# **University of Mumbai**

Practical Journal of
Big Data Analytics
&
Image Processing

M.Sc. (Information Technology) Part-I

## Submitted by

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RASAYANI, 410207
MAHARASHTRA
2022-2023

# Mahatma Education Society's Pillai Hoc College of Arts, Science & Commerce, Rasayani (Affiliated to Mumbai University) RASAYANI – MAHARASHTRA - 410207

# DEPARTMENT OF INFORMATION TECHNOLOGY

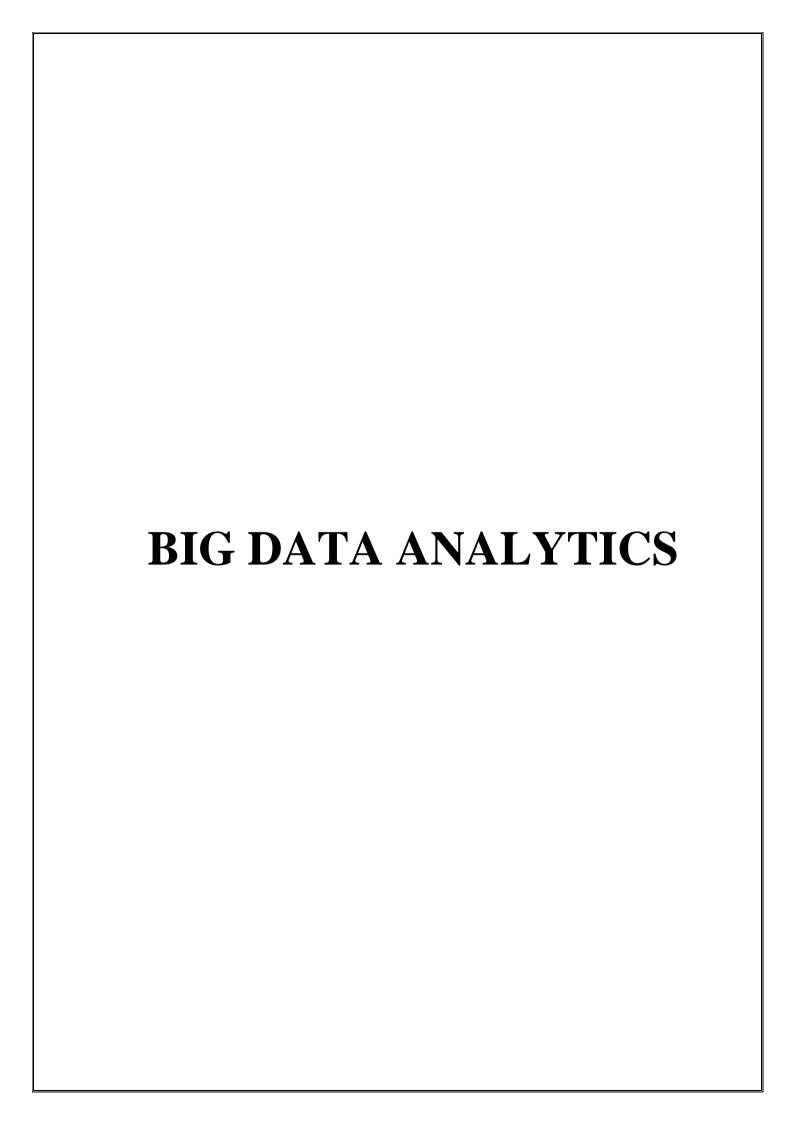


#### **CERTIFICATE**

This is to certify that the experiment work entered in this journal is as per the syllabus in **M.Sc.** (**Information Technology**) **Part-I, Semester-II**; class prescribed by University of Mumbai for the subject **Big Data Analytics** was done in computer lab of Mahatma Education Society's Pillai HOC College of Arts, Science & Commerce, Rasayani by **SHUBHAM BHOIR** during Academic year 2022-2023.

**Exam Seat No: 4133585** 

Subject In-Charge	Coordinator
External Examiner	Principal
Date:	College Seal

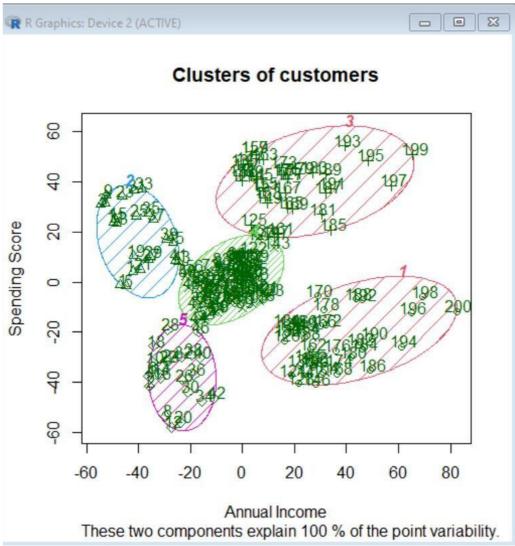


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Aim: A) Clustering algorithms for unsupervised classification. Read a datafile all\_Customers.csv and apply k-means clustering. Plot the cluster data using R visualizations.

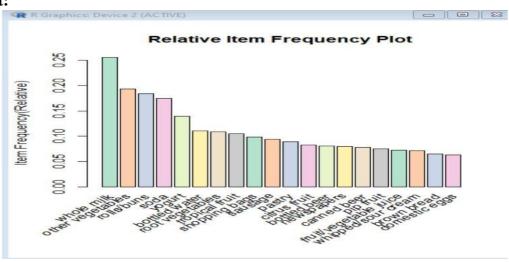
```
Code:
# K-Means Clustering
# Importing the dataset
dataset = read.csv("D:\\bda prac\\Mall_Customers.csv")
head(dataset)
dataset = dataset[4:5]
head(dataset)
wcss = vector()
for (i in 1:10) wcss[i] = sum(kmeans(dataset, i)$withinss)
plot(1:10,
   wcss,
  type = 'b',
  main = paste('The Elbow Method'),
  xlab = 'Number of clusters',
  ylab = 'WSS')
# Fitting K-Means to the dataset with no of clusters = 5
kmeans = kmeans(x = dataset, centers = 5)
y_kmeans = kmeans$cluster
# Visualising the clusters
library(cluster)
clusplot(dataset,
     y_kmeans,
     lines = 0,
     shade = TRUE,
     color = TRUE,
     labels = 2,
     main = paste('Clusters of customers'),
     xlab = 'Annual Income',
     ylab = 'Spending Score')
```



Aim: B) Implement Apriori Algorithm Recommending grocery items to a customer that is most frequently bought together, given a data set of transactions by customers of a store, using built-in Groceries file.

