



# Parameter Estimation

Chris Piech  
CS109, Stanford University

# General “Inference”



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## WebMD Symptom Checker BETA

INFO

SYMPOMS

QUESTIONS

CONDITIONS

DETAILS

TREATMENT

Add more symptoms

Type your main symptom here

AGE 30

GENDER Male

### MY SYMPTOMS

cough ×

throat irritation ×

sneezing ×

or Choose common symptoms

bloating

cough

diarrhea

dizziness

fatigue

fever

headache

muscle cramp

nausea

throat irritation

Results Strength: MODERATE



Previous

Info

Continue



## Conditions that match your symptoms

### UNDERSTANDING YOUR RESULTS

Influenza (flu) adults



 Moderate match

Pneumococcal infections



 Moderate match

H1N1 Flu Virus (Swine Flu)



 Moderate match

Bacterial Pneumonia



 Moderate match

Sepsis (blood infection)



 Moderate match

Gender Male

Age 30

Edit

My Symptoms

Edit

fever 103f to 104f, dizziness,

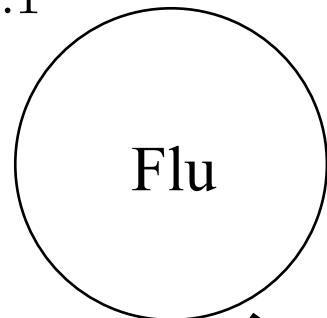
throat irritation, migraine headache



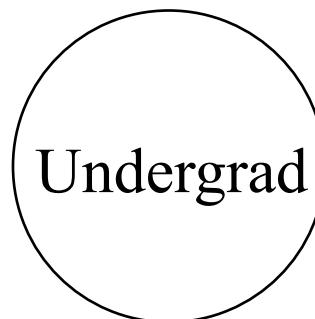
Start Over

# Probabilistic Model

$$P(Fl = 1) = 0.1$$

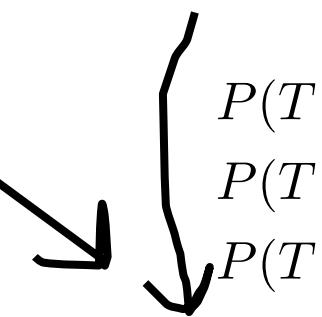
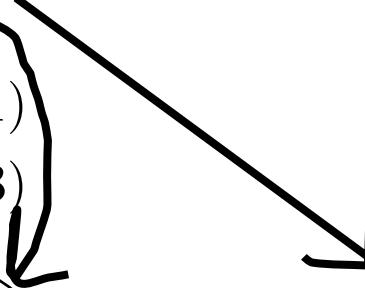
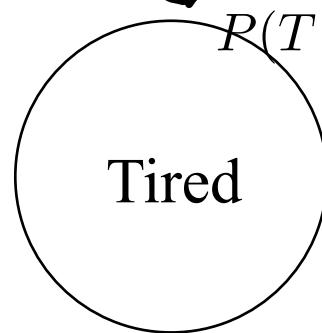
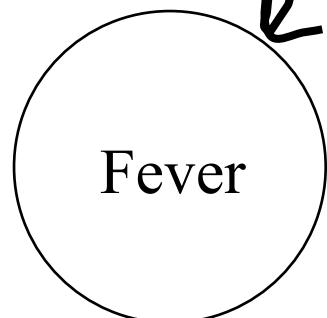


$$P(U = 1) = 0.8$$



$$Fev|Flu = 0 \sim N(100.0, 1.81)$$

$$Fev|Flu = 1 \sim N(98.25, 0.73)$$



$$P(T = 1|Flu = 0, U = 0) = 0.1$$

$$P(T = 1|Flu = 0, U = 1) = 0.8$$

$$P(T = 1|Flu = 1, U = 0) = 0.9$$

$$P(T = 1|Flu = 1, U = 1) = 1.0$$

# Alg #1: Joint Sampling

```
3 N_SAMPLES = 100000
4
5 # Program: Joint Sa
6 #
7 # we can answer any
8 # with multivariate
9 # where conditioned
10 def main():
11     obs = getObservation()
12     print 'Observation =', obs
13
14     samples = sample(N_SAMPLES)
15     prob = probFluGivenSamples(samples, obs)
16     print 'Pr(Flu | Obs) =', prob
```

[0, 0, 0, 0]  
[0, 1, 0, 1]  
[1, 0, 1, 0]  
[1, 1, 1, 1]  
[0, 1, 0, 1]  
[0, 1, 0, 0]  
[0, 0, 0, 0]  
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[0, 0, 0, 0]  
[0, 0, 0, 0]  
[1, 1, 1, 1]  
[0, 1, 0, 0]  
Observation = [None, None, None, 1]  
Pr(Flu | Obs) = 0.140635888502  
>

Each one of  
these is one  
posterior  
sample:

# Alg #2: MCMC

```
webkit -- bash -- 10x20
[1, 1, 101.0, 1]
[1, 1, 101.0, 1]
[0, 1, 101.0, 0]
[0, 0, 101.0, 0]
[1, 0, 101.0, 1]
[1, 0, 101.0, 0]
[1, 0, 101.0, 1]
[1, 0, 101.0, 1]
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[1, 1, 101.0, 1]
[1, 1, 101.0, 1]
[1, 0, 101.0, 1]
[1, 1, 101.0, 1]
[1, 1, 101.0, 1]
Pr(Flu) = 0.9773
>
```

MCMC is a way to sample  
with conditioned variables  
fixed

Each one of  
these is one  
posterior  
sample:

[Flu, Undergrad, Fever, Tired]



# Alg #2: MCMC

All Samples = []

---

Flu                  Undergrad                  Fever  
↓                  ↓                  ↓  
Initial Sample = [0, 0, 101.0, 1]                  Tired

# Alg #2: MCMC

All Samples = []

---

Flu              Undergrad      Fever      Tired  
↓              ↓              ↓              ←  
 $S^{(0)} = [0, 0, 101.0, 1]$

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓            ↓            ↓  
 $S^{(0)} = [0, 0, 101.0, 1]$       Tired



From  $S_t$  make  $S_{t+1}$

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

$$S^{(1)} = [0, 0, 101.0, 1]$$

Flu      Undergrad      Fever      Tired

The diagram illustrates the state vector  $S^{(1)}$  with four components. The first component, '0', is highlighted with a red circle. Blue arrows connect the labels 'Flu', 'Undergrad', 'Fever', and 'Tired' to the second, third, and fourth components of the vector, respectively.

$$P(Flu = 1 | \text{All others})$$

$$= P(Flu = 1 | Und = 0, Fev = 98.3, Tir = 1)$$

$$= 0.21$$

$$Flu_1 = \text{Sample} \left[ P(Flu = 1 | \text{All others}) \right]$$

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓                  ↓                  ↓  
 $S^{(1)} = [1, 0, 101.0, 1]$       Tired

$$P(Flu = 1 | \text{All others})$$

$$= P(Flu = 1 | Und = 0, Fev = 98.3, Tir = 1)$$

$$= 0.21$$

$$Flu_1 = \text{Sample} \left[ P(Flu = 1 | \text{All others}) \right]$$

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓      ↓      ↓  
 $S^{(1)} = [1, \textcircled{0}, 101.0, 1]$       Tired

$$P(Und = 1 | \text{All others})$$

$$\begin{aligned} &= P(Und = 1 | Flu = 1, Fev = 98.3, Tir = 1) \\ &= 0.91 \end{aligned}$$

$$Und_1 = \text{Sample} \left[ P(Und = 1 | \text{All others}) \right]$$

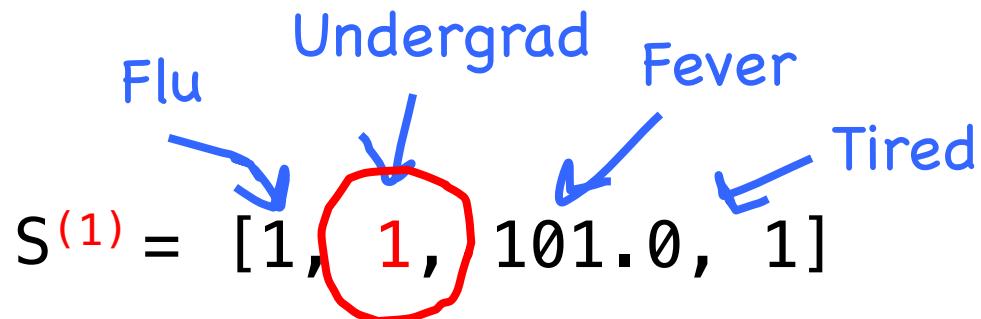
# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

$S^{(1)} = [1, 1, 101.0, 1]$

Flu      Undergrad      Fever      Tired



$$P(Und = 1 | \text{All others})$$

$$= P(Und = 1 | Flu = 1, Fev = 98.3, Tir = 1)$$

$$= 0.91$$

$$Und_1 = \text{Sample} \left[ P(Und = 1 | \text{All others}) \right]$$

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓      ↓      ↙  
 $S^{(1)} = [1, 1, 101.0, 1]$       ↙  
                Tired

Let's say you are conditioning on fever being 101.0...  
then don't change that value

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓      ↓      ↙  
 $S^{(1)} = [1, 1, 101.0, 1]$       ↙  
                Tired

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓      ↓      ↓  
 $S^{(1)} = [1, 1, 101.0, 1]$       Tired

The diagram shows a vector  $S^{(1)}$  with four elements: 1, 1, 101.0, and 1. Above the vector, four labels are positioned: 'Flu' with an arrow pointing to the first element, 'Undergrad' with an arrow pointing to the second element, 'Fever' with an arrow pointing to the third element, and 'Tired' with an arrow pointing to the fourth element. The third element, '101.0', is circled with a red marker.

# Alg #2: MCMC

All Samples =  $[S^{(0)}]$

---

Flu      Undergrad      Fever  
↓      ↓      ↙      ↙  
 $S^{(1)} = [1, 1, 101.0, 1]$       Tired

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

---

Flu      Undergrad      Fever      Tired  
↓      ↓      ↙      ↙  
 $S^{(1)} = [1, 1, 101.0, 1]$

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

---

Flu      Undergrad      Fever  
↓              ↓              ↓  
 $S^{(2)} = [1, 1, 101.0, 1]$   
Tired

$$P(Flu = 1 | \text{All others})$$

$$Flu_1 = \text{Sample} \left[ P(Flu = 1 | \text{All others}) \right]$$

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

---

Flu      Undergrad      Fever  
↓              ↓              ↓  
 $S^{(2)} = [1, 1, 101.0, 1]$

$$P(Flu = 1 | \text{All others})$$

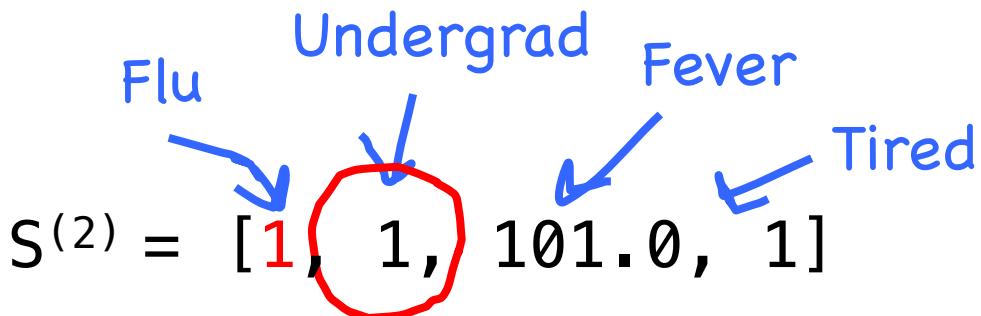
$$Flu_1 = \text{Sample} \left[ P(Flu = 1 | \text{All others}) \right]$$

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

---

Flu      Undergrad      Fever  
S<sup>(2)</sup> = [1, 1, 101.0, 1]  
Tired



$$P(Flu = 1 | \text{All others})$$

$$Flu_1 = \text{Sample} \left[ P(Flu = 1 | \text{All others}) \right]$$

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

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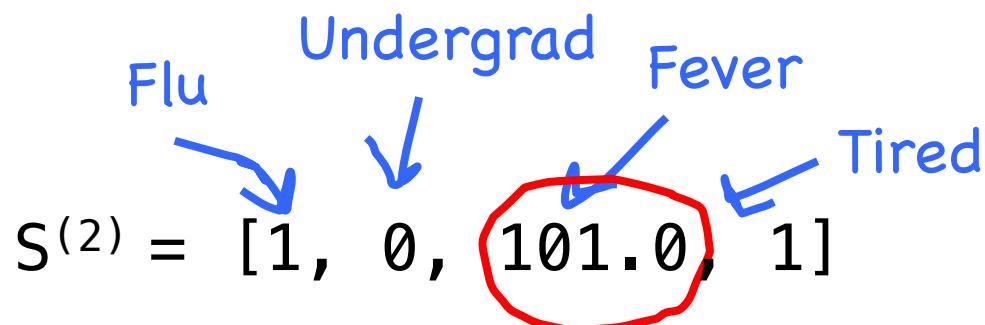
Flu      Undergrad      Fever  
↓      ↓      ↓  
 $S^{(2)} = [1, \textcircled{0}, 101.0, 1]$       Tired

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

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Flu      Undergrad      Fever  
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 $S^{(2)} = [1, 0, 101.0, 1]$       Tired



# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

---

Flu      Undergrad      Fever      Tired

$S^{(2)} = [1, 0, 101.0, 1]$

The diagram shows handwritten annotations for the MCMC samples. Above the vector  $S^{(2)}$ , four labels are written in blue: "Flu", "Undergrad", "Fever", and "Tired". Blue arrows point from each label to its respective element in the vector. The vector itself is given as  $S^{(2)} = [1, 0, 101.0, 1]$ . The number "101.0" is circled in red.

# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}]$

---

Flu      Undergrad      Fever      Tired

$S^{(2)} = [1, 0, 101.0, 1]$

The diagram illustrates the state vector  $S^{(2)}$  with handwritten labels and arrows indicating specific components. The labels are 'Flu' (pointing to the first element), 'Undergrad' (pointing to the second element), 'Fever' (pointing to the third element), and 'Tired' (pointing to the fourth element). The fourth element, '1', is circled in red.

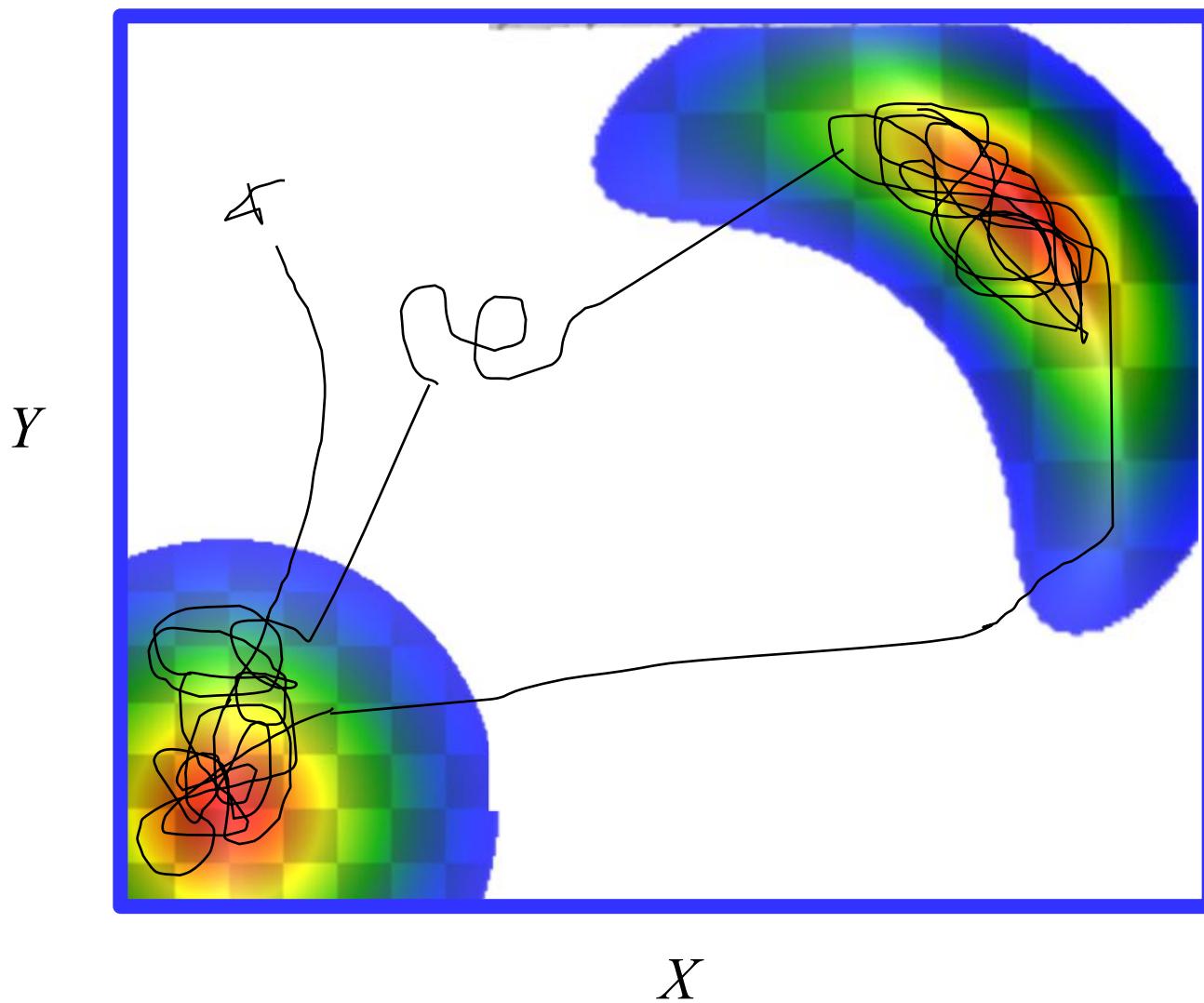
# Alg #2: MCMC

All Samples =  $[S^{(0)}, S^{(1)}, S^{(2)}]$

---

Flu      Undergrad      Fever      Tired  
↓      ↓      ↘      ↙  
 $S^{(2)} = [1, 0, 101.0, 1]$

# Alg #2: MCMC



# BAE's Theorem?

$$P(A \mid B E) = \frac{P(\textcolor{blue}{B} \mid A E) P(A \mid E)}{P(B \mid E)}$$



# $P(F = 1 \mid \text{all other rvs})$

Know:  $P(\text{symptom} \mid \text{flu, undergrad})$        $P(\text{flu})$        $P(\text{undergrad})$

Flu is independent of undergrad

Tired and fever are conditionally independent given flu, undergrad

---

$$P(F = 1 \mid \text{all other rvs})$$

$$= P(F = 1 \mid \text{symptoms}, U = u)$$

$$= \frac{P(\text{symptoms} \mid F = 1, U = u)P(F = 1 \mid U = u)}{P(\text{symptoms} \mid U = u)}$$

