred:
ftr("maybe", <selector> [, <selector_not>])
ftrs("maybe", <selector> [, <selector_not>]) # takes vector IDs, returns list of Filtors
# long form:
dict_filtors$get("maybe", <selector> [, <selector_not>])
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> SelectorMaybe
```

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# **Active bindings**

```
selector (Selector)
```

Selector being wrapped. This operator gets run with probability / proportion p\_in and generates output with probability / proportion p\_out (configuration parameters).

```
selector_not (Selector)
```

Alternative Selector being wrapped. This operator gets run with probability / proportion 1 – p\_in and generates output with probability / proportion 1 – p\_out (configuration parameters).

#### Methods

### **Public methods:**

- SelectorMaybe\$new()
- SelectorMaybe\$prime()
- SelectorMaybe\$clone()

**Method** new(): Initialize the SelectorMaybe object.

```
Usage:
```

```
SelectorMaybe$new(selector, selector_not = SelectorRandom$new())
```

Arguments:

```
selector (Selector)
```

Selector to wrap. This operator gets run with probability / fraction p\_in (Configuration parameter).

The constructed object gets a *clone* of this argument. The \$selector field will reflect this value.

```
selector_not (Selector)
```

Another Selector to wrap. This operator runs when selector is not chosen. By default, this is SelectorRandom, i.e. selecting randomly.

The constructed object gets a *clone* of this argument. The \$selector\_not field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to selector and selector\_not during construction.

```
Usage:
```

```
SelectorMaybe$prime(param_set)
```

Arguments:

```
param_set (ParamSet)
```

Passed to MiesOperator\$prime().

Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.

Usage:

```
SelectorMaybe$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

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### See Also

Other selectors: Selector, SelectorScalar, dict\_selectors\_best, dict\_selectors\_null, dict\_selectors\_proxy, dict\_selectors\_random, dict\_selectors\_sequential, dict\_selectors\_tournament Other selector wrappers: dict\_selectors\_proxy, dict\_selectors\_sequential

```
dict_selectors_null Null Selector
```

# **Description**

Selector that disregards fitness and individual values and selects individuals by order in which they are given.

## **Configuration Parameters**

• shuffle\_selection :: logical(1)
Whether to shuffle the selected output. When this is TRUE, selected individuals are returned in random order, so when this operator is e.g. used in mies\_generate\_offspring(), then subsequent recombination operators effectively operate on pairs (or larger groups) of random individuals. Otherwise they are returned in order, and recombination operates on the first batch of n\_indivs\_in returned individuals first, then the second batch etc. in order. Initialized to TRUE (recommended).

### **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("null")
sels("null") # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("null")
```

## Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> SelectorNull
```

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#### Methods

#### **Public methods:**

```
• SelectorNull$new()
```

```
• SelectorNull$clone()
```

```
Method new(): Initialize the SelectorNull object.
   Usage:
   SelectorNull$new()
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
SelectorNull$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

### See Also

```
Other selectors: Selector, SelectorScalar, dict_selectors_best, dict_selectors_maybe, dict_selectors_proxy, dict_selectors_random, dict_selectors_sequential, dict_selectors_tournament
```

# **Examples**

```
sn = sel("null")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that SelectorNull does not depend on data content
data = data.frame(x = rep(0, 5))
fitnesses = c(1, 5, 2, 3, 0)
sn$prime(p)
sn$operate(data, fitnesses, 2)
sn$operate(data, fitnesses, 4)
sn$operate(data, fitnesses, 6)
```

dict\_selectors\_proxy Proxy-Selector that Selects According to its Configuration Parameter

## **Description**

Selector that performs the operation in its operation configuration parameter. This is useful, e.g., to make OptimizerMies's selection operations fully parametrizable.

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# **Configuration Parameters**

• operation :: Selector

Operation to perform. Initialized to SelectorBest. This is primed when \$prime() of SelectorProxy is called, and also when \$operate() is called, to make changing the operation as part of self-adaption possible. However, if the same operation gets used inside multiple SelectorProxy objects, then it is recommended to \$clone(deep = TRUE) the object before assigning them to operation to avoid frequent re-priming.

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("proxy")
sels("proxy") # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("proxy")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> SelectorProxy
```

# Methods

#### **Public methods:**

- SelectorProxy\$new()
- SelectorProxy\$prime()
- SelectorProxy\$clone()

**Method** new(): Initialize the SelectorProxy object.

```
Usage:
SelectorProxy$new()
```

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the operator given to the operation configuration parameter. Note that this modifies the \$param\_set\$values\$operation object.

```
Usage:
SelectorProxy$prime(param_set)
Arguments:
```

```
param_set (ParamSet)
    Passed to MiesOperator$prime().
Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.
Usage:
SelectorProxy$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

```
Other selectors: Selector, SelectorScalar, dict_selectors_best, dict_selectors_maybe, dict_selectors_null, dict_selectors_random, dict_selectors_sequential, dict_selectors_tournament Other selector wrappers: dict_selectors_maybe, dict_selectors_sequential
```

### **Examples**

```
set.seed(1)
sp = sel("proxy")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that SelectorBest does not depend on data content
data = data.frame(x = rep(0, 5))
fitnesses = c(1, 5, 2, 3, 0)

sp$param_set$values$operation = sel("random")
sp$prime(p)
sp$operate(data, fitnesses, 2)

sp$param_set$values$operation = sel("best")
sp$operate(data, fitnesses, 2)
```

dict\_selectors\_random Random Selector

### **Description**

Random selector that disregards fitness and individual values and selects individuals randomly. Depending on the configuration parameter replace, it samples with or without replacement.

# **Configuration Parameters**

• sample\_unique :: character(1)
Whether to sample individuals globally unique ("global"), unique within groups ("groups"),
or not unique at all ("no", sample with replacement). This is done with best effort; if group\_size
(when sample\_unique is "groups") or n\_select (when sample\_unique is "global") is
greater than nrow(values), then individuals are selected with as few repeats as possible. Initialized to "groups".

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

### **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("random")
sels("random") # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("random")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> SelectorRandom
```

#### Methods

### **Public methods:**

- SelectorRandom\$new()
- SelectorRandom\$clone()

Method new(): Initialize the SelectorRandom object.

Usage:

SelectorRandom\$new()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

SelectorRandom\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

```
Other selectors: Selector, SelectorScalar, dict_selectors_best, dict_selectors_maybe, dict_selectors_null, dict_selectors_proxy, dict_selectors_sequential, dict_selectors_tournament
```

```
set.seed(1)
sr = sel("random")
p = ps(x = p_dbl(-5, 5))
# dummy data; note that SelectorRandom does not depend on data content
data = data.frame(x = rep(0, 5))
```

```
fitnesses = c(1, 5, 2, 3, 0)
sr$prime(p)
sr$operate(data, fitnesses, 2)
sr$operate(data, fitnesses, 2)
sr$operate(data, fitnesses, 2)
sr$operate(data, fitnesses, 4)
sr$operate(data, fitnesses, 4)
sr$operate(data, fitnesses, 4)
```

dict\_selectors\_sequential

Run Multiple Selection Operations in Sequence

### **Description**

Selector that wraps multiple other Selectors given during construction and uses them for selection in sequence. This makes it possible for one Selector to discard a few individuals, followed by a second Selector to discard more, etc., until n\_select individuals are remaining.

# Algorithm

Given that there are nrow(values) input individuals in an operation, and n\_select individuals requested to be selected, the operation calls selector\_i for i in 1 ... length(selectors) to reduce the number of individuals in this pipeline. The relative quantity by which the number of individuals is reduced in each step is determined by the configuration parameters reduction\_1, reduction\_2, etc., and also dependent on the sum of these values, in the following denoted, with a slight abuse of notation, by sum[reduction\_#].

Let the number of individuals passed to step i be denoted by n\_values[i], and the number of individuals requested to be selected by that step be denoted as n\_select\_[i]. In particular, n\_values[1] == nrow(values), and n\_select\_[length(selectors)] == n\_select.

n\_values[1] == nrow(values), and n\_select\_[length(selectors)] == n\_select.
When reduction\_by\_factor is TRUE, then the reduction at step i is done by a factor, meaning that

n\_values[i] / n\_select\_[i] is set (up to rounding). This factor is (nrow(values) / n\_select) ^ (reduction\_i / sum[

When reduction\_by\_factor is FALSE, then the reduction at step i is done by absolute differences, meaning that n\_values[i] - n\_select\_[i] is set (up to rounding). This difference is (nrow(values) - n\_select) \* (reduction\_i / sum[reduction\_#]), with sum[reduction\_#] as above.

In particular, this means that when all reduction\_# values are the same and reduction\_by\_factor is TRUE, then each operation reduces the number of individuals in the pipeline by the same factor. When reduction\_by\_factor is FALSE, then each operation removes the same absolute number of individuals.

While the illustrations are done with the assumption that nrow(values) >= n\_select, they hold equivalently with nrow(values) < n\_select.

All except the last Selectors are called with group\_size set to their n\_select value; the last Selector is called with the group\_size value given as input.

### **Configuration Parameters**

This operator has the configuration parameters of the Selectors that it wraps: The configuration parameters of the operator given to the selectors construction argument are prefixed with "selector\_1", "selector\_2", ... up to "selector\_#", where # is length(selectors).

Additional configuration parameters:

- reduction\_1, reduction\_2, ... :: numeric(1)
  Relative reduction done by selector\_1, selector\_2, ..., as described in the section Algorithm. The values are all initialized to 1, meaning the same factor (when reduction\_by\_factor is TRUE) or absolute number (otherwise) of reduction by each operation.
- reduction\_by\_factor :: logical(1)
   Whether to do reduction by factor (TRUE) or absolute number (FALSE), as described in Algorithm. Initialized to TRUE.

# **Supported Operand Types**

Supported Domain classes are the set intersection of supported classes of the Selectors given in selectors.

## **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("sequential", <selectors>)
sels("sequential", <selectors>) # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("sequential", <selectors>)
```

# Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Selector-> SelectorSequential
```

# **Active bindings**

```
selectors (list of Selector)
Selectors being wrapped. These operators get run sequentially in order.
```

#### Methods

#### **Public methods:**

- SelectorSequential\$new()
- SelectorSequential\$prime()
- SelectorSequential\$clone()

**Method** new(): Initialize the SelectorSequential object.

SelectorSequential\$new(selectors)

Usage:

```
Arguments:
       selectors (list of Selector)
           Selectors to wrap. The operations are run in order given to selectors. The constructed
           object gets a clone of this argument. The $selectors field will reflect this value.
     Method prime(): See MiesOperator method. Primes both this operator, as well as the wrapped
     operators given to selectors during construction.
       Usage:
       SelectorSequential$prime(param_set)
       Arguments:
       param_set (ParamSet)
           Passed to MiesOperator$prime().
       Returns: invisible self.
     Method clone(): The objects of this class are cloneable with this method.
       SelectorSequential$clone(deep = FALSE)
       Arguments:
       deep Whether to make a deep clone.
See Also
    Other selectors: Selector, SelectorScalar, dict_selectors_best, dict_selectors_maybe,
    dict_selectors_null, dict_selectors_proxy, dict_selectors_random, dict_selectors_tournament
    Other selector wrappers: dict_selectors_maybe, dict_selectors_proxy
```

### **Description**

Selector that repeatedly samples k individuals and selects the best ouf of these.

Tournament Selector

# **Configuration Parameters**

dict\_selectors\_tournament

- k :: integer(1)

  Tournament size. Must be set by the user.
- choose\_per\_tournament :: Number of individuals to choose in each tournament. Must be smaller than k. The special value 0 sets this to the group\_size hint given to the \$operate()-call (but at most k). This is equal to n\_select when used as survival-selector in mies\_survival\_plus()/mies\_survival and equal to n\_indivs\_in of a Recombinator used in mies\_generate\_offspring(). Initialized to 1.

• sample\_unique :: character(1)

Whether to sample individuals globally unique ("global", selected individuals are removed from the population after each tournament), unique within groups ("groups", individuals are replaced when group\_size individuals were sampled), unique per tournament ("tournament", individuals are replaced after each tournament), or not unique at all ("no", individuals are sampled with replacement within tournaments). This is done with best effort; if group\_size (when sample\_unique is "groups") or n\_select (when sample\_unique is "global") is greater than nrow(values), then the first nrow(values) \* floor(group\_size / nrow(values)) or nrow(values) \* floor(n\_select / nrow(values)) individuals are chosen deterministically by selecting every individual with the same frequency, followed by tournament selection for the remaining required individuals. Initialized to "groups".

# **Supported Operand Types**

```
Supported Domain classes are: p_lgl ('ParamLgl'), p_int ('ParamInt'), p_dbl ('ParamDbl'), p_fct ('ParamFct')
```

# **Dictionary**

This Selector can be created with the short access form sel() (sels() to get a list), or through the the dictionary dict\_selectors in the following way:

```
# preferred:
sel("tournament")
sels("tournament") # takes vector IDs, returns list of Selectors
# long form:
dict_selectors$get("tournament")
```

# Super classes

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> miesmuschel::SelectorScalar
-> SelectorTournament
```

# Methods

### **Public methods:**

- SelectorTournament\$new()
- SelectorTournament\$clone()

**Method** new(): Initialize the SelectorTournament object.

```
Usage:
SelectorTournament$new(scalor = ScalorSingleObjective$new())
Arguments:
scalor (Scalor)
```

Scalor to use to generate scalar values from multiple objectives, if multi-objective optimization is performed. Initialized to ScalorSingleObjective: Doing single-objective optimization normally, throwing an error if used in multi-objective setting: In that case, a Scalor needs to be explicitly chosen.

dist\_crowding 89

```
Method clone(): The objects of this class are cloneable with this method.
```

```
Usage:
```

```
SelectorTournament$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

#### See Also

```
Other selectors: Selector, SelectorScalar, dict_selectors_best, dict_selectors_maybe, dict_selectors_null, dict_selectors_proxy, dict_selectors_random, dict_selectors_sequential
```

## **Examples**

```
sb = sel("tournament", k = 4)
p = ps(x = p_dbl(-5, 5))
# dummy data; note that SelectorBest does not depend on data content
data = data.frame(x = rep(0, 7))
fitnesses = c(1, 5, 2, 3, 0, 4, 6)
sb$prime(p)
sb$operate(data, fitnesses, 2)
sb$operate(data, fitnesses, 4, group_size = 2)
```

dist\_crowding

Calculate Crowding Distance

#### **Description**

Takes a matrix of fitness values and calculates the crowding distance for individuals in that matrix. Individuals that are minimal or maximal with respect to at least one dimension are assigned infinite crowding distance.

Individuals are assumed to be in a (epsilon-) nondominated front.

# Usage

```
dist_crowding(fitnesses)
```

### **Arguments**

```
fitnesses (numeric matrix)
```

fitness matrix, with one row per individual and one column per objective

#### Value

numeric: Vector of crowding distances.

90 domhy

domhv

Calculate Dominated Hypervolume

# **Description**

Use Chan's algorithm (Chan, M T (2013). "Klee's measure problem made easy." In 2013 IEEE 54th annual symposium on foundations of computer science, 410–419. IEEE.) to calculate dominated hypervolume.

# Usage

```
domhv(fitnesses, nadir = 0, prefilter = TRUE, on_worse_than_nadir = "warn")
```

# **Arguments**

fitnesses (numeric matrix)

fitness matrix, with one row per individual and one column per objective

nadir (numeric)

Lowest fitness point up to which to calculate dominated hypervolume. May be a scalar, in which case it is used for all dimensions, or a vector, in which case its

length must match the number of dimensions. Default 0.

prefilter (logical(1))

Whether to make a first pass that filters out dominated individuals. If it can be guaranteed that all individuals are non-dominated, setting this to FALSE improves performance a bit. Otherwise the recommended value is the default

FALSE.

on\_worse\_than\_nadir

(character(1)) Action when individuals that do not dominate the nadir are found. One of "quiet" (ignore), "warn" (give warning, default), or "stop"

(throw error).

#### Value

numeric(1): The dominated hypervolume of individuals in fitnesses.

```
(fitnesses = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2))
# to see the fitness matrix, use:
## plot(fitnesses, pch = as.character(1:5))
domhv(fitnesses)
```

domhv\_contribution 91

domhv\_contribution

Calculate Hypervolume Contribution

## Description

Takes a matrix of fitness values and calculates the hypervolume contributions of individuals in that matrix.

Hypervolume contribution of an individual I is the difference between the dominated hypervolume of a set of individuals including I, where the fitness of I is increased by epsilon, and the dominated hypervolume of the same set but excluding I.

Individuals that are less than another individual more than epsilon in any dimension have hypervolume contribution of 0.

# Usage

```
domhv_contribution(fitnesses, nadir = 0, epsilon = 0)
```

### Arguments

fitnesses (numeric matrix)

fitness matrix, with one row per individual and one column per objective

nadir (numeric)

Lowest fitness point up to which to calculate dominated hypervolume. May be a scalar, in which case it is used for all dimensions, or a vector, in which case its

length must match the number of dimensions. Default 0.

epsilon (numeric)

Added to each individual before calculating its particular hypervolume contribution. epsilon may be a scalar, in which case it is used for all dimensions, or a vector, in which case its length must match the number of dimensions. Default

0.

#### Value

numeric: The vector of dominated hypervolume contributions for each individual in fitnesses.

```
(fitnesses = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2))
# to see the fitness matrix, use:
## plot(fitnesses, pch = as.character(1:5))
domhv_contribution(fitnesses)
```

92 domhv\_improvement

domhv\_improvement

Calculate Hypervolume Improvement

### **Description**

Takes a matrix of fitness values and calculates the hypervolume improvement of individuals in that matrix, one by one, over the baseline individuals.

The hypervolume improvement for each point is the measure of all points that have fitnesses that are

- greater than the respective value in nadir in all dimensions, and
- smaller than the respective value in the given point in all dimensions, and
- greater than all points in baseline in at least one dimension.

Individuals in fitnesses are considered independently of each other. A possible speedup is achieved because baseline individuals only need to be pre-filtered once.

# Usage

```
domhv_improvement(fitnesses, baseline = NULL, nadir = 0)
```

## **Arguments**

fitnesses (numeric matrix)

fitness matrix, with one row per individual and one column per objective

baseline (matrix|NULL)

Fitness-matrix with one column per objective, giving a population over which the hypervolume improvement should be calculated. If NULL, the hypervolume

of each individual in fitnesses is calculated.

nadir (numeric)

Lowest fitness point up to which to calculate dominated hypervolume. May be a scalar, in which case it is used for all dimensions, or a vector, in which case its

length must match the number of dimensions. Default 0.

#### Value

numeric: The vector of dominated hypervolume contributions for each individual in fitnesses.

```
(fitnesses = matrix(c(1, 5, 2, 3, 0, 3, 1, 0, 10, 8), ncol = 2))
# to see the fitness matrix, use:
## plot(fitnesses, pch = as.character(1:5))
domhv_improvement(fitnesses)
domhv_improvement(fitnesses, fitnesses[1, , drop = FALSE])
```

Filtor

Filtor Base Class

# Description

Base class representing filter operations, inheriting from MiesOperator.

A Filtor gets a table of individuals that are to be filtered, as well as a table of individuals that were already evaluated, along with information on the latter individuals' performance values. Furthermore, the number of individuals to return is given. The Filtor returns a vector of unique integers indicating which individuals were selected.

Filter operations are performed in ES algorithms to facilitate concentration towards individuals that likely perform well with regard to the fitness measure, without evaluating the fitness measure, for example through a surrogate model.

Fitness values are always *maximized*, both in single- and multi-criterion optimization.

Unlike most other operator types inheriting from MiesOperator, the \$operate() function has four arguments, which are passed on to \$.filter()

- values:: data.frame
   Individuals to filter. Must pass the check of the ParamSet given in the last \$prime() call and may not have any missing components.
- known\_values :: data.frame
  Individuals to use for filtering. Must pass the check of the ParamSet given in the last \$prime()
  call and may not have any missing components. Note that known\_values may be empty.
- fitnesses :: numeric | matrix Fitnesses for each individual given in known\_values. If this is a numeric, then its length must be equal to the number of rows in values. If this is a matrix, if number of rows must be equal to the number of rows in values, and it must have one column when doing single-crit optimization and one column each for each "criterion" when doing multi-crit optimization.
- n\_filter:: integer(1)

  Number of individuals to select. Some Filtors select individuals with replacement, for which this value may be greater than the number of rows in values.

The return value for an operation will be a numeric vector of integer values of ength n\_filter indexing the individuals that were selected. Filtor must always return unique integers, i.e. select every individual at most once.

### Inheriting

Filtor is an abstract base class and should be inherited from. Inheriting classes should implement the private \$.filter() function. The user of the object calls \$operate(), and the arguments are passed on to private \$.filter() after checking that the operator is primed, that the values and known\_values arguments conforms to the primed domain and that other values match.