#### **Code:**

```
install.packages("arules")
install.packages("arulesViz")
install.packages("RColorBrewer")
library(arules)
library(arulesViz)
library(RColorBrewer)
data("Groceries")
Groceries
summary(Groceries)
class(Groceries)
rules = apriori(Groceries, parameter = list(supp = 0.02, conf = 0.2))
summary(rules)
inspect(rules[1:10])
arules::itemFrequencyPlot(Groceries, topN = 20,
                col = brewer.pal(8, 'Pastel2'),
                main = 'Relative Item Frequency Plot',
                type = "relative",
                ylab = "Item Frequency(Relative)")
itemset = apriori(Groceries, parameter = list(minlen=2, maxlen=2, support=0.02,
target="frequent itemset") )
summary(itemset)
inspect(itemset[1:10])
itemsets 3 = apriori(Groceries, parameter = list(minlen=3, maxlen=3, support=0.02,
target="frequent itemset"))
summary(itemsets_3)
inspect(itemsets_3)
```

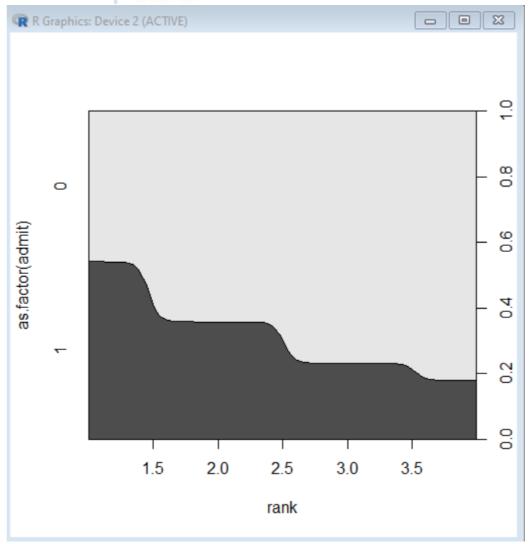


Aim: A) Import data from web storage – binary.csv. Name the dataset and do Logistic Regression to find out relation between variables that are affecting the admission of a student in an institute based on his or her GRE score, GPA obtained and rank of the student. Also check the model is fit or not.

#### Code:

```
#fetch the data
college <- read.csv("D:\\bda prac\\binary.csv")</pre>
head(college)
nrow(college)
install.packages("caTools") # For Logistic regression
library(caTools)
split <- sample.split(college, SplitRatio = 0.75)
split
training_reg <- subset(college, split == "TRUE")</pre>
test reg <- subset(college, split == "FALSE")
# Training model
fit_logistic_model <- glm(admit ~ .,
              data = training_reg,
              family = "binomial")
# Predict test data based on model
predict reg <- predict(fit logistic model,</pre>
              test reg, type = "response")
predict_reg
cdplot(as.factor(admit)~ gpa, data=college)
cdplot(as.factor(admit)~ gre, data=college)
cdplot(as.factor(admit)~ rank, data=college)
# Changing probabilities
predict_reg <- ifelse(predict_reg >0.5, 1, 0)
predict_reg
# Evaluating model accuracy
# using confusion matrix
table(test_reg$admit, predict_reg)
```

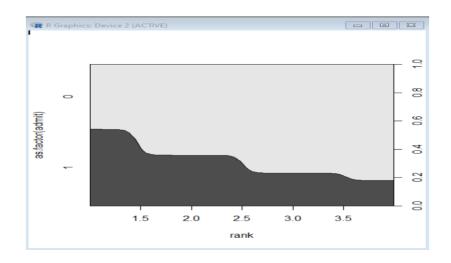
```
> table(test_reg$admit, predict_reg)
   predict_reg
    0 1
0 59 3
1 28 10
```



# Aim: B) Apply multiple regressions, if data have a continuous independent variable. Apply on above dataset – binary.csv.

```
Code:
```

```
#fetch the data
college <- read.csv("D:\\bda prac\\binary.csv")</pre>
head(college)
nrow(college)
install.packages("caTools") # For Logistic regression
library(caTools)
split <- sample.split(college, SplitRatio = 0.75)
split
training_reg <- subset(college, split == "TRUE")</pre>
test_reg <- subset(college, split == "FALSE")</pre>
# Training model
fit_MRegressor_model <- lm(formula = admit ~ gre+gpa+rank,
              data = training_reg)
# Predict test data based on model
predict_reg <- predict(fit_MRegressor_model,</pre>
              newdata = test_reg)
predict_reg
cdplot(as.factor(admit)~ gpa, data=college)
cdplot(as.factor(admit)~ gre, data=college)
cdplot(as.factor(admit)~ rank, data=college)
```

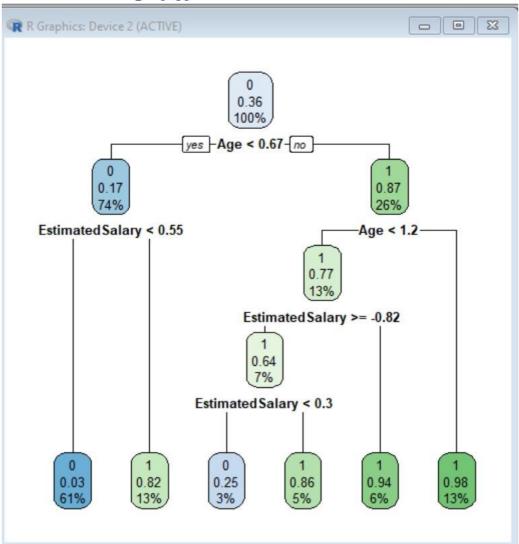


Aim: A) Implement Decision Tree classification technique using Social\_Network\_Ads.csv dataset.

```
Code:
# Decision Tree Classification
# Importing the dataset
dataset = read.csv("D:\\bda prac\\Social_Network_Ads.csv")
#print(dataset)
dataset = dataset[3:5] # columns 3 4 ad 5
print(dataset)
# Encoding the target feature as factor(just like a vector having levels
# levels to convey that only two possible values for purchased - 0 & 1
datasetPurchased = factor(datasetPurchased, levels = c(0, 1))
print (dataset$Purchased)
# Splitting the dataset into the Training set and Test set
install.packages('caTools')
library(caTools)
set.seed(123)
#split = sample.split(dataset$Purchased, SplitRatio = 0.75)
split = sample.split(dataset$Purchased, SplitRatio = 0.75)
training set = subset(dataset, split == TRUE)
test set = subset(dataset, split == FALSE)
# Feature Scaling - scale() method centers and/or scales the columns of a numeric matrix.
training set[-3] = scale(training set[-3]) # scaling first 2 columns, don't consider 3rd column
test\_set[-3] = scale(test\_set[-3])
#print(test_set[-3])
# Fitting Decision Tree Classification to the Training set
install.packages('rpart')
library(rpart) # for partitioning tree
install.packages('rpart.plot')
library(rpart.plot)
classifier = rpart(formula = Purchased ~ .,data = training_set)
# Predicting the Test set results
y_pred = predict(classifier, newdata = test_set[-3], type = 'class')
print(y_pred)
# Making the Confusion Matrix
cm = table(test_set[, 3], y_pred)
print(cm)
y_grid = predict(classifier, newdata = grid_set, type = 'class')
```

#### # Plotting the tree

#extra=106 class model with a binary response #extra=104 class model with a response having more than two levels rpart.plot(classifier, extra = 106)



