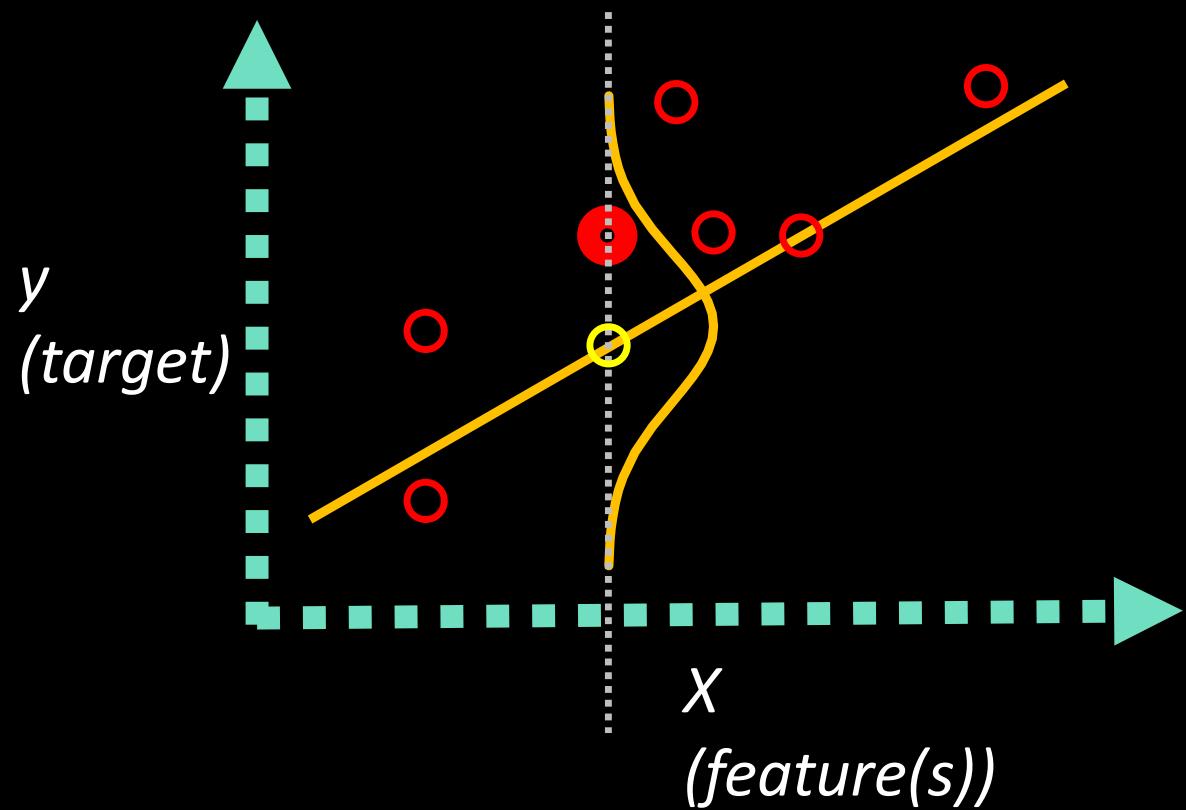


Recap: Linear Regression

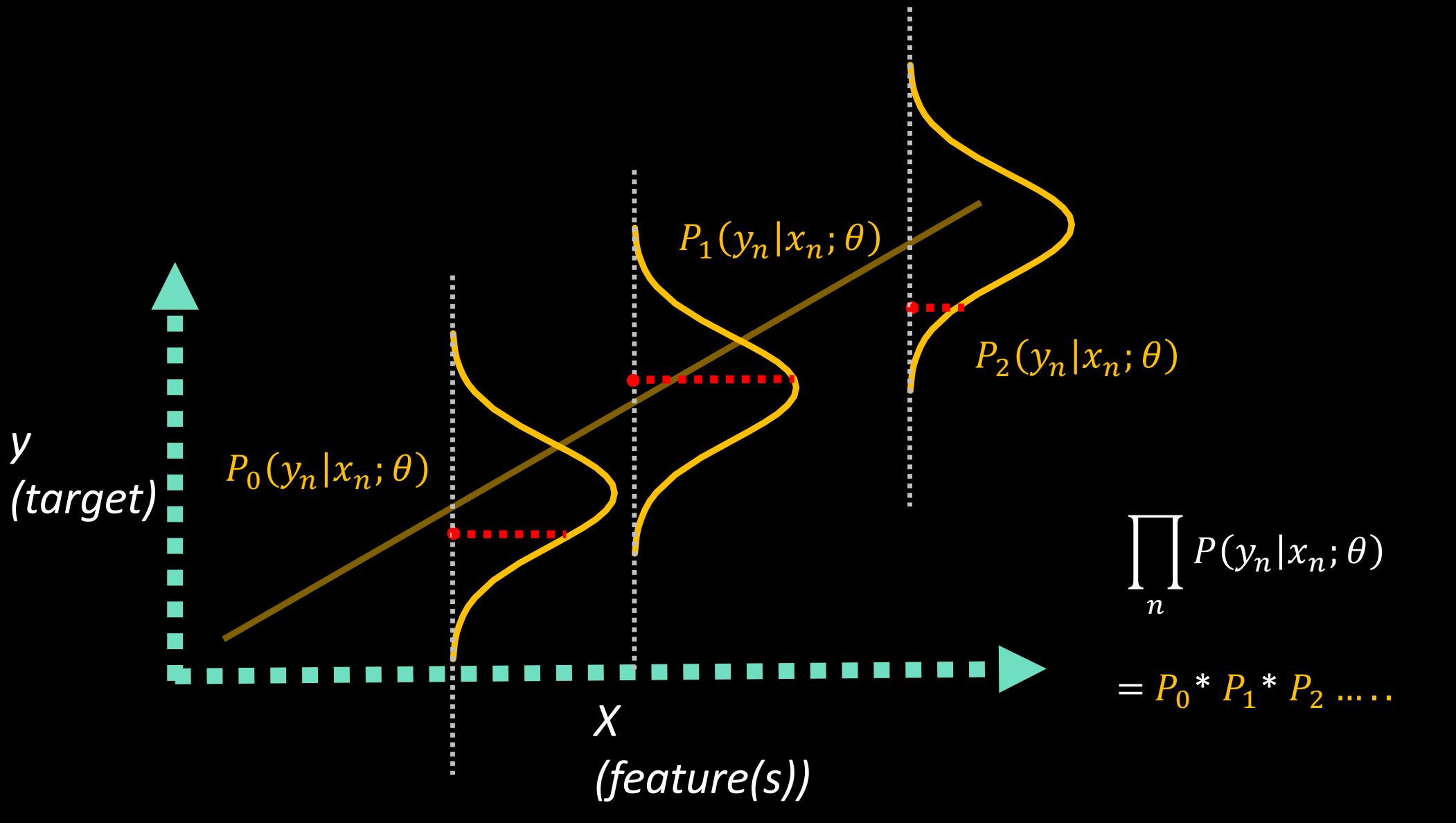
Linear Regression: Predicts continuous outcomes



Linear regression is a method of predicting an outcome based on input data.

A line represents the best-fit linear relationship between weight and size.

This line can be used to make predictions: given a new weight value, we can estimate the corresponding size.



Maximum Likelihood Estimation (MLE)

$$\theta_{MLE} = \operatorname{argmax}_{\theta} P(y|X; \theta)$$

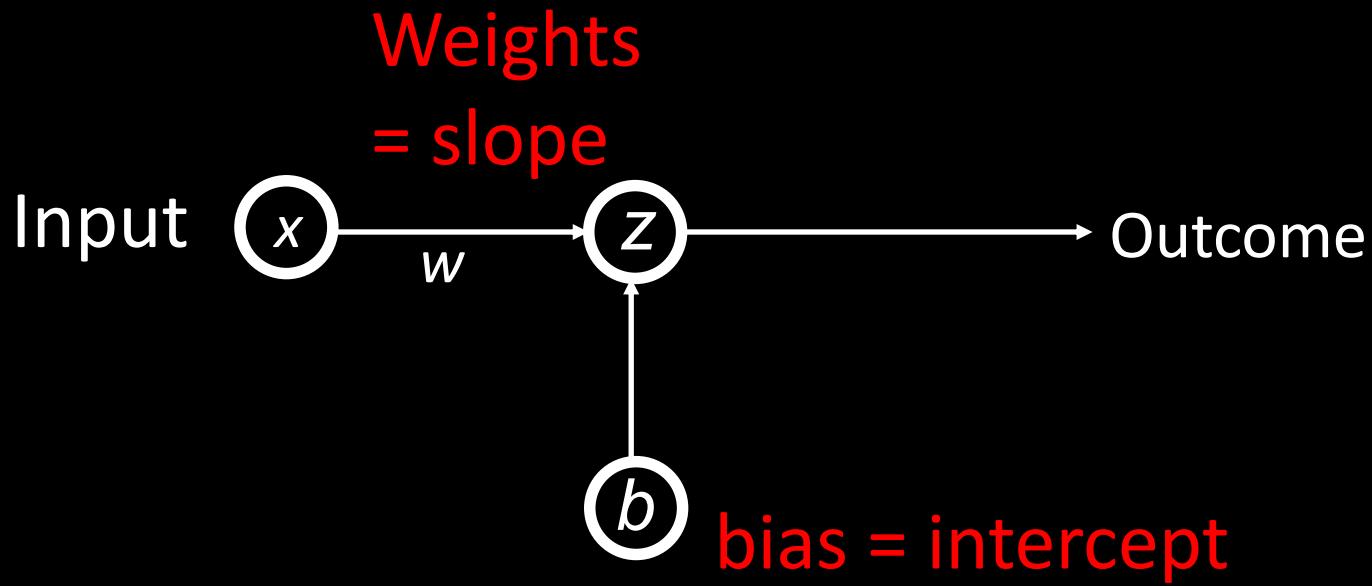
$P(y|X)$ (true) probability of seeing stroke given blood pressure

$P(y|X; \theta)$ Probability of seeing stroke given blood pressure from model with θ

$\operatorname{argmax}_{\theta}$ Find the parameters θ to maximize the given probability $P(y|X, \theta)$