The private\$.needed\_input() function should also be overloaded, it is called by the public \$needed\_input() function after initial checks; see the documentation there.

Typically, the \$initialize() function should also be overloaded, and optionally the \$prime() function; they should call their super equivalents.

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#### Super class

```
miesmuschel::MiesOperator-> Filtor
```

### **Active bindings**

```
supported (character)

Optimization supported by this Filtor, can be "single-crit", "multi-crit", or both.
```

### Methods

#### **Public methods:**

- Filtor\$new()
- Filtor\$needed\_input()
- Filtor\$clone()

**Method** new(): Initialize base class components of the Filtor.

```
Usage:
Filtor$new(
   param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
   param_set = ps(),
   supported = c("single-crit", "multi-crit"),
   packages = character(0),
   dict_entry = NULL,
   own_param_set = quote(self$param_set)
)
Arguments:
```

param\_classes (character)

Classes of parameters that the operator can handle. May contain any of "ParamLgl", "ParamInt", "ParamDbl", "ParamFct". Default is all of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

```
supported (character)
```

Subset of "single-crit" and "multi-crit", indicating wether single and / or multi-criterion optimization is supported. Default both of them.

The \$supported field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

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```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** needed\_input(): Calculate the number of values that are required to filter down to output\_size, given the current configuration parameter settings.

Usage:

Filtor\$needed\_input(output\_size)

Arguments:

output\_size (integer(1))

A positive integer indicating the number of individuals for which the needed input size should be calculated.

*Returns*: integer(1): The minimum number of rows required to filter down to output\_size. At least output\_size.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

Filtor\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

### See Also

Other base classes: FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, Selector, SelectorScalar Other filtors: FiltorSurrogate, dict\_filtors\_maybe, dict\_filtors\_null, dict\_filtors\_proxy, dict\_filtors\_surpog, dict\_filtors\_surtour

FiltorSurrogate

Abstract Surrogate Model Filtering Base Class

## Description

Abstract base class for surrogate model filtering.

A *surrogate model* is a regression model, based on an mlr3::Learner, which predicts the approximate performance of newly sampled configurations given the empirical performance of already evaluated configurations. The surrogate model can be used to propose points that have, according to the surrogate model, a relatively high chance of performing well.

The FiltorSurrogate base class can be inherited from to create different Filtors that filter based on a surrogate model, for example tournament filtering or progresive filtering.

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#### **Configuration Parameters**

FiltorSurrogateProgressive's configuration parameters are the hyperparameters of the surrogate\_learner Learner, as well as the configuration parameters of the surrogate\_selector Selector.

# **Supported Operand Types**

Supported Domain classes depend on the supported feature types of the surrogate\_learner, as reported by surrogate\_learner\$feature\_types: "ParamInt" requires "integer", "ParamDbl" requires "numeric", "ParamLgl" requires "logical", and "ParamFct" requires "factor".

### Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Filtor-> FiltorSurrogate
```

# **Active bindings**

```
surrogate_learner (mlr3::LearnerRegr)
    Regression learner for the surrogate model filtering algorithm.
surrogate_selector (Selector)
    Selector with which to select using surrogate-predicted performance
```

#### Methods

## **Public methods:**

- FiltorSurrogate\$new()
- FiltorSurrogate\$prime()
- FiltorSurrogate\$clone()

**Method** new(): Initialize the base class components of the FiltorSurrogate.

```
Usage:
FiltorSurrogate$new(
  surrogate_learner,
  surrogate_selector = SelectorBest$new(),
  param_set = ps(),
  packages = character(0),
  dict_entry = NULL
Arguments:
surrogate_learner (mlr3::LearnerRegr)
   Regression learner for the surrogate model filtering algorithm.
   The $surrogate_learner field will reflect this value.
surrogate_learner (mlr3::LearnerRegr)
   Regression learner for the surrogate model filtering algorithm.
   The $surrogate_learner field will reflect this value.
surrogate_selector (Selector) Selector for the surrogate model filtering algorithm.
   The $surrogate_selector field will reflect this value.
```

```
surrogate_selector (Selector) Selector for the surrogate model filtering algorithm.
```

The \$surrogate\_selector field will reflect this value.

```
param_set (ParamSet)
```

ParamSet of the method implemented in the inheriting class with configuration parameters that go beyond the parameters of the surrogate\_learner and surrogate\_selector.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operator given to surrogate\_selector during construction.

```
Usage:
FiltorSurrogate$prime(param_set)
Arguments:
param_set (ParamSet)
    Passed to MiesOperator$prime().
Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.
Usage:
FiltorSurrogate$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

Other base classes: Filtor, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, Selector, SelectorScalar

Other filtors: Filtor, dict\_filtors\_maybe, dict\_filtors\_null, dict\_filtors\_proxy, dict\_filtors\_surprog, dict\_filtors\_surtour

MiesOperator

Operator Base Class

## **Description**

Base class representing MIES-operators: Recombinator, Mutator, and Selector.

Operators perform a specific function within ES algorithms, and by exchanging them, the character of ES algorithms can be modified. Operators operate on collections of individuals and return modified individuals (mutated or recombined) or indices of selected individuals. Operators can be combined using MutatorCombination / RecombinatorCombination and other operators wrappers.

Before applying operators, they have to be *primed* for the domain of the individuals which they are operating on; this is done using the \$prime() function. Afterwards, the \$operate() function may be called with a data.frame of individuals that fall into this domain. \$operate() may be called multiple times after priming, and a once primed operator can be primed again for a different domain by calling \$prime() agian (which forgets the old priming).

# Inheriting

MiesOperator is an abstract base class and should be inherited from. Inheriting classes should implement the private \( \).operate() function. The user of the object calls \( \)operate(), and the arguments are passed on to private \( \).operate() after checking that the operator is primed, and that the values argument conforms to the primed domain. Typically, the \( \)initialize() and \( \)prime() functions are also overloaded, but should call their super equivalents.

In most cases, the MiesOperator class should not be inherited from, directly; instead, the operator classes (Recombinator, Mutator, Selector) or their subclasses should be inherited.

# **Active bindings**

```
param_set (ParamSet)
    Configuration parameters of the MiesOperator object. Read-only.

param_classes (character)
    Classes of parameters that the operator can handle, contains any of "ParamLgl", "ParamInt",
    "ParamDbl", "ParamFct". Read-only.

packages (character)
    Packages needed for the operator. Read-only.

dict_entry (character(1) | NULL)
    Key of this class in its respective Dictionary. Is NULL if this class it not (known to be) in a
    Dictionary. Read-only.
```

```
dict_shortaccess (character(1) | NULL)
    Name of Dictionary short-access function where an object of this class can be retrieved. Is
    NULL if this class is not (known to be) in a Dictionary with a short-access function. Read-
    only.
endomorphism (logical(1))
     Whether the output of $operate() is a data.frame / data.table in the same domain as its
    input. Read-only.
primed_ps (ParamSet | NULL)
    ParamSet on which the MiesOperator is primed. Is NULL if it has not been primed. Writing
     to this acrive binding calls $prime().
is_primed (logical(1))
     Whether the MiesOperator was primed before. Is FALSE exactly when $primed_ps is NULL.
     Read-only.
man (character(1))
    Name of this class, in the form <package>::<classname>. Used by the $help() method.
 Public methods:
```

#### Methods

- MiesOperator\$new()
- MiesOperator\$repr()
- MiesOperator\$print()
- MiesOperator\$prime()
- MiesOperator\$operate()
- MiesOperator\$help()
- MiesOperator\$clone()

Method new(): Initialize base class components of the MiesOperator.

```
Usage:
MiesOperator$new(
  param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
  param_set = ps(),
  packages = character(0),
  dict_entry = NULL,
  dict_shortaccess = NULL,
  own_param_set = quote(self$param_set),
  endomorphism = TRUE
)
Arguments:
param_classes (character)
   Classes of parameters that the operator can handle. May contain any of "ParamLg1",
   "ParamInt", "ParamDbl", "ParamFct". Default is all of them.
   The $param_classes field will reflect this value.
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to

super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
dict_shortaccess (character(1) | NULL)
```

Name of the Dictionary short access function in which the operator is registered. This is used to inform the user about how to construct a given object. Should ordinarily be one of "mut", "rec", "sel".

The \$dict\_shortaccess field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

```
endomorphism (logical(1))
```

Whether the private \$.operate() operation creates a data.table with the same columns as the input (i.e. conforming to the primed ParamSet). If this is TRUE (default), then the return value of \$.operate() is checked for this and columns are put in the correct order. The \$endomorphsim field will reflect this value.

**Method** repr(): Create a call object representing this operator.

```
Usage:
MiesOperator$repr(
    skip_defaults = TRUE,
    show_params = TRUE,
    show_constructor_args = TRUE,
    ...
)
Arguments:
skip_defaults (logical(1))
    Whether to skip construction arguments that have their default value. Default TRUE.
show_params (logical(1))
    Whether to show ParamSet values. Default TRUE.
show_constructor_args (logical(1))
    Whether to show construction args that are not ParamSet values. Default TRUE.
... (any)
    Ignored.
```

```
Method print(): Print this operator.
 MiesOperator$print(verbose = FALSE, ...)
 Arguments:
 verbose (logical(1))
     Whether to show all construction arguments, even the ones at default values. Default FALSE.
 ... (any)
     Ignored.
Method prime(): Prepare the MiesOperator to function on the given ParamSet. This must be
called before $operate(). It may be called multiple times in the lifecycle of the MiesOperator
object, and prior primings are forgotten when priming on a new ParamSet. The ParamSet on
which the MiesOperator was last primed can be read from $primed_ps.
 Usage:
 MiesOperator$prime(param_set)
 Arguments:
 param_set (ParamSet)
     The ParamSet to which all values tables passed to $operate() will need to conform to.
     May only contiain Domain objects that conform to the classes listed in $param_classes.
 Returns: invisible self.
Method operate(): Operate on the given individuals. This calls private $.operate(), which
must be overloaded by an inheriting class, passing through all function arguments after performing
some checks.
 Usage:
 MiesOperator$operate(values, ...)
 Arguments:
 values (data.frame)
     Individuals to operate on. Must pass the check of the ParamSet given in the last $prime()
     call and may not have any missing components.
 ... (any)
     Depending on the concrete class, passed on to $.operate().
 Returns: data.frame: the result of the operation. If the input was a data.table instead of a
 data. frame, the output is also data. table.
Method help(): Run utils::help() for this object.
 Usage:
 MiesOperator$help(help_type = getOption("help_type"))
 Arguments:
 help_type (character(1))
     One of "text", "html", or "pdf": The type of help page to open. Defaults to the "help_type"
     option.
 Returns: help_files_with_dopic object, which opens the help page.
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
MiesOperator$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

Other base classes: Filtor, FiltorSurrogate, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, Selector, SelectorScalar

```
mies_aggregate_generations
```

Get Aggregated Performance Values by Generation

# **Description**

Get evaluated performance values from an OptimInstance aggregated for each generation. This may either concern all individuals that were alive at the end of a given generation (survivors\_only TRUE) or at any point during a generation (survivors\_only FALSE).

The result is a single data.table object with a dob column indicating the generation, as well as one column for each aggregations entry crossed with each objective of inst.

See mies\_generation\_apply() on how to apply functions to entire fitness-matrices, not only individual objectives.

# Usage

```
mies_aggregate_generations(
   inst,
   objectives = inst$archive$codomain$ids(),
   aggregations = list(min = min, mean = mean, max = max, median = stats::median, size =
        length),
   as_fitnesses = TRUE,
   survivors_only = TRUE,
   condition_on_budget_id = NULL
)
```

### **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

objectives (character)

Objectives for which to calculate aggregates. Must be a subset of the codomain elements of inst, but when as\_fitnesses is TRUE, elements that are neither being minimized nor maximized are ignored.

aggregations (named list of function)

List containing aggregation functions to be evaluated on a vector of objective falues for each generation. These functions should take a single argument and return a scalar value.

as\_fitnesses (

(logical(1))

Whether to transform performance values into "fitness" values that are always to be maximized. This means that values that objectives that should originally be minimized are multiplied with -1, and that parts of the objective codomain that are neither being minimized nor maximized are dropped. Default TRUE.

survivors\_only (logical(1))

Whether to ignore configurations that have "eol" set to the given generation, i.e. individuals that were killed during that generation. When this is TRUE (default), then only individuals that are alive at the *end* of a generation are considered; otherwise all individuals alive at any point of a generation are considered. If it is TRUE, this leads to individuals that have "dob" == "eol" being ignored.

condition\_on\_budget\_id

(character(1) | NULL)

Budget component when doing multi-fidelity optimization. When this is given, then for each generation, only individuals with the highest value for this component are considered. If survivors\_only is TRUE, this means the highest value of all survivors of a given generation, if it is FALSE, then it is the highest value of all individuals alive at any point of a generation. To ignore possible budget-parameters, set this to NULL (default). This is inparticular necessary when fidelity is not monotonically increasing (e.g. if it is categorical).

#### Value

a data.table with the column "dob", indicating the generation, as well as further columns named by the items in aggregations. There is more on element in objectives (or more than one element not being minimized/maximized when as\_fitnesses is TRUE), then columns are named <aggregations element name>.<objective name>. Otherwise, they are named by <aggregations element name> only. To get a guarantee that elements are only named after elements in aggregations, set objectives to a length 1 character.

# See Also

Other aggregation methods: mies\_aggregate\_single\_generation(), mies\_get\_generation\_results()

```
library("bbotk")
lgr::threshold("warn")

# Define the objective to optimize
objective <- ObjectiveRFun$new(
  fun = function(xs) {
    z <- 10 - exp(-xs$x^2 - xs$y^2) + 2 * exp(-(2 - xs$x)^2 - (2 - xs$y)^2)
    list(Obj = z)
},</pre>
```

```
domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "minimize"))
)
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 6)
)
op <- opt("mies",</pre>
  lambda = 2, mu = 2,
  mutator = mut("gauss", sdev = 0.1),
  recombinator = rec("xounif"),
  parent_selector = sel("best")
set.seed(1)
op$optimize(oi)
# negates objectives that are minimized:
mies_aggregate_generations(oi)
# silly aggregation: first element
mies_aggregate_generations(oi, aggregations = list(first = function(x) x[1]))
# real objective values:
mies_aggregate_generations(oi, as_fitnesses = FALSE)
# Individuals that died are included:
mies_aggregate_generations(oi, survivors_only = FALSE)
```

mies\_aggregate\_single\_generation

Aggregate a Value for a given Generation

# Description

Applies a fitness\_aggregator function to the values that were alive in the archive at a given generation. The function is supplied with the fitness values, and optionally other data, of all individuals that are alive at that point.

### Usage

```
mies_aggregate_single_generation(
   archive,
   fitness_aggregator,
   generation = NA,
   include_previous_generations = FALSE
)
```

### Arguments

archive (Archive)

The archive over which to aggregate.

fitness\_aggregator

(function)

Aggregation function, called with information about alive individuals of each generation. See details.

generation (numeric(1))

Generation for which to aggregate the value. If include\_previous\_generations is FALSE, then an individual is considered to be alive at generation i if its dob is smaller or equal to i, and if its eol is either NA or greater than i. If include\_previous\_generations is TRUE, then all individuals with dob smaller or equal to i are considered. If this is NA, the currently alive (include\_previous\_generations FALSE) or all (include\_previous\_generations TRUE) individuals are aggregated. If multiple individuals considered "alive" with the same x\_id are found, then only the last individual is used. This excludes previous individuals that were re-evaluated with a different fidelity.

include\_previous\_generations

(logical(1))

Aggregate all individuals that were alive at generation or at any point before that. Duplicates with the same x\_id are removed, meaning that if an individual was re-evaluated with different fidelity, only the last re-evaluation is counted. However, note that individuals from different generations may still have been evaluated with different fidelity, so if Default FALSE.

#### **Details**

The fitness\_aggregator function may have any of the following arguments, which will be given the following information when fitness\_aggregator is called:

• fitnesses :: matrix

Will contain fitnesses for each alive individual. This value has one column when doing single-crit optimization and one column for each "criterion" when doing multi-crit optimization. Fitnesses are always being maximized, so if an objective is being minimized, the fitness\_aggregator function is given the objective values \* -1.

• objectives\_unscaled :: matrix

The objective values as given to fitnesses, but not multiplied by -1 if they are being minimized. It is recommended that the codomain argument is queried for "maximize" or "minimize" tags when objectives\_unscaled is used.

• budget :: scalar

If multi-fidelity evaluation is being performed, then this is the "budget" value of each individual. Otherwise it is a vector containing the value 1 for each individual.

• xdt :: data.table

The configurations that were evaluated for the alive individuals. Rows are in the same order as the values given to fitnesses or objectives\_unscaled.

• search\_space :: ParamSet

The search space of the Archive under evaluation.

• codomain :: ParamSet

The codomain of the Archive under evaluation. This is particularly useful when using objectives\_unscaled to determine minimization or maximization.

Not all of these arguments need to present, but at least one of fitnesses, objectives\_unscaled, or xdt must be.

fitness\_aggregator will never be called for an empty generation.

#### Value

The value returned by fitness\_aggregator when applied to individuals alive at generation generation. If no individuals of the requested generation are present, fitness\_aggregator is not called and mies\_aggregate\_single\_generation() returns NULL instead.

#### See Also

Other aggregation methods: mies\_aggregate\_generations(), mies\_get\_generation\_results()

```
library("bbotk")
lgr::threshold("warn")
objective <- ObjectiveRFun$new(</pre>
  fun = function(xs) {
   list(y1 = xs$x1, y2 = xs$x2)
  domain = ps(x1 = p_dbl(0, 1), x2 = p_dbl(-1, 0)),
  codomain = ps(y1 = p_dbl(0, 1, tags = "maximize"),
   y2 = p_dbl(-1, 0, tags = "minimize"))
)
oi <- OptimInstanceMultiCrit$new(objective, terminator = trm("none"))</pre>
try(mies_aggregate_single_generation(oi$archive, identity), silent = TRUE)
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses)
mies_init_population(oi, 2, budget_id = "x1", fidelity = .5)
oi$archive$data
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses)
# Notice how fitnesses are positive, since x2 is scaled with -1.
# To get the original objective-values, use objectives_unscaled:
mies_aggregate_single_generation(oi$archive,
  function(objectives_unscaled) objectives_unscaled)
# When `...` is used, all information is passed:
mies_aggregate_single_generation(oi$archive, function(...) names(list(...)))
```

```
# Generation 10 is not present, but individuals with eol `NA` are still
# considered alive:
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses,
 generation = 10)
# Re-evaluating points with higher "fidelity" (x1)
mies_step_fidelity(oi, budget_id = "x1", fidelity = 0.7)
oi$archive$data
# Lower-fidelity values are considered dead now, even for generation 1:
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses,
 generation = 1)
# This adds two new alive individuals at generation 2.
# Also the individuals from gen 1 are reevaluated with fidelity 0.8
mies_evaluate_offspring(oi, offspring = data.frame(x2 = c(-0.1, -0.2)),
 budget_id = "x1", fidelity = 0.9, reevaluate_fidelity = 0.8)
oi$archive$data
mies_aggregate_single_generation(oi$archive, function(budget, ...) budget)
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses,
 generation = 1)
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses,
 generation = 2)
# No individuals were killed, but some were fidelity-reevaluated.
# These are not present with include_previous_generations:
mies_aggregate_single_generation(oi$archive, function(fitnesses) fitnesses,
 generation = 2, include_previous_generations = TRUE)
# Typical use-case: get dominated hypervolume
mies_aggregate_single_generation(oi$archive, function(fitnesses) domhv(fitnesses))
# Get generation-wise mean fitness values
mies_aggregate_single_generation(oi$archive, function(fitnesses) {
 apply(fitnesses, 2, mean)
})
```

mies\_evaluate\_offspring

Evaluate Proposed Configurations Generated in a MIES Iteration

### **Description**

Calls \$eval\_batch of a given OptimInstance on a set of configurations as part of a MIES operation. The dob extra-info in the archive is also set properly to indicate a progressed generation.

This function can be used directly, but it is easier to use it within the OptimizerMies class if standard GA operation is desired.

Multifidelity evaluation is supported as described in vignette("mies-multifid"). For this, an extra component named after budget\_id is appended to each individual, chosen from the fidelity argument and depending on the value of survivor\_budget. budget\_id should have the same values as given to the other mies\_\* functions.

# Usage