$$\propto P(\text{symptoms} \mid F = 1, U = u)P(F = 1 \mid U = u)$$

$$\propto P(F = 1)P(\text{symptoms} \mid F = 1, U = u)$$

$$\propto P(F = 1) \prod_i P(\text{symptom}_i \mid F = 1, U = u)$$

$$P(F = 1 | \text{all other rvs}) \propto P(F = 1) \prod_i P(\text{symptom}_i | F = 1, U = u)$$

```
120 def sampleFlu(sample):
121     f1 = getFluPr1(sample)
122     f0 = getFluPr0(sample)
123     p1 = f1 / (f1 + f0)
124     return bern(p1)
125
126 def getFluPr0(sample):
127     _, und, fev, tir = sample
128     pFlu0 = 0.9
129     pFev = getPrFeverX(fev, flu=0)
130     pTir = getPrTiredX(tir, und=und, flu=0)
131     return pFlu0 * pFev * pTir
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133 def getFluPr1(sample):
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135     pFlu1 = 0.1
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$$P(F = 1 | \text{all other rvs}) \propto P(F = 1) \prod_i P(\text{symptom}_i | F = 1, U = u)$$

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$$P(F = 1 \mid \text{all other rvs}) \boxed{\propto} P(F = 1) \prod_i P(\text{symptom}_i | F = 1, U = u)$$

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```

See you soon!

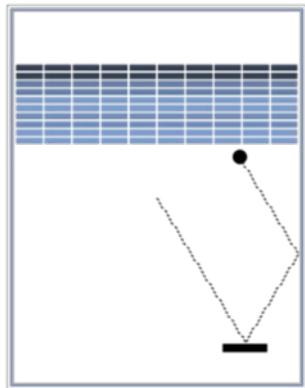
# Summary

# General Inference Summary



- **Straight Math** is fast, but can be prohibitively hard for complex models (see hw).
- **Joint Sampling** is really easy to program but fails for continuous variables (and when what you are conditioning on is rare)
- **MCMC** works well when conditioning on rare events, but is *much* harder to code / derive.
- All sampling is **slow**.

# Insight



Let  $x$  be a student's program

Let  $y$  be student mistakes

## Label Console

✓ Num Done: 8273

### Strategy

- Beeper Boundary (most people do this)
- Triangle Strategy
- Recursive Strategy

### Looping

- Correct use of looping
- Doesn't use a while
- Doesn't have correct stop condition
- Body is missing statements
- Body has extra statements
- Body order is incorrect
- Sets up initial precondition
- Does not get nesting
- Loop post condition doesn't match precondition
- Repetition of bodies

Feedback task:

$$P(y|x)$$

Hard for humans  
Hard for computers

Joint sample:

$$(\hat{x}, \hat{y}) \sim P(x, y)$$

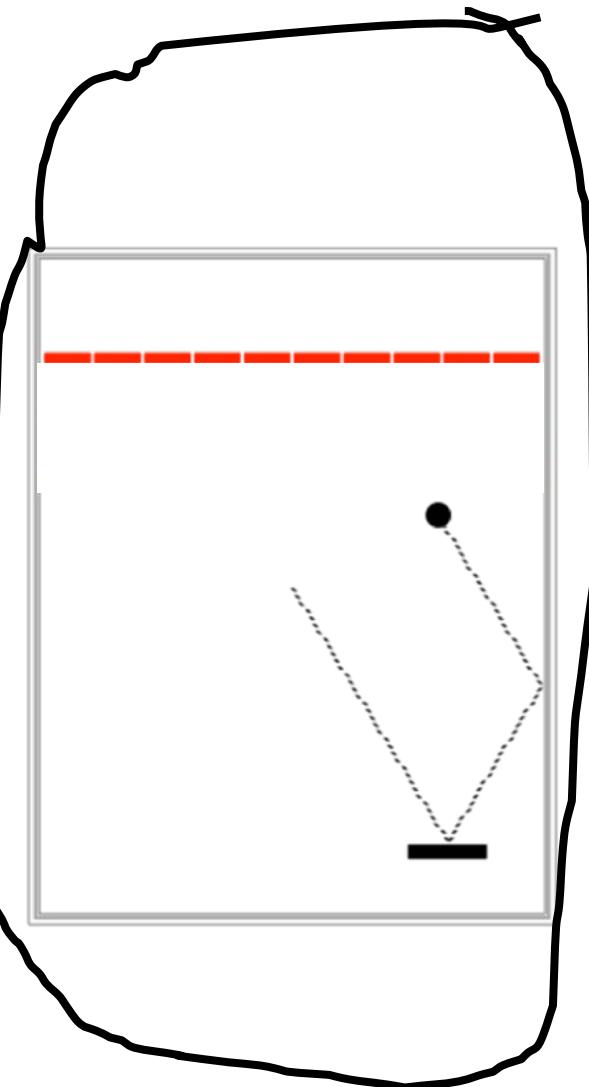
Easy for humans



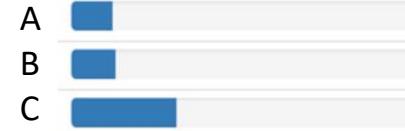
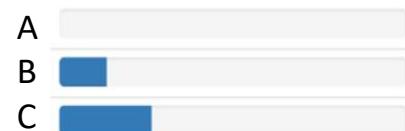
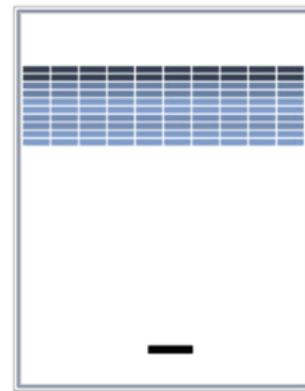
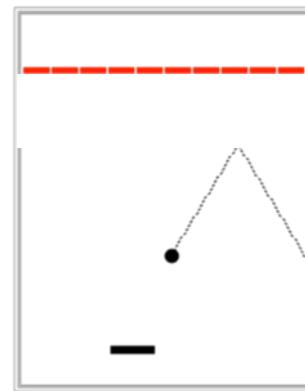
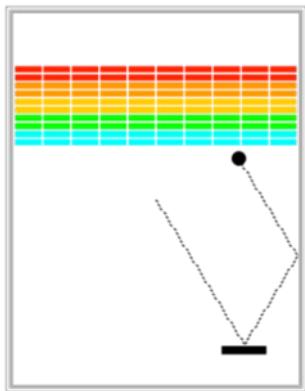
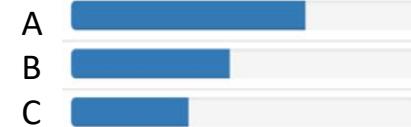
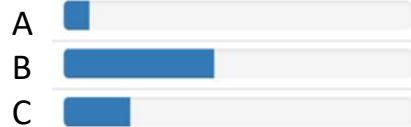
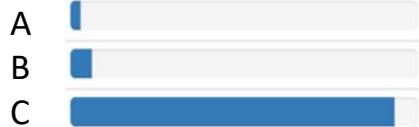
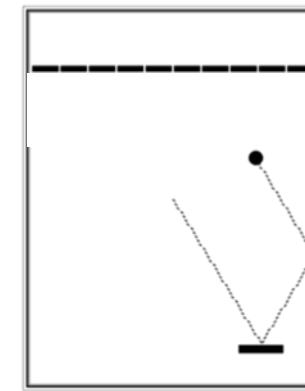
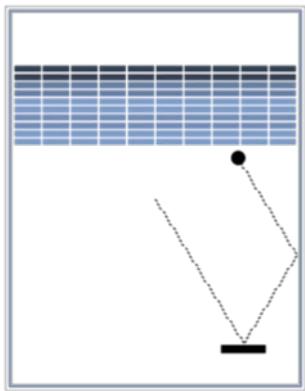
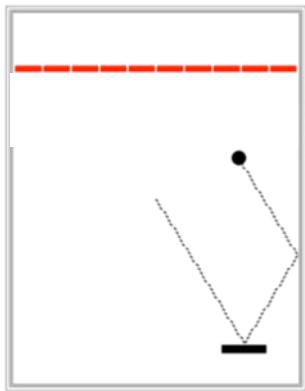
# Imagine Students

$$(\hat{x}, \hat{y}) \sim P(x, y)$$

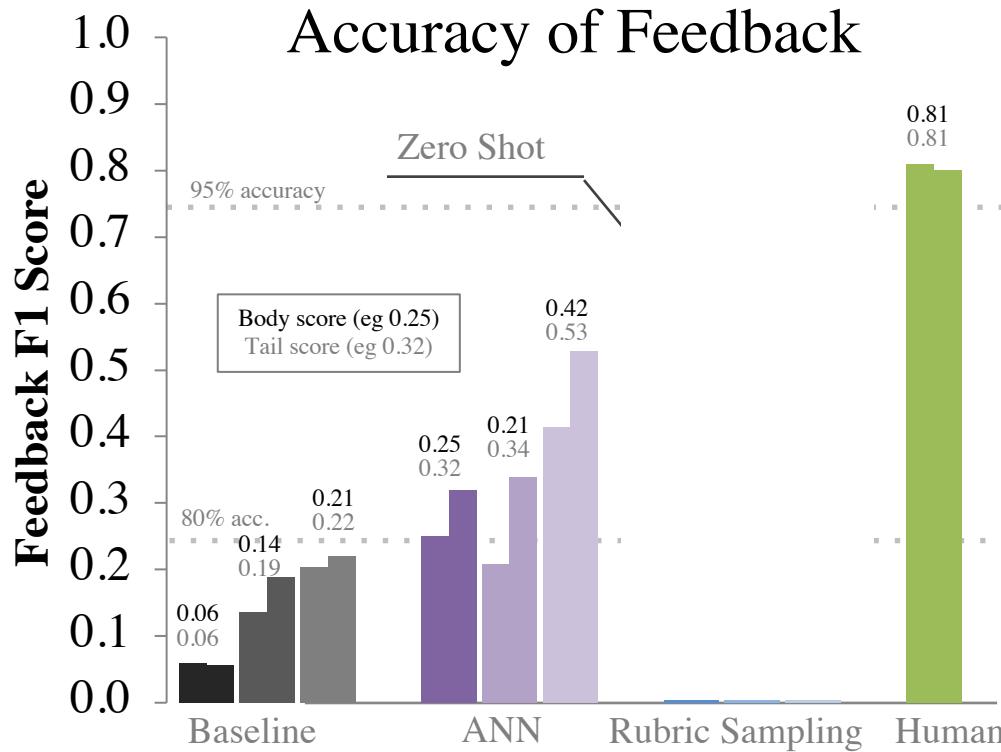
- Struggle with double for loops
