

# Package ‘mlr3viz’

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**Title** Visualizations for 'mlr3'

**Version** 0.5.10

**Description** Provides visualizations for 'mlr3' objects such as tasks, predictions, resample results or benchmark results via the autoplot() generic of 'ggplot2'. The returned 'ggplot' objects are intended to provide sensible defaults, yet can easily be customized to create camera-ready figures. Visualizations include barplots, boxplots, histograms, ROC curves, and Precision-Recall curves.

**License** LGPL-3

**URL** <https://mlr3viz.mlr-org.com>, <https://github.com/mlr-org/mlr3viz>

**BugReports** <https://github.com/mlr-org/mlr3viz/issues>

**Depends** R (>= 3.1.0)

**Imports** checkmate, data.table, ggplot2 (>= 3.3.0), mlr3misc (>= 0.7.0), scales, utils, viridis

**Suggests** bbotk, cluster, factoextra, GGally, ggfortify (>= 0.4.11), ggparty, glmnet, knitr, lgr, mlr3 (>= 0.6.0), mlr3cluster, mlr3filters, mlr3learners, mlr3tuning (>= 0.9.0), paradox, partykit, patchwork (>= 1.1.1), precrec, ranger, rpart, stats, testthat (>= 3.0.0), vdiffir (>= 1.0.2), xgboost

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**Collate** 'BenchmarkResult.R' 'Filter.R' 'LearnerClassifCVGlmnet.R'  
'LearnerClassifGlmnet.R' 'LearnerClassifRpart.R'  
'LearnerClustHierarchical.R' 'LearnerRegrCVGlmnet.R'  
'LearnerRegrGlmnet.R' 'LearnerRegrRpart.R'  
'OptimInstanceSingleCrit.R' 'Prediction.R'  
'PredictionClassif.R' 'PredictionClust.R' 'PredictionRegr.R'  
'ResampleResult.R' 'Task.R' 'TaskClassif.R' 'TaskClust.R'

'TaskRegr.R' 'TuningInstanceSingleCrit.R' 'as\_precrec.R'  
 'bibentries.R' 'helper.R' 'plot\_learner\_prediction.R'  
 'reexports.R' 'theme-mlr3.R' 'zzz.R'

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## Description

Provides visualizations for 'mlr3' objects such as tasks, predictions, resample results or benchmark results via the autoplot() generic of 'ggplot2'. The returned 'ggplot' objects are intended to provide sensible defaults, yet can easily be customized to create camera-ready figures. Visualizations include barplots, boxplots, histograms, ROC curves, and Precision-Recall curves.

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**See Also**

Useful links:

- <https://mlr3viz.mlr-org.com>
- <https://github.com/mlr-org/mlr3viz>
- Report bugs at <https://github.com/mlr-org/mlr3viz/issues>

---

as\_precrec

*Convert to 'precrec' Format*

---

**Description**

Converts to a format which is understood by `precrec::evalmod()` of package **precrec**.

**Usage**

```
as_precrec(object)

## S3 method for class 'PredictionClassif'
as_precrec(object)

## S3 method for class 'ResampleResult'
as_precrec(object)

## S3 method for class 'BenchmarkResult'
as_precrec(object)
```

**Arguments**

object            (any)  
                  Object to convert.

**Value**

Object as created by `precrec::mmdata()`.

**References**

Saito T, Rehmsmeier M (2017). "Precrec: fast and accurate precision-recall and ROC curve calculations in R." *Bioinformatics*, **33**(1), 145-147. doi:10.1093/bioinformatics/btw570.

---

autoplot.BenchmarkResult

*Plot for BenchmarkResult*

---

**Description**

Generates plots for `mlr3::BenchmarkResult`, depending on argument type:

- "boxplot" (default): Boxplots of performance measures, one box per `mlr3::Learner` and one facet per `mlr3::Task`.
- "roc": ROC curve (1 - specificity on x, sensitivity on y). The `mlr3::BenchmarkResult` may only have a single `mlr3::Task` and a single `mlr3::Resampling`. Note that you can subset any `mlr3::BenchmarkResult` with its `$filter()` method (see examples). Requires package `precrec`. Additional arguments will be passed down to the respective `autoplot()` function in package `precrec`. Arguments `calc_avg` and `cb_alpha` are passed to `precrec::evalmod()`.
- "prc": Precision recall curve. See "roc".

