

Description

`h2omlselect` retrieves the fitted model with the hyperparameter configuration you select after `h2omlgbm` and `h2omlrf` perform tuning using a [grid search](#). These estimation commands select the top-performing model, the one with the most optimal tuning performance metric, as the working model. After estimation, you can use `h2omlestat gridsummary` to see performance metrics for models with different hyperparameter configurations and to obtain an ID for each of these models. You can then select a different model to be the working model by using `h2omlselect`. `h2omlselect` selects and retrieves the fitted model; afterward, you can treat this model just as you would treat estimation results from the `h2omlgbm` and `h2omlrf` estimation commands. Subsequent postestimation commands are based on the selected model.

Quick start

After performing multiclass classification and obtaining the grid-search summary, select the model that has `id = 2`

```
h2oml rfmulticlass y x1-x20, ntrees(10(5)100) maxdepth(3(1)10)
h2omlestat gridsummary
h2omlselect id = 2
```

Menu

Statistics > H2O machine learning

Syntax

```
h2omlselect id = #
```

where `#` is a grid ID from `h2omlestat gridsummary` corresponding to the desired model configuration.

Remarks and examples

Building a machine learning model that generalizes well to new data involves choosing an appropriate method and selecting a model by tuning hyperparameters. We can perform a grid search using gradient boosting and random forest methods and then use `h2omlestat gridsummary` to report the hyperparameter configurations that achieve the top performance based on the specified metric. For example, you might use the log-loss metric to choose between models with 10, 20, and 30 trees. Typically, you would select the model that performs the best based on the chosen metric. However, you may want to explore different hyperparameter configurations that do not correspond to the best model, in which case you can use `h2omlselect` and `h2omlexplore`.

After you review the grid-search summary from `h2omlstat gridsummary`, you can select the model you are interested in by specifying the ID number with `h2omlselect`. Once you have selected a model with `h2omlselect`, you can treat the model in the same way you would treat results from the `h2oml gbm` and `h2oml rf` estimation commands. Postestimation commands will be based on the model selected by `h2omlselect`; for example, you could estimate variable importance for the selected model with `h2omlgraph varimp`. `h2omlselect` overwrites the previously stored estimation results, which can be recovered by refitting the original model or by storing the estimation results before running `h2omlselect` and then restoring them; see [\[H2OML\] h2omlest](#).

► Example 1: Selecting the second-best model

In this example, we illustrate the use of `h2omlselect` by performing random forest binary classification with the social pressure dataset discussed in [example 1](#) of [\[H2OML\] h2oml rf](#).

We start by opening the social pressure dataset in Stata and then putting the data into an H2O frame. Recall that `h2o init` initiates an H2O cluster, `_h2oframe put` loads the current Stata dataset in an H2O frame, and `_h2oframe change` makes the specified frame the current H2O frame. We use the `_h2oframe split` command to randomly split the `social` frame into a training frame (80% of observations) and a validation frame (20% of observations), which we name `train` and `valid`, respectively. We also change the current frame to `train`. For details, see [Prepare your data for H2O machine learning in Stata](#) in [\[H2OML\] h2oml](#) and see [\[H2OML\] H2O setup](#).

```
. use https://www.stata-press.com/data/r19/socialpressure
(Social pressure data)

. h2o init
(output omitted)

. _h2oframe _put, into(social)
Progress (%): 0 100

. _h2oframe _split social, into(train valid) split(0.8 0.2) rseed(19)

. _h2oframe _change train
```

We define a global macro, `predictors`, to store the names of our predictors. We perform random forest binary classification, and we specify the `maxdepth()` and `predsampvalue()` options to tune the maximum tree depth and predictor sampling rate hyperparameters. For illustration, we use the area under the precision–recall curve (AUCPR) metric for tuning.

```
. global predictors gender g2000 g2002 p2000 p2002 p2004 treatment age
. h2oml rfbinclass voted $predictors, validframe(valid) h2orseed(19)
> ntrees(200) maxdepth(3(3)12) predsampvalue(-1, 1(2)8) tune(metric(aucpr))
```

Progress (%): 0 100

Random forest binary classification using H2O

Response: voted

Frame:	Number of observations:
Training: train	Training = 183,607
Validation: valid	Validation = 45,854

Tuning information for hyperparameters

Method: Cartesian

Metric: AUCPR

Hyperparameters	Grid values		
	Minimum	Maximum	Selected
Max. tree depth	3	12	6
Pred. sampling value	-1	7	7

Model parameters

Number of trees = 200

actual = 200

Tree depth:	Pred. sampling value =	7
Input max = 6	Sampling rate =	.632
min = 6	No. of bins cat. =	1,024
avg = 6.0	No. of bins root =	1,024
max = 6	No. of bins cont. =	20
Min. obs. leaf split = 1	Min. split thresh. =	.00001

Metric summary

Metric	Training	Validation
Log loss	.5724664	.5705699
Mean class error	.3935492	.3943867
AUC	.6705554	.6734867
AUCPR	.4658395	.4725543
Gini coefficient	.3411109	.3469735
MSE	.1946923	.1935647
RMSE	.4412395	.4399599

Next we obtain the grid-search summary by using the `h2omlestat gridsummary` command. This command lists the configuration of the hyperparameters we are tuning ranked by AUCPR.

```
. h2omlestat gridsummary
```

Grid summary using H2O

ID	Max. tree depth	Pred. sampling value	AUCPR
1	6	7	.4725543
2	6	5	.4723736
3	6	3	.4714554
4	9	3	.4712076
5	6	-1	.4708614
6	12	-1	.4706606
7	9	-1	.4705794
8	9	5	.4689799
9	9	7	.4682457
10	9	1	.4674565

The top two models have very similar values of AUCPR, and they correspond to models with 7 and 5 randomly sampled predictors and a maximum tree depth of 6. As discussed in [H2OML] *h2oml rf*, using a random sample of predictors improves the ability of the model to generalize to new data, compared with using the full set of predictors, because it introduces an additional randomness to the method. Therefore, we may prefer to continue our analysis with the second-best model.

To select the second-best model, we specify `id = 2` in `h2omlselect`.

```
. h2omlselect id = 2
```

Random forest binary classification using H2O

Response: voted

Frame:

Training: train

Validation: valid

Number of observations:

Training = 183,607

Validation = 45,854

Model parameters

Number of trees = 200

actual = 200

Tree depth:

Input max = 6

min = 6

avg = 6.0

max = 6

Min. obs. leaf split = 1

Pred. sampling value = 5

Sampling rate = .632

No. of bins cat. = 1,024

No. of bins root = 1,024

No. of bins cont. = 20

Min. split thresh. = .00001

Metric summary

Metric	Training	Validation
Log loss	.57237	.5704978
Mean class error	.3979593	.3945857
AUC	.671146	.6737527
AUCPR	.4670326	.4723736
Gini coefficient	.342292	.3475054
MSE	.1946602	.1935627
RMSE	.4412031	.4399576

Now we can continue our analysis using the second-best model.



Stored results

`h2omlselect` retrieves the selected fitted model and thus stores the same results as the estimation command used.

See *Stored results* in [H2OML] [h2oml gbm](#) or [H2OML] [h2oml rf](#).

Also see

[H2OML] [h2oml](#) — Introduction to commands for Stata integration with H2O machine learning

[H2OML] [h2omlestat gridsummary](#) — Display grid-search summary

[H2OML] [h2omlexplore](#) — Explore models after grid search

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