

# Package ‘modeltuning’

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**Title** Model Selection and Tuning Utilities

**Version** 0.1.2

**Description** Provides a lightweight framework for model selection and hyperparameter tuning in R. The package offers intuitive tools for grid search, cross-validation, and combined grid search with cross-validation that work seamlessly with virtually any modeling package. Designed for flexibility and ease of use, it standardizes tuning workflows while remaining fully compatible with a wide range of model interfaces and estimation functions.

**License** MIT + file LICENSE

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**CV***Predictive Models with Cross Validation***Description**

CV allows the user to specify a cross validation scheme with complete flexibility in the model, data splitting function, and performance metrics, among other essential parameters.

**Public fields**

- learner Predictive modeling function.
- scorer List of performance metric functions.
- splitter Function that splits data into cross validation folds.

**Methods****Public methods:**

- [CV\\$fit\(\)](#)
- [CV\\$new\(\)](#)
- [CV\\$clone\(\)](#)

**Method fit():** fit performs cross validation with user-specified parameters.

*Usage:*

```
CV$fit(
  formula = NULL,
  data = NULL,
  x = NULL,
  y = NULL,
  response = NULL,
  convert_response = NULL,
  progress = FALSE
)
```

*Arguments:*

**formula** An object of class [formula](#): a symbolic description of the model to be fitted.

**data** An optional data frame, or other object containing the variables in the model. If data is not provided, how formula is handled depends on \$learner.

**x** Predictor data (independent variables), alternative interface to data with formula.

**y** Response vector (dependent variable), alternative interface to data with formula.

**response** String; In the absence of formula or y, this specifies which element of learner\_args is the response vector.

`convert_response` Function; This should be a single function that transforms the response vector. E.g. a function converting a numeric binary variable to a factor variable.

`progress` Logical; indicating whether to print progress across cross validation folds.

*Details:* `fit` follows standard R modeling convention by surfacing a formula modeling interface as well as an alternate matrix option. The user should use whichever interface is supported by the specified `$learner` function.

*Returns:* An object of class [FittedCV](#).

*Examples:*

```
if (require(e1071) && require(rpart) && require(yardstick)) {
  iris_new <- iris[sample(1:nrow(iris), nrow(iris)), ]
  iris_new$Species <- factor(iris_new$Species == "virginica")

  ### Decision Tree Example

  iris_cv <- CV$new(
    learner = rpart::rpart,
    learner_args = list(method = "class"),
    splitter = cv_split,
    scorer = list(accuracy = yardstick::accuracy_vec),
    prediction_args = list(accuracy = list(type = "class")))
  )
  iris_cv_fitted <- iris_cv$fit(formula = Species ~ ., data = iris_new)

  ### Example with multiple metric functions

  iris_cv <- CV$new(
    learner = rpart::rpart,
    learner_args = list(method = "class"),
    splitter = cv_split,
    splitter_args = list(v = 3),
    scorer = list(
      f_meas = yardstick::f_meas_vec,
      accuracy = yardstick::accuracy_vec,
      roc_auc = yardstick::roc_auc_vec,
      pr_auc = yardstick::pr_auc_vec
    ),
    prediction_args = list(
      f_meas = list(type = "class"),
      accuracy = list(type = "class"),
      roc_auc = list(type = "prob"),
      pr_auc = list(type = "prob")
    ),
    convert_predictions = list(
      f_meas = NULL,
      accuracy = NULL,
      roc_auc = function(i) i[, "FALSE"],
      pr_auc = function(i) i[, "FALSE"]
    )
  )
```

```

    )
)
iris_cv_fitted <- iris_cv$fit(formula = Species ~ ., data = iris_new)

# Print the mean performance metrics across CV folds
iris_cv_fitted$mean_metrics

# Grab the final model fitted on the full dataset
iris_cv_fitted$model

#### OLS Example

mtcars_cv <- CV$new(
  learner = lm,
  splitter = cv_split,
  splitter_args = list(v = 2),
  scorer = list("rmse" = yardstick::rmse_vec, "mae" = yardstick::mae_vec)
)

mtcars_cv_fitted <- mtcars_cv$fit(
  formula = mpg ~ .,
  data = mtcars
)

#### Matrix interface example - SVM

mtcars_x <- model.matrix(mpg ~ . - 1, mtcars)
mtcars_y <- mtcars$mpg

mtcars_cv <- CV$new(
  learner = e1071::svm,
  learner_args = list(scale = TRUE, kernel = "polynomial", cross = 0),
  splitter = cv_split,
  splitter_args = list(v = 3),
  scorer = list(rmse = yardstick::rmse_vec, mae = yardstick::mae_vec)
)
mtcars_cv_fitted <- mtcars_cv$fit(
  x = mtcars_x,
  y = mtcars_y
)
}