**Usage**

```
## S3 method for class 'BenchmarkResult'
autoplot(object, type = "boxplot", measure = NULL, ...)
```

**Arguments**

object	( <code>mlr3::BenchmarkResult</code> ).
type	(character(1)): Type of the plot. See description.
measure	( <code>mlr3::Measure</code> ) Performance measure to use.
...	(any): Additional arguments, passed down to the respective geom or plotting function.

**Value**

`ggplot2::ggplot()` object.

**Theme**

The `theme_mlr3()` and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**References**

Saito T, Rehmsmeier M (2017). “Precrec: fast and accurate precision-recall and ROC curve calculations in R.” *Bioinformatics*, **33**(1), 145-147. doi:10.1093/bioinformatics/btw570.

**Examples**

```
if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  tasks = tsks(c("pima", "sonar"))
  learner = lrns(c("classif.featureless", "classif.rpart"),
    predict_type = "prob")
  resampling = rsmpls("cv")
  object = benchmark(benchmark_grid(tasks, learner, resampling))

  head(fortify(object))
  autoplot(object)
  autoplot(object$clone(deep = TRUE)$filter(task_ids = "pima"), type = "roc")
}
```

---

 autoplot.Filter

*Plot for Filter Scores*


---

**Description**

Generates plots for `mlr3filters::Filter`, depending on argument type:

- "barplot" (default): Bar plot of filter scores.

**Usage**

```
## S3 method for class 'Filter'
autoplot(object, type = "boxplot", n = Inf, ...)
```

**Arguments**

object	( <code>mlr3filters::Filter</code> ).
type	(character(1)): Type of the plot. See description.
n	(integer(1)) Only include the first n features with highest importance. Defaults to all features.
...	(any): Additional argument, passed down to the respective geom.

**Value**

`ggplot2::ggplot()` object.

**Theme**

The `theme_ml3()` and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```
if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)
  library(mlr3filters)

  task = tsk("mtcars")
  f = flt("correlation")
  f$calculate(task)

  head(fortify(f))
  autoplot(f, n = 5)
}
```

---

`autoplot.LearnerClassifCVGlmnet`

*Plot for LearnerClassifGlmnet / LearnerRegrGlmnet / LearnerClassifCVGlmnet / LearnerRegrCVGlmnet*

---

**Description**

Visualizations for `mlr3learners::mlr_learners_classif.glmnet`, `mlr3learners::mlr_learners_regr.glmnet`, `mlr3learners::mlr_learners_classif.cv.glmnet` and `mlr3learners::mlr_learners_regr.cv.glmnet` using the package **ggfortify**.

Note that learner-specific plots are experimental and subject to change.

**Usage**

```
## S3 method for class 'LearnerClassifCVGlmnet'
autoplot(object, ...)

## S3 method for class 'LearnerClassifGlmnet'
autoplot(object, ...)

## S3 method for class 'LearnerRegrCVGlmnet'
autoplot(object, ...)

## S3 method for class 'LearnerRegrGlmnet'
autoplot(object, ...)
```

**Arguments**

object (mlr3learners::LearnerClassifGlmnet | mlr3learners::LearnerRegrGlmnet | mlr3learners::LearnerRegrCVGlmnet | mlr3learners::LearnerRegrCVGlmnet).

... (any): Additional arguments, passed down to `ggparty::autoplot.party()`.

**Value**

`ggplot2::ggplot()` object.

**Theme**

The `theme_ml3()` and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**References**

Tang Y, Horikoshi M, Li W (2016). “ggfortify: Unified Interface to Visualize Statistical Result of Popular R Packages.” *The R Journal*, **8**(2), 474–485. doi:10.32614/RJ2016060.

**Examples**

