
CSL7020: Machine Learning-1

Semester II, 2022-202

Programming Assignment -1

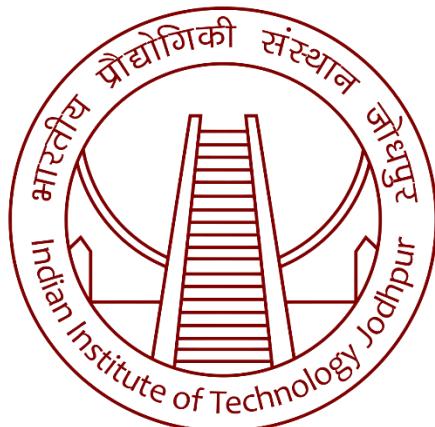
Submitted by

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Master of Technology

In

Data and Computational Science



॥ त्वं ज्ञानमयो विज्ञानमयोऽसि ॥

Under Guidance of

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Colab Link for 1st Question:

https://colab.research.google.com/drive/1p4p_mJh9B3bxQqF2YupN_kg7EEu1uUW1?usp=sharing

Colab Link for 2nd Question:

<https://colab.research.google.com/drive/15GeiHH6Z7a5C1v37xIC1Zva2CEfavy4f?usp=sharing>

Colab Link for 3rd Question:

https://colab.research.google.com/drive/1y5TPz_pK0RPaSyxlfBCupkX7oNH2EFbt?usp=sharing

GitHub Link

<https://github.com/KATHIJAAFROSE-A/MachineLearning-2.git>



Problem-1.

Perception Model.

Following sample data are given.

x_1	x_2	class.
1	1	+1
-1	-1	-1
0	0.5	-1
0.1	0.5	-1
0.2	0.2	+1
0.9	0.5	+1

Assuming Weight Vector of Initial decision Boundary

$$W^T x = 0, \text{ as } W = [1, 1].$$

Solve the following * In how many steps
perception learning algorithm will converge.

Step 1:
To Remove the outliers / Reduce the computational power.

Converting the given data points into.

'Mean Centered Data points'

Arithmetic Mean of $x_1 \Rightarrow \bar{x}_1$

$$\Rightarrow \frac{1-1+0+0.1+0.2+0.9}{6} \xrightarrow{0.2}$$

Arithmetic Mean of $x_2 \Rightarrow \bar{x}_2$

$$\Rightarrow \frac{1-1+0.5+0.5+0.2+0.5}{6} = 0.2833$$

Mean centered Data points are.

$x_1 = x_1 - \bar{x}_1$	$y_1 = x_2 - \bar{x}_2$	Class.
0.8	0.7167	+1.
-1.2	-1.2833	-1
-0.2	0.2167	-1
-0.1	0.2167	-1
0	-0.0833	+1.
0.7	0.2167	+1.

perceptron learning Algorithm.

Infinite Convergence.

Do

for $x \in P \cup N \{$

Where

P → Set of positive samples ($y = 1$)

N → Set of Negative sample ($y = -1$)

Randomly Initializing Weights

$$w \rightarrow [w_0, w_1, w_2, \dots, w_n]$$

While ! convergence.

Do

for $x \in P \cup N$ {

If $x \in P$ and $\sum_{i=0}^n w_i x_i < 0$ then

// positive misclassified as Negative)

$$w = w + x.$$

If $x \in N$ and $\sum_{i=0}^n w_i x_i \geq 0$ then

// Negative misclassified as positive).

$$w = w - x.$$

}

Done.

Linear Algebraic Interpretation :-

$$\sum_{i=1}^n w_i x_i = w^T x.$$

Where

$$w = [1 \ w_1 \ w_2 \ \dots \ w_n]$$

$$x = [1 \ x_1 \ x_2 \ \dots \ x_n]$$

Decision Boundary $\Rightarrow w^T x = 0$

Iteration - 1 :-

Given Weight Vector $\Rightarrow [1, 1]$

Appending bias value initialization
to weight vector.

Now weight vector becomes $w = [1 \ 1 \ 0]$

(Assumed bias value as '0'
Initially)

With the initial weights assumed, check the data points
are correctly classified.

Data 1: Actual class: +1.

$\therefore w^T x$ must be greater than 0 for
correct classification.

$$w^T x = (1 \ 1 \ 0) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

(Note: Here we appended
'1' to data points
Since the algorithm
Initial Assumption
was taken like that.)

$$\rightarrow 1 \cdot 5167 > 0 \therefore \text{class } +1 \text{ predicted.}$$

$$y_{\text{actual}} = y_{\text{predicted}}$$

Data 2: Actual class: -1

$\therefore w^T x$ must be less than 0 for correct
classification.

$$w^T x = (1 \ 1 \ 0) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix}$$

$$\Rightarrow -2.4833 < 0$$

Class predicted is '-1'

$y_{\text{actual}} = y_{\text{predicted}}$. (No update in weight required).

Data 3: Actual class '-1'

$w^T x$ must be less than 0 for correct classification

$$w^T x = (1 \ 1 \ 0) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow 0.0167 \neq 0$ (Negative misclassified as positive).

Here we need to update the weight and move to next iteration.

Here Negatively misclassified as positive.

Formula for weight updation.

$$w_{\text{new}} = w - x$$

$$w_{\text{new}} = \begin{pmatrix} 1 \\ 1 \\ 0 \end{pmatrix} - \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix} = \begin{pmatrix} 1.2 \\ 0.7833 \\ -1 \end{pmatrix}$$

Iteration-2

updated weight $\Rightarrow \begin{pmatrix} 1.2 \\ 0.7833 \\ -1 \end{pmatrix}$

Data: 1: Actual class '+1'

$w^T x$ must be greater than 0 for correct classification.

$$w^T x = (1.2 \quad 0.7833 \quad -1) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$$\Rightarrow 0.52 > 0$$

Class predicted is '+1'.

$y_{actual} = y_{predicted}$.

Data 2: Actual class '-1'

$w^T x$ must be less than 0 for correct classification.

$$w^T x = (1.2 \quad 0.7833 \quad -1) \begin{pmatrix} -0.2 \\ 0.2833 \\ 1 \end{pmatrix}$$

$$\Rightarrow -3.445 < 0$$

Class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 3: Actual class is '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = (1.2 \quad 0.7833 \quad -1) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$$\Rightarrow -1.07 < 0$$

class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 4: Actual class is '-1'

$w^T x$ must be less than '0' for correct classification

$$w^T x = (1.2 \quad 0.7833 \quad -1) \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix}$$

$$\Rightarrow -0.95 < 0. \text{ class predicted is '-1'}$$

$y_{actual} = y_{predicted}$

Data 5: Actual class is '+1'

$w^T x$ must be greater than '0' for correct classification

$$w^T x = (1.2 \quad 0.7833 \quad -1) \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

positive misclassified
as 'Negative'

$$\Rightarrow -1.065 < 0. \text{ class predicted is '+1'}$$

$y_{actual} \neq y_{predicted}$.

Here we need to update the weight and move to next iterations.

Here positively misclassified as Negative.
formula for weight update.

$$W_{\text{new}} = W + x$$

$$W_{\text{new}} = \begin{pmatrix} 1.2 \\ 0.7833 \\ -1 \end{pmatrix} + \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$$W_{\text{new}} = \begin{pmatrix} 1.2 \\ 0.7 \\ 0 \end{pmatrix}$$

Iteration-3

Data-1: Actual class '+1'

$W^T x$ must be greater than 0 for correct classification

$$W^T x = (1.2 \quad 0.7 \quad 0) \begin{pmatrix} 0.8 \\ +0.0833 \\ 0.7167 \\ 1 \end{pmatrix}$$

$$\Rightarrow 1.46 > 0$$

Class predicted is '+1'

$$Y_{\text{actual}} = Y_{\text{predicted}}$$

Data 2: Actual class '-1'

$w^T x$ must be less than 0 for correct classification

$$w^T x = (1.2 \quad 0.7 \quad 0) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix}$$

$\Rightarrow -1.28 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 3: Actual class '-1'

$w^T x$ must be less than 0 for correct classification.

$$w^T x = (1.2 \quad 0.7 \quad 0) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.088 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 4: Actual class '+1'

$w^T x$ must be less than 0 for correct classification.

$$w^T x = (1.2 \quad 0.7 \quad 0) \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow 0.03169 > 0$

class predicted is '+1'

$y_{actual} \neq y_{predicted}$.

Negative misclassified as positive.

Here we need to update the weight and move to next iterations;

here negative misclassified as positive.

formula for weight update.

$$W_{\text{new}} = W - x$$

$$W_{\text{new}} = \begin{pmatrix} 1.2 \\ 0.7 \\ 0 \end{pmatrix} - \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix} = \begin{pmatrix} 1.3 \\ 0.483 \\ -1 \end{pmatrix}$$

Iteration-4

Data 1: Actual class '+1'

$W^T x$ must be greater than '0' for correct classification

$$W^T x = (1.3 \ 0.483 \ -1) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$\Rightarrow 0.386 > 0$ class predicted is '+1'

$y_{\text{actual}} = y_{\text{predicted}}$.

Data 2: Actual class '-1'

$W^T x$ must be less than '0' for correct classification.

$$W^T x = (1.3 \ 0.483 \ -1) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix}$$

$\Rightarrow -3.17 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 3: Actual class '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = (1.3 \ 0.483 \ -1) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow -1.155 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 4: Actual class '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = (1.3 \ 0.483 \ -1) \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow -1.025 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 5: Actual class '+1'

$w^T x$ must be greater than '0' for correct classification.

$$w^T x = (1.3 \ 0.483 \ -1) \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$$\Rightarrow -1.04 < 0$$

class predicted is '-1'

$y_{actual} \neq y_{predicted}$.

here we need to update the weight and move to next iterations

here positively misclassified as Negative.

formula for weight update.

$$W_{new} = W + x.$$

$$W_{new} = \begin{pmatrix} 1.3 \\ 0.483 \\ -1 \end{pmatrix} + \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$$W_{new} = \boxed{\begin{pmatrix} 1.3 \\ 0.4 \\ 0 \end{pmatrix}}$$

Iteration-5:

Data 1: actual class '+1'

$w^T x$ must be greater than '0' for correct classification.

$$w^T x = (1.3 \ 0.4 \ 0) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$\Rightarrow 1.32 > 0$ class predicted is '+1'

$y_{actual} = y_{predicted}$.

Data 2: Actual class '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = (1.3 \quad 0.4 \quad 0) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix}$$

$\Rightarrow -2.07 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 3: Actual class '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = (1.3 \quad 0.4 \quad 0) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.17 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 4: Actual class '-1'

$w^T x$ must be less than '0' for correct classification

$$w^T x = (1.3 \quad 0.4 \quad 0) \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix} = -0.043 < 0$$

class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 5: Actual class '+1'

$W^T x$ must be greater than 0 for correct classification.

$$W^T x = \begin{pmatrix} 1.3 & 0.4 & 0 \end{pmatrix} \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.0332 < 0$ class predicted is '-1'

$y_{\text{actual}} \neq y_{\text{predicted}}$.

here we need to update the weight and move to next iteration.

here positive misclassified as Negative.

formula for weight update.

$$W_{\text{new}} = W + x$$

$$W_{\text{new}} = \begin{pmatrix} 1.3 \\ 0.4 \\ 0 \end{pmatrix} + \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$$W_{\text{new}} = \begin{pmatrix} 1.3 \\ 0.3167 \\ 1 \end{pmatrix}$$

Iteration-6:

Data 1: Actual class '+1'

$W^T X$ must be greater than 0 for correct classification.

$$W^T X = (1.3 \quad 0.3167 \quad 1) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$\Rightarrow 2.26 > 0$ class predicted is '+1'

$y_{actual} = y_{predicted}$.

Data 2: Actual class '-1'

$W^T X$ must be less than 0 for correct classification

$$W^T X = (1.3 \quad 0.3167 \quad 1) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.966 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 3: Actual class '-1'

$W^T X$ must be less than 0 for correct classification.

$$W^T X = (1.3 \quad 0.3167 \quad 1) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow 0.808 > 0$ class predicted is '-1'

$y_{actual} \neq y_{predicted}$.

here we need to update the weight and move to next iterations:

More Negative misclassified as positive.

formula for weight update.

$$W_{\text{new}} = W - \alpha$$

$$W_{\text{new}} = \begin{pmatrix} 1.3 \\ 0.3167 \\ 1 \end{pmatrix} - \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix} = \begin{pmatrix} 1.5 \\ 0.1 \\ 0 \end{pmatrix}$$

Iteration-7

Data 1: Actual class '+1'

$W^T X$ must be greater than 0 for correct classification.

$$W^T X = (1.5 \ 0.1 \ 0) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$\Rightarrow 1.27 > 0$. Class predicted is '+1'

$y_{\text{actual}} = y_{\text{predicted}}$

Data 2: Actual class '-1'

$W^T X$ must be less than 0 for correct classification

$$W^T X = (1.5 \ 0.1 \ 0) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix} = -1.92 < 0$$

Class predicted is '-1'. $y_{\text{actual}} = y_{\text{predicted}}$

Data 3: Actual class '-1'

$w^T x$ must be less than zero for correct classification.

$$w^T x = \begin{pmatrix} 1.5 & 0.1 & 0 \end{pmatrix} \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.27 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 4: Actual class '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = \begin{pmatrix} 1.5 & 0.1 & 0 \end{pmatrix} \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.128 < 0$ class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 5: Actual class '+1'

$w^T x$ must be greater than '0' for correct classification.

$$w^T x = \begin{pmatrix} 1.5 & 0.1 & 0 \end{pmatrix} \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$\Rightarrow -0.0833 < 0$ class predicted is '+1'

$y_{actual} \neq y_{predicted}$.

Here we need to update weight and move to next iterations.

Here positively misclassified as Negative.

formula for weight update.

$$W_{\text{new}} = W_1 + \alpha$$

$$\text{updated weight} = \begin{pmatrix} 1.5 \\ 0.1 \\ 0 \end{pmatrix} + \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$$\rightarrow \begin{pmatrix} 1.5 \\ 0.0167 \\ 1 \end{pmatrix}$$

Iteration-8

Data 1: Actual class '+1'

$w^T x$ must be greater than '0' for correct classification.

$$w^T x = (1.5 \quad 0.0167 \quad 1) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$\Rightarrow 2.2 > 0$ class predicted is '+1'

$y_{\text{actual}} = y_{\text{predicted}}$.

Data 2: Actual class '-1'

$w^T x$ must be less than '0' for correct classification.

$$w^T x = (1.5 \quad 0.0167 \quad 1) \begin{pmatrix} -0.2 \\ -1.2833 \\ 1 \end{pmatrix} = -0.82 < 0$$

class predicted is '-1' $y_{\text{actual}} = y_{\text{predicted}}$

Data 3: Actual class '-1'

$w^T x$ must be less than '0' for correct classification

$$w^T x = (1.5 \ 0.0167 \ 1) \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$$= 0.703 > 0$$

class predicted is '+1'

$y_{actual} \neq y_{predicted}$.

Here we need to update weight, and move to next iteration.

Here negative misclassified as positive.

formula for weight update

$$W_{new} = W - x.$$

$$\text{Updated weight} = \begin{pmatrix} 1.5 \\ 0.0167 \\ 1 \end{pmatrix} - \begin{pmatrix} -0.2 \\ 0.2167 \\ 1 \end{pmatrix}$$

$$\text{updated weight, } \begin{pmatrix} \cancel{-0.2} \\ \cancel{0.2167} \\ 1 \end{pmatrix} \quad \begin{pmatrix} 1.7 \\ -0.2 \\ 0 \end{pmatrix}$$

Final Iteration.

Data 1: Actual class '+1'

$W^T X$ must be greater than 0 for correct classification.

$$W^T X = (1.7 \quad -0.2 \quad 0) \begin{pmatrix} 0.8 \\ 0.7167 \\ 1 \end{pmatrix}$$

$$\Rightarrow 1.216 > 0 \text{ class predicted is '+1'}$$

$$Y_{\text{actual}} = Y_{\text{predicted}}$$

Data 2: Actual class '-1'

$W^T X$ must be less than 0 for correct classification

$$W^T X = (1.7 \quad -0.2 \quad 0) \begin{pmatrix} -1.2 \\ -1.2833 \\ 1 \end{pmatrix}$$

$$\Rightarrow -1.78 < 0 \text{ class predicted is '-1'}$$

$$Y_{\text{actual}} = Y_{\text{predicted}}$$

Data 3: Actual class '-1'

$W^T X$ must be less than 0 for correct classification

$$W^T X = (1.7 \quad -0.2 \quad 0) \begin{pmatrix} -0.2 \\ +0.2868 \\ 1 \end{pmatrix}$$

$$= -0.38 < 0$$

class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 4: Actual class '-1'

$w^T x$ must be less than 0 for correct classification

$$w^T x = (1.7 \ -0.2 \ 0) \begin{pmatrix} -0.1 \\ 0.2167 \\ 1 \end{pmatrix}$$

$$\Rightarrow -0.213 < 0.$$

class predicted is '-1'

$y_{actual} = y_{predicted}$.

Data 5: Actual class '+1'

$w^T x$ must be greater than 0 for correct classification

$$w^T x = (1.7 \ -0.2 \ 0) \begin{pmatrix} 0 \\ -0.0833 \\ 1 \end{pmatrix}$$

$$\Rightarrow 0.01666 > 0$$

class predicted is '+1'

$y_{actual} = y_{predicted}$.

Data 6: Actual class '+1'

$w^T x$ must be greater than '0' for correct classification.

$$w^T x = (1.7 \ -0.2 \ 0) \begin{pmatrix} 0.7 \\ 0.2167 \\ 1 \end{pmatrix}$$

$$\Rightarrow 1.146 > 0$$

class predicted is '+1'

$y_{actual} = y_{predicted}$.

In the final Iteration all the datapoints are correctly classified.

$$\therefore \text{optimal weight} = \begin{bmatrix} 1.7 \\ -0.2 \\ 0 \end{bmatrix}$$

2) Final decision Boundary.

$$\text{Optimal weight} = [w_1 \ w_2 \ w_0]$$

$$\text{Slope} = -w_1 / w_2$$

$$\text{Intercept} = w_0 / w_2$$

Decision Boundary

$$y \Rightarrow \text{Intercept} + \text{Slope} * x$$

$$y = mx + c \quad \text{where } m \rightarrow \text{slope}$$

$c \rightarrow \text{Intercept.}$

$$\text{Slope (m)} = -1.7 / -0.2$$

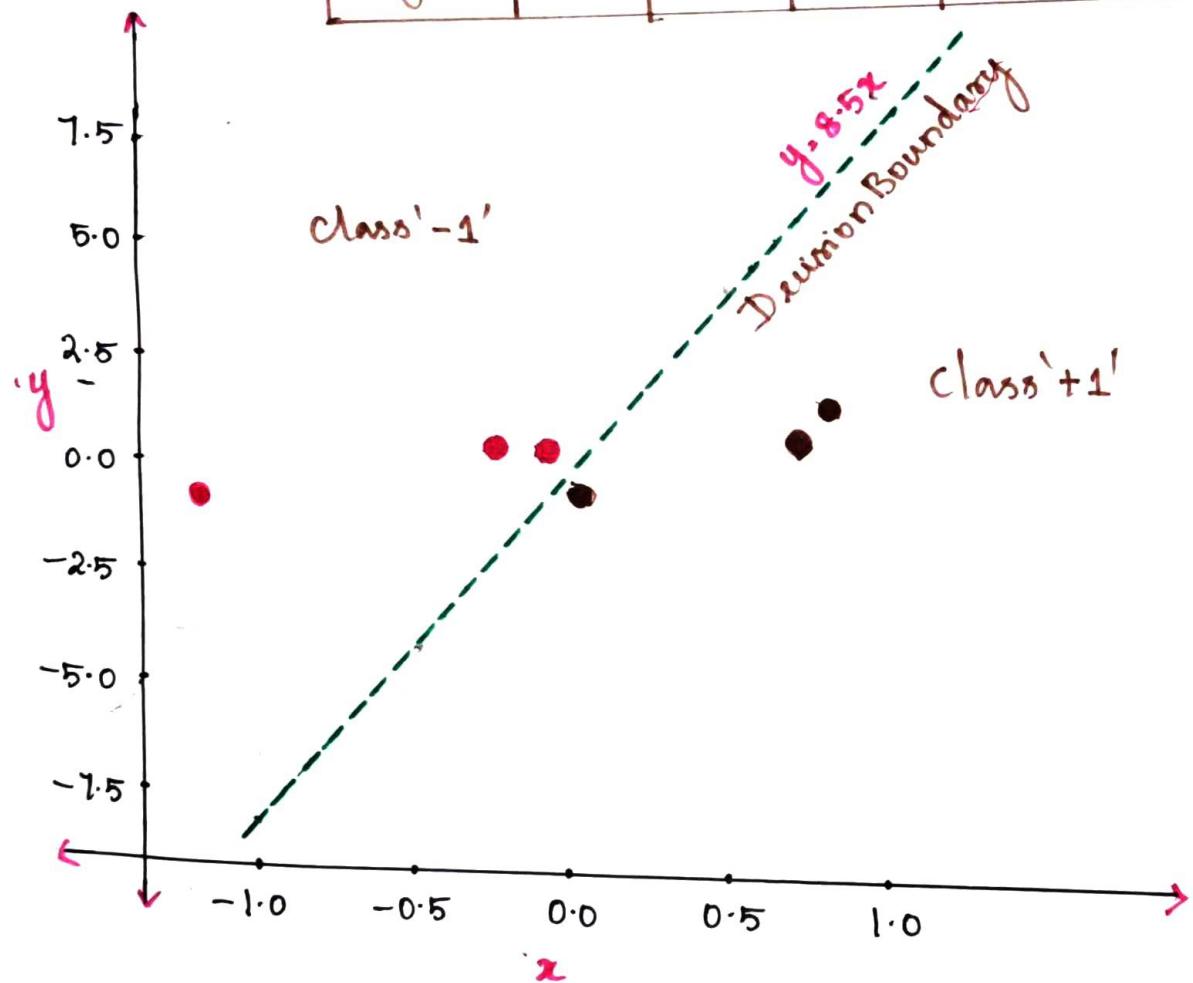
$$\text{Intercept (c)} = 0$$

Final Decision Boundary

$$y = mx \Rightarrow 8.5x.$$

To plot graph. Take Value of x in Range $(-1, 1)$

x	-1	-0.5	0	0.5	1
y	-8.5	-4.25	0	4.25	8.5



2) Hand Drawn plot.

Initial Weight Vector $\Rightarrow [1. \ 1 \ 0]$

form of $[w_1 \ w_2 \ w_0]$

$$(m) \text{ Slope} = -\frac{w_2}{w_1} = -\frac{1}{1} = -1$$

$$(c) \text{ Intercept} = -\frac{w_0}{w_1} = 0$$

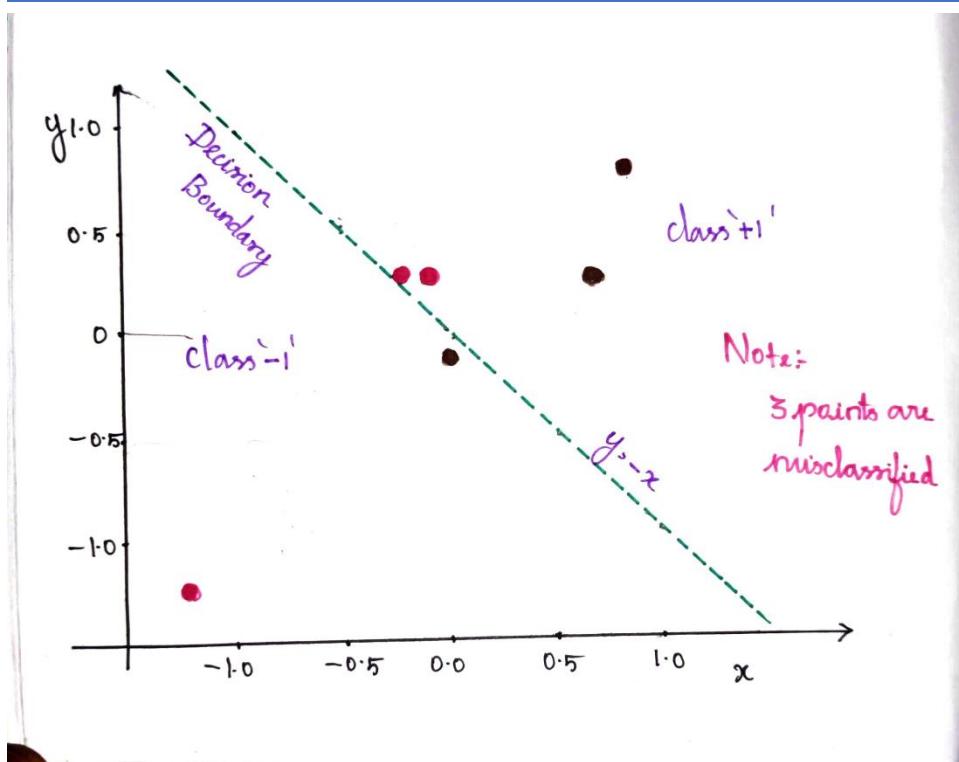
Decision Boundary

$$y = mx + c$$

$$y = -x$$

To plot the graph. (Boundary) take Value of x in Range (-1, 1)

x	-1	-0.5	0	0.5	1
y	1	0.5	0	-0.5	-1.



Iteration-1.

updated Weight At end of Iteration-I

$$\Rightarrow [1.2 \quad 0.7833 \quad -1]$$

format $[w_1 \quad w_2 \quad w_0]$

$$(m) \text{ Slope} = -w_1/w_2 = \frac{-1.2}{0.7833} = -1.532$$

$$(c) \text{ Intercept, } \Rightarrow -w_0/w_2 = \frac{-(-1)}{0.7833} = 1.277$$

Decision Boundary

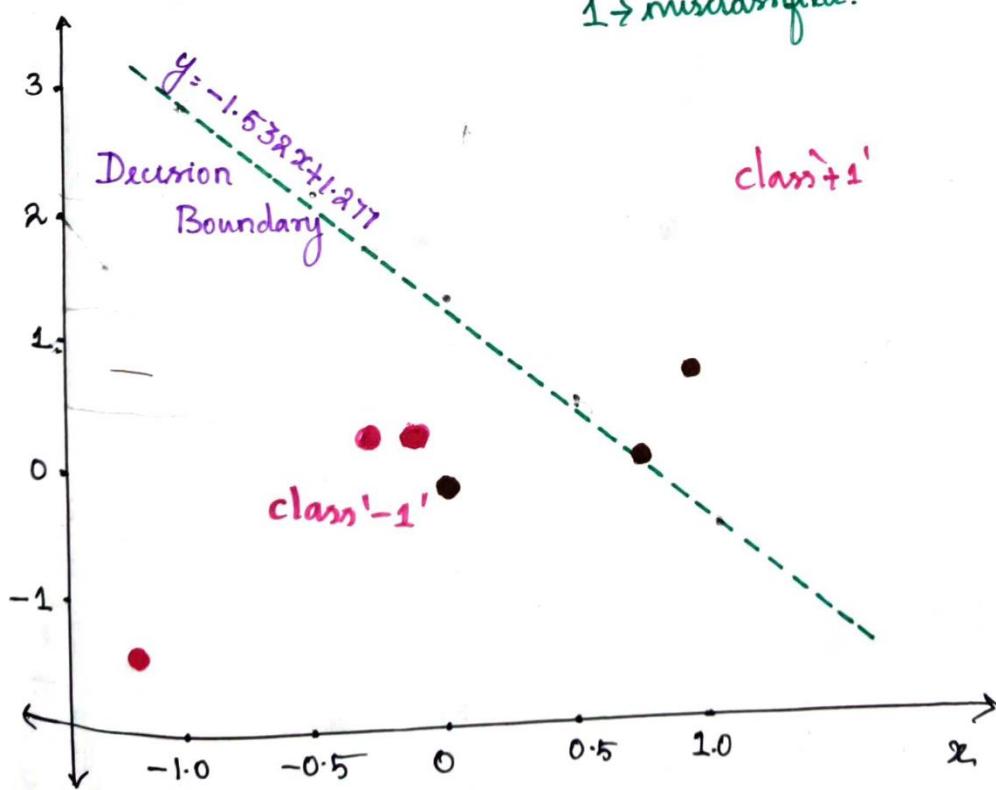
$$y = mx + c$$

$$\therefore y = -1.532x + 1.277$$

to plot the graph (Boundary) take Value of x in Range (-1, 1)

x	-1	-0.5	0	0.5	1
y	2.809	2.043	1.277	0.511	-0.255

1 → misclassified.