```

**Method new():** Create a new [CV](#) object.

*Usage:*

```
CV$new(
  learner = NULL,
  splitter = NULL,
  scorer = NULL,
```

```

    learner_args = NULL,
    splitter_args = NULL,
    scorer_args = NULL,
    prediction_args = NULL,
    convert_predictions = NULL
)

```

*Arguments:*

**learner** Function that estimates a predictive model. It is essential that this function support either a formula interface with `formula` and `data` arguments, or an alternate matrix interface with `x` and `y` arguments.

**splitter** A function that computes cross validation folds from an input data set or a pre-computed list of cross validation fold indices. If `splitter` is a function, it must have a `data` argument for the input data, and it must return a list of cross validation fold indices. If `splitter` is a list of integers, the number of cross validation folds is `length(splitter)` and each element contains the indices of the data observations that are included in that fold.

**scorer** A named list of metric functions to evaluate model performance on each cross validation fold. Any provided metric function must have `truth` and `estimate` arguments for true outcome values and predicted outcome values respectively, and must return a single numeric metric value.

**learner\_args** A named list of additional arguments to pass to `learner`.

**splitter\_args** A named list of additional arguments to pass to `splitter`.

**scorer\_args** A named list of additional arguments to pass to `scorer`. `scorer_args` must either be length 1 or `length(scoring)` in the case where different arguments are being passed to each scoring function.

**prediction\_args** A named list of additional arguments to pass to `predict`. `prediction_args` must either be length 1 or `length(scoring)` in the case where different arguments are being passed to each scoring function.

**convert\_predictions** A list of functions to convert predicted values prior to being evaluated by the metric functions supplied in `scorer`. This list should either be length 1, in which case the same function will be applied to all predicted values, or `length(scoring)` in which case each function in `convert_predictions` will correspond with each function in `scorer`.

**Returns:** An object of class **CV**.

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

*Usage:*

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

*Arguments:*

`deep` Whether to make a deep clone.

## Examples

```

## -----
## Method `CV$fit`
## -----
if (require(e1071) && require(rpart) && require(yardstick)) {

```

```

iris_new <- iris[sample(1:nrow(iris), nrow(iris)), ]
iris_new$Species <- factor(iris_new$Species == "virginica")

### Decision Tree Example

iris_cv <- CV$new(
  learner = rpart::rpart,
  learner_args = list(method = "class"),
  splitter = cv_split,
  scorer = list(accuracy = yardstick::accuracy_vec),
  prediction_args = list(accuracy = list(type = "class"))
)
iris_cv_fitted <- iris_cv$fit(formula = Species ~ ., data = iris_new)

### Example with multiple metric functions

iris_cv <- CV$new(
  learner = rpart::rpart,
  learner_args = list(method = "class"),
  splitter = cv_split,
  splitter_args = list(v = 3),
  scorer = list(
    f_meas = yardstick::f_meas_vec,
    accuracy = yardstick::accuracy_vec,
    roc_auc = yardstick::roc_auc_vec,
    pr_auc = yardstick::pr_auc_vec
  ),
  prediction_args = list(
    f_meas = list(type = "class"),
    accuracy = list(type = "class"),
    roc_auc = list(type = "prob"),
    pr_auc = list(type = "prob")
  ),
  convert_predictions = list(
    f_meas = NULL,
    accuracy = NULL,
    roc_auc = function(i) i[, "FALSE"],
    pr_auc = function(i) i[, "FALSE"]
  )
)
iris_cv_fitted <- iris_cv$fit(formula = Species ~ ., data = iris_new)

# Print the mean performance metrics across CV folds
iris_cv_fitted$mean_metrics

# Grab the final model fitted on the full dataset
iris_cv_fitted$model

### OLS Example

mtcars_cv <- CV$new(
  learner = lm,
  splitter = cv_split,