```
## Not run:
library(mlr3)
library(mlr3viz)
library(mlr3learners)

# classification
task = tsk("sonar")
learner = lrn("classif.glmnet")
learner$train(task)
autoplot(learner)

# regression
task = tsk("mtcars")
learner = lrn("regr.glmnet")
learner$train(task)
autoplot(learner)

## End(Not run)
```

---

autoplot.LearnerClassifRpart

*Plot for LearnerClassifRpart / LearnerRegrRpart*

---

**Description**

Visualize trees for `mlr3::mlr_learners_classif.rpart` and `mlr3::mlr_learners_regr.rpart` using the package **ggparty**.

Contrary to **ggparty**, boxplots are shown in the terminal nodes for regression trees.

Note that learner-specific plots are experimental and subject to change.

**Usage**

```
## S3 method for class 'LearnerClassifRpart'
autoplot(object, ...)

## S3 method for class 'LearnerRegrRpart'
autoplot(object, ...)
```

**Arguments**

`object` (`mlr3::LearnerClassifRpart` | `mlr3::LearnerRegrRpart`).  
`...` (any): Additional arguments, passed down to `ggparty::autoplot.party()`.

**Value**

`ggplot2::ggplot()` object.

**Theme**

The `theme_ml3()` and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```
if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  # classification
  task = tsk("iris")
  learner = lrn("classif.rpart", keep_model = TRUE)
  learner$train(task)
  autoplot(learner)

  # regression
  task = tsk("mtcars")
  learner = lrn("regr.rpart", keep_model = TRUE)
  learner$train(task)
  autoplot(learner)
}
```



---

`autoplot.LearnerClustHierarchical`*Plot for Hierarchical Clustering Learners*

---

## Description

Generates plots for hierarchical clusterers, depending on argument type:

- "dend" (default): dendrograms using **factoextra** package.
- "scree": scree plot that shows the number of possible clusters on x-axis and the height on the y-axis.

Note that learner-specific plots are experimental and subject to change.

## Usage

```
## S3 method for class 'LearnerClustHierarchical'  
autoplot(object, type = "dend", ...)
```

## Arguments

<code>object</code>	( <code>mlr3cluster::LearnerClustAgnes</code>   <code>mlr3cluster::LearnerClustDiana</code>   <code>mlr3cluster::LearnerClustHclust</code> ).
<code>type</code>	( <code>character(1)</code> ): Type of the plot. See description.
<code>...</code>	(any): Additional arguments, passed down to function <code>factoextra::fviz_dend()</code> in package <b>factoextra</b> .

## Value

`ggplot2::ggplot()` object.

## Theme

The `theme_ml3()` and `viridis` color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

## Examples

```
if (requireNamespace("mlr3")) {  
  library(mlr3)  
  library(mlr3cluster)  
  library(mlr3viz)  
  
  task = mlr_tasks$get("usarrests")  
  
  # agnes clustering  
  learner = mlr_learners$get("clust.agnes")  
  learner$train(task)
```

```

autoplot(learner)

# diana clustering
learner = mlr_learners$get("clust.diana")
learner$train(task)
autoplot(learner,
  k = learner$param_set$values$k, rect_fill = TRUE,
  rect = TRUE, rect_border = "red")

# hclust clustering
learner = mlr_learners$get("clust.hclust")
learner$train(task)
autoplot(learner, type = "scree")
}

```

---

```

autoplot.OptimInstanceSingleCrit
  Plot for OptimInstanceSingleCrit

```

---

## Description

Generates plots for [bbotk::OptimInstanceSingleCrit](#).