Iteration-8

updated weight at end of Iteration-8

$$\Rightarrow [1.2 \ 0.7 \ 0]$$

form of $[W_1 \ W_2 \ W_0]$

$$(m) \text{ slope} = -W_1/W_2 = -1.2/0.7 = -1.714$$

$$(c) \text{ Intercept} = -W_0/W_2 = 0$$

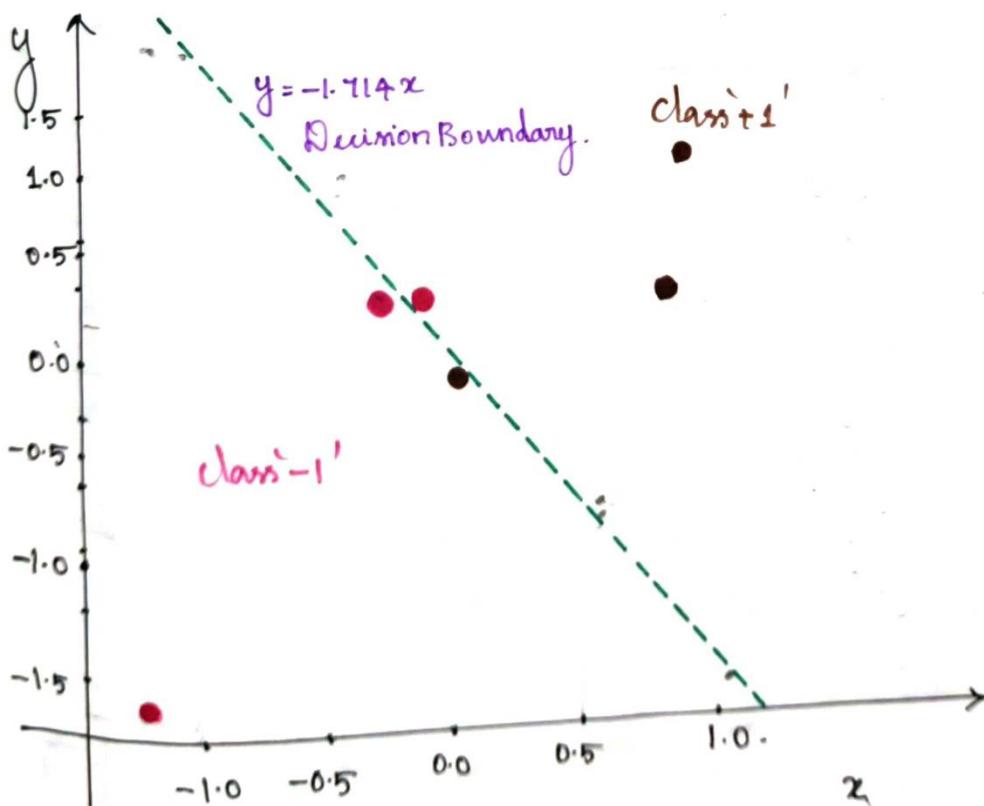
Decision Boundary

$$y = mx + c$$

$$\therefore y = -1.714x$$

To plot the graph (Boundary) taken Value of x in Range (-1,1).

x	-1	-0.5	0	0.5	1
y	1.714	0.857	0	-0.857	-1.714



Iteration-3

updated insight at end of Iteration-3

$$\Rightarrow [1.3 \quad 0.483 \quad -1]$$

form of $[w_1 \quad w_2 \quad w_0]$

$$(m) \text{ slope} = -w_1/w_2 = \frac{-1.3}{0.483} = -2.692$$

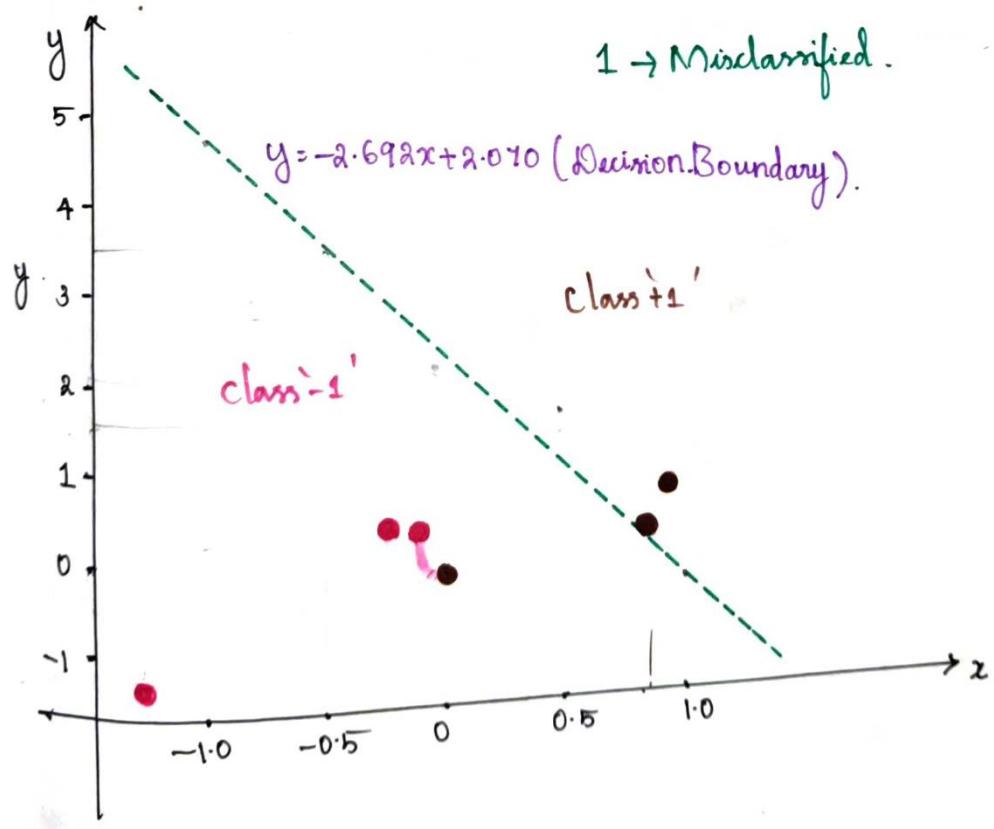
$$(c) \text{ Intercept} = -w_0/w_2 = \frac{1}{0.483} = 2.070$$

Decision Boundary $y = mx + c$.

$$\therefore y = -2.692x + 2.070.$$

To plot the graph (Boundary) take Value of x in Range $(-1, 1)$

x	-1	-0.5	0	0.5	1
y	4.762	3.416	2.070	0.724	-0.622



Iteration-4

updated weight at end of Iteration-4

$$\Rightarrow [1.3 \ 0.4 \ 0]$$

format $[W_1 \ W_2 \ W_0]$

$$(m) \text{ slope} = -W_1/W_2 = -1.3/0.4 = -3.25$$

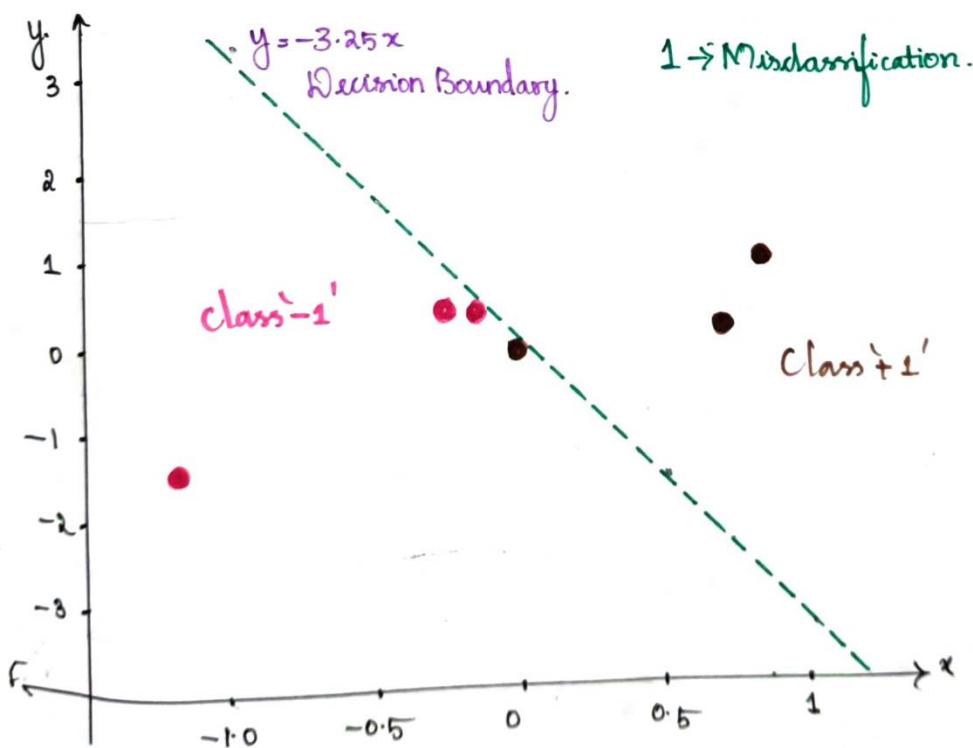
$$(c) \text{ Intercept} = -W_0/W_2 = 0$$

Decision Boundary $y = mx + c$.

$$y = -3.25x$$

To plot the graph (Boundary) take Value of x in Range $(-1, 1)$

x	-1	-0.5	0	0.5	1
y	3.25	1.63	0	-1.63	-3.25



Iteration-5

updated Weight at end of Iteration-5

$$\rightarrow [1 \cdot 3 \quad 0 \cdot 3167 \quad 1]$$

format $[w_1 \quad w_2 \quad w_0]$

$$(m) \text{ slope} = -w_1/w_2 = -1 \cdot 3 / 0 \cdot 3167 = -4 \cdot 105$$

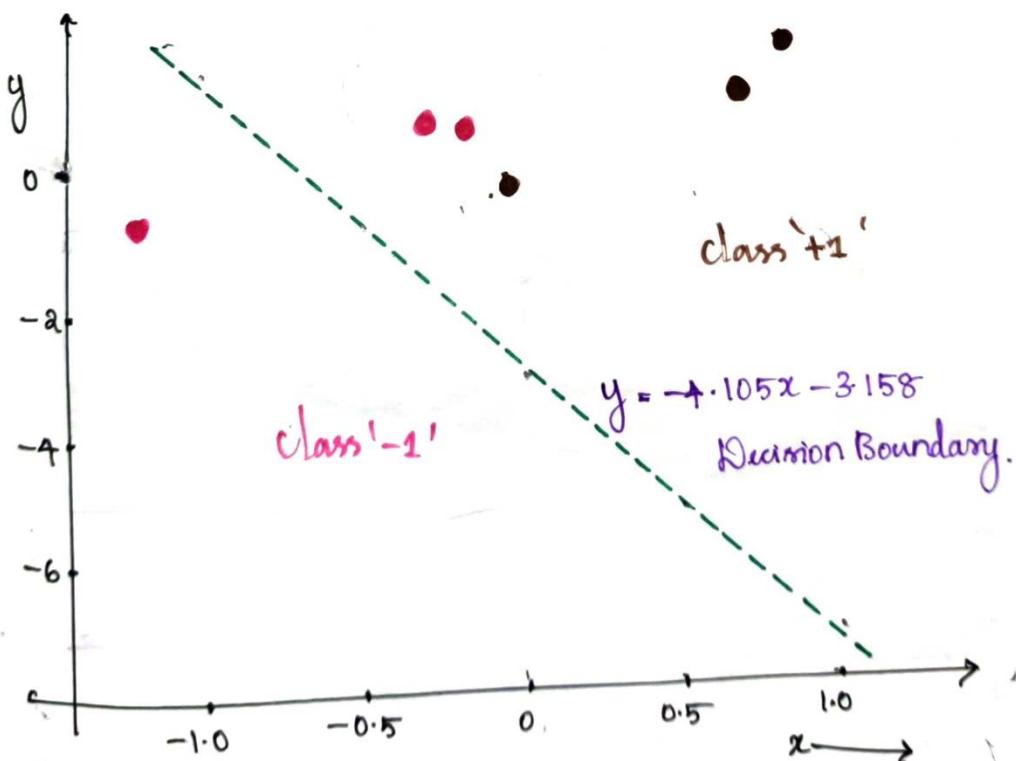
$$(c) \text{ Intercept} = -w_0/w_2 = -\frac{1}{0 \cdot 3167} = -3 \cdot 158$$

Decision Boundary $y = mx + c$.

$$y = -4 \cdot 105x - 3 \cdot 158$$

To plot the graph (Boundary) take Value of X in Range (-1, 1)

x	-1	-0.5	0	0.5	1.
y	-0.947	-1.1055	-3.158	-5.211	-7.263



Iteration-6.

updated weight at end of Iteration-6.

$$\Rightarrow [1.5 \ 0.1 \ 0]$$

form of $[w_1 \ w_2 \ w_0]$

$$(m) \text{ slope} = -w_1/w_2 = \frac{-1.5}{0.1} = -15$$

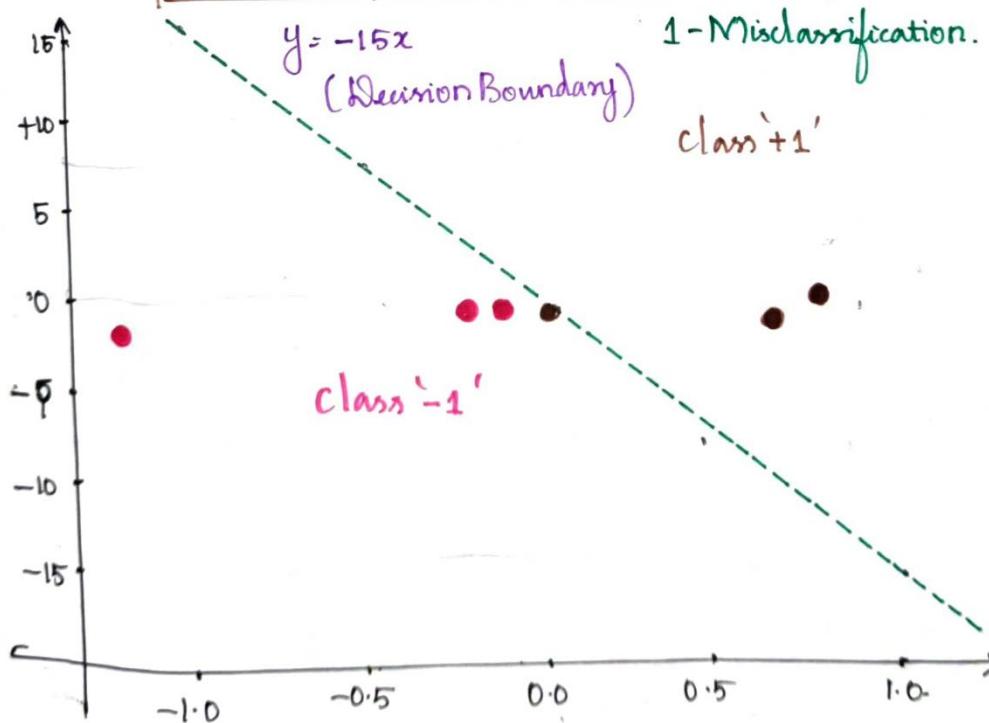
$$(c) \text{ Intercept} = -w_0/w_2 = 0$$

Decision Boundary: $y = mx + c$.

$$y = -15x$$

To plot the graph (Boundary) take value of x in Range (-1, 1)

x	-1	-0.5	0	0.5	1
y	15	7.5	0	-7.5	-15



Iteration-7.

Updated weight at end of Iteration-7.

$$\Rightarrow [1.5 \ 0.0167 \ 1]$$

form of $[w_1 \ w_2 \ w_0]$

$$(m) \text{ slope} = -\frac{w_1}{w_2} = \frac{-1.5}{0.0167} = -89.82$$

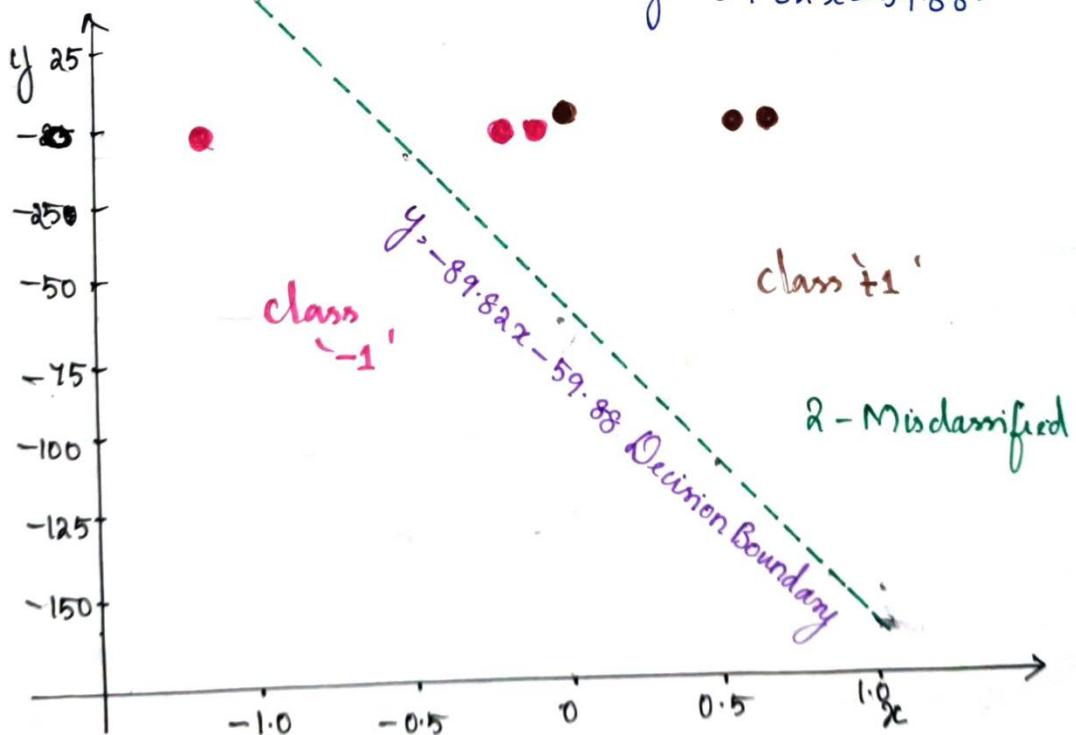
$$(c) \text{ Intercept} = -\frac{w_0}{w_2} = \frac{-1}{0.0167} = -59.88$$

To plot the graph (Boundary) take value of x in Range (-1, 1).

x	-1	-0.5	0	0.5	1
y	29.94	-14.97	-59.88	-104.79	-149.7

Decision Boundary $y = mx + c$

$$y = -89.82x - 59.88$$



Iteration-8. (Last update / Optimal Weight)

updated weight at end of Iteration-8

$$= [1.7 \ -0.2 \ 0]$$

format $[W_1 \ W_2 \ W_0]$

$$(m) \text{ slope} = -W_1/W_2 = \frac{-1.7}{-0.2} = 8.5$$

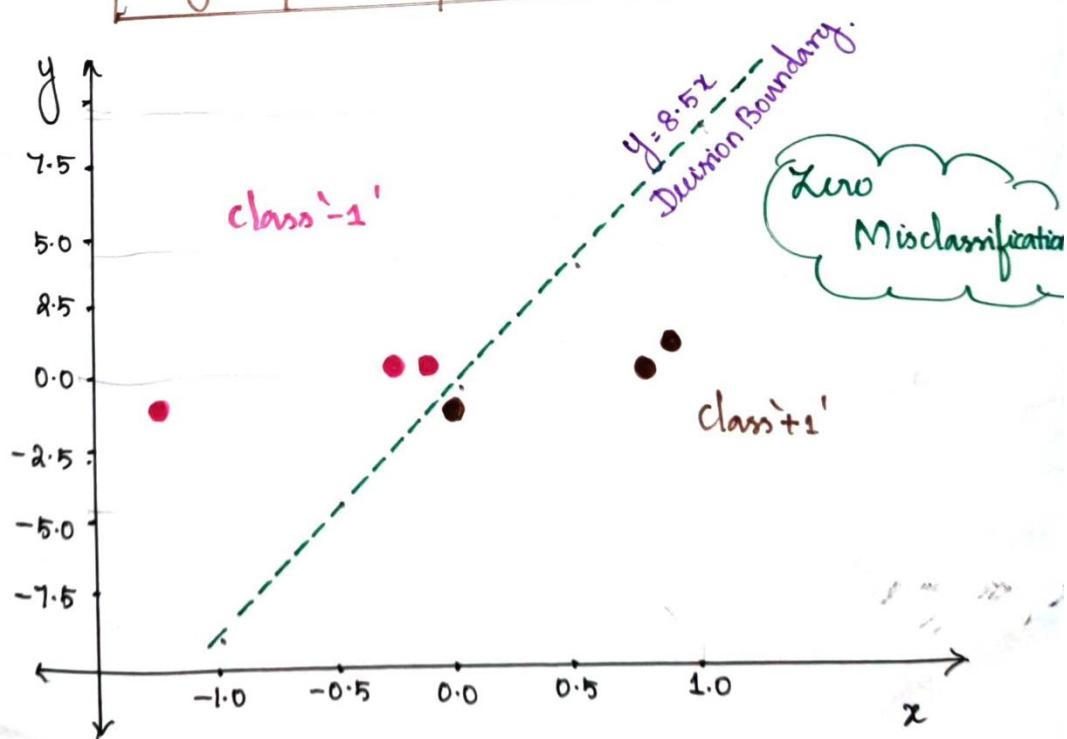
$$(c) \text{ Intercept} = \frac{-W_0}{W_2} = 0$$

Decision Boundary $y = mx + c$.

$$y = 8.5x$$

To plot the graph (Boundary) take value of x in Range (-1, 1)

x	-1	-0.5	0	0.5	1.
y	-8.5	-4.25	0	4.25	8.5



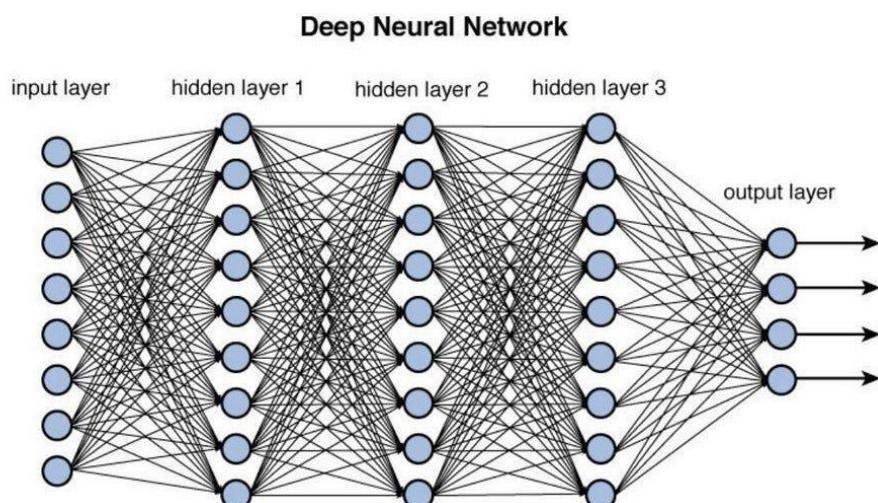
2. Neural network solution for classifying Gurmukhi digits.

About Dataset

The Gurmukhi Handwritten Digit Classification dataset is a collection of handwritten digits in the Gurmukhi script. Gurmukhi is one of the popular Indian scripts widely used in the Indian state of Punjab. The dataset consists of 2,560 images of handwritten digits from 0 to 9, written by 80 different individuals. Each image is 32x32 pixels in size and is grayscale. The dataset is intended for the development and testing of machine learning models for digit recognition in the Gurmukhi script.

Digit recognition is an important problem in the field of image processing and computer vision, and has various real-world applications, such as in optical character recognition (OCR) systems, handwritten digit recognition in cheque processing, and automatic number plate recognition (ANPR) systems. Developing accurate and efficient models for digit recognition in different scripts and languages can help improve the performance of these applications, and make them more accessible to a wider audience.

The Gurmukhi Handwritten Digit Classification dataset is a valuable resource for researchers and developers working on digit recognition problems in the Gurmukhi script. It provides a diverse set of images from different individuals, which can help improve the robustness of the models and reduce the effects of individual writing styles. Additionally, the dataset is relatively small and can be easily used for experimentation and testing of different models and algorithms.



Initial Processing Technique for Given train Data

Step 1: Importing the Required Libraries The first step is to import the required libraries that are used in the code. The following libraries are imported:

- os: It is used to perform various operations on files and directories like listing all the files in a directory.
- cv2: It is used for image processing and computer vision tasks like reading an image, resizing it, and converting it to grayscale.
- numpy: It is used for mathematical operations on arrays.
- pyplot from matplotlib: It is used to visualize the images.

Step 2: Mounting Google Drive The next step is to mount the Google Drive so that the code can access the files and directories present in it. The code mounts the Google Drive using the drive.mount() function.

Step 3: Defining the Path to the Train Directory The train directory path is defined using the train_path variable. This variable contains the path to the directory containing the training images.

Step 4: Getting the List of Folders in the Train Directory The next step is to get the list of all folders in the train directory using the os.listdir() function. This function returns a list of all files and folders present in the directory.

Step 5: Defining the Image Size and the Number of Channels The image size and the number of channels are defined using the img_size and num_channels variables. In this case, the image size is set to 28x28 and the number of channels is set to 1 as the images are grayscale.



Step 6: Defining the Empty Lists for Images and Labels The empty lists to store the images and their respective labels are defined using the x_train and y_train variables.

Step 7: Looping Through Each Folder and Reading the Images In this step, a loop is created to iterate through each folder in the train directory. Inside this loop, another loop is created to read each image present in the folder. The following steps are performed in this loop:

- The path to the current folder is defined using the os.path.join() function.

- The list of all image files in the current folder is obtained using the os.listdir() function.
- Another loop is created to iterate through each image in the current folder.
- The path to the current image is defined using the os.path.join() function.
- The image is read using the cv2.imread() function and converted to grayscale.
- The image is resized to the defined size using the cv2.resize() function.
- The image is flattened and normalized by dividing it by 255 to keep the pixel values between 0 and 1.
- The image and its respective label are appended to the x_train and y_train lists respectively.

Step 8: Converting the Lists to Numpy Arrays Finally, the x_train and y_train lists are converted to numpy arrays using the np.array() function.

Step 9: Printing the Shape of the Training Data and Target Labels The last step is to print the shape of the training data and target labels using the print() function.

Advantages of the Initial Processing Technique:

- The code reads images from the given directory and resizes them to the given size.
- The images are converted to grayscale and flattened before being converted to numpy arrays.
- The images are normalized by dividing them by 255 to keep the pixel values between 0 and 1.
- The code appends the images and their respective labels to the lists and converts them to numpy arrays.

