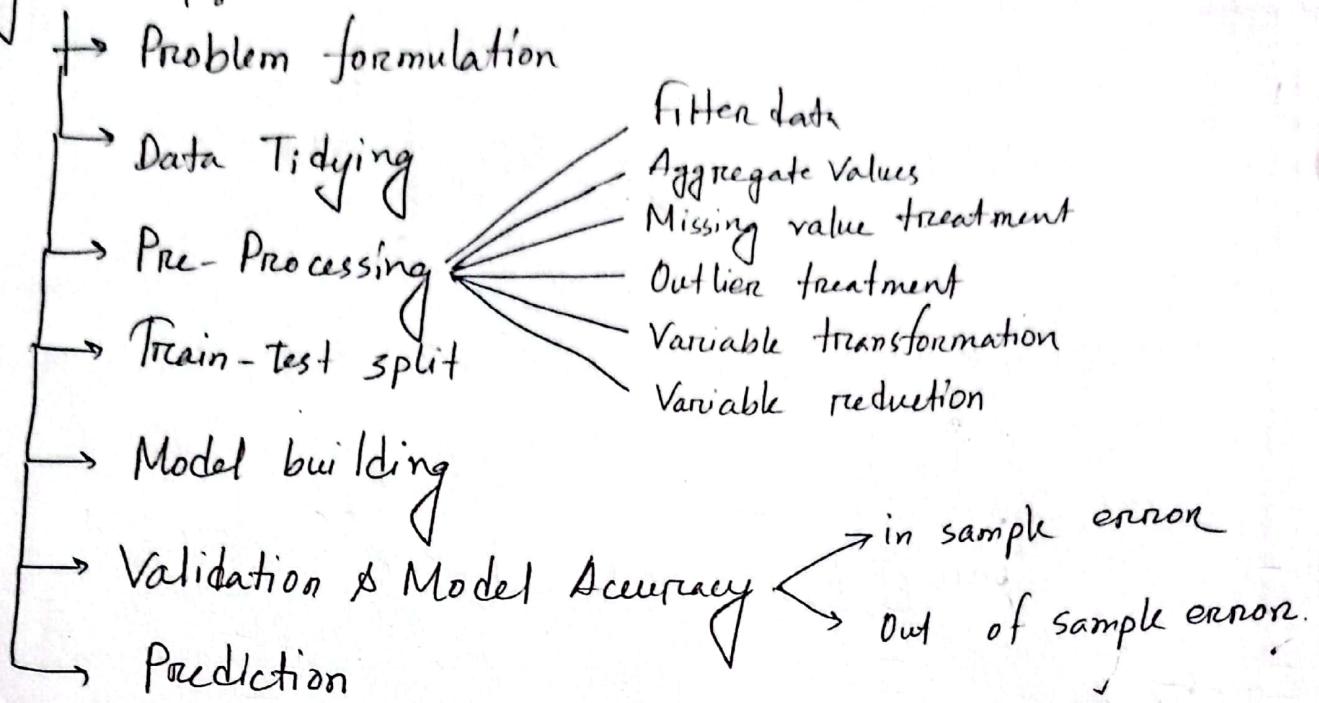


Machine learning

①

building model:



Jupyter:

pwd → working directory

df = pd.read_csv('file.csv'), header=0)

df.head()

DATA SET

df.shape
(row no, col no)

Univariate analysis:

pattern finding → 0% to 25%

EDD: (extended data dictionary)

min, max, mode, avg, 25%, 50%, 75%. It is a summary,

df.describe()

Seaborn:

seaborn

import seaborn as sns

sns.distplot(data.age) [histogram of age]

sns.distplot(data.age, kde, False) [mehrere PDF überlappend]

" " (" ", color="red")

sns.Jointplot(x="x", y="y", data="data-name")
↳ [Scatter plot (nurjed)]

sns.pairplot(data-name) → [variable variable scatter plot matrix]

outlier:

sns.Jointplots(x="n_hot_rooms", y="price", data=df)

Treating outliers: column → to see percentile to see array size
uv = np.percentile(df.n_hot_rooms, [99]) [0] 97.9%

df[(df.n_hot_rooms > uv)]

↳ (2) value uv → (273 ist der Wert)

df.no_hot_rooms[(df.n_hot_rooms > 3 * uv)] = 3 * uv

lv = np.percentile(df.maintain, [1]) [0]

df[(df.Rainfall < lv)]

df.Rainfall[(df.Rainfall < 0.3 * lv)] = 0.3 * lv

variable transformation:

Treating Missing values:

df.n_hos_beds = df.n_hos_beds.fillna(df.n_hos_beds.mean())

संस्कृत दोषो ग्रन्ति बुद्धिमत्ता :-

df = df.fillna(df.mean()) .

Bivariate analysis:

जटिल independent variable फैला मात्र।

(1) df.crime_rate = np.log(1 + df.crime_rate)

df['avg_dist'] = (df.dist1 + df.dist2 + df.dist3 + df.dist4) / 4

del df['dist1']

जटिल सम्बन्ध वर्णन :-

(2) scatter plot

(3) correlation matrix: 0.22 रेलवे संख्या, high 22 रेलवे
independent variable current independent ना, 0.87 रेलवे
रेलवे रेलवे रेलवे।

Creating dummy variable:

df = pd.get_dummies(df)

del df['airport_NO']

del df['waterbody_Non']

Correlation analysis:

df.corr()

Linear Regression:

$$Y = \beta_0 + \beta_1 X$$

Model coefficient

Residual, $e_i = y_i - \hat{y}_i$

Population regression line, sample regression line

$$[\hat{\beta}_1 - 2 \times SE(\hat{\beta}_1), \hat{\beta}_1 + 2 \times SE(\hat{\beta}_1)]$$

↳ true value of β_1 , ~~in~~ interval ≈ 276 mm.

Hypothesis test:

H_0 : No relationship between X & Y

H_a : Some n

To disapprove H_0 ,

we calculate T statistics, $t = \frac{\hat{\beta}_1 - 0}{SE(\hat{\beta}_1)}$

+ value ~~list~~ ~~list~~ ~~list~~ probability of observing any value is $= |t|$ or larger.

↳ ~~list~~ probability (≈ 0.01) \approx p value.

p value ~~list~~ ~~list~~ X ~~list~~ Y \Rightarrow ~~list~~ Relation CMΣ,

Accuracy of Model:

$$\rightarrow RSE = \sqrt{\frac{(y_i - \hat{y}_i)^2}{n-2}}$$

$$\rightarrow R^2 = \frac{TSS - RSS}{TSS} = 1 - \frac{RSS}{TSS}$$

↳ (0 (20%, 70% 28%) 1 to 100% ~~list~~ ~~list~~ ~~list~~)

Simple linear Regression in Python:

(3)

```
import statsmodels.api as sm
```

```
X = sm.add_constant(df['room-num'])
```

\downarrow
model object \downarrow
ordinary least square

 $lm = sm.OLS(df['price'], df['room-num']).fit()$

lm.summary

another method: (most used)

```
from sklearn.linear_model import LinearRegression
```

 $y = df['price'] \rightarrow 2d array$
 $X = df[['room-num']]$

lm2 = LinearRegression()

lm2.fit(X, y)

Print(lm2.intercept_, lm2.coef_)

lm2.predict(X)

sns.jointplot(x=df['room-num'], y=df['price'], data=df,
kind='reg')

Multiple linear Regression:

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p + \epsilon$$

β_0 = intercept
 p = no. of predictors
 ϵ = an error term

F statistics test for all

At p value threshold α

Fit the regression model predictor (x_i) in response

To predict price, for individual x_i value and p-value

Categorical variable

estimate bpt 227 210 value no 81.75 or 2020,

GDP P value b135 1410T no Dependency on
Relationship between result & variable.

test-train split:

from sklearn.model_selection import train_test_split

X_train, X-test, y_train, y-test = train_test_split (X-multi,
X-multi = df.drop ("price", axis=1) $\xrightarrow{\text{column drop}} \xrightarrow{\text{for row drop}}$

y-multi = df['price']

X-multi_cons = sn.add_constant (X-multi)

lm-multi = sn.OLS(y-multi, X-multi_cons).fit()

lm-multi.summary()

Another Method:

lm3 = linear Regression()

lm3.fit (X-multi, y-multi)

print (lm3.intercept_, lm3.coef_)

Test-train split:

from sklearn.model_selection import train_test_split

X_train, X-test, Y_train, Y-test = train_test_split
(X-multi, y-multi, test_size=
random_state=0)

print(x_train.shape, x_test.shape, y_train.shape, y_test.shape) ①

lm_a = LinearRegression()

lm_a.fit(X_train, Y_train)

y-test-a = lm_a.predict(X-test)

y-train-a = lm_a.predict(X-train)

from sklearn.metrics import r2_score

[r2-score?] details

r2_score(y-test, y-test-a)

r2_score(Y-train, y-train-a)

② Ridge & Lasso Regression: → ① step 2 data standardizing
from sklearn import preprocessing stores info about how to set

scalar = preprocessing.StandardScaler().fit(X-train)

X-train-s = scalar.transform(X-train)

X-test-s = scalar.transform(X-test)

from sklearn.linear_model import Ridge

lm_r = Ridge(alpha=0.3)

lm_r.fit(X-train-s, y-train)

r2_score(y-test, lm_r.predict(X-test-s))

[alpha vary w.r.t validation curve]

from sklearn.model_selection import validation_curve

param_range = np.logspace(-2, 8, 100)
↳ $10^{-2} \dots 10^8$ 100 values
create 100

train_scores, test_scores = validation_curve(Ridge(), X_train_s,

y_train, "alpha", param_range, scoring='r2')

print(train_scores) → Only one line
print(test_scores)

train_mean = np.mean(train_scores, axis=1)

test_mean = np.mean(test_scores, axis=1)

train_mean

max(test_mean)

sns.jointplot(x=np.log(param_range), y=test_mean)

np.where(test_mean == max(test_mean))

param_range[33]

lm_r_best = Ridge(alpha=param_range[33])

lm_r_best.fit(X_train_s, y_train)

r2_score(y_test, lm_r_best.predict(X_test_s))

r2_score(y_train, lm_r_best.predict(X_train_s))

Lasso:

from sklearn.linear_model import Lasso

lm_f = Lasso(alpha=0.4)

: (same as Ridge)

problem we may face is called Heteroscedasticity

(5)

$\log y \sigma \log y$ find variance among

Outlier detect $\text{np}(\text{not})$ box plot 3 we obt 20

sns.boxplot(y = "n_hot_rooms", data=df)

categorical vs not countplot

sns.countplot(x="airport", data=df)

② missing value

df.info

filter character n-hos-beds A var data are^t,
categorical & not numerical &

carat, EDD (age) 3200 min, But edd only

symm num,

Soln: df.n-hos-beds = df.n-hos-beds.fillna(df.n-hos-beds.mean())

df = df.fillna(df.mean())

③ Ver transformation & deletion

→ $\sqrt{\text{sum}(\text{dist}^2)}$ distance (norm)
 $\text{df}[\text{'avg_dist'}] = (\text{df.dist1} + \text{df.dist2} + \text{df.dist3} + \text{df.dist4}) / 4$

drop dist 4 variables delete rows 3 & 20

drop df["dist1"]

drop df["dist2"], ... [dist4]

drop var - ten col" (cont) more outliers, 0.02 & 0.03 deleted

del df["bus - ten"]

④ Non-numerical values (or numerical A string)

128) 2021 one dummy variable create 4.3 categories
 $d_f = pd \cdot get_dummies(df)$

airport 228 variable 1 is 0 if airport - yes
airport - yes & 1 if airport - no.

188

Classification algo is used for qualitative or categorical output.

→ Logistic Regression

→ Linear discriminant analysis

K nearest neighbor

(u Ex) question like ml:-

① Prediction question

→ will the house be sold within 3 months of putting

listed

② Influential Question

→ how accurately can we estimate the effect
of each of the predictor variables on the
response variable.