## Usage

```

## S3 method for class 'OptimInstanceSingleCrit'
autoplot(
  object,
  type = "marginal",
  cols_x = NULL,
  trafo = FALSE,
  learner = mlr3::lrn("regr.ranger"),
  grid_resolution = 100,
  batch = NULL,
  ...
)

```

## Arguments

object	( <a href="#">bbotk::OptimInstanceSingleCrit</a> .
type	(character(1)): Type of the plot. Available choices: <ul style="list-style-type: none"> <li>"marginal": scatter plots of x versus y The colour of the points shows the batch number.</li> <li>"performance": scatter plots of batch number versus y</li> <li>"parameter": scatter plots of batch number versus input. The colour of the points shows the y values.</li> </ul>

- "parallel" parallel coordinates plot. x values are rescaled by  $(x - \text{mean}(x)) / \text{sd}(x)$ .
- "points" - scatter plot of two x dimensions versus y. The colour of the points shows the y values.
- "surface": surface plot of two x dimensions versus y values. The y values are interpolated with the supplied [mlr3::Learner](#).
- "pairs": plots all x and y values against each other.

cols_x	(character()) Column names of x values. By default, all untransformed x values from the search space are plotted. Transformed hyperparameters are prefixed with x_domain_.
trafo	(logical(1)) Determines if untransformed (FALSE) or transformed (TRUE) x values are plotted.
learner	( <a href="#">mlr3::Learner</a> ) Regression learner used to interpolate the data of the surface plot.
grid_resolution	(numeric()) Resolution of the surface plot.
batch	(integer()) The batch number(s) to limit the plot to. Default is all batches.
...	(any): Additional arguments, possibly passed down to the underlying plot functions.

**Value**

[ggplot2::ggplot\(\)](#) object.

**Theme**

The [theme\\_ml3\(\)](#) and viridis color maps are applied by default to all autoplot() methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```
if (requireNamespace("mlr3") && requireNamespace("bbotk") && requireNamespace("patchwork")) {
  library(bbotk)
  library(paradox)

  fun = function(xs) {
    c(y = -(xs[[1]] - 2)^2 - (xs[[2]] + 3)^2 + 10)
  }
  domain = ps(
    x1 = p_dbl(-10, 10),
    x2 = p_dbl(-5, 5)
  )
  codomain = ps(
    y = p_dbl(tags = "maximize")
  )
  obfun = ObjectiveRfun$new(
```

```

    fun = fun,
    domain = domain,
    codomain = codomain
  )

instance = OptimInstanceSingleCrit$new(objective = obfun, terminator = trm("evals", n_evals = 20))

optimizer = opt("random_search", batch_size = 2)
optimizer$optimize(instance)

# plot y versus batch number
autoplot(instance, type = "performance")

# plot x1 values versus performance
autoplot(instance, type = "marginal", cols_x = "x1")

# plot parallel coordinates plot
autoplot(instance, type = "parallel")

# plot pairs
autoplot(instance, type = "pairs")
}

```

---

autoplot.PredictionClassif

*Plot for PredictionClassif*

---

## Description

Generates plots for [mlr3::PredictionClassif](#), depending on argument type:

- "stacked" (default): Stacked barplot of true and estimated class labels.
- "roc": ROC curve (1 - specificity on x, sensitivity on y). Requires package **precrec**.
- "prc": Precision recall curve. Requires package **precrec**.
- "threshold": Systematically varies the threshold of the [mlr3::PredictionClassif](#) object and plots the resulting performance as returned by *measure*. Arguments in ... are passed down to the score function of the [mlr3::Measure](#).

## Usage

```
## S3 method for class 'PredictionClassif'
autoplot(object, type = "stacked", measure = NULL, ...)
```

## Arguments

object	( <a href="#">mlr3::PredictionClassif</a> ).
type	(character(1)): Type of the plot. See description.

measure	( <a href="#">mlr3::Measure</a> ) Performance measure to use.
...	(any): Additional arguments, passed down to the respective geom, plotting function or measure.

**Value**

[ggplot2::ggplot\(\)](#) object.

**Theme**

The [theme\\_ml3\(\)](#) and viridis color maps are applied by default to all autoplot() methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**References**

Saito T, Rehmsmeier M (2017). "Precrec: fast and accurate precision-recall and ROC curve calculations in R." *Bioinformatics*, **33**(1), 145-147. doi:10.1093/bioinformatics/btw570.

**Examples**

