

# Package ‘agua’

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

**Version** 0.0.1

**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>

**Depends** parsnip

**Imports** dplyr, glue, h2o (>= 3.36.0.4), hardhat, purrr, rlang, stats, tibble, utils

**Suggests** covr, knitr, testthat (>= 3.0.0)

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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	Treatment of problematic column names: <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., <code>.name_repair = make.names</code> 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 <code>repair</code> 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	How to treat existing row names of a data frame or matrix: <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 <code>row.names</code> attribute is deleted. Read more in <a href="#">rownames</a>.</li> </ul>

**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
}
```

---

`h2o_running`*Check if h2o cluster is initialized*

---

**Description**

Check if h2o cluster is initialized

**Usage**

```
h2o_running(verbose = FALSE)
```

**Arguments**

`verbose` Print out the message if no cluster is available.

**Value**

A logical.

**Examples**

```
h2o_running()
h2o_running(verbose = TRUE)
```

h2o\_train

*Model wrappers for h2o***Description**

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

**Usage**

```
h2o_train(x, y, model, ...)
```

```
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,
  ...
)
```

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

**Arguments**

x	A data frame of predictors
y	A vector of outcomes.
model	A character string for the model. Current selections are "randomForest", "xgboost", and "glm". Use <code>h2o:h2o.xgboost.available()</code> to see if that model can be used on your OS/h2o server.
...	Other options to pass to the h2o model functions (e.g., <code>h2o:h2o.randomForest()</code> ).
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.

**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 = 1000) %>%
    set_engine("h2o") %>%
    set_mode("regression")

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

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

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