Supervised Learning with Tabular Data Gov 2018

Naijia Liu

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Regression with Continuous Outcome

- Generalized Additive Models (GAM)
- CART
- Motivation of Model Selection

Model Selection and Combination

- Model Selection
- Benign Overfitting
- Model Aggregation
- Evaluation Metrics, Cross-Validation

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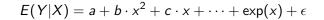
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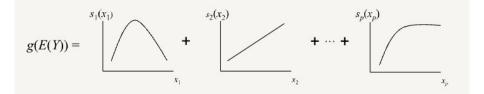
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- For example, it can be linear regression: $f(X_{i1}) = \beta_1 X_{i1}$
- Relationships between IV and DV follow some smooth patterns (linear or non-linear).
- We can estimate these smooth relationships simultaneously and then predict E(Y|X) by simply adding them up.

Example





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- Estimate by iteratively "backfitting": fit f₁(·) with regular (bivariate) smoothing procedure, fit f₂(·) on residuals, ... until convergence
- Can use a combination of linear and flexible terms (e.g. to estimate constant treatment effect while controlling flexibly for other variables)
- Also works for generalized linear models: $E[Y_i] = g^{-1} (\beta_0 + 27(X_{i1}) + \dots + f_K(X_{iK}))$, where $g(\cdot)$ is an exponential link function

GAM is a Flexible Model

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- We can penalize the objective function.

Regression with Continuous Outcome

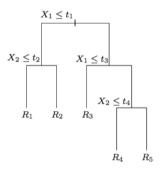
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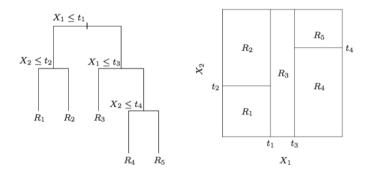
Tree-Based Methods

Classification and Regression Trees (CART) recursively partition the input space



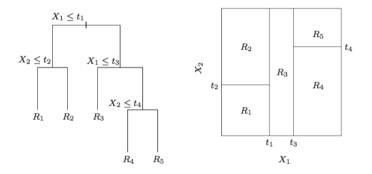
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$$f(x) = \mathbb{E}[\mathbf{y}|X] = \sum_{m=1}^{M} \bar{y}_m \mathbb{1}\{x \in R_m\}$$

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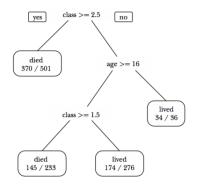
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 - ▶ Titanic example from Varian (2014, JEP): rich children survived



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- Finding the optimal partitioning is NP-complete (i.e., computation time grows too quickly as the size of the problem grows)
- Greedy algorithm (not thinking ahead, missing high-level interaction): locally optimal choice at each stage (not necessarily leads to a global optimum)

CART Algorithm

• At each stage, axis parallel splits is done by:

$$\begin{split} \min_{j,s} \left[\min_{c_1} \sum_{x_i \in R_1(j,s)} (y_i - c_1)^2 + \min_{c_2} \sum_{x_i \in R_2(j,s)} (y_i - c_2)^2 \right] \\ \text{where } R_1(j,s) = \{X | X_j \leq s\} \text{ and } R_2(j,s) = \{X | X_j \geq s\}, \text{ and } c_m \text{ is the mean of } y_i | i \in R_m(j,s). \end{split}$$

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- CART is very popular and easy to implement (rpart package in R)
- However, the greedy algorithm often leads to poor prediction. It also has a problem of over-fitting (pruning is needed), and thus highly variable.

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Model Selection

- What variables should we include in a given model?
- Ridge and Lasso regressions: the knowledge of λ value.
- GAM depends on the selection of smoothing function.
- CART requires pruning to achieve optimal performance.

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Goals

- Model assessment: Estimate generalization error
 - How good will this model be when I use it out-of-sample?
 - Is it better/worse than random choice?
 - Is it better/worse than human coders? (inter-coder reliability)
 - Than a previous approach?

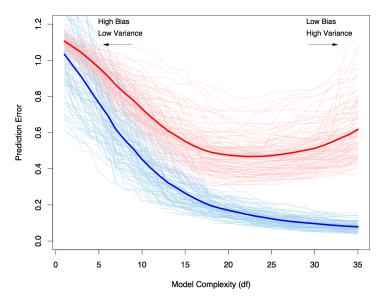
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- Model aggregation: Combining candidate models into something better

Training Error vs. Test Error



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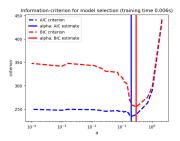
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- We don't have a luxury of having a large data as a test set
- Estimating error variance $\hat{\sigma}$ to calculate loglik is problematic with high-dimensional setting (n < p)
- Cross-validation: Holding out a subset of the training observations from the training process

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• When is this positive? All possible distributions $g(x) \neq f(x)$.

