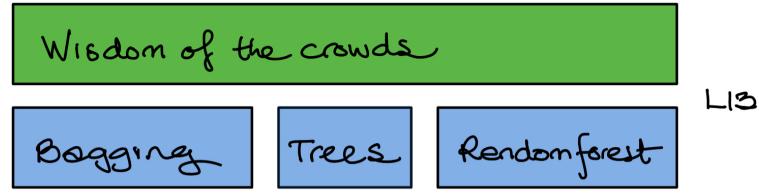
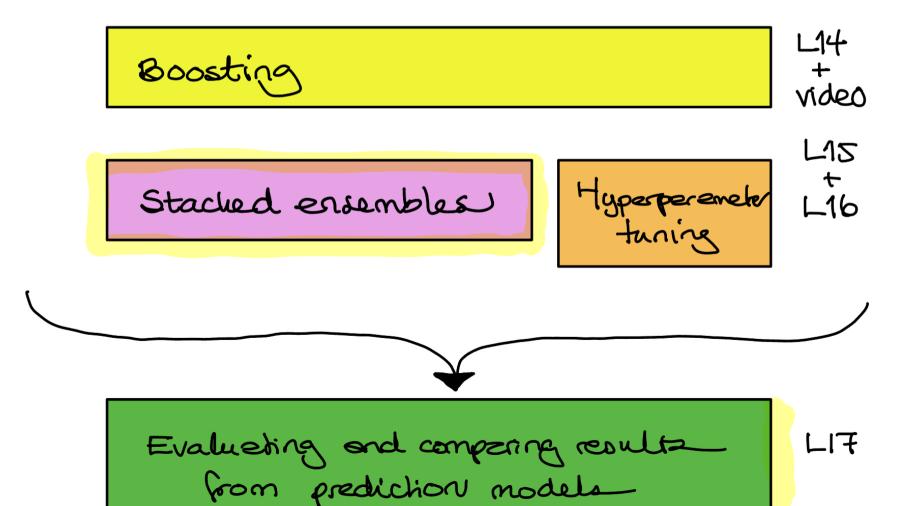
MA8701 Advanced methods in statistical inference and learning Part 3: Ensembles, L15: Stacked ensembles

Mette Langaas

3/5/23-Lechred 06.03.2023

Before we start





< ⊡ >

Stacked ensembles

The idea is to combine predictions is well-known in statistics but now should this be done in practice 2

aka super learner or generalized stacking

What is it?

The Stacked Esembles is an algorithm that combines

- multiple, (typically) diverse prediction methods (learning algorithms) called base learners (first-level) into a
- a second-level metalearner which can be seen as a single method.

 method.
 R

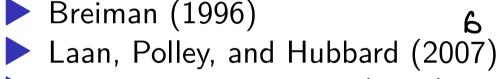
1) Base learned the l=1,.., h: from possibly differt methods

2) Metalezner
$$\overline{\Phi}$$
: an "ophinal" combination of the lose learner

Literature

Erin Le Dell (2015): Scalable Ensemble Learning and Computationally Efficient Variance Estimation. PhD Thesis, University of California, Berkeley. or https://github.com/ledell/phd-thesis. Section 2.

Supporting literature



Polley, Rose, and Laan (2011)



ESTIMATION STRATEGY

STACKED ENERBLE FITTING STRATEGY

"Any" method that produces a prediction - "all" types of problems.

- linear regression
 - lasso
- **c**art
- random forest with mtry=value 1
- random forest with mtry=value 2
- xgboost with hyperparameter set 1
- xgboost with hyperparameter set 2
- neural net with hyperparameter set 1

Simplest case: regression
$$\hat{\varphi}(x) = \hat{\chi}_1 \hat{\psi}(x) + \dots + \hat{\chi}_L \hat{\psi}(x)$$

 $\hat{\varphi}(x_i) = \hat{\chi}_1 Z_{1i} + \hat{\chi}_L Z_{1i}$

fit using elestic net \Rightarrow get $\hat{\chi}_{L}$, $\hat{\chi}_{L}$ and $\hat{\chi}_{S} \ge 0$