# Aim: B) Implement SVM Classification technique using Social\_Network\_Ads.csv dataset. Evaluate the performance of classifier.

```
Code:
# Support Vector Machine (SVM)
# Importing the dataset
dataset = read.csv("D:\\bda prac\\Social_Network_Ads.csv")
dataset = dataset[3:5]
print(dataset)
print(dataset$Purchased)
# Splitting the dataset into the Training set and Test set
install.packages('caTools')
library(caTools)
set.seed(123)
split = sample.split(dataset$Purchased, SplitRatio = 0.75)
training_set = subset(dataset, split == TRUE)
print(training set)
test set = subset(dataset, split == FALSE)
print(test_set)
# Feature Scaling
training_set[-3] = scale(training_set[-3]) # [-3] means 3rd index will be dropped
test\_set[-3] = scale(test\_set[-3])
print(training_set[-3])
print (test_set[-3])
# Fitting SVM to the Training set
install.packages('e1071')
library(e1071)
classifier = svm(formula = Purchased ~ .,
          data = training_set,
          type = 'C-classification',
          kernel = 'linear')
print (classifier)
# Predicting the Test set results
y_pred = predict(classifier, newdata = test_set[-3])
print(y_pred)
# Making the Confusion Matrix
cm = table(test_set[, 3], y_pred)
print (cm)
Output:
                         > cm = table(test_set[, 3], y_pred)
                         > print (cm)
                            y pred
```

0 57 7 1 13 23

Aim: A) Implement Naïve Bayes Classification technique using Social\_Network\_Ads.csv dataset. Evaluate the performance of classifier.

#### Code:

```
# Naive Bayes
# Importing the dataset
dataset = read.csv('C:\\2022-23\\BDA practical 2023\\Social_Network_Ads.csv')
dataset = dataset[3:5]
# Encoding the target feature as factor
dataset$Purchased = factor(dataset$Purchased, levels = c(0, 1))
# Splitting the dataset into the Training set and Test set
#install.packages('caTools')
library(caTools)
set.seed(123)
split = sample.split(dataset$Purchased, SplitRatio = 0.75)
training set = subset(dataset, split == TRUE)
test_set = subset(dataset, split == FALSE)
# Feature Scaling
training_set[-3] = scale(training_set[-3])
test set[-3] = scale(test set[-3])
# Fitting Naive Bayes to the Training set
install.packages('e1071')
library(e1071)
classifier = naiveBayes(x = training set[-3],
               y = training set$Purchased)
# Predicting the Test set results
y_pred = predict(classifier, newdata = test_set[-3])
# Making the Confusion Matrix
cm = table(test_set[, 3], y_pred)
print(cm)
```

# Aim: B) Find the confusion matrix to find restaurant review based of sentiment analysis of Natural Language processing. Use Resaurentreviews.tsv file for your study.

#### Code:

```
dataset_original = read.delim("D:\\bda prac\\Restaurant_Reviews.txt", quote = ",
stringsAsFactors = FALSE)
install.packages('tm')
install.packages('SnowballC')
library(tm)
library(SnowballC)
corpus = VCorpus(VectorSource(dataset_original$Review))
corpus = tm_map(corpus, content_transformer(tolower))
corpus = tm_map(corpus, removeNumbers)
corpus = tm map(corpus, removePunctuation)
corpus = tm_map(corpus, removeWords, stopwords())
corpus = tm_map(corpus, stemDocument)
corpus = tm map(corpus, stripWhitespace)
dtm = DocumentTermMatrix(corpus)
dtm = removeSparseTerms(dtm, 0.999)
dataset = as.data.frame(as.matrix(dtm))
dataset$Liked = dataset_original$Liked
print(dataset$Liked)
dataset$Liked = factor(dataset$Liked, levels = c(0,1))
install.packages(caTools)
library(caTools)
set.seed(123)
split = sample.split(dataset$Liked, SplitRatio = 0.8)
training set = subset(dataset, split == TRUE)
test_set = subset(dataset, split == FALSE)
install.packages('randomForest')
library(randomForest)
classifier = randomForest(x = training\_set[-692],
               y = training_set$Liked,
               ntree = 10
y_pred = predict(classifier, newdata = test_set[-692])
cm = table(test_set[,692], y_pred)
print(cm)
Output:
                      > cm = table(test_set[,692], y_pred)
                       > print (cm)
                          y pred
```

Aim: Take the inbuilt data file: iris and perform classification on that data using various classification models – Decision Tree, K Nearest Neighbour and Support Vector Machine. Find the confusion matrix for all three models and evaluate them by finding their accuracy. Find the algorithm which performs best on the given data file, out of all these three models.

```
Code:
#PBL
install.packages('rpart')
install.packages('rpart.plot')
install.packages('gmodels')
install.packages('e1071')
library(rpart)
library(rpart.plot)
library(gmodels)
library(e1071)
data(iris)
summary(iris)
#normalize the continuous variables before performing any analysis on the dataset
temp = as.data.frame(scale(iris[,1:4]))
temp$Species = iris$Species # levels: setosa versicolor virginica
summary(temp)
# Splitting the dataset into the Training set and Test set
install.packages('caTools')
library(caTools)
set.seed(123)
split = sample.split(temp$Species, SplitRatio = 0.75)
train = subset(temp, split == TRUE)
test = subset(temp, split == FALSE)
nrow(train)
nrow(test)
#1. Decision Trees
dt_classifier = rpart(formula = Species ~ .,data = train)
# Predicting the Test set results
dt_y_pred = predict(dt_classifier, newdata = test, type = 'class')
print(dt_y_pred)
# Making the Confusion Matrix for Decision Tree
cm = table(test$Species, dt_y_pred)
print(cm)
#accuracy of DT model
```

DTaccu = ((12+9+11)/nrow(test))\*100#true positive nos of 3\*3 confusion matrix

#### DTaccu

```
#2. k-Nearest Neighbours
install.packages("class")
library(class)
cl = train$Species
set.seed(1234)
knn_y_pred = knn(train[,1:4],test[,1:4],cl,k=5)
# cm of k-Nearest Neighbours
cm = table(test$Species, knn_y_pred)
print(cm)
#accuracy of KNN model
KNNaccu = ((12+11+11)/nrow(test))*100 #true positive nos of 3*3 confusion matrix
KNNaccu
#3. Support Vector Machine(SVM)
symclassifier = sym(Species \sim ., data = train)
svm_y_pred = predict(svmclassifier,newdata = test)
cm = table(test$Species, svm_y_pred)
print(cm)
#accuracy of SVM model
SVMaccu = ((12+11+11)/nrow(test))*100 #true positive nos of 3*3 confusion matrix
SVMaccu
#Decision Tree vs kNN
which(dt_y_pred != knn_y_pred)
#Decision Tree vs SVM
which(dt_y_pred != svm_y_pred)
#svm vs kNN
which(svm_y_pred != knn_y_pred) #both are equal
#Comparison of the accuracy of different models on testing dataset.
models = data.frame(Technique = c("Decision Tree", "kNN", "SVM"), Accuracy_Percentage =
c(88.88889,94.44444,94.44444))
models
```

```
> # Making the Confusion Matrix for Decision Tree
> cm = table(test$Species, dt_y_pred)
> print(cm)
           dt_y_pred
           setosa versicolor virginica
 setosa
                12
                                      0
                0
  versicolor
                             9
                                      3
 virginica
                 0
                             1
                                      11
> #accuracy of DT model
> DTaccu = ((12+9+11)/nrow(test))*100 #true positive nos of 3*3 confusion matrix
> DTaccu
[1] 88.88889
> # cm of k-Nearest Neighbours
> cm = table(test$Species, knn_y_pred)
> print(cm)
            knn_y_pred
setosa versicolor virginica
  setosa
               12
  versicolor
                 0
                            11
                                      11
  virginica
                             1
> #accuracy of KNN model
> KNNaccu
[1] 94.44444
> cm = table(test$Species, svm_y_pred)
> print(cm)
            svm_y_pred
            setosa versicolor virginica
                12
                                  0
  setosa
                           0
  versicolor
                  0
                            1.1
                                      11
  virginica
> #accuracy of SVM model
> SVMaccu = ((12+11+11)/nrow(test))*100 #true positive nos of 3*3 confusion matrix
> SVMaccu
[1] 94.44444
> #Comparison of the accuracy of different models on testing dataset.
> models = data.frame(Technique = c("Decision Tree", "kNN", "SVM"), Accuracy Percentage = c(88.88889,94.44444,94.44444))
> models
    Technique Accuracy Percentage
l Decision Tree
                88.88889
2
       kNN
                  94.44444
                  94.44444
3
       SVM
5 |
```