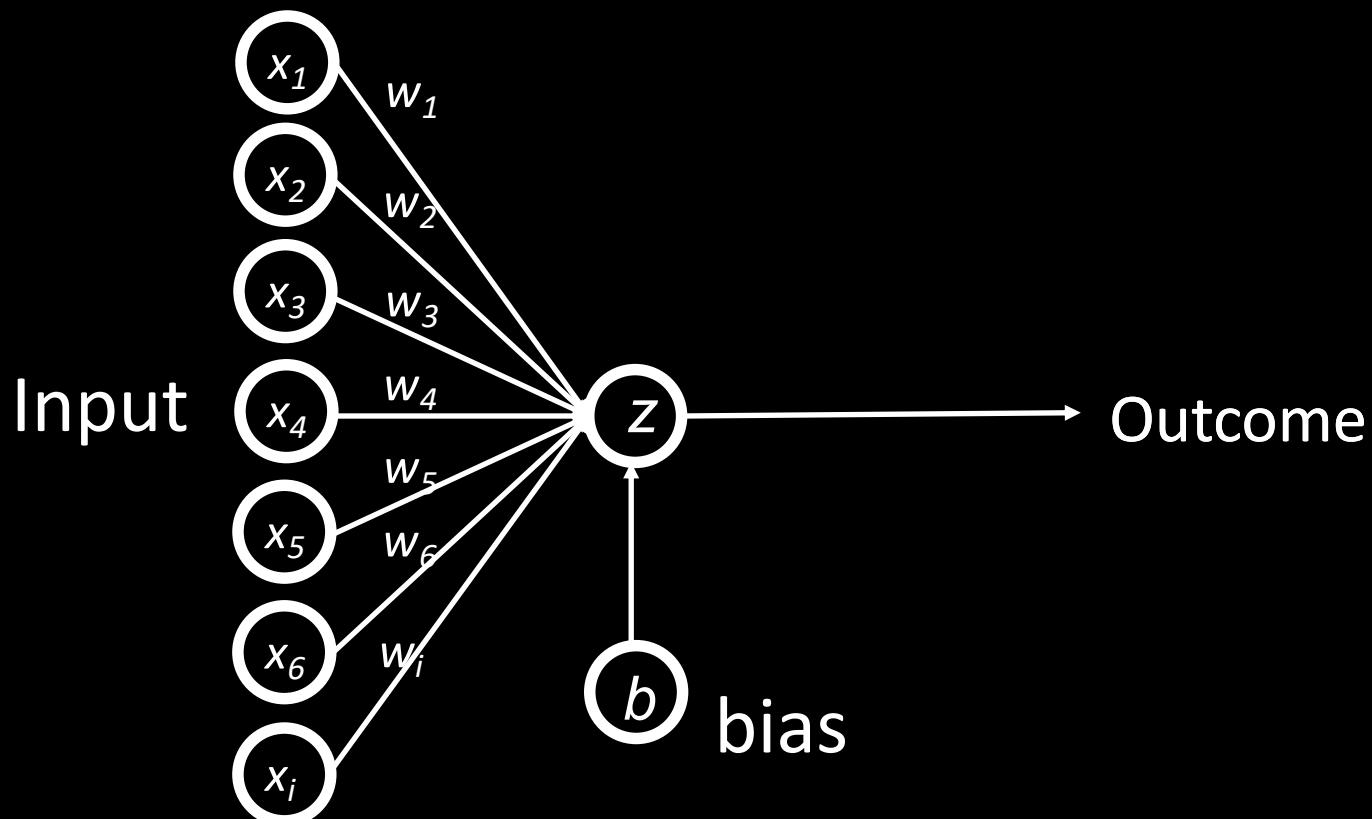
Graph of Linear Regression

$$y = WX + b$$

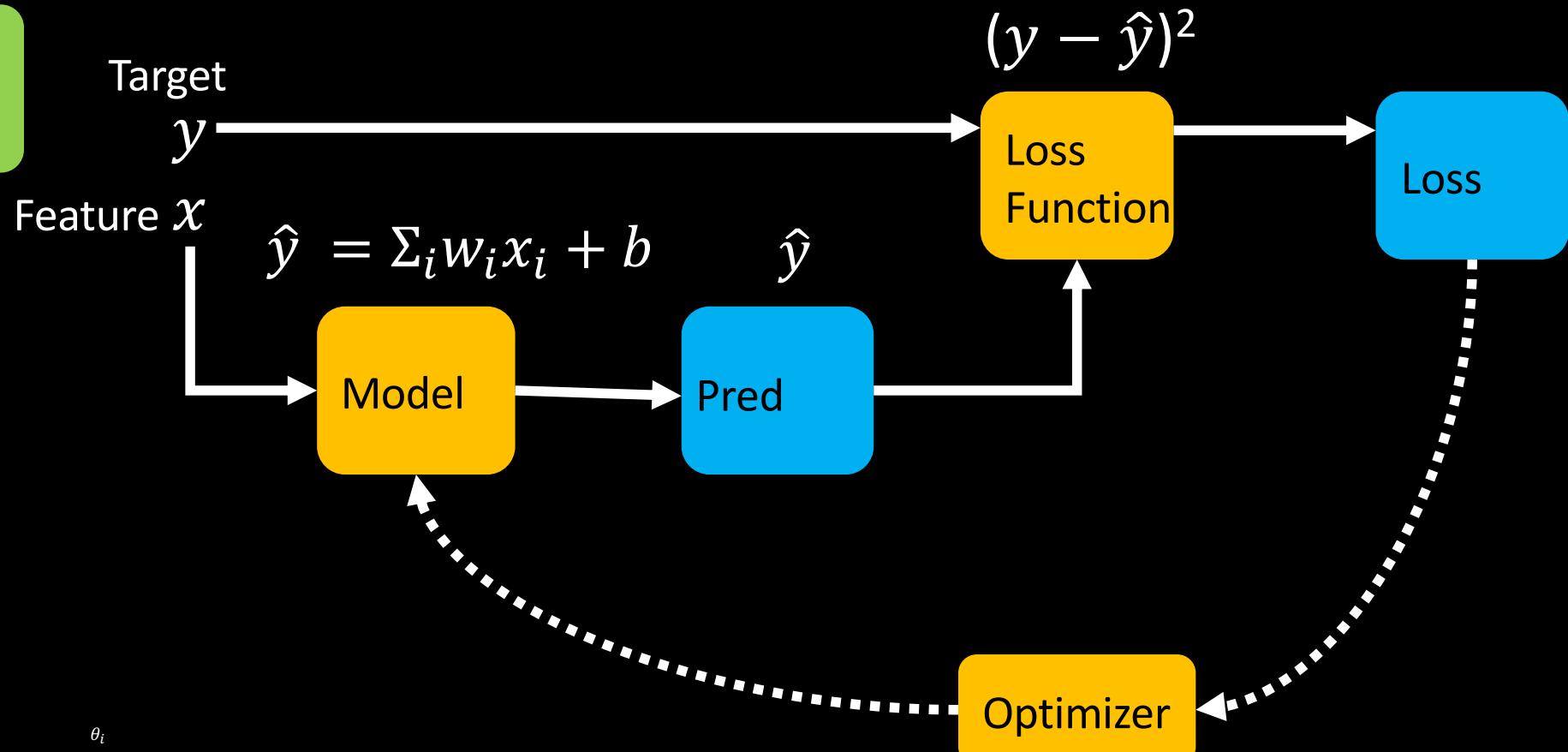


Graph of Linear Regression

$$y = w_1x_1 + w_2x_2 + \dots + b$$



Flow Chart for Linear Regression

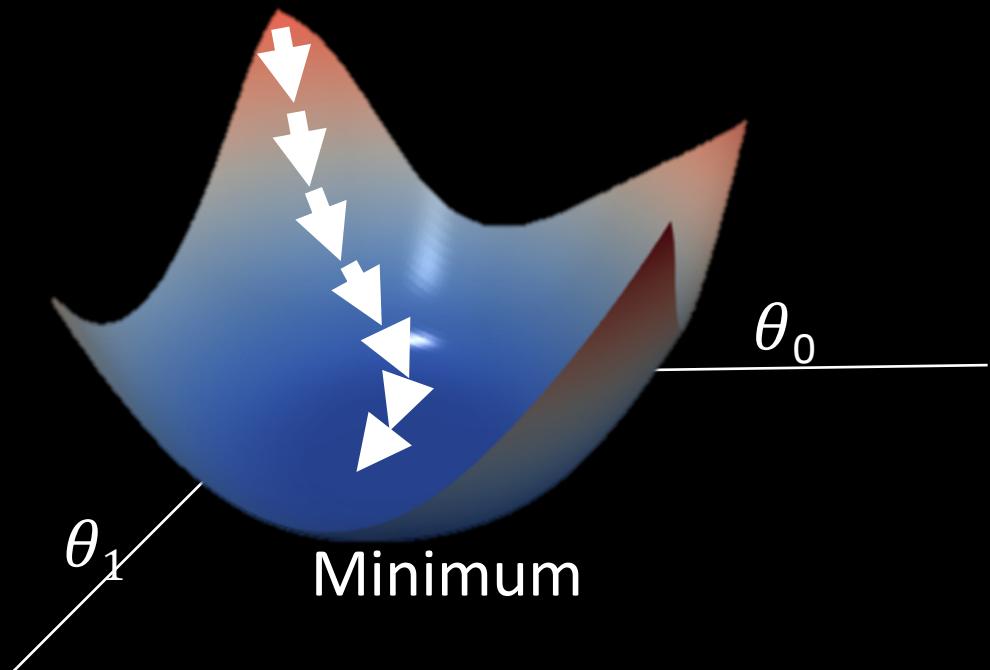


Gradient Descent

Moves in the direction of steepest descent (negative gradient) to minimize the objective function.

Steps:

1. Start with initial parameter values
2. Calculate the gradient of the objective function
3. Update parameters in the opposite direction of the gradient
4. Repeat steps 2-3 until convergence



Classification: Logistic Regression

Classification vs Regression

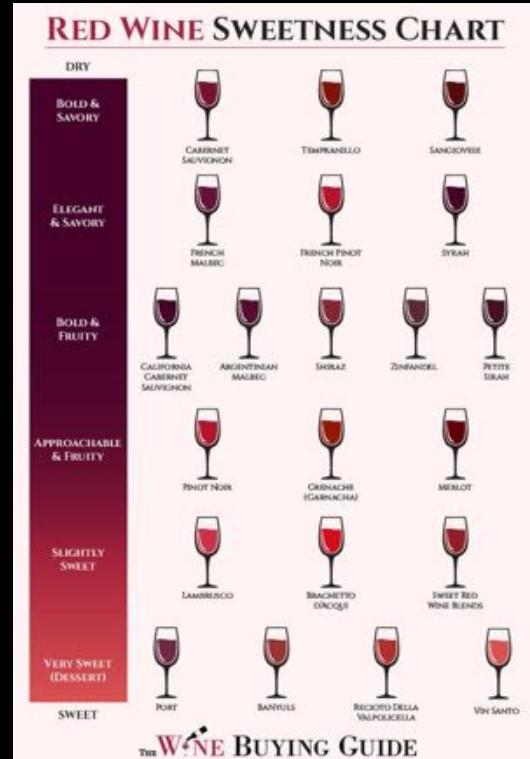
TNM 8 th - Primary tumor characteristics	
T_x	Tumor in sputum/bronchial washings but not be assessed in imaging or bronchoscopy
T₀	No evidence of tumor
T_{is}	Carcinoma in situ
T₁	≤ 3 cm surrounded by lung/visceral pleura, not involving main bronchus
T_{1a(mi)}	Minimally invasive carcinoma
T_{1a}	≤ 1 cm
T_{1b}	> 1 to ≤ 2 cm
T_{1c}	> 2 to ≤ 3 cm
T₂	> 3 to ≤ 5 cm or involvement of main bronchus without carina, regardless of distance from carina or invasion visceral pleural or atelectasis or post obstructive pneumonitis extending to hilum
T_{2a}	>3 to ≤4cm
T_{2b}	>4 to ≤5cm
T₃	>5 to ≤7cm in greatest dimension or tumor of any size that involves chest wall, pericardium, phrenic nerve or satellite nodules in the same lobe
T₄	> 7cm in greatest dimension or any tumor with invasion of mediastinum, diaphragm, heart, great vessels, recurrent laryngeal nerve, carina, trachea, oesophagus, spine or separate tumor in different lobe of ipsilateral lung
N₁	Ipsilateral peribronchial and/or hilar nodes and intrapulmonary nodes
2	Ipsilateral mediastinal and/or subcarinal nodes
3	Contralateral mediastinal or hilar; ipsilateral/contralateral scalene/ supraclavicular
M₁	Distant metastasis
M_{1a}	Tumor in contralateral lung or pleural/pericardial nodule/malignant effusion
M_{1b}	Single extrathoracic metastasis, including single non-regional lymphnode
M_{1c}	Multiple extrathoracic metastases in one or more organs

Human has a tendency (or need) to categorize things for our own perception

Table 2. Examples of the training data set (values are normalized between 0 and 1)

Fruit image	Input of the network		Output
	Mean	Variance	Fruit Grading
	0.43	0.11	'Extra'
	0.22	0.05	'Type 1'
	0.27	0.23	'Type 2'
	0.26	0.31	'Rejected'
	0.28	0.29	'Rejected'

St. Cerc. St. CICBIA 2016 17 (1)