```
mies_evaluate_offspring(
   inst,
   offspring,
   budget_id = NULL,
   fidelity = NULL,
   reevaluate_fidelity = NULL,
   fidelity_monotonic = TRUE
)
```

## **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

offspring (data.frame)

Proposed configurations to be evaluated, must have columns named after the

inst's search space, minus budget\_id if not NULL.

budget\_id (character(1) | NULL)

Budget component when doing multi-fidelity optimization. This component of the search space is added to individuals according to fidelity. Should be

NULL when no multi-fidelity optimization is performed (default).

fidelity (atomic(1)|NULL)

Atomic scalar indicating the value to be assigned to the budget\_id component of offspring. This value must be NULL if no multi-fidelity optimization is performed (the default).

reevaluate\_fidelity

(atomic(1))

Fidelity with which to evaluate alive individuals from previous generations that have a budget value below (if fidelity\_monotonic is TRUE) or different from the current fidelity value. Default NULL: Do not re-evaluate. Must be NULL when budget\_id and fidelity are NULL. See also mies\_step\_fidelity.

fidelity\_monotonic

(logical(1))

When reevaluate\_fidelity is non-NULL, then this indicates whether individuals should only ever be re-evaluated when fidelity would be increased. Default TRUE. Ignored when reevaluate\_fidelity is NULL

#### Value

invisible data.table: the performance values returned when evaluating the offspring values through eval\_batch.

#### See Also

Other mies building blocks: mies\_generate\_offspring(), mies\_get\_fitnesses(), mies\_init\_population(), mies\_select\_from\_archive(), mies\_step\_fidelity(), mies\_survival\_comma(), mies\_survival\_plus()

```
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
  fun = function(xs) {
   z \leftarrow exp(-xs$x^2 - xs$y^2) + 2 * exp(-(2 - xs$x)^2 - (2 - xs$y)^2)
   list(Obj = z)
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
mies_init_population(inst = oi, mu = 3)
# Initial state:
oi$archive
# 'offspring' is just a data.frame of values to evaluate.
# In general it should be created using 'mies_generate_offspring()'.
offspring = data.frame(x = 1:2, y = 2:1)
mies_evaluate_offspring(oi, offspring = offspring)
# This evaluated the given points and assigned them 'dob' 2.
oi$archive
# Note that at this point one would ordinarily call a 'mies_survival_*()'
# function.
# Advanced demo, making use of additional components and doing multi-fidelity
# declare 'y' the budget parameter. It will not be in the 'offspring'
# table any more.
budget_id = "y"
```

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```
# but: offspring may contain any other value that is appended to 'oi'. These
# are ignored by the objective.
offspring = data.frame(x = 0:1, z = 3)
mies_evaluate_offspring(oi, offspring = offspring, budget_id = budget_id,
  fidelity = 1)
# This now has the additional column 'z'. Values of y for the new evaluations
# are 1.
oi$archive
offspring = data.frame(x = 2, z = 3)
# Increasing the fidelity will not cause re-evaluation of existing individuals
# when `reevaluate_fidelity` is not given.
mies_evaluate_offspring(oi, offspring = offspring, budget_id = budget_id,
  fidelity = 2)
oi$archive
offspring = data.frame(x = 3, z = 3)
# Depending on the effect of fidelity, this may however have a biasing effect,
# so it may be desirable to re-evaluate surviving individuals from previous
# generations. The 'reevaluate_fidelity' may even be different from 'fidelity'
mies_evaluate_offspring(oi, offspring = offspring, budget_id = budget_id,
  fidelity = 3, reevaluate_fidelity = 2)
# In this example, only individuals with 'y = 1' were re-evaluated, since
# 'fidelity_monotonic' is TRUE.
oi$archive
```

mies\_filter\_offspring Filter Offspring

#### **Description**

Uses a Filtor to extract a subset of individuals from a given set. The individuals are either returned directly (when get\_indivs is TRUE) or in form of an index into the given individuals (when get\_indivs is FALSE).

Filtors must always select individuals without replacement, so selected individual indices are unique.

# Usage

```
mies_filter_offspring(
  inst,
  individuals,
  lambda,
  filtor = NULL,
  budget_id = NULL,
```

```
fidelity = NULL,
  get_indivs = TRUE
)
```

# Arguments

inst (OptimInstance)

Optimization instance to evaluate.

individuals (data.frame|data.table)

Individuals to filter. Must have columns according to filtor\$primed\_ps, and

must have at least filtor\$needed\_input(lambda) rows.

lambda (integer(1))

Number of individuals to filter down to.

filtor (Filtor | NULL)

Filtor operator that filters. When NULL is given, then the FiltorNull operation

is performed and the first lambda individuals are taken from individuals.

budget\_id (character(1) | NULL)

Budget component when doing multi-fidelity optimization. This component of

the search space is added to individuals according to fidelity. Should be

NULL when no multi-fidelity optimization is performed (default).

fidelity (atomic|NULL)

scalar indicating the value of the budget\_id component with which to evaluate

individuals to be filtered.

This value must be NULL when no multi-fidelity optimization is performed, but it may **also** be NULL when the maximum value of the budget\_id found in

inst\$archive should be used (the default).

get\_indivs (logical(1))

Whether to return the data. frame or data. table of selected individuals, or an

index into individuals.

# Value

If get\_indivs is TRUE: a data.frame or data.table (depending on the input type of individuals) of filtered configurations. If get\_indivs is FALSE: an integer vector indexing the filtered individuals.

mies\_generate\_offspring

Generate Offspring Through Mutation and Recombination

# **Description**

Generate new proposal individuals to be evaluated using mies\_evaluate\_offspring().

Parent individuals are selected using parent\_selector, then mutated using mutator, and thend recombined using recombinator. If only a subset of these operations is desired, then it is possible to set mutator or recombinator to the respective "null"-operators.

# Usage

```
mies_generate_offspring(
  inst,
  lambda,
  parent_selector = NULL,
  mutator = NULL,
  recombinator = NULL,
  budget_id = NULL
)
```

### Arguments

inst (OptimInstance)

Optimization instance to evaluate.

lambda (integer(1))

Number of new individuals to generate. This is not necessarily the number with which parent\_selector gets called, because recombinator could in principle need more than lambda input individuals to generate lambda output individuals.

parent\_selector

(Selector | NULL)

Selector operator that selects parent individuals depending on configuration values and objective results. When parent\_selector\$operate() is called, then objectives that are being minimized are multiplied with -1 (through mies\_get\_fitnesses()), since Selectors always try to maximize fitness. When this is NULL (default), then a SelectorBest us used.

The Selector must be primed on a superset of inst\$search\_space; this *includes* the "budget" component when performing multi-fidelity optimization. All components on which selector is primed on must occur in the archive. The given Selector *may* return duplicates.

mutator (Mutator | NULL)

Mutator operation to apply to individuals selected out of inst using parent\_selector. The Mutator must be primed on a ParamSet similar to inst\$search\_space, but without the "budget" component when budget\_id is given (multi-fidelity optimization). Such a ParamSet can be generated for example using mies\_prime\_operators. When this is NULL (default), then a MutatorNull is used, effectively disabling

mutation.

recombinator (Recombinator | NULL)

Recombinator operation to apply to individuals selected out of int using parent\_selector after mutation using mutator. The Recombinator must be primed on a ParamSet similar to inst\$search\_space, but without the "budget" component when budget\_id is given (multi-fidelity optimization). Such a ParamSet can be generated for example using mies\_prime\_operators.

When this is NULL (default), then a RecombinatorNull is used, effectively disabling recombination.

budget\_id (character(1) | NULL)

Budget component when doing multi-fidelity optimization. This component of the search space is removed from individuals sampled from the archive in inst before giving it to mutator and recombinator. Should be NULL when not doing multi-fidelity.

#### Value

data.table: A table of configurations proposed as offspring to be evaluated using mies\_evaluate\_offspring().

#### See Also

```
Other mies building blocks: mies_evaluate_offspring(), mies_get_fitnesses(), mies_init_population(), mies_select_from_archive(), mies_step_fidelity(), mies_survival_comma(), mies_survival_plus()
```

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(</pre>
  fun = function(xs) {
   z \leftarrow exp(-xs$x^2 - xs$y^2) + 2 * exp(-(2 - xs$x)^2 - (2 - xs$y)^2)
   list(Obj = z)
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
# Demo operators
m = mut("gauss", sdev = 0.1)
r = rec("xounif")
s = sel("random")
# Operators must be primed
mies_prime_operators(objective$domain, list(m), list(r), list(s))
# We would normally call mies_init_population, but for reproducibility
# we are going to evaluate three given points
oi\$eval_batch(data.table::data.table(x = 0:2, y = 2:0, dob = 1, eol = NA_real_))
# Evaluated points:
oi$archive
# Use default operators: no mutation, no recombination, parent_selctor is
# sel("best") --> get one individual, the one with highest performance in the
# archive (x = 1, y = 1).
```

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```
# (Note 'mies_generate_offspring()' does not modify 'oi')
mies_generate_offspring(oi, lambda = 1)

# Mutate the selected individual after selection. 'm' has 'sdev' set to 0.1,
# so the (x = 1, y = 1) is slightly permuted.
mies_generate_offspring(oi, lambda = 1, mutator = m)

# Recombination, then mutation.
# Even though lambda is 1, there will be two individuals selected with
# sel("best") and recombined, because rec("xounif") needs two parents. One
# of the crossover results is discarded (respecting that 'lambda' is 1),
# the other is mutated and returned.
mies_generate_offspring(oi, lambda = 1, mutator = m, recombinator = r)

# General application: select, recombine, then mutate.
mies_generate_offspring(oi, lambda = 5, parent_selector = s, mutator = m, recombinator = r)
```

mies\_generation

Get the Last Generation that was Evaluated

# **Description**

Gets the last generation that was evaluated as counted by the "dob" column in the OptimInstance's Archive.

This accepts OptimInstances that were not evaluated with miesmuschel and are therefore missing the "dob" column, returning a value of 0. However, if the "dob" column is invalid (the inferred generation is not integer numeric or not non-negative), an error is thrown.

# Usage

```
mies_generation(inst)
```

# **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

### Value

a scalar integer value indicating the last generation that was evaluated in inst. It is 0 when inst is empty, and also typically 0 if all evaluations in inst so far were performed outside of miesmuschel. Every call of mies\_init\_population that actually performs evaluations, as well as each call to mies\_evaluate\_offspring with non-empty offspring, increases the generation by 1.

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# **Examples**

```
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
 fun = function(xs) {
   z <- 10 - exp(-xs$x^2 - xs$y^2) + 2 * exp(-(2 - xs$x)^2 - (2 - xs$y)^2)
   list(Obj = z)
 },
 domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
 codomain = ps(Obj = p_dbl(tags = "minimize"))
)
oi <- OptimInstanceSingleCrit$new(objective,</pre>
 terminator = trm("evals", n_evals = 6)
op <- opt("mies",
 lambda = 2, mu = 2,
 mutator = mut("gauss", sdev = 0.1),
 recombinator = rec("xounif"),
 parent_selector = sel("best")
)
set.seed(1)
mies_generation(oi)
op$optimize(oi)
mies_generation(oi)
oi$terminator = trm("evals", n_evals = 10)
op$optimize(oi)
mies_generation(oi)
```

mies\_generation\_apply Aggregate Values for All Generations Present

# **Description**

Applies a fitness\_aggregator function to the values that were alive in the archive at at any generation. mies\_aggregate\_single\_generation() is used, see there for more information about fitness\_aggregator.

Generations for which fitness\_aggregator returns NULL, or which are not present in any dob in the archive, or which contain no alive individuals (e.g. because eol is smaller or equal dob for all of them) are ignored.

as.list() is applied to the values returned by fitness\_aggregator, and data.table::rbindlist() is called on the list of resulting values. If the first non-NULL-value returned by fitness\_aggregator, then data.table::rbindlist() is called with fill = TRUE and use.names = TRUE.

If no non-empty generations are present, or fitness\_aggregator returns NULL on every call, then the return value is data.table(dob = numeric(0)).

In contrast with mies\_aggregate\_generations(), mies\_generate\_apply() can construct aggregated values for entire fitness matrices, not only individual objectives (see examples). However, mies\_aggregate\_generations() is simpler if per-objective aggregates are desired.

# Usage

```
mies_generation_apply(
   archive,
   fitness_aggregator,
   include_previous_generations = FALSE
)
```

# **Arguments**

```
archive (Archive)
The archive over which to aggregate.
fitness_aggregator
(function)
```

Aggregation function, called with information about alive individuals of each generation. See mies\_aggregate\_single\_generation().

include\_previous\_generations

(logical(1))

Aggregate all individuals that were alive at generation or at any point before that. Duplicates with the same x\_id are removed, meaning that if an individual was re-evaluated with different fidelity, only the last re-evaluation is counted. However, note that individuals from different generations may still have been evaluated with different fidelity, so if Default FALSE.

#### Value

data.table with columns dob, next to the columns constructed from the return values of fitness\_aggregator.

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")

objective <- ObjectiveRFun$new(
  fun = function(xs) {
    list(y1 = xs$x1, y2 = xs$x2)
  },
  domain = ps(x1 = p_dbl(0, 1), x2 = p_dbl(-1, 0)),
  codomain = ps(y1 = p_dbl(0, 1, tags = "maximize"),</pre>
```

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```
y2 = p_dbl(-1, 0, tags = "minimize"))
)
oi <- OptimInstanceMultiCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 40))
op <- opt("mies",
  lambda = 4, mu = 4,
  mutator = mut("gauss", sdev = 0.1),
  recombinator = rec("xounif"),
  parent_selector = sel("random"),
  survival_selector = sel("best", scl("hypervolume"))
)
op$optimize(oi)
# Aggregated hypervolume of individuals alive in each gen:
mies_generation_apply(oi$archive, function(fitnesses) {
  domhv(fitnesses)
})
# Aggregated hypervolume of all points evaluated up to each gen
# (may be slightly more, since the domhv of more points is evaluated).
# This would be the dominated hypervolume of the result set at each
# generation:
mies_generation_apply(oi$archive, function(fitnesses) {
  domhv(fitnesses)
}, include_previous_generations = TRUE)
# The following are simpler with mies_aggregate_single_generations():
mies_generation_apply(oi$archive, function(fitnesses) {
  apply(fitnesses, 2, mean)
})
# Compare:
mies_aggregate_generations(oi, aggregations = list(mean = mean))
mies_generation_apply(oi$archive, function(objectives_unscaled) {
  apply(objectives_unscaled, 2, mean)
})
# Compare:
mies_aggregate_generations(oi, aggregations = list(mean = mean),
  as_fitnesses = FALSE)
```

mies\_get\_fitnesses

Get Fitness Values from OptimInstance

# Description

Get fitness values in the correct form as used by Selector operators from an OptimInstance. This works for both single-criterion and multi-criterion optimization, and entails multiplying objectives with -1 if they are being minimized, since Selector tries to maximize fitness.

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# Usage

```
mies_get_fitnesses(inst, rows)
```

# **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

rows optional (integer)

Indices of rows within inst to consider. If this is not given, then the entire

archive is used.

#### Value

numeric matrix with length(rows) (if rows is given, otherwise nrow(inst\$archive\$data)) rows and one column for each objective: fitnesses to be maximized.

#### See Also

```
Other mies building blocks: mies_evaluate_offspring(), mies_generate_offspring(), mies_init_population(), mies_select_from_archive(), mies_step_fidelity(), mies_survival_comma(), mies_survival_plus()
```

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
  fun = function(xs) {
    z \leftarrow \exp(-xsx^2 - xsy^2) + 2 * \exp(-(2 - xsx)^2 - (2 - xsy)^2)
    list(Obj = z)
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
mies_init_population(inst = oi, mu = 3)
oi$archive
mies_get_fitnesses(oi, c(2, 3))
###
# Multi-objective, and automatic maximization:
objective2 <- ObjectiveRFun$new(</pre>
```

```
fun = function(xs) list(Obj1 = xs$x^2, Obj2 = -xs$y^2),
  domain = ps(x = p_dbl(-2, 4), y = p_dbl(-2, 4)),
  codomain = ps(
    Obj1 = p_dbl(tags = "minimize"),
    Obj2 = p_dbl(tags = "maximize")
)

# Using MultiCrit!
oi <- OptimInstanceMultiCrit$new(objective2,
    terminator = trm("evals", n_evals = 100)
)

mies_init_population(inst = oi, mu = 3)
oi$archive

# Note Obj1 has a different sign than in the archive.
mies_get_fitnesses(oi, c(2, 3))</pre>
```

```
mies_get_generation_results
```

Get Performance Values by Generation

# Description

Get evaluated performance values from an OptimInstance for all individuals that were alive at a given generation. Depending on survivors\_only, all individuals alive at the *end* of a generation are returned, or all individuals alive at any point during a generation.

The resulting data.table object is formatted for easy manipulation to get relevant information about optimization progress. To get aggregated values per generation, use by = "dob".

#### Usage

```
mies_get_generation_results(
   inst,
   as_fitnesses = TRUE,
   survivors_only = TRUE,
   condition_on_budget_id = NULL
)
```

### **Arguments**

```
inst (OptimInstance)
```

Optimization instance to evaluate.

```
as_fitnesses (logical(1))
```

Whether to transform performance values into "fitness" values that are always to be maximized. This means that values that objectives that should originally be

minimized are multiplied with -1, and that parts of the objective codomain that are neither being minimized nor maximized are dropped. Default TRUE.

```
survivors_only (logical(1))
```

Whether to ignore configurations that have "eol" set to the given generation, i.e. individuals that were killed during that generation. When this is TRUE (default), then only individuals that are alive at the *end* of a generation are considered; otherwise all individuals alive at any point of a generation are considered. If it is TRUE, this leads to individuals that have "dob" == "eol" being ignored.

Budget component when doing multi-fidelity optimization. When this is given, then for each generation, only individuals with the highest value for this component are considered. If survivors\_only is TRUE, this means the highest value of all survivors of a given generation, if it is FALSE, then it is the highest value of all individuals alive at any point of a generation. To ignore possible budget-parameters, set this to NULL (default). This is inparticular necessary when fidelity is not monotonically increasing (e.g. if it is categorical).

#### Value

a data.table with the column "dob", indicating the generation, as well as further columns named by the OptimInstance's objectives.

#### See Also

Other aggregation methods: mies\_aggregate\_generations(), mies\_aggregate\_single\_generation()

```
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
 fun = function(xs) {
   z <-10 - exp(-xs$x^2 - xs$y^2) + 2 * exp(-(2 - xs$x)^2 - (2 - xs$y)^2)
   list(Obj = z)
 domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
 codomain = ps(Obj = p_dbl(tags = "minimize"))
)
oi <- OptimInstanceSingleCrit$new(objective,
  terminator = trm("evals", n_evals = 6)
op <- opt("mies",
 lambda = 2, mu = 2,
 mutator = mut("gauss", sdev = 0.1),
 recombinator = rec("xounif"),
```

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```
parent_selector = sel("best")
)
set.seed(1)
op$optimize(oi)

# negates objectives that are minimized:
mies_get_generation_results(oi)

# real objective values:
mies_get_generation_results(oi, as_fitnesses = FALSE)

# Individuals that died are included:
mies_get_generation_results(oi, survivors_only = FALSE)
```

# **Description**

Set up an OptimInstance for MIES optimization. This adds the dob and eol columns to the instance's archive, and makes sure there are at least mu survivors (i.e. entries with eol set to NA) present. If there are already >= mu prior evaluations present, then the last mu of these remain alive (the other's eol set to 0); otherwise, up to mu new randomly sampled configurations are evaluated and added to the archive and have eol set to NA.

# Usage

```
mies_init_population(
   inst,
   mu,
   initializer = generate_design_random,
   survival_selector = SelectorBest$new()$prime(inst$search_space),
   budget_id = NULL,
   fidelity = NULL,
   fidelity_new_individuals_only = FALSE,
   fidelity_monotonic = TRUE,
   additional_component_sampler = NULL
)
```

# Arguments

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functioning like paradox::generate\_design\_random or paradox::generate\_design\_lhs. Note that paradox::generate\_design\_grid can not be used and must be wrapped with a custom function that ensures that only n individuals are produced. The generated design must correspond to the inst's \$search\_space; for components that are not in the objective's search space, the additional\_component\_sampler is used.

survival\_selector

(Selector)

Used when the given  ${\tt OptimInstance}$  already contains more individuals than mu.

Selector operator that selects surviving individuals depending on configuration values and objective results, When survival\_selector\$operate() is called, then objectives that are being minimized are multiplied with -1 (through mies\_get\_fitnesses), since Selectors always try to maximize fitness.

