

# Package ‘agua’

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**Title** 'tidymodels' Integration with 'h2o'

**Version** 0.1.4

**Description** Create and evaluate models using 'tidymodels' and 'h2o' <<https://h2o.ai/>>. The package enables users to specify 'h2o' as an engine for several modeling methods.

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**URL** <https://agua.tidymodels.org/>, <https://github.com/tidymodels/agua>

**BugReports** <https://github.com/tidymodels/agua/issues>

**Depends** parsnip

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agua\_backend\_options    *Control model tuning via `h2o::h2o.grid()`*

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### Description

Control model tuning via `h2o::h2o.grid()`

### Usage

```
agua_backend_options(parallelism = 1)
```

### Arguments

`parallelism`    Level of Parallelism during grid model building. 1 = sequential building (default). Use the value of 0 for adaptive parallelism - decided by H2O. Any number > 1 sets the exact number of models built in parallel.

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as\_h2o                    *Data conversion tools*

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### Description

Data conversion tools

### Usage

```
as_h2o(df, destination_frame_prefix = "object")

## S3 method for class 'H2OFrame'
as_tibble(
  x,
  ...,
  .rows = NULL,
  .name_repair = c("check_unique", "unique", "universal", "minimal"),
  rownames = pkgconfig::get_config("tibble::rownames", NULL)
)
```

**Arguments**

df	A R data frame.
destination_frame_prefix	A character string to use as the base name.
x	An H2OFrame.
...	Unused, for extensibility.
.rows	The number of rows, useful to create a 0-column tibble or just as an additional check.
.name_repair	<p>Treatment of problematic column names:</p> <ul style="list-style-type: none"> <li>• "minimal": No name repair or checks, beyond basic existence,</li> <li>• "unique": Make sure names are unique and not empty,</li> <li>• "check_unique": (default value), no name repair, but check they are unique,</li> <li>• "universal": Make the names unique and syntactic</li> <li>• a function: apply custom name repair (e.g., .name_repair = make.names for names in the style of base R).</li> <li>• A purrr-style anonymous function, see <a href="#">rlang::as_function()</a></li> </ul> <p>This argument is passed on as repair to <a href="#">vctrs::vec_as_names()</a>. See there for more details on these terms and the strategies used to enforce them.</p>
rownames	<p>How to treat existing row names of a data frame or matrix:</p> <ul style="list-style-type: none"> <li>• NULL: remove row names. This is the default.</li> <li>• NA: keep row names.</li> <li>• A string: the name of a new column. Existing rownames are transferred into this column and the row.names attribute is deleted. No name repair is applied to the new column name, even if x already contains a column of that name. Use <a href="#">as_tibble(rownames_to_column(...))</a> to safeguard against this case.</li> </ul> <p>Read more in <a href="#">rownames</a>.</p>

**Value**

A tibble or, for `as_h2o()`, a list with data (an H2OFrame) and id (the id on the h2o server).

**Examples**

```
# start with h2o::h2o.init()
if (h2o_running()) {
  cars2 <- as_h2o(mtcars)
  cars2
  class(cars2$data)

  cars0 <- as_tibble(cars2$data)
  cars0
}
```

---

autoplot.workflow *Plot rankings and metrics of H2O AutoML results*

---

### Description

The `autoplot()` method plots cross validation performances of candidate models in H2O AutoML output via facets on each metric.