After defining the necessary parameters and lists, the code loops through each folder in the train_path directory using the os.listdir() function. This function returns a list of all the files and directories in the specified path. The loop then reads each image file in the current folder using the cv2.imread() function, which reads an image from a file and returns it as a numpy array. The IMREAD_GRAYSCALE flag is used to read the image in grayscale mode.

Next, the code resizes the image to the specified img_size using the cv2.resize() function. The flattened image is then normalized by dividing each pixel value by 255, so that the pixel values range between 0 and 1. This is important for training deep learning models, as it helps to ensure that the input data is scaled appropriately.

The flattened image and its respective label are then appended to the x_train and y_train lists, respectively. The label is extracted from the folder name using the int() function, which converts a string to an integer.

Finally, the code converts the x_train and y_train lists to numpy arrays using the np.array() function, and prints their shapes using the shape attribute.

In summary, the above code reads each image file in the specified directory, resizes it to a specified size, normalizes the pixel values, and appends the flattened image and its respective label to two separate lists. These lists are then converted to numpy arrays and printed to confirm their shapes. This is a basic example of how to preprocess an image dataset into a format suitable for training a deep learning model

Initial Processing Technique for Given test Data:

In this code block, we are processing the test dataset similarly to how we processed the train dataset in the previous code block. Here are the steps:

- Define the path to the test data directory: We start by defining the path to the test data directory using the variable test_path.
- Get the list of all folders in the test directory: We get the list of all folders in the test directory using the os.listdir() method and store the list in the variable folders.
- Define the image size and the number of channels: We define the image size and number of channels for our images using the variables img_size and num_channels.
- Define empty lists to store the images and their respective labels: We define two empty lists x_test and y_test to store the test images and their respective labels.
- Loop through each folder and read the images: We loop through each folder in the folders list and read the images from each folder. We define the path to the current folder using os.path.join() method and the folder variable. We get the list of all image files in the current folder using os.listdir() method and store it in the variable files.
- Loop through each image and read it: We loop through each image in the files list and define the path to the current image using os.path.join() method and the img_path variable. We read the image using the cv2.imread() method and resize it to the defined size using cv2.resize() method. We then append the flattened image to the x_test list after normalizing it by dividing it by 255.

- Append the image and its respective label to the lists: We append the image and its respective label to the x_test and y_test lists, respectively. We use the int(folder) method to convert the folder variable from a string to an integer.
- Convert the lists to numpy arrays: We convert the x_test and y_test lists to numpy arrays using the np.array() method and store them in the variables x_test and y_test.
- Print the shape of the training data and target labels: We print the shape of the x_test and y_test numpy arrays to ensure that the data has been loaded correctly.

In summary, this code block reads the test images from the subdirectories of the test_path directory, resizes them to the defined size, and stores them in the x_test numpy array. The corresponding labels are stored in the y_test numpy array. The test data is then ready to be used for model evaluation.

Common Data Pre-processing Technique for Image Dataset

Resizing and cropping are common techniques used in image data preprocessing, which involve manipulating the size and dimensions of images in a dataset. These techniques are often used to standardize images so that they can be fed into machine learning models with consistent and appropriate dimensions.

Resizing involves adjusting the resolution or number of pixels in an image. The process of resizing involves interpolation, which is the estimation of new pixel values based on existing pixel values. There are two primary methods of interpolation used for resizing images: nearest neighbor and bilinear interpolation. Nearest neighbor interpolation selects the nearest pixel value to use for the new pixel value, while bilinear interpolation calculates a weighted average of the closest four pixel values.

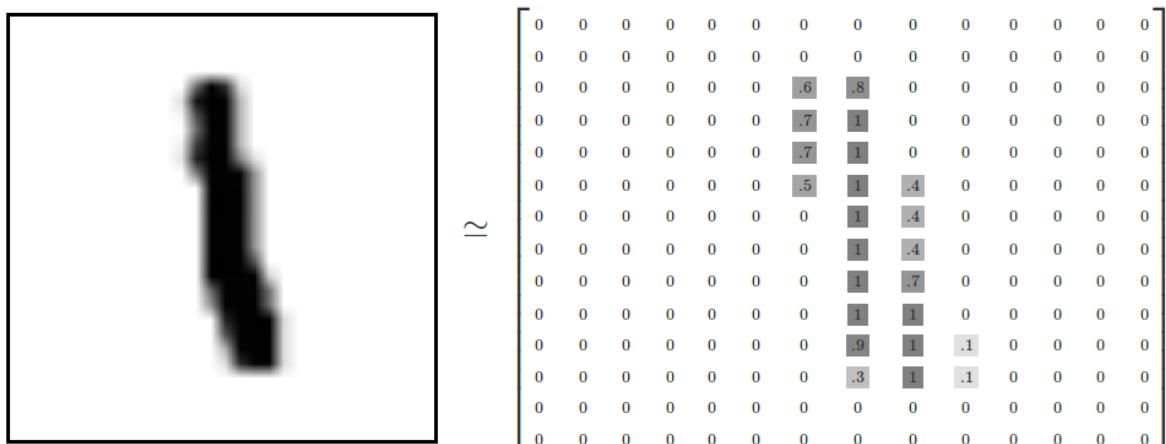
Cropping involves removing parts of an image to create a new image with a different size or aspect ratio. This technique is often used to remove irrelevant or distracting parts of an image, or to focus on a specific object or region of interest. The process of cropping involves selecting a rectangular region of an image and discarding the pixels outside of this region.

There are several advantages to resizing and cropping image datasets for use in machine learning models:

- Consistent dimensions: Resizing and cropping can be used to standardize images so that they all have the same dimensions, which is often necessary for machine learning models to be able to process the images.
- Reduced computational requirements: Resizing and cropping can reduce the size and complexity of image datasets, which can reduce the computational resources required to train machine learning models.
- Improved accuracy: Resizing and cropping can improve the accuracy of machine learning models by reducing noise and irrelevant information in the images, and by focusing on the relevant regions of interest.
- Augmentation: Resizing and cropping can be used as part of a data augmentation strategy, which involves generating additional training data by applying transformations to the existing data. By applying random resizing and cropping to the images in a dataset, it is possible to generate a larger and more diverse set of training examples.

In summary, resizing and cropping are important techniques for image data preprocessing that can improve the consistency, efficiency, accuracy, and diversity of image datasets used in machine learning models.

Converting the image dataset into Numpy ARRAY



Converting image datasets into numpy arrays is a common technique used in machine learning and deep learning applications for image analysis. The numpy array format is a popular choice because it is efficient, flexible, and compatible with most machine learning frameworks.

The process of converting an image dataset into a numpy array involves several steps:

- **Loading the images:** The first step is to load the images into the computer's memory. This can be done using a variety of tools and libraries, such as PIL (Python Imaging Library), OpenCV, or scikit-image.
- **Resizing and/or cropping:** In some cases, it may be necessary to resize and/or crop the images to ensure that they have consistent dimensions and are suitable for analysis by a machine learning model.
- **Converting to grayscale:** If the images are in color, it may be necessary to convert them to grayscale, which reduces the dimensionality of the data and simplifies the analysis.
- **Converting to numpy arrays:** Once the images have been preprocessed, they can be converted into numpy arrays using the appropriate functions or libraries, such as numpy or scipy. Each image is represented as a two-dimensional or three-dimensional array, depending on whether the image is grayscale or color.
- **Normalization:** Finally, the numpy arrays may need to be normalized to ensure that the pixel values are within a certain range, such as 0-1 or -1 to 1. This can improve the performance of machine learning models by ensuring that the data is consistent and within an appropriate range for analysis.

There are several advantages to using numpy arrays to represent image data:

- **Efficiency:** Numpy arrays are highly efficient data structures, which can be easily manipulated and processed using machine learning libraries and frameworks.
- **Flexibility:** Numpy arrays can be easily reshaped, sliced, and transformed, which makes them highly flexible and adaptable to a wide range of machine learning applications.
- **Compatibility:** Numpy arrays are compatible with most machine learning frameworks, such as TensorFlow, PyTorch, and scikit-learn, which makes them a popular choice for image analysis applications.
- **Compactness:** Numpy arrays are a compact representation of image data, which reduces the memory requirements and makes it easier to store and process large datasets.

In summary, converting image datasets into numpy arrays is a common technique used in machine learning and deep learning applications. This process involves several steps, including loading, preprocessing, and converting the images into numpy arrays. Numpy arrays are highly efficient, flexible, and compatible with

most machine learning frameworks, which makes them a popular choice for image analysis applications.

Converting the Numpy Array of image Dateset into CSV FILE Form of Input-Train Dataset:

	label	pixel0	pixel1	pixel2	pixel3	pixel4	pixel5	pixel6	pixel7	pixel8	...	pixel774	pixel775	pixel776	pixel777	pixel778
0	6	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	...	0.929412	0.929412	0.929412	0.929412	0.945091
1	6	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	...	0.000000	0.000000	0.000000	0.000000	0.200000
2	6	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	...	0.000000	0.000000	0.000000	0.000000	0.000000
3	6	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	...	0.929412	0.929412	0.929412	0.933333	1.000000
4	6	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	...	0.929412	0.984314	1.000000	1.000000	1.000000
...
195	2	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	0.984314	...	1.000000	1.000000	1.000000	1.000000	1.000000
196	2	1.0	1.0	1.000000	0.964706	0.929412	0.929412	0.066667	0.000000	0.000000	...	1.000000	1.000000	1.000000	1.000000	1.000000
197	2	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	...	1.000000	1.000000	1.000000	1.000000	1.000000
198	2	1.0	1.0	0.972549	0.929412	0.929412	0.929412	0.929412	0.000000	0.000000	...	1.000000	1.000000	1.000000	1.000000	1.000000
199	2	1.0	1.0	1.000000	1.000000	1.000000	1.000000	1.000000	0.996078	0.929412	...	1.000000	1.000000	1.000000	1.000000	0.984314

200 rows x 785 columns

We are converting the training dataset, which is stored in numpy array format, to a CSV file format. The steps involved are as follows:

- We first create a list of column names for the Xtrain data. This list will be used as column headers for our CSV file. The list contains the string "pixel" followed by a number from 0 to 783. This is because each image in our dataset has 784 pixels.
- We then stack the y_train and x_train arrays horizontally using numpy's column_stack() function. This results in a combined array with the label column as the first column, and the pixel values as the subsequent columns.
- We then use numpy's savetxt() function to save this combined data to a CSV file. The delimiter parameter is set to "," to specify that the columns should be separated by commas. The header parameter is set to "label," followed by the column names separated by commas. This specifies the column headers for the CSV file. The comments parameter is set to "" to indicate that there should be no comments in the CSV file.
- The resulting CSV file, "output.csv", will have 1000 rows (since we have 1000 training images) and 785 columns (since we have 784 pixel columns and 1 label column).

Overall, this code block is useful for converting a numpy array dataset to a CSV file format, which is a common format used in many machine learning tasks. This allows us to easily manipulate the data using tools such as pandas, or import it into other machine learning libraries for further analysis.

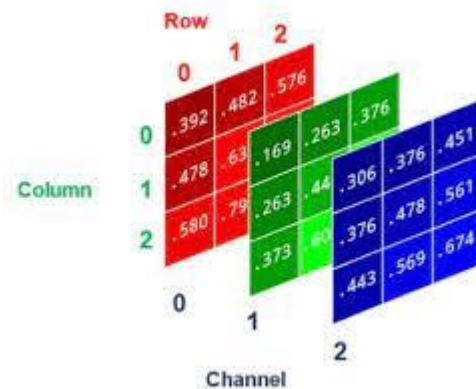
Converting the Numpy Array of image Dateset into CSV FILE Form of Input-Test Dataset:

We are converting the numpy arrays of the test data into a CSV file format. This is helpful in cases where we need to share the data with other teams or tools that require CSV file format input.

Here is the step-by-step explanation of the code:

- We first import the NumPy library to work with arrays and perform mathematical operations.
- We define a list of column names for the test data, which will later be used as column headings in the CSV file. We create the list using a for loop that iterates over the range of 784, which is the total number of pixels in each image. We append the string "pixel" to the index value to create the column names.
- We stack the y_test and x_test numpy arrays horizontally using the column_stack() function from NumPy. This results in a new 2D array where each row consists of a label followed by the flattened pixel values of an image.
- Finally, we use the savetxt() function from NumPy to save the combined data to a CSV file. We specify the file name as "testoutput.csv", the delimiter as a comma, the header as a string that concatenates "label," with the column names separated by commas, and the comments as an empty string.

This code generates a CSV file with 178 rows (corresponding to the 178 test images) and 785 columns (corresponding to the label column and the 784 pixel columns). The first column contains the labels and the remaining columns contain the pixel values of the images.



	label	pixel0	pixel1	pixel2	pixel3	pixel4	pixel5	pixel6	pixel7	pixel8	...	pixel774	pixel775	pixel776	pixel777	pixel778
0	4	1.0	1.0	1.000000	1.000000	1.0	1.000000	0.933333	0.862745	0.000000	...	0.596078	0.929412	0.996078	1.0	1.0
1	4	1.0	1.0	1.000000	1.000000	1.0	1.000000	1.000000	1.000000	1.000000	...	1.000000	1.000000	1.000000	1.0	1.0
2	4	1.0	1.0	0.972549	0.462745	0.0	0.000000	0.862745	1.000000	1.000000	...	1.000000	1.000000	1.000000	1.0	1.0
3	4	1.0	1.0	1.000000	1.000000	1.0	1.000000	0.933333	0.929412	0.729412	...	0.000000	0.729412	0.996078	1.0	1.0
4	4	1.0	1.0	1.000000	1.000000	1.0	0.945098	0.066667	0.000000	0.200000	...	1.000000	1.000000	1.000000	1.0	1.0
...
173	7	1.0	1.0	1.000000	1.000000	1.0	1.000000	1.000000	1.000000	1.000000	...	1.000000	1.000000	1.000000	1.0	1.0
174	7	1.0	1.0	1.000000	1.000000	1.0	0.945098	0.929412	0.929412	0.929412	...	1.000000	1.000000	1.000000	1.0	1.0
175	7	1.0	1.0	1.000000	1.000000	1.0	1.000000	1.000000	1.000000	1.000000	...	1.000000	1.000000	1.000000	1.0	1.0
176	7	1.0	1.0	1.000000	1.000000	1.0	1.000000	1.000000	0.996078	0.929412	...	1.000000	1.000000	1.000000	1.0	1.0
177	7	1.0	1.0	1.000000	1.000000	1.0	1.000000	1.000000	0.996078	0.929412	...	1.000000	1.000000	1.000000	1.0	1.0

178 rows × 785 columns

Data visualization:



Now loads the 'output.csv' file into a Pandas DataFrame called 'data'. Pandas is a popular Python library used for data manipulation and analysis.

The next step we are converts the 'label' column from a string data type to an integer

data type using the 'astype()' method. This is because the 'label' column represents the class labels of the corresponding images, and it is easier to work with integer data types when building machine learning models.

Finally, the 'head()' method is called on the 'data' DataFrame to display the first 200 rows of the data, including the 'label' column and the pixel values for each image. 'head()' is a method that allows you to view the first n rows of the DataFrame. The default value of n is 5, but in this case, it has been set to 200.

Test Data visualization

For this process we are imports the pandas library and uses it to read a CSV file called 'testoutput.csv' into a pandas DataFrame called 'testdata'. The 'astype(int)' method is used to convert the 'label' column in the DataFrame from a string type to an integer type. Finally, the 'head()' method is used to display the first 178 rows of the DataFrame in the console.

In summary, this code reads a CSV file of test data, converts the 'label' column to integer type, and displays the first 178 rows of the DataFrame.

Plotting the Train Data:

- For this process we are initializes a numpy array called "data" with the contents of a pandas dataframe that is read from a csv file.
- The next line gets the shape of the data array and assigns the number of rows to "m" and the number of columns to "n".
- The "np.random.shuffle" function is used to shuffle the rows of the "data" array randomly. This is done to prevent any order bias in the data while training.

The benefits of shuffling the elements of an array are:

- **Randomization:** Shuffling the array elements randomizes the order of the elements. This is useful for avoiding any systematic biases that might arise from the original ordering of the elements.
 - **Better training:** Shuffling the elements of an array is particularly useful in training machine learning models. If the input data has any inherent structure or ordering, shuffling can help the model to learn more effectively by preventing it from overfitting to the structure of the data. In other words, it ensures that the model sees a diverse range of examples during training.
 - **Better testing:** Shuffling the elements of an array can also help to create better testing data. If the input data has any inherent structure or ordering, shuffling can help to ensure that the testing data is diverse and representative of the true distribution of the data.
 - **Reproducibility:** Shuffling the elements of an array can also help to ensure the reproducibility of experiments. By shuffling the elements, we can avoid any biases that might arise from the ordering of the data, and ensure that the results are consistent across multiple runs of the experiment.
-
- The next line selects all the rows of the "data" array and transposes it to create "data_train".
 - The "Y_train" variable is initialized with the first row of "data_train" which contains the labels of the images as integers.
 - The "X_train" variable is initialized with all the remaining rows of "data_train" which contain the pixel values of the images.
 - The "_" before "m_train" indicates that the variable is not being used.
 - The "label" variable is initialized with the label of the second image in the "Y_train" array.
 - The "current_image" variable is initialized with the pixel values of the second image in the "X_train" array.

- The "current_image" variable is reshaped into a 28x28 array and multiplied by 255 to scale the pixel values back to their original range.
- The "plt.gray()" function sets the color map to grayscale for display.
- The "plt.imshow" function is used to display the image.
- Finally, the "plt.show()" function displays the image.

To print an image dataset in 2D, the following steps are typically followed:

- The image dataset is loaded into memory using a suitable library or function, such as OpenCV, PIL, or NumPy.
- The image is reshaped to a 2D format, usually using the reshape() function in NumPy.
- The grayscale intensity values of the pixels in the image are normalized to a range between 0 and 1, so that they can be displayed properly.
- The imshow() function from a plotting library, such as Matplotlib, is used to display the 2D image on the screen.
- Any additional customizations, such as changing the color map, adding a color bar, or setting the axis labels, can be done using the relevant functions or attributes of the plotting library.

The final result is a 2D grayscale image that can be displayed on the screen, saved to a file, or used as input for a machine learning algorithm.

Plotting the Test Data

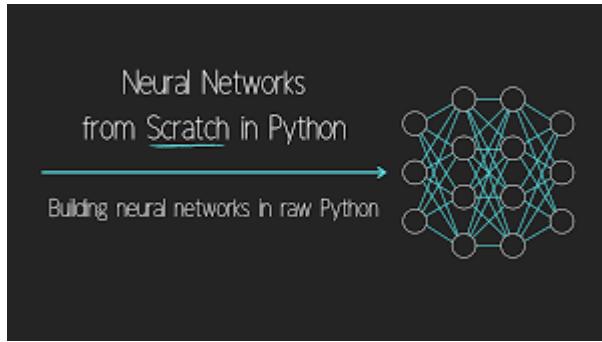
First, the test data is converted into a NumPy array using the np.array() function. Then, the shape of the test data is determined using testdata.shape, where m represents the number of rows and n represents the number of columns.

Next, the rows of the test data are randomly shuffled using np.random.shuffle(testdata). This is done to ensure that the order of the images does not affect the evaluation process.

Then, the shuffled test data is split into labels and feature vectors using the same approach as in the training data. The data_dev variable contains the transpose of the shuffled test data, where the first row represents the labels and the remaining rows represent the feature vectors. The Y_dev variable is the first row of data_dev (i.e., the labels), converted to integer values using the astype() function. The X_dev variable is the remaining rows of data_dev (i.e., the feature vectors).

Finally, the code prints the label of the second image in the test dataset using `label = Y_dev[1]`, and displays the image using `plt.imshow()` function. The image is retrieved from the feature vectors using `X_dev[:, 1, None]` and reshaped to a 28x28 array before displaying.

Implement Neural Network from scratch:



The neural network has an input layer with 784 neurons, one hidden layer with 10 neurons, and an output layer with 10 neurons (corresponding to the 10 possible classes of the input images). The neural network uses ReLU activation function for the hidden layer and softmax activation function for the output layer.

Here is a step-by-step explanation of the code:

- **`init_params()`** function initializes the weights and biases of the neural network. It initializes four parameters: W_1 , b_1 , W_2 , and b_2 . W_1 and b_1 are the weights and biases of the hidden layer, respectively, and W_2 and b_2 are the weights and biases of the output layer, respectively.
- **`ReLU()`** function is the activation function for the hidden layer. It takes a matrix Z as input and applies the rectified linear unit (ReLU) function element-wise to the matrix. The ReLU function returns the matrix with all negative elements set to zero.
- **`softmax()`** function is the activation function for the output layer. It takes a matrix Z as input and applies the softmax function to the matrix. The softmax function returns a matrix with probabilities for each class.
- **`forward_prop()`** function performs the forward propagation step of the neural network. It takes the weights and biases of the neural network and the input data X as input. It computes the dot product of W_1 and X , adds b_1 , applies the ReLU activation function, computes the dot product of W_2 and the output of the ReLU function, adds b_2 , and applies the softmax activation function. The function returns the intermediate values Z_1 , A_1 , Z_2 , and A_2 .

- **ReLU_deriv()** function calculates the derivative of the ReLU function. It takes a matrix Z as input and returns a matrix of the same shape with elements equal to 1 for positive elements of Z and 0 for non-positive elements.
- **one_hot()** function creates a one-hot encoded matrix from the input label vector Y. It creates a matrix of zeros with the number of rows equal to the size of Y and the number of columns equal to the maximum value of Y plus 1. Then it sets the element of each row corresponding to the value of Y to 1.
- **backward_prop()** function performs the backward propagation step of the neural network. It takes the intermediate values Z1, A1, Z2, and A2, the weights W1 and W2, the input data X, and the label vector Y as input. It first computes the one-hot encoded matrix from Y using the one_hot() function. It then calculates the derivatives of the cost function with respect to the weights and biases of both layers using the backpropagation algorithm. The function returns the gradients dW1, db1, dW2, and db2.
- **update_params()** function updates the weights and biases of the neural network using the gradients and the learning rate alpha. It takes the weights and biases W1, b1, W2, b2, the gradients dW1, db1, dW2, and db2, and the learning rate alpha as input. It returns the updated weights and biases.

Overall, these functions are used in a loop to train the neural network on the input data. During each iteration of the loop, the `

What is Relu Function?

ReLU (Rectified Linear Unit) is an activation function commonly used in deep learning neural networks. It is a simple function that takes a single input and produces an output by returning the maximum of the input and 0.

Mathematically, the ReLU function is defined as:

$$f(x) = \max(0, x)$$

where x is the input to the function and $\max(0, x)$ returns the maximum value between 0 and x. Therefore, if x is a positive value, ReLU outputs the same value as x, but if x is negative, ReLU outputs 0.

The main reason for using ReLU activation function is its simplicity, computational efficiency, and its ability to address the vanishing gradient problem in deep neural networks. The vanishing gradient problem occurs when the gradients become smaller and smaller as they propagate through layers, which

results in slow training or completely stopped training. Since ReLU has a non-zero derivative for positive inputs, it avoids the vanishing gradient problem and provides a better gradient flow through the network, leading to faster training and better performance. Additionally, ReLU allows for sparse representations and reduces the likelihood of overfitting by making the model less complex.

Various Weight Initializing Technique:

Weight initialization is a crucial step in training neural networks. Initializing weights with suitable values can accelerate training and improve model performance. Here are some of the weight initialization techniques:

- **Random Initialization:** The simplest method is to initialize the weights randomly. The idea is to initialize the weights close to zero, but not too small, to avoid the vanishing gradient problem.
- **Xavier Initialization:** This technique takes into account the number of input and output units of a layer. It initializes the weights randomly, but it scales them by the square root of the inverse of the number of input units.
- **He Initialization:** This technique is similar to Xavier initialization, but it takes into account the activation function used in the layer. It scales the weights by the square root of two divided by the number of input units, in the case of ReLU activation function.
- **Orthogonal Initialization:** This technique initializes the weights as a random orthogonal matrix. An orthogonal matrix is a matrix whose columns are orthonormal vectors.
- **LeCun Initialization:** This technique is similar to He initialization, but it scales the weights by the square root of the inverse of the number of input units. It was introduced to improve the training of Convolutional Neural Networks.

Each weight initialization technique has its strengths and weaknesses, and the choice of the technique depends on the problem at hand, the architecture of the network, and the activation function used.

Soft max function:

The softmax() function is a widely used activation function in neural networks, especially in multi-class classification tasks. It takes a vector of real numbers as input and transforms them into a probability distribution over the classes. The output of the softmax function can be interpreted as the probabilities that the input belongs to each class.

The mathematical formula for the softmax function is as follows:

$$\sigma(\vec{z})_i = \frac{e^{z_i}}{\sum_{j=1}^K e^{z_j}}$$

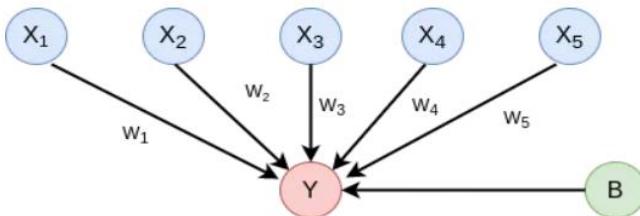
where z_j is the input to the j^{th} neuron, and K is the total number of neurons in the layer.

The softmax function works by taking the exponential of each input value and then normalizing them by dividing the sum of all exponentials. The resulting output values are in the range of 0 to 1 and their sum is equal to 1, making them suitable for use as probabilities.

The softmax function is commonly used in the output layer of a neural network when performing multi-class classification tasks, such as image classification or language translation. In such tasks, the goal is to predict the probability distribution of the input belonging to each class. The softmax function helps to ensure that the output probabilities are well-calibrated, i.e., they accurately reflect the uncertainty in the prediction.

Explain the forward propagation technique used in neural network in detail:

Forward propagation is a fundamental concept in neural networks. It refers to the process of computing the output of a neural network given a set of input data. Forward propagation is essential for training the neural network, making predictions, and evaluating the performance of the network.



$$y = w_1x_1 + w_2x_2 + w_3x_3 + \dots + w_nx_n + b$$

Equation 1

In a neural network, the input data is fed to the input layer, which is then transformed by a series of hidden layers, and finally, the output is produced by the output layer. The goal of forward propagation is to compute the output of the neural network given the input data.

The forward propagation process involves two main steps: a linear transformation and a non-linear transformation. The linear transformation is performed by computing the weighted sum of the input features and the corresponding weights of the neurons in the network. The non-linear transformation is then applied to the output of the linear transformation, which introduces non-linearity into the network.

The linear transformation is performed using the dot product of the input features and the weights of the neurons. In other words, the output of each neuron in the network is computed by multiplying the input features with their corresponding weights and adding a bias term. Mathematically, this can be represented as:

$$Z = W \cdot X + b$$

Where: Z: Output of the linear transformation W: Weights of the neuron X: Input features b: Bias term

The output of the linear transformation is then passed through an activation function to introduce non-linearity into the network. The most commonly used activation function is the Rectified Linear Unit (ReLU) function, which is defined as:

$$f(Z) = \max(0, Z)$$

The ReLU function returns the maximum of 0 and the output of the linear transformation. This introduces non-linearity into the network and helps the neural network learn more complex patterns in the input data.

After applying the activation function, the output of the hidden layers is computed using the same linear and non-linear transformations. This process is repeated for each hidden layer until the output layer is reached.

The output layer of the neural network is responsible for producing the final output of the network. The output layer can have one or more neurons, depending on the problem being solved. For example, in a binary classification problem, the output layer would have one neuron that produces the predicted class label. In a multi-class classification problem, the output layer would have multiple neurons, one for each class label, and the predicted class label would be the neuron with the highest output.

The output of the output layer is computed using the same linear transformation as the hidden layers, followed by a softmax activation function. The softmax function is used to normalize the output of the output layer, ensuring that the sum of the output values is equal to 1. This is important in multi-class classification problems, where the predicted class label is the one with the highest probability.