The Selector must be primed on inst\$search\_space; this *includes* the "budget" component when performing multi-fidelity optimization. Default is SelectorBest. The given Selector may *not* return duplicates.

budget\_id (character(1) | NULL)

Budget component when doing multi-fidelity optimization. This component of the search space is added to individuals according to fidelity. Should be NULL when no multi-fidelity optimization is performed (default).

fidelity (atomic(1)|NULL)

Atomic scalar indicating the value to be assigned to the budget\_id component of offspring. This value must be NULL if no multi-fidelity optimization is performed (the default).

fidelity\_new\_individuals\_only

(logical(1))

When fidelity is not NULL: Whether to re-evaluate individuals that are already present in inst should they have a smaller (if fidelity\_monotonic is TRUE) or different (if fidelity\_monotonic is FALSE) value from the one given to fidelity. Default FALSE. Ignored when fidelity is NULL.

fidelity\_monotonic

(logical(1))

Whether to only re-evaluate configurations for which the fidelity would increase. Default TRUE. Ignored when fidelity is NULL or when fidelity\_new\_individuals\_only is TRUE.

additional\_component\_sampler

(Sampler | NULL)

Sampler for components of individuals that are not part of inst's \$search\_space. These components are never used for performance evaluation, but they may be useful for self-adaptive OperatorCombinations. See the description of mies\_prime\_operators() on how operators need to be primed to respect additional components.

It is possible that additional\_component\_sampler is used for *more* rows than initializer, which happens when the inst's \$archive contains prior evaluations that are alive, but does not contain columns pertaining to additional columns, or contains *all* these columns but there are rows that are NA valued. If only *some* of the columns are present, or if all these columns are present but

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there are rows that are only NA valued for some columns, then an error is thrown. Default is NULL: no additional components.

#### Value

invisible OptimInstance: the input instance, modified by-reference.

#### See Also

```
Other mies building blocks: mies_evaluate_offspring(), mies_generate_offspring(), mies_get_fitnesses(), mies_select_from_archive(), mies_step_fidelity(), mies_survival_comma(), mies_survival_plus()
```

```
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
  fun = function(xs) {
    z \leftarrow \exp(-xsx^2 - xsy^2) + 2 * \exp(-(2 - xsx)^2 - (2 - xsy)^2)
   list(0bj = z)
  domain = ps(x = p_dbl(-2, 4), y = p_dbl(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
)
mies_init_population(inst = oi, mu = 3)
# 3 evaluations, archive contains 'dob' and 'eol'
oi$archive
# Advanced demo, making use of additional components and fidelity
##
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
mies_init_population(inst = oi, mu = 3, budget_id = "y", fidelity = 2,
  additional_component_sampler = Sampler1DRfun$new(
    param = ps(additional = p_dbl(-1, 1)), rfun = function(n) rep(-1, n)
  )
)
```

```
# 3 evaluations. We also have 'additional', sampled from rfun (always -1),
# which is ignored by the objective. Besides, we have "y", which is 2,
# according to 'fidelity'.
oi$archive
```

# **Description**

Prime the given MiesOperators for an optimization run with the given search space.

In its simplest form, MIES optimization only optimizes the search space of the Objective to be optimized. However, more advanced optimization may handle a "budget" parameter for multifidelity optimization differently: It is still selected by Selectors, but not mutated or recombined and instead handled separately. It is also possible to add additional components to the search space that are not evaluated by the objective function, but that are used for self-adaption by other operators.

The mies\_prime\_operators() function uses the information that the user usually has readily at hand—the Objectives search space, the budget parameter, and additional components—and primes [Mutator objects in the right way:

- Selectors are primed on a union of search\_space and additional\_components
- Mutators and Recombinators are primed on the Selector's space with the budget\_id Domain removed.

mies\_prime\_operators() is called with an arbitrary number of MiesOperator arguments; typically one Mutator, one Recombinator and at least two Selector: one for survival selection, and one parent selection. Supplied MiesOperators are primed by-reference, but they are also returned as invisible list.

If neither additional components nor multi-fidelity optimization is used, it is also possible to use the \$prime() function of hte MiesOperators directly, although using mies\_prime\_operators() gives flexibility for future extension.

# Usage

```
mies_prime_operators(
   search_space,
   mutators = list(),
   recombinators = list(),
   selectors = list(),
   filtors = list(),
   ...,
   additional_components = NULL,
   budget_id = NULL
)
```

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### **Arguments**

```
search_space
                  (ParamSet)
                 Search space of the Objective or OptimInstance to be optimized.
                 (list of Mutator)
mutators
                 Mutator objects to prime. May be empty (default).
recombinators
                 (list of Recombinator)
                 Recombinator objects to prime. May be empty (default).
selectors
                 (list of Selector)
                 Selector objects to prime. May be empty (default).
filtors
                  (list of Filtor)
                 Filtor objects to prime. May be empty (default).
                 Must not be given. Other operators may be added in the future, so the following
                 arguments should be passed by name.
additional_components
                 (ParamSet | NULL)
                  Additional components to optimize over, not included in search_space, but
                  possibly used for self-adaption. This must be the ParamSet of mies_init_population()'s
                  additional_component_sampler argument.
budget_id
                 (character(1) | NULL)
                  Budget component used for multi-fidelity optimization.
```

### Value

invisible named list with entries \$mutators (list of Mutator, primed mutators), \$recombinators (list of Recombinator, primed recombinators), and \$selectors (list of Selector, primed selectors).

```
# Search space of a potential TuningInstance for optimization:
search_space = ps(x = p_dbl(), y = p_dbl())
# Additional search space components that are not part of the TuningInstance
additional_components = ps(z = p_dbl())
# Budget parameter not subject to mutation or recombination
budget_id = "y"

m = mut("gauss")
r = rec("xounif")
s1 = sel("best")
s2 = sel("random")
f = ftr("null")

mies_prime_operators(search_space, mutators = list(m),
    recombinators = list(r), selectors = list(s1, s2), filtors = list(f),
    additional_components = additional_components, budget_id = budget_id
)
```

```
# contain search_space without budget parameter, with additional_components
m$primed_ps
r$primed_ps

# contain also the budget parameter
s1$primed_ps
s2$primed_ps
f$primed_ps
```

mies\_select\_from\_archive

Select Individuals from an OptimInstance

# **Description**

Apply a Selector operator to a subset of configurations inside an OptimInstance and return the index within the archive (when get\_indivs FALSE) or the configurations themselves (when get\_indivs is TRUE).

It is not strictly necessary for the selector to select unique individuals / individuals without replacement.

Individuals are selected independently of whether they are "alive" or not. To select only from alive individuals, set rows to inst\$archive\$data[, which(is.na(eol))].

# Usage

```
mies_select_from_archive(
   inst,
   n_select,
   rows,
   selector = SelectorBest$new()$prime(inst$search_space),
   group_size = 1,
   get_indivs = TRUE
)
```

# **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

n\_select (integer(1))

Number of individuals to select.

rows optional (integer)

Indices of rows within inst to consider. If this is not given, then the entire

archive is used.

selector (Selector)

Selector operator that selects individuals depending on configuration values and objective results. When selector\$operate() is called, then objectives that

are being minimized are multiplied with -1 (through mies\_get\_fitnesses()), since Selectors always try to maximize fitness. Defaults to SelectorBest. The Selector must be primed on a superset of inst\$search\_space; this *includes* the "budget" component when performing multi-fidelity optimization. All components on which selector is primed on must occur in the archive. The given Selector *may* return duplicates.

group\_size

(integer)

Sampling group size hint, indicating that the caller would prefer there to not be any duplicates within this group size. The Selector may or may not ignore this value, however. This may possibly happen because of certain configuration parameters, or because the input size is too small.

Must either be a scalar value or sum up to n\_select. Must be non-negative. A scalar value of 0 is interpreted the same as 1.

Default is 1.

get\_indivs

(logical(1))

Whether to return configuration values from within the archive (TRUE) or just the indices within the archive (FALSE). Default is TRUE.

#### Value

integer | data.table: Selected individuals, either index into inst or subset of archive table, depending on get\_indivs.

# See Also

Other mies building blocks: mies\_evaluate\_offspring(), mies\_generate\_offspring(), mies\_get\_fitnesses(), mies\_init\_population(), mies\_step\_fidelity(), mies\_survival\_comma(), mies\_survival\_plus()

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
  fun = function(xs) {
    z < -exp(-xs*x^2 - xs*y^2) + 2 * exp(-(2 - xs*x)^2 - (2 - xs*y)^2)
   list(Obj = z)
  },
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
)
s = sel("best")
```

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```
s$prime(oi$search_space)
mies_init_population(inst = oi, mu = 6)
oi$archive
# Default: get individuals
mies_select_from_archive(oi, n_select = 2, rows = 1:6, selector = s)
# Alternatively: get rows within archive
mies_select_from_archive(oi, n_select = 2, rows = 1:6, selector = s,
 get_indivs = FALSE)
# Rows gotten from archive are relative from *all* rows, not from archive[rows]:
mies_select_from_archive(oi, n_select = 2, rows = 3:6, selector = s,
 get_indivs = FALSE)
# When using additional components: mies_select_from_archive learns about
# additional components from primed selector.
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
 terminator = trm("evals", n_evals = 100)
mies_init_population(inst = oi, mu = 6,
 additional_component_sampler = Sampler1DRfun$new(
   param = ps(additional = p_dbl(-1, 1)), rfun = function(n) -1
)
oi$archive
# Wrong: using selector primed only on search space. The resulting
# individuals do not have the additional component.
mies_select_from_archive(oi, n_select = 2, rows = 1:6, selector = s)
# Correct: selector must be primed on search space + additional component
mies_prime_operators(oi$search_space, selectors = list(s),
 additional_components = ps(additional = p_dbl(-1, 1)))
mies_select_from_archive(oi, n_select = 2, rows = 1:6, selector = s)
```

mies\_step\_fidelity Re-Evaluate Existing Configurations with Higher Fidelity

### Description

As part of the "rolling-tide" multifidelity-setup, do reevaluation of configurations with higher fidelity that have survived lower-fidelity selection. The evaluations are done as part of the *current*  mies\_step\_fidelity 129

generation, so the dob value is not increased.

This function should only be called when doing rolling-tide multifidelity, and should not be part of the MIES cycle otherwise.

# Usage

```
mies_step_fidelity(
   inst,
   budget_id,
   fidelity,
   current_gen_only = FALSE,
   fidelity_monotonic = TRUE,
   additional_components = NULL
)
```

# **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

budget\_id (character(1))

Budget component that is set to the fidelity value.

fidelity (atomic(1))

Atomic scalar indicating the value to be assigned to the budget\_id component of offspring.

current\_gen\_only

(logical(1))

Whether to only re-evaluate survivors individuals generated in the latest generation (TRUE), or re-evaluate all currently alive individuals (FALSE). In any case, only individuals that were not already evaluated with the chosen fidelity are evaluated, so this will usually only have an effect when the fidelity of surviving individuals changed between generations.

fidelity\_monotonic

(logical(1))

Whether to only re-evaluate configurations for which the fidelity would increase. Default TRUE.

additional\_components

(ParamSet | NULL)

Additional components to optimize over, not included in search\_space, but possibly used for self-adaption. This must be the ParamSet of mies\_init\_population()'s additional\_component\_sampler argument.

### Value

invisible data.table: the performance values returned when evaluating the offspring values through eval\_batch.

mies\_step\_fidelity

# See Also

Other mies building blocks: mies\_evaluate\_offspring(), mies\_generate\_offspring(), mies\_get\_fitnesses(), mies\_init\_population(), mies\_select\_from\_archive(), mies\_survival\_comma(), mies\_survival\_plus()

```
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(</pre>
  fun = function(xs) {
   z \leftarrow \exp(-xsx^2 - xsy^2) + 2 * \exp(-(2 - xsx)^2 - (2 - xsy)^2)
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
)
budget_id = "y"
# Create an initial population with fidelity ("y") value 1
mies_init_population(oi, mu = 2, budget_id = budget_id, fidelity = 1)
oi$archive
# Re-evaluate these individuals with higher fidelity
mies_step_fidelity(oi, budget_id = budget_id, fidelity = 2)
oi$archive
# The following creates a new generation without killing the initial
# generation
offspring = data.frame(x = 0:1)
mies_evaluate_offspring(oi, offspring = offspring, budget_id = budget_id,
  fidelity = 3)
oi$archive
# Re-evaluate only individuals from last generation by setting current_gen_only
mies_step_fidelity(oi, budget_id = budget_id, fidelity = 4,
  current_gen_only = TRUE)
oi$archive
# Default: Re-evaluate all that *increase* fidelity: Only the initial
# population is re-evaluated here.
```

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```
mies_step_fidelity(oi, budget_id = budget_id, fidelity = 3)
oi$archive

# To also re-evaluate individuals with *higher* fidelity, use
# 'fidelity_monotonic = FALSE'. This does not re-evaluate the points that already have
# the requested fidelity, however.
mies_step_fidelity(oi, budget_id = budget_id, fidelity = 3, fidelity_monotonic = FALSE)
oi$archive
```

mies\_survival\_comma

Choose Survivors According to the "Mu, Lambda" ("Comma") Strategy

# **Description**

Choose survivors during a MIES iteration using the "Comma" survival strategy, i.e. selecting survivors from the latest generation only, using a Selector operator, and choosing "elites" from survivors from previous generations using a different Selector operator.

When n\_elite is greater than the number of alive individuals from previous generations, then all these individuals from previous generations survive. In this case, it is possible that more than mu - n\_elite individuals from the current generation survive. Similarly, when mu is greater than the number of alive individuals from the last generation, then all these individuals survive.

### Usage

```
mies_survival_comma(inst, mu, survival_selector, n_elite, elite_selector, ...)
```

#### **Arguments**

inst (OptimInstance)

Optimization instance to evaluate.

mu (integer(1))

Population target size, non-negative integer.

survival\_selector

(Selector)

Selector operator that selects surviving individuals depending on configuration values and objective results. When survival\_selector\$operate() is called, then objectives that are being minimized are multiplied with -1 (through mies\_get\_fitnesses), since Selectors always try to maximize fitness.

The Selector must be primed on inst\$search\_space; this *includes* the "budget" component when performing multi-fidelity optimization.

The given Selector may *not* return duplicates.

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n\_elite (integer(1))

Number of individuals to carry over from previous generations. n\_elite individuals will be selected by elite\_selector, while mu - n\_elite will be selected by survival\_selector from the most recent generation. n\_elite may be 0 (no elitism), in which case only individuals from the newest generation survive. n\_elite must be strictly smaller than mu to permit any optimization progress.

elite\_selector (Selector)

Selector operator that selects "elites", i.e. surviving individuals from previous generations, depending on configuration values and objective results. When elite\_selector\$operate() is called, then objectives that are being minimized are multiplied with -1 (through mies\_get\_fitnesses()), since Selectors always try to maximize fitness.

The Selector must be primed on inst\$search\_space; this *includes* the "budget" component when performing multi-fidelity optimization.

The given Selector may *not* return duplicates.

... (any)

Ignored, for compatibility with other  $mies\_survival\_*$  functions.

#### Value

invisible data.table: The value of inst\$archive\$data, changed in-place with eol set to the current generation for non-survivors.

#### See Also

Other mies building blocks: mies\_evaluate\_offspring(), mies\_generate\_offspring(), mies\_get\_fitnesses(), mies\_init\_population(), mies\_select\_from\_archive(), mies\_step\_fidelity(), mies\_survival\_plus()

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(</pre>
  fun = function(xs) {
    z \leftarrow \exp(-xsx^2 - xsy^2) + 2 * \exp(-(2 - xsx)^2 - (2 - xsy)^2)
    list(Obj = z)
  },
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
)
```

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```
mies_init_population(inst = oi, mu = 3)
# Usually the offspring is generated using mies_generate_offspring()
# Here shorter for demonstration purposes.
offspring = generate_design_random(oi$search_space, 3)$data
mies_evaluate_offspring(oi, offspring = offspring)
# State before: different generations of individuals. Alive individuals have
# 'eol' set to 'NA'.
oi$archive
s = sel("best")
s$prime(oi$search_space)
mies_survival_comma(oi, mu = 3, survival_selector = s,
  n_elite = 2, elite_selector = s)
# sel("best") lets only the best individuals survive.
# mies_survival_comma selects from new individuals (generation 2 in this case)
# and old individuals (all others) separately: n_elite = 2 from old,
# mu - n_elite = 1 from new.
# The surviving individuals have 'eol' set to 'NA'
oi$archive
```

mies\_survival\_plus

Choose Survivors According to the "Mu + Lambda" ("Plus") Strategy

# **Description**

Choose survivors during a MIES iteration using the "Plus" survival strategy, i.e. combining all alive individuals from the latest and from prior generations indiscriminately and choosing survivors using a survival Selector operator.

When mu is greater than the number of alive individuals, then all individuals survive.

#### Usage

```
mies_survival_plus(inst, mu, survival_selector, ...)
```

# **Arguments**

```
inst (OptimInstance)
```

Optimization instance to evaluate.

mu (integer(1))

Population target size, non-negative integer.

survival\_selector

(Selector)

Selector operator that selects surviving individuals depending on configuration values and objective results. When survival\_selector\$operate() is called, then objectives that are being minimized are multiplied with -1 (through mies\_get\_fitnesses), since Selectors always try to maximize fitness.

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```
The Selector must be primed on inst$search_space; this includes the "budget" component when performing multi-fidelity optimization.

The given Selector may not return duplicates.
```

.. (any

Ignored, for compatibility with other mies\_survival\_\* functions.

#### Value

invisible data.table: The value of inst\$archive\$data, changed in-place with eol set to the current generation for non-survivors.

#### See Also

Other mies building blocks: mies\_evaluate\_offspring(), mies\_generate\_offspring(), mies\_get\_fitnesses(), mies\_init\_population(), mies\_select\_from\_archive(), mies\_step\_fidelity(), mies\_survival\_comma()

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Define the objective to optimize
objective <- ObjectiveRFun$new(</pre>
  fun = function(xs) {
   z \leftarrow \exp(-xsx^2 - xsy^2) + 2 * \exp(-(2 - xsx)^2 - (2 - xsy)^2)
   list(Obj = z)
  domain = ps(x = p_db1(-2, 4), y = p_db1(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
)
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,
  terminator = trm("evals", n_evals = 100)
mies_init_population(inst = oi, mu = 3)
offspring = generate_design_random(oi$search_space, 2)$data
mies_evaluate_offspring(oi, offspring = offspring)
# State before: different generations of individuals. Alive individuals have
# 'eol' set to 'NA'.
oi$archive
s = sel("best")
s$prime(oi$search_space)
mies_survival_plus(oi, mu = 3, survival_selector = s)
# sel("best") lets only the three best individuals survive.
# The others have 'eol = 2' (the current generation).
oi$archive
```

```
mlr_terminators_budget
```

Terminator that Limits Total Budget Component Evaluation

# **Description**

Terminator that terminates after the sum (or similar aggregate) of a given "budget" search space component croses a threshold.

# **Dictionary**

This Terminator can be created with the short access form trm() (trms() to get a list), or through the dictionary mlr\_terminators in the following way:

```
# preferred
trm("budget")
trms("budget") # takes vector IDs, returns list of Terminators
# long form
mlr_terminators$get("budget")
```

# **Configuration Parameters**

- budget :: numeric(1)

  Total budget available, after which to stop. Not initialized and should be set to the desired value during construction.
- aggregate :: function
  Function taking a vector of values of the budget search space component, returning a scalar
  value to be compared to the budget configuration parameter. If this function returns a value
  greater or equal to budget the termination criterion is matched. Calling this function with
  NULL must return the lower bound of the budget value; percentage progress is reported as the
  progress from this lower bound to the value of budget. Initialized to sum().

# Super class

```
bbotk::Terminator -> TerminatorBudget
```

#### Methods

# **Public methods:**

- TerminatorBudget\$new()
- TerminatorBudget\$is\_terminated()
- TerminatorBudget\$clone()

Method new(): Initialize the TerminatorBudget object.

Usage:

TerminatorBudget\$new()

**Method** is\_terminated(): Is TRUE if when the termination criterion is matched, FALSE otherwise.

Usage:

TerminatorBudget\$is\_terminated(archive)

Arguments:

archive Archive Archive to check.

*Returns:* logical(1): Whether to terminate.

Method clone(): The objects of this class are cloneable with this method.

Usage:

TerminatorBudget\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

# **Examples**