90) Logistic Regression:

$$p(x) = \frac{e^{B_0 + B_1 x}}{1 + e^{B_0 + B_1 x}}$$

Linear Regression A ordinary least square method we use
logistic \rightarrow Maximum likelihood \rightarrow " "

logistic

(1) creating logistic Regression Model

→ sk learn library use \rightarrow preferred
set model u u u

sklearn

$X = df[[\text{'price}]]$ → two dimension 2501 rows

$y = df[\text{'sold}]$

single variable

from sklearn.linear_model import LogisticRegression

clf_lr = LogisticRegression()

clf_lr.fit(X, y)

to see $\beta_0 + \beta_1$

clf_lr.coef_

(β_0, β_1)

clf_lr.intercept_

multiple preditory

use $\text{df}[\text{'size}]\text{ & } \text{df}[\text{'3d}]$ (3d == sold 215 point)

22.4% 200 point, 62.8% 200 point

But

logistic use 201 into c But not main

LDA (linear discriminant analysis use not 20 main

Q5

$X = \text{df.loc[:, df.columns != 'sold']}$ \rightarrow sold one was not fair

$y = \text{df['sold']}$

Q6 clf - In 2 Logistic Regression()

clf - fn. fit(x, y)

clf - ln. coef_

$\Rightarrow (\beta_0)$

clf - ln. intercept -

(β_0)

Q7 confusion matrix

No Yes
true default status

No	Yes	True Default Status
No	9432	138 → type 2 error
Yes	235	193 (false or preferent error) \downarrow 0.001

boundary value 0.312

0.75 no

type 1

error

{
 not pregnant } {
 {
 true
 unlock }
 {
 for anyone
 not
 anyone }
 }

8) creating confusion matrix:

clf - lr. predict_proba(x)

$\begin{bmatrix} \text{Col 1} & \text{Col 2} \\ \downarrow & \downarrow \\ \text{prob of not sold} & \text{prob of sold} \end{bmatrix}$

by default boundary condition = 0.5

y-pred = clf - lr. predict(x).

y-pred = 0.3 = (clf - lr. predict_proba(x)[:, 1]) >= 0.3

accuracy func. (f1)
from sklearn.metrics import confusion_matrix
(y, y-pred)
confusion_matrix(y, y-pred - 0.3)

③ evaluating performance metrics

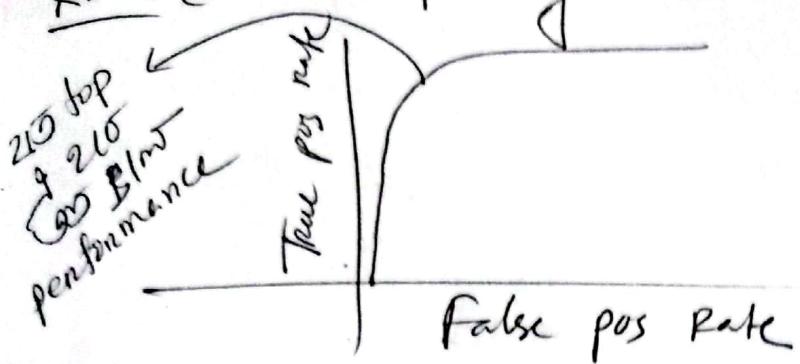
True -
 $\begin{array}{c} \downarrow \\ \text{True Neg (TN)} \\ \hline \text{False Neg (FN)} \end{array}$ → $\frac{\text{True Neg}}{\text{True Neg + False Neg}} \rightarrow N$
+ false Neg (FN) → P

N^* P^* Significance → $\approx \frac{TN}{N}$

Name	def	P^*	Significance
false pos rate	$\frac{FP}{N} \rightarrow$ type I error, 1 - specificity		
true " "	$\frac{TP}{P} \rightarrow$ type II error, power, sensitivity, recall		
pos pred value	$\frac{TP}{P^*}$ precision, 1 - false discovery proportion		
Neg "	$\frac{TN}{N}$		

Neg "

ROC: (Reciever operating characteristics)



100) evaluating model performance:

$$\text{precision} = \frac{\text{True Pos}}{\text{True pos} + \text{False pos}}$$

$$\text{recall} = \frac{\text{True Pos}}{\text{True pos} + \text{False neg}}$$

from `sklearn.metrics import precision_score, recall_score`

`precision_score(y, y-pred)`

`recall_score(y, y-pred)`

From `sklearn.metrics import roc_auc_score`

`roc_auc_score(y, y-pred)`

Linear discriminant analysis

Weight	f_{fit}	not f_{fit}	$P_{fit}(\text{pred})$ assign 215
low	13	22	~35
medium	15	25	~10
high	$\frac{20}{48}$	$\frac{5}{52}$	≈ 2.5

$P_{fit}(\text{pred})$ assign 215
No f_{fit}

- * Bayes classification: Assigns conditional probability to all classes
- * assign the class with highest probability.

$$(41)(\text{pred}) \text{ medium } f_{fit} \text{ & not } f_{fit}$$

$$= \frac{15}{100} = \frac{15}{48} \times \frac{40}{100} \times \frac{15}{40}$$

↑
 Medium
 ↑
 Fit & not
 ↓
 Medium
 ↓
 Fit & not

$$\Rightarrow \frac{15}{40} = \left(\frac{15}{48} \times \frac{40}{100} \right) \div \frac{40}{100}$$

Normal distribution \Rightarrow prediction result from sklearn.

- (103) from sklearn . discriminant analysis import LinearDiscriminantAnalysis

clf_lda = LinearDiscriminantAnalysis()

clf_lda . fit(X,y)

y_pred_lda = clf_lda . predict(X)

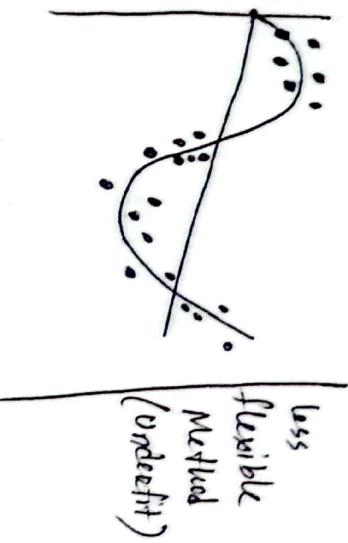
confusion_matrix(y,y_pred_lda)

- (105) K-nearest neighbors classifier

test - train split

Training error \rightarrow Performance of model on the previous seen data.

Test error \rightarrow performance of model on the unseen data.



Split techniques

- Validation set approach
 - (80:20)
 - increase of large no of observations
- Leave one out cross validation
 - ↪ first observation keep out ($n-1$) & use rest training
- K-fold validation
 - divide into K set
 - 1 fold test & other $K-1$ training & fit

(d) from `sklearn.model_selection import train_test_split`

`X_train, X_test, y_train, y_test, train_test_split(X, y, test_size=0.2, random_state=0)`

`clf_LR = LogisticRegression()`

`clf_LR.fit(X_train, y_train)`

`y_test_pred = clf_LR.predict(X_test)`

`from sklearn.metrics import accuracy_score, confusion_matrix`
`confusion_matrix(y_test, y_test_pred) \rightarrow \begin{bmatrix} 36 & 22 \\ 13 & 31 \end{bmatrix} \rightarrow \left\{ \begin{array}{l} \text{actual} \\ \text{class} \end{array} \right. \begin{array}{l} \text{predicted} \\ \text{class} \end{array}`
`accuracy_score(y_test, y_test_pred)`

KNN

Optimum value of K choose upto 25.

scale matter no, cause 1% diff salary, the cost a lot of diff
GDP salary 1000 diff ~~2000~~ ~~2000~~ ~~1000~~, but KNN
will consider this 1000 as very high. And
standardize ratio 25 independent (so)

⑩9) from sklearn import preprocessing
scaler = preprocessing.StandardScaler().fit(X_train)

X_train_s = scaler.transform(X_train)

scaler = preprocessing.StandardScaler().fit(X_test)

X_test_s = scaler.transform(X_test)

from sklearn.neighbors import KNeighborsClassifier

clf_knn_1 = KNeighborsClassifier(n_neighbors=1)

clf_knn_1.fit(X_train_s, y_train)

confusion_matrix(y_test, clf_knn_1.predict(X_test_s))

accuracy_score(y_test, clf_knn_1.predict(X_test_s))

$K=3$ ~~is~~ ~~large~~ is similarly (24 ml 0 smf),

But ~~when~~ $K=22$, value $\sqrt{0.1}$ & 22 model create 22 to
create 22, 21 \hat{y} 1 \hat{y}_1 model create 22 to

Column

~~110~~

110 from sklearn.model_selection import GridSearchCV

params = { 'n_neighbors': [1, 2, 3, 4, 5, 6, 7, 8, 9, 10] }

grid_search_cv = GridSearchCV(KNeighborsClassifier(), params)

grid_search_cv.fit (X-train-s, y-train)
grid-search-cv. best_params_ { (1-10) & (best parameter) }
optimised - KNN = grid-search-cv. best_estimator_-

y-test-pred = optimised - KNN. predict (X-test-s)
confusion - matrix (y-test, y-test-pred)
accuracy - matrix (y-test, y-test-pred)

112 $P_n(Y=1)$ & value \neq result \rightarrow Result \rightarrow impact to other
Estimate variable \rightarrow its impact to other, (-) sign \neq 1
there inverse relation.

\hookrightarrow 128 Logistic Regression & LDA (both ways), KNN

& CART algorithm.

Method $\begin{cases} \text{Logistic}, \text{LDA}, \text{KNN} \\ \text{Accuracy} \end{cases}$

65%

66.6%

55%

LDA

KNN($k=3$)

• All the steps just now

① Data collection (relevant, noisy)

② a) pre-processing

- ↳ outlier treatment (single, multiple)
- ↳ missing value imputation
- ↳ Variable transformation

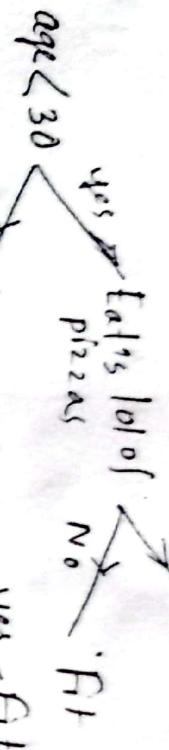
③ Model - training

- ↳ test-train split
- ↳ Use derived template to train
- ↳ Different combination use & compare their
- ↳ test set & apply their result compare

④ select best model

- ↳ For prediction purposes the model with best accuracy
- ↳ For interpretation purposes (like finding out the relation between different variables) look at the coefficient values of parametric models.

(114) Basics of decision tree



Types:

→ Regression tree: for continuous quantitative target variable.

ex: Predicting rainfall, revenue, marks, etc.

→ Classification tree: for discrete categorical target variables

ex: High or low, win or loss, healthy or unhealthy etc

(115) Regression tree

Factors at Salient Basis of Node 4 Q15 आज़?

$$\sum_{j=1}^J \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2$$

actual Response/
predicted value

RSS (Residual sum of squares)

no.	score	hour	mid
1	35	6	42
2	38	5	65
3	40	7	35
4	45	6	75
5	35	8	60
6	65	11	50
7	70	12	45
8	75	18	40
9	80	19	80
10	85	12	82

Mean score

$$RSS = (35-39)^2 + (38-39)^2 + \dots + (85-39)^2$$

$$RSS = (65-75)^2 + (70-75)^2 + \dots + (85-75)^2$$

Ans 5, 5 O.P.M. (2, 8), (3, 7), (1, 6), (6, 4), (7, 3) (9, 2)
एवं सभी combination में RSS (7.7, 7.7) 122/225
एवं RSS 2/21 (25/21) 1/25/21

एक एक mean PI score का एक Midterm
वे एक नियम , वर्गीकृत variable अपनी ओर
बिंदु पर दो बड़ी approach होती हैं।
→ top down, greedy approach - that is known as recursive

binary splitting.