For classification
$$10, 13$$

let $\hat{J}(x) = \hat{\alpha}_{1} \hat{f}^{1}(x) + \dots + \hat{\alpha} \hat{f}^{1}(x) + \hat{\alpha}_{0}$
or maybe $\hat{J}(x) = \hat{\alpha}_{1} \log_{1}(\hat{f}^{1}(x)) + \dots + \hat{\alpha}_{1}\log_{1}(\hat{f}^{1}(x)) + \hat{\alpha}_{0}$
end fit logistic elastic net
 $\hat{f}(x) = \frac{e^{\hat{f}(x)}}{1 + e^{\hat{f}(x)}} = exp_{1}(\hat{f}(x) + o \operatorname{get} \hat{d}_{1,\dots,n} dn)$
prediction
BUT, also use othe methods, for example:

the mean (bagging)
constructed by minimizing the squared loss (ordinary least squares) or
non-negative least squares (most popular) < rest page
ridge or lasso regression or elected
logistic regression (for binary classification) < defended
constructed by minimizing 1-ROC-AUC
Le Dell phD < "suited for class inhole ree"

< ₽ >

2. Why Non-negativity Constraints Work

Only partial answers are available. Suppose that the $\{v_k(x)\}$ are strongly correlated and the $\{\alpha_k\}$ are chosen using least squares or ridge regression. Then there is no guarantee that the resulting predictor $\sum_k \alpha_k v_k(x)$ will stay near the range $[\min_k v_k(x), \max_k v_k(x)]$ and generalization may be poor.

Λ **(** Ψ(×)

L. BREIMAN

Breinen

1996

Now consider imposing the non-negativity constraints on the $\{\alpha_k\}$ together with the additional constraint $\sum \alpha_k = 1$. For any $\{\alpha_k\}$ satisfying the constraints $\alpha_k \ge 0$, $\sum_k \alpha_k = 1$,

$$v(\boldsymbol{x}) = \sum_{k} \alpha_{k} v_{k}(\boldsymbol{x})$$

is an "interpolating" predictor. That is, for every value of x,

$$\min_{k} v_k(\boldsymbol{x}) \leq v(\boldsymbol{x}) \leq \max_{k} v_k(\boldsymbol{x}).$$

So, what our procedure does is to find the best "interpolating" predictor.

52



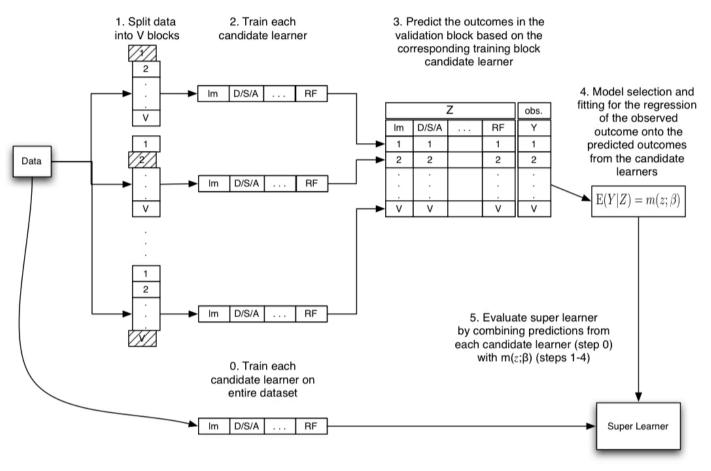
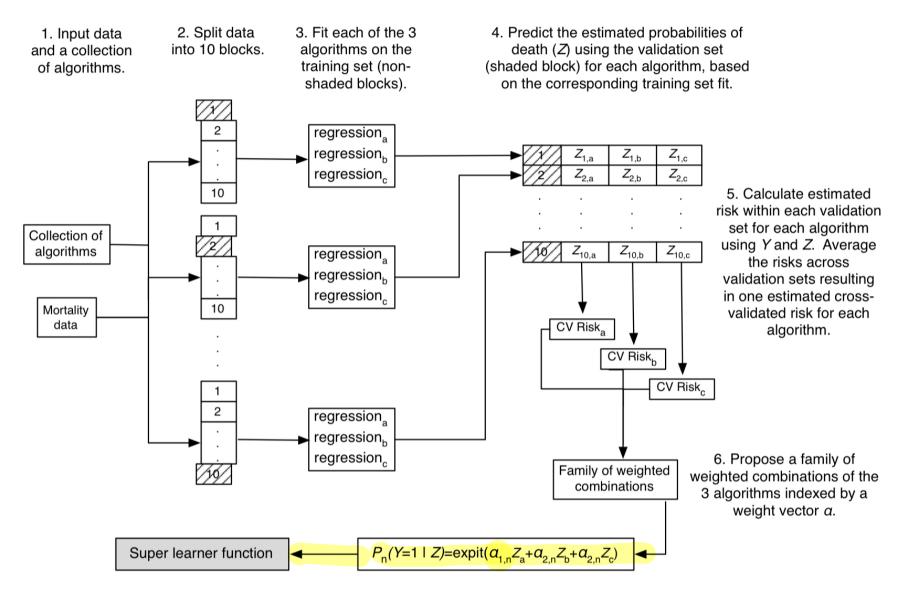