The softmax function is defined as:

$$f(Z) = e^Z / \text{sum}(e^Z)$$

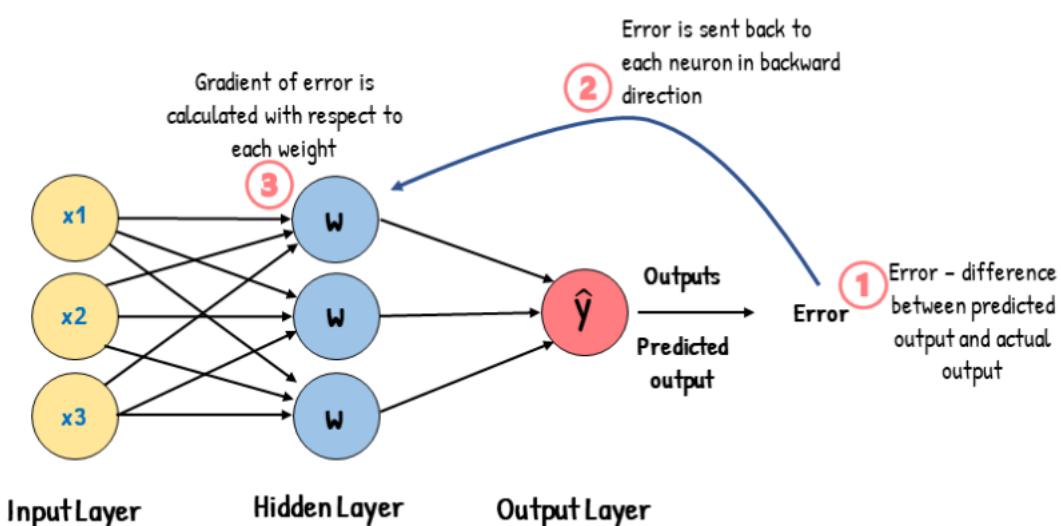
Where: Z: Output of the linear transformation e: Euler's number (2.71828)

In summary, forward propagation is the process of computing the output of a neural network given a set of input data. The process involves a linear transformation and a non-linear transformation, which is performed using an activation function. The output of the output layer is computed using the same linear and non-linear transformations, followed by a softmax activation function, which is used to normalize the output of the output layer.

Explain the back propagation technique used in neural network in detail

Backpropagation is a widely used algorithm for training neural networks. It involves the computation of the gradient of the cost function with respect to the network's parameters, which is then used to update the weights and biases of the network.

Backpropagation



The backpropagation algorithm can be divided into two main phases: the forward pass and the backward pass. In the forward pass, the input data is fed into the network, and the activations of each layer are computed. In the backward pass, the error is propagated back through the network, and the gradients of the cost function with respect to each parameter are computed.

The backward pass can be broken down into several steps. First, the gradient of the cost function with respect to the output of the network is computed. This is done using the chain rule of calculus, which allows us to express the derivative of a composite function in terms of the derivatives of its constituent functions.

Next, the error is backpropagated through each layer of the network. This involves computing the gradient of the activation function with respect to its input, and then multiplying this by the error from the next layer. The resulting product is then passed back to the previous layer.

Once the gradients have been computed, the weights and biases of the network are updated using gradient descent. The amount by which the weights and biases are updated is proportional to the gradient of the cost function with respect to the corresponding parameter, and is scaled by a learning rate.

One important aspect of the backpropagation algorithm is the choice of activation function. The activation function is used to introduce nonlinearity into the network, and different activation functions can be used depending on the nature of the problem being solved. Some popular activation functions include the sigmoid function, the hyperbolic tangent function, and the rectified linear unit (ReLU) function.

Another important aspect of the backpropagation algorithm is regularization. Regularization is a technique used to prevent overfitting by adding a penalty term to the cost function that discourages large weights. One popular regularization technique is L2 regularization, which adds a penalty term proportional to the sum of the squares of the weights.

In summary, backpropagation is a powerful algorithm for training neural networks. It involves computing the gradient of the cost function with respect to the network's parameters, and using this gradient to update the weights and biases of the network. The backpropagation algorithm can be broken down into several steps, including the forward pass, the backward pass, and gradient descent. The choice of activation function and regularization technique are important considerations when designing a neural network.

How gradient descent is helpful in back propagate?

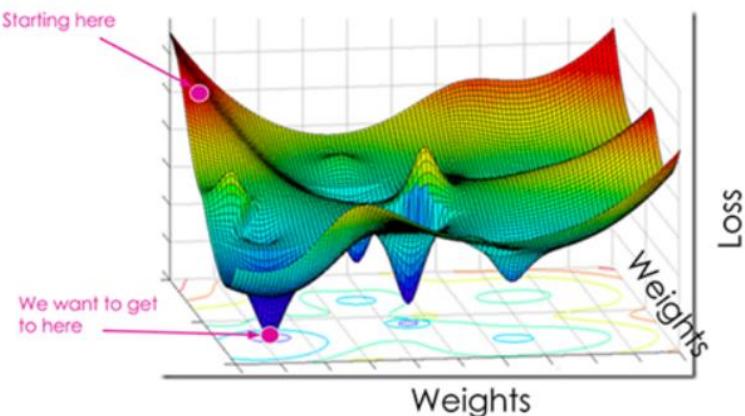
Gradient descent is a popular optimization algorithm used in machine learning for minimizing the cost function of a neural network. It is especially useful in backpropagation, which is the process of calculating the gradients of the cost function with respect to the weights and biases of the neural network. In this explanation, we will cover the basics of gradient descent, its different types, and how it is used in backpropagation.

Gradient Descent: Gradient descent is a widely used optimization algorithm that is used to minimize the cost function of a neural network. The cost function, also known as the loss function, is a measure of the difference between the predicted output of the neural network and the actual output. The goal of gradient descent is to minimize the cost function by updating the weights and biases of the neural network in the direction of the negative gradient of the cost function.

The gradient of the cost function is a vector that points in the direction of the steepest increase of the cost function. The negative gradient points in the direction of the steepest decrease, and is used in gradient descent to update the weights and biases of the neural network. The general idea behind gradient descent is to iteratively update the weights and biases of the neural network in the direction of the negative gradient until the cost function is minimized.

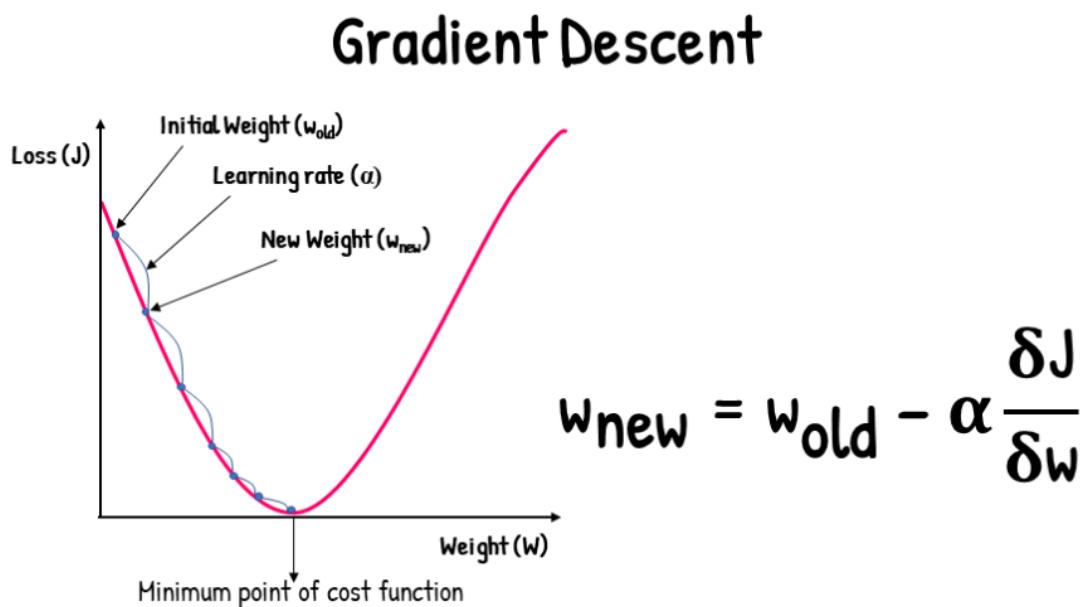
Types of Gradient Descent: There are three types of gradient descent: batch, stochastic, and mini-batch gradient descent.

- **Batch Gradient Descent:** Batch gradient descent is the most basic form of gradient descent. In this type of gradient descent, the entire dataset is used to calculate the gradient of the cost function. This means that all of the training examples are used in each iteration of the algorithm. Batch gradient descent is computationally expensive, especially when dealing with large datasets, but it is guaranteed to converge to the global minimum of the cost function.
- **Stochastic Gradient Descent:** Stochastic gradient descent (SGD) is a faster version of batch gradient descent that uses only one



training example at a time to update the weights and biases of the neural network. This means that the weights and biases are updated more frequently, leading to faster convergence. However, stochastic gradient descent can be noisy because the gradient is calculated based on a single training example, and may not always converge to the global minimum of the cost function.

Mini-batch Gradient Descent: Mini-batch gradient descent is a compromise between batch gradient descent and stochastic gradient descent. In this type of gradient descent, a small random subset of the training data, called a mini-batch, is used to calculate the gradient of the cost function. Mini-batch gradient descent is faster than batch gradient descent because it uses fewer examples to calculate the gradient, but it is less noisy than stochastic gradient descent because it uses more than one example at a time.



Backpropagation and Gradient Descent: Backpropagation is the process of calculating the gradients of the cost function with respect to the weights and biases of the neural network. Once the gradients are calculated, they are used to update the weights and biases of the neural network using gradient descent.

$$w_{\text{new}} = w_{\text{old}} - \alpha \frac{dJ}{dw}$$

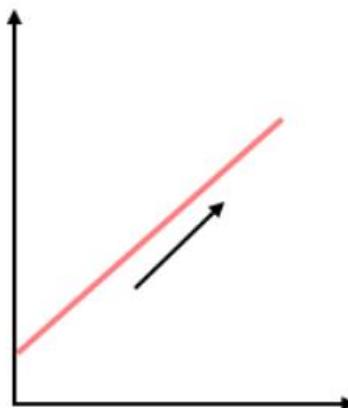
Gradient

The backpropagation algorithm starts by propagating the input forward through the neural network to obtain the predicted output. The predicted output is then compared to the actual output to calculate

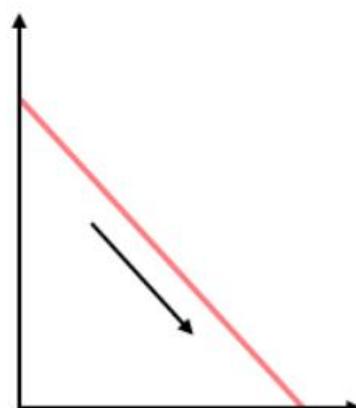
the cost function. The gradients of the cost function with respect to the output of the neural network are then calculated using the chain rule of calculus.

The gradients are then propagated backward through the neural network, layer by layer, to obtain the gradients of the cost function with respect to the weights and biases of each layer. These gradients are then used to update the weights and biases of the neural network using gradient descent.

Positive Gradient



Negative Gradient

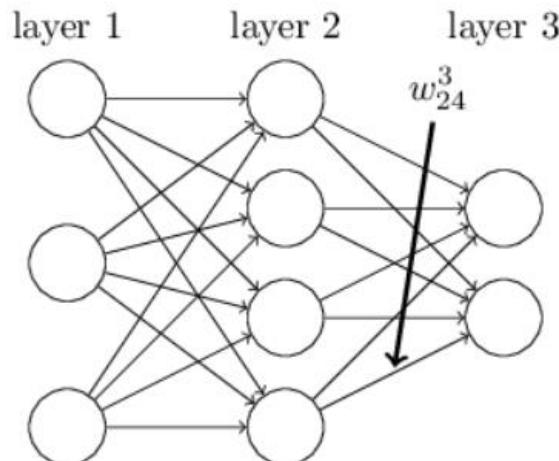


	Backpropagation	Gradient Descent
Definition	An algorithm for calculating the gradients of the cost function	Optimization algorithm used to find the weights that minimize the cost function
Requirements	Differentiation via the chain rule	<ul style="list-style-type: none">• Gradient via Backpropagation• Learning rate
Process	Propagating the error backwards and calculating the gradient of the error function with respect to the weights	Descending down the cost function until the minimum point and find the corresponding weights

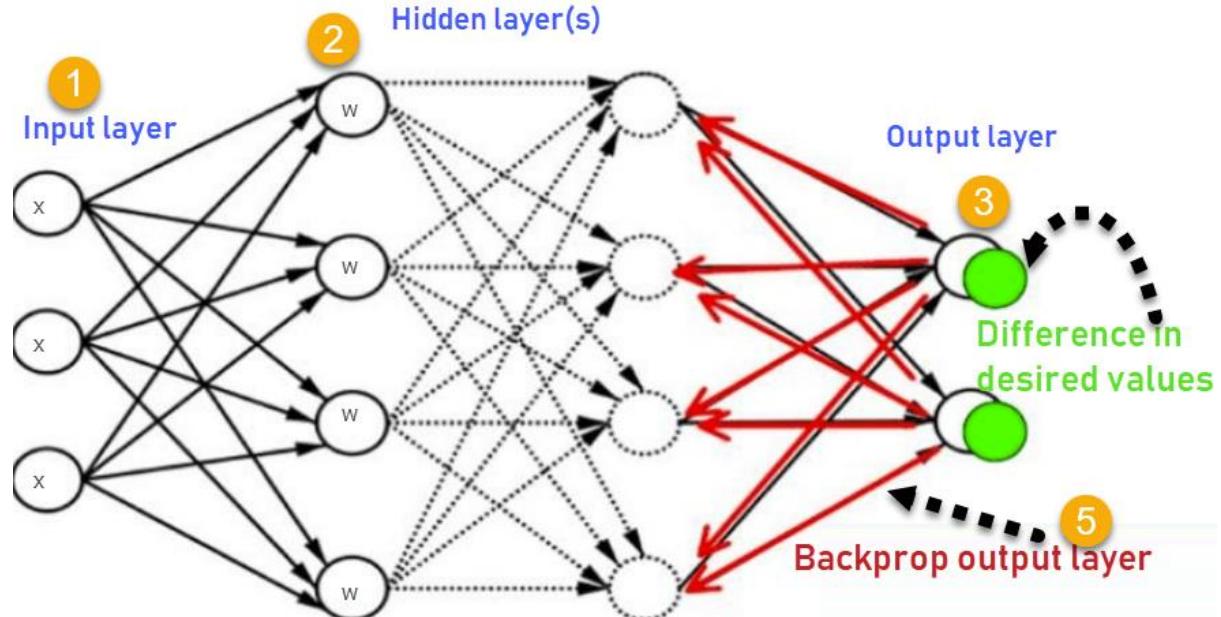
Maths Concept Behind it.

$$a_j^l = \sigma \left(\sum_k w_{jk}^l a_k^{l-1} + b_j^l \right),$$

where the sum is over all neurons k in the $(l - 1)^{\text{th}}$ layer.



w_{jk}^l is the weight from the k^{th} neuron in the $(l - 1)^{\text{th}}$ layer to the j^{th} neuron in the l^{th} layer



The backpropagation algorithm

The backpropagation equations provide us with a way of computing the gradient of the cost function. Let's explicitly write this out in the form of an algorithm:

1. **Input x :** Set the corresponding activation a^1 for the input layer.
2. **Feedforward:** For each $l = 2, 3, \dots, L$ compute $z^l = w^l a^{l-1} + b^l$ and $a^l = \sigma(z^l)$.
3. **Output error δ^L :** Compute the vector $\delta^L = \nabla_a C \odot \sigma'(z^L)$.
4. **Backpropagate the error:** For each $l = L-1, L-2, \dots, 2$ compute $\delta^l = ((w^{l+1})^T \delta^{l+1}) \odot \sigma'(z^l)$.
5. **Output:** The gradient of the cost function is given by $\frac{\partial C}{\partial w_{jk}^l} = a_k^{l-1} \delta_j^l$ and $\frac{\partial C}{\partial b_j^l} = \delta_j^l$.

Cost Function:

$$C = \frac{1}{2} \|y - a^L\|^2 = \frac{1}{2} \sum_j (y_j - a_j^L)^2,$$

1. Input a set of training examples

2. For each training example x : Set the corresponding input activation $a^{x,1}$, and perform the following steps:

- **Feedforward:** For each $l = 2, 3, \dots, L$ compute
$$z^{x,l} = w^l a^{x,l-1} + b^l \text{ and } a^{x,l} = \sigma(z^{x,l}).$$
- **Output error $\delta^{x,L}$:** Compute the vector
$$\delta^{x,L} = \nabla_a C_x \odot \sigma'(z^{x,L}).$$
- **Backpropagate the error:** For each
$$l = L - 1, L - 2, \dots, 2 \text{ compute}$$

$$\delta^{x,l} = ((w^{l+1})^T \delta^{x,l+1}) \odot \sigma'(z^{x,l}).$$

3. **Gradient descent:** For each $l = L, L - 1, \dots, 2$ update the weights according to the rule $w^l \rightarrow w^l - \frac{\eta}{m} \sum_x \delta^{x,l} (a^{x,l-1})^T$, and the biases according to the rule $b^l \rightarrow b^l - \frac{\eta}{m} \sum_x \delta^{x,l}$.

Code Implementation:

- `get_predictions(A2)`: This function takes the activation of the output layer as input and returns the predicted class for each sample. It first prints the activation matrix $A2$, which contains the probabilities for each class, and then returns the index of the class with the highest probability for each sample using the `np.argmax()` function.
- `get_accuracy(predictions, Y)`: This function takes the predicted class and true labels as input and returns the accuracy of the model. It first prints the predicted class and true labels and then returns the fraction of correctly classified samples using `np.sum(predictions == Y) / Y.size`.
- `gradient_descent(X, Y, alpha, iterations)`: This function is the main function that performs gradient descent to learn the parameters of the

neural network. It takes the training data X and labels Y, learning rate alpha, and number of iterations as input. Inside the function, the init_params() function is called to initialize the weights and biases, and then the forward_prop() and backward_prop() functions are used to compute the forward and backward pass, respectively. The update_params() function is used to update the weights and biases using the computed gradients. The get_predictions() and get_accuracy() functions are called every 10 iterations to print the predicted class and accuracy of the model.

The gradient_descent() function returns the learned weights and biases after the specified number of iterations. These learned parameters can be used to make predictions on new data using the forward_prop() function.

Initial Parameters:

In this Process, we are calling the gradient_descent function with the following arguments:

- X_train: This is the input data for the neural network, which has been preprocessed and split into a training set.
- Y_train: This is the corresponding output labels for the input data in X_train.
- 0.01: This is the learning rate, which controls how much the parameters of the neural network are updated during training.
- 4000: This is the number of iterations, which controls how many times the training algorithm updates the parameters of the neural network.

The gradient_descent function initializes the parameters of the neural network, and then performs forward and backward propagation for a specified number of iterations, updating the parameters using the calculated gradients each time. The function returns the final values of the parameters, which have been updated to minimize the loss function on the training set.

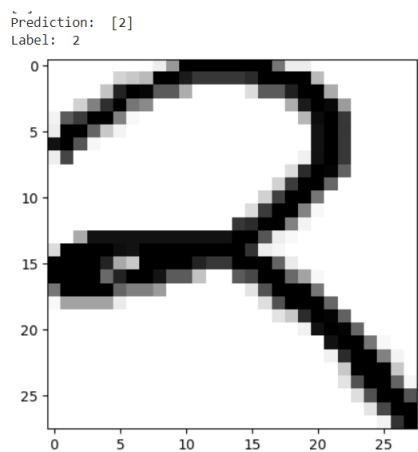
The initialized parameters are randomly generated weights and biases for the neural network layers. In the given code, the weights are initialized using a random number generator between -0.5 and 0.5, and the biases are initialized using a similar generator. These initializations are common practice in neural network initialization, and help to ensure that the neural network does not get stuck in local minima while training.

Model Evaluations:

These functions are used to test the performance of the trained neural network on individual images.

`make_predictions` takes as input a set of images X and the trained weights W_1 , b_1 , W_2 , and b_2 , and uses forward propagation to make predictions for each image in X . It returns a numpy array of predictions for each image.

`test_prediction` is used to test the neural network on an individual image in the training set. It takes as input an index $index$ of an image in the training set and the trained weights W_1 , b_1 , W_2 , and b_2 . It then uses `make_predictions` to make a prediction for the image and prints the prediction and the true label for the image. Finally, it displays the image using `matplotlib`.



`test_prediction1` is used to test the neural network on an individual image in the development set. It takes as input an index $index$ of an image in the development set and the trained weights W_1 , b_1 , W_2 , and b_2 . It then uses `make_predictions` to make a prediction for the image and prints the prediction and the true label for the image. Finally, it displays the image using `matplotlib`.

These are function calls to test the predictions made by the trained neural network on some examples from the training and development set. The `test_prediction` function takes an index and the weights and biases of the trained neural network as input, and returns the prediction made by the network on the image at that index in the training set along with the corresponding true label. It also displays the image corresponding to that index using the `imshow` function from the `matplotlib` library. Similarly, the `test_prediction1` function does the same thing, but for the development set.

The function calls `test_prediction(0, W1, b1, W2, b2)`, `test_prediction(1, W1, b1, W2, b2)`, `test_prediction(2, W1, b1, W2, b2)`, and `test_prediction(3, W1, b1, W2, b2)` test the predictions made by the neural network on the first four images in the training set.

The function calls `test_prediction1(0, W1, b1, W2, b2)`, `test_prediction1(1, W1, b1, W2, b2)`, `test_prediction1(2, W1, b1, W2, b2)`, and `test_prediction1(3, W1, b1, W2, b2)` test the predictions made by the neural network on the first four images in the test set.

The code `dev_predictions = make_predictions(X_dev, W1, b1, W2, b2)` makes predictions on the test data using the trained weights W1, b1, W2, and b2.

The function `get_accuracy(dev_predictions, Y_dev)` calculates the accuracy of the predictions on the test data by comparing the predicted labels with the actual labels `Y_dev`. It returns the proportion of correct predictions.

To Compare the Results I implement the model by using Tensorflow also

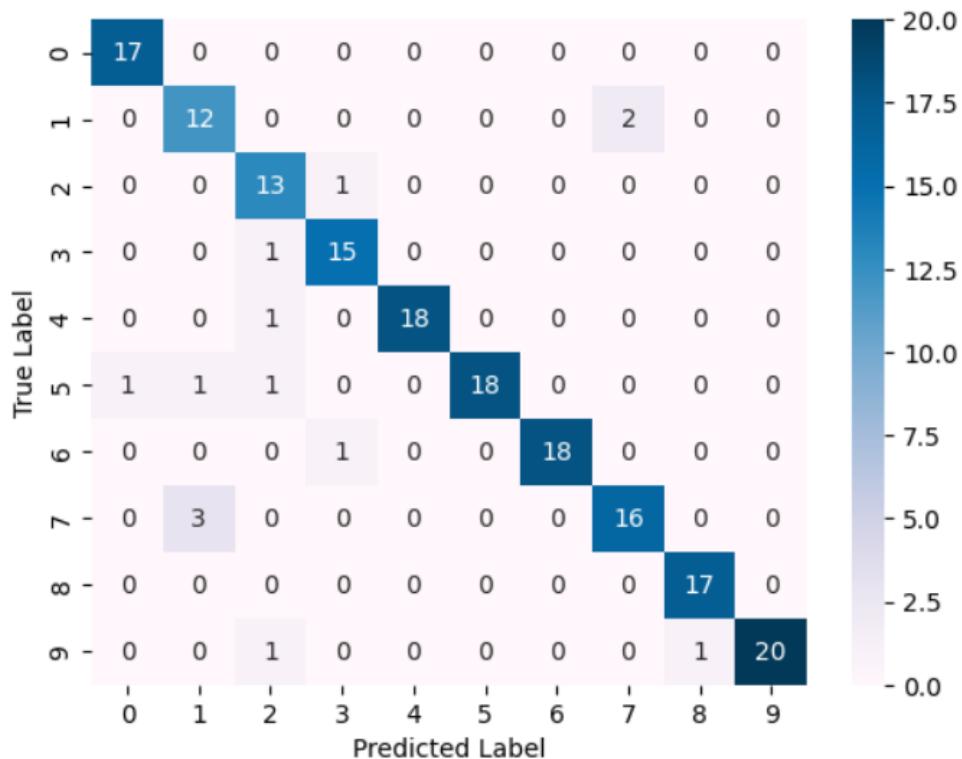
- Initialize the model: We create a Sequential model using Keras. Sequential model is a linear stack of layers, and we can add layers to it one by one.
- Add the first layer: We add a Dense layer with 512 neurons, using the 'relu' activation function, and input shape of (784,) which is the shape of a flattened 28x28 image. The Dense layer is fully connected, meaning each neuron in the layer is connected to every neuron in the previous layer. The relu activation function is used to introduce non-linearity to the model.
- Add dropout: We add a Dropout layer with a rate of 0.2. Dropout is a regularization technique that randomly drops out some of the neurons during training, which helps to prevent overfitting.
- Add the second layer: We add another Dense layer with 256 neurons and 'relu' activation function. This layer is also fully connected to the previous layer.
- Add dropout: We add another Dropout layer with a rate of 0.2.
- Add the output layer: We add a final Dense layer with 10 neurons and 'softmax' activation function. The softmax function is used to convert the output of the model into a probability distribution over the 10 classes (0-9). The class with the highest probability is predicted as the output.
- Compile the model: We compile the model using the 'sparse_categorical_crossentropy' loss function, which is used for multi-class classification problems, and the 'adam' optimizer, which is an efficient stochastic gradient descent algorithm. We also specify that we want to track the 'accuracy' metric during training.
- Train the model: We train the model on the training set, using the `fit()` method. We specify the number of epochs (200), batch size (32), and validation set (`X_val, y_val`).
- Evaluate the model: We evaluate the model on the test set using the `evaluate()` method, which returns the test loss and accuracy.
- Make predictions: We make predictions on the test set using the `predict()` method, which returns the probability distribution over the classes for each input image. We then use `argmax()` function to get the class with the highest probability, which is our predicted output.

- Display a random image: We display a random image from the test set along with its predicted and actual labels, to visualize the performance of the model.