```

```

splitter_args = list(v = 2),
scorer = list("rmse" = yardstick::rmse_vec, "mae" = yardstick::mae_vec)
)

mtcars_cv_fitted <- mtcars_cv$fit(
  formula = mpg ~ .,
  data = mtcars
)

### Matrix interface example - SVM

mtcars_x <- model.matrix(mpg ~ . - 1, mtcars)
mtcars_y <- mtcars$mpg

mtcars_cv <- CV$new(
  learner = e1071::svm,
  learner_args = list(scale = TRUE, kernel = "polynomial", cross = 0),
  splitter = cv_split,
  splitter_args = list(v = 3),
  scorer = list(rmse = yardstick::rmse_vec, mae = yardstick::mae_vec)
)
mtcars_cv_fitted <- mtcars_cv$fit(
  x = mtcars_x,
  y = mtcars_y
)
}

```

**cv\_split***Generate cross-validation fold indices***Description**

Splits row indices of a data frame or matrix into k folds for cross-validation.

**Usage**

```
cv_split(data, v = 5, seed = NULL)
```

**Arguments**

- |      |  |
|------|--|
| data | A data frame or matrix.                            |
| v    | Integer. Number of folds. Defaults to 5.           |
| seed | Optional integer. Random seed for reproducibility. |

**Value**

A list of length v, where each element is a vector of row indices for that fold.

## Examples

```
folds <- cv_split(mtcars, v = 5)
str(folds)
```

**FittedCV**

*Fitted, Cross-Validated Predictive Models*

## Description

**FittedCV** is a fitted, cross-validated predictive model object that is returned by `CV$fit()` and contains relevant model components, cross-validation metrics, validation set predicted values, etc.

## Public fields

`folds` A list of length `$nfolds` where each element contains the indices of the observations contained in that fold.  
`model` Predictive model fitted on the full data set.  
`mean_metrics` Numeric list; Cross-validation performance metrics averaged across folds.  
`metrics` Numeric list; Cross-validation performance metrics on each fold.  
`nfolds` An integer specifying the number of cross-validation folds.  
`predictions` A list containing the predicted hold-out values on every fold.

## Methods

### Public methods:

- [FittedCV\\$new\(\)](#)
- [FittedCV\\$clone\(\)](#)

**Method new():** Create a new [\*\*FittedCV\*\*](#) object.

*Usage:*

```
FittedCV$new(folds, model, metrics, nfolds, predictions)
```

*Arguments:*

`folds` A list of length `$nfolds` where each element contains the indices of the observations contained in that fold.  
`model` Predictive model fitted on the full data set.  
`metrics` Numeric list; Cross-validation performance metrics on each fold.  
`nfolds` An integer specifying the number of cross-validation folds.  
`predictions` A list containing the predicted hold-out values on every fold.  
*Returns:* An object of class [\*\*FittedCV\*\*](#).

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

*Usage:*

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

*Arguments:*

`deep` Whether to make a deep clone.

---

**FittedGridSearch***Fitted Models across a Tuning Grid of Hyper-parameters*

---

**Description**

`FittedGridSearch` is an object containing fitted predictive models across a tuning grid of hyper-parameters returned by `GridSearch$fit()` as well as relevant model information such as the best performing model, best hyper-parameters, etc.

**Public fields**

`best_idx` An integer specifying the index of `$models` that contains the best-performing model.  
`best_metric` The performance metric of the best model on the validation data.  
`best_model` The best performing predictive model.  
`best_params` A named list of the hyper-parameters that result in the optimal predictive model.  
`tune_params` Data.frame of the full hyper-parameter grid.  
`models` List of predictive models at every value of `$tune_params`.  
`metrics` Numeric list; Cross-validation performance metrics on each fold.  
`predictions` A list containing the predicted hold-out values on every fold.

**Methods****Public methods:**

- `FittedGridSearch$new()`
- `FittedGridSearch$clone()`

**Method** `new()`: Create a new `FittedGridSearch` object.

*Usage:*

`FittedGridSearch$new(tune_params, models, metrics, predictions, optimize_score)`

*Arguments:*

`tune_params` Data.frame of the full hyper-parameter grid.

`models` List of predictive models at every value of `$tune_params`.

`metrics` List of performance metrics on the validation data for every model in `$models`.

