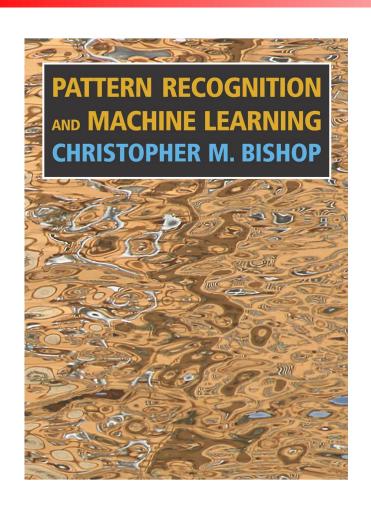
# Regression: Overfitting and Curse of Dimensionality

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Office Hour: 3-4pm Tuesday

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#### Reference



Today's class *roughly* follows Chapter 1.

Pattern Recognition and Machine Learning

Christopher Bishop, 2006

#### LS with Feature Transform

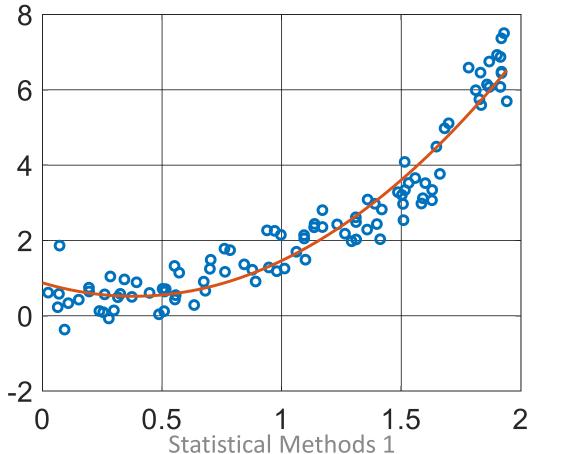
$$\mathbf{w}_{\mathrm{LS}} \coloneqq \underset{\mathbf{w}}{\operatorname{argmin}} \sum_{i \in D_0} [y_i - f(\mathbf{x}_i; \mathbf{w})]^2$$

$$f(\mathbf{x}; \mathbf{w}) \coloneqq \langle \mathbf{w}_1, \boldsymbol{\phi}(\mathbf{x}) \rangle + w_0, \mathbf{w} \coloneqq [\mathbf{w}_1, w_0]^{\mathsf{T}}$$

- $\phi(x)$  can be a collection of polynomial functions:
- $\boldsymbol{\phi}(x) \coloneqq \left[x^1, x^2, x^3 \dots x^b\right]^{\mathsf{T}}$ .
- b is called the degree of  $\phi(x)$ .

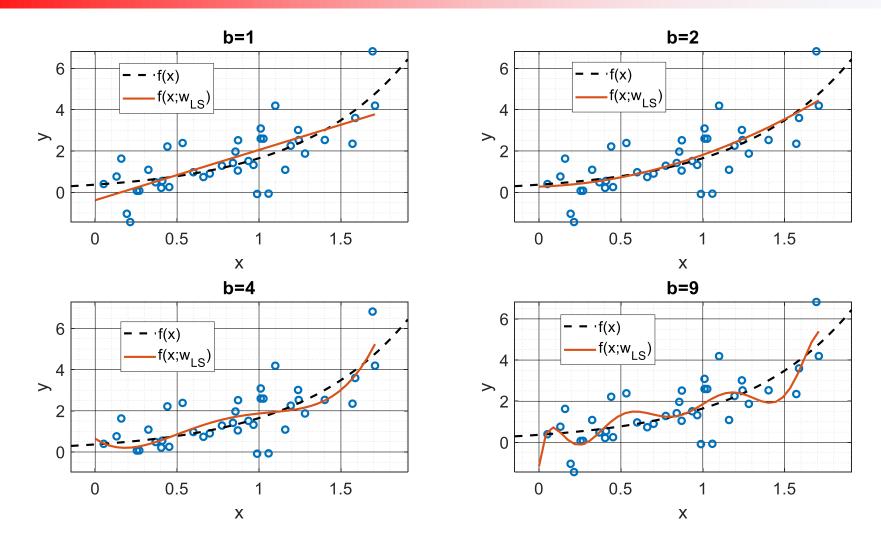
### LS with Polynomial Transform (b = 2)