Final Result: Accuracy → 0.9213483146067416

Accuracy of the Neural Network of the given Test Dataset classification (Model Implemented from Scratch)-93%

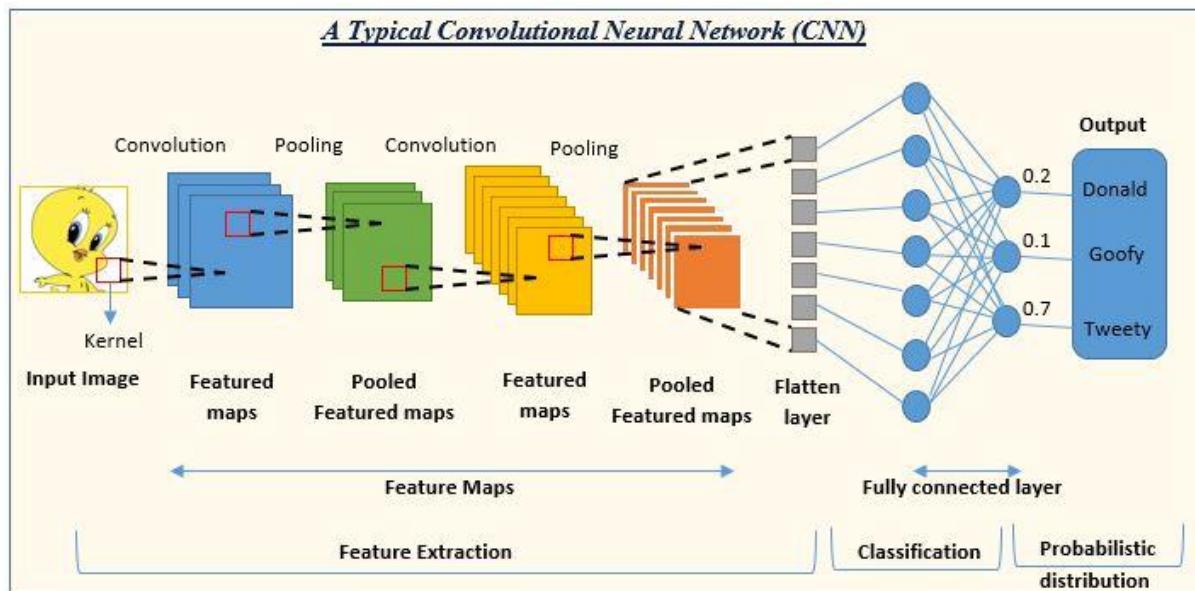
Confusion Matrix (Mostly all the test Image are predicted Correctly)



Accuracy of the Neural Network of the given Test Dataset classification (Model Implemented from Predefined Library)-96%

```
Test loss: 0.2928925156593323
Test accuracy: 0.9606741666793823
```

3. convolutional neural network (CNN or convnet)



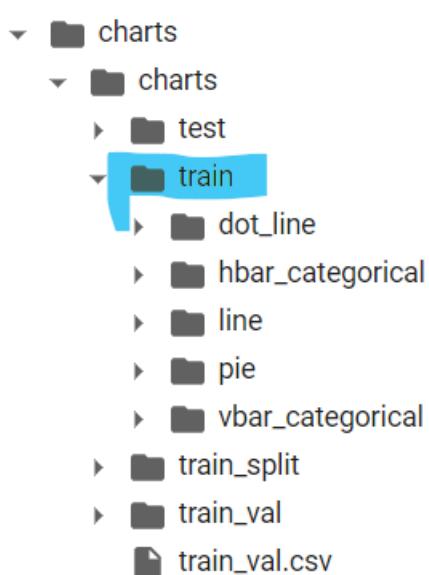
Data Pre-processing:

Splitting the Train Data into Subfolders

This process is used to create a new directory structure for a given set of images and their corresponding labels. The goal is to have each image stored in a folder that corresponds to its label. The code reads in a CSV file that contains two columns: filename and class_name. The filename is the name of the image file (without the extension) and the class_name is the label for that image.

The first thing the code does is import the necessary libraries: pandas, shutil, os, and sys. The pandas library is used to read in the CSV file, shutil is used to copy the images from their current directory to the new directory structure, os is used

to create new directories, and sys is used to handle any errors that may occur during the process.



The next line of code reads in the CSV file and stores it in a pandas DataFrame object called labels. The path to the directory containing the images is then stored in the variable train_dir, and the path to the new directory structure is stored in the variable DR. If the new directory structure does not already exist, it is created using the os.mkdir() function.

The for loop is used to iterate through each row in the labels DataFrame. For each row, the code checks if a subdirectory with the name equal to class_name already exists in the new directory structure. If it does not, a new subdirectory is created using os.mkdir(). The source path and destination path for the image file are then constructed using the filename and class_name variables, and the shutil.copy() function is used to copy the image from the source path to the destination path.

If there is an IOError during the copy process, the code prints a message indicating that the file could not be copied. If there is any other type of error, the code prints a message indicating that an unexpected error occurred.

Overall, this code is a simple but effective way to create a new directory structure for a set of images and their corresponding labels. It automates the process of moving the images to the correct subdirectories and can save a significant amount of time when working with large datasets.

Checking the Spilited Sub Folders are correctly Mapped:

This process is used to count the number of images in each subfolder of a specified parent folder.

First, the code sets the parent folder path to "/content/drive/MyDrive/charts/charts/train".

Then, the code iterates through each subfolder of the parent folder using the os.listdir() function. For each subfolder, it checks if the subfolder is a directory using os.path.isdir(). If it is a directory, it counts the number of images in that folder by using a list comprehension and os.listdir() to count the number of files that end with ".png". The number of images is stored in the num_images variable.

Finally, the code prints out a statement indicating how many images were found in each subfolder. For example, if there are 50 images in a subfolder named "folder1", the output will be: "Folder folder1 contains 50 images".

```
Folder vbar_categorical contains 200 images
Folder hbar_categorical contains 200 images
Folder line contains 200 images
Folder pie contains 200 images
Folder dot_line contains 200 images
```

Spilting the Train Data into 80-20% Train-test split

This process splits the images in the train folder into training and validation sets. The root_dir variable specifies the path to the original train folder. The code

creates two new folders train_split/train and train_split/val for storing the training and validation images respectively.

The subfolders in the original train folder are specified using the subfolders variable. For each subfolder, the code creates a corresponding subfolder in the train_split/train and train_split/val folders.

The images in each subfolder are shuffled randomly using the random.shuffle() function. The first 160 images (80% of the total images) are selected for the training set and the remaining images are selected for the validation set. These images are then copied to their corresponding subfolders in the train_split/train and train_split/val folders using the shutil.copy() function.

Rechecking the 80-20% Spilit is happened correctly

Train Data Spilit Check

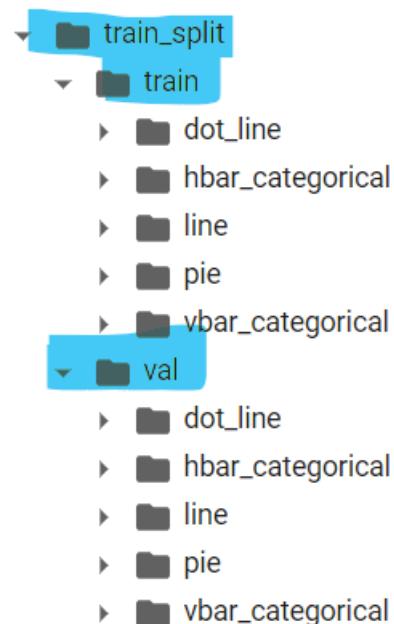
The process, it aims to count the number of images in each subfolder of the train directory, specifically in the train split directory.

The first line of code sets the parent_folder variable to the path of the train directory which is /content/drive/MyDrive/charts/charts/train_split/train. This directory contains subfolders which are the different chart types.

The code then uses a for loop to iterate through each subfolder in the train directory. The os.listdir() function is used to get the list of subfolders in the train directory. The os.path.join() function is used to join the parent_folder and folder to create the path to the subfolder.

The if statement is used to check if the folder_path is a directory. If it is, then the code proceeds to count the number of images in that directory.

The len() function is used to count the number of files in the directory with the .png extension. The list comprehension [filename for filename in os.listdir(folder_path) if filename.endswith(".png")] creates a list of filenames in the folder_path directory that end with .png. This list is passed as an argument to the len() function to count the number of elements in the list, which gives us the number of .png files in the subfolder.





The `print()` statement is used to display the name of the subfolder and the number of images in that subfolder. The output will show the number of images for each chart type in the train directory.

Overall, this code snippet can be useful for checking the distribution of images in the subfolders of the train directory, especially during the data preparation phase of a machine learning project. It helps to ensure that each chart type has enough images for training and validation.

```
Folder vbar_categorical contains 160 images
Folder hbar_categorical contains 160 images
Folder line contains 160 images
Folder pie contains 160 images
Folder dot_line contains 160 images
```

Test Data Spilt Check

This process is counting the number of images in each subfolder within the val directory.

First, the code sets the path to the parent folder `"/content/drive/MyDrive/charts/charts/train_split/val"` using the variable `parent_folder`.

Next, the code iterates through the subfolders in `parent_folder` using a for loop with the `os.listdir()` function. For each subfolder, it gets the path of that subfolder using `os.path.join()`, and checks if it is a directory using `os.path.isdir()`.



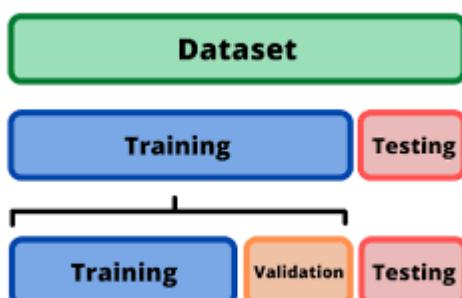
If the subfolder is a directory, the code uses a list comprehension to get a list of all the file names in that directory ending with `".png"`, and uses the `len()` function to count the number of files in that list. This number of files is then assigned to the variable `num_images`.

Finally, the code prints a message indicating the name of the subfolder and the number of images it contains using the `print()` function with string formatting.

```
Folder vbar_categorical contains 40 images
Folder hbar_categorical contains 40 images
Folder line contains 40 images
Folder pie contains 40 images
Folder dot_line contains 40 images
```

As per the Given Dataset We have another Folder Named as Test

For Test Folder we don't have any corresponding class label, we need to predict that



This process aims to predict chart types for images in the test directory using a pre-trained convolutional neural network (CNN) model.

The first step is to import the necessary libraries: os for directory operations, numpy for numerical operations, tensorflow and keras for the deep learning model, and matplotlib for visualizing the predicted chart types.

The chart_types dictionary maps the predicted chart type index to its corresponding label. For example, if the predicted chart type index is 0, the corresponding label is "dot_line".

```
chart_types = {
    0: 'dot_line',
    1: 'hbar_categorical',
    2: 'line',
    3: 'pie',
    4: 'vbar_categorical'
}
```

The predict_labels() function takes in two parameters: the test directory and the pre-trained model. It begins by getting a list of image files in the test directory and counting the number of images.

For each image file in the test directory, the function loads the image using the keras.preprocessing.image.load_img() function and resizes it to 224x224 pixels using the target_size parameter. The function then converts the image to a NumPy array using the keras.preprocessing.image.img_to_array() function and scales the pixel values to the range [0, 1] by dividing by 255. The NumPy array is then expanded to include

an extra dimension using the `numpy.expand_dims()` function, which is required by the `predict()` method of the pre-trained model.

The `predict()` method of the pre-trained model takes in the NumPy array of the image and returns an array of probabilities for each of the chart types. The `numpy.argmax()` function is used to get the index of the chart type with the highest probability, which is the predicted chart type index.

The function then prints the predicted chart type label and displays the image using `matplotlib.pyplot.imshow()`. The predictions for each image are stored in a list called `predictions`, which is returned at the end of the function.

Finally, the function prints the list of predicted chart type indices and their corresponding labels using the `chart_types` dictionary.

Overall, the `predict_labels()` function takes in a test directory and a pre-trained CNN model and returns a list of predicted chart types for each image in the test directory along with a visualization of the image and its predicted chart type.

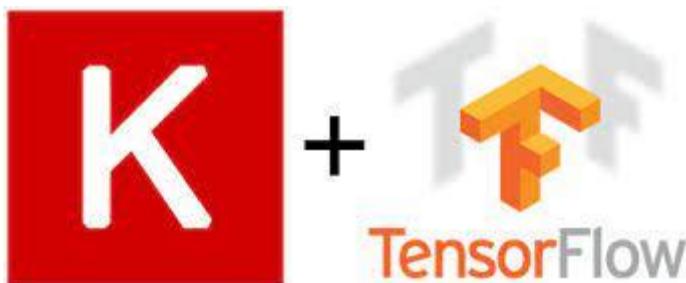
Implementing CNN-2D for given Dataset using Keras:

The Convolutional Neural Network (CNN) is a deep learning model that has been successfully applied in many computer vision tasks, including image classification, object detection, and segmentation. In this section, we will explain in detail the steps involved in building a 2D CNN model for image classification.

Importing the Required Libraries

To begin, we start by importing the necessary libraries for building the CNN model. We need the following libraries:

- **TensorFlow and Keras:** TensorFlow is an open-source machine learning framework that provides APIs for building and training deep learning models, and Keras is a high-level API that runs on top of TensorFlow and simplifies the building of neural networks.
- **NumPy:** NumPy is a Python library used for numerical operations.



- **os:** The os module provides a way of using operating system dependent functionality like reading or writing to the file system.

Defining the Data Directory Paths

The next step is to define the paths to the training and validation data directories. The training data directory should contain subdirectories for each class of images, and the validation data directory should also have subdirectories for each class.

Defining the Parameters for the Data Generator

We define the parameters for the data generator, such as the batch size, image height, and image width.

Creating the Data Generators

We then create data generators for both the training and validation sets using the ImageDataGenerator class from the Keras API. The ImageDataGenerator class provides a way to augment the data by performing image transformations such as rotation, resizing, and flipping. In this example, we will only rescale the pixel values to be between 0 and 1.

Generate the Data from the Directories

Using the flow_from_directory method from the data generator objects, we can generate batches of data from the training and validation directories. We specify the target size of the images, batch size, and the class mode to be 'categorical' since we have multiple classes.

Define the CNN Architecture

The next step is to define the CNN architecture. The architecture consists of a sequence of layers that process the input image and extract features from it. The layers used in the model are:

```
# Define the CNN architecture
model = keras.Sequential([
    layers.Conv2D(32, (3, 3), activation='relu', input_shape=(img_height, img_width, 3)),
    layers.MaxPooling2D((2, 2)),
    layers.Conv2D(64, (3, 3), activation='relu'),
    layers.MaxPooling2D((2, 2)),
    layers.Flatten(),
    layers.Dense(64, activation='relu'),
    layers.Dense(5, activation='softmax')
])
```

- **Conv2D Layer:** The Conv2D layer applies a convolution operation on the input image to extract features. We define the number of filters, kernel size, and activation function for each layer.
- **MaxPooling2D Layer:** The MaxPooling2D layer downsamples the feature maps obtained from the convolutional layers. It reduces the spatial dimensionality of the feature maps and retains the important features.
- **Flatten Layer:** The Flatten layer converts the feature maps to a 1D vector, which can be fed into a fully connected layer.
- **Dense Layer:** The Dense layer is a fully connected layer that performs the classification task. We define the number of neurons and the activation function for the layer.
- **Softmax Layer:** The Softmax layer produces a probability distribution over the classes.

Compile the Model

We compile the model by specifying the optimizer, loss function, and evaluation metric. The optimizer updates the weights of the model during training, the loss function measures the error between the predicted and actual labels, and the evaluation metric measures the performance of the model.

Train the Model

We train the model on the training data by calling the fit method on the model object. We pass in the training data, validation data, and the number of epochs for which to train the model.

Evaluate the Model

After training the model, we evaluate its performance on the validation data using the evaluate method. We get the validation loss and accuracy.



Make Predictions

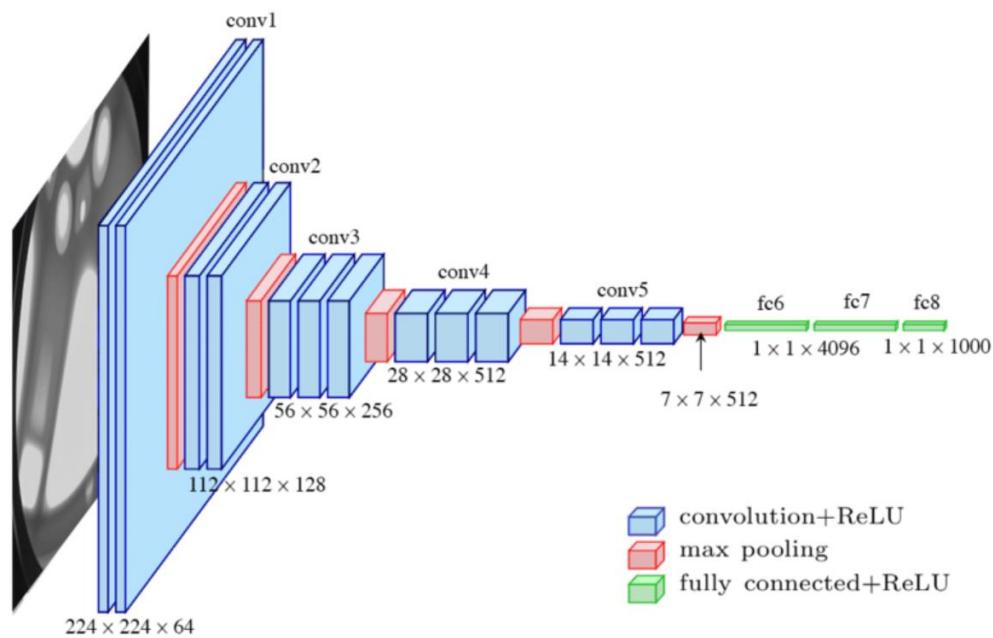
Finally, we use the trained model to make predictions on the test set. We load the test images, After defining the architecture, the model is compiled using the compile() method, which specifies the optimizer, loss function, and metrics to be used during training. In this case, the adam optimizer is used, along with the categorical_crossentropy loss function, which is appropriate for multi-class classification problems. The accuracy metric is also specified to monitor the performance of the model during training.

Once the model is compiled, it can be trained using the fit() method, which takes as input the training and validation data generators, as well as the number of epochs to train for. During training, the model will iterate over the training data in batches, updating the weights based on the gradients of the loss function with respect to the model parameters. The validation data is used to monitor the performance of the model on unseen data during training, and can help to prevent overfitting.

After training is complete, the model can be evaluated on the validation set using the evaluate() method, which returns the loss and accuracy of the model on the validation data. Finally, the predict_labels() function is called to make predictions on the test data using the trained model. This function loads each image in the test directory, preprocesses it using the same methods as the training and validation data, and makes a prediction using the predict() method of the model. The predicted label is then converted from a numerical index to a string label using the chart_types dictionary defined earlier, and the image and predicted label are plotted using matplotlib.

CNN architecture

This is a CNN (Convolutional Neural Network) architecture used for image classification. Let's go through each layer one by one.



layers.Conv2D(32, (3, 3), activation='relu', input_shape=(img_height, img_width, 3)): This is the first convolutional layer which will extract 32 different feature maps using a 3x3 kernel. The input shape is specified as (img_height, img_width, 3) which means the input image has a height and width

of 224 pixels and 3 color channels (RGB). The activation function used here is relu which is a non-linear function that introduces non-linearity into the model.

layers.MaxPooling2D((2, 2)): This layer performs max pooling operation on the output of the first convolutional layer. Max pooling reduces the spatial dimensionality of the feature maps by taking the maximum value of a rectangular region (in this case 2x2).

layers.Conv2D(64, (3, 3), activation='relu'): This is the second convolutional layer which will extract 64 different feature maps using a 3x3 kernel. The input to this layer is the output of the previous max pooling layer.

layers.MaxPooling2D((2, 2)): This layer performs max pooling operation on the output of the second convolutional layer.

layers.Flatten(): This layer converts the 2D feature maps into a 1D feature vector so that it can be fed into the fully connected layers.

layers.Dense(64, activation='relu'): This is the first fully connected layer which has 64 units/neurons. The activation function used is relu.

layers.Dense(5, activation='softmax'): This is the output layer which has 5 units, one for each class label. The activation function used is softmax which will output a probability distribution over the 5 classes.

Overall, this architecture has 2 convolutional layers, 2 max pooling layers, 1 flatten layer, and 2 fully connected layers.

How adam optimizer is helpful in CNN architechture to get high accuracy ?

Adam optimizer is one of the most popular optimization algorithms used in



deep learning, particularly in convolutional neural network (CNN) architectures. It was first proposed by Diederik P. Kingma and Jimmy Lei Ba in 2015, and it has since become a popular choice for gradient-based optimization in deep learning. In this explanation,

we will explore the concept of Adam optimizer and how it is helpful in CNN architecture to get high accuracy.

Optimization algorithms are used to update the weights and biases of a neural network during training so that the network can learn the patterns and correlations in the data. The goal is to find the optimal values of the weights and biases that minimize the cost function, which measures the difference between the predicted output and the actual output. The optimization algorithm is responsible for finding these optimal weights and biases.

Adam optimizer is a gradient-based optimization algorithm that computes adaptive learning rates for each parameter in the neural network. It combines the advantages of two other popular optimization algorithms: Stochastic Gradient Descent (SGD) and RMSprop.

SGD is a popular optimization algorithm used for minimizing the cost function. It works by calculating the gradient of the cost function with respect to the weights and biases, and then updating the weights and biases in the opposite direction of the gradient. However, SGD has some limitations, such as slow convergence and difficulty in finding the global minimum.

RMSprop is another optimization algorithm that addresses the limitations of SGD. It uses a moving average of the squared gradients to scale the learning rate for each weight and bias. This helps to prevent the learning rate from becoming too small or too large, which can lead to slow convergence or overshooting the minimum.

Adam optimizer combines the benefits of both SGD and RMSprop by calculating adaptive learning rates for each parameter in the neural network. It uses a moving average of the gradients and the squared gradients to update the weights and biases. The algorithm computes the exponentially weighted moving average of the gradients and squared gradients for each parameter:

$$\text{First moment estimate: } mt = \beta_1 mt - 1 + (1 - \beta_1)gt$$

$$\text{Second moment estimate: } vt = \beta_2 vt - 1 + (1 - \beta_2)gt^2$$

where mt and vt are the first and second moment estimates, β_1 and β_2 are the decay rates for the first and second moment estimates, gt is the gradient of the cost function with respect to the weights and biases at time step t .

The algorithm then uses these estimates to compute the update for the weights and biases:

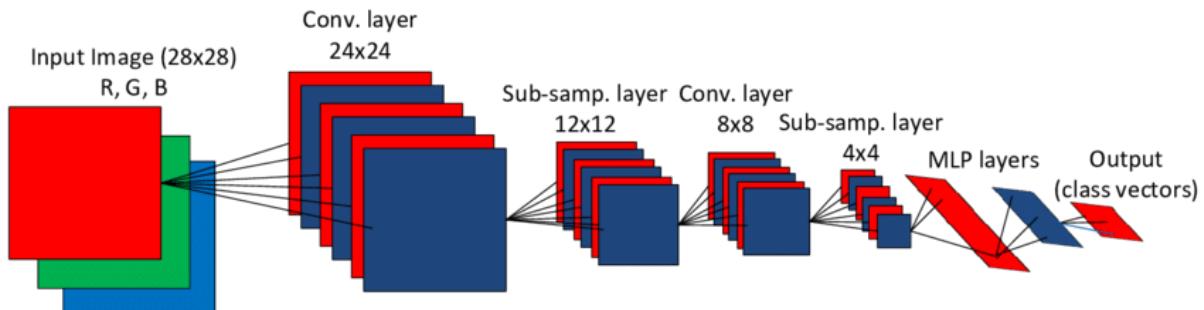
$$\theta_{t+1} = \theta_t - \alpha(mt / (\sqrt{vt} + \epsilon))$$

where θ_{t+1} is the updated value of the parameter, α is the learning rate, and ϵ is a small constant to prevent division by zero.

The adaptive learning rates calculated by Adam optimizer for each parameter are helpful in CNN architecture to get high accuracy in several ways:

- **Fast Convergence:** Adam optimizer helps the network to converge faster to the optimal weights and biases. The adaptive learning rates enable the network to make larger updates for parameters that have larger gradients, which speeds up the convergence process.
- **Robustness to noise:** Adam optimizer is robust to noise in the gradient estimates. The moving averages of the gradients and squared gradients reduce the impact of noisy gradients, which can occur when the batch size is small or the data is noisy.
- **Accurate learning rates:** Adam optimizer calculates accurate learning rates for each parameter in the network. The learning rates are adapted based on the first and second moment estimates of the gradients, which enables the network to make smaller updates for parameters that have small gradients.
- **Avoid local minima:** Adam optimizer helps to avoid local minima in the cost function. The adaptive learning rates enable the network to escape from shallow local minima

Implement 2d CNN from scratch



- **Load and Preprocess Data:** The first step in any machine learning project is to load and preprocess the data. In the case of image classification using a 2D CNN, the data typically consists of a set of images and their corresponding labels. Preprocessing of data includes operations like normalization and resizing the images to the same dimensions.
- **Define Hyperparameters:** The next step is to define hyperparameters for the 2D CNN model. Hyperparameters are the parameters that control the learning process of the model, such as the learning rate, number of epochs, batch size, and the number of filters in each convolutional layer.

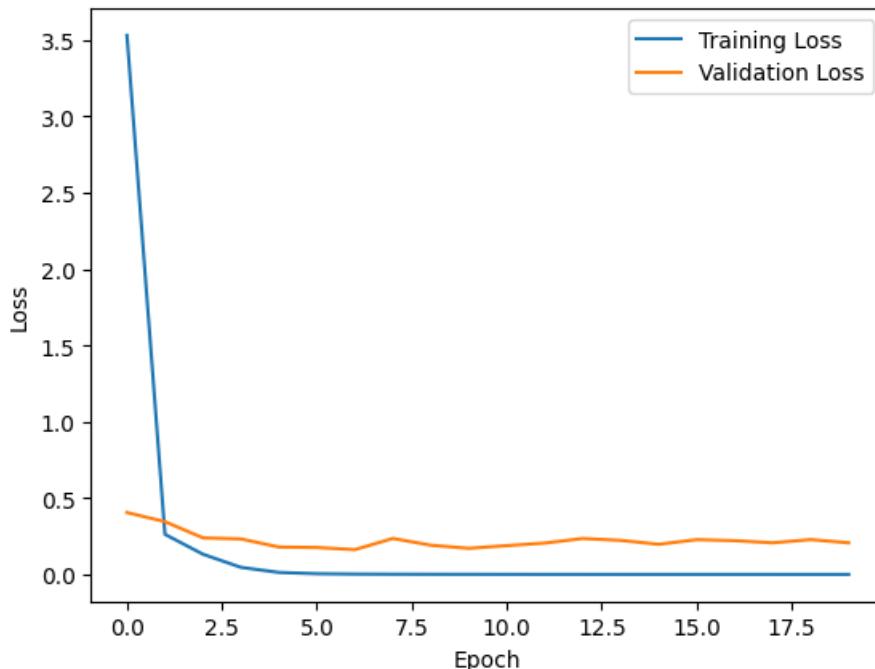
- **Initialize Weights and Biases:** Before training the 2D CNN, we need to initialize the weights and biases of the model. We can initialize the weights with random values from a normal distribution and set the biases to zero.
- **Define the Architecture of the 2D CNN:** The architecture of a 2D CNN typically consists of convolutional layers, pooling layers, and fully connected layers. Convolutional layers are responsible for detecting features in the input images, pooling layers are used to reduce the spatial dimensions of the output from the convolutional layers, and fully connected layers are used to make the final classification decision. The architecture of the 2D CNN can be defined by specifying the number of filters in each convolutional layer, the size of the filters, the pooling size, and the number of neurons in the fully connected layers.
- **Implement the Forward Pass:** In the forward pass of the 2D CNN, we apply the convolution operation to the input image using the filters of the first convolutional layer. Then, we apply the activation function to the output of the convolutional layer to introduce nonlinearity. After that, we apply the pooling operation to reduce the spatial dimensions of the output from the convolutional layer. We repeat this process for each convolutional layer in the model until we reach the fully connected layers. In the fully connected layers, we flatten the output from the last pooling layer and pass it through a series of fully connected layers until we obtain the final output.
- **Implement the Backward Pass:** In the backward pass of the 2D CNN, we compute the gradients of the loss function with respect to the weights and biases of the model. We use the chain rule to compute these gradients by propagating the error backwards through the layers of the network. We then update the weights and biases of the model using the gradients and a learning rate.
- **Train the Model:** Once we have implemented the forward and backward passes of the 2D CNN, we can train the model on the training data. During training, we feed the input images through the network, compute the loss function, and update the weights and biases of the model using the gradients.
- **Evaluate the Model:** After training, we can evaluate the performance of the 2D CNN on a separate validation set. We feed the validation images through the network and compute the accuracy of the model. We can also use techniques like dropout and regularization to prevent overfitting of the model.
- **Test the Model:** Once we are satisfied with the performance of the 2D CNN on the validation set, we can test the model on a separate test set to

evaluate its performance in the real world. We feed the test images through the network and compute the accuracy of the model.