```
if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  task = tsk("spam")
  learner = lrn("classif.rpart", predict_type = "prob")
  object = learner$train(task)$predict(task)

  head(fortify(object))
  autoplot(object)
  autoplot(object, type = "roc")
  autoplot(object, type = "prc")
}
```

---

autoplot.PredictionClust

*Plot for PredictionClust*

---

**Description**

Generates plots for [mlr3cluster::PredictionClust](#), depending on argument type:

- "scatter" (default): scatterplot with correlation values and colored cluster assignments.
- "sil": Silhouette plot with mean silhouette value as a reference line. Requires package [ggfortify](#).
- "pca": Perform PCA on data and color code cluster assignments. Inspired by and uses [ggfortify::autoplot.kmeans](#).

**Usage**

```
## S3 method for class 'PredictionClust'
autoplot(object, task, row_ids = NULL, type = "scatter", ...)
```

**Arguments**

object	( <a href="#">mlr3cluster::PredictionClust</a> ).
task	( <a href="#">mlr3cluster::TaskClust</a> ).
row_ids	row ids to subset task data to ensure that only the data used to make predictions are shown in plots.
type	(character(1)): Type of the plot. See description.
...	(any): Additional arguments, passed down to the respective geom.

**Value**

[ggplot2::ggplot\(\)](#) object.

**Theme**

The [theme\\_mlr3\(\)](#) and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**References**

Tang Y, Horikoshi M, Li W (2016). “ggfortify: Unified Interface to Visualize Statistical Result of Popular R Packages.” *The R Journal*, **8**(2), 474–485. doi:[10.32614/RJ2016060](https://doi.org/10.32614/RJ2016060).

**Examples**

```
if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3cluster)
  library(mlr3viz)

  task = tsk("usarrests")
  learner = lrn("clust.kmeans", centers = 3)
  object = learner$train(task)$predict(task)

  head(fortify(object))
  autoplot(object, task)
}
```

---

```
autoplot.PredictionRegr
```

*Plot for PredictionRegr*

---

## Description

Generates plots for [mlr3::PredictionRegr](#), depending on argument type:

- "xy" (default): Scatterplot of "true" response vs. "predicted" response. By default a linear model is fitted via `geom_smooth(method = "lm")` to visualize the trend between x and y (by default colored blue).
  - In addition `geom_abline()` with `slope = 1` is added to the plot.
  - Note that `geom_smooth()` and `geom_abline()` may overlap, depending on the given data.
- "histogram": Histogram of residuals:  $r = y - \hat{y}$ .
- "residual": Plot of the residuals, with the response  $\hat{y}$  on the "x" and the residuals on the "y" axis.
  - By default a linear model is fitted via `geom_smooth(method = "lm")` to visualize the trend between x and y (by default colored blue).

## Usage

```
## S3 method for class 'PredictionRegr'  
autoplot(object, type = "xy", ...)
```

## Arguments

object	( <a href="#">mlr3::PredictionRegr</a> ).
type	(character(1)): Type of the plot. See description.
...	(any): Additional arguments, passed down to the respective geom.

## Value

`ggplot2::ggplot()` object.

## Theme

The `theme_ml3()` and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```

if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  task = tsk("boston_housing")
  learner = lrn("regr.rpart")
  object = learner$train(task)$predict(task)

  head(fortify(object))
  autoplot(object)
  autoplot(object, type = "histogram", binwidth = 1)
  autoplot(object, type = "residual")
}

```

---

autoplot.ResampleResult

*Plot for ResampleResult*

---

**Description**

Generates plots for [mlr3::ResampleResult](#), depending on argument type:

- "boxplot" (default): Boxplot of performance measures.
- "histogram": Histogram of performance measures.
- "roc": ROC curve (1 - specificity on x, sensitivity on y). The predictions of the individual [mlr3::Resamplings](#) are merged prior to calculating the ROC curve (micro averaged). Requires package **precrec**. Additional arguments will be passed down to the respective [autoplot\(\)](#) function in package **precrec**. Arguments `calc_avg` and `cb_alpha` are passed to [precrec::evalmod\(\)](#).
- "prc": Precision recall curve. See "roc".
- "prediction": Plots the learner prediction for a grid of points. Needs models to be stored. Set `store_models = TRUE` for `[mlr3::resample]`. For classification, we support tasks with exactly two features and learners with `predict_type=` set to "response" or "prob". For regression, we support tasks with one or two features. For tasks with one feature we can print confidence bounds if the predict type of the learner was set to "se". For tasks with two features the predict type will be ignored.