- $x \sim \text{uniform}(0,2)$
- $y = f(x) + \epsilon$ ,  $f(x) = \exp(1.5x 1)$ ,  $\epsilon \sim N(0, .64)$



10/2/2023

# Poly. Transform with various b



# Poly. Feature with various b

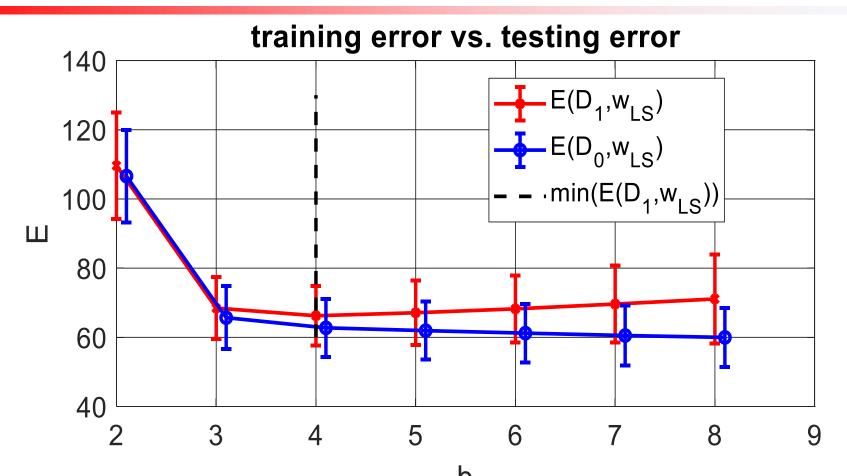
• The higher the b, the more flexible our f(x; w) is.

- However, when increasing b,
  - The fit of  $f(x; \mathbf{w}_{LS})$  first got better (b = 2).
  - then got worse (b = 4, b = 9).
  - $f(x; \mathbf{w}_{LS})$  become too "squiggly", when b is large.
  - $f(x; w_{LS})$  almost tried "too hard" to fit our data.
- Is this a general pattern?
  - We design an experiment to find out.

- We randomly split our dataset D into  $D_0$  and  $D_1$ .
  - ullet assuming D contains IID pairs.
- $w_{\rm LS}$  is fitted using  $D_0$  only.
- Define an error  $E(D', \mathbf{w}) = \sum_{i \in D} [y_i f(\mathbf{x}_i; \mathbf{w})]^2$ .
- It tells how well f(x; w) fits a specific dataset D'.
- We can have two performance metrics:
- $E(D_0, \mathbf{w}_{LS})$  is usually referred to as training error.
- $E(D_1, \mathbf{w}_{LS})$  is usually referred to as testing error.

- We do not care  $E(D_0, w_{LS})!$
- We have already seen the output in  $D_0$  during the training.
- We care performance of  $f(x; w_{LS})$  on unseen dataset  $D_1$ !
- The ability of getting low  $E(D_1, w_{\rm LS})$  is called generalization.
- Generalization is a key goal in statistical decision making.

- Go back to the example,
- As b increases, how  $E(D_0, w_{\rm LS})$  and  $E(D_1, w_{\rm LS})$  change?



Results are averaged from 100 times run with independent  $D=D_0\cup D_1$  generated by different random seeds, and are plotted with standard deviation

- Training error keeps reducing.
- $f(x; w_{LS})$  fit  $D_0$  better and better as b increases.
- Testing error drops then goes up again.
- $f(x; w_{LS})$  does not fit unseen  $D_1$  well, when b is too large.
- The problem:
- Generalization of  $f(x; w_{LS})$  deteriorates when b is too large.
- The phenomenon  $f(x; w_{\rm LS})$  fits too well on training set while underperforming on unseen datasets, is called

# Overfitting.

# Selecting b

- b should not be too small, so f is flexible enough!
- b should not be too large, so f is **not too flexible**!