- Confuses logic for deleting bricks



# Start Imagining Students



# Zero Shot Learning



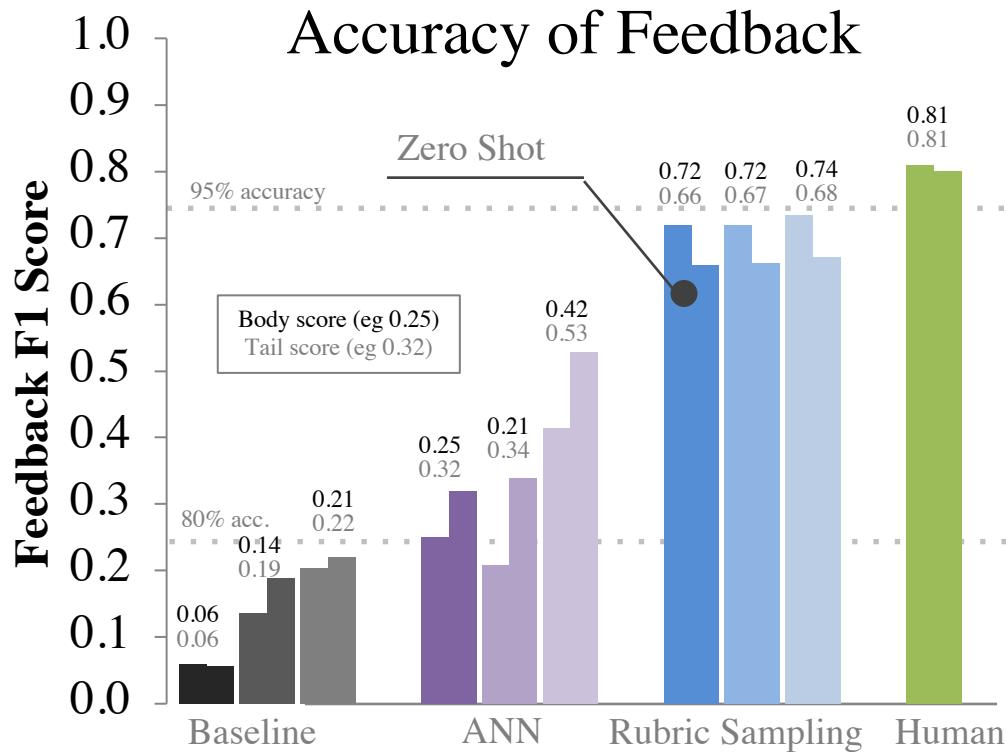
- Majority Vote
- Syntactic Analysis
- Program Output \*

- FNN
- RNN
- MVAE

■ Expert Human



# Zero Shot Learning



■ Majority Vote  
■ Syntactic Analysis  
■ Program Output \*

■ FNN  
■ RNN  
■ MVAE

■ Rubric Sampling, Zero shot  
■ Rubric Sampling, Learned  $\theta$   
■ Rubric Sampling, MVAE

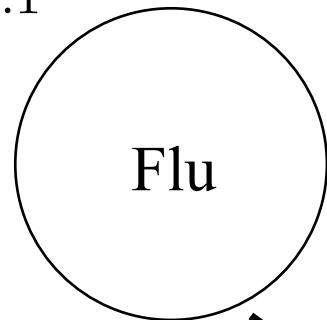
■ Expert Human

Model	Amount of Correct Feedback
Predicting from output	1,483,157 (86.0%)
Rubric sampling with MVAE	<b>1,610,020 (93.7%)</b>
Expert human	1,658,162 (96.2%)

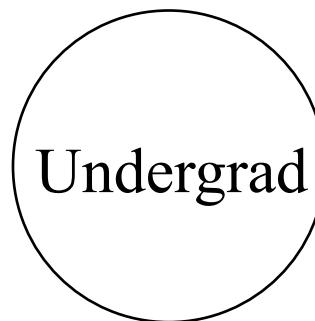


# Where Do The Numbers Come From?

$$P(Fl = 1) = 0.1$$

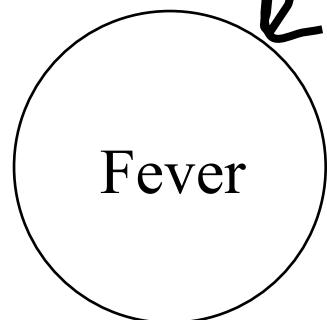


$$P(U = 1) = 0.8$$



$$Fev|Flu = 0 \sim N(100.0, 1.81)$$

$$Fev|Flu = 1 \sim N(98.25, 0.73)$$



$$P(T = 1|Flu = 0, U = 0) = 0.1$$

$$P(T = 1|Flu = 0, U = 1) = 0.8$$

$$P(T = 1|Flu = 1, U = 0) = 0.9$$

$$P(T = 1|Flu = 1, U = 1) = 1.0$$

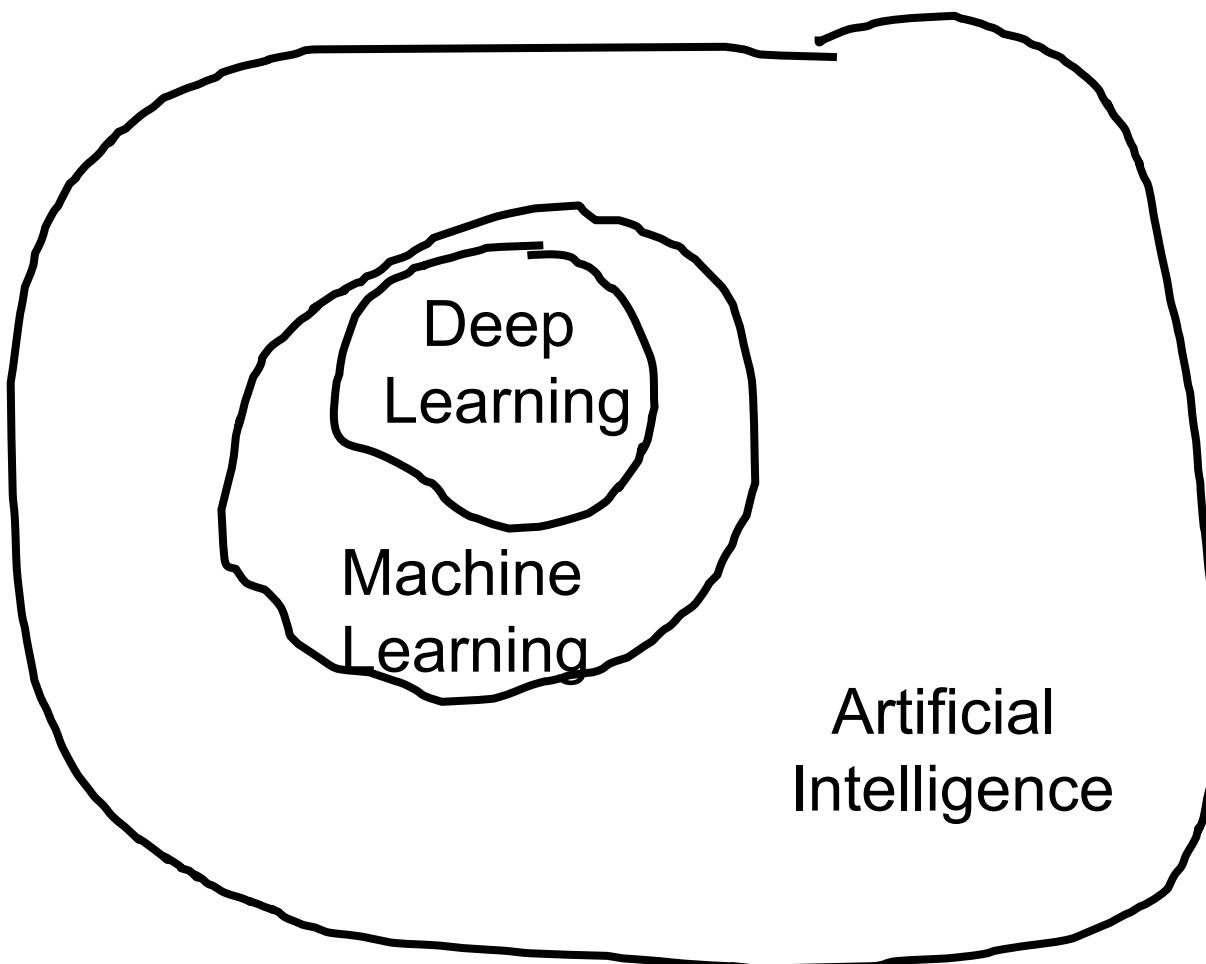
# Pedagogical Pause

At this point, if you are given a *model*,  
with all the involved probabilities, you  
can make predictions

But what if you want to *learn* the probabilities in the model?

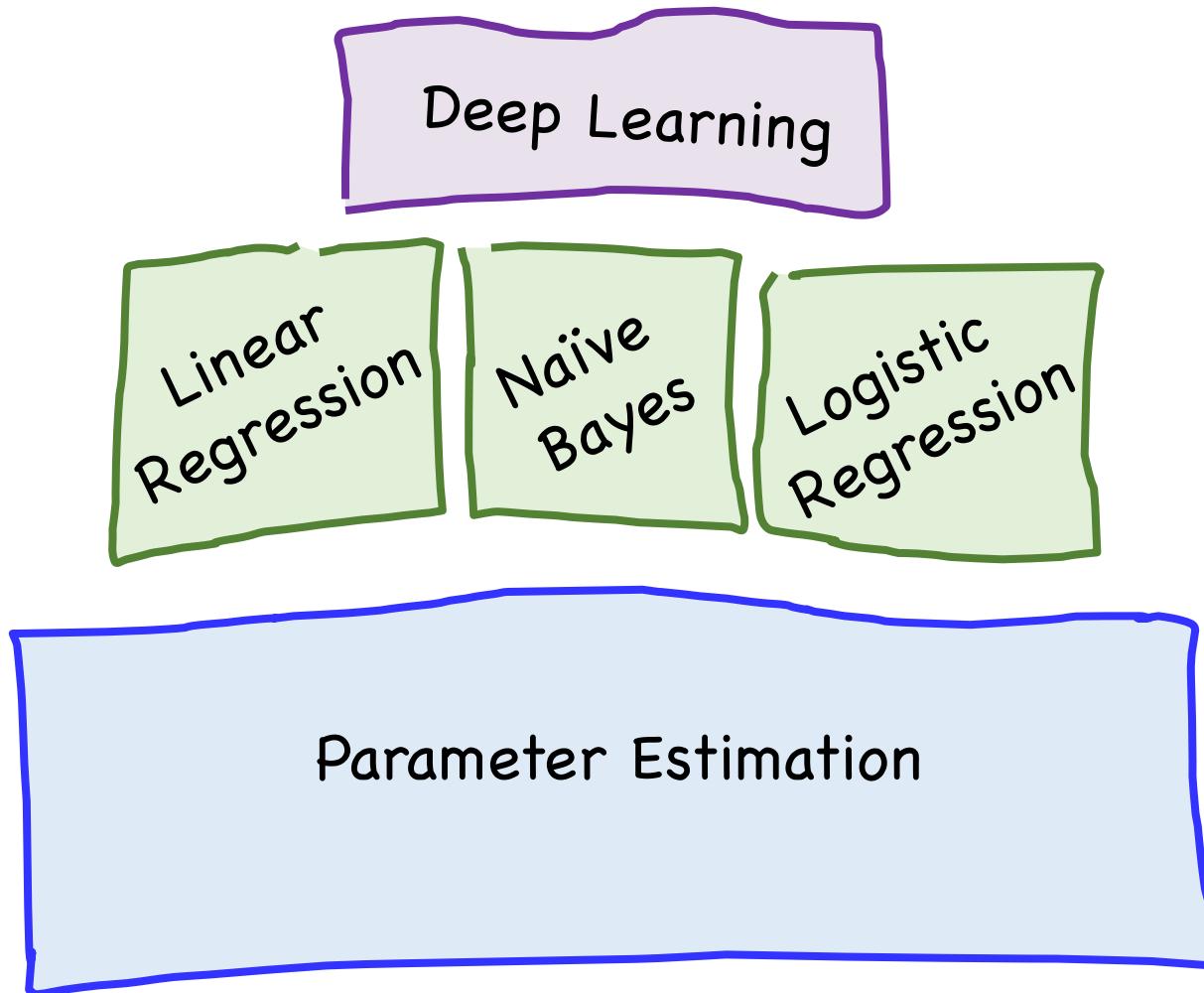
# Machine Learning

# AI and Machine Learning

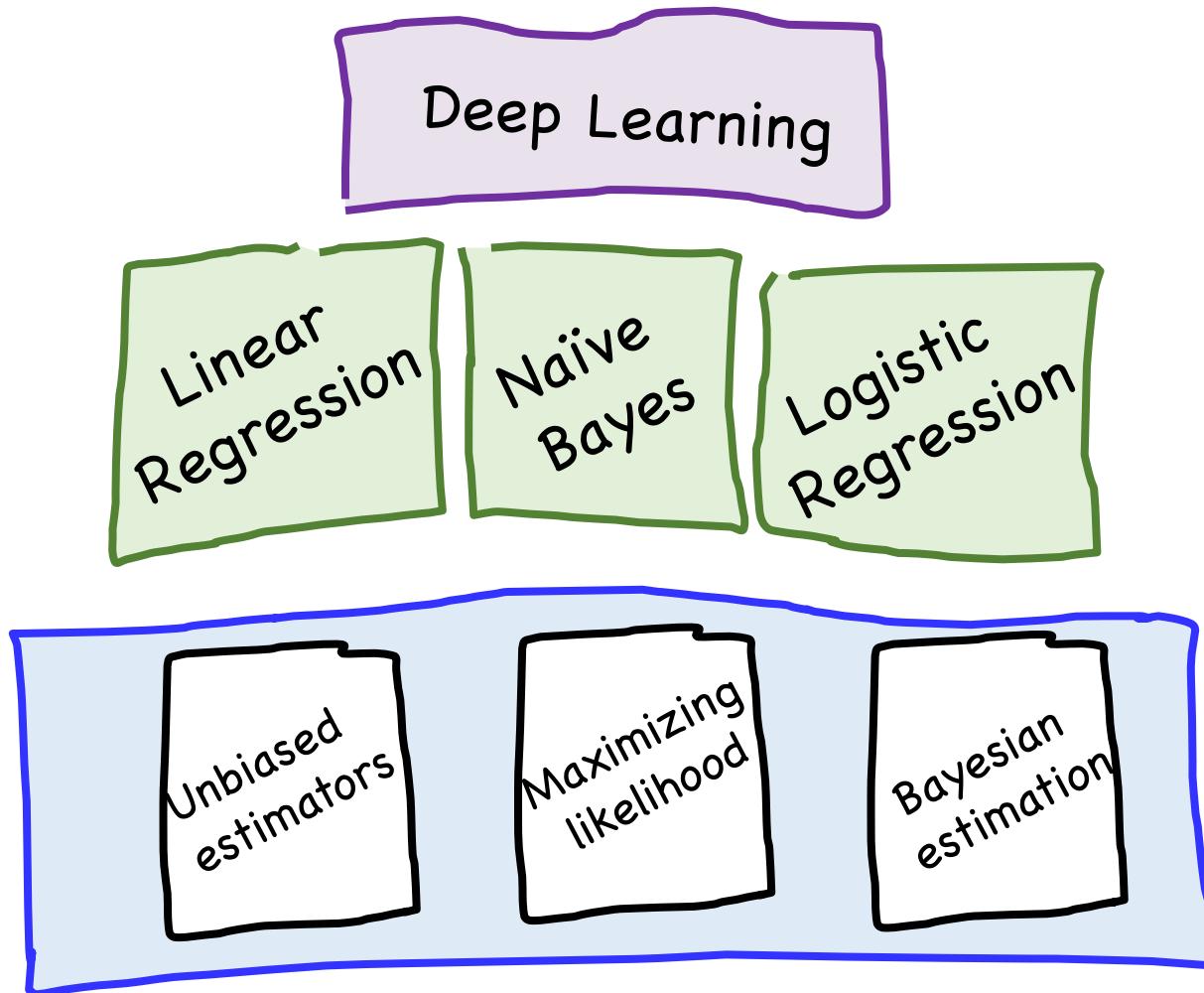


ML: Rooted in probability theory

# Our Path



# Our Path



# Jump Straight to Deep Learning?

Tensor Flow



# Jump Straight to Deep Learning?



Understand the theory to help you debug

But another reason...

# Machine Learning Uses a Lot of Data



# One Shot Learning

Single training example:

କୁ

Test set:

a	ଶ	ଅ	ଶ
କୁ	ଅ	ପ୍ର	କୁ
ମ	କୁ	ହେ	କୁ
ମ	ଅ	କୁ	ନ୍ତର

# One Shot Learning

Single  
training  
example:



Computers struggle...

... especially for **human** problems.

Understand the theory  
to push on the **grand challenges**

A silhouette of the iconic Disney castle is positioned behind the text, its spires and towers reaching towards the top of the frame.

WALT DISNEY  
PICTURES



Once upon a time...

...there was parameter estimation

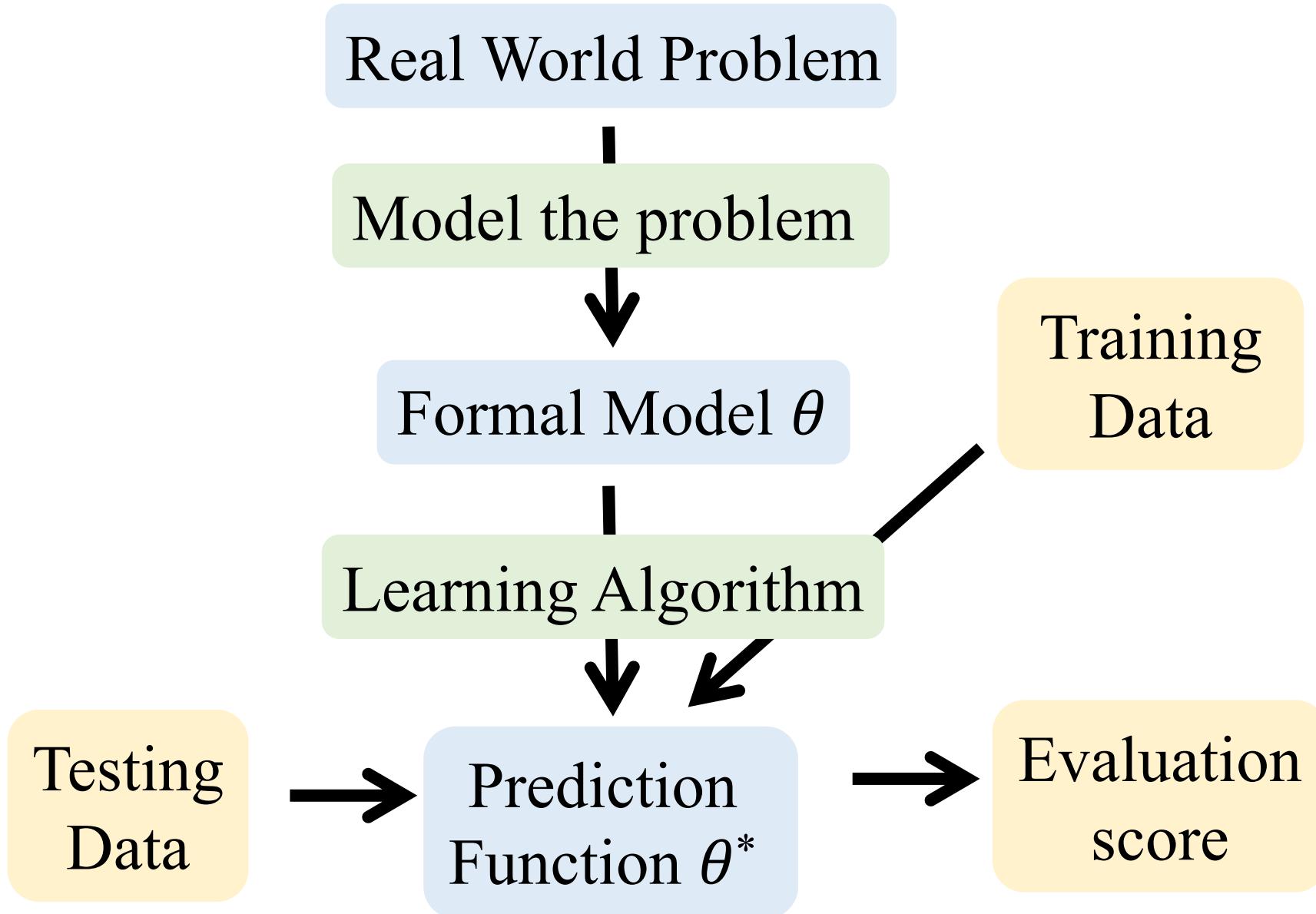
# What are Parameters?

- Consider some probability distributions:
  - $\text{Ber}(p)$   $\theta = p$
  - $\text{Poi}(\lambda)$   $\theta = \lambda$
  - $\text{Uni}(\alpha, \beta)$   $\theta = (\alpha, \beta)$
  - $\text{Normal}(\mu, \sigma^2)$   $\theta = (\mu, \sigma^2)$
  - $Y = mX + b$   $\theta = (m, b)$
  - etc...
- Call these “parametric models”
- Given model, **parameters** yield actual distribution
  - Usually refer to parameters of distribution as  $\theta$
  - Note that  $\theta$  that can be a vector of parameters

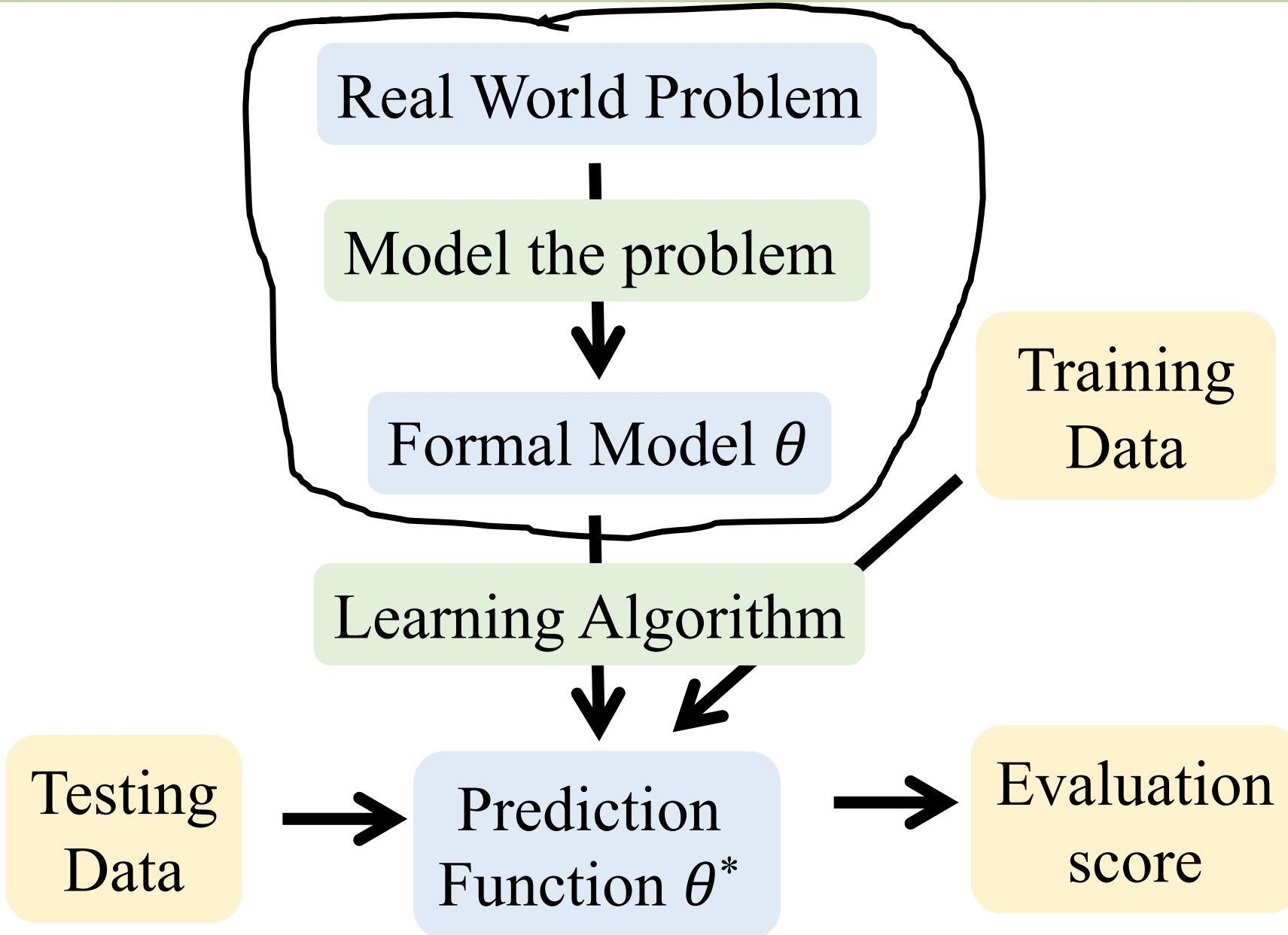
# Why Do We Care?

- In real world, don't know "true" parameters
  - But, **we do get to observe data**