Figure 1: Flow Diagram for Super Learner

(Class notes: Study Figure 3.2 from Polley, Rose, and Laan (2011) and/or Figure 1 from Laan, Polley, and Hubbard (2007))



8. Fit each of the algorithms on the complete data set. Combine these fits with the weights obtained in the previous step to generate the super learner predictor function.

 Use the probabilities (*Z*) to predict the outcome *Y* and estimate the vector *a*, thereby determining the combination that minimizes the crossvalidated risk over the family of weighted combinations.

Fig. 3.2 Super learner algorithm for the mortality study example

The metalearning

Some observations

- only one $\hat{\alpha}_{L} \neq 0$ The term discrete super learner is used if the base learner with the lowest risk (i.e. CV-error) is selected.
- Since the predictions from multiple base learners may be highly correlated - the chosen method should perform well in that case (i.e. ridge and lasso).
- When minimizing the squared loss it has been found that adding a non-negativity constraint $\alpha_l \ge 0$ works well,
- > and also the additivity constraint $\sum_{l=1}^{L} \alpha_l = 1$ the ensemble is a *convex combination* of the base learners.
- Non-linear optimization methods may be employed for the metalearner if no existing algorithm is available
- Historically a regularized linear model has "mostly" been used For classification the logistic response function can be used on the linear combination of base learners (Figure 3.2 Polley, Rose, and Laan (2011)).

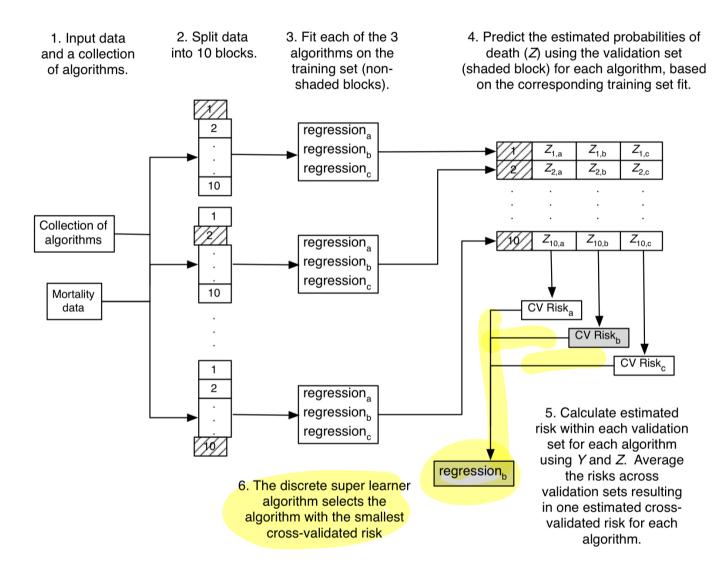


Fig. 3.1 Discrete super learner algorithm for the mortality study example where $\bar{Q}_n^b(A, W)$ is the algorithm with the smallest cross-validated risk