#### Aim: Install, configure and run Hadoop and HDFS and explore HDFS on Windows

#### **Steps to Install Hadoop**

- 1. Install Java JDK 1.8
- 2. Download Hadoop and extract and place under C drive
- 3. Set Path in Environment Variables
- 4. Config files under Hadoop directory
- 5. Create folder datanode and namenode under data directory
- 6. Edit HDFS and YARN files
- 7. Set Java Home environment in Hadoop environment
- 8. Setup Complete. Test by executing start-all.cmd

#### There are two ways to install Hadoop, i.e.

- 9. Single node
- 10. Multi node

Here, we use multi node cluster.

#### 1. Install Java

- 11. Java JDK Link to download https://www.oracle.com/java/technologies/javase-jdk8-downloads.html
- 12. extract and install Java in C:\Java
- 13. open cmd and type -> javac -version

```
C:\Users>cd Beena

C:\Users\Beena>java -version
java version "1.8.0_361"

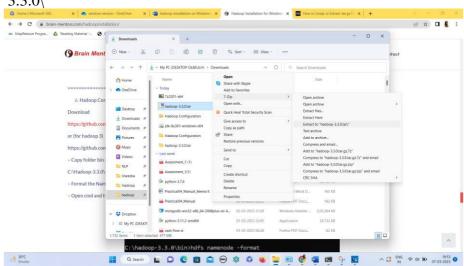
Java(TM) SE Runtime Environment (build 1.8.0_361-b09)

Java HotSpot(TM) 64-Bit Server VM (build 25.361-b09, mixed mode)
```

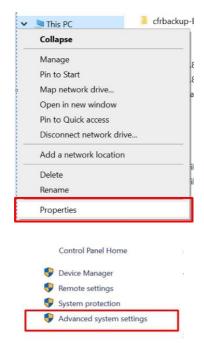
#### 2. Download Hadoop

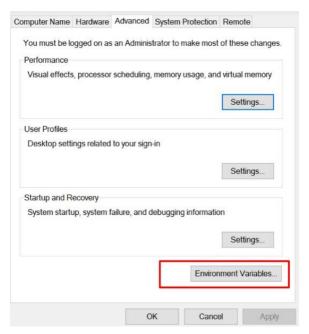
https://www.apache.org/dyn/closer.cgi/hadoop/common/hadoop-3.3.0/hadoop-3.3.0.tar.gz

• right click .rar.gz file -> show more options -> 7-zip->and extract to C:\Hadoop-3.3.0\

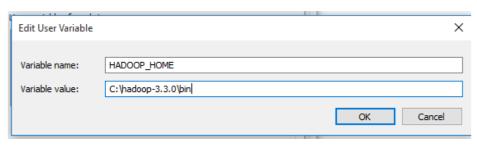


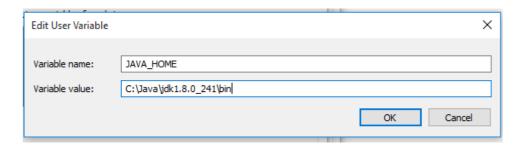
- 3. Set the path JAVA\_HOME Environment variable
- 4. Set the path HADOOP\_HOME Environment variable



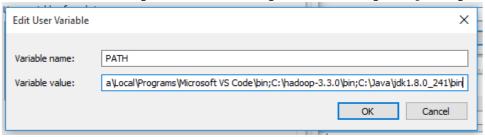


Click on New to both user variables and system variables.





Click on user variable -> path -> edit-> add path for Hadoop and java upto 'bin'



Click Ok, Ok, Ok.

#### 5. Configurations

Edit file C:/Hadoop-3.3.0/etc/hadoop/core-site.xml,

Rename "mapred-site.xml.template" to "mapred-site.xml" and edit this file C:/Hadoop-3.3.0/etc/hadoop/mapred-site.xml, paste xml code and save this file.

<configuration>
<configuration>

<name>mapreduce.framework.name
<value>yarn</value>

```
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                                Big Data Analytics
 </configuration>
Create folder "data" under "C:\Hadoop-3.3.0"
Create folder "datanode" under "C:\Hadoop-3.3.0\data"
Create folder "namenode" under "C:\Hadoop-3.3.0\data"
Edit file C:\Hadoop-3.3.0/etc/hadoop/hdfs-site.xml,
paste xml code and save this file.
<configuration>
cproperty>
   <name>dfs.replication</name>
   <value>1</value>
 cproperty>
   <name>dfs.namenode.name.dir</name>
   <value>/hadoop-3.3.0/data/namenode
 cproperty>
   <name>dfs.datanode.data.dir</name>
   <value>/hadoop-3.3.0/data/datanode
 </configuration>
Edit file C:/Hadoop-3.3.0/etc/hadoop/yarn-site.xml,
paste xml code and save this file.
<configuration>
```

```
M.Sc.IT Sem-II
                               Big Data Analytics
 cproperty>
        <name>yarn.nodemanager.aux-services</name>
        <value>mapreduce_shuffle</value>
 cproperty>
       <name>yarn.nodemanager.auxservices.mapreduce.shuffle.class</name>
        <value>org.apache.hadoop.mapred.ShuffleHandler</value>
 cproperty>
            <name>yarn.resourcemanager.address</name>
            <value>127.0.0.1:8032</value>
 cproperty>
            <name>yarn.resourcemanager.scheduler.address</name>
            <value>127.0.0.1:8030
  cproperty>
            <name>yarn.resourcemanager.resource-tracker.address</name>
            <value>127.0.0.1:8031
  </configuration>
6. Edit file C:/Hadoop-3.3.0/etc/hadoop/hadoop-env.cmd
Find "JAVA HOME=%JAVA HOME%" and replace it as
set JAVA_HOME="C:\Java\jdk1.8.0_361"
```

#### 7. Download "redistributable" package

#### Download and run VC redist.x64.exe

This is a "redistributable" package of the Visual C runtime code for 64-bit applications, from Microsoft. It contains certain shared code that every application written with Visual C expects to have available on the Windows computer it runs on.