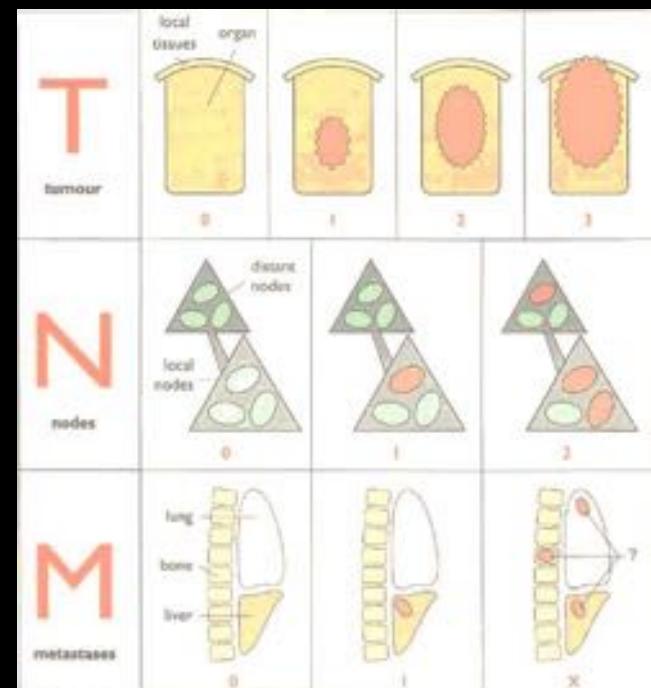


82

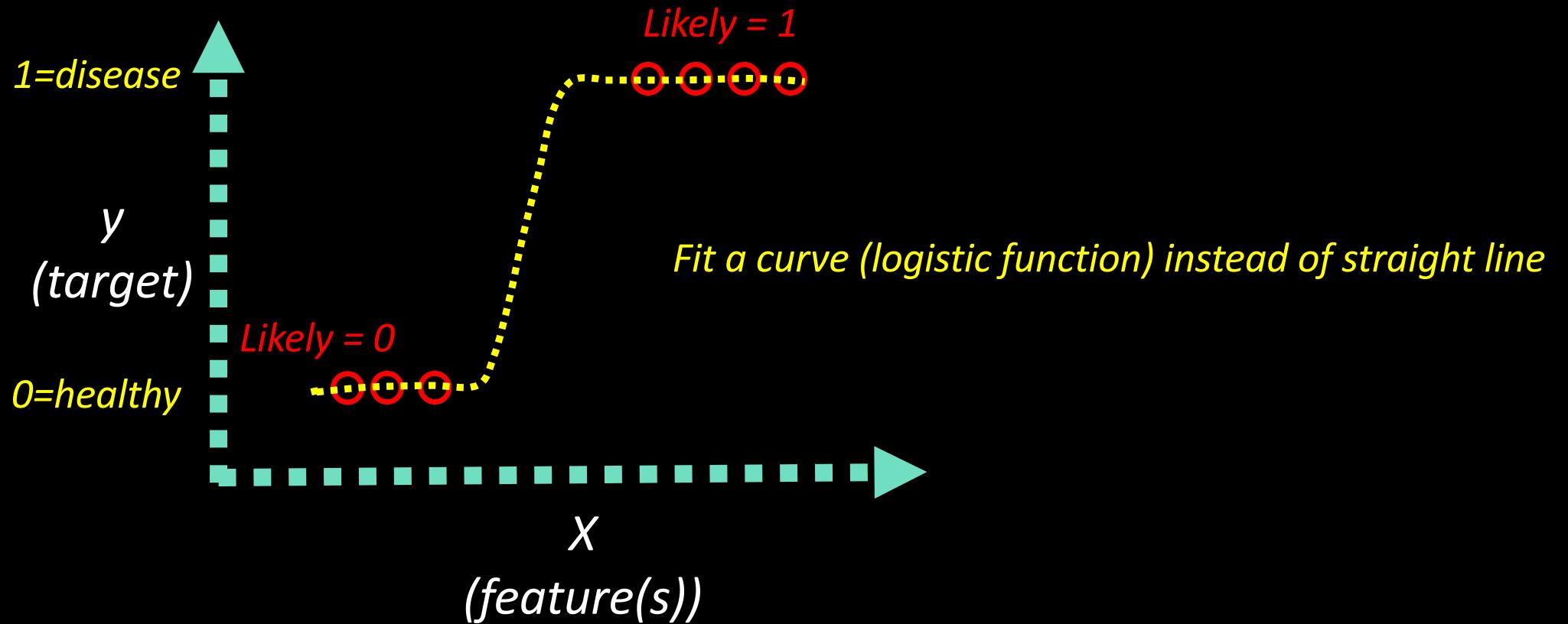
2023 Federal Tax Brackets			
TAX BRACKET/RATE	SINGLE	MARRIED FILING JOINTLY	HEAD OF HOUSEHOLD
10%	\$0 - \$11,000	\$0 - \$22,000	\$0 - \$15,700
12%	\$11,001 - \$44,725	\$22,001 - \$89,450	\$15,701 - \$59,850
22%	\$44,726 - \$95,375	\$89,451 - \$190,750	\$59,851 - \$95,350
24%	\$95,376 - \$182,100	\$190,751 - \$364,200	\$95,351 - \$182,100
32%	\$182,101 - \$231,250	\$364,201 - \$462,500	\$182,101 - \$231,250
35%	\$231,251 - \$578,125	\$462,501 - \$693,750	\$231,251 - \$578,100
37%	\$578,126+	\$693,751+	\$578,101+

© THE COLLEGE INVESTOR

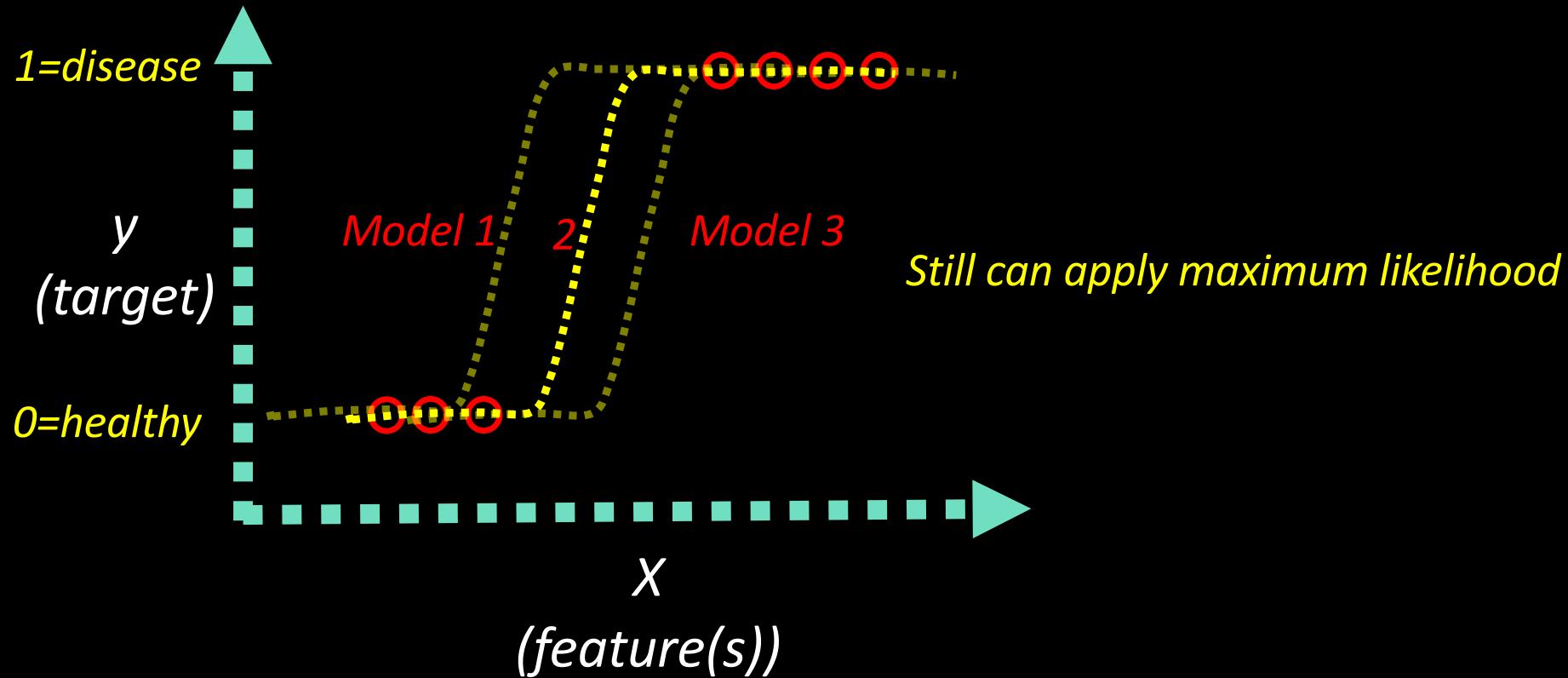
Source: TheCollegeInvestor.com



Logistic Regression: Predicts categorical outcomes



Logistic Regression: Predicts categorical outcomes



Regression -> classification, how?



regression $\sum_i w_i x_i + b$ $-\infty \sim \infty$

y_i $-\infty \sim \infty$

classification $0 < P(Y = truck) < 1$ $P(Y = car) = 1 - P(Y = truck)$

We want to find a way to map $-\infty \sim \infty$ to 0-1

Chance of wolf (=1)



0 (very sure)

0.2 (not sure)

0.7 (not sure)

1 (very sure)

y
(Output Probability)

$P=0.5$



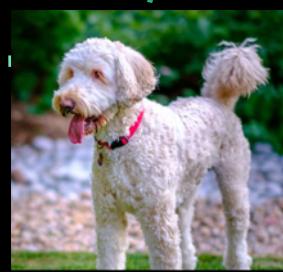
$P=0.7$



$P=1.0$



$P=0$



$P=0.2$

X
feature(s)

y
(Output Probability)

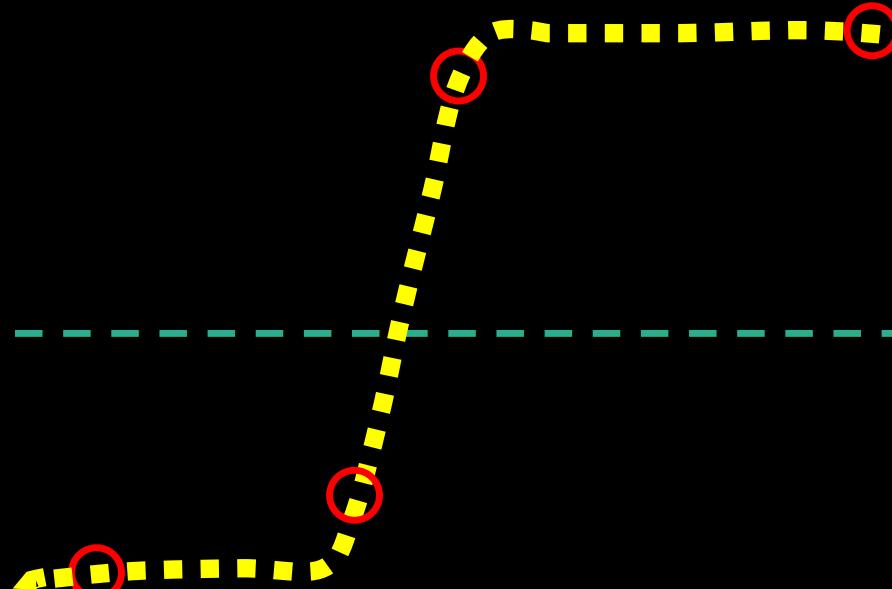
$$P=0.5$$

X
feature(s)