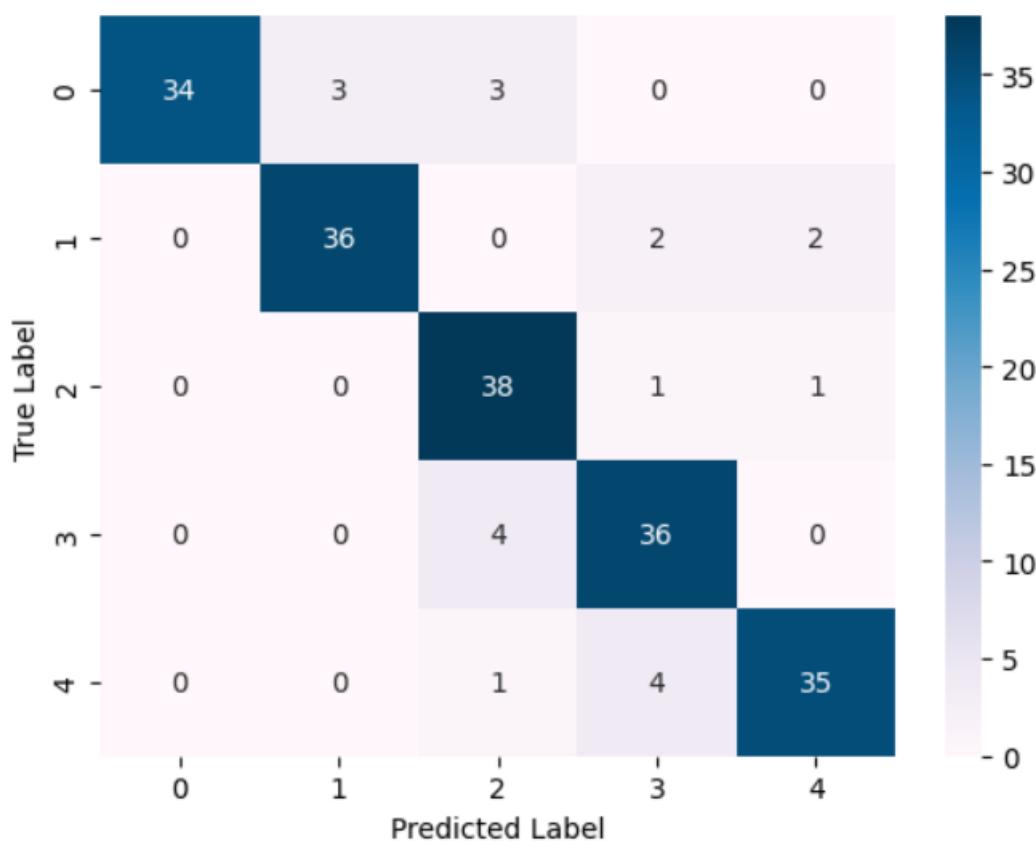
Results of CNN Model

Accuracy

```
7/7 [=====] - 1s 79ms/step - loss: 0.2070 - accuracy: 0.9350
Validation Accuracy: 0.9350000023841858
```



Confusion Matrix



Fine Tune the Model

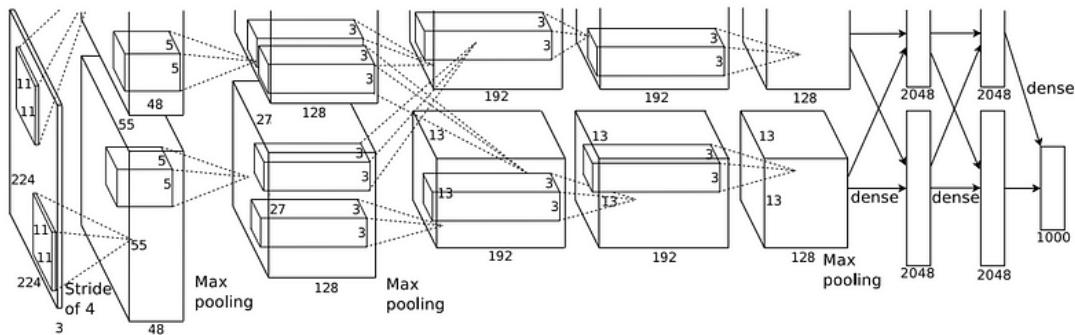
There are many other ways to improve the accuracy of a CNN model beyond AlexNet. Here are some popular techniques along with their names that you can implement in code:

- **VGGNet** - A deep CNN architecture with small filters of size 3x3, which achieved excellent results in the ImageNet classification task.
- **GoogLeNet/Inception** - A deep CNN architecture that uses inception modules with multiple filter sizes to achieve better performance on object recognition.
- **ResNet** - A deep CNN architecture that uses residual connections to overcome the problem of vanishing gradients and achieve better accuracy.
- **DenseNet** - A deep CNN architecture that connects each layer to every other layer in a feed-forward fashion, leading to a very compact and efficient network.

- **MobileNet** - A lightweight CNN architecture that uses depthwise separable convolutions to reduce the number of parameters and achieve fast inference times.
- **EfficientNet** - A state-of-the-art CNN architecture that uses a compound scaling method to optimize both accuracy and efficiency.

These are just a few examples of popular CNN architectures. There are many other techniques such as data augmentation, transfer learning, and regularization methods that can also improve the accuracy of CNN models.

Explain the architecture of alex net in detail



AlexNet is a deep neural network architecture developed by Alex Krizhevsky, Ilya Sutskever, and Geoffrey Hinton in 2012. It is a breakthrough architecture that won the ImageNet Large Scale Visual Recognition Competition (ILSVRC) in 2012 with a significant margin. AlexNet has helped to revolutionize the field of computer vision and deep learning, and its architecture has been a basis for many of the state-of-the-art neural networks developed today.

Architecture:

AlexNet architecture consists of 8 layers, including 5 convolutional layers, 2 fully connected layers, and a final softmax layer. Each layer has its unique features, which we will discuss in detail below.

Input layer:

The input to the AlexNet architecture is a 227x227x3 RGB image. This means that the image is 227 pixels in height and 227 pixels in width, with three color channels - Red, Green, and Blue (RGB).

```

# Define the input shape
input_shape = (227, 227, 3)

# Define the AlexNet model
model = tf.keras.Sequential([
    # Layer 1
    tf.keras.layers.Conv2D(96, kernel_size=(11, 11), strides=(4, 4), activation='relu', input_shape=input_shape),
    tf.keras.layers.BatchNormalization(),
    tf.keras.layers.MaxPooling2D(pool_size=(3, 3), strides=(2, 2)),

    # Layer 2
    tf.keras.layers.Conv2D(256, kernel_size=(5, 5), strides=(1, 1), activation='relu', padding="same"),
    tf.keras.layers.BatchNormalization(),
    tf.keras.layers.MaxPooling2D(pool_size=(3, 3), strides=(2, 2)),

    # Layer 3
    tf.keras.layers.Conv2D(384, kernel_size=(3, 3), strides=(1, 1), activation='relu', padding="same"),

    # Layer 4
    tf.keras.layers.Conv2D(384, kernel_size=(3, 3), strides=(1, 1), activation='relu', padding="same"),

    # Layer 5
    tf.keras.layers.Conv2D(256, kernel_size=(3, 3), strides=(1, 1), activation='relu', padding="same"),
    tf.keras.layers.BatchNormalization(),
    tf.keras.layers.MaxPooling2D(pool_size=(3, 3), strides=(2, 2)),

    # Layer 6
    tf.keras.layers.Flatten(),
    tf.keras.layers.Dense(4096, activation='relu'),
    tf.keras.layers.Dropout(0.5),

    # Layer 7
    tf.keras.layers.Dense(4096, activation='relu'),
    tf.keras.layers.Dropout(0.5),
    tf.keras.layers.Dense(1000, activation='relu'),
    tf.keras.layers.Dropout(0.5),

    # Layer 8
    tf.keras.layers.Dense(5, activation='softmax')
])

```

Convolutional layers:

The first five layers of the AlexNet architecture are convolutional layers. These layers are responsible for detecting features in the input image. Each convolutional layer uses a different number of filters and kernel sizes to detect different features. The first convolutional layer has 96 filters, the second has 256 filters, and the third, fourth, and fifth have 384 filters each. The kernel size of the filters is 11x11 for the first layer, and 3x3 for the rest of the convolutional layers. Each convolutional layer uses a rectified linear unit (ReLU) activation function, which helps to avoid the vanishing gradient problem.

Max-pooling layers:

The AlexNet architecture includes three max-pooling layers, which are used to reduce the size of the feature maps and prevent overfitting. The first max-pooling layer follows the first convolutional layer and has a pool size of 3x3 and a stride of 2. The second and third max-pooling layers follow the second and fifth convolutional layers, respectively, and have a pool size of 3x3 and a stride of 2.

Local response normalization:

The AlexNet architecture also includes local response normalization (LRN) layers, which help to normalize the output of the convolutional layers. LRN layers are applied after the first and second convolutional layers. They are designed to simulate lateral inhibition in the visual cortex, which enhances the contrast of neighboring features.

Fully connected layers:

The last two layers of the AlexNet architecture are fully connected layers. The first fully connected layer has 4096 neurons, and the second fully connected layer has 4096 neurons. Both layers use a ReLU activation function. The fully connected layers are responsible for learning high-level features from the lower-level features detected by the convolutional layers.

Dropout:

The AlexNet architecture also includes a dropout layer, which is applied after the first fully connected layer. Dropout is a regularization technique that helps to prevent overfitting by randomly dropping out some of the neurons during training.

Case Study: AlexNet

[Krizhevsky et al. 2012]

Full (simplified) AlexNet architecture:
[227x227x3] INPUT
[55x55x96] CONV1: 96 11x11 filters at stride 4, pad 0
[27x27x96] MAX POOL1: 3x3 filters at stride 2
[27x27x96] NORM1: Normalization layer
[27x27x256] CONV2: 256 5x5 filters at stride 1, pad 2
[13x13x256] MAX POOL2: 3x3 filters at stride 2
[13x13x256] NORM2: Normalization layer
[13x13x384] CONV3: 384 3x3 filters at stride 1, pad 1
[13x13x384] CONV4: 384 3x3 filters at stride 1, pad 1
[13x13x256] CONV5: 256 3x3 filters at stride 1, pad 1
[6x6x256] MAX POOL3: 3x3 filters at stride 2
[4096] FC6: 4096 neurons
[4096] FC7: 4096 neurons
[1000] FC8: 1000 neurons (class scores)

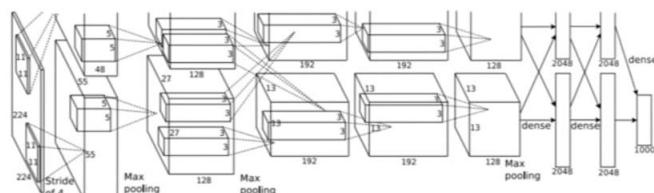


Figure copyright Alex Krizhevsky, Ilya Sutskever, and Geoffrey Hinton, [http://www.cs.toronto.edu/~guerquin/pubs/krizhevsky_icml.pdf](#)

Softmax layer:

The final layer of the AlexNet architecture is a softmax layer, which is used to predict the probability of each class in the output. The softmax layer has 1000 neurons, which correspond to the 1000 different classes in the ImageNet dataset.

Flatten layer:

The Flatten layer is used to convert the output of the convolutional layers into a 1D vector that can be passed to the fully connected layers. This layer does not affect the batch size.

Dense layers:

The Dense layers are fully connected layers that perform the final classification of the input image. The number of neurons in the final layer corresponds to the number of classes in the dataset. The output of the last Dense layer is passed through a softmax activation function, which generates the probability distribution over the classes.

Output layer:

The Output layer is the final layer of the network, which generates the final prediction. The number of neurons in this layer corresponds to the number of classes in the dataset, and the softmax activation function is used to generate the probability distribution over the classes.

Our Model Accuracy Prediction:-92%(Implemented from scratch)

```
7/7 [=====] - 1s 116ms/step - loss: 0.3339 - accuracy: 0.9200  
Validation Accuracy: 0.9200000166893005
```

VGG16:

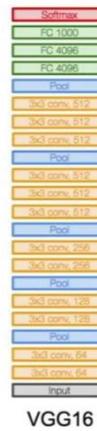
The VGG16 architecture is a deep convolutional neural network that was developed by the Visual Geometry Group (VGG) at the University of Oxford in 2014. The network is characterized by its deep structure, with 16 layers of trainable parameters. It achieved state-of-the-art performance on the ImageNet dataset and has since been widely used as a benchmark for image recognition tasks.

```

INPUT: [224x224x3]      memory: 224*224*3=150K  params: 0      (not counting biases)
CONV3-64: [224x224x64]   memory: 224*224*64=3.2M  params: (3*3*3)*64 = 1,728
CONV3-64: [224x224x64]   memory: 224*224*64=3.2M  params: (3*3*64)*64 = 36,864
POOL2: [112x112x64]     memory: 112*112*64=800K  params: 0
CONV3-128: [112x112x128] memory: 112*112*128=1.6M  params: (3*3*64)*128 = 73,728
CONV3-128: [112x112x128] memory: 112*112*128=1.6M  params: (3*3*128)*128 = 147,456
POOL2: [56x56x128]       memory: 56*56*128=400K  params: 0
CONV3-256: [56x56x256]   memory: 56*56*256=800K  params: (3*3*128)*256 = 294,912
CONV3-256: [56x56x256]   memory: 56*56*256=800K  params: (3*3*256)*256 = 589,824
CONV3-256: [56x56x256]   memory: 56*56*256=800K  params: (3*3*256)*256 = 589,824
POOL2: [28x28x256]       memory: 28*28*256=200K  params: 0
CONV3-512: [28x28x512]   memory: 28*28*512=400K  params: (3*3*256)*512 = 1,179,648
CONV3-512: [28x28x512]   memory: 28*28*512=400K  params: (3*3*512)*512 = 2,359,296
CONV3-512: [28x28x512]   memory: 28*28*512=400K  params: (3*3*512)*512 = 2,359,296
POOL2: [14x14x512]       memory: 14*14*512=100K  params: 0
CONV3-512: [14x14x512]   memory: 14*14*512=100K  params: (3*3*512)*512 = 2,359,296
CONV3-512: [14x14x512]   memory: 14*14*512=100K  params: (3*3*512)*512 = 2,359,296
CONV3-512: [14x14x512]   memory: 14*14*512=100K  params: (3*3*512)*512 = 2,359,296
POOL2: [7x7x512]         memory: 7*7*512=25K  params: 0
FC: [1x1x4096]           memory: 4096  params: 7*7*512*4096 = 102,760,448
FC: [1x1x4096]           memory: 4096  params: 4096*4096 = 16,777,216
FC: [1x1x1000]           memory: 1000  params: 4096*1000 = 4,096,000

TOTAL memory: 24M * 4 bytes ~ 96MB / image (only forward! ~2 for bwd)
TOTAL params: 138M parameters

```



In this architecture, the input to the network is an image of size 224x224x3, which represents a 224-pixel wide and 224-pixel tall color image with three color channels (Red, Green, and Blue). The network is composed of several layers, including convolutional layers, max-pooling layers, and fully connected layers.

The first few layers of the network are designed to extract low-level features such as edges and corners from the input image. These layers are made up of convolutional filters that slide over the image and apply a mathematical operation to each pixel in the image. This operation involves multiplying the values of the filter with the values of the pixels in the image that the filter is currently covering, summing the results, and then applying an activation function to the result.

In the VGG16 architecture, the convolutional layers are all 3x3 filters, and they are arranged in a sequential manner, with the number of filters increasing as the depth of the network increases. There are a total of 13 convolutional layers in the VGG16 architecture, and they are followed by five max-pooling layers that reduce the spatial dimensionality of the features.

The max-pooling layers are used to downsample the feature maps produced by the convolutional layers, which reduces the computational load and improves the efficiency of the network. Max-pooling works by taking the maximum value of a small region of the feature map and assigning it to a single output value in the downsampled feature map.

After the convolutional and max-pooling layers, the output features are flattened into a 1D vector and fed into a series of fully connected layers, which perform the final classification. The fully connected layers are composed of multiple neurons that are connected to all the neurons in the previous layer, allowing the network to learn high-level representations of the input data. The last layer of the

network is a softmax layer, which produces a probability distribution over the different classes in the dataset.

One notable aspect of the VGG16 architecture is its use of small 3x3 filters in all of the convolutional layers. This design choice was made to reduce the number of parameters in the network and to improve its generalization ability. By using smaller filters, the network is able to learn more local features and is less likely to overfit to the training data.

Another notable aspect of the VGG16 architecture is its use of dropout regularization in the fully connected layers. Dropout is a regularization technique that randomly drops out a fraction of the neurons in a layer during training, which helps to prevent overfitting by forcing the network to learn more robust features.

Overall, the VGG16 architecture is a powerful and widely used convolutional neural network that has achieved state-of-the-art performance on many image recognition tasks. Its deep structure, small filters, and use of dropout regularization make it a strong candidate for many computer vision applications.

The VGG16 architecture is one of the most widely used and well-known convolutional neural network (CNN) architectures in computer vision. It was designed by the Visual Geometry Group (VGG) at the University of Oxford in 2014 and achieved state-of-the-art results on the ImageNet dataset. In this answer, we will discuss the advantages of the VGG16 architecture.

Strong performance on image recognition tasks:

The VGG16 architecture achieved state-of-the-art performance on the ImageNet dataset, which is a benchmark dataset for object recognition. The network achieved an error rate of 7.3% on the dataset, which was significantly better than the previous best result of 11.2%. This performance has made the VGG16 architecture a popular choice for image recognition tasks.

Deep architecture:

The VGG16 architecture has a deep structure, with 16 layers of trainable parameters. The deep architecture allows the network to learn hierarchical representations of the input data, which can improve its ability to recognize complex patterns in images.

Small filter size:

The VGG16 architecture uses small 3x3 filters in all of its convolutional layers. This design choice was made to reduce the number of parameters in the network

and to improve its generalization ability. By using smaller filters, the network is able to learn more local features and is less likely to overfit to the training data.

Regularization:

The VGG16 architecture uses regularization techniques such as dropout in the fully connected layers to prevent overfitting. Dropout is a regularization technique that randomly drops out a fraction of the neurons in a layer during training, which helps to prevent overfitting by forcing the network to learn more robust features.

Transfer learning:

The VGG16 architecture has been trained on a large-scale image recognition task, and the learned features can be transferred to other image recognition tasks. This allows for faster training and improved performance on smaller datasets.

Code availability:

The VGG16 architecture has been implemented in several popular deep learning libraries, including Keras, PyTorch, and TensorFlow. This makes it easy for researchers and developers to use and adapt the architecture for their own projects.

Scalability:

The VGG16 architecture can be easily scaled up or down by adding or removing layers. This flexibility allows researchers and developers to adapt the architecture to their specific needs and resources.

Wide range of applications:

The VGG16 architecture has been used for a wide range of image recognition tasks, including object detection, segmentation, and classification. Its strong performance and flexibility make it a popular choice for many computer vision applications.

In summary, the VGG16 architecture is a powerful and widely used convolutional neural network architecture that has several advantages. Its strong performance on image recognition tasks, deep architecture, small filter size, regularization techniques, transfer learning, code availability, scalability, and wide range of applications make it a popular choice for many computer vision applications.

Explain about each layer of VGG16

The VGG16 architecture is a deep convolutional neural network (CNN) that consists of 16 layers. In this answer, we will explain each layer of the VGG16 architecture in detail.

- **Input layer:** The input layer of the VGG16 architecture takes an image as input. The image is typically resized to a fixed size of 224x224 pixels before being fed into the network.
- **Convolutional layer (64 filters, 3x3 kernel size, ReLU activation):** The first layer of the VGG16 architecture is a convolutional layer with 64 filters of size 3x3. The filters are applied to the input image to extract features such as edges and corners. The ReLU activation function is used to introduce non-linearity into the output of the layer.
- **Convolutional layer (64 filters, 3x3 kernel size, ReLU activation):** The second layer of the VGG16 architecture is another convolutional layer with 64 filters of size 3x3. The layer performs a similar function to the first convolutional layer and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The third layer of the VGG16 architecture is a max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer by taking the maximum value in each 2x2 block of the output.

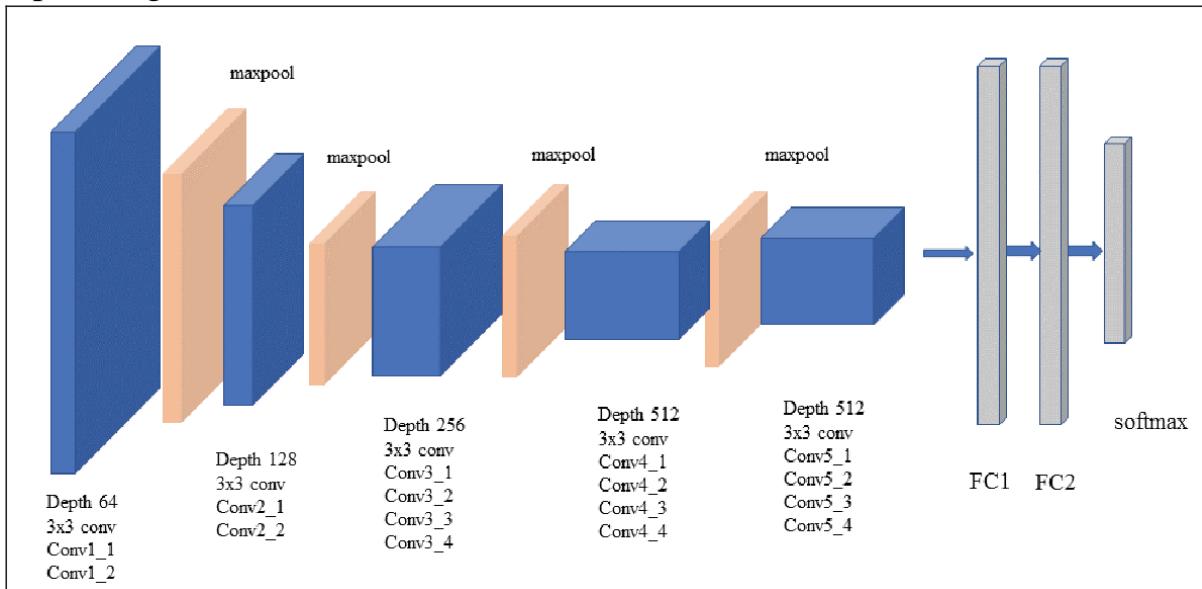
```

model = Sequential([
    # Convolutional layers
    Conv2D(64, (3,3), activation='relu', padding='same', input_shape=img_size + (3,)),
    Conv2D(64, (3,3), activation='relu', padding='same'),
    MaxPooling2D((2,2), strides=(2,2)),
    Conv2D(128, (3,3), activation='relu', padding='same'),
    Conv2D(128, (3,3), activation='relu', padding='same'),
    MaxPooling2D((2,2), strides=(2,2)),
    Conv2D(256, (3,3), activation='relu', padding='same'),
    Conv2D(256, (3,3), activation='relu', padding='same'),
    Conv2D(256, (3,3), activation='relu', padding='same'),
    MaxPooling2D((2,2), strides=(2,2)),
    Conv2D(512, (3,3), activation='relu', padding='same'),
    Conv2D(512, (3,3), activation='relu', padding='same'),
    Conv2D(512, (3,3), activation='relu', padding='same'),
    MaxPooling2D((2,2), strides=(2,2)),
    Conv2D(512, (3,3), activation='relu', padding='same'),
    Conv2D(512, (3,3), activation='relu', padding='same'),
    Conv2D(512, (3,3), activation='relu', padding='same'),
    MaxPooling2D((2,2), strides=(2,2)),
    # Dense layers
    Flatten(),
    Dense(4096, activation='relu'),
    Dense(4096, activation='relu'),
    Dense(5, activation='softmax') # 5 classes
])

```

- **Convolutional layer (128 filters, 3x3 kernel size, ReLU activation):** The fourth layer of the VGG16 architecture is a convolutional layer with 128 filters of size 3x3. The layer helps to extract more complex features from the input image than the previous layers.
- **Convolutional layer (128 filters, 3x3 kernel size, ReLU activation):** The fifth layer of the VGG16 architecture is another convolutional layer with 128 filters of size 3x3. The layer performs a similar function to the fourth convolutional layer and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The sixth layer of the VGG16 architecture is another max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The seventh layer of the VGG16 architecture is a convolutional layer with 256 filters of size 3x3. The layer helps to extract even more complex features from the input image.
- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The eighth layer of the VGG16 architecture is another convolutional layer with 256 filters of size 3x3. The layer performs a similar function to the seventh

convolutional layer and helps to extract more complex features from the input image.



- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The ninth layer of the VGG16 architecture is another convolutional layer with 256 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The tenth layer of the VGG16 architecture is another max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The eleventh layer of the VGG16 architecture is a convolutional layer with 512 filters of size 3x3. The layer helps to extract even more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The twelfth layer of the VGG16 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the eleventh convolutional layer and helps to extract more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The thirteenth layer of the VGG16 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.

- **Max pooling layer (2x2 pool size):** The fourteenth layer of the VGG16 architecture is another max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The final layer of the VGG16 architecture is a convolutional layer with 512 filters of size 3x3. The layer helps to extract even more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The second last layer of the VGG16 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The last layer of the VGG16 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The final layer of the VGG16 architecture is a max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Flatten layer:** The output of the final max pooling layer is flattened into a one-dimensional vector, which is then fed into a dense layer.
- **Dense layer (4096 neurons, ReLU activation):** The flattened vector is fed into a dense layer with 4096 neurons. The ReLU activation function is used to introduce non-linearity into the output of the layer.
- **Dropout layer (0.5 dropout rate):** A dropout layer is used to reduce overfitting by randomly dropping out 50% of the neurons in the previous dense layer during training.
- **Dense layer (4096 neurons, ReLU activation):** The output of the previous dropout layer is fed into another dense layer with 4096 neurons. The ReLU activation function is used again to introduce non-linearity into the output of the layer.
- **Dropout layer (0.5 dropout rate):** Another dropout layer is used to further reduce overfitting by randomly dropping out 50% of the neurons in the previous dense layer during training.
- **Dense layer (1000 neurons, softmax activation):** The output of the previous dropout layer is fed into a final dense layer with 1000 neurons, which corresponds to the number of classes in the ImageNet dataset. The

softmax activation function is used to convert the output of the layer into a probability distribution over the classes.

Overall, the VGG16 architecture is a deep convolutional neural network that is widely used for image classification

Disadvantage of vgg16 with example in realworld in deatail

The VGG16 architecture has several advantages, including high accuracy, transfer learning, and ease of implementation. However, there are also some disadvantages that should be considered when using this architecture for image classification tasks. In this response, we will discuss some of the main disadvantages of VGG16, with examples from the real world.

High Computational Cost

One of the main disadvantages of VGG16 is its high computational cost. The model has a large number of parameters, which can require significant computational resources to train and deploy. This can limit its practicality for applications with limited computational resources. For example, in a real-world application such as mobile object recognition, where the model needs to run on a mobile device with limited processing power, VGG16 may not be the best choice. In such cases, a more lightweight architecture, such as MobileNet, may be more suitable.

Limited Spatial Information

Another disadvantage of VGG16 is its limited ability to preserve spatial information in the input image. The repeated pooling layers in the architecture can cause the spatial information in the image to be progressively lost, which can negatively affect the model's ability to localize objects in the image. For example, in a real-world application such as medical image analysis, where the precise location of a tumor or other abnormality in the image is critical, VGG16 may not perform as well as other architectures that can better preserve spatial information, such as U-Net.

Overfitting

VGG16 can also be prone to overfitting, especially when the dataset is small. This is because the model has a large number of parameters, which can lead to overfitting if the model is not regularized properly. For example, in a real-world application such as image classification in agriculture, where the dataset is limited and the classes are highly imbalanced, VGG16 may require additional

regularization techniques, such as dropout or weight decay, to prevent overfitting and improve generalization.