#### 8. Hadoop Configurations

Download bin folder from

https://github.com/s911415/apache-hadoop-3.1.0-winutils

- Copy the bin folder to c:\hadoop-3.3.0. Replace the existing bin folder.
- 9. copy "hadoop-yarn-server-timelineservice-3.0.3.jar" from ~\hadoop-3.0.3\share\hadoop\yarn\timelineservice to ~\hadoop-3.0.3\share\hadoop\yarn folder.
- 10. Format the NameNode
- Open cmd 'Run as Administrator' and type command "hdfs namenode -format"

```
Administrator: Command Prompt

Microsoft Windows [Version 10.0.22621.1265]
(c) Microsoft Corporation. All rights reserved.

C:\Windows\System32>cd\hadoop-3.3.0\bin

C:\hadoop-3.3.0\bin>hdfs namenode -format
```

#### 11. Testing

- Open cmd 'Run as Administrator' and change directory to C:\Hadoop-3.3.0\sbin
- type start-all.cmd

#### OR

- type start-dfs.cmd
- type start-yarn.cmd

```
C:\hadoop-3.3.0\sbin>start-all.cmd
This script is Deprecated. Instead use start-dfs.cmd and start-yarn.cmd
The filename, directory name, or volume label syntax is incorrect.
The filename, directory name, or volume label syntax is incorrect.
starting yarn daemons
The filename, directory name, or volume label syntax is incorrect.
```

- You will get 4 more running threads for Datanode, namenode, resouce manager and node manager

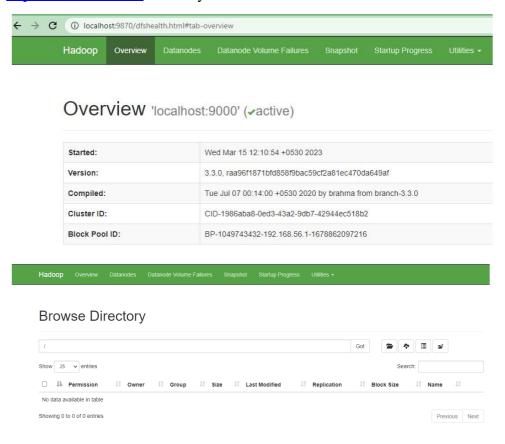
```
Apache Hadoop Distribution - hadoop namenode
2023-03-07 20:33:00,395 INFO ipc.Server: Starting Socket Reader #1 for port 9000
2023-03-07 20:33:00,547 INFO namenode.FSNamesystem: Registered FSNamesystemState, ReplicatedBlocksState and ECBlockGroup
sState MBeans.
2023-03-07 20:33:00,549 INFO common.Util: Assuming 'file' scheme for path /hadoop-3.3.0/data/namenode in configuration.
2023-03-07 20:33:00,554 INFO namenode.LeaseManager: Number of blocks under construction: 0
2023-03-07 20:33:00,563 INFO blockmanagement.DatanodeAdminDefaultMonitor: Initialized the Default Decommission and Maint
 enance monitor
2023-03-07 20:33:00,566 INFO blockmanagement.BlockManager: initializing replication queues
2023-03-07 20:33:00,567 INFO hdfs.StateChange: STATE* Leaving safe mode after 0 secs
2023-03-07 20:33:00,567 INFO hdfs.StateChange: STATE* Network topology has 0 racks and 0 datanodes 2023-03-07 20:33:00,569 INFO hdfs.StateChange: STATE* UnderReplicatedBlocks has 0 blocks
2023-03-07 20:33:00,574 INFO blockmanagement.BlockManager: Total number of blocks
2023-03-07 20:33:00,575 INFO blockmanagement.BlockManager: Number of invalid blocks
                                                                                                                                                                           = 0
                                                                                                                                                                           = 0
2023-03-07 20:33:00,575 INFO blockmanagement.BlockManager: Number of under-replicated blocks = 0
2023-03-07 20:33:00,576 INFO blockmanagement.BlockManager: Number of over-replicated blocks = 0
2023-03-07 20:33:00,576 INFO blockmanagement.BlockManager: Number of blocks being written = 0
2023-03-07 20:33:00,576 INFO hdfs.StateChange: STATE* Replication Queue initialization scan for invalid, over- and under
 replicated blocks completed in 9 msec
2023-03-07 20:33:00,607 INFO ipc.Server: IPC Server Responder: starting
2023-03-07 20:33:00,607 INFO ipc.Server: IPC Server listener on 9000: starting 2023-03-07 20:33:00,611 INFO ipc.MameNode: NameNode RPC up at: localhost/127.0.0.1:9000
2023-03-07 20:33:00,614 INFO namenode.FSNamesystem: Starting services required for active state
2023-03-07 20:33:00,614 INFO namenode.FSDirectory: Initializing quota with 4 thread(s)
2023-03-07 20:33:00,622 INFO namenode.FSDirectory: Quota initialization completed in 7 milliseconds
 name space=1
storage space=0
storage types=RAM_DISK=0, SSD=0, DISK=0, ARCHIVE=0, PROVIDED=0
2023-03-07 20:33:00,626 INFO blockmanagement.CacheReplicationMonitor: Starting CacheReplicationMonitor with interval 300
 00 milliseconds
```

#### **Output:**

12. Type JPS command to start-all.cmd command prompt, you will get following output.

```
C:\hadoop-3.3.0\sbin>jps
5632 Jps
7572 DataNode
3752 ResourceManager
7992 NameNode
8028 NodeManager
```

## 13. Run <a href="http://localhost:9870/">http://localhost:9870/</a> from any browser



Aim: Implement an application that stores big data in Hbase / MongoDB and manipulate it using R / Python

#### Requirements

- a. PyMongo
- b. Mongo Database

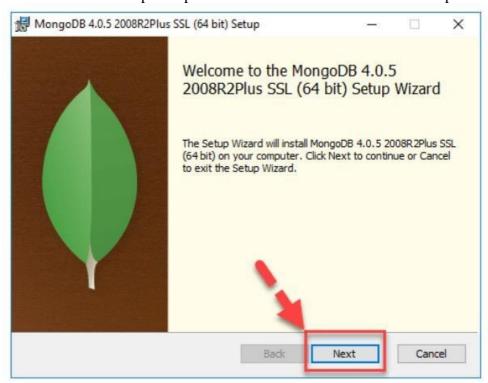
#### Step A: Install Mongo database

Step 1) Go to (https://www.mongodb.com/download-center/community) and Download MongoDB Community Server. We will install the 64-bit version for Windows.