    - E.g., number of times coin comes up heads, lifetimes of disk drives produced, number of visitors to web site per day, etc.
  - Need to estimate model parameters from data
  - "Estimator" is random variable estimating parameter
- Estimate of parameters allows:
  - Better understanding of process producing data
  - Future **predictions** based on model
  - Simulation of processes

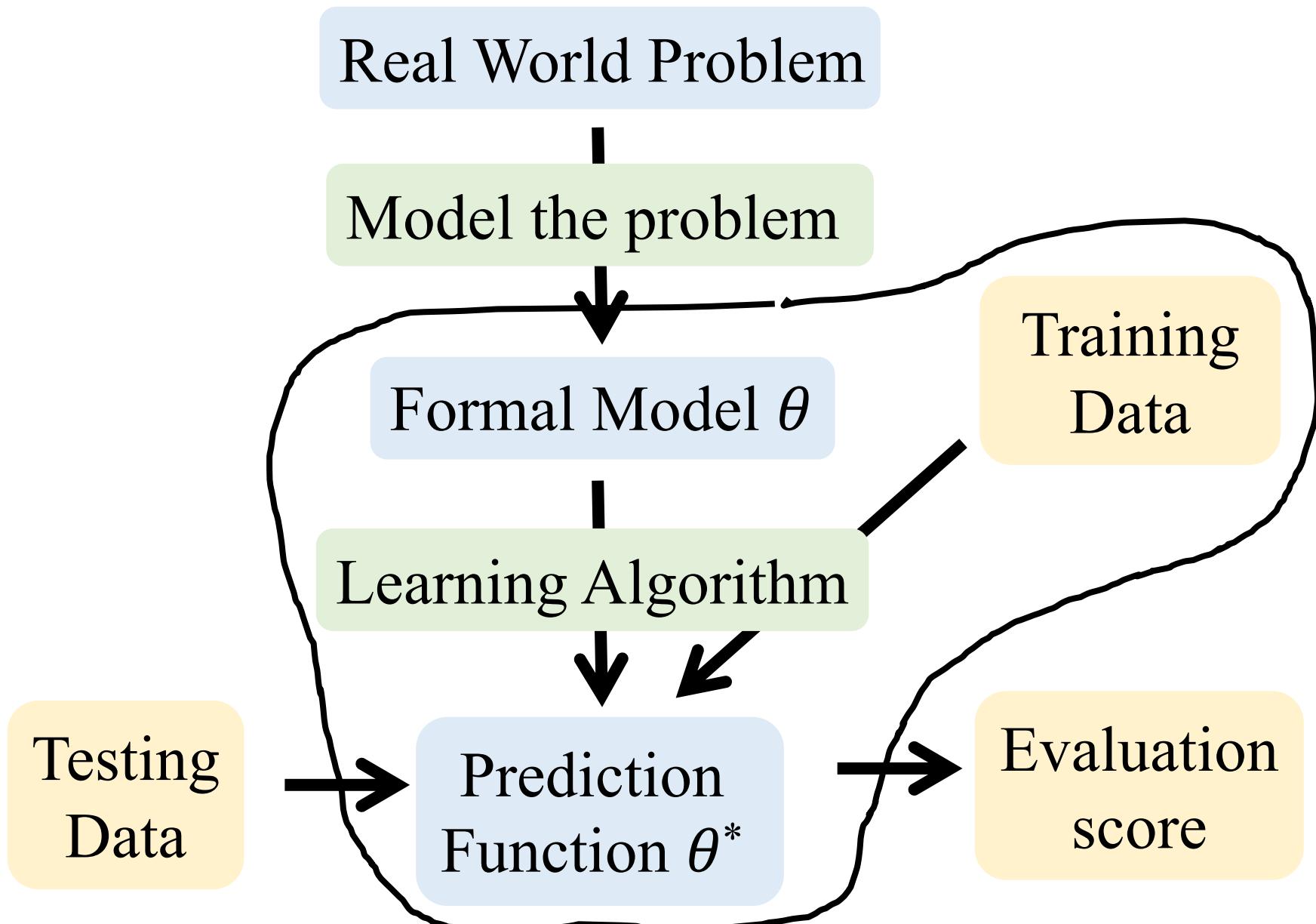
# Supervised Learning



# Modelling



# Training



# Testing

Real World Problem

Model the problem

Formal Model  $\theta$

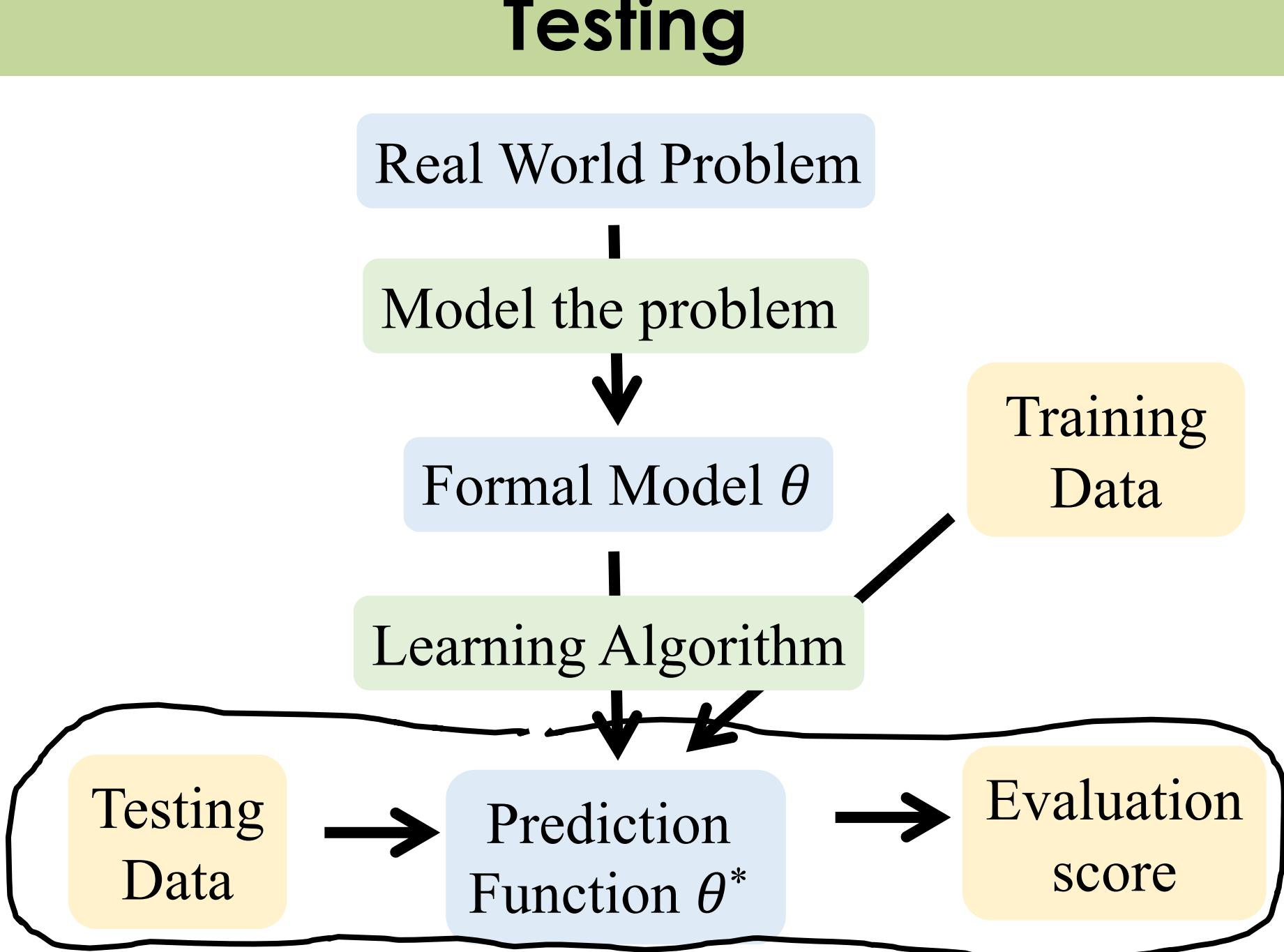
Training  
Data

Learning Algorithm

Testing  
Data

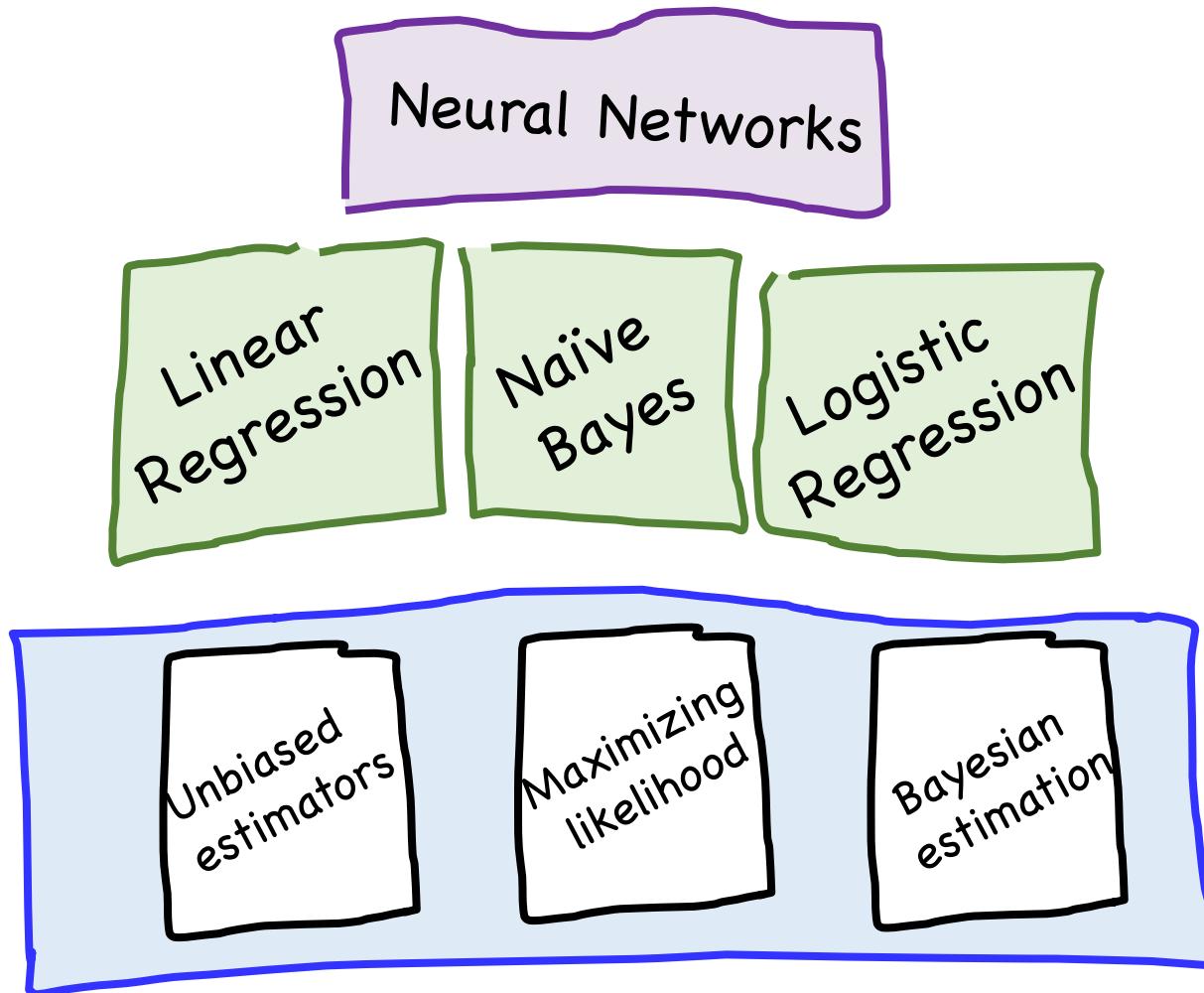
Prediction  
Function  $\theta^*$

Evaluation  
score

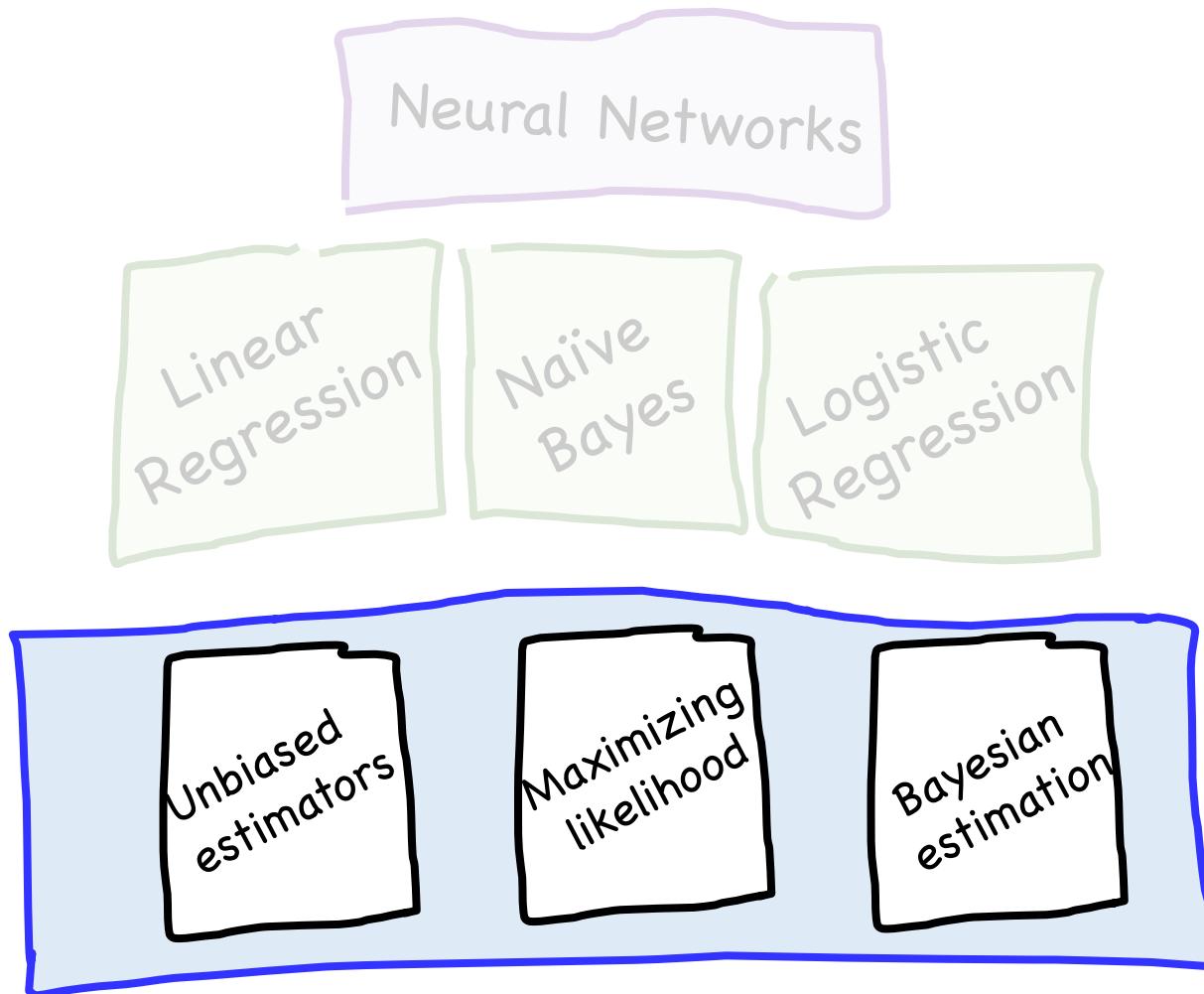


# Basis for learning from data

# Our Path



# Parameter Estimation



# Recall Sample Mean + Variance?

- Consider  $n$  I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i$  have distribution  $F$  with  $E[X_i] = \mu$  and  $\text{Var}(X_i) = \sigma^2$
  - We call sequence of  $X_i$  a sample from distribution  $F$
  - Recall sample mean:  $\bar{X} = \sum_{i=1}^n \frac{X_i}{n}$  where  $E[\bar{X}] = \mu$   
$$\bar{X} \sim N\left(\mu, \frac{\sigma^2}{n}\right) \text{ as } n \rightarrow \infty$$
  - Recall sample variance:

$$S^2 = \sum_{i=1}^n \frac{(X_i - \bar{X})^2}{n-1} = \text{undefined}$$

Estimate parameters for  
Bernoulli and Normal

Limited tool: how could we use that for fitting a “Mixture of Gaussians”?

# Great idea in Machine Learning

# Likelihood of Data

- Consider  $n$  I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i$  is a sample from density function  $f(X_i | \theta)$ 
    - Note: now explicitly specify parameter  $\theta$  of distribution



Likelihood question:  
How likely is the data given the samples?

$$\text{Likelihood}(\theta) = f(\text{Samples}|\theta)$$

[Demo](#)





# Likelihood of Data

- Consider  $n$  I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i$  is a sample from density function  $f(X_i | \theta)$ 
    - Note: now explicitly specify parameter  $\theta$  of distribution
  - We want to determine how “likely” the observed data  $(x_1, x_2, \dots, x_n)$  is based on density  $f(X_i | \theta)$
  - Define the Likelihood function,  $L(\theta)$ :

$$L(\theta) = \prod_{i=1}^n f(X_i | \theta)$$

- This is just a product since  $X_i$  are I.I.D.
- Intuitively: what is probability of observed data using density function  $f(X_i | \theta)$ , for some choice of  $\theta$

# Maximum Likelihood Estimator

- The Maximum Likelihood Estimator (MLE) of  $\theta$ , is the value of  $\theta$  that maximizes  $L(\theta)$ 
  - More formally:  $\theta_{MLE} = \arg \max_{\theta} L(\theta)$



Likelihood (of data given parameters):

$$L(\theta) = \prod_{i=1}^n f(X_i \mid \theta)$$

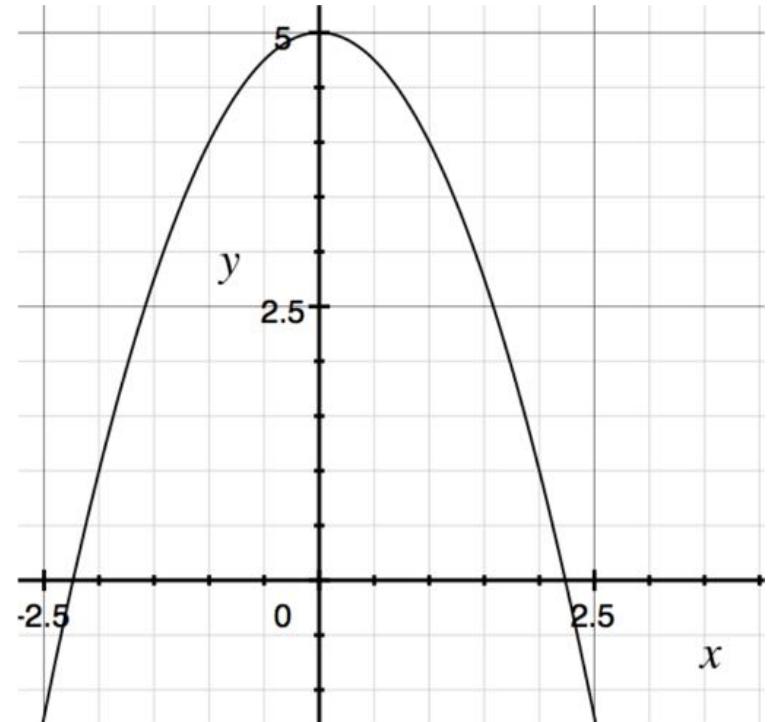