`predictions` A list containing the predicted values on the validation data for every model in `$models`.

`optimize_score` Either "max" or "min" indicating whether or not the specified performance metric was maximized or minimized to find the optimal predictive model.

*Returns:* An object of class `FittedGridSearch`.

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

*Usage:*

`FittedGridSearch$clone(deep = FALSE)`

*Arguments:*

`deep` Whether to make a deep clone.

---

FittedGridSearchCV	<i>Fitted Models with Cross Validation across a Tuning Grid of Hyper-parameters</i>
--------------------	---

---

## Description

`FittedGridSearchCV` is an object containing fitted predictive models across a tuning grid of hyper-parameters returned by `GridSearchCV$fit()` as well as relevant model information such as the best performing model, best hyper-parameters, etc.

## Public fields

- `best_idx` An integer specifying the index of `$models` that contains the best-performing model.
- `best_metric` The average performance metric of the best model across cross-validation folds.
- `best_model` The best performing predictive model.
- `best_params` A named list of the hyper-parameters that result in the optimal predictive model.
- `folds` A list of length `$models` where each element contains a list of the cross-validation indices for each fold.
- `tune_params` A [data.frame](#) of the full hyper-parameter grid.
- `models` List of predictive models at every value of `$tune_params`.
- `metrics` Numeric list; Cross-validation performance metrics for every model in `$models`.
- `predictions` A list containing the cross-validation fold predictions for each model in `$models`.

## Methods

### Public methods:

- [FittedGridSearchCV\\$new\(\)](#)
- [FittedGridSearchCV\\$clone\(\)](#)

**Method** `new()`: Create a new [FittedGridSearchCV](#) object.

*Usage:*

```
FittedGridSearchCV$new(
  tune_params,
  models,
  folds,
  metrics,
  predictions,
  optimize_score
)
```

*Arguments:*

- `tune_params` Data.frame of the full hyper-parameter grid.
- `models` List of predictive models at every value of `$tune_params`.
- `folds` List of cross-validation indices at every value of `$tune_params`.

`metrics` List of cross-validation performance metrics for every model in `$models`.  
`predictions` A list containing the predicted values on the cross-validation folds for every model in `$models`.  
`optimize_score` Either "max" or "min" indicating whether or not the specified performance metric was maximized or minimized to find the optimal predictive model.

*Returns:* An object of class [FittedGridSearchCV](#).

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

*Usage:*

`FittedGridSearchCV$clone(deep = FALSE)`

*Arguments:*

`deep` Whether to make a deep clone.

## Description

GridSearch allows the user to specify a Grid Search schema for tuning predictive model hyper-parameters with complete flexibility in the predictive model and performance metrics.

## Public fields

`learner` Predictive modeling function.  
`scorer` List of performance metric functions.  
`tune_params` Data.frame of full hyper-parameter grid created from `$tune_params`

## Methods

### Public methods:

- [GridSearch\\$fit\(\)](#)
- [GridSearch\\$new\(\)](#)
- [GridSearch\\$clone\(\)](#)

**Method** `fit()`: fit tunes user-specified model hyper-parameters via Grid Search.

*Usage:*

```
GridSearch$fit(
  formula = NULL,
  data = NULL,
  x = NULL,
  y = NULL,
  progress = FALSE
)
```

*Arguments:*

**formula** An object of class **formula**: a symbolic description of the model to be fitted.  
**data** An optional data frame, or other object containing the variables in the model. If **data** is not provided, how **formula** is handled depends on **\$learner**.  
**x** Predictor data (independent variables), alternative interface to **data** with **formula**.  
**y** Response vector (dependent variable), alternative interface to **data** with **formula**.  
**progress** Logical; indicating whether to print progress across cross validation folds.

**Details:** **fit** follows standard R modeling convention by surfacing a formula modeling interface as well as an alternate matrix option. The user should use whichever interface is supported by the specified **\$learner** function.