KL divergence is always positive

• In inequality:

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• Proof:

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KL divergence is not a proper distance measure

- Kullback-Leibler divergence is not a distance metric!
- Distances must be nonnegative, symmetric, and satisfy triangle inequality: distance from A to C must be less than (A to B) + (B to C)

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- Useful for comparing images (represented as distribution of pixel values in multivariate color space)
- Or documents, if they are represented as distributions of word coordinates in a continuous embedding space

"The limited but real influence of elite rhetoric in the 2009–2010 health care debate" , Hopkins, 2018, Political behavior

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- The problem: we don't know the true population distribution

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• Interpretation: KL divergence is directly related to the expected log likelihood (under the candidate model) of a randomly drawn observation from the true DGP

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- As sample size goes to infinity, size of validation set generally also goes to infinity (not true for leave-one-out CV)

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Benign Overfitting

- A new statistical phenomenon: good prediction with zero training error (such as a deep learning method) .
- When a perfect fit to training data in linear regression is compatible with accurate prediction.
- Over-parameterization is essential for benign overfitting in some setting: more features than sample size.
- "Benign overfitting in linear regression", Bartlett et al, PNAS, 2020.
- Most of situations we don't have benign overfitting unless you can justify the choice.
- Al models such as Chat GPT relies on massive over-fitting that are benign.

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 - ▶ Use some aggregation rule (e.g. mean, majority vote) to combine

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- Place a prior over all models (e.g. uniform $\pi = [1/K, \dots, 1/K]^{\top})$
- Assumed DGP: first randomly choose a model, then generate all data points from that model

$$Z \sim \mathsf{Cat}(\pi)$$

 $Y_i \sim f_Z(y|X_i = x_i)$

taking \boldsymbol{X} as given

• Posterior belief $P(Z = k | \mathbf{Y}, \mathbf{X})$ depends on π_k and how well $f_k()$ explains observed \mathbf{Y}

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- Out-of-sample prediction:
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 - Take weighted average of predictions

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- However, bagging procedure normally results in highly correlated predictors

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- Improves variance reduction of bagging by reducing the correlation between models
- Easy to implement (randomForest package in R) and often very good off-the-shelf performance

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Bootstrapping: More Perspectives

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- So bootstrap draws each come with a free validation set!

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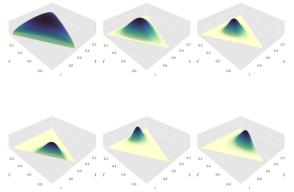
• Observation *i* is in validation set with probability $(1 - 1/N)^N \approx 1 - 1/e \approx 0.632$

• So bootstrap draws each come with a free validation set! Each bootstrap draw is a reweighting of the original data

- For each draw, $w_i \in \{0, 1/N, \cdots, 1\})$
- Weights must sum to 1
- In other words, weights are a distribution over the N-1 simplex

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Bootstrapping: More Perspectives



Bayesian bootstrap:

• The Dirichlet distribution is also a distribution over the N-1 simplex

•
$$\mathbf{w} = [w_1, \cdots, w_N]$$

- $\mathbf{w} \sim \mathsf{Dirichlet}([1, \dots, 1]^{\top})$
- $E[w] = [1/N, ..., 1/N]^{\top}$
- Procedure: Sample w, then do the original analysis. Repeat.

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- These bases can be combined to form an even better approximation to the true conditional expectation function

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- Why? What happens if the true model is among the candidates?
- Asymptotically at least as accurate as the best possible input

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• For out-of-sample data, generate final predictions as $\hat{\mathbf{Y}} = \sum_{m} \alpha_{m} \hat{\mathbf{Y}}^{m}$

- Again, the sequential models $m = 1, \ldots, M$ create basis functions
- We greedily approximate the true conditional expectation with these bases
- Make the best approximation possible using m = 1
- Then fine-tune it with $m = 2, \ldots$
- Models evolve in an adaptive way to remove bias
- Later models focus on examples that were misclassified in earlier rounds (hard to classify, e.g. near best decision boundary)

Regression with Continuous Outcome

- Generalized Additive Models (GAM)
- CART
- Motivation of Model Selection

Model Selection and Combination

- Model Selection
- Benign Overfitting
- Model Aggregation
- Evaluation Metrics, Cross-Validation

Creating out-of-sample, in sample.