Select the server you would like to run:

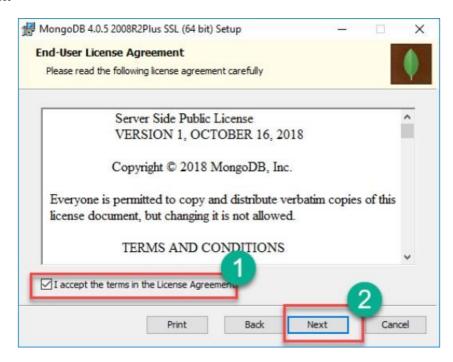


Step 2) Once download is complete open the msi file. Click Next in the start up screen

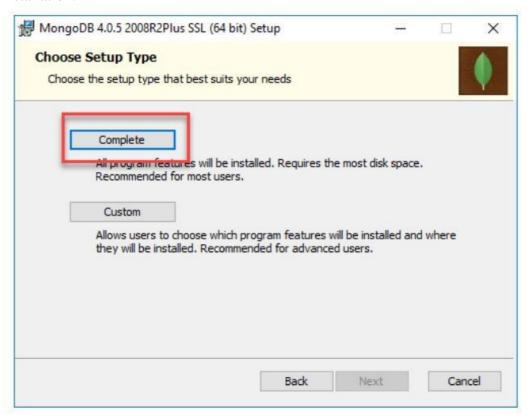


#### Step 3)

- 1. Accept the End-User License Agreement
- 2. Click Next



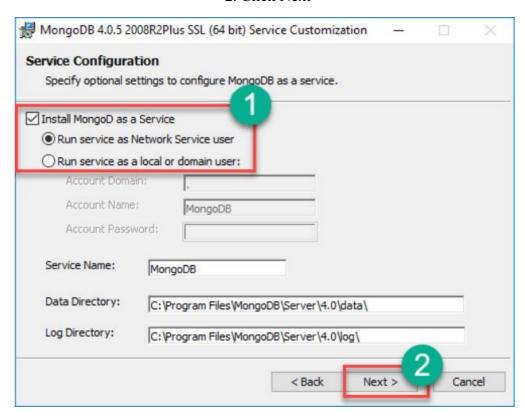
Step 4) Click on the "complete" button to install all of the components. The custom option can be used to install selective components or if you want to change the location of the installation.



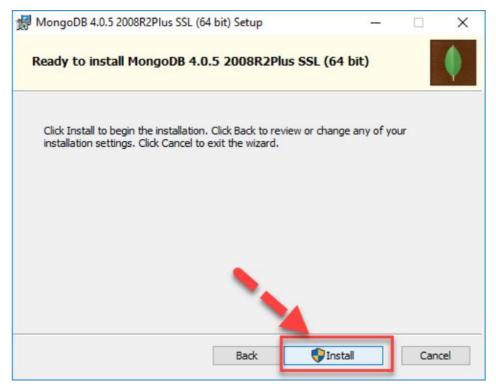
#### Step 5)

1. Select "Run service as Network Service user". make a note of the data directory, we"ll need this later.

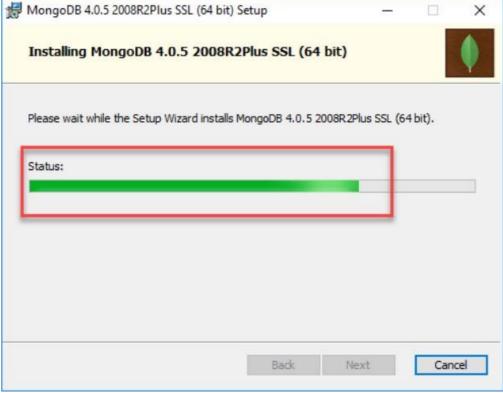
#### 2. Click Next



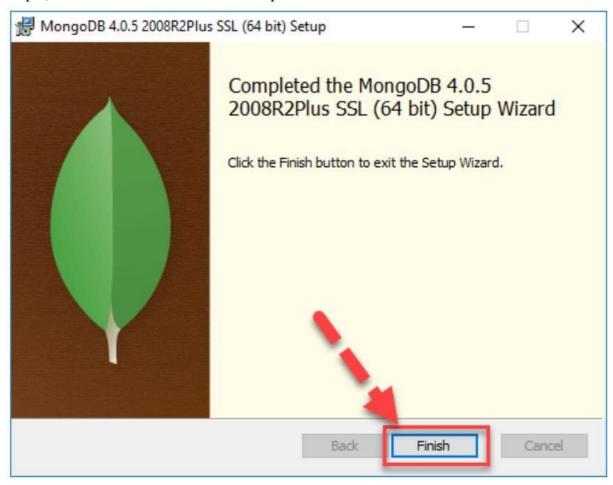
Step 6) Click on the Install button to start the installation.



Step 7) Installation begins. Click Next once completed.



Step 8) Click on the Finish button to complete the installation.



#### **Test Mongodb**

**Step 1**) Go to "C:\Program Files\MongoDB\Server\4.0\bin" and double click on **mongo.exe.** Alternatively, you can also click on the MongoDB desktop icon.

• Create the directory where MongoDB will store its files.

Open command prompt window and apply following commands

C:\users\admin> cd\ C:\>md data\db

#### **Step 2) Execute mongodb**

Open another command prompt window.

C:\> cd C:\Program Files\MongoDB\Server\4.0\bin

C:\Program Files\MongoDB\Server\4.0\bin> mongod

*In case if it gives an error then run the following command:* 

 $C: \Program\ Files \MongoDB \Server \4.0 \bin>mongod-repair$ 

#### Step 3) Connect to MongoDB using the Mongo shell

Let the MongoDB daemon to run.

Open another command prompt window and run the following commands:

C:\users\admin> cd C:\Program Files\MongoDB\Server\4.0\bin

C:\Program Files\MongoDB\Server\4.0\bin>mongo

#### Step 4) Install PyMongo

Open another command prompt window and run the following commands:

Check the python version on your desktop / laptop and copy that path from window explorer

C:\users\admin>cd C:\Program Files\Python311\Scripts

C:\Program Files\<Python38>\Scripts > python -m pip install pymongo

Note: # -m option is for <module-name>
Now you have downloaded and installed a mongoDB driver.

#### Step 5) Test PyMongo

Run the following command from python command prompt

import pymongo

Now, either create a file in Python IDLE or run all commands one by one in sequence on Python cell

#### Program 1: Creating a Database: create\_dp.py

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
print(myclient.list_database_names())
['admin', 'config', 'local']
```

### Progam 2: Creating a Collection: create\_collection.py

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol=mydb["student"]
print(mydb.list_collection_names())
[]
```

#### Progam 3: Insert into Collection: insert into collection.py

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol=mydb["student"]
mydict={"name":"Beena", "address":"Mumbai"}
x=mycol.insert_one(mydict) # insert_one(containing the name(s) and value(s) of each field
```

#### Program 4: Insert Multiple data into Collection: insert\_many.py

```
import pymongo
myclient = pymongo.MongoClient("mongodb://localhost:27017/")
mydb = myclient["mybigdata"]
mycol=mydb["student"]
mylist=[{"name":"Khyati", "address":"Mumbai"}, {"name":"Kruti", "address":"Mumbai"},
{"name":"Nidhi", "address":"Pune"}, {"name":"Komal", "address":"Pune"},]
x=mycol.insert_many(mylist)
```

#### Step 6) Test in Mongodb to check database and data inserted in collection

a. If you want to check your database list, use the command show dbs in mongo command prompt

> show dbs

```
admin 0.000GB
config 0.000GB
local 0.000GB
mybigdata 0.000GB
```

- b. If you want to use a database with name mybigdata, then use database statement would be as follow:
- > use mybigdata

