

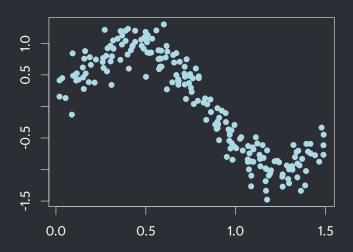
Model evaluation: training and test sets

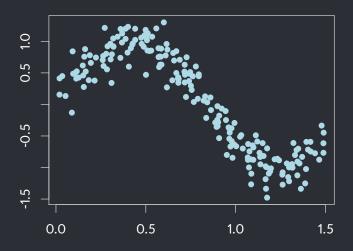
Lecture 21

STA 371G

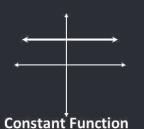
Overfitting

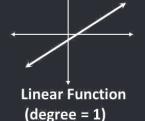
- A related problem to p-hacking is the issue of overfitting: creating a model that fits your sample very well but does not generalize well to the larger population.
- In other words, an overfit model is one where the R² (for linear regression) or prediction accuracy (for logistic regression) is high, but the model will not work as well as expected when given new data.
- Let's consider a simple problem of predicting Y from one X, and fitting a polynomial curve to the data.

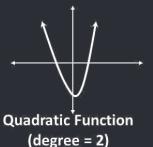




As we increase the degree (the highest power of x) of the polynomial, the fit improves, since the curve can zig and zag to capture more of the idiosyncrasies of the sample.

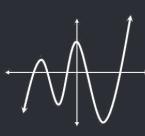










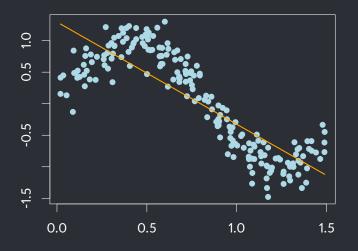


Cubic Function (deg. = 3)

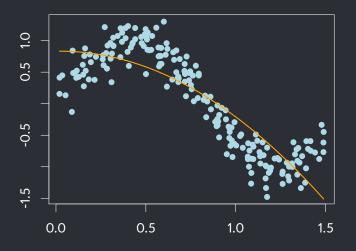


Quintic Function (deg. = 5)

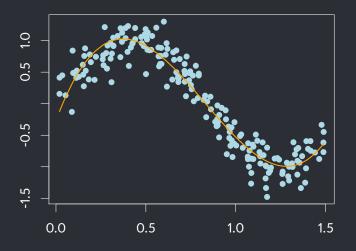
Degree 1 polynomial



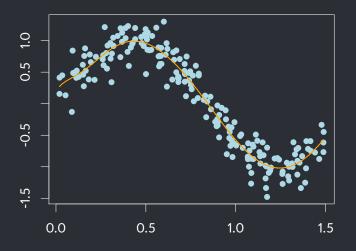
Degree 2 polynomial



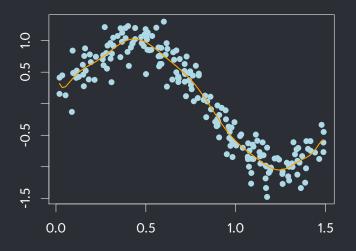
Degree 3 polynomial



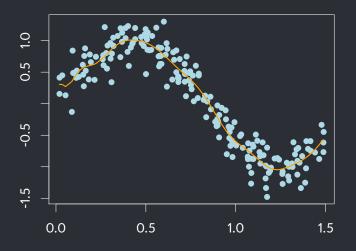
Degree 10 polynomial



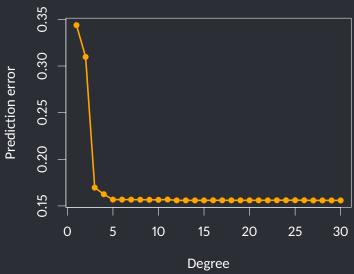
Degree 20 polynomial



Degree 30 polynomial



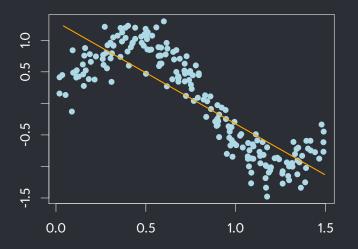
As we fit increasingly complex polynomials, the average prediction error decreases:



Parsimony

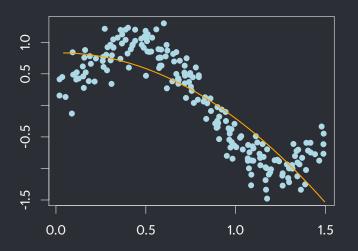
- We've talked a lot about the need to select models that are parsimonious, i.e., those that strike a good balance between fitting well and being simple.
- Besides being easier to communicate, there's another reason to prefer simpler models—they tend to generalize better to new data.
- Let's see how these models perform on a new sample from the same population!