# Argmax

$$f(x) = -x^2 + 5$$

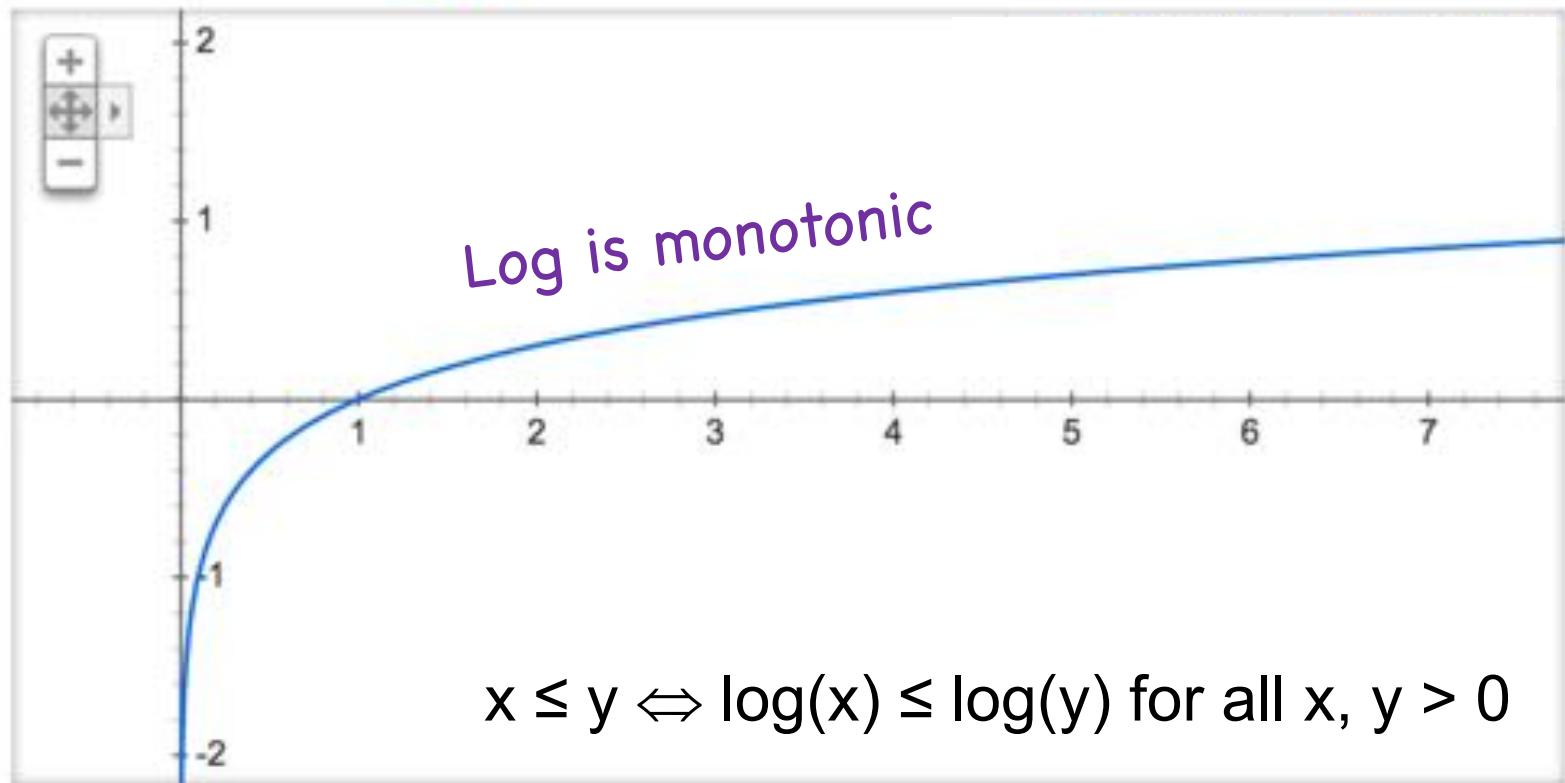
$$\max_x -x^2 + 5 = 5$$

$$\operatorname{argmax}_x -x^2 + 5 = 0$$



# Argmax of Log

Graph for  $\log(x)$



Claim:  $\operatorname{argmax}_x f(x) = \operatorname{argmax}_x \log f(x)$

# Argmax of Log

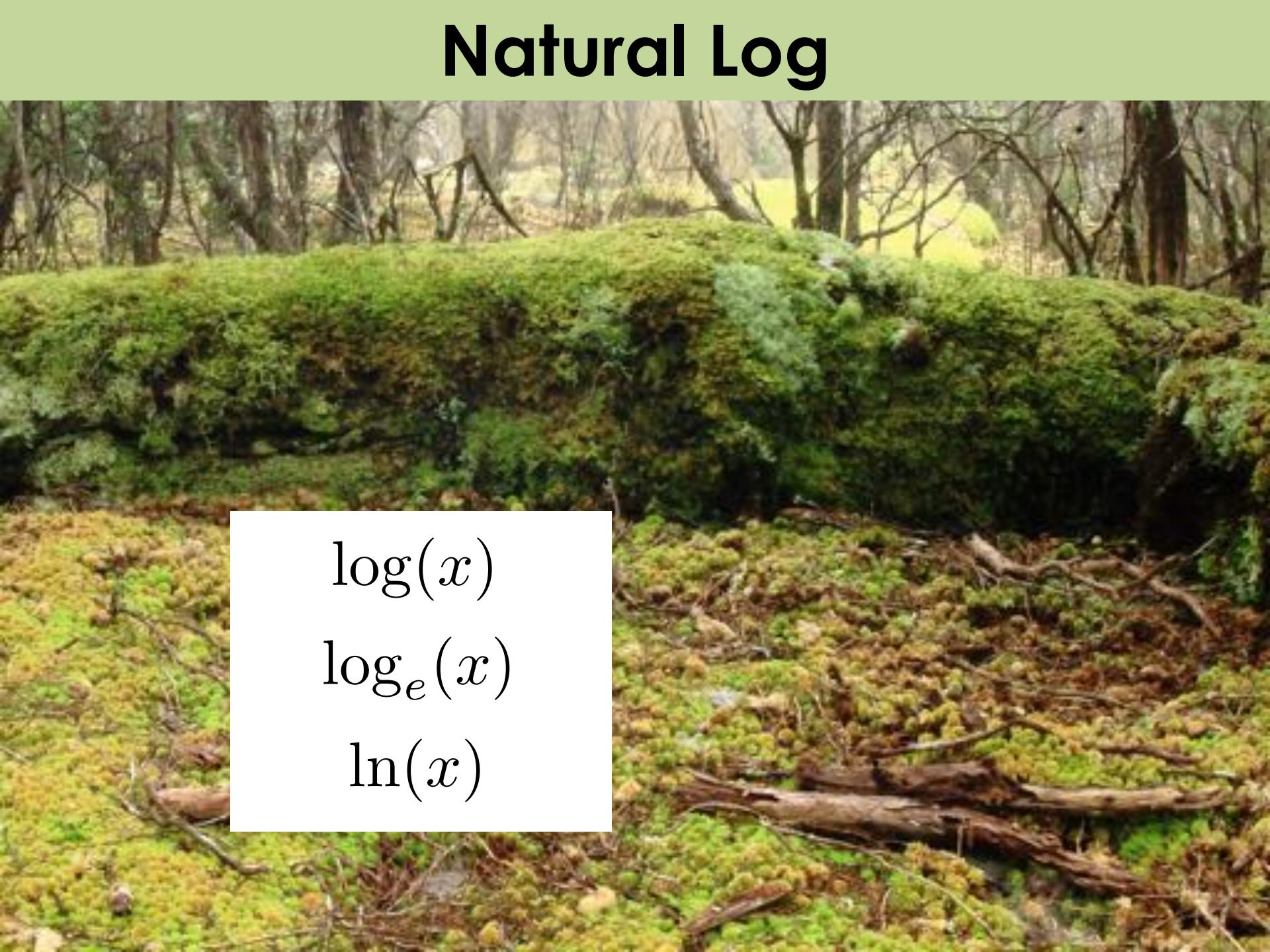


$$\operatorname{argmax}_x f(x) = \operatorname{argmax}_x \log f(x)$$

# Log I Love You

$$\log(ab) = \log(a) + \log(b)$$

# Natural Log



$\log(x)$

$\log_e(x)$

$\ln(x)$

# Maximum Likelihood Estimator

- The Maximum Likelihood Estimator (MLE) of  $\theta$ , is the value of  $\theta$  that maximizes  $L(\theta)$ 
  - More formally:  $\theta_{MLE} = \arg \max_{\theta} L(\theta)$
  - More convenient to use log-likelihood function,  $LL(\theta)$ :

$$LL(\theta) = \log L(\theta) = \log \prod_{i=1}^n f(X_i | \theta) = \sum_{i=1}^n \log f(X_i | \theta)$$

- $\theta$  that maximizes  $LL(\theta)$  also maximizes  $L(\theta)$ 
  - Formally:  $\arg \max_{\theta} LL(\theta) = \arg \max_{\theta} L(\theta)$
  - Similarly, for any positive constant  $c$  (not dependent on  $\theta$ ):

$$\arg \max_{\theta} (c \cdot LL(\theta)) = \arg \max_{\theta} LL(\theta) = \arg \max_{\theta} L(\theta)$$

Story so far: We can chose parameters by  
finding the argmax of the log likelihood of our  
data



# Maximum Likelihood

$$L(\theta) = \prod_{i=1}^n f(X_i | \theta)$$

$$LL(\theta) = \sum_{i=1}^n \log f(X_i | \theta)$$

$$\hat{\theta} = \operatorname{argmax}_{\theta} LL(\theta)$$





But how do we compute argmax?

# Option #1: Straight optimization

# Computing the MLE

- General approach for finding MLE of  $\theta$ 
  - Determine formula for  $LL(\theta)$
  - Differentiate  $LL(\theta)$  w.r.t. (each)  $\theta$ :  $\frac{\partial LL(\theta)}{\partial \theta}$
  - To maximize, set  $\frac{\partial LL(\theta)}{\partial \theta} = 0$
  - Solve resulting (simultaneous) equations to get  $\theta_{MLE}$ 
    - Make sure that derived  $\hat{\theta}_{MLE}$  is actually a maximum (and not a minimum or saddle point). E.g., check  $LL(\theta_{MLE} \pm \varepsilon) < LL(\theta_{MLE})$ 
      - This step often ignored in expository derivations
      - So, we'll ignore it here too (and won't require it in this class)

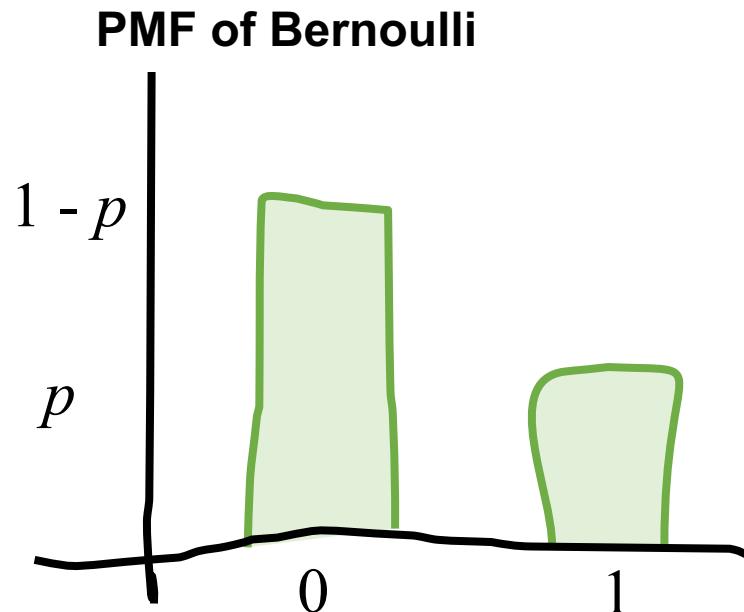
# Maximizing Likelihood with Bernoulli

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim \text{Ber}(p)$
  - Probability mass function,  $f(X_i | p)$ :

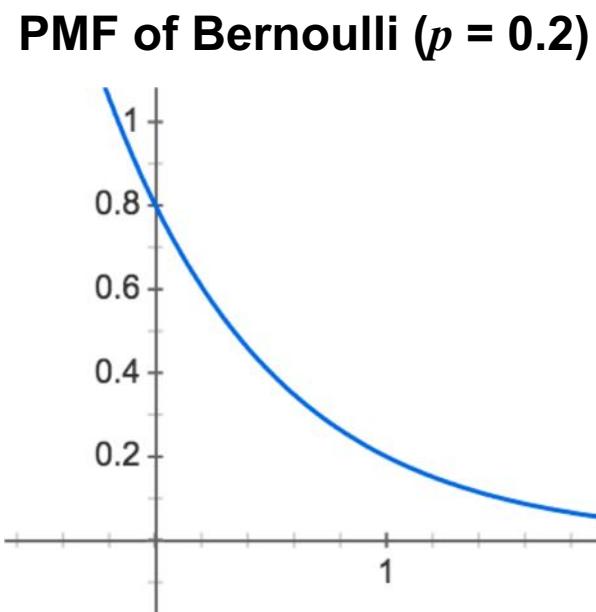


# Maximizing Likelihood with Bernoulli

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim \text{Ber}(p)$
  - Probability mass function,  $f(X_i | p)$ :



$$f(X_i | p) = p^{x_i} (1-p)^{1-x_i}$$



$$f(x) = 0.2^x (1 - 0.2)^{1-x}$$

# Bernoulli PMF

$$X \sim \text{Ber}(p)$$



$$f(X = x|p) = p^x(1 - p)^{1-x}$$

# Maximizing Likelihood with Bernoulli

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim \text{Ber}(p)$