```
library("bbotk")
# Evaluate until sum of budget component of evaluated configs is >= 100
trm("budget", budget = 100)
# Evaluate until sum of two to the power of budget component is >= 100
trm("budget", budget = 1024, aggregate = function(x) sum(2 ^ x))
```

mlr\_terminators\_genperfreached

Terminator That Stops When a Generation-Wise Aggregated Value Reaches a Target

# **Description**

Terminator that terminates when a value, aggregated over generations, reaches a target value.

The user-supplied fitness\_aggregator function is called whenever the archive of evaluated configurations contains a new generation. The function is supplied with the fitness values, and optionally other data, of all individuals that are alive at that point (include\_previous\_generations = FALSE) or at any point (include\_previous\_generations = TRUE). Its result is saved inside the \$data\_extra field of the Archive object. Termination is then signaled when the aggregated value meets or exceeds level.

The mies\_aggregate\_single\_generation() function is used, see the documentation there for the functioning of fitness\_aggregator. The fitness\_aggregator functions used for termination must return a scalar value or NULL, if a generation should be ignored. The value returned by fitness\_aggregator should be increasing for better performance, even if the underlying objective is being minimized.

### **Multi-Fidelity Optimization**

Multi-fidelity optimization can introduce a few edge-cases because the individuals inside the generation(s) being aggregated may have been evaluated with different fidelity values, which can give biased results.

When OptimizerMies is constructed with multi\_fidelity set to TRUE, it typically evaluates some configurations multiple times, at first with a lower fidelity, followed by an evaluation at "full" fidelity. fitness\_aggregator will only be called for generations containing entirely full-fidelity-evaluations will be aggregated.

This is achieved by caching aggregated fitness values in the \$data\_extra field of the Archive and only ever calling fitness\_aggregator for a generation that does not have a cached value. Since mies\_step\_fidelity() will count low-fidelity evaluations as part of the "previous" generation, fitness\_aggregator will not see them. Note, however that if fitness\_aggregator returns NULL, it will be called again should a second evaluation occur in the same generation, since NULL is not cached and instead treated as absent.

It is possible for fitness\_aggregator to see fitness values that were evaluated with different fidelities when using OptimizerMies, and

- 1. fidelity\_monotonic is set to TRUE and fidelity decreases (unlikely setup), or
- 2. if fidelity\_current\_gen\_only is set to FALSE (advanced usage), or
- The value returned by the fidelity configuration parameter (not fidelity\_offspring)
   changes over the course of optimization and include\_previous\_generations of TerminatorGenerationStagnation
   is set to TRUE.

(1) and (2) only need consideration in advanced scenarios, but (3) may be a common, e.g. when doing multi-fidelity optimization and stopping on reaching an overall dominated hypervolume target. In this case, it may be necessary to inspect the budget value given to fitness\_aggregator and to remove all individuals evaluated with a different than the current fidelity.

When using a custom-written optimization loop, case (1) relates to fidelity\_monotonic argument of mies\_step\_fidelity() and mies\_init\_population(), and case (2) relates to the current\_gen\_only argument of mies\_step\_fidelity() and the fidelity\_new\_individuals\_only argument of mies\_init\_population(). Case (3) relates to changing the fidelity given to mies\_step\_fidelity() if that function is used, or to changing the fidelity given to mies\_evaluate\_offspring() if mies\_step\_fidelity() is not used.

# **Dictionary**

This Terminator can be created with the short access form trm() (trms() to get a list), or through the dictionary mlr\_terminators in the following way:

```
# preferred
trm("genperfreached")
trms("genperfreached") # takes vector IDs, returns list of Terminators
# long form
mlr_terminators$get("genperfreached")
```

# **Configuration Parameters**

- fitness\_aggregator :: function
  - Aggregation function, called with information about alive individuals of each generation. This argument is passed to mies\_aggregate\_single\_generation(), see there for more details. The aggregated values returned by fitness\_aggregator should be maximized, so a larger value must be returned to indicate improvement in a generation, even if an underlying objective is being minimized. The return value must be a scalar numeric(1).
- include\_previous\_generations :: logical(1)
  Whether to aggregate over all individuals that were evaluated (TRUE), or only the individuals alive in the current generation (FALSE). If multi-fidelity optimization is being performed and individuals were re-evaluated with a different fidelity, their x\_id will be the same and only the last fidelity-reevaluation will be given to fitness\_aggregator. However, individuals from different generations may still have been evaluated with different fidelity and it may be necessary to inspect the budget value given to fitness\_aggregator if include\_previous\_generations is TRUE in a multi-fidelity-setting. See the "Multi-Fidelity Optimization" section for more.
- level :: numeric(1)

  Minimum aggregated value for which to terminate.

### Super class

bbotk::Terminator -> TerminatorGenerationPerfReached

#### Methods

### **Public methods:**

- TerminatorGenerationPerfReached\$new()
- TerminatorGenerationPerfReached\$is\_terminated()
- TerminatorGenerationPerfReached\$clone()

Method new(): Initialize the TerminatorGenerationPerfReached object.

Usage:

TerminatorGenerationPerfReached\$new()

**Method** is\_terminated(): Is TRUE if when the termination criterion is matched, FALSE otherwise.

Usage:

 $Terminator {\tt GenerationPerfReached\$is\_terminated} (archive)$ 

Arguments:

archive Archive Archive to check.

*Returns:* logical(1): Whether to terminate.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

TerminatorGenerationPerfReached\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

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# **Examples**

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Terminate when hypervolume with nadir c(0, 0, ...)
# does not improve for 3 generations by at least 0.1:
tg <- trm("genperfreached",</pre>
  fitness_aggregator = function(fitnesses) domhv(fitnesses),
  include_previous_generations = TRUE,
  level = 1
)
set.seed(1)
objective <- ObjectiveRFun$new(
  fun = function(xs) {
    list(y1 = xs$x1, y2 = xs$x2)
  },
  domain = ps(x1 = p_dbl(0, 1), x2 = p_dbl(-1, 0)),
  codomain = ps(y1 = p_dbl(0, 1, tags = "maximize"),
    y2 = p_dbl(-1, 0, tags = "minimize"))
)
oi <- OptimInstanceMultiCrit$new(objective, terminator = tg)</pre>
op <- opt("mies",
  lambda = 4, mu = 4,
  mutator = mut("gauss", sdev = 0.1),
  recombinator = rec("xounif"),
  parent_selector = sel("random"),
  survival_selector = sel("best", scl("hypervolume"))
)
op$optimize(oi)
# the observed aggregated values:
\verb|oi\$| archive\$| data\_| extra\$| Terminator Generation PerfReached|
# ... or as calculated by mies_generation_apply
mies_generation_apply(oi$archive, function(fitnesses) {
  domhv(fitnesses)
}, include_previous_generations = TRUE)
#' @export
```

# **Description**

Terminator that terminates after a given number of generations have passed in OptimizerMies.

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If OptimizerMies is started on an archive that already has evaluated configurations, these evaluations count as generation 0. If an initial, randomly sampled generation is generated by OptimizerMies, it has generation number 1. Setting generation to 1 therefore terminates after the evaluation of the initial sample, *unless* no initial sample is generated by OptimizerMies and instead found in the archive. generation set to 0 avoids any evaluation within OptimizerMies (but is ignored if no dob column is in the archive).

When doing multi-fidelity optimization, and fidelity of a configuration is increased because of a step in the fidelity schedule, or because they were sampled new and survived, then this fidelity refinement happens as part of an already started generation. This means termination at this fidelity refinement step is avoided.

# **Dictionary**

This Terminator can be created with the short access form trm() (trms() to get a list), or through the dictionary mlr\_terminators in the following way:

```
# preferred
trm("gens")
trms("gens") # takes vector IDs, returns list of Terminators
# long form
mlr_terminators$get("gens")
```

# **Configuration Parameters**

• generations :: integer(1)

Number of generations to evaluate, after which to stop. Not initialized and should be set to the desired value during construction.

#### Super class

```
bbotk::Terminator -> TerminatorGenerations
```

### Methods

### **Public methods:**

- TerminatorGenerations\$new()
- TerminatorGenerations\$is\_terminated()
- TerminatorGenerations\$clone()

**Method** new(): Initialize the TerminatorGenerations object.

Usage:

TerminatorGenerations\$new()

**Method** is\_terminated(): Is TRUE if when the termination criterion is matched, FALSE otherwise.

Usage:

TerminatorGenerations\$is\_terminated(archive)

```
Arguments:
archive Archive Archive to check.

Returns: logical(1): Whether to terminate.

Method clone(): The objects of this class are cloneable with this method.

Usage:
TerminatorGenerations$clone(deep = FALSE)

Arguments:
deep Whether to make a deep clone.
```

# **Examples**

```
library("bbotk")
trm("gens", generations = 10)
```

```
mlr_terminators_genstag
```

Terminator That Stops When a Generation-Wise Aggregated Value Stagnates

# **Description**

Terminator that terminates when a value, aggregated over generations, stagnates.

The user-supplied fitness\_aggregator function is called whenever the archive of evaluated configurations contains a new generation. The function is supplied with the fitness values, and optionally other data, of all individuals that are alive at that point (include\_previous\_generations = FALSE) or at any point (include\_previous\_generations = TRUE). Its result is saved inside the \$data\_extra field of the Archive object. Termination is then signaled when the aggregated value has stagnated, or not improved enough, for a given number of generations.

The mies\_aggregate\_single\_generation() function is used, see the documentation there for the functioning of fitness\_aggregator. The fitness\_aggregator functions used for termination must return a scalar value or NULL, if a generation should be ignored. The value returned by fitness\_aggregator should be increasing for better performance, even if the underlying objective is being minimized.

Termination is triggered in generation i when the aggregated value, returned by fitness\_aggregator, of generation i - patience is not greater than the maximum of all later generations (i - patience + 1 .. i) by at least min\_delta.

If the aggregated value for generation i - patience is not known, for example because fewer than patience + 1 generations have been evaluated so far, because fitness\_aggregator returned NULL, or because previous generations were evaluated when TerminatorGenerationStagnation was not used, then termination is not triggered, regardless of whether values are known for generations before generation i - patience.

# **Multi-Fidelity Optimization**

Multi-fidelity optimization can introduce a few edge-cases, for one because the individuals inside the generation(s) being aggregated may have been evaluated with different fidelity values, which can give biased results. Besides that, it may also be problematic that increase of fidelity could lead to "worse" aggregated results (e.g. because of reduced noise making max-aggregation worse), triggering an undesired termination.

# **Termination from fidelity changes:**

Higher fidelity values can sometimes lead to worse aggregated fitness values, which can trigger undesired termination. However, in many multi-fidelity-setups, terminating before the last fidelity increase, controlled e.g. by the fidelity configuration parameter of OptimizerMies, may be undesirable to begin with.

If the fidelity increase follows a fixed schedule based on evaluations or generations, one solution may be to use a TerminatorCombo together with a TerminatorEvals or TerminatorGenerations that prevents premature termination. Termination should happen at least patience generations after the last switch to the highest fidelity if termination from biased values because of fidelity changes should be avoided.

Otherwise it may be necessary to check whether the budget value given to fitness\_aggregator reached the desired level, and to prevent comparisons by letting fitness\_aggregator return NULL if not.

In both cases one may still have a problem with biased aggregations within an aggregated set of individuals if include\_previous\_generations is TRUE.

# **Biases within aggregated generation(s):**

When OptimizerMies is constructed with multi\_fidelity set to TRUE, it typically evaluates some configurations multiple times, at first with a lower fidelity, followed by an evaluation at "full" fidelity. fitness\_aggregator will only be called for generations containing entirely full-fidelity-evaluations will be aggregated.

This is achieved by caching aggregated fitness values in the \$data\_extra field of the Archive and only ever calling fitness\_aggregator for a generation that does not have a cached value. Since mies\_step\_fidelity() will count low-fidelity evaluations as part of the "previous" generation, fitness\_aggregator will not see them. Note, however that if fitness\_aggregator returns NULL, it will be called again should a second evaluation occur in the same generation, since NULL is not cached and instead treated as absent.

It is possible for fitness\_aggregator to see fitness values that were evaluated with different fidelities when using OptimizerMies, and

- 1. fidelity\_monotonic is set to TRUE and fidelity decreases (unlikely setup), or
- 2. if fidelity\_current\_gen\_only is set to FALSE (advanced usage), or
- 3. The value returned by the fidelity configuration parameter (not fidelity\_offspring) changes over the course of optimization and include\_previous\_generations of TerminatorGenerationStagnation is set to TRUE.

(1) and (2) only need consideration in advanced scenarios, but (3) may be a common, e.g. when doing multi-fidelity optimization and stopping on overall dominated hypervolume stagnation. In this case, it may be necessary to inspect the budget value given to fitness\_aggregator and to remove all individuals evaluated with a different than the current fidelity.

When using a custom-written optimization loop, case (1) relates to fidelity\_monotonic argument of mies\_step\_fidelity() and mies\_init\_population(), and case (2) relates to the

current\_gen\_only argument of mies\_step\_fidelity() and the fidelity\_new\_individuals\_only argument of mies\_init\_population(). Case (3) relates to changing the fidelity given to mies\_step\_fidelity() if that function is used, or to changing the fidelity given to mies\_evaluate\_offspring() if mies\_step\_fidelity() is not used.

### **Dictionary**

This Terminator can be created with the short access form trm() (trms() to get a list), or through the dictionary mlr\_terminators in the following way:

```
# preferred
trm("genstag")
trms("genstag") # takes vector IDs, returns list of Terminators
# long form
mlr_terminators$get("genstag")
```

# **Configuration Parameters**

- fitness\_aggregator :: function
  Aggregation function, called with information about alive individuals of each generation. This
  argument is passed to mies\_aggregate\_single\_generation(), see there for more details.
  The aggregated values returned by fitness\_aggregator should be maximized, so a larger
  value must be returned to indicate improvement in a generation, even if an underlying objective is being minimized. The return value must be a scalar numeric(1).
- include\_previous\_generations :: logical(1)
  Whether to aggregate over all individuals that were evaluated (TRUE), or only the individuals alive in the current generation (FALSE). If multi-fidelity optimization is being performed and individuals were re-evaluated with a different fidelity, their x\_id will be the same and only the last fidelity-reevaluation will be given to fitness\_aggregator. However, individuals from different generations may still have been evaluated with different fidelity and it may be necessary to inspect the budget value given to fitness\_aggregator if include\_previous\_generations is TRUE in a multi-fidelity-setting. See the "Multi-Fidelity Optimization" section for more.
- min\_delta:: numeric(1)
  Minimum positive change of aggregated value to count as improvement. This value may also be negative, resulting in termination only when aggregated value *decreases* by at least the given amount. However, depending on the survival setup, or on include\_previous\_generations, it is possible that aggregate values never decrease; in this case, setting min\_delta to a negative value may never trigger termination. Initialized to 0.
- patience :: integer(1)

  Number of generations with no improvement better than min\_delta after which to terminate.

  Initialized to 1.

### Super class

bbotk::Terminator -> TerminatorGenerationStagnation

# Methods

#### **Public methods:**

- TerminatorGenerationStagnation\$new()
- TerminatorGenerationStagnation\$is\_terminated()
- TerminatorGenerationStagnation\$clone()

**Method** new(): Initialize the TerminatorGenerationStagnation object.

Usage:

TerminatorGenerationStagnation\$new()

**Method** is\_terminated(): Is TRUE if when the termination criterion is matched, FALSE otherwise.

Usage:

TerminatorGenerationStagnation\$is\_terminated(archive)

Arguments:

archive Archive Archive to check.

*Returns:* logical(1): Whether to terminate.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

TerminatorGenerationStagnation\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

```
set.seed(1)
library("bbotk")
lgr::threshold("warn")
# Terminate when hypervolume with nadir `c(0, 0, ...)`
# does not improve for 3 generations by at least 0.1:
tg <- trm("genstag",</pre>
 fitness_aggregator = function(fitnesses) domhv(fitnesses),
 include_previous_generations = TRUE,
 min_delta = 0.1,
 patience = 3
)
set.seed(1)
objective <- ObjectiveRFun$new(
 fun = function(xs) {
   list(y1 = xs$x1, y2 = xs$x2)
 domain = ps(x1 = p_dbl(0, 1), x2 = p_dbl(-1, 0)),
 codomain = ps(y1 = p_dbl(0, 1, tags = "maximize"),
   y2 = p_dbl(-1, 0, tags = "minimize"))
```

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```
)
oi <- OptimInstanceMultiCrit$new(objective, terminator = tg)</pre>
op <- opt("mies",
  lambda = 4, mu = 4,
  mutator = mut("gauss", sdev = 0.1),
  recombinator = rec("xounif"),
  parent_selector = sel("random"),
  survival_selector = sel("best", scl("hypervolume"))
)
op$optimize(oi)
# the observed aggregated values:
\verb"oi$archive$data\_extra$TerminatorGenerationStagnation"
# ... or as calculated by mies_generation_apply
mies_generation_apply(oi$archive, function(fitnesses) {
  domhv(fitnesses)
}, include_previous_generations = TRUE)
#' @export
```

mut

Short Access Forms for Operators

# **Description**

These functions complement dict\_mutators, dict\_recombinators, dict\_selectors with functions in the spirit of mlr3::mlr\_sugar.

# Usage

```
mut(.key, ...)
muts(.keys, ...)
rec(.key, ...)
recs(.key, ...)
sel(.key, ...)
scl(.key, ...)
scl(.key, ...)
```

Mutator Mutator

```
ftr(.key, ...)
ftrs(.key, ...)
```

### **Arguments**

```
.key (character(1))
Key passed to the respective dictionary to retrieve the object.
... (any)
Additional arguments.
.keys (character())
Keys passed to the respective dictionary to retrieve multiple objects.
```

#### Value

- Mutator for mut()
- list of Mutator for muts()
- Recombinator for rec().
- list of Recombinator for recs().
- Selector for sel().
- list of Selector for sels().
- Scalor for scl().
- list of Scalor for scls().

# See Also

Other dictionaries: dict\_filtors, dict\_mutators, dict\_recombinators, dict\_scalors, dict\_selectors

# **Examples**

```
mut("gauss", sdev = 0.5)
rec("xounif")
sel("random")
scl("nondom")
```

Mutator

Mutator Base Class

# **Description**

Base class representing mutation operations, inheriting from MiesOperator.

Mutations get a table of individuals as input and return a table of modified individuals as output. Individuals are acted on as individuals: every line of output corresponds to the same line of input, and presence or absence of other input lines does not affect the result.

Mutation operations are performed in ES algorithms to facilitate exploration of the search space around individuals.

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# **Inheriting**

Mutator is an abstract base class and should be inherited from. Inheriting classes should implement the private \(\frac{1}{2}\) mutate() function. The user of the object calls \(\frac{1}{2}\) perate(), and the arguments are passed on to private \(\frac{1}{2}\). mutate() after checking that the operator is primed, and that the values argument conforms to the primed domain. Typically, the \(\frac{1}{2}\) initialize() function should also be overloaded, and optionally the \(\frac{1}{2}\)prime() function; they should call their super equivalents.

In many cases, it is advisable to inherit from one of the abstract subclasses, such as MutatorNumeric, or MutatorDiscrete.

# Super class

```
miesmuschel::MiesOperator -> Mutator
```

#### Methods

### **Public methods:**

- Mutator\$new()
- Mutator\$clone()

**Method** new(): Initialize base class components of the Mutator.

```
Usage:
Mutator$new(
  param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
  param_set = ps(),
  packages = character(0),
  dict_entry = NULL,
  own_param_set = quote(self$param_set)
)
Arguments:
```

param\_classes (character)

Classes of parameters that the operator can handle. May contain any of "ParamLg1", "ParamInt", "ParamDb1", "ParamFct". Default is all of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators,

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dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

Mutator\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

#### See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, Selector, SelectorScalar

Other mutators: MutatorDiscrete, MutatorNumeric, OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_erase, dict\_mutators\_gauss, dict\_mutators\_maybe, dict\_mutators\_null, dict\_mutators\_proxy, dict\_mutators\_sequential, dict\_mutators\_unif

MutatorDiscrete

Discrete Mutator Base Class

# **Description**

Base class for mutation operations on discrete individuals, inheriting from Mutator.

MutatorDiscrete operators perform mutation on discrete (logical and factor valued) individuals. Inheriting operators implement the private \$.mutate\_discrete() function that is called once for each individual and is given a character vector.

# **Inheriting**

MutatorDiscrete is an abstract base class and should be inherited from. Inheriting classes should implement the private \( \). mutate\_discrete() function. During \( \)operate(), the \( \). mutate\_discrete() function is called once for each individual, with the parameters values (the individual as a single character vector), and levels (a list of character containing the possible values for each element of values). Typically, \( \)initialize() should also be overloaded.

# Super classes

miesmuschel::MiesOperator -> miesmuschel::Mutator -> MutatorDiscrete

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### Methods

#### **Public methods:**

- MutatorDiscrete\$new()
- MutatorDiscrete\$clone()

**Method** new(): Initialize base class components of the MutatorNumeric.