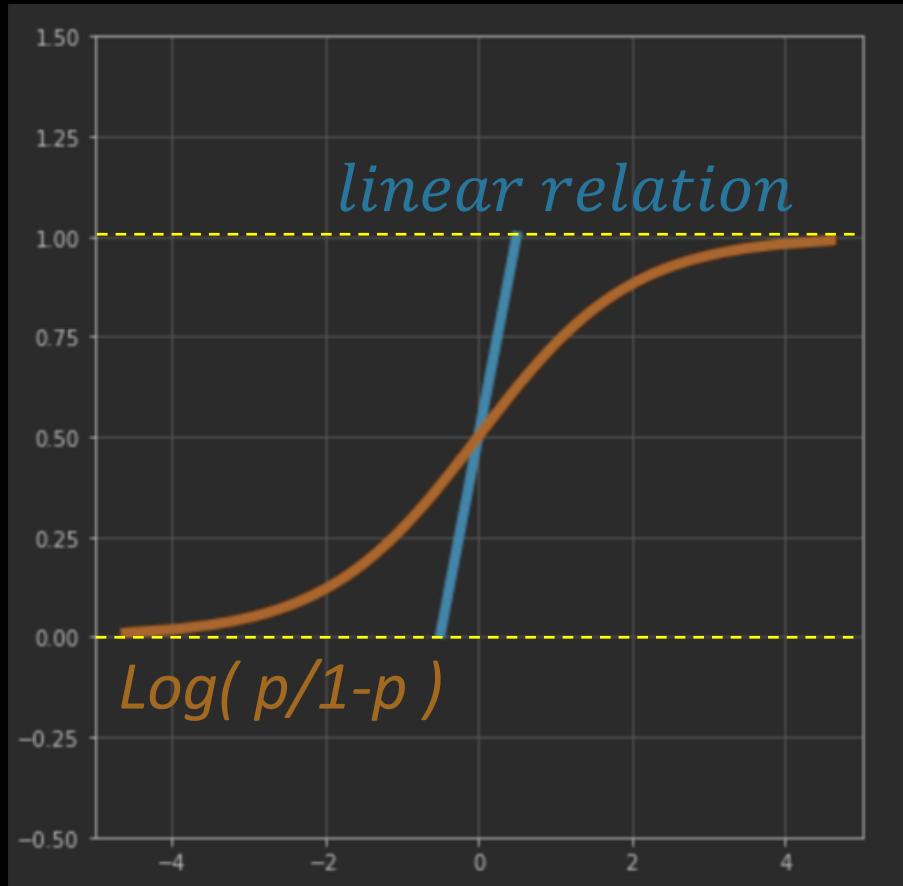


$P \geq 0.5$ Class = 1

$P < 0.5$ Class = 0



Probability
0-1



feature – $-\infty \sim \infty$

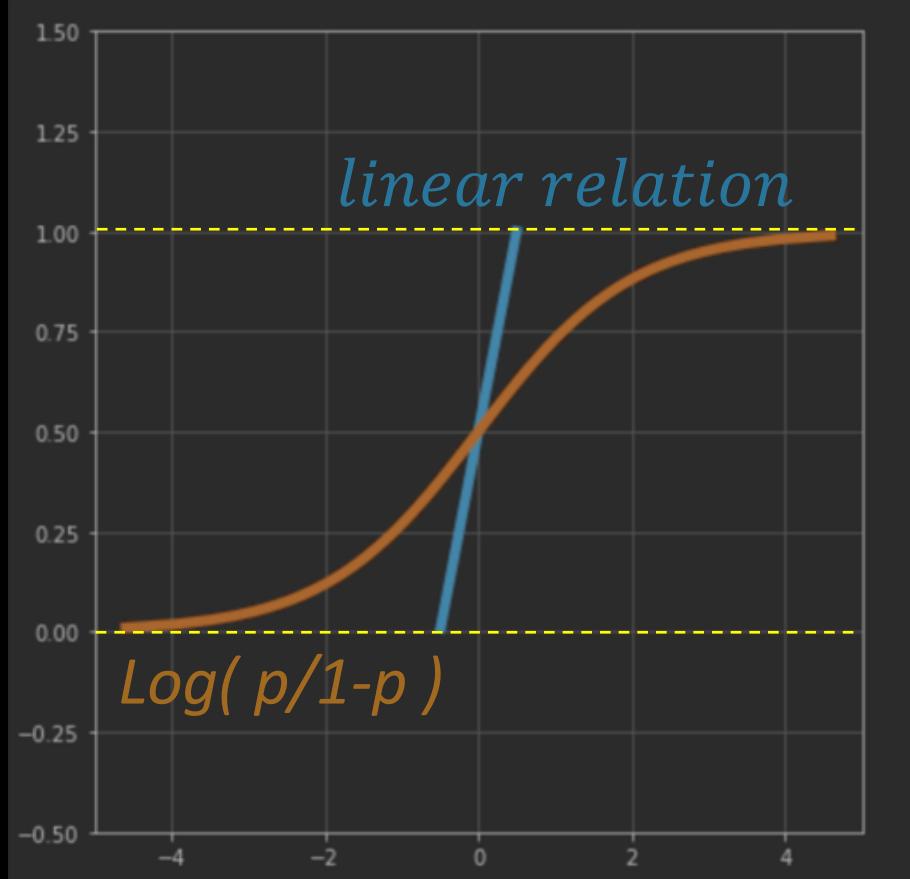
How to predict probability?

Remember we got
output of range

$-\infty \sim \infty$

From regression

Probability
0-1



feature – $\infty\sim\infty$

Probability p $0\sim 1$

Logit $\log\left(\frac{p}{1-p}\right)$ $-\infty\sim\infty$

same range!

linear regression $\sum_i w_i x_i + b$ $-\infty\sim\infty$

(Odd Ratio) $\frac{p}{1-p}$ $0\sim\infty$

	No. of case patients	No. of control subjects	Odds ratio	95% confidence interval	P for trend*
Nonsmokers	117	1750	1.0	Referent	
Duration of tobacco use, y					
0.1–20.0	3	33	1.3	0.4–4.5	
20.1–32.0	7	33	3.4	1.4–8.0	
32.1–44.0	21	33	13.3	7.2–24.9	
≥44.1	30	30	19.1	10.4–35.1	<.0001
Average consumption of tobacco, g/day					
0.1–3.5	2	10	2.2	0.5–10.4	
3.6–5.0	22	54	7.9	4.3–14.3	
5.1–10.7	6	18	4.8	1.9–12.6	
≥10.8	31	47	12.4	7.2–21.4	.1
Cumulative consumption of tobacco, g/day × y					
1–71	3	31	1.3	0.4–4.4	
72–157	13	34	7.6	3.8–15.4	
158–382	15	32	8.0	4.0–15.7	
≥383	30	32	18.3	10.2–32.8	.0006
Age at start of tobacco use, y					
≤19	27	48	9.6	5.6–16.7	
20–26	20	52	6.3	3.5–11.2	
≥27	14	29	8.2	4.1–16.3	.4

*Test for linear trend, two-sided *P* value (considered statistically significant for *P*<.05).

Now we have a good target for regression:

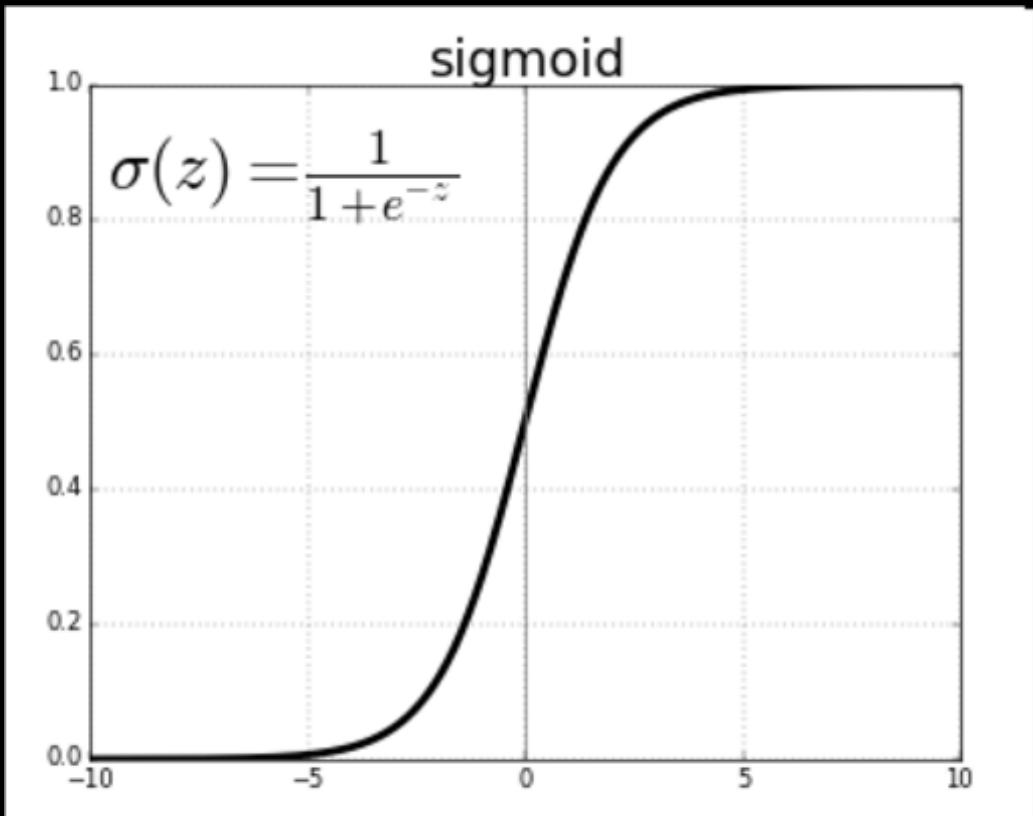
$$\sum_i w_i x_i + b \quad -\infty \sim \infty \qquad \text{to} \qquad \textit{Logit} \quad \log\left(\frac{p}{1-p}\right) \quad -\infty \sim \infty$$

$$p = \frac{1}{1 + \exp(-(\sum_i w_i x_i + b))} = \text{Sigmoid}(w_i x_i + b) = S(\beta X)$$

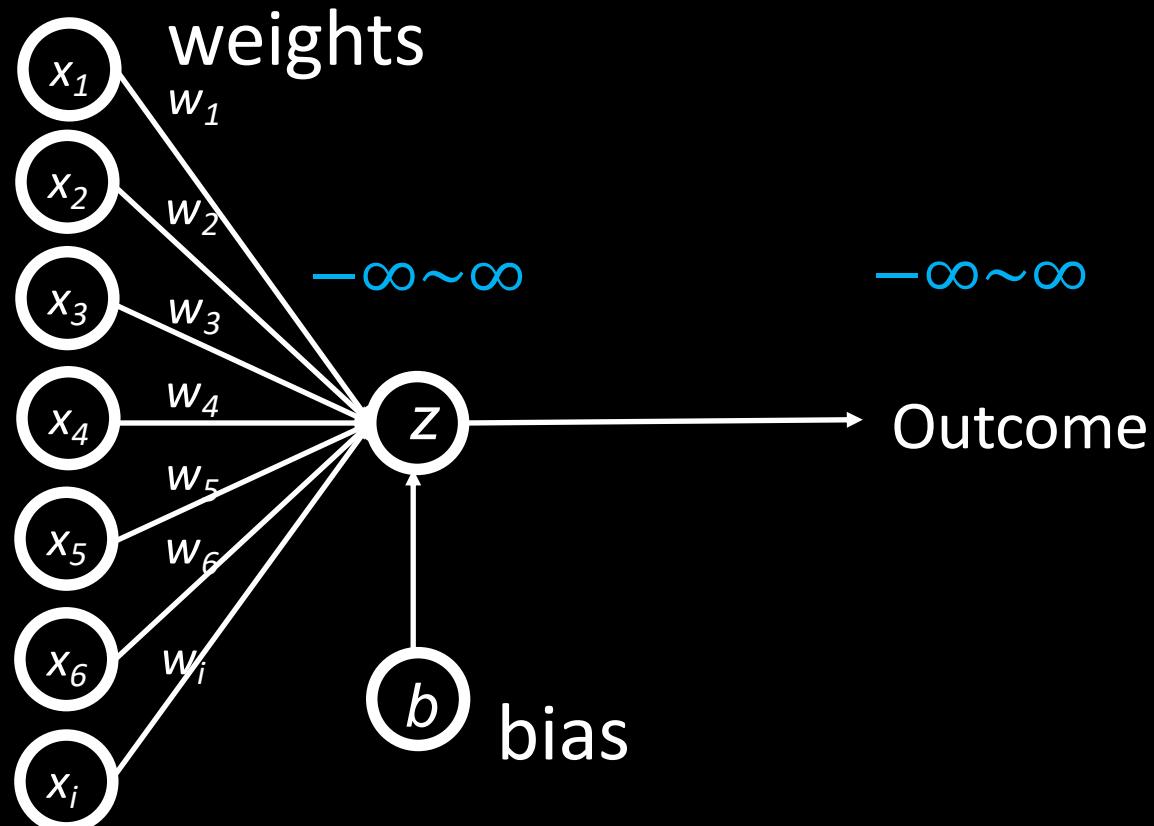
$$P = S(\beta X)$$

Logistic Regression = Regression Between Probability and Sigmoid of (weighted features)

```
def sigmoid(x):  
    return 1 / (1 + np.exp(-x))  
  
# Create x values from -6 to 6  
x = np.linspace(-6, 6, 1000)  
  
# Calculate sigmoid values  
y_sigmoid = sigmoid(x)  
# Plot sigmoid  
plt.plot(x, y_sigmoid)
```