**Returns:** An object of class **FittedGridSearch**.

**Examples:**

```
if (require(e1071) && require(rpart) && require(yardstick)) {
  iris_new <- iris[sample(1:nrow(iris), nrow(iris)), ]
  iris_new$Species <- factor(iris_new$Species == "virginica")
  iris_train <- iris_new[1:100, ]
  iris_validate <- iris_new[101:150, ]

  ### Decision Tree example

  iris_grid <- GridSearch$new(
    learner = rpart::rpart,
    learner_args = list(method = "class"),
    tune_params = list(
      minsplit = seq(10, 30, by = 5),
      maxdepth = seq(20, 30, by = 2)
    ),
    evaluation_data = list(x = iris_validate[, 1:4], y = iris_validate$Species),
    scorer = list(accuracy = yardstick::accuracy_vec),
    optimize_score = "max",
    prediction_args = list(accuracy = list(type = "class"))
  )
  iris_grid_fitted <- iris_grid$fit(
    formula = Species ~ .,
    data = iris_train
  )

  ### Example with multiple metric functions

  iris_grid <- GridSearch$new(
    learner = rpart::rpart,
    learner_args = list(method = "class"),
    tune_params = list(
      minsplit = seq(10, 30, by = 5),
      maxdepth = seq(20, 30, by = 2)
    ),
    evaluation_data = list(x = iris_validate, y = iris_validate$Species),
    scorer = list(

```

```
accuracy = yardstick::accuracy_vec,
auc = yardstick::roc_auc_vec
),
optimize_score = "max",
prediction_args = list(
  accuracy = list(type = "class"),
  auc = list(type = "prob")
),
convert_predictions = list(
  accuracy = NULL,
  auc = function(i) i[, "FALSE"]
)
)
iris_grid_fitted <- iris_grid$fit(
  formula = Species ~ .,
  data = iris_train,
)

# Grab the best model
iris_grid_fitted$best_model

# Grab the best hyper-parameters
iris_grid_fitted$best_params

# Grab the best model performance metrics
iris_grid_fitted$best_metric

### Matrix interface example - SVM

mtcars_train <- mtcars[1:25, ]
mtcars_eval <- mtcars[26:nrow(mtcars), ]

mtcars_grid <- GridSearch$new(
  learner = e1071::svm,
  tune_params = list(
    degree = 2:4,
    kernel = c("linear", "polynomial")
  ),
  evaluation_data = list(x = mtcars_eval[, -1], y = mtcars_eval$mpg),
  learner_args = list(scale = TRUE),
  scorer = list(
    rmse = yardstick::rmse_vec,
    mae = yardstick::mae_vec
  ),
  optimize_score = "min"
)
mtcars_grid_fitted <- mtcars_grid$fit(
  x = mtcars_train[, -1],
```

```

    y = mtcars_train$mpg
)
}

}

```

**Method new():** Create a new [GridSearch](#) object.

*Usage:*

```

GridSearch$new(
  learner = NULL,
  tune_params = NULL,
  evaluation_data = NULL,
  scorer = NULL,
  optimize_score = c("min", "max"),
  learner_args = NULL,
  scorer_args = NULL,
  prediction_args = NULL,
  convert_predictions = NULL
)

```

*Arguments:*

**learner** Function that estimates a predictive model. It is essential that this function support either a formula interface with `formula` and `data` arguments, or an alternate matrix interface with `x` and `y` arguments.

**tune\_params** A named list specifying the arguments of `$learner` to tune.

**evaluation\_data** A two-element list containing the following elements: `x`, the validation data to generate predicted values with; `y`, the validation response values to evaluate predictive performance.

**scorer** A named list of metric functions to evaluate model performance on `evaluation_data`.

Any provided metric function must have `truth` and `estimate` arguments, for true outcome values and predicted outcome values respectively, and must return a single numeric metric value. The last metric function will be the one used to identify the optimal model from the Grid Search.

**optimize\_score** One of "max" or "min"; Whether to maximize or minimize the metric defined in `scorer` to find the optimal Grid Search parameters.

**learner\_args** A named list of additional arguments to pass to `learner`.

**scorer\_args** A named list of additional arguments to pass to `scorer`. `scorer_args` must either be length 1 or `length(scorer)` in the case where different arguments are being passed to each scoring function.

**prediction\_args** A named list of additional arguments to pass to `predict`. `prediction_args` must either be length 1 or `length(scorer)` in the case where different arguments are being passed to each scoring function.

**convert\_predictions** A list of functions to convert predicted values prior to being evaluated by the metric functions supplied in `scorer`. This list should either be length 1, in which case the same function will be applied to all predicted values, or `length(scorer)` in which case each function in `convert_predictions` will correspond with each function in `scorer`.