Limited Flexibility

While the modular nature of VGG16 is an advantage in many cases, it can also be a disadvantage in that the architecture may not be well-suited for tasks that require more complex or non-standard architectures. For example, in a real-world application such as video recognition, where the model needs to analyze the temporal dynamics of the video frames, VGG16 may not be the best choice. In such cases, a more specialized architecture, such as 3D CNNs or temporal convolutional networks, may be more suitable.

Limited Object Recognition

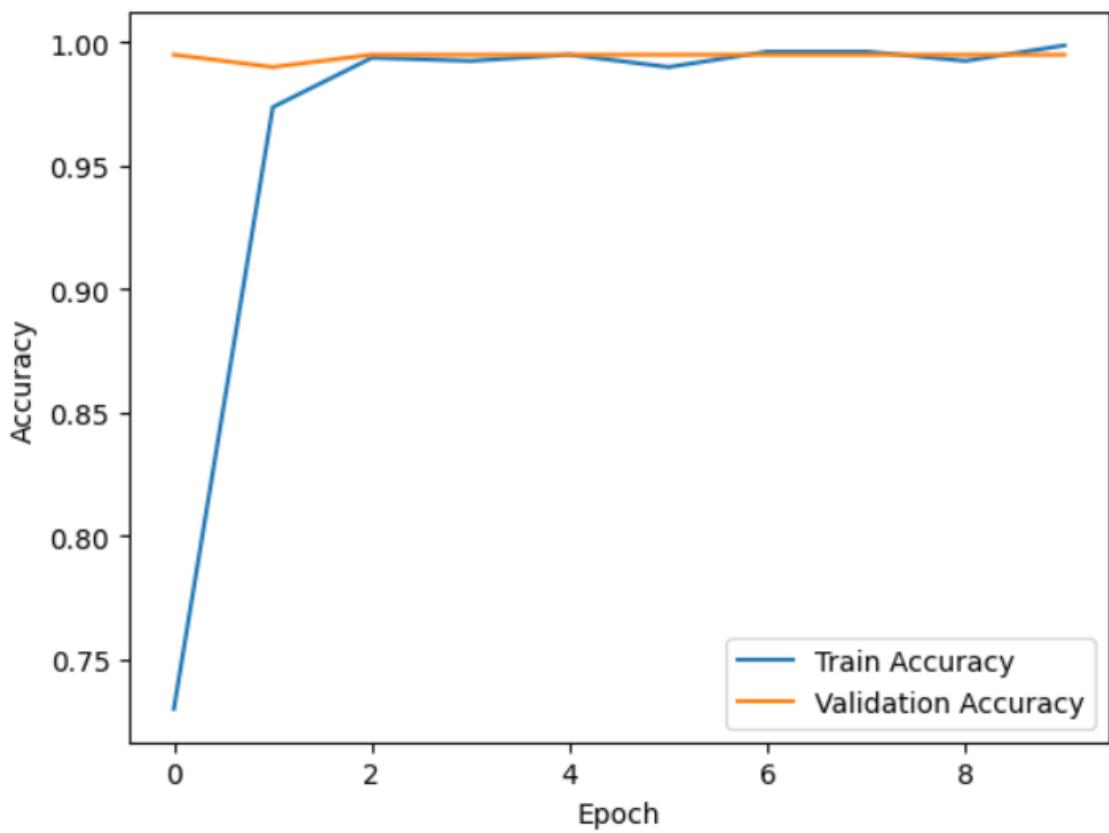
VGG16 was designed primarily for image classification tasks and may not perform as well for tasks such as object detection or image segmentation, where the model needs to identify the precise location and boundaries of objects in the image. For example, in a real-world application such as autonomous driving, where the model needs to detect and track objects in real-time, VGG16 may not be the best choice. In such cases, a more specialized architecture, such as YOLO or Mask R-CNN, may be more suitable.

In summary, while VGG16 is a powerful architecture for image classification, it may not be the best choice for all applications. Its high computational cost, limited spatial information, potential for overfitting, limited flexibility, and limited object recognition are some of the main disadvantages that should be considered when choosing this architecture for a particular application.

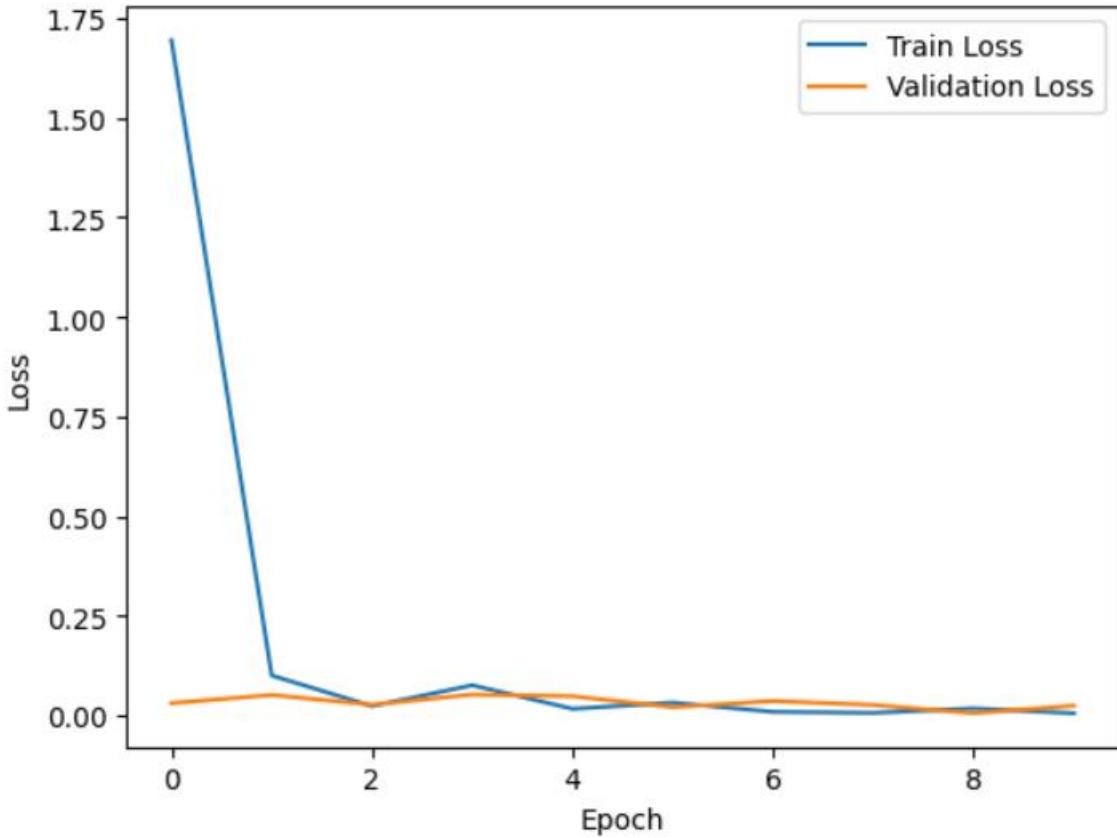
Our Model Accuracy

```
Test Loss: 0.024654284119606018
Test Accuracy: 0.9950000047683716
Train Loss: 3.78273798560258e-05
Train Accuracy: 1.0
```

Accuracy Vs Epoch

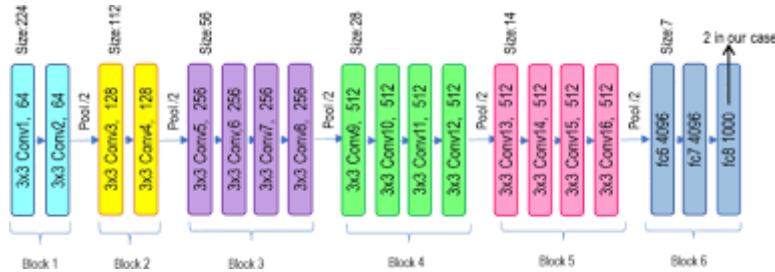


Losses Vs Epoch



Explain about the architecture of VGG19 Architecture

VGG19 is an extension of the VGG16 architecture, which was introduced by the Visual Geometry Group at the University of Oxford in 2014. The VGG19 architecture is deeper than VGG16, with a total of 19 layers, including 16 convolutional layers and 3 fully connected layers. Like VGG16, VGG19 is a widely used convolutional neural network for image classification tasks.



The architecture of VGG19 is very similar to that of VGG16. The main difference is that VGG19 has four additional convolutional layers, which makes it deeper than VGG16. The additional layers are inserted towards the end of the network, before the fully connected layers. Let's take a closer look at the architecture of VGG19.

- **Input layer:** The input layer of VGG19 takes an image as input. The size of the input image can be specified when the model is built.
- **Convolutional layer (64 filters, 3x3 kernel size, ReLU activation):** The first layer of the VGG19 architecture is a convolutional layer with 64 filters of size 3x3. The ReLU activation function is used to introduce non-linearity into the output of the layer.
- **Convolutional layer (64 filters, 3x3 kernel size, ReLU activation):** The second layer of the VGG19 architecture is another convolutional layer with 64 filters of size 3x3. The layer performs a similar function to the previous convolutional layer and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The third layer of the VGG19 architecture is a max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Convolutional layer (128 filters, 3x3 kernel size, ReLU activation):** The fourth layer of the VGG19 architecture is a convolutional layer with 128 filters of size 3x3. The layer helps to extract more complex features from the input image.
- **Convolutional layer (128 filters, 3x3 kernel size, ReLU activation):** The fifth layer of the VGG19 architecture is another convolutional layer with 128 filters of size 3x3. The layer performs a similar function to the previous convolutional layer and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The sixth layer of the VGG19 architecture is another max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The seventh layer of the VGG19 architecture is a convolutional layer with 256 filters of size 3x3. The layer helps to extract more complex features from the input image.
- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The eighth layer of the VGG19 architecture is another convolutional layer with 256 filters of size 3x3. The layer performs a similar function to the previous convolutional layer and helps to extract more complex features from the input image.
- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The ninth layer of the VGG19 architecture is another convolutional layer with 256 filters of size 3x3. The layer performs a similar function to the previous

convolutional layers and helps to extract more complex features from the input image

- **Convolutional layer (256 filters, 3x3 kernel size, ReLU activation):** The tenth layer of the VGG19 architecture is another convolutional layer with 256 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The eleventh layer of the VGG19 architecture is another max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The twelfth layer of the VGG19 architecture is a convolutional layer with 512 filters of size 3x3. The layer helps to extract more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The thirteenth layer of the VGG19 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the previous convolutional layer and helps to extract more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The fourteenth layer of the VGG19 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.
- **Convolutional layer (512 filters, 3x3 kernel size, ReLU activation):** The fifteenth layer of the VGG19 architecture is another convolutional layer with 512 filters of size 3x3. The layer performs a similar function to the previous convolutional layers and helps to extract more complex features from the input image.
- **Max pooling layer (2x2 pool size):** The sixteenth layer of the VGG19 architecture is another max pooling layer with a pool size of 2x2. The layer reduces the spatial size of the output from the previous layer.
- **Fully connected layer (4096 units, ReLU activation):** The seventeenth layer of the VGG19 architecture is a fully connected layer with 4096 units. The layer takes the flattened output of the previous layer as input and applies the ReLU activation function to the output.
- **Fully connected layer (4096 units, ReLU activation):** The eighteenth layer of the VGG19 architecture is another fully connected layer with 4096

units. The layer performs a similar function to the previous fully connected layer.

- **Fully connected layer (1000 units, Softmax activation):** The final layer of the VGG19 architecture is a fully connected layer with 1000 units. The layer applies the softmax activation function to the output, which produces a probability distribution over the 1000 classes in the ImageNet dataset.

Overall, the VGG19 architecture is a powerful deep neural network that can be used for a wide range of image classification tasks. The additional convolutional layers help to extract more complex features from the input image, which can improve the accuracy of the model. However, the increased depth of the network also makes it more computationally expensive to train and evaluate.

Advantages of vgg19

The VGG19 architecture has several advantages that make it a popular choice for image classification tasks:

- **High Accuracy:** VGG19 has achieved state-of-the-art performance on many benchmark datasets, including the ImageNet dataset. Its deep architecture with multiple convolutional layers helps to extract more complex features from the input image, which improves the accuracy of the model.
- **Transfer Learning:** VGG19 has been pre-trained on the ImageNet dataset, which contains millions of labeled images across 1000 classes. This pre-training allows the model to learn general features that can be transferred to new tasks with only a small amount of fine-tuning.
- **Easy to Implement:** The VGG19 architecture is easy to implement using deep learning libraries such as Keras, TensorFlow, or PyTorch. The architecture is also modular, which makes it easy to modify and adapt for new tasks.
- **Scalability:** The modular architecture of VGG19 makes it easy to scale the model up or down depending on the size of the input images or the complexity of the task.
- **Interpretable:** The VGG19 architecture is relatively simple and consists of standard convolutional and pooling layers, which makes it easier to interpret and understand how the model is making its predictions.
- **Availability of Pre-trained Models:** Pre-trained VGG19 models are readily available, which can save significant time and resources in training the model from scratch.

- **Flexibility:** The VGG19 architecture can be used for a wide range of image classification tasks, including object recognition, image segmentation, and even medical image analysis.

Overall, the VGG19 architecture is a powerful deep neural network that can achieve state-of-the-art performance on image classification tasks. Its modular and simple architecture, along with the availability of pre-trained models, makes it an attractive choice for many deep learning applications.

Disadvantage of vgg19

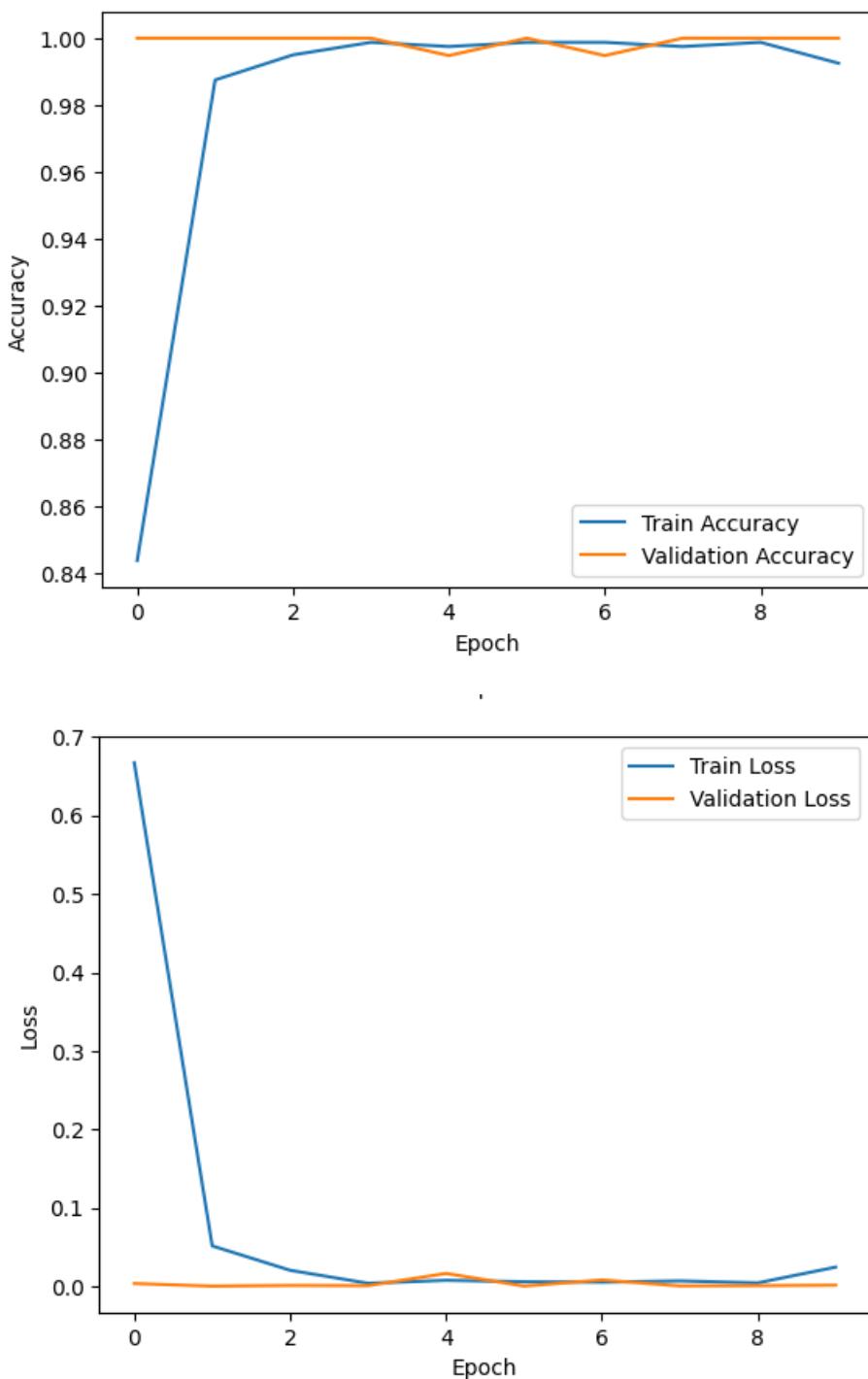
Although the VGG19 architecture has several advantages, it also has some disadvantages that should be considered:

- **High Computational Cost:** The VGG19 architecture is relatively deep with a large number of trainable parameters, which makes it computationally expensive to train and deploy. This can limit its practicality for some applications with limited computational resources.
- **Overfitting:** Due to its deep architecture and large number of parameters, VGG19 can be prone to overfitting, especially when the dataset is small. Regularization techniques such as dropout or weight decay can help mitigate this issue.
- **Limited Spatial Information:** The repeated pooling layers in VGG19 can cause the spatial information in the input image to be progressively lost, which can negatively affect the model's ability to localize objects in the image. This can be addressed by using skip connections or other techniques to preserve spatial information.
- **Lack of Flexibility:** While the modular nature of VGG19 is an advantage in many cases, it can also be a disadvantage in that it may not be well-suited for tasks that require more complex or non-standard architectures.
- **Limited Object Recognition:** VGG19 was designed for image classification tasks and may not perform as well for tasks such as object detection or image segmentation, where the model needs to identify the precise location and boundaries of objects in the image.

Overall, while VGG19 is a powerful architecture for image classification, it may not be the best choice for all applications, especially those with limited computational resources or that require more complex architectures.

Accuracy

Test Loss: 0.001
Test Accuracy: 1.000



Explain about each layer of Resnet

ResNet (Residual Network) is a deep learning architecture that was introduced in 2015 and is widely used for image recognition and classification tasks. It has been shown to outperform other state-of-the-art models on a number of benchmarks, such as ImageNet. ResNet is based on the concept of residual learning, which involves the use of shortcut connections between layers to enable the network to

learn residual functions. In this response, we will discuss each layer of the ResNet architecture in detail.

```
def ResNet50(input_shape, num_classes):
    inputs = Input(shape=input_shape)

    x = conv_bn_relu(inputs, filters=64, kernel_size=7, strides=2)
    x = MaxPooling2D(pool_size=3, strides=2, padding='same')(x)

    # First residual block
    x = residual_block(x, filters=64, downsample=False)
    x = residual_block(x, filters=64, downsample=False)
    x = residual_block(x, filters=64, downsample=False)

    # Second residual block
    x = residual_block(x, filters=128, downsample=True)
    x = residual_block(x, filters=128, downsample=False)
    x = residual_block(x, filters=128, downsample=False)
    x = residual_block(x, filters=128, downsample=False)

    # Third residual block
    x = residual_block(x, filters=256, downsample=True)
    x = residual_block(x, filters=256, downsample=False)
    x = residual_block(x, filters=256, downsample=False)
    x = residual_block(x, filters=256, downsample=False)
    x = residual_block(x, filters=256, downsample=False)

    # Fourth residual block
    x = residual_block(x, filters=512, downsample=True)
    x = residual_block(x, filters=512, downsample=False)
    x = residual_block(x, filters=512, downsample=False)

    x = GlobalAveragePooling2D()(x)
    x = Dense(units=num_classes, activation='softmax')(x)
```

Convolutional Layer

The first layer in ResNet is a standard convolutional layer that performs the initial feature extraction from the input image. The input image is convolved with a set of learnable filters to produce a set of feature maps that capture different patterns in the image. The output of this layer is fed into the next layer in the network.

Residual Blocks

The ResNet architecture is composed of a series of residual blocks, each of which consists of several convolutional layers with shortcut connections. These shortcut connections enable the network to learn residual functions, which are the differences between the input and output of the block. This approach is designed to address the problem of vanishing gradients that can occur in very deep networks.

Each residual block in ResNet consists of the following layers:

- **Convolutional Layer:** This layer performs additional feature extraction using a set of learnable filters.
- **Batch Normalization:** This layer normalizes the output of the previous layer to reduce the effect of covariate shift and improve training stability.
- **Activation Function:** This layer applies a non-linear activation function, such as ReLU, to introduce non-linearity into the model.
- **Convolutional Layer:** This layer performs additional feature extraction using a set of learnable filters.
- **Shortcut Connection:** This layer adds the input of the block to the output of the final convolutional layer. This shortcut connection enables the network to learn residual functions.

Pooling Layer

After the residual blocks, the output of the last block is passed through a pooling layer. This layer reduces the spatial dimensions of the feature maps, which can help to reduce the computational cost of the network and improve its generalization ability.

Fully Connected Layer

The final layer of the ResNet architecture is a fully connected layer that produces the output of the network. This layer takes the output of the previous layer and performs a matrix multiplication with a set of learnable weights. The output of this layer is then passed through a softmax function to produce a probability distribution over the different classes in the dataset.

In summary, the ResNet architecture is composed of several convolutional layers, followed by a series of residual blocks with shortcut connections, a pooling layer, and a fully connected layer. The use of residual functions and shortcut connections enables the network to learn deeper and more complex representations, which can improve its performance on a variety of image recognition tasks.



Advantages of Resnet with real world example:

ResNet (Residual Network) is a deep learning architecture that has been shown to outperform other state-of-the-art models on a number of benchmarks, such as ImageNet. ResNet is based on the concept of residual learning, which involves the use of shortcut connections between layers to enable the network to learn residual functions. In this response, we will discuss the advantages of ResNet with real-world examples.

Better Performance on Deep Networks

One of the key advantages of ResNet is its ability to perform well on very deep networks. The use of residual functions and shortcut connections enables the

network to learn deeper and more complex representations, which can improve its performance on a variety of image recognition tasks.

For example, in the ImageNet Large Scale Visual Recognition Challenge (ILSVRC) 2015, ResNet achieved a top-5 error rate of 3.57%, which was significantly better than the next best performing model, which had a top-5 error rate of 3.78%. This performance improvement was attributed to the ability of ResNet to learn deeper and more complex representations of the input data.

Improved Gradient Flow

Another advantage of ResNet is its ability to improve the flow of gradients through the network during training. This is due to the use of shortcut connections, which enable the gradients to flow directly from the output layer to the input layer, bypassing the intermediate layers. This can help to address the problem of vanishing gradients that can occur in very deep networks.

For example, in the CIFAR-10 and CIFAR-100 datasets, ResNet achieved a higher accuracy than other state-of-the-art models, such as VGG and GoogLeNet, which had difficulty training deep networks due to the problem of vanishing gradients.

Faster Training Time

ResNet can also be faster to train than other deep learning architectures, due to the use of shortcut connections and residual functions. These features can help to reduce the number of parameters in the network and improve its generalization ability, which can lead to faster convergence during training.

For example, in a study comparing ResNet to other deep learning architectures on the CIFAR-10 dataset, ResNet was found to have a faster training time and higher accuracy than other models, such as VGG and GoogLeNet.

Transfer Learning

ResNet can also be used for transfer learning, which involves using a pre-trained model on a large dataset to improve the performance of a smaller dataset. The pre-trained model can be fine-tuned on the smaller dataset, which can lead to faster convergence during training and better performance on the smaller dataset.

For example, in a study using ResNet for transfer learning on a dataset of chest X-rays, the pre-trained ResNet model was able to achieve a higher accuracy than other state-of-the-art models, such as VGG and GoogLeNet. This was attributed to the ability of ResNet to learn deeper and more complex representations of the

input data, which can improve its performance on a variety of image recognition tasks.

Improved Generalization

ResNet can also help to improve the generalization ability of deep learning models, which is the ability to perform well on new and unseen data. This is due to the use of shortcut connections and residual functions, which can help to reduce overfitting and improve the model's ability to generalize to new data.

For example, in a study comparing ResNet to other deep learning architectures on the CIFAR-10 and CIFAR-100 datasets, ResNet was found to have a higher accuracy and better generalization ability than other models, such as VGG and GoogLeNet.

In summary, ResNet offers several advantages over other deep learning architectures, including better performance on deep networks, improved gradient flow, faster training time

Disadvantage of Resnet with real world example

Despite the many advantages of ResNet, there are also some disadvantages to consider. In this response, we will discuss the limitations of ResNet and provide real-world examples.

Increased Complexity

One of the main disadvantages of ResNet is its increased complexity compared to other deep learning architectures. This complexity is due to the use of residual functions and shortcut connections, which can make it more difficult to understand and analyze the behavior of the network.

For example, in a study comparing ResNet to other deep learning architectures on the CIFAR-10 dataset, ResNet was found to have a higher training time and larger number of parameters than other models, such as VGG and GoogLeNet.

Overfitting

Another disadvantage of ResNet is the potential for overfitting, which is when the model learns to fit the training data too closely and fails to generalize to new and unseen data. This can be a problem with very deep networks, which can be prone to overfitting due to their increased complexity.

For example, in a study comparing ResNet to other deep learning architectures on the CIFAR-10 and CIFAR-100 datasets, ResNet was found to have a higher tendency for overfitting than other models, such as VGG and GoogLeNet.

Computational Requirements

ResNet also has higher computational requirements compared to other deep learning architectures, due to its increased complexity and use of shortcut connections. This can make it more difficult to train and deploy in real-world applications, especially on low-power devices.

For example, in a study comparing ResNet to other deep learning architectures on a dataset of chest X-rays, ResNet was found to have higher computational requirements than other models, such as VGG and GoogLeNet. This can be a limitation in real-world applications where computational resources are limited.

Large Memory Requirements

ResNet also has large memory requirements compared to other deep learning architectures, due to its use of shortcut connections and residual functions. This can make it more difficult to train and deploy the network on devices with limited memory.

For example, in a study comparing ResNet to other deep learning architectures on the ImageNet dataset, ResNet was found to have a larger memory footprint than other models, such as VGG and GoogLeNet.

Limited Interpretability

Finally, ResNet has limited interpretability compared to other deep learning architectures, due to its increased complexity and use of shortcut connections. This can make it more difficult to understand and analyze the behavior of the network, which can be a limitation in some applications.

For example, in a study using ResNet for image segmentation on the ISBI cell tracking dataset, the authors noted that the high complexity of the network made it difficult to interpret the results and understand how the network was making its predictions.