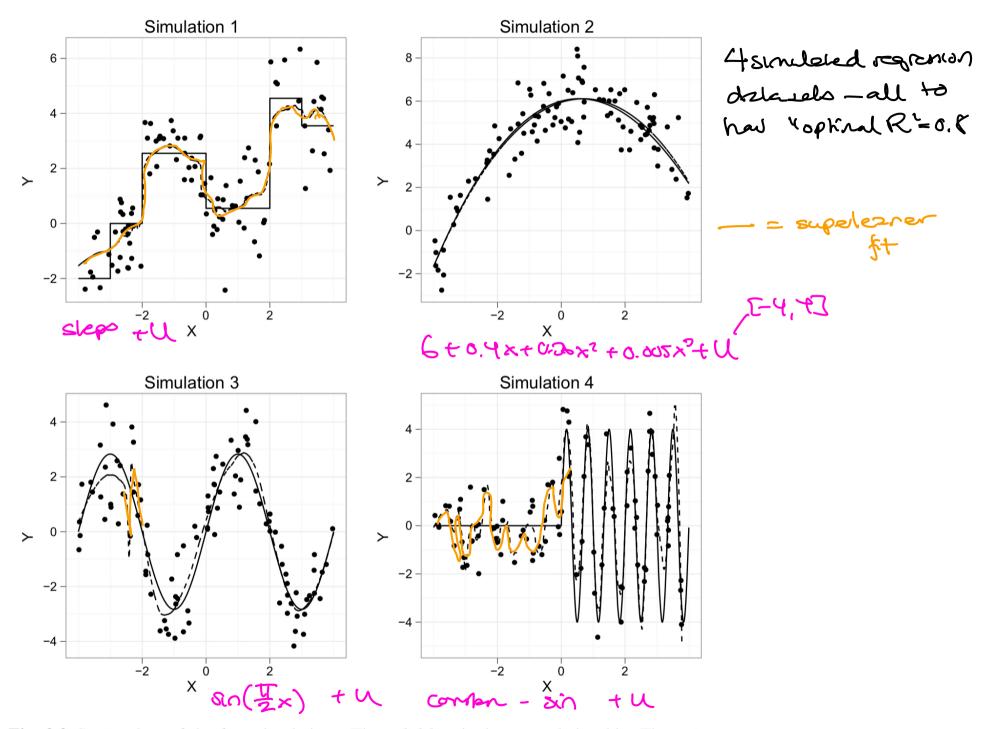


Fig. 3.3 Scatterplots of the four simulations. The *solid line* is the true relationship. The *points* represent one of the simulated data sets of size n = 100. The *dashed line* is the super learner fit for the shown data set

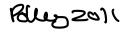


Table 3.2 Results for four simulations. Average R^2 based on 100 simulations and the corresponding standard errors

	Algorithm	Sim 1		Sim 2		Sim 3		Sim 4					
		R^2	$SE(R^2)$	R^2	$SE(R^2)$	R^2	$SE(R^2)$	R^2	$SE(R^2)$	R2-	1_	$\frac{\Sigma(Y,-\hat{Y})}{\Sigma(Y,-\bar{Y})^2}$	2
	Super learner	0.741	0.032	0.754	0.025	0.760	0.025	0.496	0.122		1-	524:-9)?	2
	Discrete SL	0.729	0.079	0.758	0.029	0.757	0.055	0.509	0.132				
	SL.glm	0.422	0.012	0.189	0.016	0.107	0.016	-0.018	0.021				
	SL.interaction	0.428	0.016	0.769	0.011	0.100	0.020	-0.018	0.029				
	SL.randomForest	0.715	0.021	0.702	0.027	0.724	0.018	0.460	0.109				
	SL.bagging(0.01)	0 <mark>.751</mark>	0.022	0.722	0.036	0.723	0.018	0.091	0.054				
	SL.bagging(0.1)	0.635	0.120	0.455	0.195	0.661	0.029	0.020	0.025				
	SL.bagging(0.0)	0.752	0.021	0.722	0.034	0.727	0.017	0.102	0.060				
	SL.bagging(ms5)	<mark>0.747</mark>	0.020	0.727	0.030	0.741	0.016	0.369	0.104				
	SL.gam(2)	0.489	0.013	0.649	0.026	0.213	0.029	-0.014	0.023				
	SL.gam(3)	0.535	0.033	0.748	0.024	0.412	0.037	-0.017	0.029				
	SL.gam(4)	0.586	0.027	0.759	0.020	0.555	0.022	-0.020	0.034				
	SL.gbm	0.717	0.035	0.694	0.038	0.679	0.022	0.063	0.040				
	SL.nnet(2)	0.476	0.235	0.591	0.245	0.283	0.285	-0.008	0.030				
	SL.nnet(3)	0.700	0.096	0.700	0.136	0.652	0.218	0.009	0.035				
	SL.nnet(4)	0.719	0.077	0.730	0.062	0.738	0.102	0.032	0.052				
	SL.nnet(5)	0.705	0.079	0.716	0.070	0.731	0.077	0.042	0.060				
	SL.polymars	0.704	0.033	0.733	0.032	0.745	0.034	0.003	0.040				
	SL.bart	0.740	0.015	0.737	0.027	0.764	0.014	0.077	0.034				
	SL.loess(0.75)	0.599	0.023	0.761	0.019	0.487	0.028	-0.023	0.033				
	SL.loess(0.50)	0.695	0.018	0.754	0.022	0.744	0.029	-0.033	0.038				
	SL.loess(0.25)	0.729	0.016	0.738	0.025	0.772	0.015	-0.076	0.068				
,	SL.loess(0.1)	0.690	0.044	0.680	0.064	0.699	0.039	0.544	0.118				