### Usage

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

## S3 method for class 'H2OAutoML'
autoplot(
  object,
  type = c("rank", "metric"),
  metric = NULL,
  std_errs = qnorm(0.95),
  ...
)
```

### Arguments

<code>object</code>	A fitted <code>auto_ml()</code> model.
<code>...</code>	Other options to pass to <code>autoplot()</code> .
<code>type</code>	A character value for whether to plot average ranking ("rank") or metrics ("metric").
<code>metric</code>	A character vector or NULL for which metric to plot. By default, all metrics will be shown via facets.
<code>std_errs</code>	The number of standard errors to plot.

### Value

A ggplot object.

### Examples

```
if (h2o_running()) {
  auto_fit <- auto_ml() %>%
    set_engine("h2o", max_runtime_secs = 5) %>%
    set_mode("regression") %>%
    fit(mpg ~ ., data = mtcars)

  autoplot(auto_fit)
}
```

---

h2o_predict	<i>Prediction wrappers for h2o</i>
-------------	------------------------------------

---

**Description**

Prediction wrappers for fitted models with h2o engine that include data conversion, h2o server cleanup, and so on.

**Usage**

```
h2o_predict(object, new_data, ...)

h2o_predict_classification(object, new_data, type = "class", ...)

h2o_predict_regression(object, new_data, type = "numeric", ...)

## S3 method for class `~_H2OAutoML`
predict(object, new_data, id = NULL, ...)
```

**Arguments**

object	An object of class <code>model_fit</code> .
new_data	A rectangular data object, such as a data frame.
...	Other options passed to <a href="#">h2o::h2o.predict()</a>
type	A single character value or NULL. Possible values are "numeric", "class", "prob", "conf_int", "pred_int", "quantile", "time", "hazard", "survival", or "raw". When NULL, <code>predict()</code> will choose an appropriate value based on the model's mode.
id	Model id in AutoML results.

**Details**

For AutoML, prediction is based on the best performing model.

**Value**

For type `!= "raw"`, a prediction data frame with the same number of rows as `new_data`. For type `== "raw"`, return the result of [h2o::h2o.predict\(\)](#).

**Examples**

```
if (h2o_running()) {
  spec <-
    rand_forest(mtry = 3, trees = 100) %>%
    set_engine("h2o") %>%
    set_mode("regression")
}
```

```
set.seed(1)
mod <- fit(spec, mpg ~ ., data = mtcars)
h2o_predict_regression(mod$fit, new_data = head(mtcars), type = "numeric")

# using parsnip
predict(mod, new_data = head(mtcars))
}
```

---

h2o\_start

*Utility functions for interacting with the h2o server*

---

## Description

Utility functions for interacting with the h2o server

## Usage

```
h2o_start()

h2o_end()

h2o_running(verbose = FALSE)

h2o_remove(id)

h2o_remove_all()

h2o_get_model(id)

h2o_get_frame(id)

h2o_xgboost_available()
```

## Arguments

verbose	Print out the message if no cluster is available.
id	Model or frame id.

## Examples

```
## Not run:
if (!h2o_running()) {
  h2o_start()
}

## End(Not run)
```

**Description**

Basic model wrappers for h2o model functions that include data conversion, seed configuration, and so on.

**Usage**

```
h2o_train(  
  x,  
  y,  
  model,  
  weights = NULL,  
  validation = NULL,  
  save_data = FALSE,  
  ...  
)
```

```
h2o_train_rf(x, y, ntrees = 50, mtries = -1, min_rows = 1, ...)
```

```
h2o_train_xgboost(  
  x,  
  y,  
  ntrees = 50,  
  max_depth = 6,  
  min_rows = 1,  
  learn_rate = 0.3,  
  sample_rate = 1,  
  col_sample_rate = 1,  
  min_split_improvement = 0,  
  stopping_rounds = 0,  
  validation = NULL,  
  ...  
)
```

```
h2o_train_gbm(  
  x,  
  y,  
  ntrees = 50,  
  max_depth = 6,  
  min_rows = 1,  
  learn_rate = 0.3,  
  sample_rate = 1,  
  col_sample_rate = 1,  
  min_split_improvement = 0,
```

```

    stopping_rounds = 0,
    ...
)

h2o_train_glm(x, y, lambda = NULL, alpha = NULL, ...)

h2o_train_nb(x, y, laplace = 0, ...)