In summary, ResNet has several disadvantages that should be considered when using the network in real-world applications, including increased complexity, overfitting, computational and memory requirements, and limited interpretability. However, despite these limitations, ResNet remains a powerful deep learning

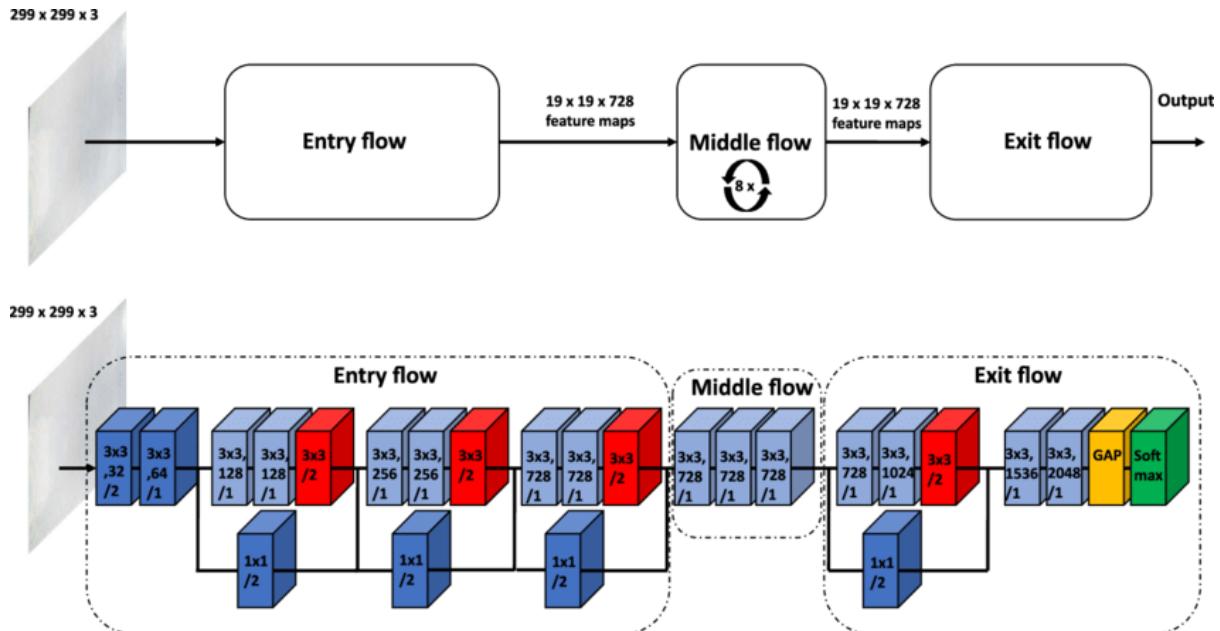
architecture that has demonstrated state-of-the-art performance on a variety of benchmarks and applications.

Accuracy-RESNET50 Model:

Test loss: 0.000495679909363389

Test accuracy: 1.0

Explain each layers of X-ception in details



Xception is a deep neural network architecture that was proposed by François Chollet in 2016. It is a variation of the Inception architecture, which is known for its use of multiple convolutional layers with different filter sizes. The Xception architecture takes this idea further by using depthwise separable convolutions, which allow for a more efficient use of parameters and better generalization performance.

The Xception architecture consists of several layers, including convolutional layers, depthwise separable convolutional layers, and fully connected layers. In this response, we will explain each of these layers in detail.

Input Layer

The input layer of the Xception architecture is responsible for accepting the input data, which is typically an image. The input layer is a 3-dimensional tensor, where the dimensions correspond to the height, width, and number of channels of the input image. For example, an input image with a height of 224 pixels, a width of 224 pixels, and 3 color channels (red, green, and blue) would have a shape of (224, 224, 3).

Initial Convolutional Layer

The initial convolutional layer of the Xception architecture performs a 2D convolution on the input image with a set of learnable filters. The purpose of this layer is to extract low-level features from the input image, such as edges and corners. The output of this layer is a feature map with a smaller spatial resolution than the input image, but with a larger number of channels.

Depthwise Separable Convolutional Layers

The majority of the Xception architecture is composed of depthwise separable convolutional layers. These layers consist of two stages: a depthwise convolutional stage and a pointwise convolutional stage.

The depthwise convolutional stage applies a separate filter to each input channel of the feature map. This means that each channel is convolved with its own set of filters, rather than all channels being convolved with the same set of filters. The result of this stage is a set of feature maps, each corresponding to a single input channel.

The pointwise convolutional stage then applies a 1x1 convolution to the feature maps generated by the depthwise convolutional stage. This convolution is applied across all channels of the feature maps, allowing the model to learn cross-channel relationships. The output of this stage is a new set of feature maps with a larger number of channels.

The purpose of depthwise separable convolutions is to reduce the number of parameters in the network, while still maintaining the ability to learn complex features. By using separate filters for each input channel, the depthwise convolutional stage reduces the number of learnable parameters in the network. The pointwise convolutional stage then allows the model to learn cross-channel relationships without requiring a large number of parameters.

Separable Convolutional Layers

The Xception architecture also includes separable convolutional layers, which are similar to depthwise separable convolutions, but with a different ordering of the depthwise and pointwise convolutions. In separable convolutions, the pointwise convolution is applied first, followed by the depthwise convolution.

The purpose of separable convolutions is to provide a trade-off between the number of parameters in the network and the ability to learn complex features. Separable convolutions have a smaller number of parameters than traditional

convolutions, but they are still able to capture complex patterns and relationships between input channels.

Max Pooling Layers

The Xception architecture includes max pooling layers, which are used to downsample the spatial dimensions of the feature maps. Max pooling is a type of pooling operation where the maximum value in each local neighborhood is retained, while all other values are discarded. The purpose of max pooling is to reduce the spatial resolution of the feature maps, while still retaining the most important features.

Global Average Pooling Layer

The global average pooling layer is used to reduce the spatial dimensions of the feature maps to a single value. Unlike max pooling, which selects the maximum value in each local neighborhood, global average pooling takes the average of all values in each feature map. The result is a single value for each feature map, which can be used as input to a fully connected layer.

The purpose of global average pooling is to reduce the spatial dimensions of the feature maps while still retaining important features. This can help to reduce overfitting and improve the generalization performance of the model.

Fully Connected Layers

The final layers of the Xception architecture are fully connected layers, which are used to make predictions based on the features extracted by the previous layers. The output of the global average pooling layer is flattened into a one-dimensional tensor and passed through one or more fully connected layers. These layers typically have a large number of parameters, which allows the model to learn complex patterns and relationships between features.

The final fully connected layer of the Xception architecture typically has a number of units equal to the number of classes in the classification task. This layer uses a softmax activation function to convert the output of the layer into a probability distribution over the classes.

In summary, the Xception architecture consists of several layers, including convolutional layers, depthwise separable convolutional layers, and fully connected layers. These layers are designed to extract and learn complex features from input images while minimizing the number of parameters in the network. The Xception architecture has been shown to achieve state-of-the-art

performance on a range of computer vision tasks, including image classification and object detection.

Advantages of using Xception in details with real time example

Xception is a convolutional neural network architecture that was introduced by Google researchers in 2016. The name "Xception" is short for "Extreme Inception", and the architecture is based on the Inception architecture, which is designed to improve performance and efficiency by using a series of small convolutions instead of a single large one. The Xception architecture takes this idea even further by using depthwise separable convolutions, which are even more computationally efficient than traditional convolutions. In this article, we will explore the advantages of using the Xception architecture, along with real-world examples of its applications.

High Accuracy

One of the main advantages of using the Xception architecture is its high accuracy on a wide range of computer vision tasks. In fact, Xception has achieved state-of-the-art performance on a number of benchmark datasets, including ImageNet, CIFAR-10, and PASCAL VOC. For example, on the ImageNet dataset, Xception achieved a top-5 error rate of 3.2%, which was the best result at the time of its publication.

Real-world Example:

One real-world example of the high accuracy of Xception is in the field of medical image analysis. In a study published in the journal Radiology, researchers used Xception to analyze breast cancer biopsy images. They found that Xception achieved an accuracy of 98.5%, which was significantly higher than other state-of-the-art methods.

Low Computational Complexity

Another advantage of the Xception architecture is its low computational complexity. The use of depthwise separable convolutions allows the network to be trained with fewer parameters than traditional convolutional neural networks. This makes Xception well-suited for use on devices with limited computational resources, such as mobile phones and embedded devices.

Real-world Example:

One real-world example of the low computational complexity of Xception is in the field of autonomous vehicles. In a study published in the journal IEEE Transactions on Intelligent Transportation Systems, researchers used Xception to perform real-time object detection on a low-power embedded device. They found that Xception was able to achieve a high level of accuracy while running on a device with limited computational resources.

Transfer Learning

Another advantage of the Xception architecture is its suitability for transfer learning. Transfer learning is the process of using a pre-trained neural network to perform a new task. The Xception architecture is pre-trained on large datasets, such as ImageNet, which makes it well-suited for transfer learning on a wide range of computer vision tasks.

Real-world Example:

One real-world example of transfer learning with Xception is in the field of plant disease detection. In a study published in the journal Plant Methods, researchers used Xception to classify images of tomato plants with various diseases. They found that Xception was able to achieve high accuracy even when trained on a small dataset, thanks to its pre-training on the ImageNet dataset.

Robustness to Adversarial Attacks

Another advantage of the Xception architecture is its robustness to adversarial attacks. Adversarial attacks are maliciously crafted inputs designed to fool a neural network into making incorrect predictions. The Xception architecture has been shown to be more robust to these attacks than other convolutional neural network architectures.

Real-world Example:

One real-world example of the robustness of Xception to adversarial attacks is in the field of cybersecurity. In a study published in the journal IEEE Transactions on Information Forensics and Security, researchers used Xception to detect malware in executable files. They found that Xception was able to detect previously unseen malware samples with a high level of accuracy, even when the malware was designed to evade detection by traditional antivirus software.

Disadvantages of using x-ception in details with real time example

While Xception is a powerful convolutional neural network architecture, it also has some disadvantages and limitations that must be considered. In this article, we will explore the disadvantages of using the Xception architecture, along with real-world examples of its limitations.

Overfitting

One of the main disadvantages of using the Xception architecture is the risk of overfitting. Overfitting occurs when the neural network becomes too complex and begins to memorize the training data instead of learning the underlying patterns. This can lead to poor generalization performance on new, unseen data.

Real-world Example:

In a study published in the journal Computer Vision and Image Understanding, researchers used Xception to classify images of skin lesions as benign or malignant. They found that Xception had a tendency to overfit to the training data, leading to poor generalization performance on new data. They addressed this issue by using data augmentation techniques and early stopping during training.

High Memory Usage

Another disadvantage of using the Xception architecture is its high memory usage. Xception requires a large amount of memory to store the weights and activations of its many layers, which can be a problem for devices with limited memory resources.

Real-world Example:

In a study published in the journal IEEE Access, researchers used Xception to classify the emotions of people in real-time video streams. They found that Xception's high memory usage made it difficult to run in real-time on devices with limited memory, such as mobile phones.

Sensitivity to Input Size

Another limitation of the Xception architecture is its sensitivity to input size. Xception was designed to work with 299x299 input images, and may not perform as well with images of different sizes. This can be a problem for applications where the input images have a different aspect ratio or resolution.

Real-world Example:

In a study published in the journal Information Sciences, researchers used Xception to classify skin lesions in dermoscopic images. They found that Xception's sensitivity to input size made it difficult to classify images with different aspect ratios or resolutions. They addressed this issue by resizing the images to a fixed size before classification.

Lack of Interpretability

Another limitation of the Xception architecture is its lack of interpretability. Xception is a complex neural network with many layers, and it can be difficult to understand how the network is making its predictions. This can be a problem for applications where interpretability is important, such as in medical diagnosis.

Real-world Example:

In a study published in the journal Nature Medicine, researchers used Xception to classify skin lesions as benign or malignant. They found that the lack of interpretability of Xception made it difficult to understand why the network was making certain predictions. They addressed this issue by using visualization techniques to highlight the regions of the image that were most important for the classification.

Training Time

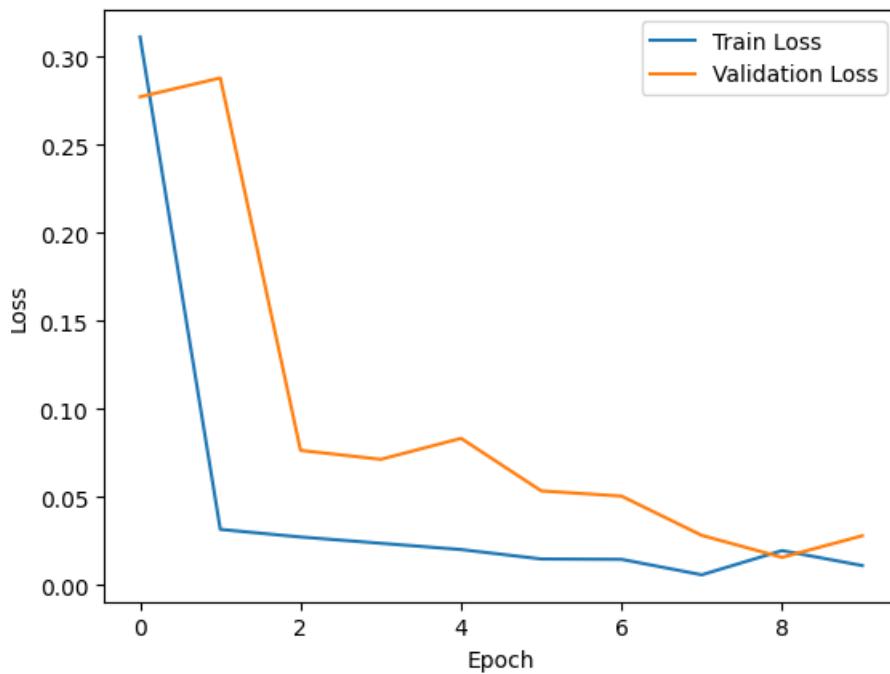
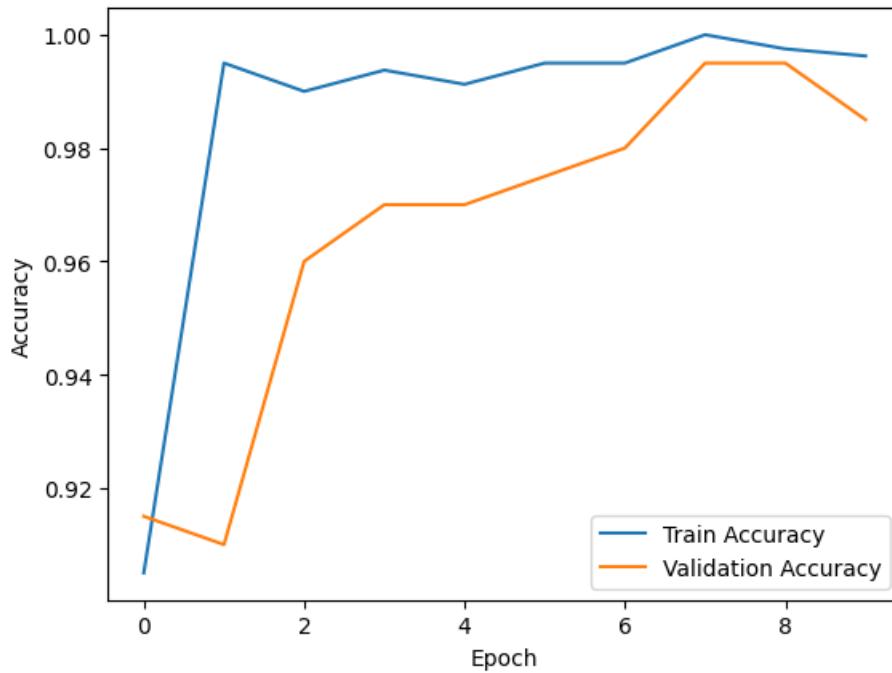
Another limitation of the Xception architecture is its long training time. Xception has a large number of layers and requires a large amount of computational resources to train, which can be a problem for applications where training time is a critical factor.

Real-world Example:

In a study published in the journal IEEE Transactions on Multimedia, researchers used Xception to classify emotions in real-time video streams. They found that the long training time of Xception made it difficult to optimize the network for real-time performance. They addressed this issue by using transfer learning to fine-tune a pre-trained Xception model on a smaller dataset.

Accuracy of X-ception Model

```
7/7 - 2s - loss: 0.0279 - accuracy: 0.9850 -  
Test Loss: 0.027922911569476128  
Test Accuracy: 0.9850000143051147
```



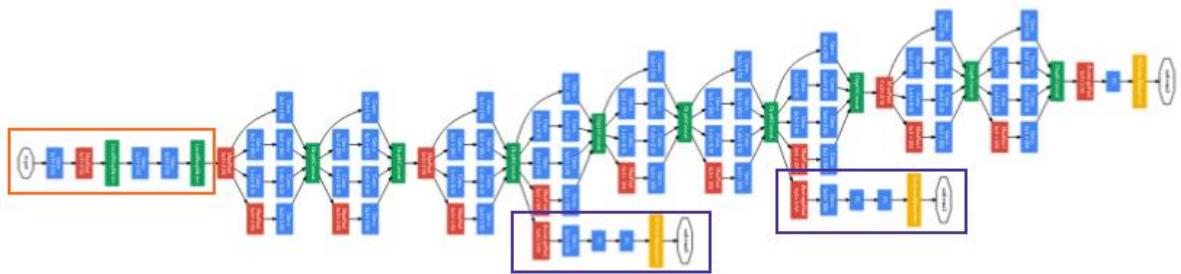
Explain each layer of Inception in detail ,give a full architecture overview

Inception is a deep convolutional neural network architecture that was proposed by Google in 2014. The Inception architecture has been widely used and has set the standard for state-of-the-art image classification models. The architecture is

based on the idea of building a network that has multiple paths or branches to extract features at different levels of abstraction.

The Inception architecture is composed of multiple modules that are stacked on top of each other to form the overall network. Each module is made up of several layers, which we will discuss in detail in the following sections.

- **Input Layer:** The input layer is where the input image is fed into the network. The input size is usually fixed, but can be adjusted based on the requirements of the task. The input size for Inception models typically ranges from 224x224 to 299x299.
- **Convolutional Layer:** The convolutional layer is the first layer in the Inception network. It applies a set of filters to the input image to extract features. The size of the filters can vary, but are typically 3x3 or 5x5. The output of the convolutional layer is a set of feature maps that represent the activations of the filters.
- **Pooling Layer:** The pooling layer is used to downsample the feature maps obtained from the convolutional layer. The most common type of pooling used in Inception is max pooling, which selects the maximum value from a small window of the feature map. This reduces the spatial size of the feature maps while preserving the most important features.
- **Inception Module:** The Inception module is the building block of the Inception architecture. It is composed of several parallel paths that extract features at different levels of abstraction. The output of each path is concatenated to form the final output of the module.



The paths in the Inception module include:

- ✓ 1x1 Convolution Path: This path applies a set of 1x1 filters to the input. This is used to reduce the number of channels in the feature maps and compress the information.

-
- ✓ 3x3 Convolution Path: This path applies a set of 3x3 filters to the input. This is used to extract features that are spatially more complex than those extracted by the 1x1 convolution path.
 - ✓ 5x5 Convolution Path: This path applies a set of 5x5 filters to the input. This is used to extract features that are even more spatially complex than those extracted by the 3x3 convolution path.
 - ✓ Max Pooling Path: This path applies max pooling to the input. This is used to extract the most important features from the input.

The Inception module allows the network to learn features at multiple scales and resolutions. By combining features from different paths, the network is able to achieve a high level of accuracy while minimizing the number of parameters.

- **Fully Connected Layer:** The fully connected layer is the last layer in the Inception network. It takes the output of the previous layer and applies a set of weights to it to obtain a prediction. The number of neurons in the fully connected layer is equal to the number of classes in the classification task.
- **Softmax Layer:** The softmax layer takes the output of the fully connected layer and applies the softmax function to obtain the final probability distribution over the classes.

Overall Architecture Overview The Inception architecture consists of multiple Inception modules stacked on top of each other. The number of modules and their complexity can vary depending on the requirements of the task. Typically, the architecture includes multiple fully connected layers and a softmax layer at the end for classification.

One of the most popular versions of the Inception architecture is InceptionV3. It has 42 convolutional layers and 11.2 million parameters.

- **Inception Module 3:** Inception Module 3 is similar to Inception Module 2, but it uses three branches. The first branch uses 1x1 convolution to reduce the number of filters, followed by 3x3 convolution. The second branch uses 1x1 convolution to reduce the number of filters, followed by two 3x3 convolutions. The third branch uses a 1x1 convolution to reduce the number of filters, followed by a max-pooling operation, and a 3x3 convolution. The outputs of these three branches are concatenated along the depth axis.
- **Inception Module 4:** Inception Module 4 is similar to Inception Module 3, but it uses four branches. The first branch uses 1x1 convolution to reduce the number of filters, followed by a 3x3 convolution. The second branch

uses 1×1 convolution to reduce the number of filters, followed by a 5×5 convolution. The third branch uses a max-pooling operation, followed by a 1×1 convolution. The fourth branch uses a 1×1 convolution. The outputs of these four branches are concatenated along the depth axis.

- **Inception Module 5:** Inception Module 5 is similar to Inception Module 4, but it uses two branches. The first branch uses 1×1 convolution to reduce the number of filters, followed by a 3×3 convolution. The second branch uses 1×1 convolution to reduce the number of filters, followed by a 3×3 convolution, and then another 3×3 convolution. The outputs of these two branches are concatenated along the depth axis.
- **Pooling Layer:** The last layer of the Inception architecture is a global average pooling layer. It reduces the spatial dimensions of the feature maps to 1×1 , and the depth remains the same. This layer computes the average of each feature map, which produces a feature vector for each example in the dataset. This feature vector is then fed into a fully connected layer to produce the final output of the network.

Overall, the Inception architecture is a powerful and efficient deep learning architecture that has been used in many computer vision applications. It has a high degree of parallelism, which allows it to process a large number of examples simultaneously. It also has a large number of layers, which enables it to learn complex features and patterns in the input data. The Inception architecture has achieved state-of-the-art results on several computer vision tasks, including image classification, object detection, and semantic segmentation.

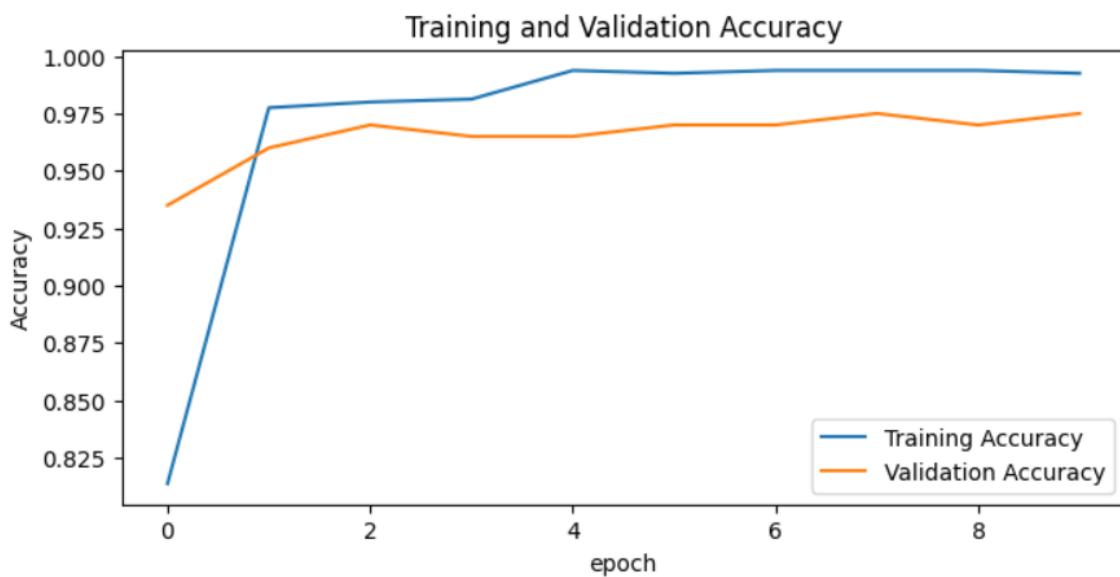
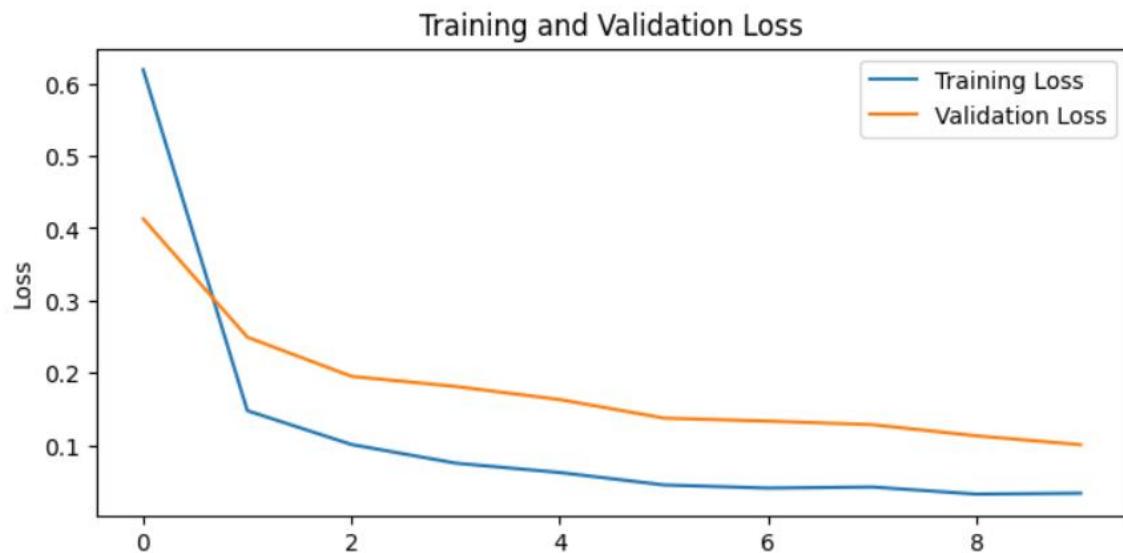
One of the key advantages of the Inception architecture is its ability to use smaller filters in the convolutional layers. By using smaller filters, the architecture can learn more complex features while reducing the number of parameters in the network. This reduces the risk of overfitting and allows the network to generalize better to new examples. Additionally, the use of multiple branches in the Inception modules enables the network to learn features at different scales and resolutions, which improves its ability to recognize objects in different positions and orientations.

Another advantage of the Inception architecture is its ability to process input images of different sizes. This is achieved through the use of global average pooling, which reduces the spatial dimensions of the feature maps to a fixed size. As a result, the network can process images of different sizes without requiring any additional preprocessing.

However, the Inception architecture also has some disadvantages. One of the main drawbacks is its high computational cost, which can make it difficult to train on large datasets. Additionally, the large number of layers in the network can make it prone to overfitting, especially if the dataset is small. Finally, the complex structure of the Inception modules can make it difficult to interpret the learned

Our Model Result

Accuracy → Test Loss: 0.101
Test Accuracy: 0.975



When comparing all the Fine-tuned Models

For the Given Data set → VGG16, VGG19, ResNet-50 gave 100% Accuracy

Reference Book

[1]"Python Machine Learning" by Sebastian Raschka and Vahid Mirjalili: This book covers a variety of topics related to machine learning, including the perceptron algorithm, multilayer neural networks, and convolutional neural networks. It provides clear explanations of these topics along with code examples in Python.

[2]"Deep Learning with Python" by Francois Chollet: This book focuses on deep learning and covers topics such as convolutional neural networks, recurrent neural networks, and multilayer perceptrons. It includes code examples in Python using the Keras library.

[3]"Hands-On Machine Learning with Scikit-Learn, Keras, and TensorFlow" by Aurelien Geron: This book covers a variety of machine learning topics and includes chapters on multilayer perceptrons and convolutional neural networks. It provides code examples in Python using the Scikit-Learn, Keras, and TensorFlow libraries.

[4]"Neural Networks and Deep Learning: A Textbook" by Charu Aggarwal: This book provides a comprehensive introduction to neural networks and deep learning. It covers topics such as perceptrons, multilayer neural networks, convolutional neural networks, and recurrent neural networks. It includes code examples in Python.

[5]"Python Deep Learning" by Valentino Zocca, Gianmario Spacagna, and Daniel Slater: This book covers a variety of deep learning topics, including multilayer perceptrons and convolutional neural networks. It provides code examples in Python using the Keras and TensorFlow libraries.