  - Probability mass function,  $f(X_i | p)$ , can be written as:

$$f(X_i | p) = p^{x_i} (1-p)^{1-x_i} \quad \text{where } x_i = 0 \text{ or } 1$$

- Likelihood:  $L(\theta) = \prod_{i=1}^n p^{X_i} (1-p)^{1-X_i}$

- Log-likelihood:

$$\begin{aligned} LL(\theta) &= \sum_{i=1}^n \log(p^{X_i} (1-p)^{1-X_i}) = \sum_{i=1}^n [X_i (\log p) + (1-X_i) \log(1-p)] \\ &= Y(\log p) + (n-Y)\log(1-p) \quad \text{where } Y = \sum_{i=1}^n X_i \end{aligned}$$

- Differentiate w.r.t.  $p$ , and set to 0:

$$\frac{\partial LL(p)}{\partial p} = Y \frac{1}{p} + (n-Y) \frac{-1}{1-p} = 0 \quad \Rightarrow \quad p_{MLE} = \frac{Y}{n} = \frac{1}{n} \sum_{i=1}^n X_i$$

Isn't that the same as  
unbiased estimator?

Yes. For Bernoulli.



# Maximum Likelihood Algorithm

1. Decide on a model for the distribution of your samples. Define the PMF / PDF for your sample.

2. Write out the log likelihood function.

3. State that the optimal parameters are the argmax of the log likelihood function.

4. Use an optimization algorithm to calculate argmax



# Maximizing Likelihood with Poisson

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim \text{Poi}(\lambda)$
  - PMF:  $f(X_i | \lambda) = \frac{e^{-\lambda} \lambda^{x_i}}{x_i!}$       Likelihood:  $L(\theta) = \prod_{i=1}^n \frac{e^{-\lambda} \lambda^{X_i}}{X_i!}$
  - Log-likelihood:
$$\begin{aligned} LL(\theta) &= \sum_{i=1}^n \log\left(\frac{e^{-\lambda} \lambda^{X_i}}{X_i!}\right) = \sum_{i=1}^n [-\lambda \log(e) + X_i \log(\lambda) - \log(X_i!)] \\ &= -n\lambda + \log(\lambda) \sum_{i=1}^n X_i - \sum_{i=1}^n \log(X_i!) \end{aligned}$$
  - Differentiate w.r.t.  $\lambda$ , and set to 0:

$$\frac{\partial LL(\lambda)}{\partial \lambda} = -n + \frac{1}{\lambda} \sum_{i=1}^n X_i = 0 \Rightarrow \lambda_{MLE} = \frac{1}{n} \sum_{i=1}^n X_i$$

Its so general!

# Maximizing Likelihood with Normal

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim N(\mu, \sigma^2)$
  - PDF:  $f(X_i | \mu, \sigma^2) = \frac{1}{\sqrt{2\pi}\sigma} e^{-(X_i - \mu)^2 / (2\sigma^2)}$
  - Log-likelihood:

$$LL(\theta) = \sum_{i=1}^n \log\left(\frac{1}{\sqrt{2\pi}\sigma} e^{-(X_i - \mu)^2 / (2\sigma^2)}\right) = \sum_{i=1}^n \left[ -\log(\sqrt{2\pi}\sigma) - (X_i - \mu)^2 / (2\sigma^2) \right]$$

- First, differentiate w.r.t.  $\mu$ , and set to 0:

$$\frac{\partial LL(\mu, \sigma^2)}{\partial \mu} = \sum_{i=1}^n 2(X_i - \mu) / (2\sigma^2) = \frac{1}{\sigma^2} \sum_{i=1}^n (X_i - \mu) = 0$$

- Then, differentiate w.r.t.  $\sigma$ , and set to 0:

$$\frac{\partial LL(\mu, \sigma^2)}{\partial \sigma} = \sum_{i=1}^n -\frac{1}{\sigma} + 2(X_i - \mu)^2 / (2\sigma^3) = -\frac{n}{\sigma} + \sum_{i=1}^n (X_i - \mu)^2 / (\sigma^3) = 0$$

# Being Normal, Simultaneously

- Now have two equations, two unknowns:

$$\frac{1}{\sigma^2} \sum_{i=1}^n (X_i - \mu) = 0 \quad -\frac{n}{\sigma} + \sum_{i=1}^n (X_i - \mu)^2 / (\sigma^3) = 0$$