Degree 1 polynomial



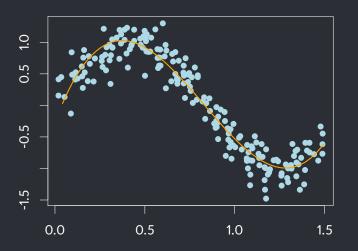
average absolute prediction error = 0.3339 (was 0.3438 in the original sample)

Degree 2 polynomial



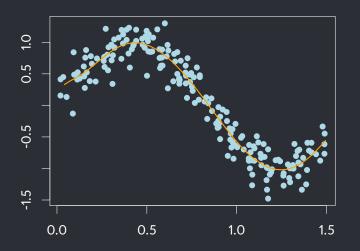
average absolute prediction error = 0.3688 (was 0.3097 in the original sample)

Degree 3 polynomial



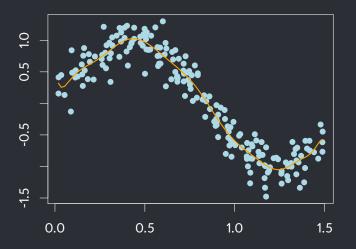
average absolute prediction error = 0.2718 (was 0.1696 in the original sample)

Degree 10 polynomial



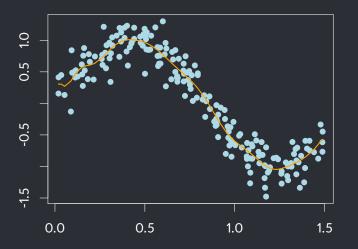
average absolute prediction error = 0.2927 (was 0.1565 in the original sample)

Degree 20 polynomial



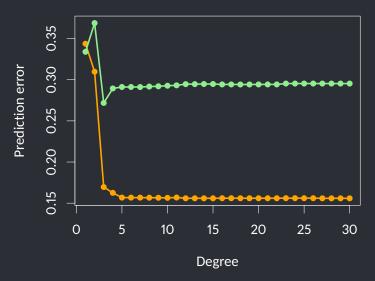
average absolute prediction error = 0.2927 (was 0.156 in the original sample)

Degree 30 polynomial



average absolute prediction error = 0.2927 (was 0.1559 in the original sample)

The increasingly complex polynomials do not perform as well on the new data as the simple polynomials, because they had overfit the idiosyncrasies of the original data.



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- In general, a model's performance on the data used to build the model will usually not be as strong as its performance on new data.
- We can use the concept of training and test sets to address both p-hacking and overfitting.

Review of training and test sets

| Original data set | |
|-------------------|----------|
| Training set | Test set |

- Split the data into a training set and a test set (a typical split is 70% training set / 30% test set).
- We use the training set to build the model, and then evaluate the quality of the model on how well it predicts Y in the test set.

The housing data set

Let's look at three different models to predict housing prices, with three different sets of predictors:

- Model 1: living area only (Living.Area)
- Model 2: living area and whether the house is on the water (Living. Area and Waterfront)
- Model 3: living area, whether the house is on the water, and whether the house uses electric, gas, or oil (Living.Area, Waterfront, and Fuel.Type)

Let's split our data into a training and test set with a rough 70/30 split. (N = 1728, and 30% of 1728 is approximately 518.)

```
# Select which row numbers will correspond to test cases
test.cases <- sample(1:1728, 518)
# Build the test set from those row numbers
test.set <- houses[test.cases,]</pre>
# Remaining row numbers will correspond to training cases
training.cases <- setdiff(1:1728, test.cases)</pre>
# Build the training set from those row numbers
training.set <- houses[training.cases,]</pre>
```

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- But this would lead to the same overfitting and p-hacking issues as before!
- Instead, set aside the test set and don't look at it until you have a final model: what you think is the most parsimonious model.
- Evaluate model performance in the test set as a way to get a fair measurement of how well your model will perform on new data.

Let's start with a simple model that uses living area only:

```
model <- lm(Price ~ Living.Area, data=training.set)</pre>
```

The training set average error is:

```
mean(abs(resid(model)))
[1] 46371.65
```

The test set average error comes from manually computing the prediction error for each case in the test set:

```
price.hat <- predict(model, test.set)
mean(abs(test.set$Price - price.hat))</pre>
```

[1] 50596.29

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```
cor(test.set$Price, price.hat)^2
[1] 0.4817818
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```
cor(test.set$Price, price.hat)^2
[1] 0.4817818
```

Compare this to what R² is in the training set:

```
summary(model)$r.squared
[1] 0.5200801
```

Using training and test sets in logistic regression

- The logic of training and test sets is identical for logistic regression.
- If we build a logistic model to predict something, using a test set for evaluation will give us a more realistic estimate of prediction accuracy on new data.
- Let's try this out on a data set of passengers from the *Titanic*.

The data set

The *Titanic* data set has data on 756 passengers on the *Titanic*; we'll predict the categorical variable survival from age of the passenger. Let's segment the data into training and test sets, as before:

```
# Select which row numbers will correspond to test cases
test.cases <- sample(1:756, 227)
# Build the test set from those row numbers
test.set <- titanic[test.cases,]</pre>
# Remaining row numbers will correspond to training cases
training.cases <- setdiff(1:756, test.cases)
# Build the training set from those row numbers
training.set <- titanic[training.cases,]</pre>
```

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model <- glm(Survived ~ Age, data=training.set)</pre>

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model <- glm(Survived ~ Age, data=training.set)
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Then we can evaluate prediction accuracy as usual:

```
predicted <- (predict(model, type="response") >= 0.5)
actual <- (training.set$Survived == 1)
sum(predicted == actual) / nrow(training.set)
[1] 0.5879017</pre>
```

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```

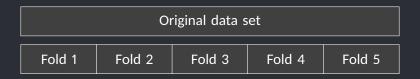
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sum(predicted == actual) / nrow(training.set)
[1] 0.5879017</pre>
```

To evaluate prediction accuracy on the training set, we use the same model, but apply it to the test set:

```
predicted <- (predict(model, test.set, type="response") >= 0.5)
actual <- (test.set$Survived == 1)
sum(predicted == actual) / nrow(test.set)
[1] 0.5814978</pre>
```

Cross-validation



- Split the data into k "folds" (here k = 5).
- For each fold, use that fold as a test set and the other folds together as a training set.
- Average the prediction accuracy across all *k* folds as your best estimate of prediction accuracy.
- Cross-validation reduces the impact of the random chance from your selection of training and test sets.