21(2) base bære Tehe home mensage: - the best algo (baselearner) is not known in advance and will change = dependent on problem& deba - adding more base (corners -> superleaner does better end get R²= 0.76 optimally

Regresson $RE = \frac{MSE}{MSE(UN)}$ geo rezo of Bresuls SuperLearner discreteSL bart gam(3)gam(4) gam gam(5) polymars step.interaction glm bayesglm -Method 200-654 glmnet(.75) -3-18 glmnet(1) glmnet(.50) Name Source п р/ glmnet(.25) ais 202 10 Cook and Weisberg (1994) DSA diamond 308 17 Chu (2001) step cps78 550 18 Berndt (1991) ridge cps85 534 17 Berndt (1991) gbm(2) -. 209 6 Kibler et al. (1989) cpu randomForest Rosner (1999) **FEV** 654 4 gbm(1) -392 7 Newman et al. (1998) Pima svm 200 10 laheart Afifi and Azen (1979) 201 3 mussels Cook (1998) 0.5 10 12 14 2 4 8 6 258 6 enroll Liu and Stengos (1999) **Relative MSE** 252 14 Penrose et al. (1985) fat diabetes 366 15 Harrell (2001) Newman et al. (1998) 506 13 house

Fig. 3.4 Tenfold cross-validated relative mean squared error compared to glm across 13 real data sets. Sorted by geometric mean, denoted by the plus (+) sign

```
13 details
```

Theoretical result

- LeDell (2015) (page 6)
 - Oracle selector: the estimator among all possible weighted combinations of the base prediction function that minimizes the risk under the *true data generating distribution*.
 - The oracle result was established for the Super Learner by Laan, Polley, and Hubbard (2007)
 - If the true prediction function cannot be represented by a combination of the base learners (available), then "optimal" will be the closest linear combination that would be optimal if the true data-generating function was known.
 - The oracle result require an uniformly bounded loss function. Using the convex restriction (sum alphas =1) implies that if each based learner is bounded so is the convex combination. In practice: truncation of the predicted values to the range of the outcome in the training set is sufficient to allow for unbounded loss functions

Other issues

- Many different implementations available, and much work on parallell processing and speed and memory efficient execution.
- Super Learner implicitly can handle hyperparameter tuning by including the same base learner with different model parameter sets in the ensemble.
- Speed and memory improvements for large data sets involves subsampling, and the R subsemble package is one solution, the H2o package another.

Super Learner Algorithm

H20 help progr

The steps below describe the individual tasks involved in training and testing a Super Learner ensemble. H2O automates most of the steps below so that you can quickly and easily build ensembles of H2O models.

1. Set up the ensemble.

- a. Specify a list of L base algorithms (with a specific set of model parameters).
- b. Specify a metalearning algorithm.

2. Train the ensemble.

- a. Train each of the L base algorithms on the training set.
- b. Perform k-fold cross-validation on each of these learners and collect the cross-validated predicted values from each of the L algorithms.
- c. The N cross-validated predicted values from each of the L algorithms can be combined to form a new N x L matrix. This matrix, along with the original response vector, is called the "level-one" data. (N = number of rows in the training set.)
- d. Train the metalearning algorithm on the level-one data. The "ensemble model" consists of the L base learning models and the metalearning model, which can then be used to generate predictions on a test set.
- 3. Predict on new data.
 - a. To generate ensemble predictions, first generate predictions from the base learners.
 - b. Feed those predictions into the metalearner to generate the ensemble prediction.