→ top down तथा root node

→ binary भेजना जहाँ नोड बड़े लगते

→ greedy भूल उसले तो वह अधिक से

(2) node का और (3) node का तरीका best

split choose करो।

इसमें Node का लक्ष तोहाँ को depend करें stopping Criteria
करना चाहिए।

116 Stopping Criteria

→ There should be a minimum number of observation required for further split.

→ There should be a minimum number of observation needed at each node after splitting. oddepth
→ There should be maximum layers of tree , possible.

117 Get required Data set (movie budget / collection in box office)

118 Regression tree code.

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import math
df = pd.read_csv("Movie-regression.csv", header=0)
df.head()
df.info()
```

```
shift cut
shift + entry current cell no.
shift + entry + next cell ready
Alt+Enter current cell no.
+ reinsert empty
shift + tab Enter
Alt+Forward slash
```

(120) df.info() Data with movie time take up 506 rows 212511 bytes.

Data after 120th missing value impute, 200 rows or mode or zero comment long story short mode.

120 df['Time Taken'].mean()

df['Time taken'].fillna(value = df['Time taken'].mean(), inplace = True)

df.info()

(121) categorical (3D, movie genre) (so numerical so transform 200 rows)

df.pop('get_dummies(df, columns = ["3D available", "Genre"], drop_first = True))

df.head()

(122) X-y split 200 rows

collection OR ~~X~~ \rightarrow all rows except collection

X = df.loc[:, df.columns != "Collection"]

X.shape()

y = df["Collection"]

y.head() / shape()

constant 0/1/1
constant 0/1/1
constant 0/1/1
constant 0/1/1
constant 0/1/1

(123) // Test-train split 200

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, testsize=0.2, random_state=0)

(25) // creating decision tree
from sklearn import tree

regtree = tree.DecisionTreeRegressor(max_depth=3)

regtree.fit(X_train, y_train)

$y\text{-train-pred} = \text{regtree}\cdot\text{predict}(x\text{train})$

$y\text{-test-pred} = u \cdot u (x\text{-test})$

(127) // evaluation

(127) from sklearn.metrics import mean_squared_error, r2_score

mean_squared_error(y-test, y-test-pred)

r2_score(y-train, y-train-pred)

r2_score(y-test, y-test-pred)

(128) How to plot decision tree //

↳ create dot file
↳ convert dot file to image
↳ use tex the image to graph

dot_data = tree.export_graphviz(regtree, out_file=None)

from IPython.display import Image

import pydotplus

graph = pydotplus.graph_from_dot_data(dot_data)

graph = pydotplus.graph_from_dot_data(dot_data)

Image(graph.create_png())

pip install pydotplus

pip install pydotplus

(129) Max depth & gdp better solution using mgo (mgo)

tree pruning not & v.

$$\sum_{i=1}^n \sum_{x_i \in R_m} (y_i - \hat{y}_{R_m})^2 + \alpha |T| \quad \left\{ \begin{array}{l} \text{Hence} \\ \alpha_2 \text{ Tuning parameter} \\ T = \text{number of leaf nodes} \end{array} \right. \quad \text{RSS}$$

Root (all x & y all) (A not fit with (x RSS 913.0))

2nd best Regression tree etc.

130

Controlling tree growth

i) Maximum number of levels in tree

regtree1 = tree.DecisionTreeRegression(max_depth=3)

regtree1.fit(X_train, y_train)

dot_data = tree.export_graphviz(regtree1, out_file=None, feature_name=X_train.columns, filled=True)

graph1 = pydotplus.graph_from_dot_data(dot_data)

Image(graph1.create_png())

ii) minimum observation at internal node

regtree2 = tree.DecisionTreeRegressor(min_samples_split=10)

... we same.

iii) minimum observations at leaf node

regtree3 = tree.DecisionTreeRegression(min_samples_leaf=25)

... we same.

if for condition to monitor apply root imp. of 2.28

regtree4 = tree.DecisionTreeRegressor(min_samples_leaf=25, max_depth=4)

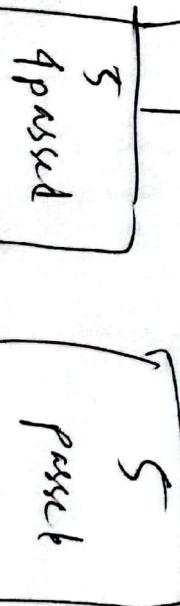
132 Classification Tree:

Regression vs mean we say 20.01 answer mode.

A² total 10 A² 9.933333

A² pass hours < 5 → total fail.

A² fail



Recursive binary splitting we do. But

22% not correctly use 25%

Classification error rate \rightarrow worst example 1 option 1 75%
Total 2 on total $\frac{3}{10} \times 100 = 30\%$
misclassified. error = 30%.

Gini index

Gini index 1 or 0 will always pure class. (Gini 0%)

(133) star_tech_oscar area 50% (30%) predict 20%.

(134) // code:

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt

df = pd.read_csv('movie-classification.csv', header=0)
df.head() /inf0()

// missing value imputation.
df['Time-taken'].fillna(value=df['Time-taken'].mean(), inplace=True)

// dummy
df = pd.get_dummies(df, columns=['3D-available', 'Genre'])
drop first = True

// Xy split
X_2 = df.loc[:, df.columns != "start_Tech_Oscar"]
y = df['start_Tech_Oscar']

// Test train split
```

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.4, random_state=0)

// training
from sklearn import tree
clf_tree = tree.DecisionTreeClassifier(max_depth=3)
clf_tree.fit(X_train, y_train)

// predicting values
y_train_pred = clf_tree.predict(X_train)
y_test_pred = clf_tree.predict(X_test)

// Model performance
from sklearn.metrics import accuracy_score, confusion_matrix
confusion_matrix(y_train, y_train_pred)
confusion_matrix(y_test, y_test_pred)
accuracy_score(y_test, y_test_pred)

// plotting decision tree
dot_data = tree.export_graphviz(clf_tree, out_file=None, filled=True,
                                feature_names=X_train.columns)

from IPython.display import Image
import pydotplus
graph = pydotplus.graph_from_dot_data(dot_data)
Image(graph.create_png())

// controlling tree growth (similarly)
clf_tree2 = tree.DecisionTreeClassifier(min_samples_leaf=20,
                                         max_depth=9)
```

(137) ad is disadvantage of decision tree

- ad is disadvantage of decision tree
- easy to explain
- more human like
- graphically shown
- dummy create
- not suitable qualitative data handle predictor handle rare events

disadvantage of predictive model
→ same level of predictive accuracy

(138) Ensemble Methods

- Bagging
- Random forest

→ Boosting

A 20% training set A 20% model use all the 20% data set B 20% test set model apply on 20% data set B 20% test set model accuracy on 20% data set.

Bootstrap → Bootstrapping → pick and add training set (20%) randomly Data pick and add same size as original data.

Bagging
→ pruning is not done, full length trees are grown
→ individual trees have high variance & low bias,
→ averaging reduces the variance.

Bagging
→ pruning is not done, full length trees are grown
→ individual trees have high variance & low bias,
→ averaging reduces the variance.
→ In regression, we take the average of predicted values.
→ In case of classification, we take majority vote i.e. most predicted class will be taken as the final prediction.

139 // Bagging code (Classification)

```
from sklearn import tree  
clf_tree = tree.DecisionTreeClassifier()  
from sklearn.ensemble import BaggingClassifier  
bag_clf = BaggingClassifier(base_estimator=clf_tree, n_estimators=1000,  
bootstrap=True, n_jobs=-1, random_state=42)
```

with
subset replacement
from bag to base
to get 10% process
multiple parallel
at best

```
bag_clf.fit(X_train, y_train)
```

```
confusion_matrix(y_test, bag_clf.predict(X_test))
```

```
accuracy_score(y_test, u^v)
```

140 // Random forest

→ improvement over bagging

→ can't use best predictor variable (use not correlated, known
Randomly ~~not~~ "select not correlated,

→ robust Predictor value → no outliers?
for Regression, $M = \sqrt{P}$

for Classification, $M = \sqrt{P}$

→ If the variables are highly correlated, smaller value of
 M should be used.

142 // Random forest Code (Classification)

```
from sklearn.ensemble import RandomForestClassifier
```

```
rf_clf = RandomForestClassifier(n_estimators=1000, n_jobs=-1,  
random_state=42)
```

```
rf_clf.fit(X_train, y_train)
```

```
confusion_matrix(y_test, rf_clf.predict(X_test))
```

```
accuracy_score(y_test, rf_clf.predict(X_test))
```

(4.3) Optimising Random Forest

from sklearn.model_selection import GridSearchCV
rf_clf = RandomForestClassifier(n_estimators=250, random_state=42)

params_grid = {
 "max_features": [1, 5, 6, 7, 8, 9, 10],
 "min_samples_split": [2, 3, 10] }

grid_search = GridSearchCV(rf_clf, params_grid, n_jobs=-1, cv=5, scoring="accuracy")
cross validation

grid_search.fit(X_train, y_train)

grid_search.best_params_ max: 7, min: 3

cvt_clf = grid_search.best_estimator_

accuracy_score(y_test, cvt_clf.predict(X_test))

(4.5) Boosting

→ each tree is grown from using information of previously grown tree.

Tree boosting

→ Residuals from fitting a linear model

→ Gradient Boost → add one more tree, ignore in previous tree.

→ Ada Boost

→ XG Boost → similar as gradient but better perform because regularization is used.

Regularization:

The cost function we are trying to optimise (MSE in regression etc) also contains a penalty term for number of variables. In a way, we want to minimize the number of variables in final model along with the MSE or accuracy. This helps in avoiding overfitting.

Q // Gradient Boosting

```
from sklearn.ensemble import GradientBoostingClassifier
```

```
gbc_clf = GradientBoostingClassifier()
```

```
gbc_clf.fit(X_train, y_train)
```

```
accuracy_score(y_test, gbc_clf.predict(X_test))
```

```
gbc_clf2 = GradientBoostingClassifier(learning_rate=0.02, max_depth=1,
```

n_estimators=1000).