- First, solve for  $\mu_{MLE}$ :

$$\frac{1}{\sigma^2} \sum_{i=1}^n (X_i - \mu) = 0 \Rightarrow \sum_{i=1}^n X_i = n\mu \Rightarrow \mu_{MLE} = \frac{1}{n} \sum_{i=1}^n X_i$$

- Then, solve for  $\sigma^2_{MLE}$ :

$$-\frac{n}{\sigma} + \sum_{i=1}^n (X_i - \mu)^2 / (\sigma^3) = 0 \Rightarrow n\sigma^2 = \sum_{i=1}^n (X_i - \mu)^2$$

$$\sigma_{MLE}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \mu_{MLE})^2$$

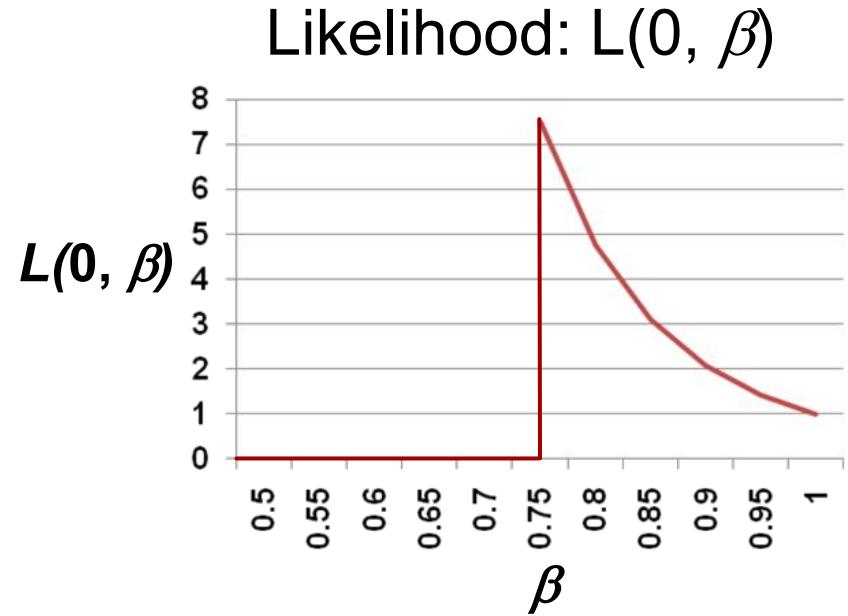
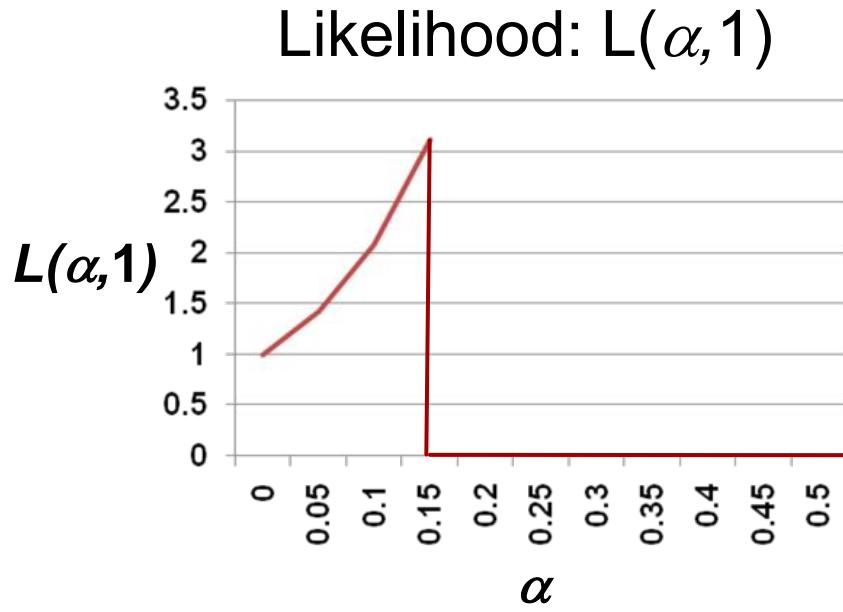
- Note:  $\mu_{MLE}$  unbiased, but  $\sigma^2_{MLE}$  biased

# Maximizing Likelihood with Uniform

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim \text{Uni}(\alpha, \beta)$
  - PDF:  $f(X_i | \alpha, \beta) = \begin{cases} \frac{1}{\beta - \alpha} & \alpha \leq x_i \leq \beta \\ 0 & \text{otherwise} \end{cases}$
  - Likelihood:  $L(\theta) = \begin{cases} \left(\frac{1}{\beta - \alpha}\right)^n & \alpha \leq x_1, x_2, \dots, x_n \leq \beta \\ 0 & \text{otherwise} \end{cases}$ 
    - Constraint  $\alpha \leq x_1, x_2, \dots, x_n \leq \beta$  makes differentiation tricky
    - Intuition: want interval size  $(\beta - \alpha)$  to be as small as possible to maximize likelihood function for each data point
    - But need to make sure all observed data contained in interval
      - If all observed data not in interval, then  $L(\theta) = 0$
  - Solution:  $\alpha_{MLE} = \min(x_1, \dots, x_n) \quad \beta_{MLE} = \max(x_1, \dots, x_n)$

# Understanding MLE with Uniform

- Consider I.I.D. random variables  $X_1, X_2, \dots, X_n$ 
  - $X_i \sim \text{Uni}(0, 1)$
  - Observe data:
    - 0.15, 0.20, 0.30, 0.40, 0.65, 0.70, 0.75



# Small Samples = Problems

- How do small samples affect MLE?
  - In many cases,  $\mu_{MLE} = \frac{1}{n} \sum_{i=1}^n X_i$  = sample mean
    - Unbiased. Not too shabby...
  - As seen with Normal,  $\sigma_{MLE}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \mu_{MLE})^2$ 
    - Biased. Underestimates for small  $n$  (e.g., 0 for  $n = 1$ )
  - As seen with Uniform,  $\alpha_{MLE} \geq \alpha$  and  $\beta_{MLE} \leq \beta$ 
    - Biased. Problematic for small  $n$  (e.g.,  $\alpha = \beta$  when  $n = 1$ )
  - Small sample phenomena intuitively make sense:
    - Maximum likelihood  $\Rightarrow$  best explain data we've seen
    - Does not attempt to generalize to unseen data

# Properties of MLE

- Maximum Likelihood Estimators are generally:
  - **Consistent:**  $\lim_{n \rightarrow \infty} P(|\hat{\theta} - \theta| < \varepsilon) = 1$  for  $\varepsilon > 0$
  - Potentially biased (though asymptotically less so)
  - **Asymptotically optimal**
    - Has smallest variance of “good” estimators for large samples
  - **Often used in practice** where sample size is large relative to parameter space
    - But be careful, there are some very large parameter spaces