```
Usage:
```

```
MutatorDiscrete$new(
  param_classes = c("ParamLgl", "ParamFct"),
  param_set = ps(),
  packages = character(0),
  dict_entry = NULL,
  own_param_set = quote(self$param_set)
)
```

Arguments:

param\_classes (character)

Classes of parameters that the operator can handle. May contain any of "ParamLg1", "ParamFct". Default is both of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

```
MutatorDiscrete$clone(deep = FALSE)
```

Arguments:

deep Whether to make a deep clone.

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#### See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, Selector, SelectorScalar

Other mutators: Mutator, MutatorNumeric, OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_erase, dict\_mutators\_gauss, dict\_mutators\_maybe, dict\_mutators\_null, dict\_mutators\_proxy, dict\_mutators\_sequential, dict\_mutators\_unif

MutatorNumeric

Numeric Mutator Base Class

# **Description**

Base class for mutation operations on numeric and integer valued individuals, inheriting from Mutator.

MutatorNumeric operators perform mutation on numeric (integer and real valued) individuals. Inheriting operators implement the private \$.mutate\_numeric() function that is called once for each individual and is given a numeric vector.

# **Inheriting**

MutatorNumeric is an abstract base class and should be inherited from. Inheriting classes should implement the private \( \). mutate\_numeric() function. During \( \)operate(), the \( \). mutate\_numeric() function is called once for each individual, with the parameters values (the individual as a single numeric vector), lowers and uppers (numeric vectors, the lower and upper bounds for each component of values). Typically, \( \)initialize() should also be overloaded.

MutatorNumerics that perform real-valued operations, such as e.g. MutatorGauss, operate on integers by widening the lower and upper bounds of integer components by 0.5, applying their operation, and rounding resulting values to the nearest integer (while always staying inside bounds).

# Super classes

```
miesmuschel::MiesOperator-> miesmuschel::Mutator-> MutatorNumeric
```

# Methods

#### **Public methods:**

- MutatorNumeric\$new()
- MutatorNumeric\$clone()

**Method** new(): Initialize base class components of the MutatorNumeric.

Usage:

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```
MutatorNumeric$new(
    param_classes = c("ParamInt", "ParamDbl"),
    param_set = ps(),
    packages = character(0),
    dict_entry = NULL,
    own_param_set = quote(self$param_set)
 )
 Arguments:
 param_classes (character)
     Classes of parameters that the operator can handle. May contain any of "ParamInt",
     "ParamDb1". Default is both of them.
     The $param_classes field will reflect this value.
 param_set (ParamSet | list of expression)
     Strategy parameters of the operator. This should be created by the subclass and given to
     super$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet
     directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate
     to ParamSets, possibly referencing self and private. These ParamSet are then combined
     using a ParamSetCollection. Default is the empty ParamSet.
     The $param_set field will reflect this value.
 packages (character) Packages that need to be loaded for the operator to function. This
     should be declared so these packages can be loaded when operators run on parallel in-
     stances. Default is character(0).
     The $packages field will reflect this values.
 dict_entry (character(1) | NULL)
     Key of the class inside the Dictionary (usually one of dict_mutators, dict_recombinators,
     dict_selectors), where it can be retrieved using a short access function. May be NULL if
     the operator is not entered in a dictionary.
     The $dict_entry field will reflect this value.
 own_param_set (language)
     An expression that evaluates to a ParamSet indicating the configuration parameters that are
     entirely owned by this operator class (and not proxied from a construction argument object).
     This should be quote(self$param_set) (the default) when the param_set argument is not
     a list of expressions.
Method clone(): The objects of this class are cloneable with this method.
 Usage:
 MutatorNumeric$clone(deep = FALSE)
```

# See Also

Arguments:

deep Whether to make a deep clone.

 $Other \ base \ classes: Filtor, Filtor Surrogate, Mies Operator, Mutator, Mutator Discrete, Operator Combination, Recombinator, Recombinator Pair, Scalor, Selector, Selector Scalar$ 

Other mutators: Mutator, MutatorDiscrete, OperatorCombination, dict\_mutators\_cmpmaybe, dict\_mutators\_erase, dict\_mutators\_gauss, dict\_mutators\_maybe, dict\_mutators\_null, dict\_mutators\_proxy, dict\_mutators\_sequential, dict\_mutators\_unif

OperatorCombination Self-Adaptive Operator Combinations

# **Description**

Combines multiple operators and makes operator-configuration parameters self-adaptive.

The OperatorCombination operators combine operators for different subspaces of individuals by wraping other MiesOperators given during construction. Different MiesOperators are assigned to different components or sets of components and operate on them independently of the rest of the components or the other operators. An operator can be assigned to a single component by giving it in operators with the name of the component, or to multiple components by giving it in operators with the name of a *group*. Groups are created by the groups argument, but several default groups that catch components by type exist.

#### **Details**

Operators can be made *self-adaptive* by coupling their configuration parameter values to values in individuals. This is done by giving functions in adaptions; these functions are executed for each individual before an operator is applied, and the result given to a named operator configuration parameter.

OperatorCombination is the base class from which MutatorCombination and RecombinatorCombination inherit. The latter two are to be used for Mutator and Recombinator objects, respectively.

Besides groups created with the groups construction argument, there are special groups that all unnamed operators fall into based on their Domain class: "ParamInt", "ParamDbl", "ParamFct", and "ParamLgl". A component of an individual that is not named directly in operators or made part of a group in groups is automatically in one of these special groups. There is furthermore a special catch-all group "ParamAny", which catches all components that are are not operated directly, not in a group, and not in another special group that is itself named directly or in a group. I.e., all components that would otherwise have no assigned operation.

RecombinatorCombination can only combine operators where \$n\_indivs\_in and \$n\_indivs\_out can be combined. This is currently supported either when \$n\_indivs\_in and \$n\_indivs\_out for each operator are the same (but \$n\_indivs\_in may be unequal \$n\_indivs\_out in each of them); or when \$n\_indivs\_in is equal to \$n\_indivs\_out for each operator and the set of all \$n\_indivs\_in that occur contains 1 and one more integer. \$n\_indivs\_in and \$n\_indivs\_out for the resulting RecombinatorCombination operator will be set the maximum of occuring \$n\_indivs\_in and \$n\_indivs\_out, respectively.

# **Supported Operand Types**

Supported Domain classes are calculated based on the supported classes of the wrapped operators. They are frequently just the set union of supported classes, unless inference can be drawn from type-specific groups that an operator is assigned to. If e.g. an operator that supports p\_dbl and p\_int is assigned to group "ParamInt", and an operator that supports p\_lgl is assigned to component "a", then the result will support p\_lgl and p\_int only.

### **Configuration Parameters**

The OperatorCombination has the configuration parameters of all encapsulated MiesOperators, minus the configuration parameters that are named in the adaptions. Configuration parameter names are prefixed with the name of the MiesOperator in the operators list.

# **Dictionary**

This Mutator can be created with the short access form mut() (muts() to get a list), or through the the dictionary dict\_mutators in the following way:

```
# preferred:
mut("combine", <operators>, ...)
muts("combine", <operators>, ...) # takes vector IDs, returns list of Mutators
# long form:
dict_mutators$get("combine", <operators>, ...)

This Recombinator can be created with the short access form rec() (recs() to get a list), or through the dictionary dict_recombinators in the following way:

# preferred:
rec("combine", <operators>, ...)
recs("combine", <operators>, ...) # takes vector IDs, returns list of Recombinators
# long form:
dict_recombinators$get("combine", <operators>, ...)
```

### Super class

```
miesmuschel::MiesOperator -> OperatorCombination
```

### **Active bindings**

```
operators (named list of MiesOperator)
```

List of operators to apply to components of individuals, as set during construction. Read-only.

groups (named list of character)

List of groups that operators can act on, as set during construction. Read-only.

```
adaptions (named list of function)
```

List of functions used for self-adaption of operators, as set during construction. Read-only.

```
binary_fct_as_logical (logical(1))
```

Whether to treat binary p\_fct components of ParamSets as p\_lgl with respect to the special groups "ParamLgl" and "ParamFct", as set during construction. Read-only.

```
on_type_not_present (character(1))
```

Action to perform during \$prime() when an operator is assigned to a type special group but there is no component available that falls in this group. See the construction argument. Can be changed during the object's lifetime.

```
on_name_not_present (character(1))
```

Action to perform during \$prime() when an operator is assigned to a specifically named component, but the component is not present. See the construction argument. Can be changed during the object's lifetime.

#### Methods

#### **Public methods:**

- OperatorCombination\$new()
- OperatorCombination\$prime()
- OperatorCombination\$clone()

**Method** new(): Initialize the OperatorCombination object.

```
Usage:
OperatorCombination$new(
  operators,
  groups = list(),
  adaptions = list(),
  binary_fct_as_logical = FALSE,
  on_type_not_present = "warn",
  on_name_not_present = "stop",
  granularity = 1,
  dict_entry = NULL,
  dict_shortaccess = NULL
)
Arguments:
operators (named list of MiesOperator)
```

List of operators to apply to components of individuals. Names are either names of individual components, or group names which are either as defined through groups or special groups. Individual components can only be member of either a (non-special) group or named in operators, so a name that occurs in operators may not be a member of a group as defined in groups.

The \$operators field will reflect this value.

```
groups (named list of character)
```

List of groups that operators can act on. Names of this list define new groups. The content of each list element contains the names of components or special groups (a Domain subclass name or "ParamAny") to subsume under the group. Individual components can only be member of either a (non-special) group or named in operators, so a name that occurs in operators may not be a member of a group as defined in groups. The default is the empty list.

The \$groups field will reflect this value.

```
adaptions (named list of function)
```

List of functions used for self-adaption of operators. The names of the list must be names of configuration parameters of wrapped operators, prefixed with the corresponding name in the operators list. This is the same name as the configuration parameter would otherwise have if exposed by the OperatorCombination object. The values in the list must be functions that receive a single input, the individual or individuals being operated on, as a data.table.

It must return a value that is then assigned to the configuration parameter of the operator to which it pertains. Note that MutatorCombination adaption functions are always called with a data.table containing a single row, while RecombinatorCombination adaption functions are called with data.tables with multiple rows according to \$n\_indivs\_in. In both cases, the return value must be a scalar. The default is the empty list.

The \$adaption field will reflect this value.

```
binary_fct_as_logical (logical(1))
```

Whether to treat binary p\_fct components of ParamSets as p\_lgl with respect to the special groups "ParamLgl" and "ParamFct". This does *not* perform any conversion, so a MiesOperator assigned to the "ParamLgl" special group when binary\_fct\_as\_logical is TRUE and there are binary p\_fcts present will receive a factorial value and must also support p\_fct in this case. This is checked during \$prime(), but not during construction. Default is FALSE.

The \$binary\_fct\_as\_logical field will reflect this value.

```
on_type_not_present (character(1))
```

Action to perform during \$prime() when an operator is assigned to a type special group but there is no component available that falls in this group, either because no components of the respective type are present, or because all these components are also directly named in operators or in groups. One of "quiet" (do nothing), "warn" (give warning, default), or "stop" (generate an error).

The writable <code>\$on\_type\_not\_present</code> field will reflect this value.

```
on_name_not_present (character(1))
```

Action to perform during \$prime() when an operator is assigned to a specifically named component, but the component is not present. One of "quiet" (do nothing), "warn" (give warning), or "stop" (generate an error, default).

The writable <code>\$on\_name\_not\_present</code> field will reflect this value.

```
granularity (integer(1))
```

At what granularity to query adaptions for sets of individuals. Functions in adaptions are always called once per granularity individuals in input values, and the function argument in these calls will then have granularity number of rows. This is used internally, it is set to 1 for MutatorCombination, and to  $n_i$  for RecombinatorCombination.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
dict_shortaccess (character(1) | NULL)
```

Name of the Dictionary short access function in which the operator is registered. This is used to inform the user about how to construct a given object. Should ordinarily be one of "mut", "rec", "sel".

The \$dict\_shortaccess field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operators given to operators during construction. Priming of wrapped operators happens according to component assignments to wrapped operators.

Usage:

```
OperatorCombination$prime(param_set)
```

```
Arguments:
       param_set (ParamSet)
          Passed to MiesOperator$prime().
       Returns: invisible self.
     Method clone(): The objects of this class are cloneable with this method.
       OperatorCombination$clone(deep = FALSE)
      Arguments:
       deep Whether to make a deep clone.
Super classes
    miesmuschel::MiesOperator->miesmuschel::OperatorCombination->MutatorCombination
Methods
     Public methods:
       • MutatorCombination$new()
       • MutatorCombination$clone()
     Method new(): Initialize the MutatorCombination object.
       Usage:
      MutatorCombination$new(
         operators = list(),
         groups = list(),
         adaptions = list(),
         binary_fct_as_logical = FALSE,
         on_type_not_present = "warn",
         on_name_not_present = "stop"
       )
      Arguments:
       operators see above.
       groups see above.
       adaptions see above.
       binary_fct_as_logical see above.
      on_type_not_present see above.
       on_name_not_present see above.
     Method clone(): The objects of this class are cloneable with this method.
       Usage:
      MutatorCombination$clone(deep = FALSE)
      Arguments:
       deep Whether to make a deep clone.
```

# Super classes

```
miesmuschel::MiesOperator->miesmuschel::OperatorCombination->RecombinatorCombination
```

# **Active bindings**

```
n_indivs_in (integer(1))
    Number of individuals to consider at the same time. When operating, the number of input individuals must be divisible by this number.
n_indivs_out (integer(1))
```

Number of individuals produced for each group of \$n\_indivs\_in individuals.

## Methods

# **Public methods:**

- RecombinatorCombination\$new()
- RecombinatorCombination\$clone()

Method new(): Initialize the RecombinatorCombination object.

```
Usage:
 RecombinatorCombination$new(
   operators = list(),
   groups = list(),
   adaptions = list(),
   binary_fct_as_logical = FALSE,
   on_type_not_present = "warn",
   on_name_not_present = "stop"
 )
 Arguments:
 operators see above.
 groups see above.
 adaptions see above.
 binary_fct_as_logical see above.
 on_type_not_present see above.
 on_name_not_present see above.
Method clone(): The objects of this class are cloneable with this method.
 Usage:
 RecombinatorCombination$clone(deep = FALSE)
 Arguments:
 deep Whether to make a deep clone.
```

#### See Also

```
Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, Recombinator, RecombinatorPair, Scalor, Selector, SelectorScalar
```

```
Other mutators: Mutator, MutatorDiscrete, MutatorNumeric, dict_mutators_cmpmaybe, dict_mutators_erase, dict_mutators_gauss, dict_mutators_maybe, dict_mutators_null, dict_mutators_proxy, dict_mutators_sequential, dict_mutators_unif
```

```
Other mutator wrappers: dict_mutators_cmpmaybe, dict_mutators_maybe, dict_mutators_proxy, dict_mutators_sequential
```

Other recombinators: Recombinator, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convedict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_null, dict\_recombinators\_prox\_dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swap, dict\_recombinators\_xonadict\_recombinators\_xounif

Other recombinator wrappers: dict\_recombinators\_cmpmaybe, dict\_recombinators\_maybe, dict\_recombinators\_proxy, dict\_recombinators\_sequential

# **Examples**

```
set.seed(1)
data = data.frame(x = 0, y = 0, a = TRUE, b = "a",
  stringsAsFactors = FALSE) # necessary for R <= 3.6
p = ps(x = p_dbl(-1, 1), y = p_dbl(-1, 1), a = p_lgl(), b = p_fct(c("a", "b")))
# Demo operators:
m0 = mut("null") # no mutation
msmall = mut("gauss", sdev = 0.1) # mutates to small value around 0
mbig = mut("gauss", sdev = 100) # likely mutates to +1 or -1
mflip = mut("unif", can_mutate_to_same = FALSE) # flips TRUE/"a" to FALSE/"b"
# original:
data
# operators by name
op = mut("combine", operators = list(x = msmall, y = mbig, a = m0, b = mflip))
op$prime(p)
op$operate(data)
# operators by type
op = mut("combine",
  operators = list(ParamDbl = msmall, ParamLgl = m0, ParamFct = mflip)
)
op$prime(p)
op$operate(data)
# the binary ParamFct 'b' counts as 'ParamLgl' when
# 'binary_fct_as_logical' is set to 'TRUE'.
op = mut("combine",
  operators = list(ParamDbl = msmall, ParamLgl = m0),
  binary_fct_as_logical = TRUE
op$prime(p)
```

```
op$operate(data)
# operators by type; groups can be mixed types
op = mut("combine",
 operators = list(group1 = m0, group2 = msmall, group3 = mflip),
  groups = list(group1 = c("a", "x"), group2 = "y", group3 = "b")
)
op$prime(p)
op$operate(data)
# Special type-groups can be used inside groups.
op = mut("combine",
  operators = list(group1 = m0, b = mflip),
  groups = list(group1 = c("ParamDbl", "a"))
op$prime(p)
op$operate(data)
# Type-groups only capture all parameters that were not caught by name.
# The special 'ParamAny' group captures all that is left.
op = mut("combine",
  operators = list(ParamAny = m0, ParamDbl = msmall, x = mbig)
)
op$prime(p)
op$operate(data)
# Configuration parameters are named by names in the 'operators' list.
op$param_set
###
# Self-adaption:
# In this example, the 'ParamDbl''s operation is changed depending on the
# value of 'b'.
op = mut("combine",
  operators = list(ParamAny = m0, ParamLgl = mflip, ParamDbl = msmall),
  adaptions = list(ParamDbl.sdev = function(x) if (x\$a) 100 else 0.1)
op$prime(p)
data2 = data[c(1, 1, 1, 1), ]
data2$a = c(TRUE, TRUE, FALSE, FALSE)
data2
# Note the value of x$a gets used line-wise, and that it is used *before*
# being flipped here. So the first two lines get large mutations, even though
# they have 'a' 'FALSE' after the operation.
op$operate(data2)
```

OptimInstanceMultiCrit

OptimInstanceMultiCrit Class

# **Description**

bbotk's OptimInstanceMultiCrit class. Re-exported since bbotk will change the name.

# Super classes

```
bbotk::OptimInstance->bbotk::OptimInstanceBatch->bbotk::OptimInstanceBatchMultiCrit
-> OptimInstanceMultiCrit
```

# Methods

#### **Public methods:**

• OptimInstanceMultiCrit\$clone()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

OptimInstanceMultiCrit\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

OptimInstanceSingleCrit

OptimInstanceSingleCrit Class

# Description

bbotk's OptimInstanceSingleCrit class. Re-exported since bbotk will change the name.

### Super classes

```
bbotk::OptimInstance->bbotk::OptimInstanceBatch->bbotk::OptimInstanceBatchSingleCrit
-> OptimInstanceSingleCrit
```

# Methods

#### **Public methods:**

• OptimInstanceSingleCrit\$clone()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

OptimInstanceSingleCrit\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

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Optimizer

Optimizer Class

### **Description**

bbotk's Optimizer class. Re-exported since bbotk will change the name.

# Super classes