**Usage**

```

## S3 method for class 'ResampleResult'
autoplot(object, type = "boxplot", measure = NULL, predict_sets = "test", ...)

```



**Arguments**

object	(mlr3::ResampleResult).
type	(character(1)): Type of the plot. See description.
measure	(mlr3::Measure) Performance measure to use.
predict_sets	(character()) Only for type set to "prediction". Which points should be shown in the plot? Can be a subset of ("train", "test") or empty.
...	(any): Additional arguments, passed down to the respective geom or plotting function.

**Value**

ggplot2::ggplot() object.

**Theme**

The `theme_ml3()` and viridis color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**References**

Saito T, Rehmsmeier M (2017). "Precrec: fast and accurate precision-recall and ROC curve calculations in R." *Bioinformatics*, **33**(1), 145-147. doi:10.1093/bioinformatics/btw570.

**Examples**

```
if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  task = tsk("sonar")
  learner = lrn("classif.rpart", predict_type = "prob")
  resampling = rsmpl("cv")
  object = resample(task, learner, resampling)

  head(fortify(object))

  # Default: boxplot
  autoplot(object)

  # Histogram
  autoplot(object, type = "histogram", bins = 30)

  # ROC curve, averaged over resampling folds:
  autoplot(object, type = "roc")

  # ROC curve of joint prediction object:
```

```

autoplot(object$prediction(), type = "roc")

# Precision Recall Curve
autoplot(object, type = "prc")

# Prediction Plot
task = tsk("iris")$select(c("Sepal.Length", "Sepal.Width"))
resampling = rsmpl("cv", folds = 3)
object = resample(task, learner, resampling, store_models = TRUE)
autoplot(object, type = "prediction")
}

```

---

autoplot.TaskClassif *Plot for Classification Tasks*

---

## Description

Generates plots for `mlr3::TaskClassif`, depending on argument type:

- "target" (default): Bar plot of the target variable (default).
- "duo": Passes data and additional arguments down to `GGally::ggduo()`. `columnsX` is target, `columnsY` is features.
- "pairs": Passes data and additional arguments down to `GGally::ggpairs()`. Color is set to target column.

## Usage

```

## S3 method for class 'TaskClassif'
autoplot(object, type = "target", ...)

```

## Arguments

<code>object</code>	( <code>mlr3::TaskClassif</code> ).
<code>type</code>	( <code>character(1)</code> ): Type of the plot. See description.
<code>...</code>	( <code>any</code> ): Additional argument, possibly passed down to the underlying plot functions.

## Value

`ggplot2::ggplot()` object.

## Theme

The `theme_ml3()` and `viridis` color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```

if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  task = tsk("iris")

  head(fortify(task))
  autoplot(task)
  autoplot(task$clone()$select(c("Sepal.Length", "Sepal.Width")),
    type = "pairs")
  autoplot(task, type = "duo")
}

```

---

autoplot.TaskClust      *Plot for Clustering Tasks*

---

**Description**

Generates plots for [mlr3cluster::TaskClust](#), depending on argument type:

- "pairs": Passes data and additional arguments down to [GGally::ggpairs\(\)](#) (default).

**Usage**