How do we select?

- We can split full dataset D into  $D_0$  and  $D_1$ .
- Use  $D_0$  to fit  $f_{LS}(b)$  and use  $D_1$  to compute  $E(D_1, f_{LS}(b))$ .
- Select a b such that  $E(D_1, f_{LS}(b))$  is the lowest.
- Fit  $f_{\rm LS}$  again using the selected b on the full dataset.

# Selecting b (better approach)

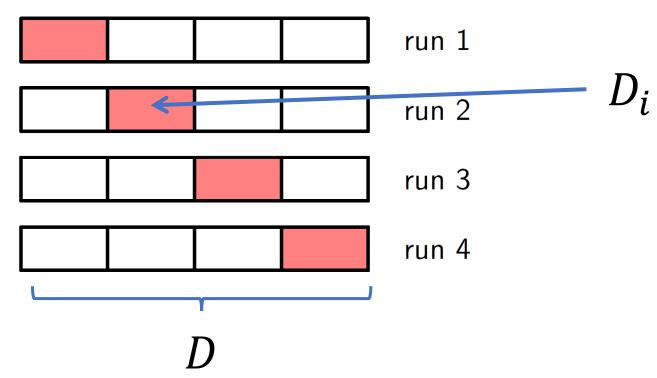
#### Problem of splitting D into $D_0$ and $D_1$ :

- 1. However, we have wasted  $D_1$  for validation.
  - What if  $D_1$  contains info that is beneficial for choosing b?
- 2.  $E(D_1, f_{LS}(b))$  is random, the selection may be random.
  - We want to take average and reduce the randomness.
- Split D into  $D_0$  and  $D_1$ , compute  $E(D_1, f_{LS}(b))$
- Swap the role of  $D_0$  and  $D_1$ , compute  $E(D_0, f'_{LS}(b))$ 
  - $f'_{LS}(b)$  is fitted using  $D_1$
- Select b that minimizes  $\frac{E\left(D_1, f_{\mathrm{LS}}(b)\right)}{2} + \frac{E\left(D_0, f'_{\mathrm{LS}}(b)\right)}{2}$

#### Cross-validation

- The extension of above idea gives rise to a commonly used model selection method: Cross-validation.
- Split D into **disjoint**  $D_0 \dots D_k$ ,
- For i = 0 to k
  - Fit  $f_{LS}^{(i)}(b)$  on all subsets but  $D_i$ ,  $\forall b$
  - Compute  $E\left(D_i, f_{\mathrm{LS}}^{(i)}(b)\right)$ ,  $\forall b$
- Select b that minimizes  $\frac{\sum_{i} E^{(i)}}{k+1}$
- k can go as high as n-1: leave-one-out-validation

#### **Cross-validation**



- PRML, Figure 1.18
- Read Chapter PRML 1.3

#### Problem of Cross-validation

- The implementation of cross-validation is easy,
- But the computational cost is high.
  - $f_{LS}^{(i)}(x; w)$  must be fitted and validated for all splits.
- The effectiveness of cross-validation depends on the IID assumption of our dataset  ${\cal D}$ .
  - Validation set and the training set must be IID!
  - Which may not hold in reality: e.g. stock price dataset.

 Can we avoid overfitting without splitting our dataset for validation? We will discuss this in the future.