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- Then, fit model on all labeled data (training and validation)
- Presumably you have a good proxy for generalization performance from "in-sample" (all labeled data) data to "out-of-sample" (test data, new observations to predict, or true DGP)

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- \bullet Why? These steps doesn't involve access to \boldsymbol{Y}_{test}

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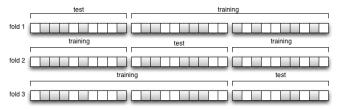
CV estimate of MSE
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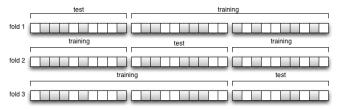
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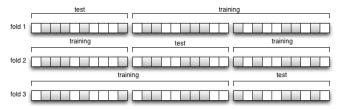
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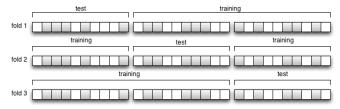


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Naijia Liu

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- Equivalently, maximizing with a proper scoring rule encourages well-calibrated predictions

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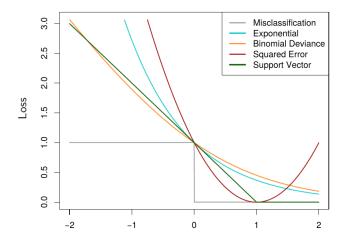
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 - Any objective functions that assigns higher loss to misclassification (e.g. is more willing to overpredict cancer than miss it)



• Loss for observation with $y_i = +1$ (vs -1). x-axis is $y \cdot f(x)$. This is also related to Hinge Loss.

• Confusion matrix: contingency table of true and predicted classes (in validation or test set)

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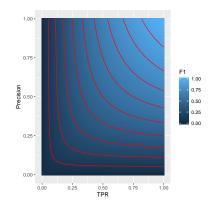
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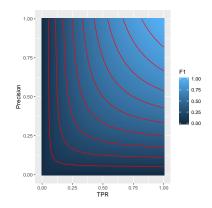
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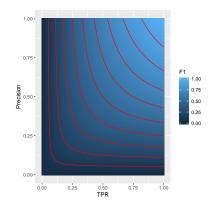
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- False negative rate: FNR = B/(B + D) = 1 TPR
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- Accuracy: (A + D)/(A + B + C + D)



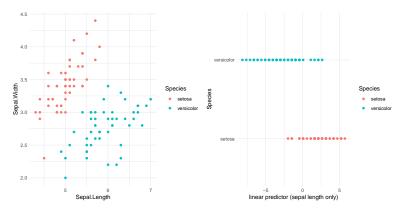
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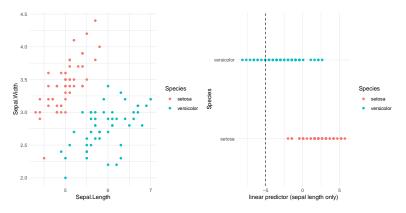
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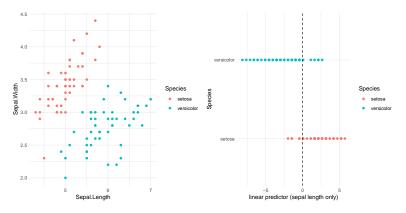
- F_1 score: An ad-hoc objective rewarding both TPR and precision
- Harmonic mean: $F_1 = \left(\frac{TPR^{-1} + Precision^{-1}}{2}\right)^{-1}$
- Greatest gain can be made by improving whichever performance measure is currently worst



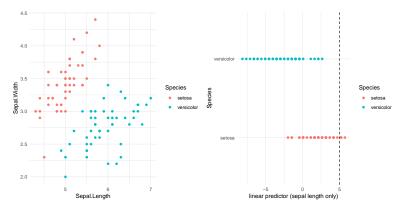
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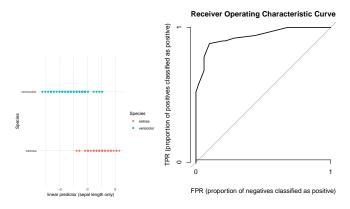
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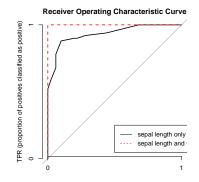
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FPR (proportion of negatives classified as positive)

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- ROC will be better if we use more variables.

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$$\begin{aligned} AUC &= \int_{-\infty}^{\infty} P(\hat{Y}_i > \theta | Y_i = +1) \cdot (-f(\theta | Y_i = -1)) d\theta \\ &= \int_{\infty}^{-\infty} P(\hat{Y}_i > \theta | Y_i = +1) \cdot f(\theta | Y_i = -1) d\theta \\ &= E[Z > Z'] \end{aligned}$$

Where Z(Z') is randomly drawn predictions from positive (negative) class and reversing sign is because increasing θ decreases FPR

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 - Then the hard confusion matrix is simply $\hat{\mathbf{Y}}^{\top}\mathbf{Y}$

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 - ► For more, see Nadeau, Claude and Yoshua Bengio. 2003. "Inference for the Generalization Error." *Machine Learning*, 53(3).

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