```
accuracy_score(y_train, gbc_clf2.predict(X_train))
```

```
v  
v (y-test, . . . . (X-test))
```

④ // Ada Boosting

```
from sklearn.ensemble import AdaBoostClassifier
```

```
ada_clf = AdaBoostClassifier(learning_rate=0.02, n_estimators=5000)
```

```
ada_clf.fit(X_train, y_train)
```

```
accuracy_score(y_train, ada_clf.predict(X_train))
```

```
accuracy_score(y_test, . . . . (X-test))
```

```
ada_clf2 = AdaBoostClassifier(n_estimators=500, learning_rate=0.05, n_estimators=500)
```

```
ada_clf2.fit(X_train, y_train)
```

```
accuracy_score(y_test, ada_clf2.predict(X_test))
```

⑤ // XGBoost

```
first XGBoost install pip & conda (xgboost) and (5)  
[pip install xgboost]
```

```
import xgboost as xgb
```

```
xgb_clf = xgb.XGBClassifier(max_depth=5, n_estimators=10000,  
learning_rate=0.3, n_jobs=-1)
```

```
xgb_clf.fit(X_train, y_train)
```

accuracy-score(y-test, xgb_clf.predict(x-test))

xgb.plot_importance(xgb_clf)

// Grid search via grid optimise

```
xgb_clf = xgb.XGBClassifier(n_estimators=250, learning_rate=0.1,  
random_state=42)
```

```
param_test1 = {  
    'max_depth': range(3, 10, 2),
```

```
    'gamma': [0.1, 0.2, 0.3],
```

```
    'subsample': [0.8, 0.9],
```

```
    'colsample_bytree': [0.8, 0.9],
```

```
    'reg_alpha': [1e-2, 0.1, 1]
```

```
grid_search2 = GridSearchCV(xgb_clf, param_test1, cv=5, n_jobs=-1,  
scoring='accuracy')
```

grid-search.fit(X_train, y_train)

xgb_clf = grid_search2.best_estimator_

```
accuracy_score(y-test, xgb_clf.predict(x-test))
```

grid-search.best_params_

52) // Out of Box Machine learning techniques (Support Vector Machin

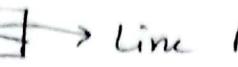
→ Maximal Margin Classifier [Support vector machine is a generalization of this classifier]

→ Support vector classifiers

↳ " " Machine

↓
classes are linearly
separable

Hyperplane :- plane that divides P dimensional space into two parts.

- For one 1D space →  point into divide
- For 2D space →  Line into divide
- " 3D space →  plane into divide

So there is a ($P-1$) plane that divide P space into two parts.

Predictor variable = 2D/3D/20D space

(14) we have to find the hyperplane that divides this spaces.

Data 2 for perfectly separable & there infinite no of hyperplane 2530.

- calculate the perpendicular distance of observations from hyperplane
- Minimum value of distance is called Margin.
- Choose the hyperplane with maximum value of margin

Support Vectors

- The observation which fall on margin are known as support vectors
- These classifiers depends on support vectors only.
- That is why this technique is different from conventional ML

(55) Limitation

- If the two classes are not separable by a hyperplane Then Maximal Margin classifier can't be used.
- very sensitive to points near to the line (support vectors)

(6) Support Vector Classifier

- soft margin classifier (Any point between margin is correct)
- $\pm \epsilon$ error allow inf. misclassified (incorrectly) correctly classified but on the wrong side.

Steps:

- we create a misclassification budget (B) to sum limit of such that
- wrong side a wrong point to distance $(x_1+x_2+x_3+\dots) < B$

→ try to maximize margin while trying to stay within budget
→ software packages like libSVM, Octo.

Impact of C

- when C is small, margins will be wide & there will be many support vectors & many misclassified observations.
- when C is large, margins will be narrow & there will be fewer support vectors & fewer misclassified values.
 - low cost value prevents overfitting & may give better loss function performance
 - optimal value of C find acc use cross val.

↓
<https://cs.stanford.edu/~karpathy/svmdemo/>

157 Limitation

- abt works when support vector classifier is linearly separable
- abt overcome not of the SVM current non linear separator (kernels)

158 Kernel Based Support Vector Machines

→ it is an extension of SVC which uses kernel to create non linear boundary

Kernels:

- g+ is some functional relationship between two observations.
- linear → $K(x_i, x_i') = \sum_{j=1}^p x_{ij} x_{i'j}$
- polynomial → $K(x_i, x_i') = (1 + \sum_{j=1}^p x_{ij} x_{i'j})^d$
- Radial → $K(x_i, x_i') = \exp(-\gamma \sum_{j=1}^p (x_{ij} - x_{i'j})^2)$

↓
Gamma defines how much influence a single training example has. The larger the gamma is, the closer other examples must be to be affected.

11 Code - 169 Regression Line in one line error

(168) // SVM based Regression Model

```
from sklearn.svm import SVR
```

```
svr = SVR(kernel='linear', C=1000)
```

```
y-test-pred = svr.predict(X-test-std)
```

```
y-train-pred = svr.predict(X-train-std)
```

from sklearn.metrics import mean_squared_error, r2_score

```
mean_squared_error(y-test, y-test-pred)
```

```
r2-score(y'-train, y-train-pred)
```

```
r2-score(y-test, y-test-pred)
```

168 - 167 → cars12.npz

(169) Standardizing data

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler().fit(X-train)
```

```
X-train-std = sc.transform(X-train)
```

```
X-test-std = sc.transform(X-test)
```

```
X-test-std
```

// SVM based Classification Model

```
from sklearn import svm
```

```
clf-svm-1 = svm.SVC(kernel='linear', C=0.01)
```

```
clf-svm-1.fit(X-train-std, y-train)
```

```
y-train-pred = clf-svm-1.predict(X-train-std)
```

```
y-test-pred = u . u (X-test-std)
```

```
from sklearn.metrics import accuracy_score, confusion-
```

```
Matrix
```

[r2 value ranges between -1 & 1. -1 = 1 + perfect fit
0 = not fit, should be 0.5~~50%~~]

Confusion matrix (y-test, y-test pred)

accuracy-score (u, u)

clf_svm_1.n_support_

{
↳ element shows 22(0, 2228)
↳ class 0 support vector, 2nd 81 2nd best}

(170) // Hyper Parameter Tuning

from sklearn.model_selection import GridSearchCV

params = { 'C': (0.001, 0.003, 0.01, 0.03, 0.1, 0.5, 1, 5, 10, 50, 100, 500, 1000) }

clf_svm_1 = svm.SVC(kernel='linear')

svm_grid_Lin = GridSearchCV(clf_svm_1, params, n_jobs=-1, cv=10,
scoring='accuracy', verbose=1)

svm_grid_Lin.fit(X_train_std, y_train)

svm_grid_Lin.best_params_

linsvm_clf = svm_grid_Lin.best_estimator_

accuracy_score(y_test, linsvm_clf.predict(X_test_std))

(171) // SVM class with polynomial kernel.

clf_svm_p3 = svm.SVC(kernel='poly', degree=2, C=0.1)

clf_svm_p3.fit(X_train_std, y_train)

y_train_pred = clf_svm_p3.predict(X_train_std)

y_test_pred = clf_svm_p3.predict(X_test_std)

accuracy_score(y_test, y_test_pred)

clf_svm_p3.n_support_

(172) // SVM class with radial

clf_svm_r = svm.SVC(kernel='rbf', gamma=0.5, C=10)

• • • • . } Same as
• - - - - . } before

clf_svm_r.n_support_

parameters = {'C': (0.01, 0.05, 0.1, 0.5, 1, 5, 10, 50), 'gamma': (0.001, 0.01, 0.1, 0.5, 1)}

clf_svm_r = svm.SVC(kernel='rbf')

svm_grid_rad = GridSearchCV(clf_svm_r, parameters, n_jobs=-1, cv=3, verbose=1, scoring='accuracy')

svm_grid_rad.fit(X_train_std, y_train)

svm_grid_rad.best_params_

~~svm_grid_rad~~ svm_clf = svm_grid_rad.best_estimator_

accuracy_score(y_test, svm_clf.predict(X_test_std))

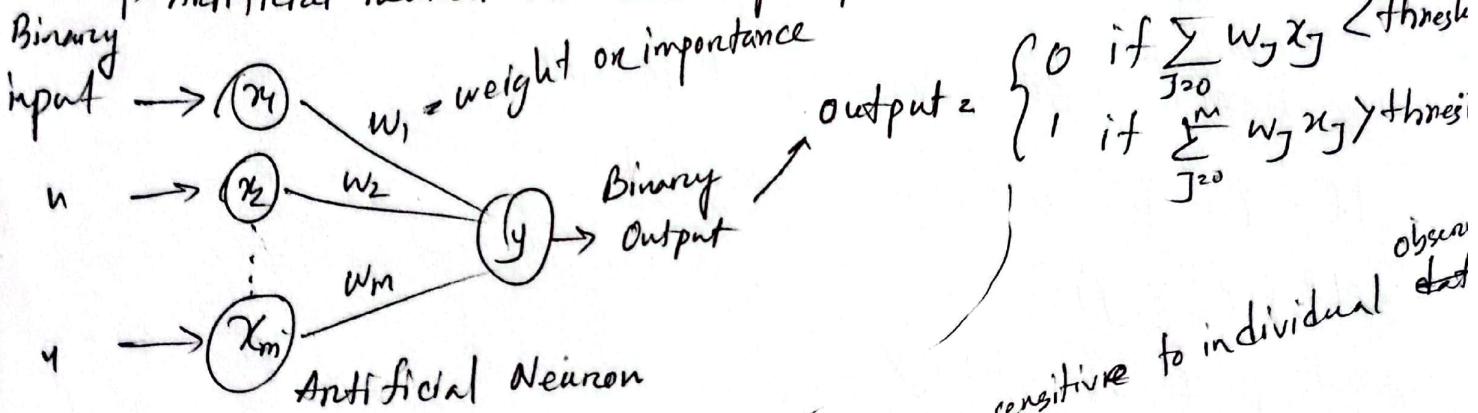
(181) Introduction to deep learning

Dataset

The dataset used here is fashion MNIST

(182) Perceptron

→ Artificial neuron are called perceptron.



(183) Activation functions

$$\text{Output} = \begin{cases} 0 & \sum_{j=0}^m w_j x_j + b < 0 \\ 1 & \sum_{j=0}^m w_j x_j + b > 0 \end{cases} \quad \text{bias is called the bias}$$

more sensitive to individual observation

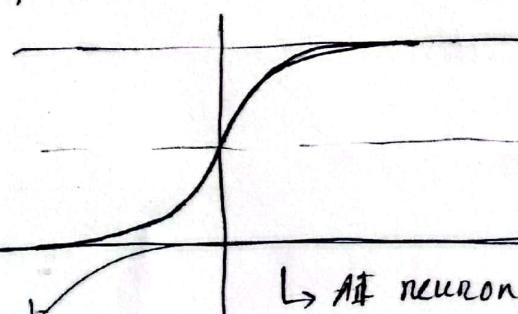
There are different kinds of activation function.

→ step

→ sigmoid

$$\text{output} = \frac{1}{1 + \exp(-\sum w_j x_j - b)}$$

$\downarrow z = \text{input}$



$$\Phi(z) = \frac{1}{1 + e^{-z}}$$

→ A neuron with sigmoid function called sigmoid or logistic neuron

184 // Creating Single Perceptron Model

if sklearn install not in

pip install -U scikit-learn

or conda install scikit-learn

```
from sklearn.datasets import load_iris
```

```
iris = load_iris()
```

// petal length & Petal width seems Depend on Iris type anticipate setosa or virginica.

```
X = iris.data[:, (2, 3)]
```

```
iris.target
```

// lets convert the target variable

```
y = (iris.target == 0) # bobl A no
```

```
y = (~y).astype(np.int) # int A no
```

```
from sklearn.linear_model import Perceptron
```

```
perc_clf = Perceptron(random_state=42)
```

```
perc_clf.fit(X, y)
```

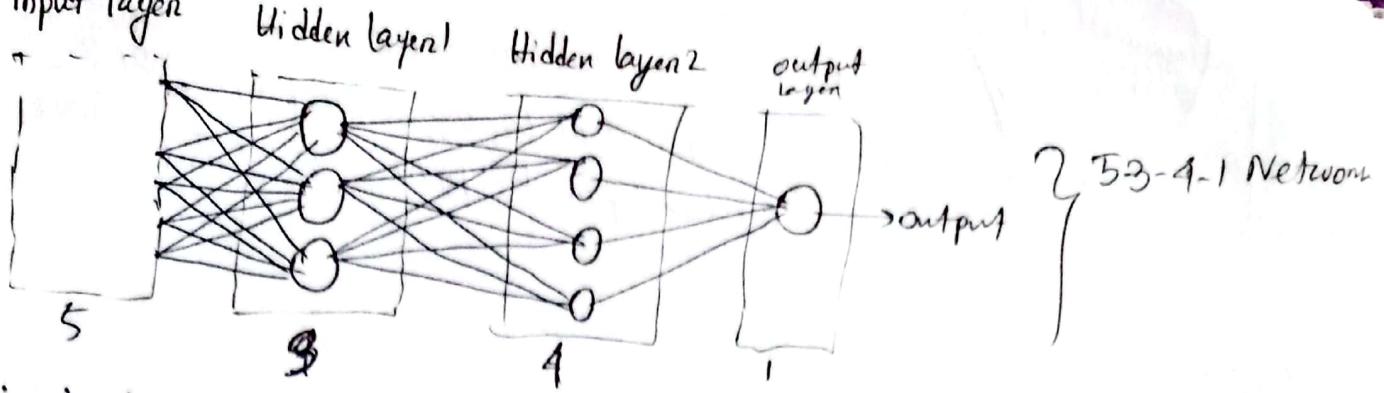
```
y_pred = perc_clf.predict(X)
```

from sklearn.metrics import accuracy_score
accuracy_score(y, y_pred)

```
perc_clf.coef_
```

```
perc_clf.intercept_
```