h2o_train_mlp(
  x,
  y,
  hidden = 200,
  l2 = 0,
  hidden_dropout_ratios = 0,
  epochs = 10,
  activation = "Rectifier",
  validation = NULL,
  ...
)

h2o_train_rule(
  x,
  y,
  rule_generation_ntrees = 50,
  max_rule_length = 5,
  lambda = NULL,
  ...
)

h2o_train_auto(x, y, verbosity = NULL, save_data = FALSE, ...)

```

### Arguments

x	A data frame of predictors.
y	A vector of outcomes.
model	A character string for the model. Current selections are "automl", "randomForest", "xgboost", "gbm", "glm", "deeplearning", "rulefit" and "naiveBayes". Use <a href="#">h2o_xgboost_available()</a> to see if xgboost can be used on your OS/h2o server.
weights	A numeric vector of case weights.
validation	An integer between 0 and 1 specifying the <i>proportion</i> of the data reserved as validation set. This is used by h2o for performance assessment and potential early stopping. Default to 0.
save_data	A logical for whether training data should be saved on the h2o server, set this to TRUE for AutoML models that needs to be re-fitted.
...	Other options to pass to the h2o model functions (e.g., <a href="#">h2o: :h2o.randomForest()</a> ).
ntrees	Number of trees. Defaults to 50.



mtries	Number of variables randomly sampled as candidates at each split. If set to -1, defaults to $\sqrt{p}$ for classification and $p/3$ for regression (where $p$ is the # of predictors Defaults to -1.
min_rows	Fewest allowed (weighted) observations in a leaf. Defaults to 1.
max_depth	Maximum tree depth (0 for unlimited). Defaults to 20.
learn_rate	(same as eta) Learning rate (from 0.0 to 1.0) Defaults to 0.3.
sample_rate	Row sample rate per tree (from 0.0 to 1.0) Defaults to 0.632.
col_sample_rate	(same as colsample_bylevel) Column sample rate (from 0.0 to 1.0) Defaults to 1.
min_split_improvement	Minimum relative improvement in squared error reduction for a split to happen Defaults to $1e-05$ .
stopping_rounds	Early stopping based on convergence of stopping_metric. Stop if simple moving average of length $k$ of the stopping_metric does not improve for $k$ :stopping_rounds scoring events (0 to disable) Defaults to 0.
lambda	Regularization strength
alpha	Distribution of regularization between the L1 (Lasso) and L2 (Ridge) penalties. A value of 1 for alpha represents Lasso regression, a value of 0 produces Ridge regression, and anything in between specifies the amount of mixing between the two. Default value of alpha is 0 when SOLVER = 'L-BFGS'; 0.5 otherwise.
laplace	Laplace smoothing parameter Defaults to 0.
hidden	Hidden layer sizes (e.g. [100, 100]). Defaults to c(200, 200).
l2	L2 regularization (can add stability and improve generalization, causes many weights to be small. Defaults to 0.
hidden_dropout_ratios	Hidden layer dropout ratios (can improve generalization), specify one value per hidden layer, defaults to 0.5.
epochs	How many times the dataset should be iterated (streamed), can be fractional. Defaults to 10.
activation	Activation function. Must be one of: "Tanh", "TanhWithDropout", "Rectifier", "RectifierWithDropout", "Maxout", "MaxoutWithDropout". Defaults to Rectifier.
rule_generation_ntrees	Specifies the number of trees to build in the tree model. Defaults to 50. Defaults to 50.
max_rule_length	Maximum length of rules. Defaults to 3.
verbosity	Verbosity of the backend messages printed during training; Must be one of NULL (live log disabled), "debug", "info", "warn", "error". Defaults to NULL.

**Value**

An h2o model object.

## Examples

```
# start with h2o::h2o.init()
if (h2o_running()) {
  # -----
  # Using the model wrappers:
  h2o_train_glm(mtcars[, -1], mtcars$mpg)