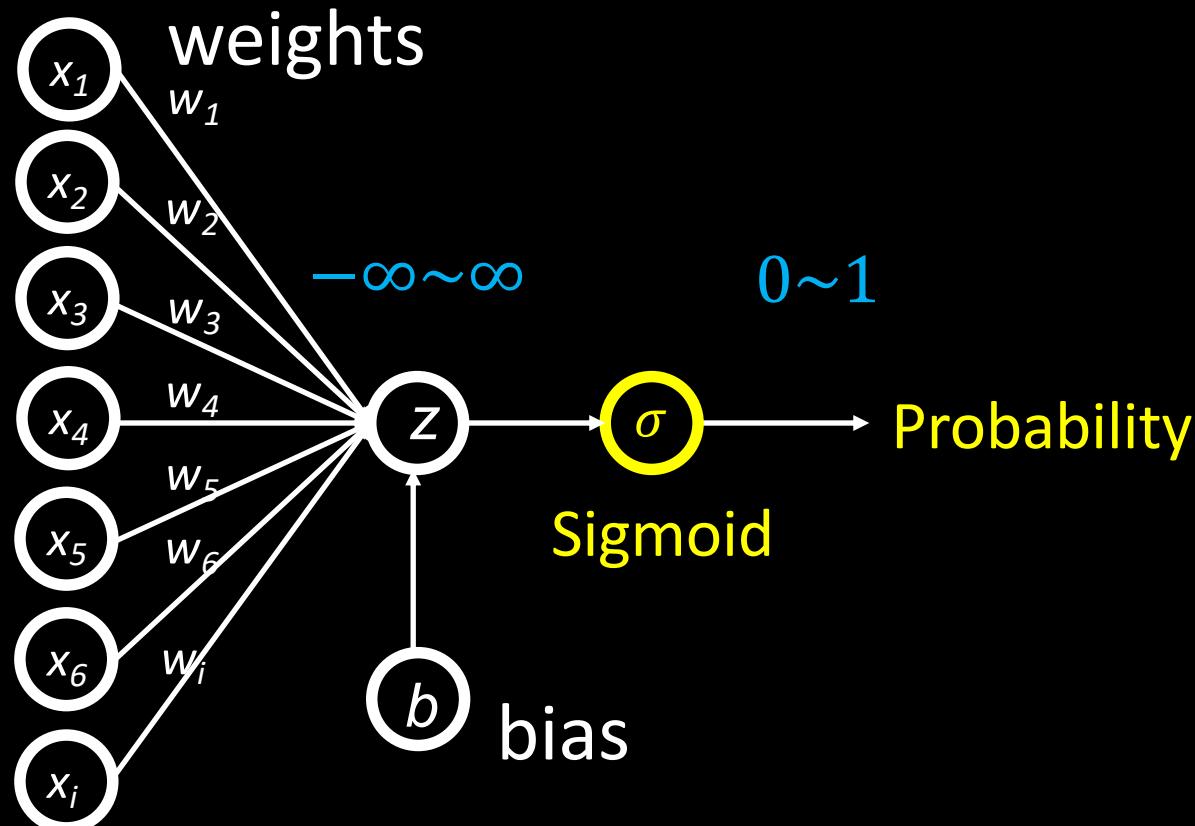


Graph of Linear Regression

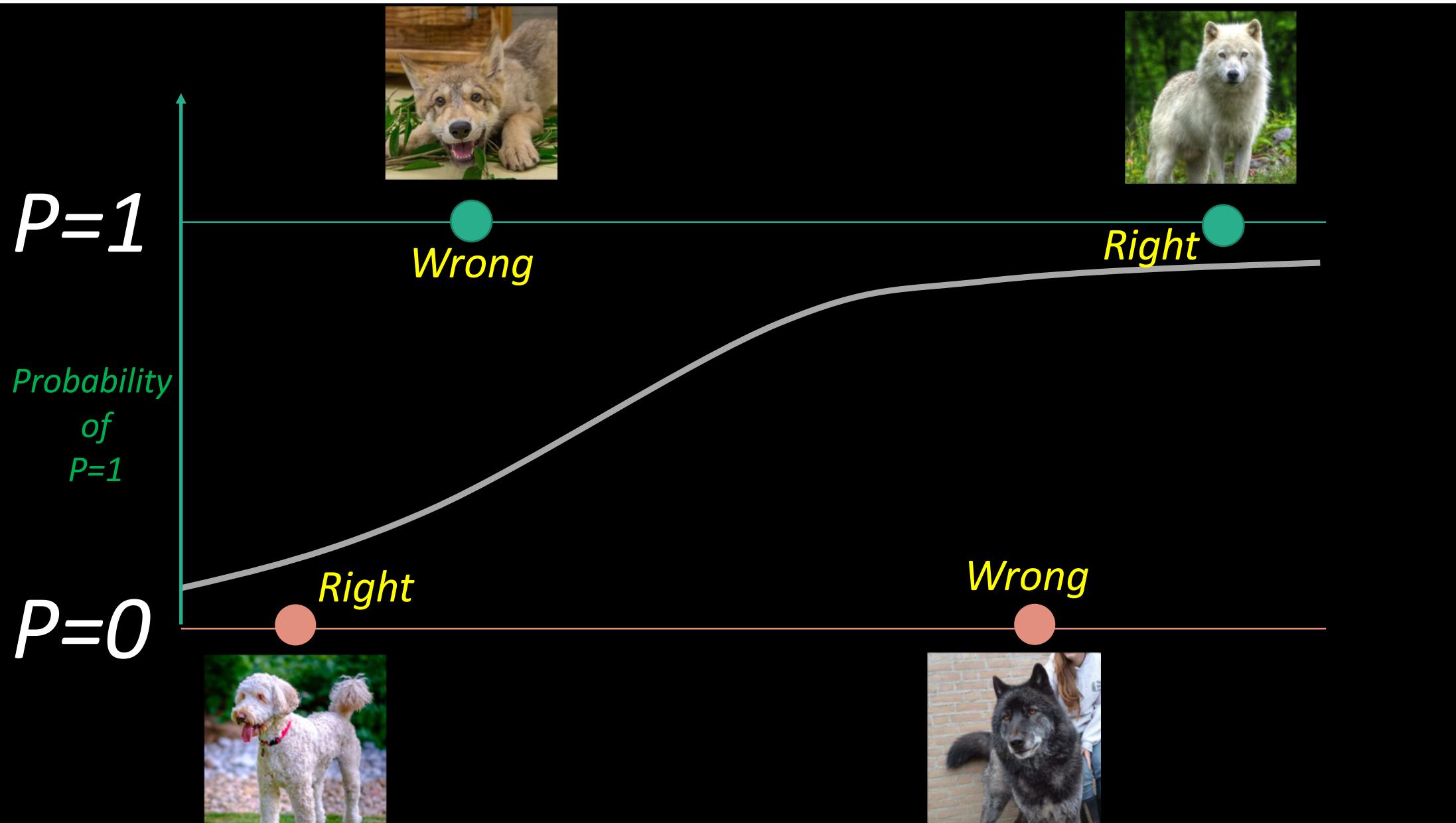


Graph of Logistic Regression

$$p = \frac{1}{1 + \exp(-(\sum_i w_i x_i + b))} = \text{Sigmoid}(\sum_i w_i x_i + b) = S(WX)$$



Loss Function for Classification



$P=1$



Probability of $P=0$

$P=0$



Probability of $P=1$

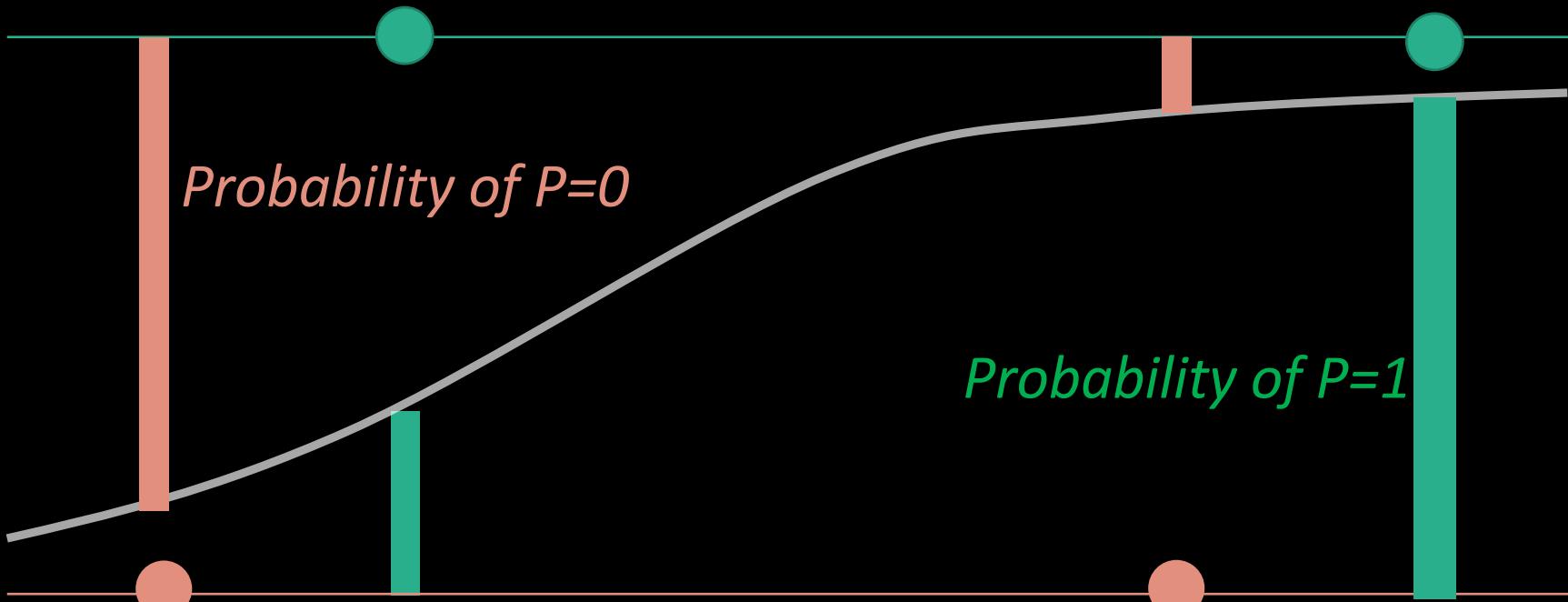
How right are we?

$P=1$

Probability of $P=0$

$P=0$

Probability of $P=1$

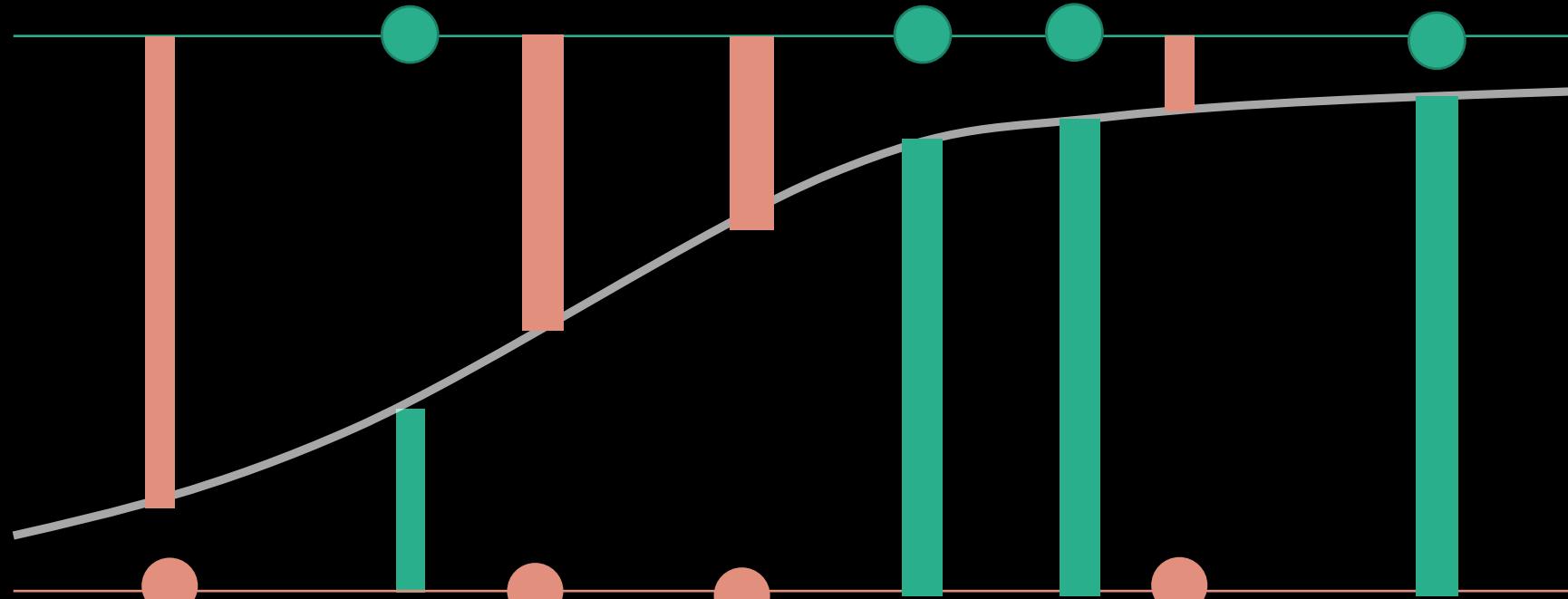


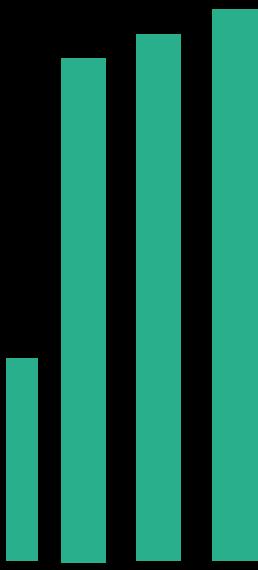
How right we are?