35 // Making networks
two ways to stack cells
→ parallel
→ sequential



5-3-4-1 Network

This is also called Feed forward Network \rightarrow one directional processing

~~if the~~ ~~the~~ layer is a perceptron ~~then~~ and perceptron ~~is~~ input and output of a cyclic Network ~~is~~: Recurrent Neural Network (RNN) is an example. (NLP 13/14)

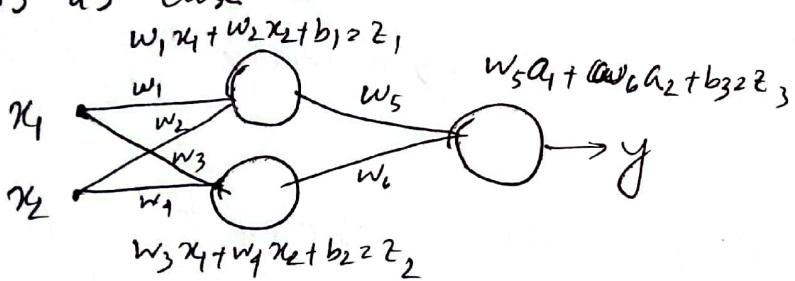
Fully connected Network - ~~as~~ layer ~~is~~ want input Next layer ~~is~~:

(186) How neural network work

$$\text{Sigmoid function, } \sigma(z) = \frac{1}{1+e^{-z}} = \frac{1}{1+\exp(-\sum w_j x_j - b)} \quad \begin{matrix} \downarrow \\ \text{weights} \end{matrix} \quad \begin{matrix} \rightarrow \\ \text{bias} \end{matrix}$$

problem statement:

- Establish the values of weights & biases so that predicted output is as close to actual output as possible.



Variables to be established
weights: w_1, \dots, w_7
biases: b_1, b_2, b_3

Gradient descent:

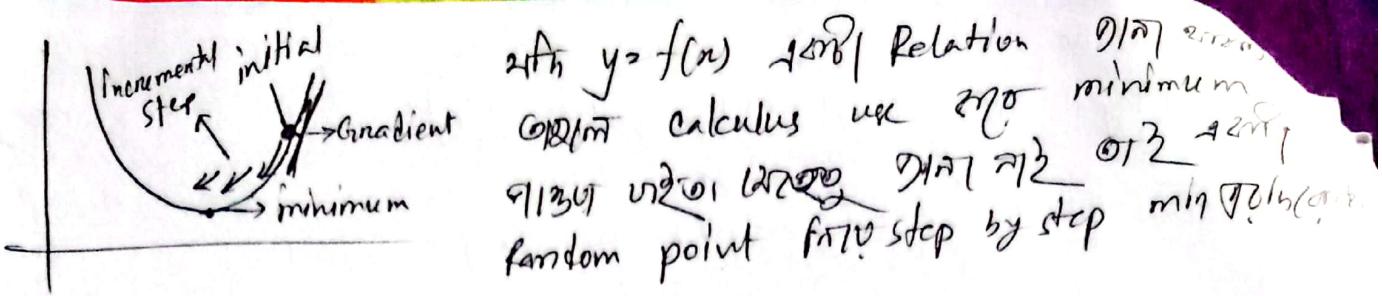
The method which is used to find the values of w_i 's & b 's.
 \rightarrow it is an optimization technique to find minimum of a function.
 \rightarrow better than OLS (Ordinary least square) as we have large no. of features & complex relationship.

Steps for gradient descent

- Assign Random w & B values [Initialization]
- calculate final output using these values [Forward propagation]
- Estimate error using error function [Backward propagation]
- find those w & B which can reduce this error
- Update w & B & repeat → [Implementation of a GD]

fig: 1





187 // Back propagation

let say,

assume predicted output = 0.3, actual output = 0

$$\text{Distance} = 0 - 0.3 = 0.3$$

$$\text{error function, } = |0.3| = 0.3$$

$$n^2 = (0.3)^2 = 0.09$$

Square function works well with regression but not with classification.

So we use, Cross Entropy Error function = $-y \log(y') - (1-y) \log(1-y')$

Here, y = actual value, y' = predicted value

This looks like fig: 1 which is necessary for gradient descent

$$\text{CEEF} = -y \log(y') - (1-y) \log(1-y')$$

if $y = 0$ then $\text{CEEF} = -(1-y) \log(1-y) \Rightarrow y'$ should be as close to 0 as possible.

$$\text{if } y = 1 \text{ then, } \text{CEEF} = -\log(y')$$

To minimize this, we have to minimize $-\log(y')$ or maximize $\log(y') \Rightarrow$ maximize y'

Since y' lies between 0 & 1. So y' should be as close to 1 as possible

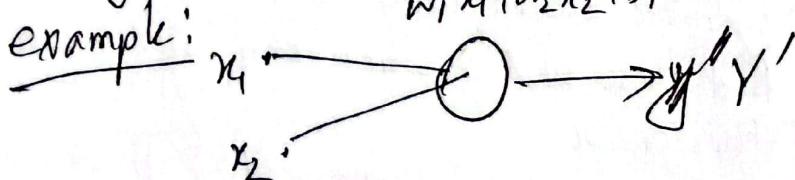
$$\text{step 1: } w \rightarrow w - \alpha \Delta w$$

$$b \rightarrow b - \alpha \Delta b$$

α is learning rate, Δw & Δb are unit steps.

α determines the number of steps we take.

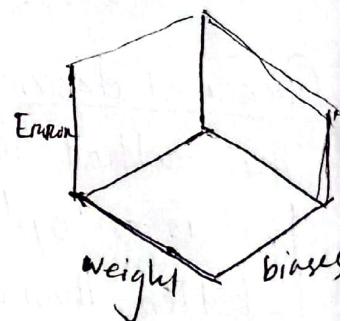
Δw & Δb from what?
through back propagation



$$\sigma(z) = \frac{1}{1+e^{-z}}$$

Error function

$$-y \log(y') - (1-y) \log(1-y')$$



$w_1 \quad w_2 \quad b$

2 3 -1 $\xrightarrow{?}$ Random

Step 2: forward propagation

example data (training data)
 x_1 x_2 $y \rightarrow$ Actual output

10 -1 1

$$z = w_1 x_1 + w_2 x_2 + b \\ z = 4, \quad \sigma(z) = 0.982$$

Step 3: Error calculation

Step 3: (Error Calculation)

$$y' \quad \frac{y}{1} \quad \left\{ \begin{array}{l} y \log(y') \\ \log(y) \end{array} \right.$$

$$E = 0.0079$$

Step 4: Back propagation

$$\frac{\partial E}{\partial y'} \quad f_{y'1} = \frac{\partial \log(y')}{\partial y'} = -\frac{1}{y'}$$

$$\frac{\partial y'}{\partial z} = \frac{\partial}{\partial z} \left(\frac{1}{1+e^{-z}} \right) = \frac{e^{-z}}{(1+e^{-z})^2}$$

$$\frac{\partial z}{\partial w_1} = x_1 = 10, \quad \frac{\partial z}{\partial w_2} = x_2 = -1, \quad \frac{\partial z}{\partial b} = 1$$

$$\text{Now, } \frac{\partial E}{\partial w_1} = \frac{\partial E}{\partial y'} \times \frac{\partial y'}{\partial z} \times \frac{\partial z}{\partial w_1} = -0.188$$

$$\text{Similarly, } \frac{\partial E}{\partial w_2}, \frac{\partial E}{\partial y'}, \frac{\partial y'}{\partial z} \times \frac{\partial z}{\partial w_2} = 0.0796$$

$$\therefore \frac{\partial E}{\partial b} = \frac{\partial E}{\partial y'} \times \frac{\partial y'}{\partial z} \times \frac{\partial z}{\partial b} = -0.018$$

Step 5: updating w & b

$$\alpha \quad \text{Rate of learning}$$

$$w_1 = w_1 - \alpha \Delta w_1 = 2 - 5 \times (-0.188) = 2.93$$

$$w_2 = w_2 - \alpha \Delta w_2 = 3 - 5 \times 0.0796 = 2.627$$

$$b = b - \alpha \Delta b = -1 - 5 \times (-0.018) = -3.907$$

Step 6: Repeat

$x_1 \quad x_2 \quad y$

10 -1 1

$$z = 2.9 \times 10 + 2.6 \times (-1) + (-3.9) = 14.7$$

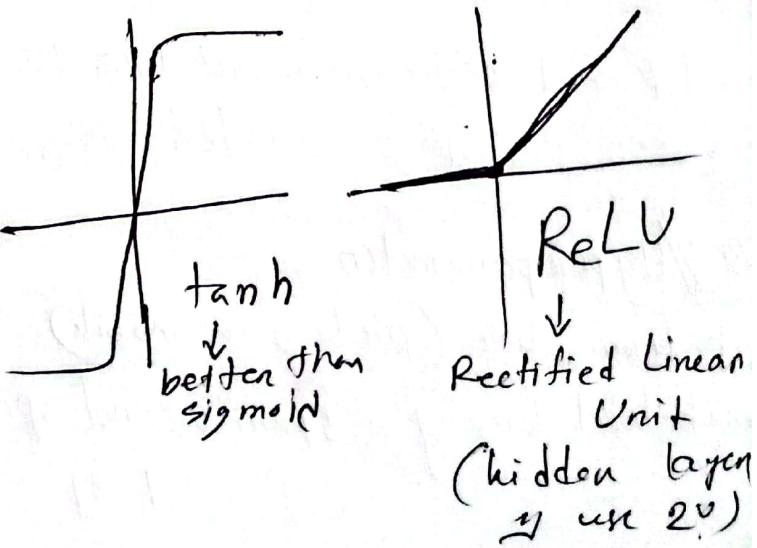
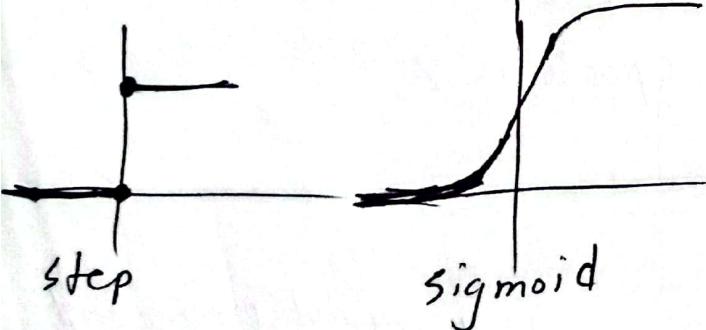
$$\sigma(z) = 0.999 = y'$$