  # -----
  # using parsnip:

  spec <-
    rand_forest(mtry = 3, trees = 500) %>%
    set_engine("h2o") %>%
    set_mode("regression")

  set.seed(1)
  mod <- fit(spec, mpg ~ ., data = mtcars)
  mod

  predict(mod, head(mtcars))
}
```

---

rank\_results.workflow *Tools for working with H2O AutoML results*

---

## Description

Functions that returns a tibble describing model performances.

- `rank_results()` ranks average cross validation performances of candidate models on each metric.
- `collect_metrics()` computes average statistics of performance metrics (summarized) for each model, or raw value in each resample (unsummarized).
- `tidy()` computes average performance for each model.
- `member_weights()` computes member importance for stacked ensemble models, i.e., the relative importance of base models in the meta-learner. This is typically the coefficient magnitude in the second-level GLM model.

`extract_fit_engine()` extracts single candidate model from `auto_ml()` results. When `id` is null, it returns the leader model.

`refit()` re-fits an existing AutoML model to add more candidates. The model to be re-fitted needs to have engine argument `save_data = TRUE`, and `keep_cross_validation_predictions = TRUE` if stacked ensembles is needed for later models.

**Usage**

```

## S3 method for class 'workflow'
rank_results(x, ...)

## S3 method for class '`_H2OAutoML`'
rank_results(x, ...)

## S3 method for class 'H2OAutoML'
rank_results(x, n = NULL, id = NULL, ...)

## S3 method for class 'workflow'
collect_metrics(x, ...)

## S3 method for class '`_H2OAutoML`'
collect_metrics(x, ...)

## S3 method for class 'H2OAutoML'
collect_metrics(x, summarize = TRUE, n = NULL, id = NULL, ...)

## S3 method for class '`_H2OAutoML`'
tidy(x, n = NULL, id = NULL, keep_model = TRUE, ...)

get_leaderboard(x, n = NULL, id = NULL)

member_weights(x, ...)

## S3 method for class '`_H2OAutoML`'
extract_fit_parsnip(x, id = NULL, ...)

## S3 method for class '`_H2OAutoML`'
extract_fit_engine(x, id = NULL, ...)

## S3 method for class 'workflow'
refit(object, ...)

## S3 method for class '`_H2OAutoML`'
refit(object, verbosity = NULL, ...)

```

**Arguments**

...	Not used.
n	An integer for the number of top models to extract from AutoML results, default to all.
id	A character vector of model ids to retrieve.
summarize	A logical; should metrics be summarized over resamples (TRUE) or return the values for each individual resample.

keep_model	A logical value for if the actual model object should be retrieved from the server. Defaults to TRUE.
object, x	A fitted <code>auto_ml()</code> model or workflow.
verbosity	Verbosity of the backend messages printed during training; Must be one of NULL (live log disabled), "debug", "info", "warn", "error". Defaults to NULL.

### Details

H2O associates with each model in AutoML an unique id. This can be used for model extraction and prediction, i.e., `extract_fit_engine(x, id = id)` returns the model and `predict(x, id = id)` will predict for that model. `extract_fit_parsnip(x, id = id)` wraps the h2o model with `parsnip` `parsnip` model object is discouraged.

The `algorithm` column corresponds to the model family H2O use for a particular model, including `xgboost` ("XGBOOST"), gradient boosting ("GBM"), random forest and variants ("DRF", "XRT"), generalized linear model ("GLM"), and neural network ("deeplearning"). See the details section in `h2o::h2o.automl()` for more information.

### Value

A `tibble::tibble()`.

### Examples

```
if (h2o_running()) {
  auto_fit <- auto_ml() %>%
    set_engine("h2o", max_runtime_secs = 5) %>%
    set_mode("regression") %>%
    fit(mpg ~ ., data = mtcars)

  rank_results(auto_fit, n = 5)
  collect_metrics(auto_fit, summarize = FALSE)
  tidy(auto_fit)
  member_weights(auto_fit)
}
```

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