# Polynomial Transform on Higher Dimensional Dataset

- So far, we only considered polynomial transform on one dimensional dataset, i.e.,  $x \in R$
- What about  $x \in \mathbb{R}^d$ , when the output y depends on multiple inputs?
- When  $x \in \mathbb{R}^d$ ,
  - $\phi(x) := [h(x^{(1)}), h(x^{(2)}), ..., h(x^{(d)})]^{\top}$ .
  - $h(t) := [t^1, t^2, ..., t^b] \in R^b$ .
  - $\phi(x) \in R^{db}$ , which means  $w_1 \in R^{db}$ .
- This does not include cross-dimension polynomials.
  - e.g.,  $x^{(1)}$   $x^{(2)}$ ,  $x^{(1)}$   $x^{(2)}$   $x^{(3)}$ , ...
  - These can be useful as the output value may depends jointly on several inputs. e.g. blood pressure <- (weight, height)</li>

# Polynomial Transform on Higher Dimensional Dataset

- To include **pairwise** cross-dimension polynomials, we can slightly redesign  $\phi(x)$ :
  - $\phi(x) := [h(x^{(1)}), ..., h(x^{(d)}), \forall_{u < v} x^{(u)} x^{(v)}]$
  - $\phi(x) \in R^{db+\binom{d}{2}}$ ,
- Similarly, we can include all the triplets:
  - $\phi(x)$ : =  $[h(x^{(1)}), ..., h(x^{(d)}), \forall_{u < v} x^{(u)} x^{(v)}, \forall_{u < v < w} x^{(u)} x^{(v)} x^{(w)}]$
  - $\phi(x) \in R^{db+\binom{d}{2}+\binom{d}{3}}$ ,
- and we can go on to include quadruplets...

# **Curse of Dimensionality**

- We can include cross terms all the way up to d-plets.
- Unfortunately, we know

$$\bullet \binom{d}{1} + \binom{d}{2} + \binom{d}{3} + \binom{d}{4} + \cdots \binom{d}{d} = 2^d$$

- We have not yet included cross terms like:
  - $[x^{(u)}]^2 x^{(v)} ...$
- The output dimension of  $\phi(x)$  can grow exponentially with dimensionality d and this is a bad news...

# **Curse of Dimensionality**

• We have seen in yesterday's homework, the number of observations n, needs to at least match the output dimension of  $\phi(x)$ , otherwise, we cannot obtain  $w_{LS}$ !

- It means we need to grow n exponentially with d!
- Imagine a problem with d=100.
  - A terabyte-data on hard-drive contains  $2^{40}$  bytes.

# **Curse of Dimensionality**

- The phenomenon, that the number of observations needed to solve a problem grows exponentially with d exists in many statistical learning tasks.
- They are collectively called "Curse of Dimensionality".

• This phenomenon forbids us solving highdimensional problems.

#### Conclusion

- We introduce poly. transform to our prediction func. f.
- This increases the flexibility of f, but we also see this additional complexity caused two major problems:

#### Overfitting

- The generalization of f is poor.
- Curse of Dimensionality
  - n needs to grows exponentially with the dimensionality of x.
- Next week, we will introduce a way to reduce the flexibility of f to combat overfitting and the probabilistic idea behind it.

#### Literature Review

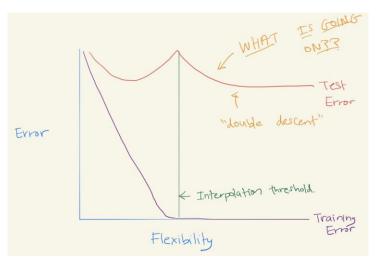
- Read the paper (stay at high level):
- 1906.11300.pdf (arxiv.org)

Answer the questions in plain words (or pictures):

- 1. What is the statistical learning problem that authors try to study?
- 2. What is Benign Overfitting?
- 3. Why does benign overfitting happen?

# Um... Actually...

• In recent years, people realize (particularly in the context of deep neural network models), there is a phenomenon called double descent:



The testing error **comes back and drops again** as the flexibility of your prediction function increases!



Read this <u>twitter thread</u> by Daniela Witten.

# Computing Lab

- Download "Prostate Cancer dataset", description, dataset.
- Implement a Least-square solver using R. Do not use builtin functions.
- Fit f(x; w) using classic linear least squares.
- Calculate the cross-validation error.
- How does the cross-validation testing error change if you remove one of the features?
  - How do you explain this using what we have learned today?