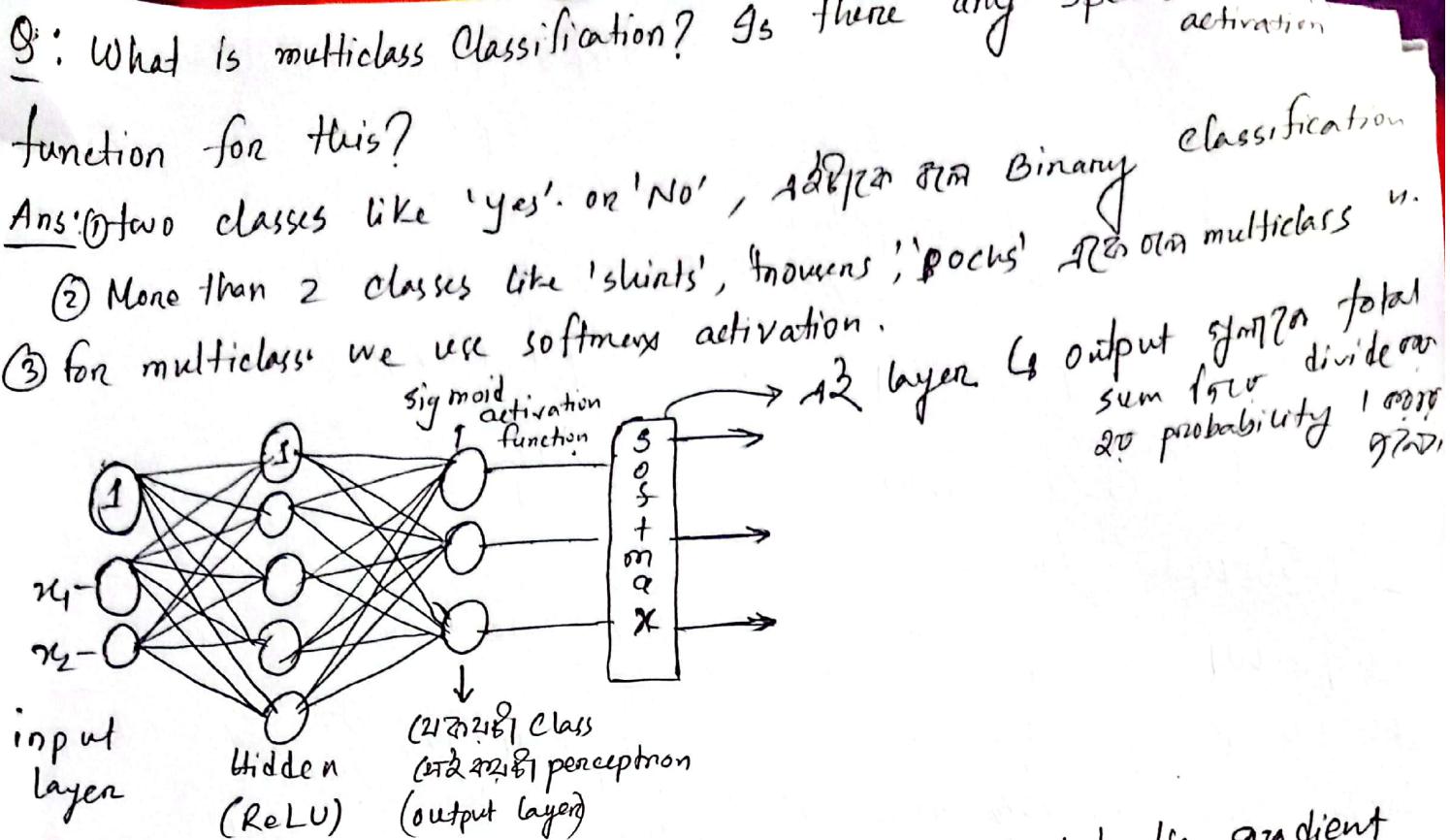
(188) // some important concepts

Q: Why do we use activation function?

Ans: to put special boundary condition, to get non linear & complex patterns. Otherwise output would be, $z = w_1 x_1 + w_2 x_2 + b$

Q: What are the different types of activation function?





Q: What is the difference between Gradient descent & stochastic gradient descent

- stochastic: → each individual training record for forward & backward propagation w, b to update fast & fast but converge slow.
- gradient: → full training set for record for OBT update slow & slow but converges well.
- Mini batch gradient: → small batch of training set for "n".

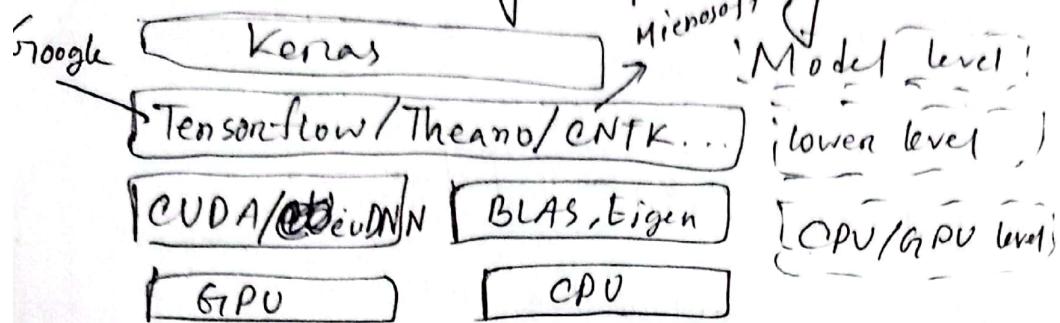
Q: What is an Epoch

- Epoch is one cycle through the full training data. Epochs mean we want the full training data to be fed 5 times. different from iteration.
- idea behind it is, to retest its performance on the same data.
- 1 Record \rightarrow input from 1000 Record \rightarrow 1000 iterations.
- 1000 Record & 1000 fed \rightarrow epoch is 2.

189 // hyperparameter

- from slide (picture in mobile)
- multilabel binary = is spam or not spam, important on not important
- 1. 87 + 2. 87 Neuron

Keras is a model level library, providing high level building blocks for developing deep learning models.



(B1) // Installing tensorflow & keras

```
import numpy as np  
import pandas as pd  
%matplotlib inline  
import matplotlib as mpl  
import matplotlib.pyplot as plt
```

```
// tensorflow install  
conda install tensorflow
```

```
conda u pip
```

```
pip install --upgrade tensorflow==2.0.0rc1
```

```
import tensorflow as tf  
from tensorflow import keras
```

[Keras. -- version --]
[tf. -- version --]

(B2) // Dataset for classification

// problem statement & dataset in the mobile. 28x28 grayscale image of pants, skirt etc (total 10 things). total 70000 dataset, 60000 train, 10000 test

```
// import dataset
```

```
fashion_mnist = keras.datasets.fashion_mnist  
(x_train_full, y_train_full), (x-test, y-test) = fashion_mnist.load_data()  
plt.imshow(x_train_full[0]) // showing first image  
y_train_full[1] // category [0..9 (0..228010)]
```

```
// category 0-9 78x72 (72x72 64x64) list create 28x28
```

class_names = ["T-shirt/top", "Trouser", "Pullover", "Dress", "Coat", "Sandal",
"Shirt", "Sneaker", "Bag", "Ankle boot"]

class_names [y_train_full[1]]

X_train_full[10] // 28x28 pixel value show 20%

⑯ // Normalization (using Gradient descent in 93) ↗ Test-train Split.

X_train_n = X_train_full / 255.0 // 0.0~1 data from 0~255 go to 0~1
X_test_n = X_test / 255.0 // 0.0~1, 255.0 max floating b/c

// training set (for our) training + validation ↗ 75% 25% (x, y) 28x28

X_valid, X_train = X_train_n[:5000], X_train_n[5000:]

y_valid, y_train = y_train_n[:5000], y_train_n[5000:]

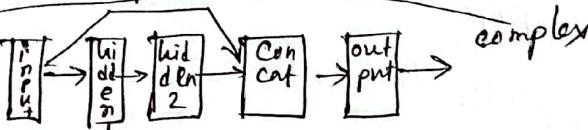
X_test = X_test_n

⑰ // Different ways to create ANN using Keras

There are two APIs for defining a model in keras

↳ Sequential model API → simple layers by layer

↳ Functional API → complex



⑲ // Building the neural network using Keras

np.random.seed(42) ↗ Same as random state

tf.random.set_seed(42)

// sequential model API

model = keras.models.Sequential()

model.add(keras.layers.Flatten(input_shape=[28, 28]))

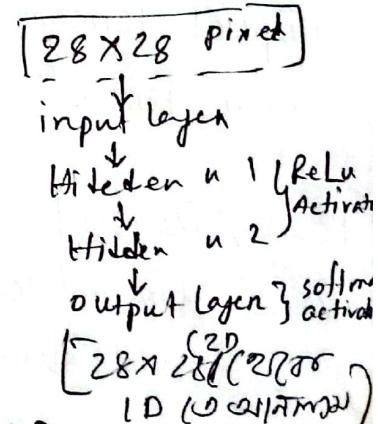
model.add(keras.layers.Dense(32, activation="relu"))

model.add(Dense(100, u = u))

model.add(Dense(10, u = u))

model.summary() ↗ // install tensorflow pip install pydot/conda install pydot

import pydot
Keras.utils.plot_model(model)



weights, biases = model.layers[1].get_weights()
weights.shape
biases
biases.shape

denonon class weights distribution over mle
stochastic gradient descent

196) // Compiling & training the neural network model

model.compile(loss="sparse_categorical_crossentropy", optimizer="sgd", metrics=[accuracy])

model_history = model.fit(X_train, y_train, epochs=30,

model_history.history

validation_data=(X_valid, y_valid))

// plotting

import pandas as pd

pd.DataFrame(model_history.history).plot(figsize=(8,5))

plt.grid(True)

plt.gca().set_ylim(0,1)

plt.show()

17) // Evaluating performance & predicting using keras

// Evaluate

model.evaluate(X_test, y_test)

X_new = X_test[:3]

round of to 2 digit

y_proba = model.predict(X_new) } probability score assigned to each class.

y_proba.round(2)

y_pred = model.predict_classes(X_new) } predicting the class

y_pred

np.array(class_names)[y_pred]

// checking if the prediction were correct or not

print(plt.imshow(X_test[0]))

print(plt.imshow(X_test[1]))

print(plt.imshow(X_test[2]))

// Building Neural Network
// Dataset that will be used is California housing using Bindep
// Objective here is to predict the price of house

```
import numpy as np
import pandas as pd
%matplotlib inline
import matplotlib as mpl
import matplotlib.pyplot as plt.
import tensorflow as tf
from tensorflow import keras
from sklearn.datasets import fetch_california_housing
housing = fetch_california_housing()
```

// Data set characteristics

Number of instances: 20640
u u attributes: 8 numeric, predictive attributes & the target

Attribute info:

- MedInc → median income in block
- HouseAge → u house age in block
- AveRooms → average no. of rooms.
- AveBedrms → " no. of bedrooms.
- Population → Block population
- AveOccup → average house occupancy
- Latitude → " block latitude
- Longitude → " " longitude

Target: the target variable is the median house value in units of 100,000 for California districts.

part 2 // code

print(housing.feature_names)

```
from sklearn.model_selection import train_test_split
x_train_full, x_test, y_train_full, y_test = train_test_split(housing.data,
                                                               housing.target,
                                                               random_state=42)
```

independent
↑ dependent

$X_train, X_valid, y_train, y_valid = train_test_split(X_train_full, y_train_full, random_state=42)$

// Standardizing the data $\rightarrow \left\{ \frac{\text{data} - \text{mean}}{\text{variance}} \cdot \text{Standardized data} \right\}$

from sklearn.preprocessing import StandardScaler

scaler = StandardScaler()

$X_train = \text{scaler}.fit_transform(X_train)$

$X_valid = \text{scaler}.transform(X_valid)$

$X_test = \text{scaler}.transform(X_test)$

np.random.seed(42)

tf.random.set_seed(42)

$X_train.shape$

model = keras.models.Sequential([

Keras.layers.Dense(30, activation="relu", input_shape=[8]),
Keras.layers.Dense(30, activation="relu"),
Keras.layers.Dense(1, activation="relu")])

])

2) Fit the model and run 20 epochs

learning Rate
default 0.01

model.summary()

model.compile(loss="mean_squared_error",
optimizer=keras.optimizers.SGD(lr=1e-3),
metrics=['mae'])

odel_history = model.fit(X_train, y_train, epochs=20, validation_data=(X_valid, y_valid))

mae_test = model.evaluate(X_test, y_test)

model_history.history

pd.DataFrame(model_history.history).plot(figsize=(8,5))

plt.grid(True)

plt.gca().set_ylim(0, 1)

plt.show()