- metalearner_algorithm (Optional) Specify the metalearner algorithm type. Options include:
 - "AUTO" (GLM with non negative weights & standardization turned off, and if validation_frame is present, then lambda_search is set to True; may change over time). This is the default.
 - "glm" (GLM with default parameters)
 - ["gbm"] (GBM with default parameters)
 - ["drf"] (Random Forest with default parameters)
 - "deeplearning" (Deep Learning with default parameters)
 - "naivebayes" (NaiveBayes with default parameters)
 - "xgboost" (if available, XGBoost with default parameters)
- metalearner_params: (Optional) If a metalearner_algorithm is specified, then you can also specify a list of customized parameters for that algorithm (for example, a GBM with ntrees=100, max_depth=10, etc.)
- metalearner_nfolds: (Optional) Specify the number of folds for cross-validation of the metalearning algorithm. Defaults to 0 (no cross-validation). If you want to compare the crossvalidated performance of the ensemble model to the cross-validated performance of the base learners or other algorithms, you should make use of this option.
- metalearner_fold_assignment: (Optional; Applicable only if a value for metalearner_nfolds is specified) Specify the cross-validation fold assignment scheme for the metalearner. The available options are AUTO (which is Random), Random, Modulo, or Stratified (which will stratify the folds based on the response variable for classification problems). This value defaults to AUTO.
- metalearner_fold_column: (Optional; Cannot be used at the same time as nfolds.) Specify the name of the column that contains the cross-validation fold assignment per observation for cross-validation of the metalearner. The column can be numeric (e.g. fold index or other integer value) or it can be categorical. The number of folds is equal to the number of unique values in this column.
- metalearner_transform: (Optional) Specify the transformation used on predictions from the base models in order to make a level one frame. Options include:
 - "NONE" (no transform applied)
 - "Logit" (applicable only to classification tasks, use logit transformation on the predicted probabilities)

R example from H2o-package

https://docs.h2o.ai/h2o/latest-stable/h2o-docs/data-science/stacked-ensembles.html

Python examples available from the same page

The Higgs boson data is used - but which version is not specified, maybe this https://archive.ics.uci.edu/ml/datasets/HIGGS or a specifically made data set. The problem is binary, so maybe to detect signal vs noise.

h2o.init()

Connection successful!

```
R is connected to the H2O cluster:

H2O cluster uptime: 1 days 3 hours

H2O cluster timezone: Europe/Oslo

H2O data parsing timezone: UTC

H2O cluster version: 3.40.0.1

H2O cluster version age: 25 days

H2O cluster name: H2O_started_from_R_mettela_bze126

H2O cluster total nodes: 1
```

```
# Identify predictors and response
  y <- "response"
  x <- setdiff(names(train), y)</pre>
  # For binary classification, response should be a factor
  train[, y] <- as.factor(train[, y])</pre>
  test[, y] <- as.factor(test[, y])</pre>
  print(dim(train))
[1] 10000
                29
  # Number of CV folds (to generate level-one data for stacking)
  nfolds <- 5
  # There are a few ways to assemble a list of models to stack toegether:
  # 1. Train individual models and put them in a list
  # 1. Generate a 2-model ensemble (GBM + RF)
  # Train & Cross-validate a GBM
  my_gbm <- h2o.gbm(x = x,
                   y = y,
                   training_frame = train,
                   distribution = "bernoulli",
                   ntrees = 10,
                   max_depth = 3,
                   min_rows = 2,
                   learn_rate = 0.2,
                   nfolds = nfolds,
                   keep_cross_validation_predictions = TRUE,
                   seed = 1)
```

16

keep_cross_validation_predictions = TRUE,
seed = 1)

Now the default metalearner

Default metalearner: Options include 'AUTO' (GLM with non negative weights; if validation_frame is present, a lambda search is performed)

```
# default metalearner_transform should be NONE
#print(summary(ensemble))
#ensemble@model
# Eval ensemble performance on a test set
perf <- h2o.performance(ensemble, newdata = test)
# Compare to base learner performance on the test set
perf_gbm_test <- h2o.performance(my_gbm, newdata = test)
perf_rf_test <- h2o.performance(my_rf, newdata = test)
baselearner_best_auc_test <- max(h2o.auc(perf_gbm_test), h2o.auc(perf_r</pre>
```

```
Adding transform "logit"
```