Probability of all the cases= $p_0 * p_1 * p_2 * p_3 * p_4$

$P=1$

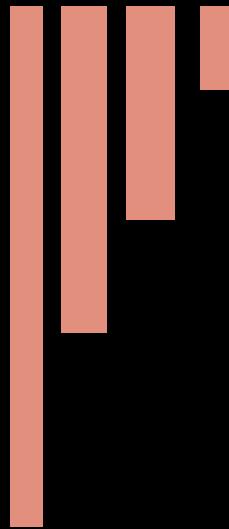
$P=0$





Probability of all the cases

$$(+)\ p_0 * p_1 * p_2 * \dots \dots$$



$$(-)\ p_0 * p_1 * p_2 * \dots \dots$$

Remember

$$\prod_n P(y_n|x_n; \theta)$$

VS

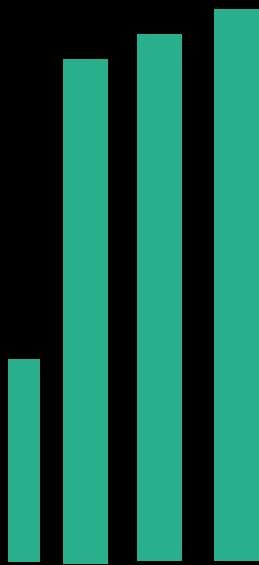
$$\sum_i \log P(y_n|x_n; \theta)$$

Binary Cross Entropy (BCE) loss

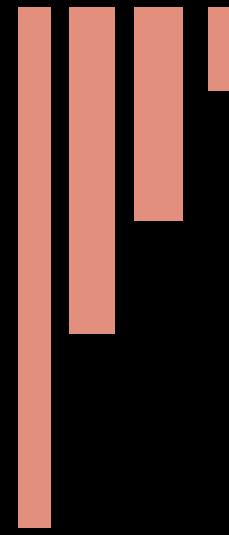
$y = 1$

$y = 0$

$$BCE = -\sum_n y \cdot \log(p(\hat{y}))$$



$$BCE = -\sum_n (1 - y) \cdot \log(p(1 - \hat{y}))$$



Binary Cross Entropy (BCE) loss

$BCE =$

$$-\sum_n [y \cdot \log(p(\hat{y})) + (1 - y) \cdot \log(p(1 - \hat{y}))]$$

Loss if $y=1$

Loss if $y=0$

Compare these
Models

Assuming
Truth = 1

Very Correct

P(0)	P(1)
0.3	0.7

Should be rewarded

Correct

P(0)	P(1)
0.45	0.55

Wrong

P(0)	P(1)
0.7	0.3

Should be punished

Very Wrong

P(0)	P(1)
0.9	0.1

Compare these
Models

Assuming
Truth = 1

Very Correct

P(0)	P(1)
0.3	0.7

Should be rewarded more

Correct

P(0)	P(1)
0.45	0.55

Should be rewarded less

Wrong

P(0)	P(1)
0.7	0.3

Should be punished less

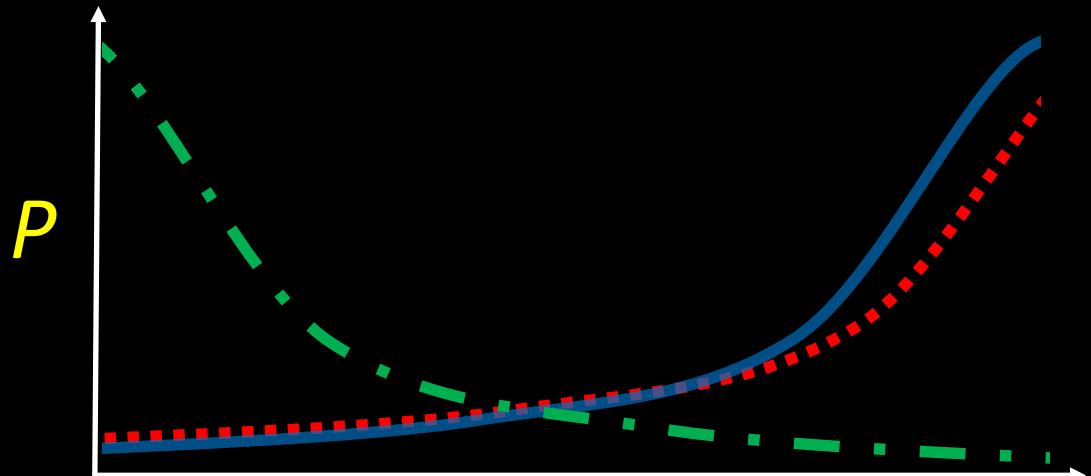
Very Wrong

P(0)	P(1)
0.9	0.1

Should be punished more

Again, we can use Maximum Likelihood Estimation (MLE)

$$\theta_{ML} = \operatorname{argmax}_{\theta} P(y|X; \theta)$$



Q, Real Probability

$P(y|X; \theta)$, good prediction

$P(y|X; \theta)$, bad prediction

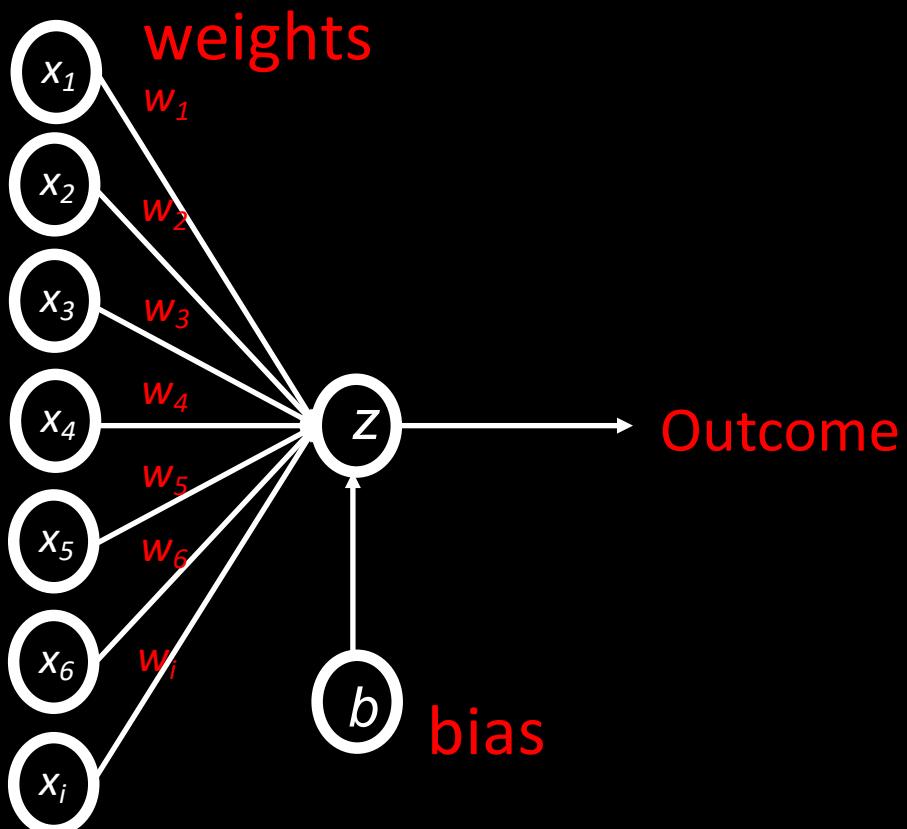
Cross Entropy:
how different they are?

Cross Entropy:
how different they are?

	<i>Q</i>	<i>P</i>		
<i>Good model</i>	0.9	0.9	(0.09)	> 0.32 <i>Good</i>
	0.1	0.1	(0.23)	
<i>Bad model</i>	0.9	0.1	(2.07)	> 2.08 <i>Bad</i>
	0.1	0.9	(0.01)	

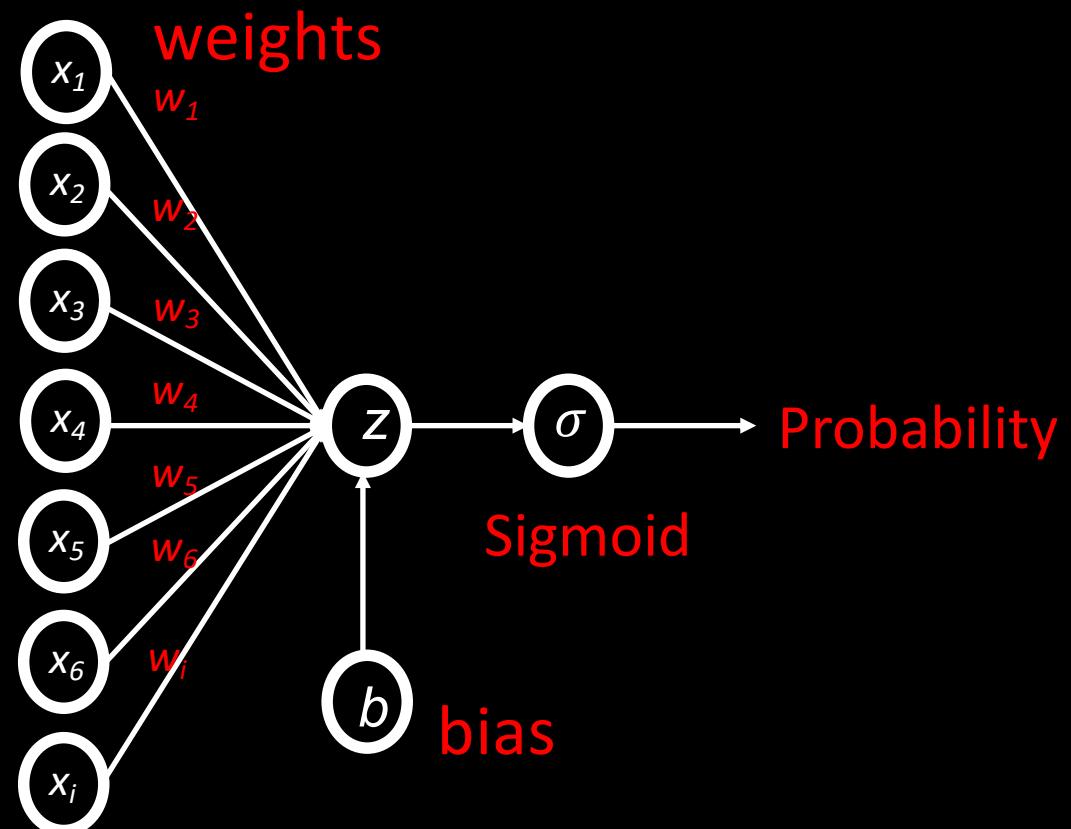
$$H(Q, P) = - \sum_n Q_i \cdot \log(P_i)$$

Regression



Loss: Mean Square error

Classification (logistic regression)



Loss: Cross Entropy

In practice!

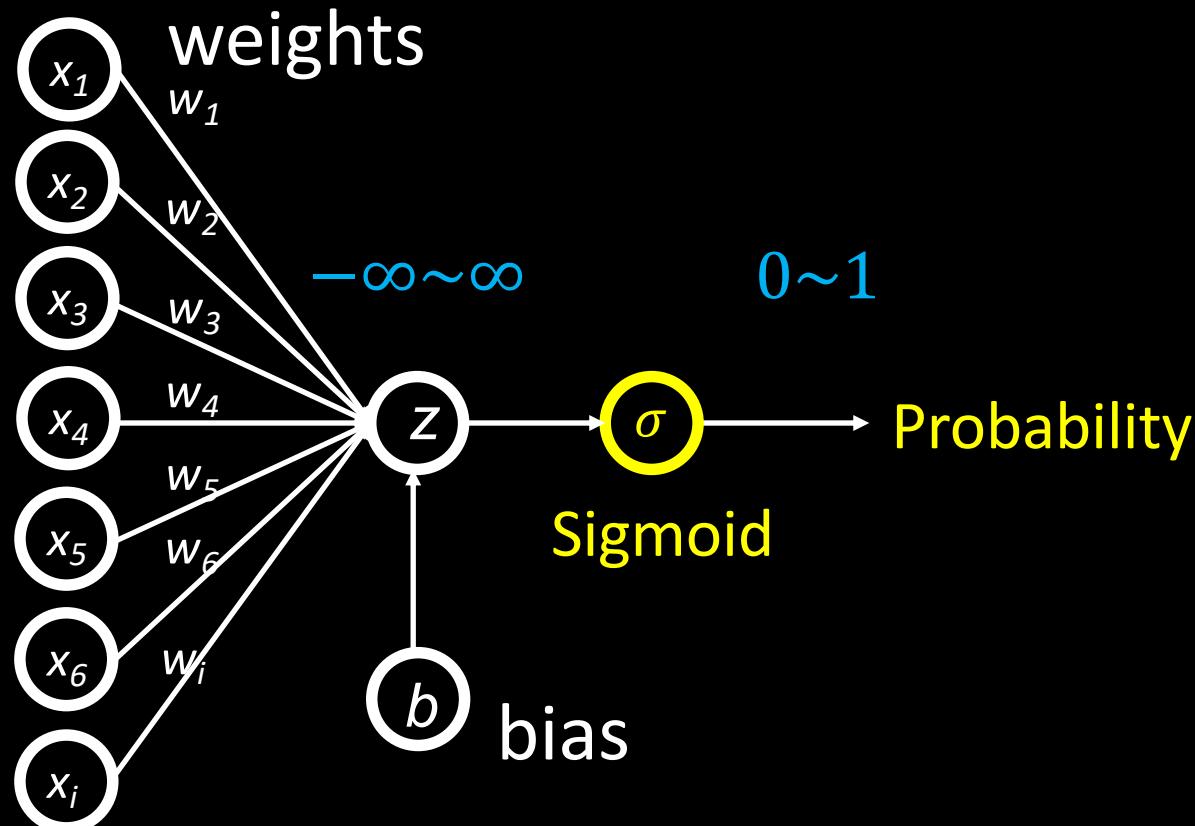
```
model = LinearRegression()  
model.fit(weights, dosages)
```

```
log_model = LogisticRegression()  
log_model.fit(cholesterol, heart_disease)
```