*Returns:* An object of class [GridSearch](#).

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

*Usage:*

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

*Arguments:*

deep Whether to make a deep clone.

## Examples

```
## -----
## Method `GridSearch$fit`
## -----
```

```
if (require(e1071) && require(rpart) && require(yardstick)) {
  iris_new <- iris[sample(1:nrow(iris), nrow(iris)), ]
  iris_new$Species <- factor(iris_new$Species == "virginica")
  iris_train <- iris_new[1:100, ]
  iris_validate <- iris_new[101:150, ]
```

```
### Decision Tree example
```

```
iris_grid <- GridSearch$new(
  learner = rpart::rpart,
  learner_args = list(method = "class"),
  tune_params = list(
    minsplit = seq(10, 30, by = 5),
    maxdepth = seq(20, 30, by = 2)
  ),
  evaluation_data = list(x = iris_validate[, 1:4], y = iris_validate$Species),
  scorer = list(accuracy = yardstick::accuracy_vec),
  optimize_score = "max",
  prediction_args = list(accuracy = list(type = "class")))
)
iris_grid_fitted <- iris_grid$fit(
  formula = Species ~ .,
  data = iris_train
)
### Example with multiple metric functions
```

```
iris_grid <- GridSearch$new(
  learner = rpart::rpart,
  learner_args = list(method = "class"),
  tune_params = list(
    minsplit = seq(10, 30, by = 5),
    maxdepth = seq(20, 30, by = 2)
  ),
  evaluation_data = list(x = iris_validate, y = iris_validate$Species),
  scorer = list(
    accuracy = yardstick::accuracy_vec,
    auc = yardstick::roc_auc_vec
  ),
  optimize_score = "max",
  prediction_args = list(
```

```

accuracy = list(type = "class"),
auc = list(type = "prob")
),
convert_predictions = list(
  accuracy = NULL,
  auc = function(i) i[, "FALSE"]
)
)
iris_grid_fitted <- iris_grid$fit(
  formula = Species ~ .,
  data = iris_train,
)

# Grab the best model
iris_grid_fitted$best_model

# Grab the best hyper-parameters
iris_grid_fitted$best_params

# Grab the best model performance metrics
iris_grid_fitted$best_metric

### Matrix interface example - SVM

mtcars_train <- mtcars[1:25, ]
mtcars_eval <- mtcars[26:nrow(mtcars), ]

mtcars_grid <- GridSearch$new(
  learner = e1071::svm,
  tune_params = list(
    degree = 2:4,
    kernel = c("linear", "polynomial")
  ),
  evaluation_data = list(x = mtcars_eval[, -1], y = mtcars_eval$mpg),
  learner_args = list(scale = TRUE),
  scorer = list(
    rmse = yardstick::rmse_vec,
    mae = yardstick::mae_vec
  ),
  optimize_score = "min"
)
mtcars_grid_fitted <- mtcars_grid$fit(
  x = mtcars_train[, -1],
  y = mtcars_train$mpg
)

}

}

```

## Description

GridSearchCV allows the user to specify a Grid Search schema for tuning predictive model hyper-parameters with Cross-Validation. GridSearchCV gives the user complete flexibility in the predictive model and performance metrics.

## Public fields

- `learner` Predictive modeling function.
- `scorer` List of performance metric functions.
- `splitter` Function that splits data into cross validation folds.
- `tune_params` Data.frame of full hyper-parameter grid created from `$tune_params`

## Methods

### Public methods:

- `GridSearchCV$fit()`
- `GridSearchCV$new()`
- `GridSearchCV$clone()`

**Method fit():** fit tunes user-specified model hyper-parameters via Grid Search and Cross-Validation.

*Usage:*

```
GridSearchCV$fit(
  formula = NULL,
  data = NULL,
  x = NULL,
  y = NULL,
  response = NULL,
  convert_response = NULL,
  progress = FALSE
)
```

*Arguments:*

`formula` An object of class `formula`: a symbolic description of the model to be fitted.

`data` An optional data frame, or other object containing the variables in the model. If `data` is not provided, how `formula` is handled depends on `$learner`.

`x` Predictor data (independent variables), alternative interface to `data` with `formula`.

`y` Response vector (dependent variable), alternative interface to `data` with `formula`.