Train a stacked ensemble using the GBM and RF above
ensemble <- h2o.stackedEnsemble(x = x,</pre>

```
y = y,
training_frame = train,
base_models = list(my_gbm, my_rf),
metalearner_transform = "Logit")
```

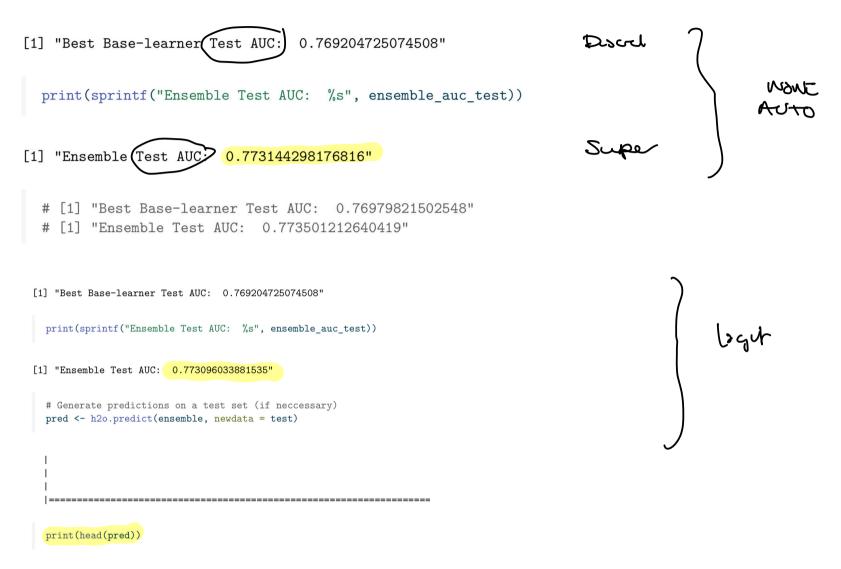
```
#print(summary(ensemble))
#print(ensemble@model)
```

Eval ensemble performance on a test set
perf <- h20.performance(ensemble, newdata = test)</pre>

```
# Compare to base learner performance on the test set
perf_gbm_test <- h20.performance(my_gbm, newdata = test)
perf_rf_test <- h20.performance(my_rf, newdata = test)
baselearner_best_auc_test <- max(h20.auc(perf_gbm_test), h20.auc(perf_r
ensemble auc test <- h20.auc(perf)</pre>
```

```
$metalearner_model
Model Details:
_____
H2OBinomialModel: glm
Model ID:
          metalearner_AUTO_StackedEnsemble_model_R_1677945156774_1830
GLM Model: summary
    family link
                                               regularization number o
1 binomial logit Elastic Net (alpha = 0.5, lambda = 3.885E-4)
                                               training_frame
1 levelone_training_StackedEnsemble_model_R_1677945156774_1830
Coefficients: glm coefficients
                          names coefficients standardized_coefficients
                                                              0.154528
1
                      Intercept
                                  -0.053725
2 GBM_model_R_1677945156774_1086 0.791767
                                                              0.515081
3 DRF_model_R_1677945156774_1214 0.845217
                                                              0.731991
```

$$\hat{\psi}(x) = -0.05 \pm 0.79 \cdot \log(\hat{\psi}_{68n}(x)) + 0.85 \cdot \log(\hat{\psi}_{68n}(x))$$