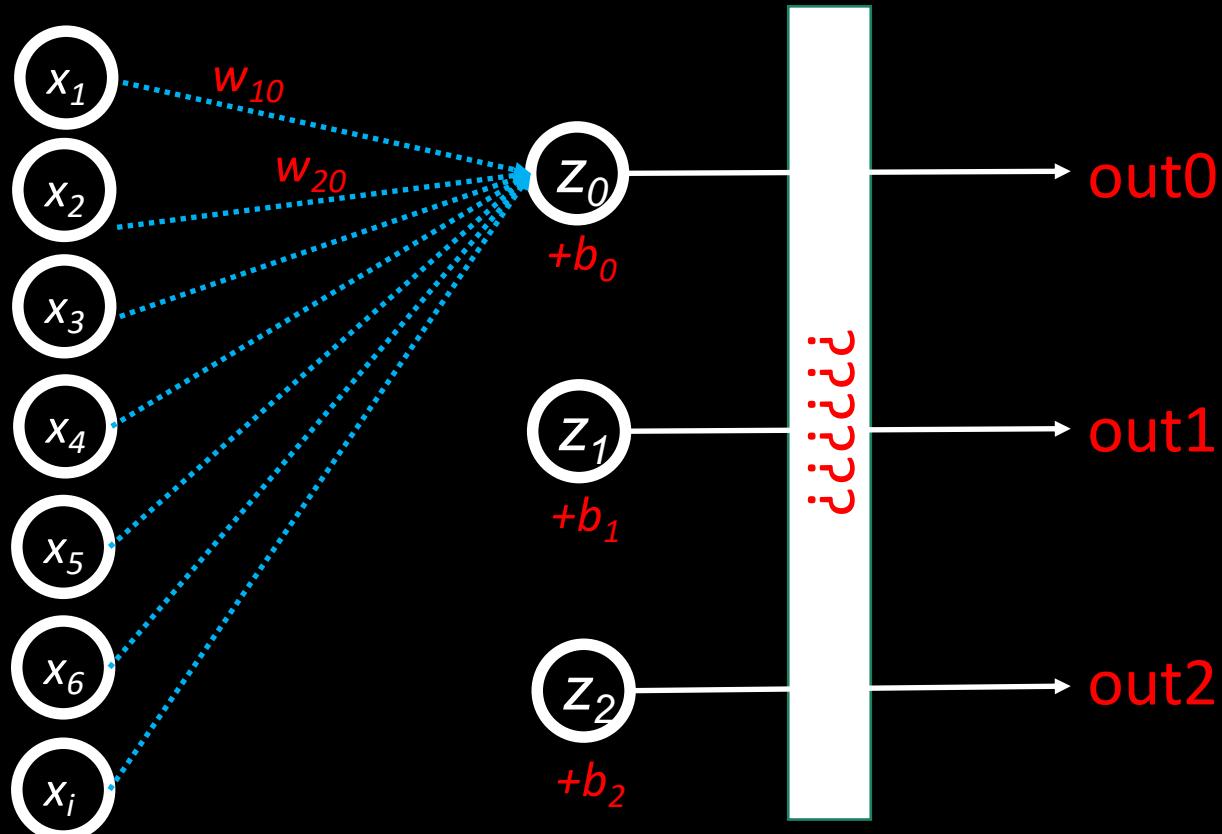
Multi-class Logistic Regression

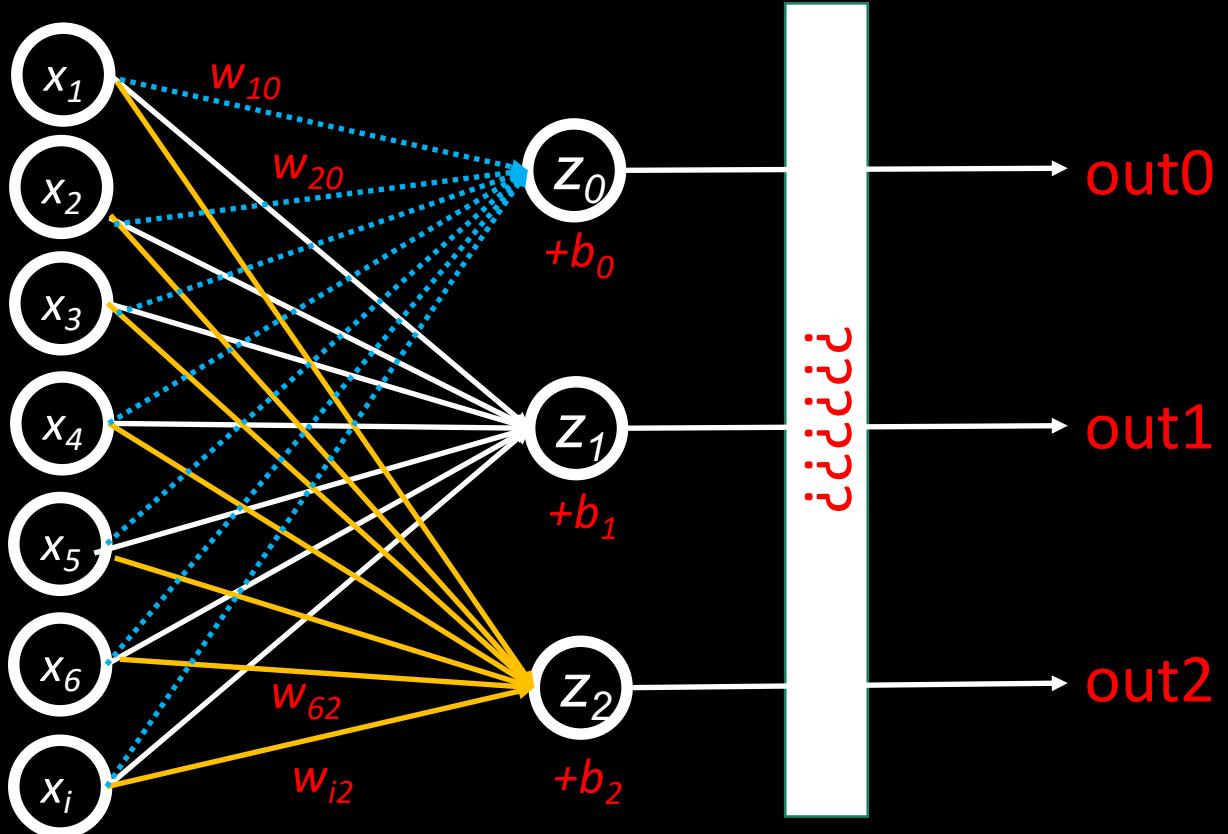
Single Class Logistic Regression

$$p = \frac{1}{1 + \exp(-(\sum_i w_i x_i + b))} = \text{Sigmoid}(\sum_i w_i x_i + b) = S(WX)$$

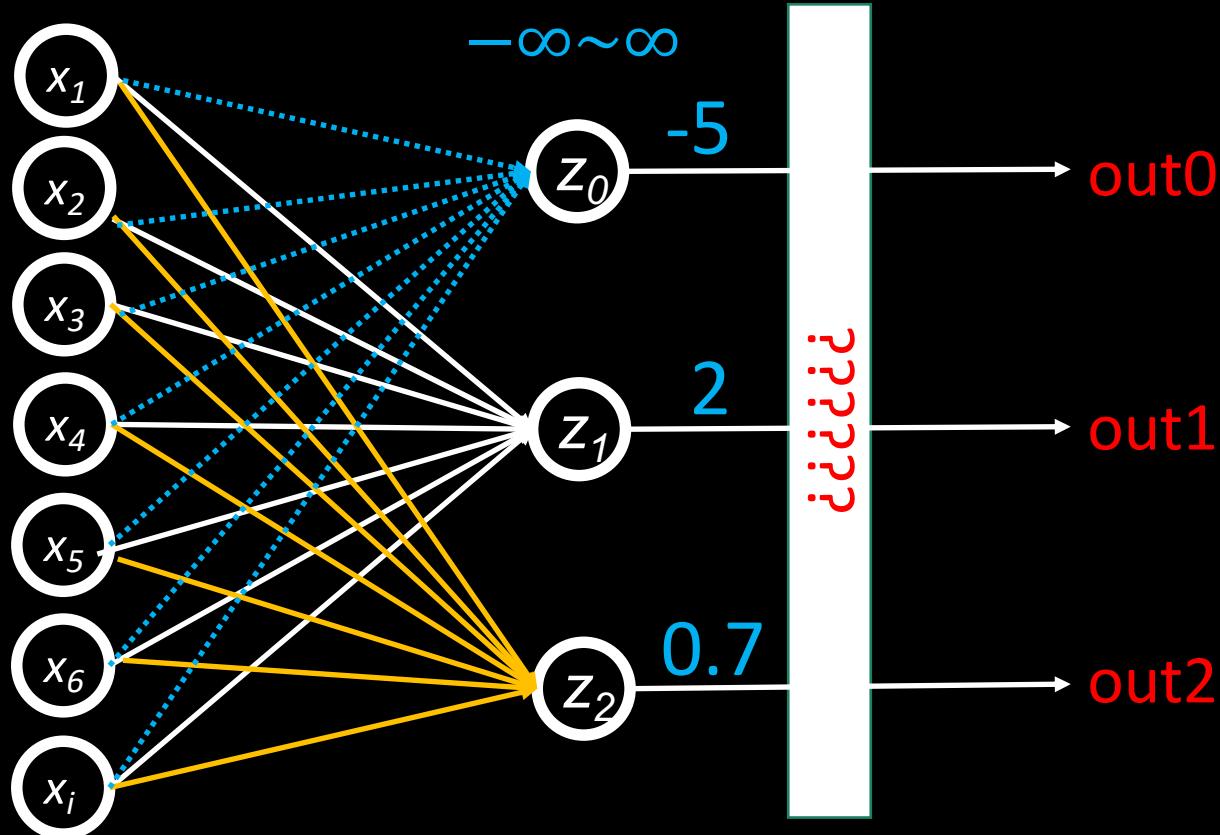


$$\text{out}0 = w_{10}x_1 + w_{20}x_2 + w_{30}x_3 + \dots$$





$$\text{out}_j = w_{1j}x_1 + w_{2j}x_2 + w_{3j}x_3 + \dots$$



Probability
0~1
&
(out0+out1+out2+....=1)

$$p(y = N) = \text{Softmax}(W_N X) = \frac{\exp-(W_N X)}{\sum_n \exp-(W_n X)}$$

Softmax ==
Sigmoid for multi-class

Data	Output (Wx)	$\exp(Wx)$	argmax	Softmax $\exp(Wx) / \sum(\exp(Wx))$
3	-0.84	0.43	0	0.02
4	2.69	14.73	1	0.68
5	1.76	5.81	0	0.27
6	-0.69	0.50	0	0.02



$$p = \frac{1}{1 + \exp - (\sum_i w_i x_i + b)} = \text{Sigmoid}(w_i x_i + b) = S(WX)$$