`response` String; In the absence of `formula` or `y`, this specifies which element of `learner_args` is the response vector.

`convert_response` Function; This should be a single function that transforms the response vector. E.g. a function converting a numeric binary variable to a factor variable.

`progress` Logical; indicating whether to print progress across the hyper-parameter grid.

*Details:* `fit` follows standard R modeling convention by surfacing a formula modeling interface as well as an alternate matrix option. The user should use whichever interface is supported by the specified `$learner` function.

*Returns:* An object of class [FittedGridSearchCV](#).

*Examples:*

```
\donttest{
if (require(e1071) && require(rpart) && require(yardstick)) {
  iris_new <- iris[sample(1:nrow(iris), nrow(iris)), ]
  iris_new$Species <- factor(iris_new$Species == "virginica")
  iris_train <- iris_new[1:100, ]
  iris_validate <- iris_new[101:150, ]

  ### Decision Tree example

  iris_grid_cv <- GridSearchCV$new(
    learner = rpart::rpart,
    learner_args = list(method = "class"),
    tune_params = list(
      minsplit = seq(10, 30, by = 5),
      maxdepth = seq(20, 30, by = 2)
    ),
    splitter = cv_split,
    splitter_args = list(v = 3),
    scorer = list(accuracy = yardstick::accuracy_vec),
    optimize_score = "max",
    prediction_args = list(accuracy = list(type = "class")))
  )
  iris_grid_cv_fitted <- iris_grid_cv$fit(
    formula = Species ~.,
    data = iris_train
  )

  ### Example with multiple metric functions

  iris_grid_cv <- GridSearchCV$new(
    learner = rpart::rpart,
    learner_args = list(method = "class"),
    tune_params = list(
      minsplit = seq(10, 30, by = 5),
      maxdepth = seq(20, 30, by = 2)
    ),
    splitter = cv_split,
    splitter_args = list(v = 3),
    scorer = list(
      accuracy = yardstick::accuracy_vec,
      auc = yardstick::roc_auc_vec
    ),
    optimize_score = "max",
    prediction_args = list(
      accuracy = list(type = "class"),
      auc = list(type = "prob"))
  )
}
```

```

),
convert_predictions = list(
    accuracy = NULL,
    auc = function(i) i[, "FALSE"]
)
)
iris_grid_cv_fitted <- iris_grid_cv$fit(
    formula = Species ~ .,
    data = iris_train,
)
# Grab the best model
iris_grid_cv_fitted$best_model

# Grab the best hyper-parameters
iris_grid_cv_fitted$best_params

# Grab the best model performance metrics
iris_grid_cv_fitted$best_metric

### Matrix interface example - SVM

mtcars_train <- mtcars[1:25, ]
mtcars_eval <- mtcars[26:nrow(mtcars), ]

mtcars_grid_cv <- GridSearchCV$new(
    learner = e1071::svm,
    tune_params = list(
        degree = 2:4,
        kernel = c("linear", "polynomial")
    ),
    splitter = cv_split,
    splitter_args = list(v = 2),
    learner_args = list(scale = TRUE),
    scorer = list(
        rmse = yardstick::rmse_vec,
        mae = yardstick::mae_vec
    ),
    optimize_score = "min"
)
mtcars_grid_cv_fitted <- mtcars_grid_cv$fit(
    x = mtcars_train[, -1],
    y = mtcars_train$mpg
)
}

}

```

**Method new():** Create a new [GridSearchCV](#) object.

*Usage:*

```
GridSearchCV$new(
  learner = NULL,
  tune_params = NULL,
  splitter = NULL,
  scorer = NULL,
  optimize_score = c("min", "max"),
  learner_args = NULL,
  splitter_args = NULL,
  scorer_args = NULL,
  prediction_args = NULL,
  convert_predictions = NULL
)
```

*Arguments:*

**learner** Function that estimates a predictive model. It is essential that this function support either a formula interface with `formula` and `data` arguments, or an alternate matrix interface with `x` and `y` arguments.

**tune\_params** A named list specifying the arguments of `$learner` to tune.

**splitter** A function that computes cross validation folds from an input data set or a pre-computed list of cross validation fold indices. If `splitter` is a function, it must have a `data` argument for the input data, and it must return a list of cross validation fold indices. If `splitter` is a list of integers, the number of cross validation folds is `length(splitter)` and each element contains the indices of the data observations that are included in that fold.