```
switched to db mybigdata
```

c. If you want to check collection in mongodb use the command show collections > show collections

student

d. If you want to display the first row from collection: db.collection\_name.find() > db.student.findOne()

e. If you want to display all the data from collection: db.collection\_name.find()

> db.student.find()

```
> db.student.find()
{ "_id" : ObjectId("640178face663db608cef72f"), "name" : "Beena", "address" : "Mumbai" }
{ "_id" : ObjectId("640179336ce317082c266dc1"), "name" : "Khyati", "address" : "Mumbai" }
{ "_id" : ObjectId("640179336ce317082c266dc2"), "name" : "Kruti", "address" : "Mumbai" }
{ "_id" : ObjectId("640179336ce317082c266dc3"), "name" : "Nidhi", "address" : "Pune" }
{ "_id" : ObjectId("640179336ce317082c266dc4"), "name" : "Komal", "address" : "Pune" }
```

f. count number of rows in a collection

> db.student.count()



#### **Site for R packages documentation:**

https://cran.r-project.org/web/packages/available\_packages\_by\_name.html

# **University of Mumbai**

Practical Journal of
Big Data Analytics
&
Image Processing

M.Sc. (Information Technology) Part-I

## Submitted by

### **BHOIR SHUBHAM MAHENDRA**

Seat No: 4133585



DEPARTMENT OF INFORMATION TECHNOLOGY
PILLAI HOC COLLEGE OF ARTS, SCIENCE & COMMERCE, RASAYANI
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MAHARASHTRA
2022-2023

# Mahatma Education Society's Pillai Hoc College of Arts, Science & Commerce, Rasayani (Affiliated to Mumbai University) RASAYANI – MAHARASHTRA - 410207

# DEPARTMENT OF INFORMATION TECHNOLOGY

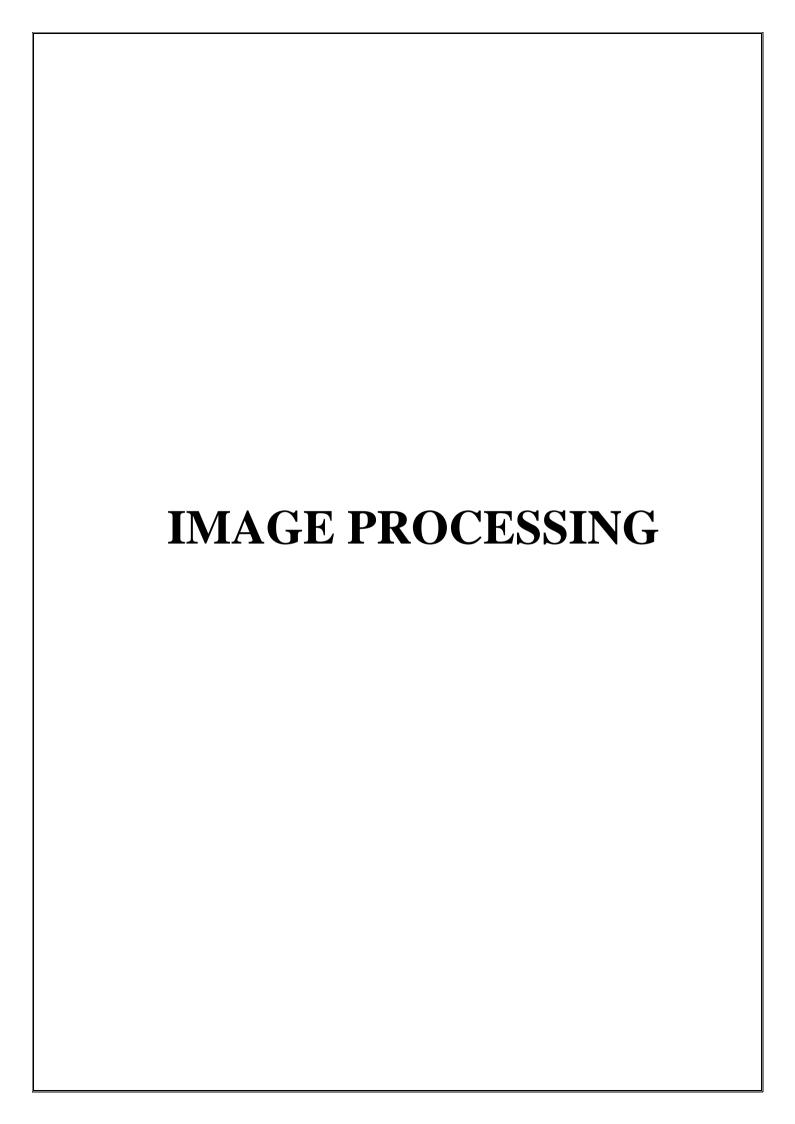


#### **CERTIFICATE**

This is to certify that the experiment work entered in this journal is as per the syllabus in **M.Sc.** (**Information Technology**) **Part-I, Semester-II**; class prescribed by University of Mumbai for the subject **Image Processing** was done in computer lab of Mahatma Education Society's Pillai HOC College of Arts, Science & Commerce, Rasayani by **SHUBHAM BHOIR** during Academic year 2022-2023.

**Exam Seat No: 4133585** 

Subject In-Charge	Coordinator
External Examiner	Principal
Date:	College Seal



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Aim: A) Program to calculate a number of samples required for the image.

### **Code:**

clc;

close;

m=4;

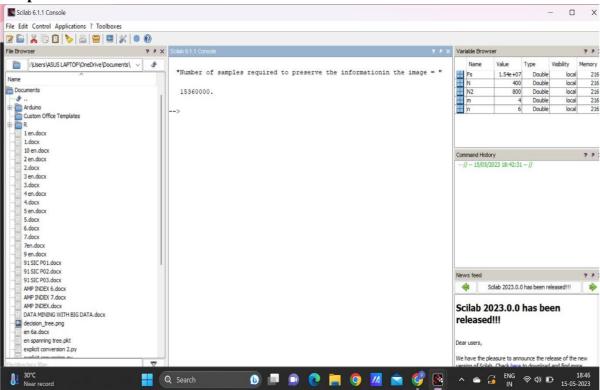
n=6;

N=400;

N2=2\*N;

Fs=m\*N2\*n\*N2;

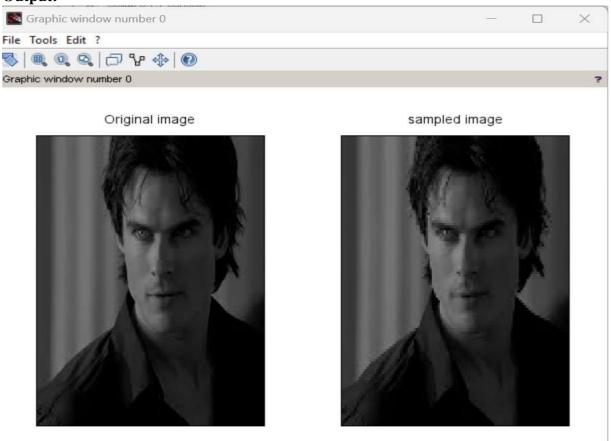
disp('Number of samples required to preserve the information in the image = ',Fs);



# Aim: B) Program to study the effects of reducing the spatial resolution of a digital image.

### **Code:**