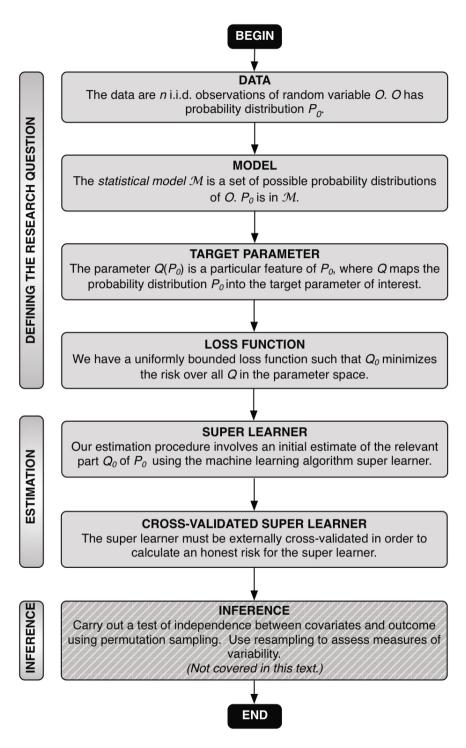


	predict	p0	p1		
1	0	0.6739209	0.3260791		
2	1	0.5814741	0.4185259		
3	1	0.5826643	0.4173357		
4	1	0.1971804	0.8028196		
5	1	0.4561659	0.5438341		
6	1	0.3365841	0.6634159		

Uncertainty in the ensemble

(Class notes: Study "Road map" 2 from Polley, Rose, and Laan (2011))

- Add an outer (external) cross validation loop (where the super learner loop is inside). Suggestion: use 20-fold, especially when small sample size.
- Overfitting? Check if the super learner does as well or better than any of the base learners in the ensemble.
- Results using *influence functions* for estimation of the variance for the Super Learner are based on asymptotic variances in the use of V-fold cross-validation (see Ch 5.3 of LeDell (2015))



Ensembles - overview

(ELS Ch 16.1)

With ensembles we want to build *one prediction model* which combines the strength of *a collection of models*.

These models may be simple base models - or more elaborate models.

We have studied bagging - where we use the bootstrap to repeatedly fit a statistical model, and then take a simple average of the predictions (or majority vote). Here the base models can be trees - or other type of models.

Random forest is a version of bagging with trees, with trees made to be different (decorrelated).

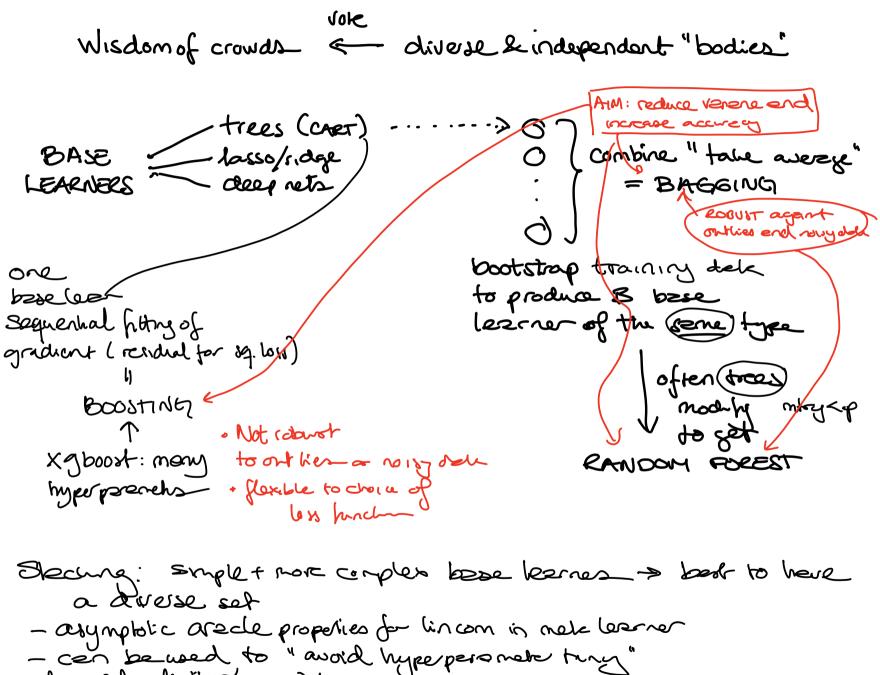
We have studied boosting, where the models are trained on sequentially different data - from residuals or gradients of loss functions - and the ensemble members cast weighted votes (downweighted by a learning rate). We have observed that there are many hyperparameters that need to be to tuned to optimize performance.

And to day - we have barned about the stacked another

END Before we start Wisdom of the crowds LIS Trees Rendomforest Bogging L14 Boosting + video L15 Hyperperenter L16 Stacked ensembles Evaluating and comparing results_ LIF from prediction models

< ⊡ >

ensemble methoda!



- avoid noced selection - no worke