V S

$$p(y = 0) = SX(W_0 X) = \frac{\exp - (W_0 X)}{\sum_n \exp - (W_n X)}$$

For multi-classes, you will have one W for each class

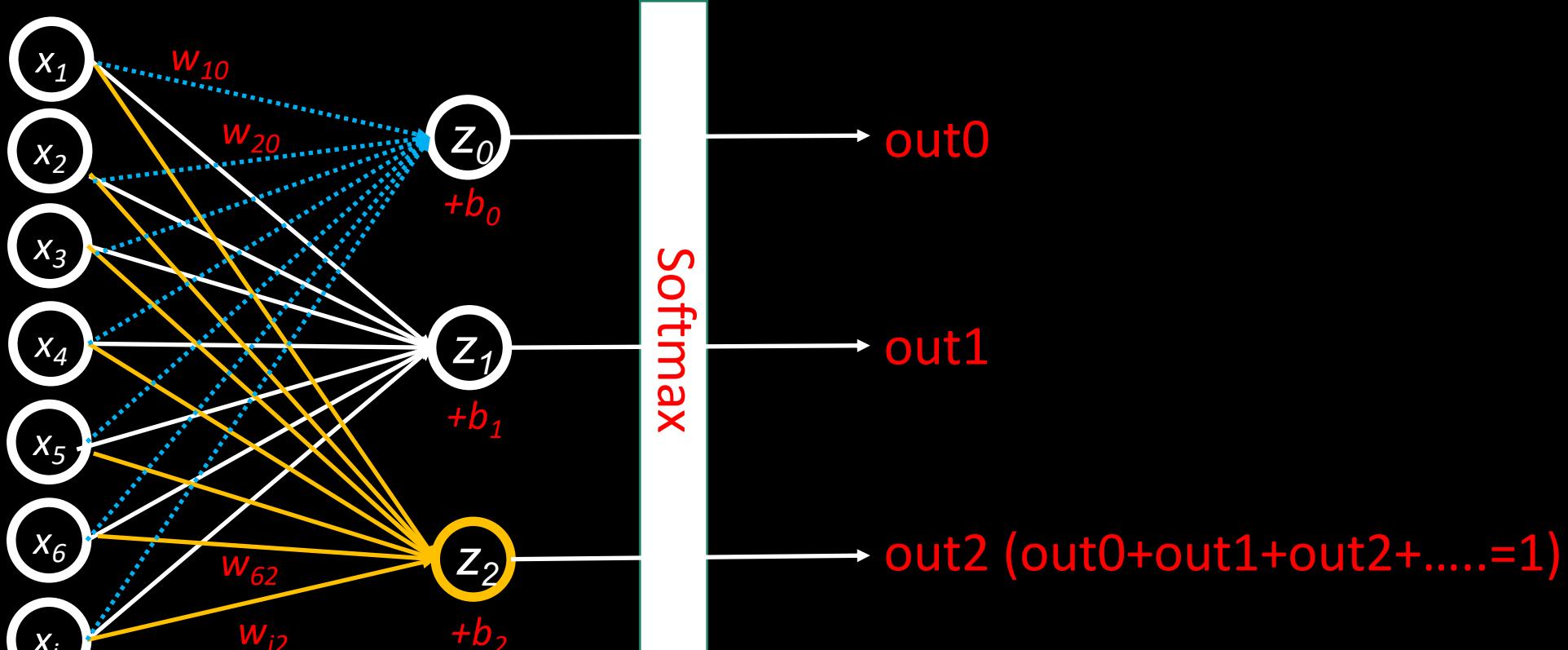
$$p(y = 1) = SX(W_1 X) = \frac{\exp - (W_1 X)}{\sum_n \exp - (W_n X)}$$

•

•

$$p(y = N) = SX(W_N X) = \frac{\exp - (W_N X)}{\sum_n \exp - (W_n X)}$$

The summation for all the P
Should be 1



Multi-class Logistic Regression ==

Linear regression between probability and Softmax(WX)

Binary Cross Entropy

$$-\sum_n [y \cdot \log(p(\hat{y})) + (1 - y) \cdot \log(p(1 - \hat{y}))]$$

Multi-Class Cross Entropy

$$-\sum_n y \cdot \log(p(\hat{y}))$$

$$\begin{matrix} y & \hat{y} \\ \hline 0 & 0.6 \\ 1 & 0.3 \\ 0 & 0.1 \end{matrix}$$

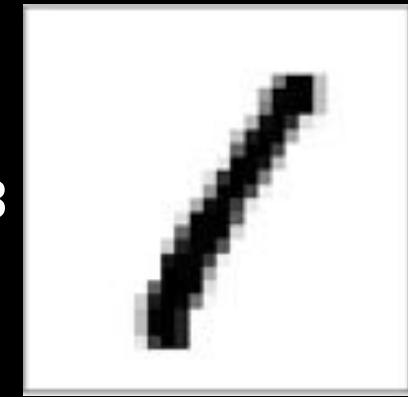
$$= -1 * \log(0.3)$$

Image Classification by Logistic Regression



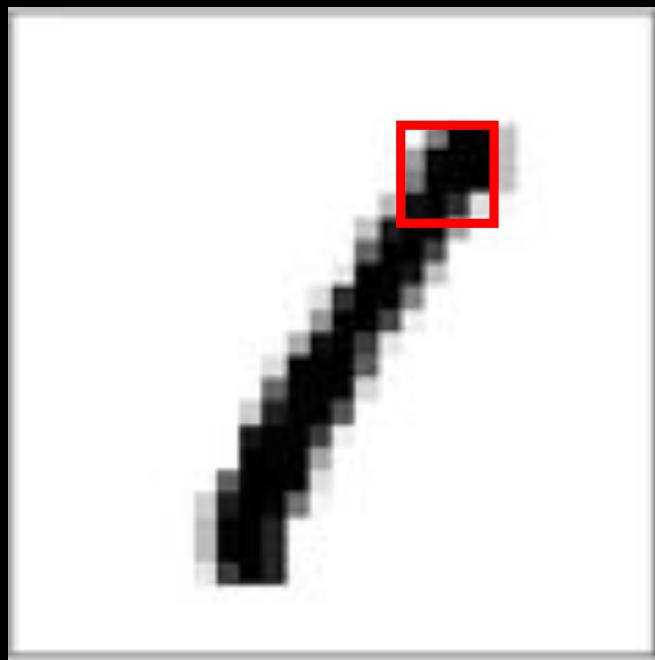
MNIST dataset

28



28

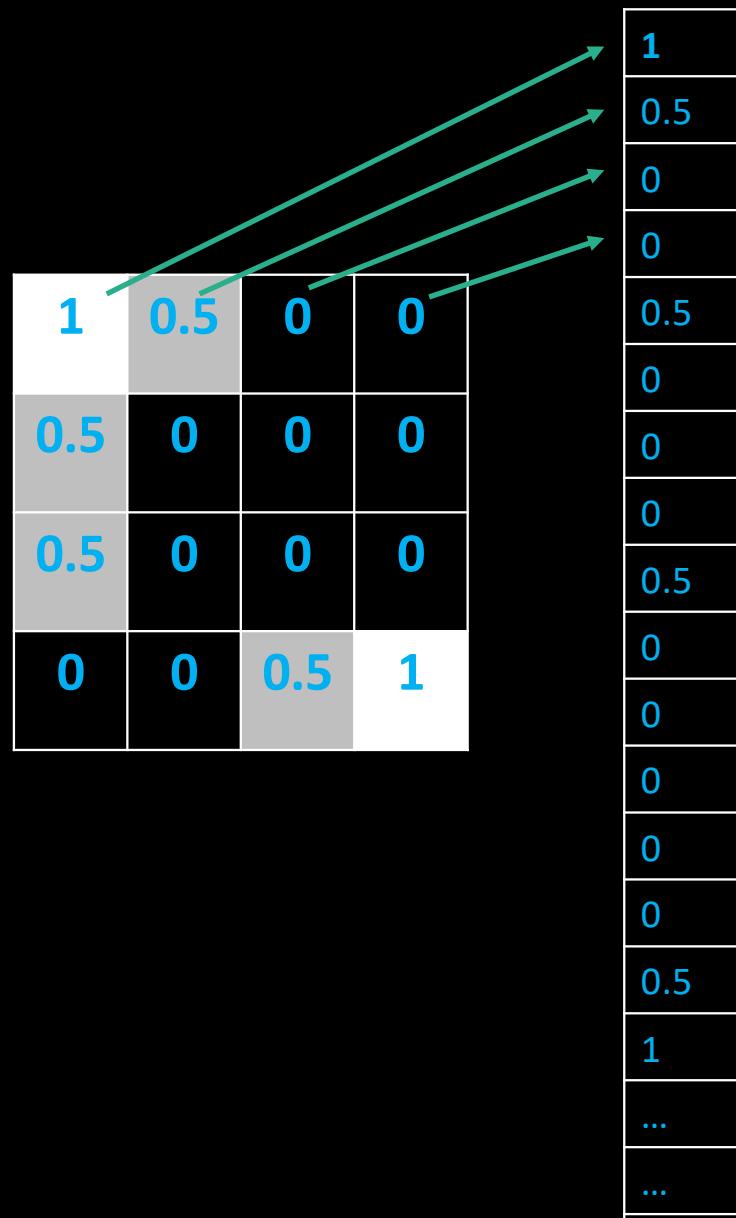
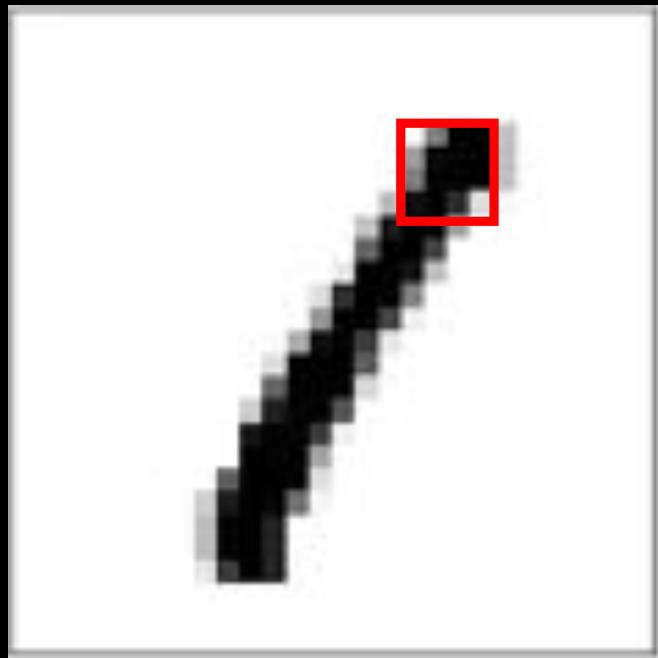
Image to grayscale pixel value



white	light gray	black	black
gray	black	black	black
gray	black	black	black
black	black	light gray	white

1	0.5	0	0
0.5	0	0	0
0.5	0	0	0
0	0	0.5	1

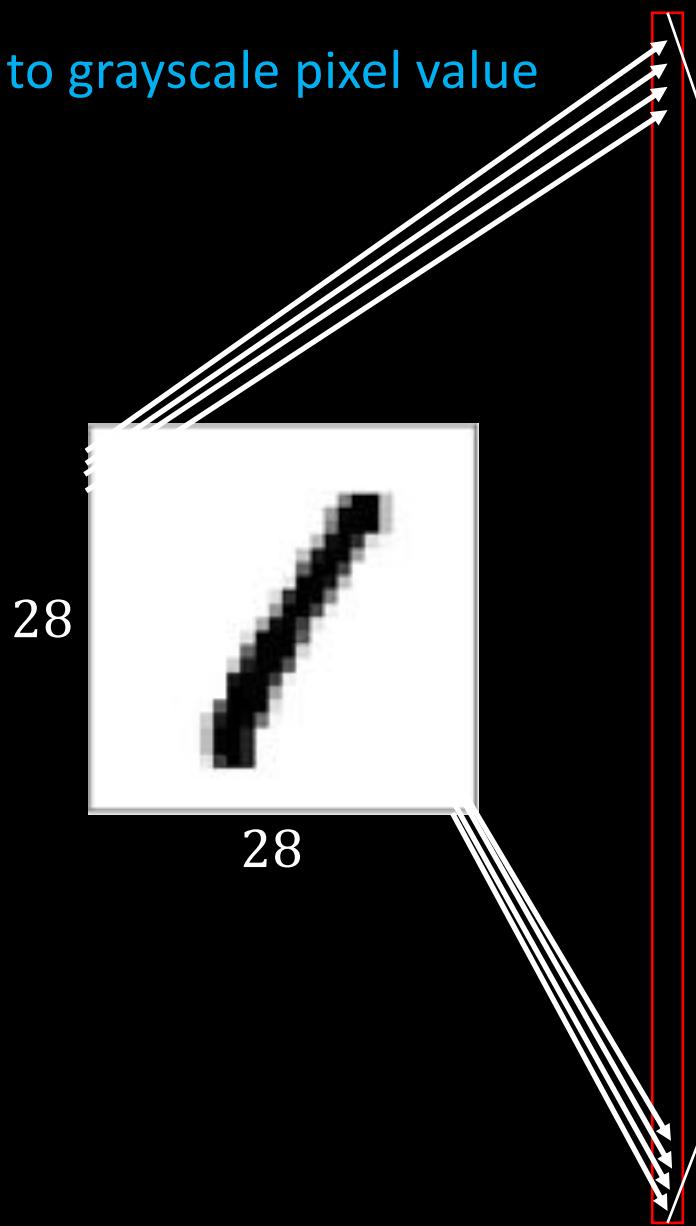
Image to grayscale pixel value



`numpy.ndarray.flatten`

`torch.array.view(-1)`

Image to grayscale pixel value



x a vector with length of 784 (28^2)

y “1”

$x \rightarrow y$ $784 \rightarrow 1$