**scorer** A named list of metric functions to evaluate model performance on `evaluation_data`. Any provided metric function must have `truth` and `estimate` arguments, for true outcome values and predicted outcome values respectively, and must return a single numeric metric value. The last metric function will be the one used to identify the optimal model from the Grid Search.

**optimize\_score** One of "max" or "min"; Whether to maximize or minimize the metric defined in `scorer` to find the optimal Grid Search parameters.

**learner\_args** A named list of additional arguments to pass to `learner`.

**splitter\_args** A named list of additional arguments to pass to `splitter`.

**scorer\_args** A named list of additional arguments to pass to `scorer`. `scorer_args` must either be length 1 or `length(scoring)` in the case where different arguments are being passed to each scoring function.

**prediction\_args** A named list of additional arguments to pass to `predict`. `prediction_args` must either be length 1 or `length(scoring)` in the case where different arguments are being passed to each scoring function.

**convert\_predictions** A list of functions to convert predicted values prior to being evaluated by the metric functions supplied in `scorer`. This list should either be length 1, in which case the same function will be applied to all predicted values, or `length(scoring)` in which case each function in `convert_predictions` will correspond with each function in `scorer`.

*Returns:* An object of class [GridSearch](#).

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

*Usage:*

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

*Arguments:*

deep Whether to make a deep clone.

## Examples

```
## -----
## Method `GridSearchCV$fit`
## -----
```

  

```
if (require(e1071) && require(rpart) && require(yardstick)) {
  iris_new <- iris[sample(1:nrow(iris), nrow(iris)), ]
  iris_new$Species <- factor(iris_new$Species == "virginica")
  iris_train <- iris_new[1:100, ]
  iris_validate <- iris_new[101:150, ]
```

  

```
### Decision Tree example
```

  

```
iris_grid_cv <- GridSearchCV$new(
  learner = rpart::rpart,
  learner_args = list(method = "class"),
  tune_params = list(
    minsplit = seq(10, 30, by = 5),
    maxdepth = seq(20, 30, by = 2)
  ),
  splitter = cv_split,
  splitter_args = list(v = 3),
  scorer = list(accuracy = yardstick::accuracy_vec),
  optimize_score = "max",
  prediction_args = list(accuracy = list(type = "class")))
iris_grid_cv_fitted <- iris_grid_cv$fit(
  formula = Species ~ .,
  data = iris_train
)
### Example with multiple metric functions
```

  

```
iris_grid_cv <- GridSearchCV$new(
  learner = rpart::rpart,
  learner_args = list(method = "class"),
  tune_params = list(
    minsplit = seq(10, 30, by = 5),
    maxdepth = seq(20, 30, by = 2)
  ),
  splitter = cv_split,
  splitter_args = list(v = 3),
  scorer = list(
    accuracy = yardstick::accuracy_vec,
    auc = yardstick::roc_auc_vec
  ),
```

```

    optimize_score = "max",
    prediction_args = list(
        accuracy = list(type = "class"),
        auc = list(type = "prob")
    ),
    convert_predictions = list(
        accuracy = NULL,
        auc = function(i) i[, "FALSE"]
    )
)
iris_grid_cv_fitted <- iris_grid_cv$fit(
    formula = Species ~ .,
    data = iris_train,
)

# Grab the best model
iris_grid_cv_fitted$best_model

# Grab the best hyper-parameters
iris_grid_cv_fitted$best_params

# Grab the best model performance metrics
iris_grid_cv_fitted$best_metric

### Matrix interface example - SVM

mtcars_train <- mtcars[1:25, ]
mtcars_eval <- mtcars[26:nrow(mtcars), ]

mtcars_grid_cv <- GridSearchCV$new(
    learner = e1071::svm,
    tune_params = list(
        degree = 2:4,
        kernel = c("linear", "polynomial")
    ),
    splitter = cv_split,
    splitter_args = list(v = 2),
    learner_args = list(scale = TRUE),
    scorer = list(
        rmse = yardstick::rmse_vec,
        mae = yardstick::mae_vec
    ),
    optimize_score = "min"
)
mtcars_grid_cv_fitted <- mtcars_grid_cv$fit(
    x = mtcars_train[, -1],
    y = mtcars_train$mpg
)
}
```

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