' predict the value in new dataset

12 graph 2020 (5000 215 227 200)
converge 200 212 612 200 . 12 code
0.000 Run 2020 90 epochs 200

```
X_new = X_test[:3]
```

```
y_pred = model.predict(X_new)  
print(y_pred)  
print(y_test[:3])
```

(19) // Using functional API for complex Architectures

to delete the previous model

```
del model  
Keras.backend.clear_session()
```

```
input_ = Keras.layers.Input(shape=X_train.shape[1:])
```

```
hidden1 = Keras.layers.Dense(30, activation='relu')(input_)
```

```
hidden2 = u . u . u (30, u = u )(hidden1)
```

```
concat = u . u . concatenate([input_, hidden2])
```

```
output = u . u . Dense(1)(concat)
```

```
model = Keras.models.Model(inputs=[input_], outputs=[output])
```

```
model.summary()
```

```
model.compile(loss="mean-squared-error",  
optimizer=Keras.optimizers.SGD(lr=1e-3),  
metrics=['mae'])
```

```
model_history = model.fit(X_train, y_train, epochs=40, validation_data=  
(X_valid, y_valid))
```

```
mae_test = model.evaluate(X_test, y_test)
```

```
mse_test = u . u (u , u )
```

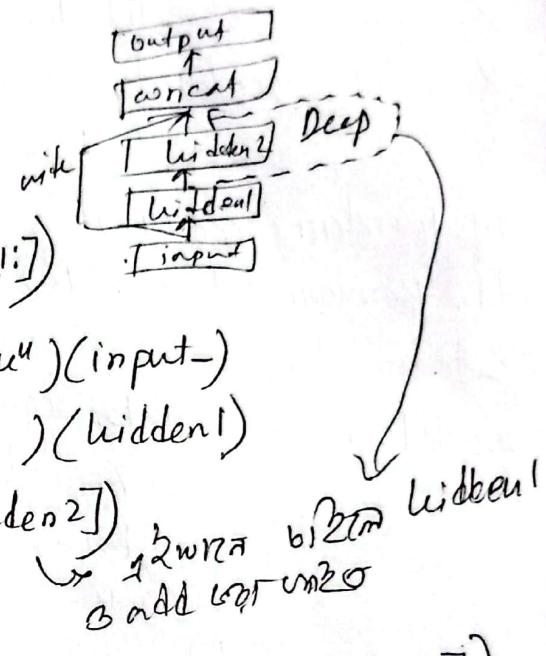
```
model_history.history
```

```
pd.DataFrame(model_history.history).plot(figsize=(8,5))
```

```
plt.grid(True)
```

```
plt.gca().set_ylim(0, 1)
```

```
plt.show()
```



Restoring Models & Using Callbacks

model.save("my_func-model.h5") → name I want to give
%pwd → to check the present working directory

%cd C:\Users\pukhr → for changing directory

del model → deleting

Keras.backend.clean_session()

model = keras.models.load_model("my_func-model.h5")

model.summary() → same model → load 200,

y-pred = ~~model.predict~~ model.predict(x)

variable file
name model-01.h5
ex: first file = model-01.h5

// Data train 200 & 20 with 10% validation for epoch 100

save every 20, or after 100 (50%)

model.compile(..., save_weights_only=True, checkpoint=True)

Checkpoint_cb = keras.callbacks.ModelCheckpoint("Model-{epoch:02d}.h5")

history = model.fit(X_train, y_train, epochs=10,
validation_data=(X_valid, y_valid),
callbacks=[checkpoint_cb])

del model

Keras.backend.clean_session()

model = keras.models.load_model("Model-10.h5")

mse_test = model.evaluate(X_test, y_test)

// ~~Model Save~~ is unnecessary. Best model by default 200

→ 10(1) 20(0) 200.

del model

Keras.backend.clean_session()

model = keras.models.Sequential([keras.layers.Dense(30, activation="relu", input_shape=[8]),
keras.layers.Dense(30, activation="relu"),
keras.layers.Dense(1)])

model.compile(loss="mse", optimizer=keras.optimizers.SGD(),
checkpoint_cb=keras.callbacks.ModelCheckpoint("Best-Model.h5",
save_best_only=True))

history = model.fit(x_train, y_train, epochs=10,
validation_data=(x_valid, y_valid),
callbacks=[checkpoint_cb])

model = keras.models.load_model("Best-Model.h5")

mse_test = model.evaluate(x_test, y_test)

// 10/20 epoch from Best result নিয়ে রাখি, But 200/300 ফল (MSE)
ক্ষেত্র ও ক্ষেত্র মাঝে পরিবর্তন, তবে best result ক্ষেত্র হবে Stop এর ফল 20,
১৮ টি গড় save best callback দ্বাৰা ২১/২১ ও ১৭/২১ এবং ১ ক্ষেত্র মাঝে পরিবর্তন
হওয়া গড় save best callback দ্বাৰা ২১/২১ ও ১৭/২১ এবং ১ ক্ষেত্র মাঝে পরিবর্তন

del model

keras.backend.clear_session()

model = keras.models.Sequential([

keras.layers.Dense(30, activation="relu", input_shape=[8]),
 keras.layers.Dense(30, activation="relu"),
 keras.layers.Dense(30, activation="relu"),
 keras.layers.Dense(1)])

model.compile(loss="mse", optimizer=keras.optimizers.SGD(lr=1e-3))
checkpoint_cb = keras.callbacks.ModelCheckpoint("early-stop-model.h5",
 save_best_only=True)

early_stopping_cb = keras.callbacks.EarlyStopping(patience=10,
 restore_best_weights=True)

history = model.fit(x_train, y_train, epochs=200,
 validation_data=(x_valid, y_valid),
 callbacks=[checkpoint_cb, early_stopping_cb])

model = keras.models.load_model("early-stop-model.h5")

mse_test = model.evaluate(x_test, y_test)

- Number of Layers
 - " " neurons per layer
 - Type of activation function
 - Learning rate
 - Weight initialization logic etc
- most of problems single layer is sufficient
But going deeper increases parameter efficiency
1st or 2nd layer for ANN model & rest MLP
transfer learning.

flexibility of ANN is also a drawback

Transfer learning: If a model which has already face recognition
ognition to use with another system design can
be used to recognize different objects like faces.
of the face recognition model to first & second layer
(and vice versa), it is called transfer learning.

Number of neurons per hidden layer:

- input & output neuron & the problem
- pyramid structure (128, 64, 32, 16, 8, 4, 2, 1 Neuron) preferred
- size of all hidden layers & same number of neurons preferred
- neuron size (eqn depth 1431 at 20 iterations)

Learning rate:

In general the optimum learning rate is half of maximum learning rate (the learning rate at which the training goes diverges). Simple approach can be:-

- start with a large value.
- then divide this value with 3 & try again
- repeat this until the training algo stops diverging

Batch size: It should not be too small. It could be less than 32 & ^{training} iteration is very fast. Small batch size ensures that each iteration takes advantage of hardware & more than 20 then it helps take advantage of hardware optimization (particular for matrix multiplication).

ch:

- Number of Layers
 - n neurons per layer
 - Type of activation function
 - Learning rate
 - Weight initialization logic etc
- most of problems single layer is sufficient.
But going deeper increases parameter efficiency
1st & 2nd layer for training model to use softmax
for transfer learning.

flexibility of ANN is also a drawback

Transfer learning: यदि कोई model कोरिए ही already face recognition
definition के लिए use करता है तो उसके लिए system design करना
जैसे Hairstyle recognize के लिए उसी model का use करता है (BERT)
जैसे यह face recognition model के first & second layer
योंके लिए इसी पर्याय, ऐसी ही & transfer learning.

Number of neurons per hidden layer:

- input & output neuron जैसा WDT का problem का है,
- कोरिए pyramid structure (जैसा 300, 100, 200, 100 Neuron) preferred.
- लेकिन all hidden layer के same number of neuron preferred.
- neuron जैसा 100 (जैसा depth 3142 तक अप्रैक्टो में 20 (लेटे,

learning rate:

In general the optimum learning rate is half of maximum learning rate (The learning rate at which the training algo diverges). Simple approach can be:-

- start with a large value.
- then divide this value with 3 & try again
- repeat this until the training algo stops diverging

Batch size:
should be less than 32 & don't be too small.
small batch size ensures that each iteration is very fast
if more than 20 then it helps take advantage of hardware & software optimization (particular for matrix multiplication).

Epoch:

(210) CNN introduction

→ better than ANN in image & speech.

→ ANN a particular pixel has no context. But human Brain particular pixel has a lot of group of pixel has more CNN is better.

arbitrary mobile a pic (গুরু)

(211) // stride

→ first neuron (or group of pixel) (or input feature filter, 7x7)

Neuron এন্টের শিফট হবে n n n n n n n n

মাত্র 2^o stride. (এ হচ্ছে 2^o shift এবং 2^o stride stride of 2). 2 pixel shift এবং stride of 2, stride 2^o (কাজে 2^o overlap হলে মাত্র এবং no. of neurons needed হবে কম।

⇒ mobile pic

(212) // padding

stride 3 filter

Large stride হলে শুধুমাত্র গুরু বিলুপ্ত পিলেজ পিল না কভার হবে এবং 16x16 image (or 5x5 এর window হিসেবে) 28 pixel column একটি (১৭৮x১৭৮), তবে solution হলো first value 1 by col. হবে last (১৭৮ - 1 by col) হবে (১৫৩), 14x14 এর তারে, 14x14 (or 5x5 window এর) stride of 3 হলে 27 কভার হবে। তাহলে ১০ padding.

মাত্র solution হলো মুদ্রণ অর্থাৎ artificially 1 by column of pixel add হবে padding, যেখানে same padding.

mobile pic for visualizing.

(213) // filter & Feature Maps

→ window size এর সাইজ dimension এর একটি matrix (for this case 5x5). মাত্র Data of value 212x212 এর 25 window এর pixel এর মান (১৫৩) এর 1 by value হিসেবে ১০, Filter layer এর ১০ এর ১০x10 এর ১০ এর 2^o.

Data এর ১০x10 এর ১০ এর 2^o feature map.

এই ক্ষেত্রে (or) output of ১০x10 এর ১০ এর 2^o feature map.

Vertical filter

0	1	0
0	1	0
0	1	0

 ১০x10 vertical line clean

Horizontal filter

0	0	0
1	1	1
0	0	0

 ১০x10 Horizontal line clean

feature map for 2×2 pixels screenshot 1.
Input layer also have multiple layer of information which are called channels.

21) // Channels

→ the input to our network when we are doing image recognition
image.
→ grayscale image / black & white → each pixel contains only one info. that is how wide the pixel is. 0-255 e.g. 0 means black. 255 means completely white. In between are shades of grey.
→ colored / RGB
↓
each pixel have 3 values rather than 1.
They are R, G, B. All value 255 is white.
 $R(255, 255, 0)$ yellow, $(0, 255, 255)$ cyan

Input layer 4×4 RGB (RGB) value for 1×1 input layer 1 info per pixel
gray scale image \rightarrow 2×2 2 P1 channel.
For P1 color \rightarrow color perceive only color pie only one example
screenshot 2. \rightarrow 4×4 image \rightarrow RGB image.

22) // Pooling Layer

→ use for computational load, memory usage & number of parameters in terms of weight.
→ 4×4 input and 1 P1 layer 2×2 (max) weight 2×2 1 way
Pooling layer \rightarrow window 2×2 select max based on {best}
→ max pooling: max pixel of that window.
→ Arg n : Arg n of u u ,
the example (screenshot) and here have a stride of 2.
feature map 4×4 & 1 pooling layer 2×2 \rightarrow 4. \rightarrow 2×2 output
It is a tradeoff. We give away some input info to reduce the computational load on our system.