```
bbotk::Optimizer -> bbotk::OptimizerBatch -> Optimizer
```

#### Methods

#### **Public methods:**

• Optimizer\$clone()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

Optimizer\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

OptimizerMies

Mixed Integer Evolution Strategies Optimizer

### **Description**

Perform optimization using evolution strategies. OptimizerMies and TunerMies implement a standard ES optimization algorithm, performing initialization first, followed by a loop of performance evaluation, survival selection, parent selection, mutation, and recombination to generate new individuals to be evaluated. Currently, two different survival modes ("comma" and "plus") are supported. Multi-fidelity optimization, similar to the "rolling-tide" algorithm described in Fieldsend (2014), is supported. The modular design and reliance on MiesOperator objects to perform central parts of the optimization algorithm makes this Optimizer highly flexible and configurable. In combination with OperatorCombination mutators and recombinators, an algorithm as presented in Li (2013) can easily be implemented.

OptimizerMies implements a standard evolution strategies loop:

- 1. Prime operators, using mies\_prime\_operators()
- 2. Initialize and evaluate population, using mies\_init\_population()
- 3. Generate offspring by selecting parents, recombining and mutating them, using mies\_generate\_offspring()
- 4. Evaluate performance, using mies\_evaluate\_offspring()

5. Select survivors, using either mies\_survival\_plus() or mies\_survival\_comma(), depending on the survival\_strategy configuration parameter

- 6. Optionally, evaluate survivors with higher fidelity if the multi-fidelity functionality is being used
- 7. Jump to 3.

# **Terminating**

As with all optimizers, Terminators are used to end optimization after a specific number of evaluations were performed, time elapsed, or other conditions are satisfied. Of particular interest is TerminatorGenerations, which terminates after a number of generations were evaluated in OptimizerMies. The initial population counts as generation 1, its offspring as generation 2 etc.; fidelity refinements (step 6. in the algorithm description above) are always included in their generation, TerminatorGenerations avoids terminating right before they are evaluated. Other terminators may, however, end the optimization process at any time.

# **Multi-Fidelity**

miesmuschel provides a simple multi-fidelity optimization mechanism that allows both the refinement of fidelity as the optimization progresses, as well as fidelity refinement within each generation. When multi\_fidelity is TRUE, then one search space component of the OptimInstance must have the "budget" tag, which is then optimized as the "budget" component. This means that the value of this component is determined by the fidelity/fidelity\_offspring parameters, which are functions that get called whenever individuals get evaluated. The fidelity function is evaluated before step 2 and before every occurrence of step 6 in the algorithm, it returns the value of the budget search space component that all individuals that survive the current generation should be evaluated with. fidelity\_offspring is called before step 4 and determines the fidelity that newly sampled offspring individuals should be evaluated with; it may be desirable to set this to a lower value than fidelity to save budget when preliminarily evaluating newly sampled individuals that may or may not perform well compared to already sampled individuals. Individuals that survive the generation and are not removed in step 5 will be re-evaluated with the fidelity-value in step 6 before the next loop iteration.

fidelity and fidelity\_offspring must have arguments inst, budget\_id, last\_fidelity and last\_fidelity\_offspring. inst is the OptimInstance bein optimized, the functions can use it to determine the progress of the optimization, e.g. query the current generation with mies\_generation. budget\_id identifies the search space component being used as budget parameter. last\_fidelity and last\_fidelity\_offspring contain the last values given by fidelity/fidelity\_offspring. Should the offspring-fidelity (as returned by fidelity\_offspring always be the same as the parent generation fidelity (as returned by fidelity), for example, then fidelity\_offspring can be set to a function that just returns last\_fidelity; this is actually the behaviour that fidelity\_offspring is initialized with.

OptimizerMies avoids re-evaluating individuals if the fidelity parameter does not change. This means that setting fidelity and fidelity\_offspring to the same value avoids re-evaluating individuals in step 6. When fidelity\_monotonic is TRUE, re-evaluation is also avoided should the desired fidelity parameter value decrease. When fidelity\_current\_gen\_only is TRUE, then step 6 only re-evaluates individuals that were created in the current generation (in the previous step 4) and sets the fidelity for individuals that are created in step 6, but it does not re-evaluate individuals

that survived from earlier generations or were already in the OptimInstance when optimization started; it is recommended to leave this value at TRUE which it is initialized with.

### **Additional Components**

The search space over which the optimization is performed is fundamentally tied to the Objective, and therefore to the OptimInstance given to OptimizerMies\$optimize(). However, some advanced Evolution Strategy based algorithms may need to make use of additional search space components that are independent of the particular objective. An example is self-adaption as implemented in OperatorCombination, where one or several components can be used to adjust operator behaviour. These additional components are supplied to the optimizer through the additional\_component\_sampler configuration parameter, which takes a Sampler object. This object both has an associated ParamSet which represents the additional components that are present, and it provides a method for generating the initial values of these components. The search space that is seen by the MiesOperators is then the union of the OptimInstance's ParamSet, and the Sampler's ParamSet.

# **Configuration Parameters**

OptimizerMies has the configuration parameters of the mutator, recombinator, parent\_selector, survival\_selector, init\_selector, and, if given, elite\_selector operator given during construction, and prefixed according to the name of the argument (mutator's configuration parameters are prefixed "mutator." etc.). When using the construction arguments' default values, they are all "proxy" operators: MutatorProxy, RecombinatorProxy and SelectorProxy. This means that the respective configuration parameters become mutator.operation, recombinator.operation etc., so the operators themselves can be set via configuration parameters in this case.

Further configuration parameters are:

- lambda:: integer(1)
  Offspring size: Number of individuals that are created and evaluated anew for each generation.
  This is equivalent to the lambda parameter of mies\_generate\_offspring(), see there for more information. Must be set by the user.
- mu :: integer(1)
   Population size: Number of individuals that are sampled in the beginning, and which are selected with each survival step. This is equivalent to the mu parameter of mies\_init\_population(), see there for more information. Must be set by the user.
- survival\_strategy :: character(1)
  May be "plus", or, if the elite\_selector construction argument is not NULL, "comma":
  Choose whether mies\_survival\_plus() or mies\_survival\_comma() is used for survival selection. Initialized to "plus".
- n\_elite::integer(1)
  Only if the elite\_selector construction argument is not NULL, and only valid when survival\_strategy is "comma": Number of elites, i.e. individuals from the parent generation, to keep during "Comma" survival. This is equivalent to the n\_elite parameter of mies\_survival\_comma(), see there for more information.
- initializer:: function
  Function that generates the initial population as a Design object, with arguments param\_set
  and n, functioning like paradox::generate\_design\_random or paradox::generate\_design\_lhs.

This is equivalent to the initializer parameter of mies\_init\_population(), see there for more information. Initialized to generate\_design\_random().

• additional\_component\_sampler:: Sampler | NULL Additional components that may be part of individuals as seen by mutation, recombination, and selection MiesOperators, but that are not part of the search space of the OptimInstance being optimized. This is equivalent to the additional\_component\_sampler parameter of mies\_init\_population(), see there for more information. Initialized to NULL (no additional components).

# • fidelity :: function

Only if the multi\_fidelity construction argument is TRUE: Function that determines the value of the "budget" component of surviving individuals being evaluated when doing multi-fidelity optimization. It must have arguments named inst, budget\_id, last\_fidelity and last\_fidelity\_offspring, see the "Multi-Fidelity"-section for more details. Its return value is given to mies\_init\_population() and mies\_step\_fidelity(). When this configuration parameter is present (i.e. multi\_fidelity is TRUE), then it is initialized to a function returning the value 1.

• fidelity\_offspring :: function

Only if the multi\_fidelity construction argument is TRUE: Function that determines the value of the "budget" component of newly sampled offspring individuals being evaluated when doing multi-fidelity optimization. It must have arguments named inst, budget\_id, last\_fidelity and last\_fidelity\_offspring, see the "Multi-Fidelity"-section for more details. Its return value is given to mies\_evaluate\_offspring(). When this configuration parameter is present (i.e. multi\_fidelity is TRUE), then it is initialized to a function returning the value of last\_fidelity, i.e. the value returned by the last call to the fidelity configuration parameter. This is the recommended value when fidelity should not change within a generation, since this means that survivor selection is performed with individuals that were evaluated with the same fidelity (at least if fidelity\_current\_gen\_only is also set to FALSE).

• fidelity\_current\_gen\_only :: logical(1)
Only if the multi\_fidelity construction argument is TRUE: When doing fidelity refinement in mies\_step\_fidelity(), whether to refine all individuals with different budget component, or

mies\_step\_fidelity(), whether to refine all individuals with different budget component, or
only individuals created in the current generation. This is equivalent to the current\_gen\_only
parameter of mies\_step\_fidelity(), see there for more information.

When this configuration parameter is present (i.e. multi\_fidelity is TRUE), then it is initialized to FALSE, the recommended value.

• fidelity\_monotonic :: logical(1)

Only if the multi\_fidelity construction argument is TRUE: Whether to only do fidelity refinement in mies\_step\_fidelity() for individuals for which the budget component value would *increase*. This is equivalent to the monotonic parameter of mies\_step\_fidelity(), see there for more information.

When this configuration parameter is present (i.e. multi\_fidelity is TRUE), then it is initialized to TRUE. When optimization is performed on problems that have a categorical "budget" parameter, then this value should be set to FALSE.

# Super classes

# **Active bindings**

```
mutator (Mutator)
     Mutation operation to perform during mies_generate_offspring().
recombinator (Recombinator)
     Recombination operation to perform during mies_generate_offspring().
parent_selector (Selector)
    Parent selection operation to perform during mies_generate_offspring().
survival_selector (Selector)
     Survival selection operation to use in mies_survival_plus() or mies_survival_comma().
elite_selector (Selector | NULL)
     Elite selector used in mies_survival_comma().
init_selector (Selector)
     Selection operation to use when there are more than mu individuals present at the beginning of
     the optimization.
param_set (ParamSet)
    Configuration parameters of the optimization algorithm.
```

#### Methods

#### **Public methods:**

- OptimizerMies\$new()
- OptimizerMies\$clone()

**Method** new(): Initialize the OptimizerMies object.

```
Usage:
OptimizerMies$new(
  mutator = MutatorProxy$new(),
  recombinator = RecombinatorProxy$new(),
  parent_selector = SelectorProxy$new(),
  survival_selector = SelectorProxy$new(),
  elite_selector = NULL,
  init_selector = survival_selector,
  multi_fidelity = FALSE
)
Arguments:
mutator (Mutator)
```

Mutation operation to perform during mies\_generate\_offspring(), see there for more information. Default is MutatorProxy, which exposes the operation as a configuration parameter of the optimizer itself.

The \$mutator field will reflect this value.

```
recombinator (Recombinator)
```

Recombination operation to perform during mies\_generate\_offspring(), see there for more information. Default is RecombinatorProxy, which exposes the operation as a configuration parameter of the optimizer itself. Note: The default RecombinatorProxy has \$n\_indivs\_in set to 2, so to use recombination operations with more than two inputs, or to

use population size of 1, it may be necessary to construct this argument explicitly. The \$recombinator field will reflect this value.

```
parent_selector (Selector)
```

Parent selection operation to perform during mies\_generate\_offspring(), see there for more information. Default is SelectorProxy, which exposes the operation as a configuration parameter of the optimizer itself.

The \$parent\_selector field will reflect this value.

```
survival_selector (Selector)
```

Survival selection operation to use in mies\_survival\_plus() or mies\_survival\_comma() (depending on the survival\_strategy configuration parameter), see there for more information. Default is SelectorProxy, which exposes the operation as a configuration parameter of the optimizer itself.

The \$survival\_selector field will reflect this value.

```
elite selector (Selector | NULL)
```

Elite selector used in mies\_survival\_comma(), see there for more information. "Comma" selection is only available when this argument is not NULL. Default NULL.

The \$elite\_selector field will reflect this value.

```
init_selector (Selector)
```

Survival selection operation to give to the survival\_selector argument of mies\_init\_population(); it is used if the OptimInstance being optimized already contains more (alive) individuals than mu. Default is the value given to survival\_selector. The \$init\_selector field will reflect this value.

```
multi_fidelity (logical(1))
```

Whether to enable multi-fidelity optimization. When this is TRUE, then the <code>OptimInstance</code> being optimized must contain a <code>Domain</code> tagged "budget", which is then used as the "budget" search space component, determined by fidelity and fidelity\_offspring instead of by the <code>MiesOperators</code> themselves. For multi-fidelity optimization, the fidelity, fidelity\_offspring, fidelity\_current\_gen\_only, and fidelity\_monotonic configuration parameters must be given to determine multi-fidelity behaviour. (While the initial values for most of these are probably good for most cases in which more budget implies higher fidelity, at least the fidelity configuration parameter should be adjusted in most cases). Default is FALSE.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
```

OptimizerMies\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

### Super classes

```
mlr3tuning::Tuner-> mlr3tuning::TunerBatch-> mlr3tuning::TunerBatchFromOptimizerBatch
-> TunerMies
```

#### Methods

# **Public methods:**

• TunerMies\$new()

### • TunerMies\$clone()

```
Method new(): Initialize the TunerMies object.
 Usage:
 TunerMies$new(
   mutator = MutatorProxy$new(),
   recombinator = RecombinatorProxy$new(),
   parent_selector = SelectorProxy$new(),
   survival_selector = SelectorProxy$new(),
   elite_selector = NULL,
   init_selector = survival_selector,
   multi_fidelity = FALSE
 Arguments:
 mutator (Mutator)
 recombinator (Recombinator)
 parent_selector (Selector)
 survival_selector (Selector)
 elite_selector (Selector | NULL)
 init_selector (Selector)
 multi_fidelity (logical(1))
Method clone(): The objects of this class are cloneable with this method.
 Usage:
 TunerMies$clone(deep = FALSE)
 Arguments:
 deep Whether to make a deep clone.
```

# References

Fieldsend, E J, Everson, M R (2014). "The rolling tide evolutionary algorithm: A multiobjective optimizer for noisy optimization problems." *IEEE Transactions on Evolutionary Computation*, **19**(1), 103–117.

Li, Rui, Emmerich, TM M, Eggermont, Jeroen, B"ack, Thomas, Sch"utz, Martin, Dijkstra, Jouke, Reiber, HC J (2013). "Mixed integer evolution strategies for parameter optimization." *Evolutionary computation*, **21**(1), 29–64.

# **Examples**

```
lgr::threshold("warn")

op.m <- mut("gauss", sdev = 0.1)
op.r <- rec("xounif", p = .3)
op.parent <- sel("random")
op.survival <- sel("best")
######</pre>
```

```
# Optimizing a Function
library("bbotk")
# Define the objective to optimize
objective <- ObjectiveRFun$new(
  fun = function(xs) {
   z \leftarrow exp(-xs$x^2 - xs$y^2) + 2 * exp(-(2 - xs$x)^2 - (2 - xs$y)^2)
   list(Obj = z)
  domain = ps(x = p_dbl(-2, 4), y = p_dbl(-2, 4)),
  codomain = ps(Obj = p_dbl(tags = "maximize"))
# Get a new OptimInstance
oi <- OptimInstanceSingleCrit$new(objective,</pre>
  terminator = trm("evals", n_evals = 100)
)
# Create OptimizerMies object
mies_opt <- opt("mies", mutator = op.m, recombinator = op.r,</pre>
  parent_selector = op.parent, survival_selector = op.survival,
  mu = 10, lambda = 5)
# mies_opt$optimize performs MIES optimization and returns the optimum
mies_opt$optimize(oi)
#####
# Optimizing a Machine Learning Method
# Note that this is a short example, aiming at clarity and short runtime.
# The settings are not optimal for hyperparameter tuning. The resampling
# in particular should not be "holdout" for small datasets where this gives
# a very noisy estimate of performance.
library("mlr3")
library("mlr3tuning")
# The Learner to optimize
learner = lrn("classif.rpart")
# The hyperparameters to optimize
learner$param_set$values[c("cp", "maxdepth")] = list(to_tune())
# Get a TuningInstance
ti = TuningInstanceSingleCrit$new(
  task = tsk("iris"),
  learner = learner,
  resampling = rsmp("holdout"),
  measure = msr("classif.acc"),
  terminator = trm("gens", generations = 10)
```

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```
# Create TunerMies object
mies_tune <- tnr("mies", mutator = op.m, recombinator = op.r,
   parent_selector = op.parent, survival_selector = op.survival,
   mu = 10, lambda = 5)
# mies_tune$optimize performs MIES optimization and returns the optimum
mies_tune$optimize(ti)</pre>
```

ParamSetShadow

ParamSetShadow

# **Description**

Wraps another ParamSet and shadows out a subset of its Domains. The original ParamSet can still be accessed through the \$origin field; otherwise, the ParamSetShadow behaves like a ParamSet where the shadowed Domains are not present.

# Super class

```
paradox::ParamSet -> ParamSetShadow
```

# **Active bindings**

```
params (named list())
```

Table of rows identifying the contained Domains

params\_unid (named list of Param) List of Param that are members of the wrapped ParamSet with the shadowed Params removed. This is a field mostly for internal usage that has the \$ids set to invalid values but avoids cloning overhead.

Deprecated by the upcoming paradox package update and will be removed in the future.

```
deps (data.table)
```

Table of dependencies, as in ParamSet. The dependencies that are related to shadowed parameters are not exposed. This data.table should be seen as read-only and not modified in-place; instead, the \$origin's \$deps should be modified.

```
values (named list)
```

List of values, as in ParamSet, with the shadowed values removed.

```
set_id (data.table)
```

Id of the wrapped ParamSet. Changing this value will also change the wrapped ParamSet's \$set\_id accordingly.

```
origin (ParamSet)
```

ParamSet being wrapped. This object can be modified by reference to influence the ParamSetShadow object itself.

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# Methods

```
Public methods:
  ParamSetShadow$new()
  • ParamSetShadow$test_constraint()
  • ParamSetShadow$add_dep()
  • ParamSetShadow$clone()
Method new(): Initialize the ParamSetShadow object.
 Usage:
 ParamSetShadow$new(set, shadowed)
 Arguments:
 set (ParamSet)
     ParamSet to wrap.
 shadowed (character)
     Ids of Domains to shadow from sets, must be a subset of set$ids().
Method test_constraint(): Checks underlying ParamSet's constraint. It uses the underlying
$values for shadowed values.
 Usage:
 ParamSetShadow$test_constraint(x, ...)
 Arguments:
 x (named list) values to test
 ... Further arguments passed to ParamSet's $test_constraint() function.
 Returns: logical(1).
Method add_dep(): Adds a dependency to the unterlying ParamSet.
 Usage:
 ParamSetShadow$add_dep(id, on, cond, allow_dangling_dependencies = FALSE, ...)
 Arguments:
 id (character(1))
 on (character(1))
 cond (Condition)
 allow_dangling_dependencies (logical(1)): Whether to allow dependencies on parame-
     ters that are not present.
 ... Further arguments passed to ParamSet's $add_dep() function.
```

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

ParamSetShadow\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

Returns: invisible(self).

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# **Examples**

```
p1 = ps(x = p_dbl(0, 1), y = p_lgl())
p1$values = list(x = 0.5, y = TRUE)
print(p1)

p2 = ParamSetShadow$new(p1, "x")
print(p2$values)

p2$values$y = FALSE
print(p2)
print(p2$origin$values)
```

rank\_nondominated

Perform Nondominated Sorting

# **Description**

Assign elements of fitnesses to nondominated fronts.

The first nondominated front is the set of individuals that is not dominated by any other individual with respect to any fitness dimension, i.e. where no other individual exists that has all fitness values greater or equal, with at least one fitness value strictly greater.

The n'th nondominated front is the set of individuals that is not dominated by any other individual that is not in any nondominated front with smaller n.

Fitnesses are *maximized*, so the individuals in lower numbered nondominated fronts tend to have higher fitness values.

#### **Usage**

```
rank_nondominated(fitnesses, epsilon = 0)
```

# Arguments

fitnesses (numeric matrix)

fitness matrix, with one row per individual and one column per objective

epsilon (numeric)

Epsilon-vaue for non-dominance. A value is epsilon-dominated by another if it is at least epsilon smaller than the other in all dimensions, and more than epsilon smaller than the other in one dimension. epsilon may be a scalar, in which case it is used for all dimensions or a vector, in which case its length must match the number of dimensions. Default 0.

#### Value

list: \$front: Vector assigning each individual in fitnesses its nondominated front. \$domcount: Length N vector counting the number of individuals that dominate the given individual.

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Recombinator

Recombinator Base Class

# Description

Base class representing recombination operations, inheriting from MiesOperator.

Recombinators get a table of individuals as input and return a table of modified individuals as output. Individuals are acted on by groups: every \$n\_indivs\_out lines of output corresponds to a group of \$n\_indivs\_in lines of input, and presence or absence of other input groups does not affect the result.

Recombination operations are performed in ES algorithms to facilitate exploration of the search space that combine partial solutions.

# **Inheriting**

Recombinator is an abstract base class and should be inherited from. Inheriting classes should implement the private \( \). recombine() function. The user of the object calls \( \)operate(), which calls \( \). recombine() for each \( \)n\_indivs\_in sized group of individuals after checking that the operator is primed, that the values argument conforms to the primed domain. \( \). recombine() should then return a table of \( \)n\_indivs\_out individuals for each call. Typically, the \( \)initialize() function should also be overloaded, and optionally the \( \)prime() function; they should call their super equivalents.

# Super class

```
miesmuschel::MiesOperator -> Recombinator
```

### **Active bindings**

```
n_indivs_in (integer(1))
```

Number of individuals to consider at the same time. When operating, the number of input individuals must be divisible by this number.

```
n_indivs_out (integer(1))
```

Number of individuals produced for each group of \$n\_indivs\_in individuals.

#### Methods

# **Public methods:**

- Recombinator\$new()
- Recombinator\$clone()

Method new(): Initialize base class components of the Recombinator.

Usage:

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```
param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
  param_set = ps(),
  n_{indivs_{in}} = 2,
  n_indivs_out = n_indivs_in,
  packages = character(0),
  dict_entry = NULL,
  own_param_set = quote(self$param_set)
Arguments:
param_classes (character)
    Classes of parameters that the operator can handle. May contain any of "ParamLg1",
    "ParamInt", "ParamDbl", "ParamFct". Default is all of them.
    The $param_classes field will reflect this value.
param set (ParamSet | list of expression)
    Strategy parameters of the operator. This should be created by the subclass and given to
    super$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet
    directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate
    to ParamSets, possibly referencing self and private. These ParamSet are then combined
    using a ParamSetCollection. Default is the empty ParamSet.
    The $param_set field will reflect this value.
n_indivs_in (integer(1))
    Number of individuals to consider at the same time. When operating, the number of input
    individuals must be divisible by this number. Default 2.
    The $n indivs in field will reflect this value.
n_indivs_out (integer(1))
    Number of individuals that result for each n_indivs_in lines of input. The number of re-
    sults from the recombinator will be nrow(values) / n_indivs_in * n_indivs_out. De-
    fault equal to n_indivs_in.
```

The \$n\_indivs\_out field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

Recombinator\$new(

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

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```
Recombinator$clone(deep = FALSE)

Arguments:
deep Whether to make a deep clone.
```

# See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, RecombinatorPair, Scalor, Selector, SelectorScalar

Other recombinators: OperatorCombination, RecombinatorPair, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swadict\_recombinators\_xounif

RecombinatorPair

Pair Recombinator Base Class

# **Description**

Base class for recombination that covers the common case of combining two individuals, where two (typically complementary) child individuals could be taken as the result, such as bitwise crossover or SBX crossover.

This is a relatively lightweight class, it adds the keep\_complement active binding and sets \$n\_indivs\_in and \$n\_indivs\_out appropriately.

### **Inheriting**

RecombinatorPair is an abstract base class and should be inherited from. Inheriting classes should implement the private \(\frac{1}{2}\) recombine\_pair() function. During \(\frac{1}{2}\) operate(), the \(\frac{1}{2}\) recombine\_pair() function is called with the same input as the \(\frac{1}{2}\) recombine() function of the Recombinator class. It should return a data.table of two individuals.

Constructors of inheriting classes should have a keep\_complement argument.

# **Super classes**

```
miesmuschel::MiesOperator -> miesmuschel::Recombinator -> RecombinatorPair
```

# **Active bindings**

```
keep_complement (logical(1))
```

Whether the operation keeps both resulting individuals of the operation or discards the complement.

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### Methods

#### **Public methods:**

- RecombinatorPair\$new()
- RecombinatorPair\$clone()

**Method** new(): Initialize base class components of the RecombinatorPair.