```

## S3 method for class 'TaskClust'
autoplot(object, type = "pairs", ...)

```

**Arguments**

object	( <a href="#">mlr3cluster::TaskClust</a> ).
type	(character(1)): Type of the plot. See description.
...	(any): Additional argument, passed down to the underlying geom or plot functions.

**Value**

[ggplot2::ggplot\(\)](#) object.

**Theme**

The [theme\\_ml3\(\)](#) and viridis color maps are applied by default to all autoplot() methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```

if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3cluster)
  library(mlr3viz)

  task = mlr_tasks$get("usarrests")

  head(fortify(task))
  autoplot(task)
}

```

---

autoplot.TaskRegr      *Plot for Regression Tasks*

---

**Description**

Generates plots for [mlr3::TaskRegr](#), depending on argument type:

- "target": Box plot of target variable (default).
- "pairs": Passes data and additional arguments down to [GGally::ggpairs\(\)](#). Color is set to target column.

**Usage**

```

## S3 method for class 'TaskRegr'
autoplot(object, type = "target", ...)

```

**Arguments**

object	( <a href="#">mlr3::TaskRegr</a> ).
type	(character(1)): Type of the plot. See description.
...	(any): Additional argument, passed down to the underlying geom or plot functions.

**Value**

[ggplot2::ggplot\(\)](#) object.

**Theme**

The [theme\\_ml3\(\)](#) and viridis color maps are applied by default to all autoplot() methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```

if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  task = tsk("mtcars")
  task$select(c("am", "carb"))

  head(fortify(task))
  autoplot(task)
  autoplot(task, type = "pairs")
}

```

---

```

autoplot.TuningInstanceSingleCrit
      Plot for TuningInstanceSingleCrit

```

---

**Description**

Generates plots for [mlr3tuning::TuningInstanceSingleCrit](#).

**Usage**

```

## S3 method for class 'TuningInstanceSingleCrit'
autoplot(
  object,
  type = "marginal",
  cols_x = NULL,
  trafo = FALSE,
  learner = mlr3::lrn("regr.ranger"),
  grid_resolution = 100,
  ...
)

```

**Arguments**

object	<a href="#">(mlr3tuning::TuningInstanceSingleCrit</a> .
type	(character(1)): Type of the plot. Available choices: <ul style="list-style-type: none"> <li>• "marginal": scatter plots of hyperparameter versus performance. The color of the points shows the batch number.</li> <li>• "performance": scatter plots of batch number versus performance.</li> <li>• "parameter": scatter plots of batch number versus hyperparameter. The color of the points shows the performance.</li> <li>• "parallel" parallel coordinates plot. Parameter values are rescaled by <math>(x - \text{mean}(x)) / \text{sd}(x)</math>.</li> <li>• "points" - scatter plot of two hyperparameters versus performance. The color of the points shows the performance.</li> </ul>

	<ul style="list-style-type: none"> <li>• "surface": surface plot of 2 hyperparameters versus performance. The performance values are interpolated with the supplied <code>mlr3::Learner</code>.</li> <li>• "pairs": plots all hyperparameters and performance values against each other.</li> </ul>
cols_x	(character()) Column names of hyperparameters. By default, all untransformed hyperparameters are plotted. Transformed hyperparameters are prefixed with <code>x_domain_</code> .
trafo	(logical(1)) Determines if untransformed (FALSE) or transformed (TRUE) hyperparameters are plotted.
learner	( <code>mlr3::Learner</code> ) Regression learner used to interpolate the data of the surface plot.
grid_resolution	(numeric()) Resolution of the surface plot.
...	(any): Additional arguments, possibly passed down to the underlying plot functions.

**Value**

`ggplot2::ggplot()` object.

**Theme**

The `theme_ml3()` and `viridis` color maps are applied by default to all `autoplot()` methods. To change this behavior set `options(mlr3.theme = FALSE)`.

**Examples**

```
if (requireNamespace("mlr3tuning") && requireNamespace("patchwork")) {
  library(mlr3tuning)

  learner = lrn("classif.rpart")
  learner$param_set$values$cp = to_tune(0.001, 0.1)
  learner$param_set$values$minsplit = to_tune(1, 10)

  instance = TuningInstanceSingleCrit$new(
    task = tsk("iris"),
    learner = learner,
    resampling = rsm("holdout"),
    measure = msr("classif.ce"),
    terminator = trm("evals", n_evals = 10))

  tuner = tnr("random_search")

  tuner$optimize(instance)