Model (Preprocessing)

```
import numpy as np  
import pandas as pd  
%matplotlib inline  
import matplotlib as mpl  
import matplotlib.pyplot as plt.  
import tensorflow as tf  
from tensorflow import keras  
// fashion MNIST data use API.
```

```
fashion_mnist = keras.datasets.fashion_mnist
```

```
(X_train_full, y_train_full), (X_test, y_test) = fashion_mnist.load_data()  
class_names = ["T-shirt/top", "Trouser", "Pullover", "Dress", "Coat", "Sandal",  
    "Shirt", "Sneaker", "Bag", "Ankle boot"]
```

// Data Reshape

ANN (1D input MNIST) for CNN (3D input MNIST)
X_train_full = X_train_full.reshape((60,000, 28, 28, 1)) → Dataset
X_test = X_test.reshape((10000, 28, 28, 1)) → channel

// Data Normalization

```
X_train_n = X_train_full / 255
```

```
X_test_n = X_test / 255
```

// Splitting the data

X_valid, X_train = X_train_n[:5000], X_train_n[5000:]
y_valid, y_train = y_train_full[:5000], y_train_full[5000:]
X_test = X_test_n

// Creating model architecture

```
np.random.seed(42)
```

```
tf.random.set_seed(42)
```

training data → training 90%
test " → test 10%
Validation data → used for
tuning the hyperparameters
& evaluate the models.

(27) // Structure & Compile

ANN ~~has~~ flatten cars input to 2D

Pooling Layer add 2D ~~avg~~

$(13 \times 13 \times 32)$ pooling $\rightarrow 13 \times 13 \text{ max}$

model = keras.models.Sequential()

model.add(keras.layers.Conv2D(filters=32, kernel_size=(3, 3), strides=1, padding='valid', activation='relu', input_shape=(28, 28, 3)))

model.add(keras.layers.MaxPooling2D((2, 2)))

// 3D flatten 2D 1D 28×28

model.add(keras.layers.Flatten())

model.add(Dense(300, activation='relu'))

model.add(Dense(100, activation='relu'))

model.add(Dense(10, activation='softmax'))

model.summary() multilabel classification problem 128×128

// compiling

model.compile(loss='sparse_categorical_crossentropy',
optimizer='sgd', metrics=['accuracy'])

(28) // Training & results

model_history = model.fit(x_train, y_train, epochs=30, batch_size=64,
validation_data=(x_valid, y_valid))

import pandas as pd

pd.DataFrame(model_history.history).plot(figsize=(8, 5))

plt.grid(True)

plt.gca().set_ylim(0, 1)

plt.show()

ev = model.evaluate(x_test_n, y_test)

ev

X_new = X_test[:3]

y_pred = model.predict_classes(X_new)

y_pred

y_test[:3]

print(plt.imshow(X_test[0].reshape(28, 28)))

Keras backend.clear_session

del model.

(219) // Comparison - Pooling vs without Pooling

→ same after coding with & without pooling layer. No comparison b/w w/o
pooling use regular PReLU and maxout. But accuracy is 20% less.

(220) // Project (Creating CNN from Scratch)

→ classify colored images of cats & dogs (Data from kaggle)

→ It is a Binary classification problem

→ colored images (So 3 channel)

→ No standard dimension

→ Here they are using subset of 50000 kaggle data

Data inside the folder is order in a particular manner (screenshort).

Process:

→ CNN with small dataset & of convo network & pooling

→ u u data Augmentation → is the process of creating artificial datasets for small datasets.

→ Transfer learning

like zooming in/out
cropping, rotating etc

transferring
two images

(228) // Project - preprocessing

import numpy as np

import pandas as pd

import matplotlib inline

import matplotlib as plt

import matplotlib.pyplot as plt

import os

import tensorflow as tf

from u import keras

// Saving the directories, first a warning line to tell forward slash
train_dir = r'C:\Users\putih\re\y\z\cats-dogs small\train'
validation_dir = r'C:\....\validation'
test_dir = r'C:\....\test'
<https://veras.io/preprocessing/images>

//Data preprocessing steps

- read the J picture files
 - Decode the Jpeg content to RBG grids of pixels
 - convert these into floating point tensors.
 - Rescale the pixel values (between 0 & 255) to the [0,1] interval.
 - ~~12.5.8.5.11 step 3(a) of 81 module Q1CE, keras ImageDataGenerator~~
from tensorflow.keras.preprocessing.image import ImageDataGenerator

//Generating batches of tensor image data

```
train_datagen = ImageDataGenerator(rescale=1./255)
```

train_datagen = ImageDataGenerator(rescale=1.0/255)
test_datagen = Image Data Generator

```
test_datagen = ImageDataGenerator()  
train_generator = train_datagen.flow_from_directory(  
    'dataset/training_set',  
    target_size=(150, 150),  
    batch_size=32,  
    class_mode='binary')
```

↓
output of this will
be in the form of x, y.
y labels will depend on
the name of the
folder. In this case
cats & dogs.

train_dir,
target_size = (150, 150),
batch_size = 20,
class_mode = 'binary'
Validation_generator = test_datagen.flow_from_directory(
validation_dir, target_size = (150, 150), batch_size = 20, class_mode = 'binary')

229 // Project-training

```
from tensorflow.keras import layers
```

model = model. sequential (

```

model = model. sequential()
model.add(layers.Conv2D(32, (3,3), activation='relu', input_shape=(150,100,3)))
model.add(layers.MaxPooling2D((2,2)))
model.add(layers.Conv2D(64, (3,3), activation='relu'))
model.add(layers.MaxPooling2D((2,2)))
model.add(layers.Conv2D(128, (3,3), activation='relu')) 128 Ftrs, same conv as  
total 4 flr layer.
same
same
same
same
same

```

model. add (layers, flatten)()

model.add(Dense(512, activation='sigmoid'))

model.add(Dense(1, activation='sigmoid'))

from tensorflow.keras import optimizers
 model.compile (loss='binary_crossentropy',
 SGD to Cuda Better
 optimizer=optimizers.RMSprop (lr=1e-4),
 in case of image processing.
 metrics=['acc'])
 history = model.fit_generator (
 history just fit use GPU. But it was
 train_generator, steps_per_epoch=100, epochs=20,
 validation_data=validation_generator, validation_steps=50)

~~pd. Dataframe (history, history). plot (figsize=(8,5))~~

~~plt. grid (True)~~

~~plt. gca(). set_ylim(0,1)~~

~~plt. show()~~

~~model.save("model.h5")~~

85120 (7244841200)
 overfit 210 (51151000)
 epoch 7137 training
 accu 913% 0% validation
 accu 913% 0% validation

(222020 for 22)
 total image no.
 batch size
 validation image no.
 batch size

~~from tensorflow.keras import backend as K~~

~~K.clear_session()~~

~~del model.~~

Q37) // Project - Data augmentation preprocessing

~~----- OUTSTANDING~~

train_datagen = ImageDataGenerator (

rescale=1./255,

rotation_range=90,

width_shift_range=0.2,

height_shift_range=0.2,

shear_range=0.2,

horizontal_flip=True)

test_datagen = ImageDataGenerator (rescale=1./255)

train_generator = train_datagen.flow_from_directory (

train_dir, target_size=(150, 150),

batch_size=32, class_mode='binary')

```
validation_generator = test_datagen.flow_from_directory(  
    validation_dir, target_size=(150, 150),  
    batch_size=32, class_mode='binary')
```

(238) // Data Augmentation - training & results

```
model = models.Sequential()
```

```
4 by conv layer
```

```
4 by pooling layer
```

```
flatten layer
```

```
model.add(layers.Dropout(0.5))
```

```
1 by Dense layer relu 80
```

```
1 by output layer Sigmoid 80.
```

```
model.compile(loss='binary_crossentropy',
```

```
optimizer=optimizers.RMSprop(lr=1e-4),
```

```
metrics=['acc'])
```

each epoch → 50% de activate neurons
very effective for avoiding overfitting

// training

```
history = model.fit_generator(train_generator, steps_per_epoch=100,  
epochs=100, validation_data=validation_generator,  
validation_steps=50)
```

```
model.save("project-cnn.h5")
```

```
d = Dataframe(history.history).plot(figsize=(8,5))
```

```
plt.grid(True)
```

```
plt.gca().set_ylim(0,1)
```

```
plt.show()
```

What are CNN? After CNN model structure comes out from graph
Data to use more than 1000, 1000 and 800 transfer learning.
Q2 Architecture of competition (2012 2013 2014) for
ImageNet Large Scale Visual Recognition Challenge (ILSVRC).
Different kind of architecture like LeNet (1998), AlexNet (2012),
ZFNet (2013), GoogleNet (2014), VGGNet (2014), ResNet (2015).
More data in the screenshot.

240) // LeNet

↳ simple archi with 3 convo layer & 2 Avg Pooling layer.
More about this in the screenshot.

241) // VGG

→ 138 million param
→ convo layers use 3x3 convo block use size 2x2, 3x3 2x2
→ ReLU use size 2x2 or softmax.

More about this in the screenshot.

242) // Google Net

→ A new concept use first 224x224, 256x256 or 224x inception m
→ very complex network.
More info about this in the screenshot.

243) // Transfer learning

→ Mostly we use the convolution base of other network.
→ If you want a new classifier add new output
model else no, But convolution base can be used
and it is used for Data Preprocess and feature
of 12 feature size 32x32 128x128
same feature size 32x32 128x128.

Q4) Project - Transfer learning

↳ VGG16 weights & to download & to apply `apply` `weights`.

from tensorflow. keras.preprocessing.image import `ImageDataGenerator`

```
train_datagen = ImageDataGenerator(  
    rescale=1/255, rotation_range=40, width_shift_range=0.2,  
    height_shift_range=0.2, shear_range=0.2, zoom_range=0.2,  
    horizontal_flip=True)  
  
test_datagen = ImageDataGenerator(rescale=1/255)
```

```
train_generator = train_datagen.flow_from_directory(  
    train_dir, target_size=(150, 150), batch_size=20, class_mode='binary')  
  
validation_generator = test_datagen.flow_from_directory(  
    validation_dir, target_size=(150, 150), batch_size=20, class_mode='binary')
```

```
from tensorflow.keras.applications import VGG16  
  
conv_base = VGG16(weights='imagenet', include_top=False, input_shape=(150, 150, 3))  
  
conv_base.summary()  
  
from tensorflow.keras import models  
u u u o u u
```

```
model = models.Sequential()  
model.add(conv_base)  
model.add(layers.Flatten())  
model.add(Dense(256, activation='relu'))  
model.add(Dense(1, activation='sigmoid'))  
  
model.summary()  
  
[conv_base.trainable=False]  
// compiling
```

from tensorflow.keras import optimizers
model.compile(loss='binary_crossentropy',
optimizer=optimizers.RMSprop(lr=2e-5), metrics=['acc'])

→ `lr` of `RMSprop` is 2×10^{-5}

→ `acc` of `accuracy`

→ `val_acc` of `val_accuracy`

```
history = model.fit_generator(
```

```
    train_generator,  
    steps_per_epoch=100,  
    epochs=30, validation_data=validation_generator,  
    validation_steps=50,  
    callbacks=[checkpoint_callback])
```

```
pd.DataFrame(history.history).plot(figsize=(8,5))
```

```
plt.grid(True)
```

```
plt.gca().set_ylim(0,1)
```

```
plt.show()
```

```
hist_df = pd.DataFrame(history.history) # saving for  
hist_csv_file = 'history.csv'  
hist_df.to_csv(hist_csv_file, mode='w') as f:  
    with open('list-csv-file', mode='w') as f:  
        list_df.to_csv(f)
```

```
test_generator = test_datagen.flow_from_directory(  
    test_dir,
```

```
target_size=(150,150),
```

```
batch_size=20,
```

```
class_mode='binary')
```

total image
batch size
↑

```
model.evaluate_generator(test_generator, steps=200)
```

→ 200 Directory over for 012 evaluate
generator against of evaluate use own