```
Usage:
```

```
RecombinatorPair$new(
  keep_complement = TRUE,
  param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
  param_set = ps(),
  packages = character(0),
  dict_entry = NULL,
  own_param_set = quote(self$param_set)
)
```

Arguments:

```
keep_complement (logical(1))
```

Whether the operation should keep both resulting individuals (TRUE), or only the first and discard the complement (FALSE). Default TRUE. The \$keep\_complement field will reflect this value.

```
param_classes (character)
```

Classes of parameters that the operator can handle. May contain any of "ParamLgl", "ParamInt", "ParamDbl", "ParamFct". Default is all of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet|list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** clone(): The objects of this class are cloneable with this method.

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```
Usage:
```

RecombinatorPair\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

# See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, Scalor, Selector, SelectorScalar

Other recombinators: OperatorCombination, Recombinator, dict\_recombinators\_cmpmaybe, dict\_recombinators\_convex, dict\_recombinators\_cvxpair, dict\_recombinators\_maybe, dict\_recombinators\_nudict\_recombinators\_proxy, dict\_recombinators\_sbx, dict\_recombinators\_sequential, dict\_recombinators\_swadict\_recombinators\_xonary, dict\_recombinators\_xounif

repr

Create a 'call' Object Representation

# **Description**

repr() creates a call object representing obj, if possible. Evaluating the call should come close to recreating the original object.

In the most trivial cases, it should be possible to recreate objects from their representation by evaluating them using eval(). Important exceptions are:

- Functions are represented by their source code, if available, and by their AST if not. This drops the context from their environments and recreated objects will not work if they contain functions that depend on specific environments
- environments are not represented.
- R6 objects are only represented if they have a \$repr() function. This function may have arbitrary arguments, and should have a ... argument to capture ignored arguments.

Objects that can not be represented are currently mapped to the call stop("<###>"), where ### is a short description of the non-representable object.

# Usage

```
repr(obj, ...)
```

# **Arguments**

```
obj (any)
```

Object to create a representation of.

... (any)

Further arguments to be passed to class methods. Currently in use are:

• skip\_defaults (logical(1)) whether to skip construction arguments that have their default value. Default TRUE.

- show\_params (logical(1)) whether to show ParamSet values. Default TRUF
- show\_constructor\_args (logical(1)) whether to show construction args that are not ParamSet values. Default TRUE.

#### Value

call: A call that, when evaluated, tries to re-create the object.

SamplerRandomWeights Sampler for Projection Weights

# **Description**

Sampler for a single p\_uty that samples weight-matrices as used by ScalorFixedProjection.

# Super class

```
paradox::Sampler -> SamplerRandomWeights
```

# **Active bindings**

```
nobjectives (numeric(1))
    Number of objectives for which weights are generated.

nweights (numeric(1))
    Number of weight vectors generated for each configuration.

weights_component_id (numeric(1))
    search space component identifying the weights by which to scalarize.
```

# Methods

# **Public methods:**

- SamplerRandomWeights\$new()
- SamplerRandomWeights\$clone()

Method new(): Initialize the SamplerRandomWeights object.

```
Usage:
SamplerRandomWeights$new(
  nobjectives = 2,
  nweights = 1,
  weights_component_id = "scalarization_weights"
)
Arguments:
nobjectives (numeric(1))
  Number of objectives for which weights are generated.
```

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```
nweights (numeric(1))
   Number of weight vectors generated for each configuration.
weights_component_id (character(1))
   Id of the p_uty. Default is "scalarization_weights". Can be changed arbitrarily but should match the ScalorFixedProjection's weights_component_id.
```

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
SamplerRandomWeights$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

# **Examples**

```
set.seed(1)
```

Scalarizer

Scalarizer

# **Description**

Scalarizer objects are functions taking a fitness-matrix fitnesses (Nindivs x Nobjectives, with higher values indicating higher desirability) and a list of weight matrices weights (Nindivs elements of Nobjectives x Nweights matrices; positive weights indicate a positive contribution to scale) and returns a matrix of scalarizations (Nindivs x Nweights, with higher values indicating greater desirability).

Any other function conforming to these requirements can also be used in place of a Scalarizer, but the provided Scalarizer functions cover the most common use cases.

Scalarizers are constructed from constructor-functions, such as scalarizer\_linear() or scalarizer\_chebyshev().

## See Also

Other Scalarizers: scalarizer\_chebyshev(), scalarizer\_linear()

scalarizer\_chebyshev 179

```
scalarizer_chebyshev Chebyshev Scalarizer
```

# **Description**

Constructs a Scalarizer that does Chebyshev scalarization, as employed in ParEGO by Knowles (2006).

The Chebyshev scalarization for a single individual with fitness values f and given weight vector w is min(w \* f) + rho \* sum(w \* f), where rho is a hyperparameter given during construction.

# Usage

```
scalarizer_chebyshev(rho = 0.05)
```

# **Arguments**

```
rho (numeric(1))
Small positive value.
```

# Value

```
a Scalarizer object.
```

#### References

Knowles, Joshua (2006). "ParEGO: A hybrid algorithm with on-line landscape approximation for expensive multiobjective optimization problems." *IEEE Transactions on Evolutionary Computation*, **10**(1), 50–66.

### See Also

```
Other Scalarizers: Scalarizer, scalarizer_linear()
```

# **Examples**

```
# fitnesses: three rows (i.e. thee indivs) with two objective values each
fitnesses <- matrix(0:5, ncol = 2)

# weights: contains one matrix for each row of 'fitnesses' (i.e. each indiv)
# which get multiplied with their respective row.
weights <- list(
    matrix(c(1, 0, 0, 1), ncol = 2),
    matrix(c(1, 2, 0, 0), ncol = 2),
    matrix(c(0, 1, 0, 1), ncol = 2)
)
sc <- scalarizer_chebyshev()</pre>
```

# The resulting row-vectors are the different scalarizations according to the

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```
# columns in the 'weights' matrices.
sc(fitnesses, weights)
sc <- scalarizer_chebyshev(rho = 0.1)
sc(fitnesses, weights)</pre>
```

scalarizer\_linear

Linear Scalarizer

# Description

Constructs a linear Scalarizer, which performs linear scalarization for ScalorFixedProjection.

# Usage

```
scalarizer_linear()
```

#### Value

a Scalarizer object.

# See Also

Other Scalarizers: Scalarizer, scalarizer\_chebyshev()

# **Examples**

```
# fitnesses: three rows (i.e. thee indivs) with two objective values each
fitnesses <- matrix(0:5, ncol = 2)

# weights: contains one matrix for each row of 'fitnesses' (i.e. each indiv)
# which get multiplied with their respective row.
weights <- list(
    matrix(c(1, 0, 0, 1), ncol = 2),
    matrix(c(1, 2, 0, 0), ncol = 2),
    matrix(c(0, 1, 0, 1), ncol = 2)
)

sc <- scalarizer_linear()

# The resulting row-vectors are the different scalarizations according to the
# columns in the 'weights' matrices.
sc(fitnesses, weights)</pre>
```

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Scalor

Scalor Base Class

#### **Description**

Base class representing ranking operations, inheriting from MiesOperator.

A Scalor gets a table of individuals as input, along with information on the individuals' performance values and returns a vector of a possible scalarization of individuals' fitness (or other qualities).

Scalors can be used by Selectors as a basis to select individuals by. This way it is possible to have tournament selection (SelectorTournament) or elite selection (SelectorBest) based on different, configurable qualities of individuals.

Unlike most other operator types inheriting from MiesOperator, the \$operate() function has two arguments, which are passed on to \$.scale()

- values :: data.frame
  Individuals to operate on. Must pass the check of the ParamSet given in the last \$prime()
  call and may not have any missing components.
- fitnesses:: numeric|matrix
  Fitnesses for each individual given in values. If this is a numeric, then its length must be equal to the number of rows in values. If this is a matrix, if number of rows must be equal to the number of rows in values, and it must have one column when doing single-crit optimization and one column each for each "criterion" when doing multi-crit optimization. Note that fitness values are always maximized, both in single- and multi-criterion optimization, so objective output is multiplied with -1 if it is tagged as "minimize".

The return value of an operation should be a numeric vector with one finite value for each entry of values, assigning high values to individuals in some way more "desirable" than others with low values.

#### **Inheriting**

Scalor is an abstract base class and should be inherited from. Inheriting classes should implement the private \$.scale() function. The user of the object calls \$operate(), and the arguments are passed on to private \$.scale() after checking that the operator is primed, that the values argument conforms to the primed domain and that other values match. Typically, the \$initialize() function should also be overloaded, and optionally the \$prime() function; they should call their super equivalents.

#### Super class

```
miesmuschel::MiesOperator-> Scalor
```

#### **Active bindings**

```
supported (character)
```

Optimization supported by this Scalor, can be "single-crit", "multi-crit", or both.

Scalor Scalor

## Methods

#### **Public methods:**

- Scalor\$new()
- Scalor\$clone()

**Method** new(): Initialize base class components of the Mutator.

```
Usage:
Scalor$new(
  param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
  param_set = ps(),
  supported = c("single-crit", "multi-crit"),
  packages = character(0),
  dict_entry = NULL,
  own_param_set = quote(self$param_set)
)
Arguments:
```

param\_classes (character)

Classes of parameters that the operator can handle. May contain any of "ParamLgl", "ParamInt", "ParamDbl", "ParamFct". Default is all of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

```
supported (character)
```

Subset of "single-crit" and "multi-crit", indicating wether single and / or multi-criterion optimization is supported. Default both of them.

The \$supported field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** clone(): The objects of this class are cloneable with this method.

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```
Usage:
Scalor$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

#### See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Selector, SelectorScalar

Other scalors: dict\_scalors\_aggregate, dict\_scalors\_domcount, dict\_scalors\_fixedprojection, dict\_scalors\_hypervolume, dict\_scalors\_nondom, dict\_scalors\_one, dict\_scalors\_proxy, dict\_scalors\_single

Selector

Selector Base Class

## **Description**

Base class representing selection operations, inheriting from MiesOperator.

A Selector gets a table of individuals as input, along with information on the individuals' performance values and the number of individuals to select, and returns a vector of integers indicating which individuals were selected.

Selection operations are performed in ES algorithms to facilitate concentration towards individuals that perform well with regard to the fitness measure.

Fitness values are always *maximized*, both in single- and multi-criterion optimization.

Unlike most other operator types inheriting from MiesOperator, the \$operate() function has three arguments, which are passed on to \$.select()

- values:: data.frame
   Individuals to operate on. Must pass the check of the Domain given in the last \$prime() call and may not have any missing components.
- fitnesses :: numeric | matrix Fitnesses for each individual given in values. If this is a numeric, then its length must be equal to the number of rows in values. If this is a matrix, if number of rows must be equal to the number of rows in values, and it must have one column when doing single-crit optimization and one column each for each "criterion" when doing multi-crit optimization. The fitnesses-value passed on to \$.select() is always a matrix.
- n\_select :: integer(1)

  Number of individuals to select. Some Selectors select individuals with replacement, for which this value may be greater than the number of rows in values.
- group\_size :: integer
  Sampling group size hint, indicating that the caller would prefer there to not be any duplicates within this group size, e.g. because the Selector is called to select individuals to be given to a Recombinator with a certain n\_indivs\_in, or because it is called as a survival\_selector

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in mies\_survival\_comma() or mies\_survival\_plus(). The Selector may or may not ignore this value, however. This may possibly happen because of certain configuration parameters, or because the input size is too small.

Must either be a scalar value or sum up to n\_select. Must be non-negative. A scalar value of 0 is interpreted the same as 1.

If not given, this value defaults to 1.

The return value for an operation will be a numeric vector of integer values of length n\_select indexing the individuals that were selected. Some Selectors select individuals with replacement, for which the return value may contain indices more than once.

## **Inheriting**

Selector is an abstract base class and should be inherited from. Inheriting classes should implement the private \$.select() function. The user of the object calls \$operate(), and the arguments are passed on to private \$.select() after checking that the operator is primed, that the values argument conforms to the primed domain and that other values match. Typically, the \$initialize() function should also be overloaded, and optionally the \$prime() function; they should call their super equivalents.

## Super class

```
miesmuschel::MiesOperator -> Selector
```

## **Active bindings**

```
supported (character)

Optimization supported by this Selector, can be "single-crit", "multi-crit", or both.
```

## Methods

#### **Public methods:**

- Selector\$new()
- Selector\$clone()

**Method** new(): Initialize base class components of the Selector.

```
Usage:
Selector$new(
   is_deterministic = FALSE,
   param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
   param_set = ps(),
   supported = c("single-crit", "multi-crit"),
   packages = character(0),
   dict_entry = NULL,
   own_param_set = quote(self$param_set)
)

Arguments:
```

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```
is_deterministic (logical(1))
```

Whether the Selector is deterministic. Setting this to TRUE adds a configuration parameter shuffle\_selection (initialized to TRUE) that causes the selection to be shuffled.

```
param_classes (character)
```

Classes of parameters that the operator can handle. May contain any of "ParamLgl", "ParamInt", "ParamDbl", "ParamFct". Default is all of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

```
supported (character)
```

Subset of "single-crit" and "multi-crit", indicating wether single and / or multi-criterion optimization is supported. Default both of them.

The \$supported field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

```
own_param_set (language)
```

An expression that evaluates to a ParamSet indicating the configuration parameters that are entirely owned by this operator class (and not proxied from a construction argument object). This should be quote(self\$param\_set) (the default) when the param\_set argument is not a list of expressions.

**Method** clone(): The objects of this class are cloneable with this method.

```
Usage:
Selector$clone(deep = FALSE)
Arguments:
```

Argumenis.

deep Whether to make a deep clone.

## See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, SelectorScalar

Other selectors: SelectorScalar, dict\_selectors\_best, dict\_selectors\_maybe, dict\_selectors\_null, dict\_selectors\_proxy, dict\_selectors\_random, dict\_selectors\_sequential, dict\_selectors\_tournament

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SelectorScalar

Selector making use of Scalors

## **Description**

Base class inheriting from Selector for selection operations that make use of scalar values, generated by Scalor.

#### **Inheriting**

SelectorScaling is an abstract base class and should be inherited from. Inheriting classes should implement the private \$.select\_scalar() function. During \$operate(), the \$.select\_scalar() function is called, it should have three arguments, similar to Selector's \$.select() function. values and n\_select are as given to \$.select() of the Selector. The fitnesses argument is first scaled by the associated Scalor and then passed on as a numeric vector.

Typically, \$initialize() should also be overloaded when inheriting.

## **Super classes**

```
miesmuschel::MiesOperator -> miesmuschel::Selector -> SelectorScalar
```

## **Active bindings**

```
scalor (Scalor)

Scalor used to scalarize fitnesses for selection.
```

#### Methods

## **Public methods:**

- SelectorScalar\$new()
- SelectorScalar\$prime()
- SelectorScalar\$clone()

**Method** new(): Initialize base class components of the SelectorScalar.

```
Usage:
SelectorScalar$new(
    scalor = ScalorSingleObjective$new(),
    is_deterministic = FALSE,
    param_classes = c("ParamLgl", "ParamInt", "ParamDbl", "ParamFct"),
    param_set = ps(),
    supported = scalor$supported,
    packages = character(0),
    dict_entry = NULL
)
Arguments:
```

```
scalor (Scalor)
```

Scalor to use to generate scalar values from multiple objectives, if multi-objective optimization is performed. Initialized to ScalorSingleObjective: Doing single-objective optimization normally, throwing an error if used in multi-objective setting: In that case, a Scalor needs to be explicitly chosen.

```
is_deterministic (logical(1))
```

Whether the Selector is deterministic. Setting this to TRUE adds a configuration parameter shuffle\_selection (initialized to TRUE) that causes the selection to be shuffled.

```
param_classes (character)
```

Classes of parameters that the operator can handle. May contain any of "ParamLgl", "ParamInt", "ParamDbl", "ParamFct". Default is all of them.

The \$param\_classes field will reflect this value.

```
param_set (ParamSet | list of expression)
```

Strategy parameters of the operator. This should be created by the subclass and given to super\$initialize(). If this is a ParamSet, it is used as the MiesOperator's ParamSet directly. Otherwise it must be a list of expressions e.g. created by alist() that evaluate to ParamSets, possibly referencing self and private. These ParamSet are then combined using a ParamSetCollection. Default is the empty ParamSet.

The \$param\_set field will reflect this value.

```
supported (character)
```

Subset of "single-crit" and "multi-crit", indicating wether single and / or multi-criterion optimization is supported. Default to the supported set of scalor.

The \$supported field will reflect this value.

packages (character) Packages that need to be loaded for the operator to function. This should be declared so these packages can be loaded when operators run on parallel instances. Default is character(0).

The \$packages field will reflect this values.

```
dict_entry (character(1) | NULL)
```

Key of the class inside the Dictionary (usually one of dict\_mutators, dict\_recombinators, dict\_selectors), where it can be retrieved using a short access function. May be NULL if the operator is not entered in a dictionary.

The \$dict\_entry field will reflect this value.

**Method** prime(): See MiesOperator method. Primes both this operator, as well as the wrapped operator given to scalor during construction.

```
Usage:
SelectorScalar$prime(param_set)
Arguments:
param_set (ParamSet)
    Passed to MiesOperator$prime().
Returns: invisible self.

Method clone(): The objects of this class are cloneable with this method.
Usage:
SelectorScalar$clone(deep = FALSE)
Arguments:
deep Whether to make a deep clone.
```

## See Also

Other base classes: Filtor, FiltorSurrogate, MiesOperator, Mutator, MutatorDiscrete, MutatorNumeric, OperatorCombination, Recombinator, RecombinatorPair, Scalor, Selector

Other selectors: Selector, dict\_selectors\_best, dict\_selectors\_maybe, dict\_selectors\_null, dict\_selectors\_proxy, dict\_selectors\_random, dict\_selectors\_sequential, dict\_selectors\_tournament

terminator\_get\_generations

Get the Numger of Generations that a Terminator Allows

## **Description**

Get the number of generations of a TerminatorGenerations. When the TerminatorGenerations is wrapped in a TerminatorCombo, then the minimum number of generations allowed by it are retrieved. This is the minimum of all terminator\_get\_generations if \$any is set to TRUE, and the maximum if \$any is set to FALSE.

The number of generations allowed by other Terminators is infinity.

#### **Usage**

```
terminator_get_generations(x)
```

## **Arguments**

x (Terminator)
Terminator to query.

#### Value

numeric(1): The theoretical maximum number of generations allowed by the Terminator.

TuningInstanceMultiCrit

TuningInstanceMultiCrit Class

## **Description**

mlr3tuning's TuningInstanceMultiCrit class. Re-exported since mlr3tuning will change the name.

## Super classes

```
bbotk::OptimInstance->bbotk::OptimInstanceBatch->bbotk::OptimInstanceBatchMultiCrit
->mlr3tuning::TuningInstanceBatchMultiCrit -> TuningInstanceMultiCrit
```

## Methods

#### **Public methods:**

• TuningInstanceMultiCrit\$clone()

Method clone(): The objects of this class are cloneable with this method.

Usage:

TuningInstanceMultiCrit\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

TuningInstanceSingleCrit

TuningInstanceSingleCrit Class

## Description

mlr3tuning's TuningInstanceSingleCrit class. Re-exported since mlr3tuning will change the name.

## Super classes

```
bbotk::OptimInstance->bbotk::OptimInstanceBatch->bbotk::OptimInstanceBatchSingleCrit
->mlr3tuning::TuningInstanceBatchSingleCrit -> TuningInstanceSingleCrit
```

## Methods

## **Public methods:**

• TuningInstanceSingleCrit\$clone()

**Method** clone(): The objects of this class are cloneable with this method.

Usage:

TuningInstanceSingleCrit\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

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