  # plot performance versus batch number
  autoplot(instance, type = "performance")
}
```

```

# plot cp values versus performance
autoplot(instance, type = "marginal", cols_x = "cp")

# plot transformed parameter values versus batch number
autoplot(instance, type = "parameter", trafo = TRUE)

# plot parallel coordinates plot
autoplot(instance, type = "parallel")

# plot pairs
autoplot(instance, type = "pairs")
}

```

---

plot\_learner\_prediction

*Plot for Learner Predictions*


---

## Description

Generates a plot for the [mlr3::Prediction](#) of a single [mlr3::Learner](#) on a single [mlr3::Task](#).

- For classification we support tasks with exactly two features and learners with `predict_type` set to "response" or "prob".
- For regression we support tasks with one or two features. For tasks with one feature we print confidence bounds if the `predict_type` of the learner was set to "se". For tasks with two features the `predict_type` will be ignored.

Note that this function is a wrapper around `autoplot.ResampleResult()` for a temporary [mlr3::ResampleResult](#) using [mlr3::mlr\\_resamplings\\_holdout](#) with ratio 1 (all observations in training set).

## Usage

```
plot_learner_prediction(learner, task, grid_points = 100L, expand_range = 0)
```

## Arguments

learner	( <a href="#">mlr3::Learner</a> ).
task	( <a href="#">mlr3::Task</a> ).
grid_points	( <code>integer(1)</code> ) Resolution of the grid. For factors, ordered and logicals this value is ignored.
expand_range	( <code>numeric(1)</code> ) Expand the prediction range for numerical features.

## Value

`ggplot2::ggplot()` object.

**Examples**

```

if (requireNamespace("mlr3")) {
  library(mlr3)
  library(mlr3viz)

  task = mlr3::tsk("pima")$select(c("age", "glucose"))
  learner = lrn("classif.rpart", predict_type = "prob")
  p = plot_learner_prediction(learner, task)
  print(p)
}

```

---

predict_grid	<i>Generates a data.table of evenly distributed points.</i>
--------------	---

---

**Description**

For each point we have the predicted class / regression value in column response. If the learner predicts probabilities, a column ".prob.response" is added that contains the probability of the predicted class

**Usage**

```
predict_grid(learners, task, grid_points, expand_range)
```

**Arguments**

learners	list of trained learners, each learner belongs to one resampling iteration
task	the task all learners are trained on
grid_points	(int): see sequenize
expand_range	see sequenize

---

theme_mlr3	<i>mlr-org ggplot2 theme</i>
------------	------------------------------

---

**Description**

The theme is heavily influenced and partly based on `ggpubr::theme_pubr()`. This theme is applied by default to all `autoplot()` methods in the mlr3 ecosystem. If you do not like it and want to use the default ggplot2 theme, you can add `+ theme_gray()` to the `autoplot()` call.



**Usage**

```
theme_mlr3(  
  base_size = 12,  
  base_family = "",  
  border = FALSE,  
  margin = TRUE,  
  legend = c("top", "bottom", "left", "right", "none"),  
  x.text.angle = 0  
)
```

**Arguments**

base_size	[integer]	Text font size.
base_family	[character]	Font family.
border	[logical]	If TRUE, adds a panel border.
margin	[logical]	If FALSE, reduces the plot margin(s).
legend	[character]	Specifies the legend position. Allowed values are one of c("top", "bottom", "left", "right", "none"). Default is "top".
x.text.angle	[numeric]	Rotation angle of x axis tick labels. Default value is 0. Use 90 for vertical text.

**Examples**

```
library("ggplot2")  
p = ggplot(mtcars, aes(x = wt, y = mpg)) +  
  geom_point(aes(color = gear))  
  
# Default plot  
p  
  
# theme_mlr3()  
p + theme_mlr3()
```

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