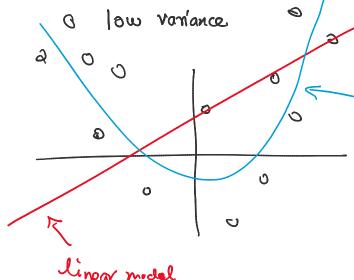
Monday, November 6, 2023

10:02 AM

Simple models -> high bias

e.g. just linear



the underlying generative process

linear model

Complex Model: Tit anything low bias high variance

 $d = (y - \hat{y})^2$

y = twe targer

y - predicted value

Squared error

Por

goal: min expected lass (on test cases)

 $E[(y-y)^{2}] = E[(y-uy)^{2}]$ expected law (vorionce of the target)

(x, y) - given

Moenge voncourse cons

Ensemble methods

Rootstrop Aggregation (Bagging)

> reducing variance

Bogging (D, Algo)

$$D_i = beotstrop sample (D)$$

poststoop sampling

Sample with replacement

$$|D| = n$$

M. Mr. . Mr.

K. different models

græn any point 2

2 closes

clamfication: majority whe { \$9,92, - \$2}

H A L L L

L

regression: $E[\mathcal{G}] = \frac{1}{k} (\mathcal{G}_1 + \mathcal{G}_2 + \cdots + \mathcal{G}_k)$

Veduce voniance: better applies to a more complex model

(low bias)

Will help lower variance

Randon Fores:

Ensemble of decision trees on bootstrap samples

To sample the attribute to split on

(A. A.)

ole the attribute to split or at every hode

(A, Az - Ad)

sample m<<d attributes

As Az -

Boosting

s reduce bias via enremble

I generally effective for simple much that have high bias

Adaphie Boosting (Adaboost)

> biases sampling

> pick points where previous round makes mistakes

Adaboust (D, Algo, K)

$$\widetilde{\omega} = \left(\frac{1}{n}, \frac{1}{n}, \frac{1}{n}\right) = \widetilde{\omega}$$

for i=1 . .. k

prob vector

wi = prob that point xi will be sampled

) Di = Sample D according to 1 g size N

Mi = Algo (Di)

 $\mathcal{E}_i = \text{enor on } \mathcal{D}$ $\mathcal{E}_i = \text{enor rate} = \underbrace{\# \mathcal{D}}_{\text{Nistake}}$

 $d_i = ln\left(\frac{1}{\epsilon_i} - 1\right) = score i Classifier Mi$

reweighting the point

$$\forall j = 1 \cdots n$$

if hi makes no mistate on x;

then we stays the same

else: 1 M' makes a Midrake

$$\omega_{j} = \omega_{j} \cdot \exp(\alpha_{i})$$

$$= \omega_{j} \times \exp(\ln(\frac{1}{\epsilon_{i}} - 1))$$

$$= \omega_{j} \cdot (\frac{1}{\epsilon} - 1)$$

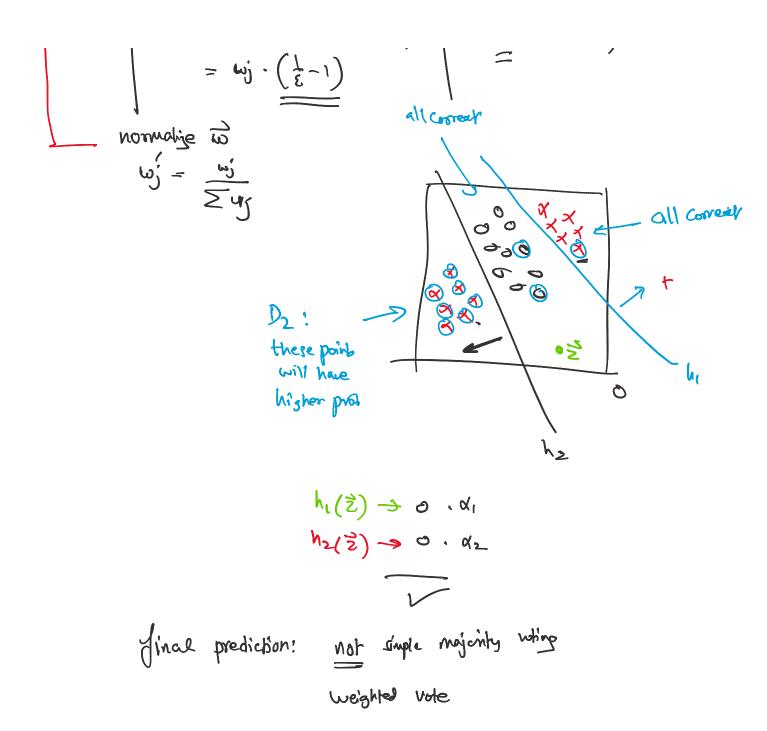
$$\mathcal{E}_{i} \leq 0.5$$

$$\mathcal{E}_{h} \left(\frac{1}{0.5} - 1 \right) = \mathcal{Q}_{h}(2-1) = \ln(1)$$

$$\mathcal{E}_{i} \leq 0.25 = \frac{1}{4}$$

$$\mathcal{Q}_{h} \left(\frac{1}{1/4} - 1 \right) = \mathcal{Q}_{h}(3-1) = \frac{1}{100}$$

$$\mathcal{Q}_{i} \leq \frac{1}{100}$$



Unsupervised Methods

Pattern Mining Clustering

Pattern Mining

tronsaction danser

$$T = d \quad \text{set } q \text{ lience } S = d \quad A, B, C, D, E S$$

$$T = d \quad \text{Subsolv } q \quad \text{lience } S = 1. \quad \{A, B, D, E\}$$

$$X = \{A, B\} = AB$$

$$2. \quad \{B, C, E\}$$

$$3. \quad \{A, B, C, E\}$$

$$4. \quad \{B, C, E\}$$

$$4. \quad \{A, B, C, E\}$$

$$4. \quad \{A, B, C, E\}$$

$$5. \quad \{A, B, C, E\}$$

$$6. \quad \{A, B, C, E\}$$

Ti C I

Q: what are the frequently occurring subset?

$$sup(x) = absolute suppor(x) = count(x)$$

$$\int_{N}^{N} (x) dx = \sum_{n=0}^{N} (x) dx$$

bup of X

(joint prod of all
the Herw in X)
relative suppor = P(X)

|I| is usually very large |I| |I|

$$|D| = n$$
 > very large

XCI X b Some Henser (sbrer of I

Abrolute repoère 3.5