```
n = input('Enter the input samples');
img=rgb2gray(imread('D:\damon.jpeg'));
a=size(img);
w=a(2);
h=a(1);
im=zeros(100);
for i=1:n:h
for j=1:n:w
for k=0:n-1
for l=0:n-1
im(i+k,j+l)=img(i,j);
end
end
end
end
subplot(1,2,1);
imshow(uint8(img));title('Original image');
<u>subplot(1,2,2);</u>
imshow(uint8(im));title('sampled image');
```

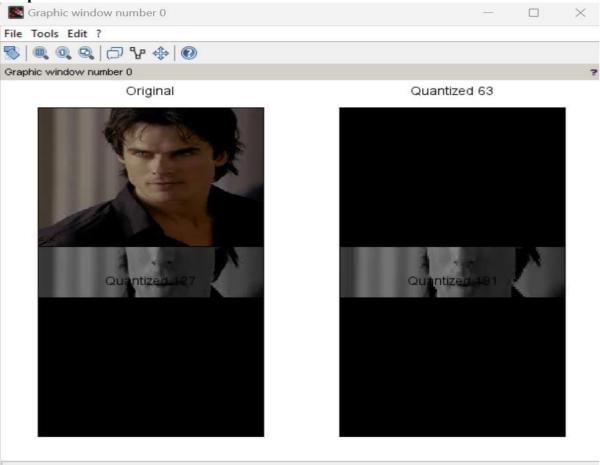


Aim: WAP to study the effect of reducing the quantization values and spatial resolution.

1) Quantization

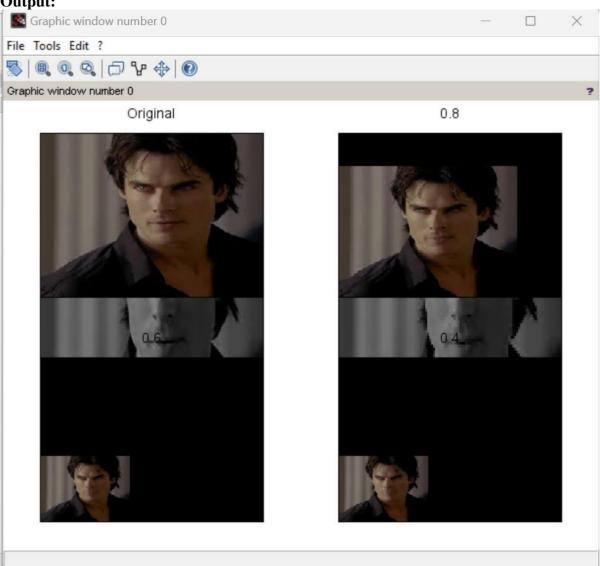
### **Code:**

```
a=imread('D:\damon.jpeg');
[m,n]=size(a);
for i=1:m
for j=1:n
b(i,j)=(a(i,j))/255*63;
c(i,j)=(a(i,j))/255*127;
d(i,j)=(a(i,j))/255*191;
end
end
subplot(2,2,1),imshow(a),title('Original');
subplot(2,2,2),imshow(b),title('Quantized 63');
subplot(2,2,3),imshow(c),title('Quantized 127');
subplot(2,2,4),imshow(d),title('Quantized 191');
```

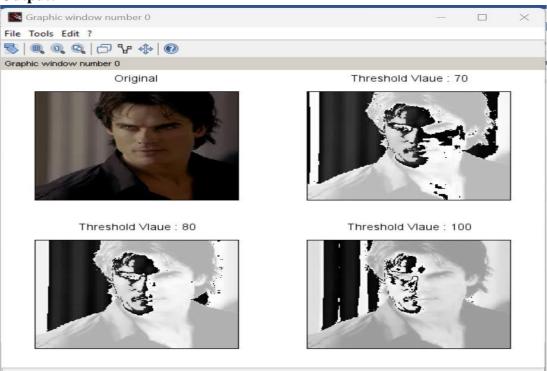


# 2) Spatial Resolution Code:

```
i=imread('D:\damon.jpeg');
a=imresize(i,0.8);
b=imresize(i,0.6);
c=imresize(i,0.4);
subplot(2,2,1),imshow(i),title('Original');
subplot(2,2,2),imshow(a),title('0.8');
subplot(2,2,3),imshow(c),title('0.6');
subplot(2,2,4),imshow(c),title('0.4');
```



```
Aim: Image Enhancement
A) Thresholding
code:
a=<u>imread</u>('D:\damon.jpeg');
[m,n]=size(a);
for i=1:m
for j=1:n
x=a(i,j);
if x > = 128
b(i,j)=a(i,j)+70;
c(i,j)=a(i,j)+80;
d(i,j)=a(i,j)+100;
else
b(i,j)=a(i,j)-70;
c(i,j)=a(i,j)-80;
d(i,j)=a(i,j)-100;
end
end
end
subplot(2,2,1),imshow(a),title('Original');
subplot(2,2,2),imshow(b),title('Threshold Vlaue : 70');
subplot(2,2,3),imshow(c),title('Threshold Vlaue : 80');
subplot(2,2,4),imshow(d),title('Threshold Vlaue : 100');
```



# B) Contrast Adjustment: Code:

```
a=imread('D:\damon.jpeg');
r1=100;
r2=140;
s1=150;
s2=240;
1=s1/r1;
m=(s2-s1)/(r2-r1);
n=(255-s2)/(255-r2);
s=size(a);
for i=1:s(1)
for j=1:s(2)
if ((a(i,j) > 0) && (a(i,j) < r1))
b(i,j) = a(i,j)*l;
end
if ((a(i,j) > r1) && (a(i,j) < r2))
b(i,j) = (m*(a(i,j)-120))+s1;
if ((a(i,j) > r2) && (a(i,j) < 256))
b(i,j) = (n*(a(i,j)-150))+s2;
end
end
end
subplot(1,2,1),imshow(a),title('Original Image');
subplot(1,2,2),imshow(uint8(b)),title('Contrast Image');
```



Original Image



Contrast Image



# C) Brightness Adjustment:

### Code:

```
a=\underline{imread}('D:\damon.jpeg');\\[m,n]=size(a);\\for i=1:m\\for j=1:n\\b(i,j)=a(i,j)-50;\\c(i,j)=a(i,j)-100;\\d(i,j)=a(i,j)+50;\\end\\end\\subplot(2,2,1),\underline{imshow}(a),\underline{title}('Original');\\subplot(2,2,2),\underline{imshow}(b),\underline{title}('Reduced By 50');\\subplot(2,2,3),\underline{imshow}(c),\underline{title}('Reduced By 100');\\subplot(2,2,4),\underline{imshow}(d),\underline{title}('Increase By 50');\\
```



Original





Reduced By 50



Increase By 50



# D) Gray Level Slicing: code:

```
a=imread('D:\damon.jpeg');
[m,n]=size(a);
min = 100;
max = 200;
for i=1:m
for j=1:n
x=a(i,j);
if x > \min \&\& x < \max
b(i,j)=a(i,j);
elseif x > max
b(i,j)=255;
else
b(i,j)=0;
end
end
end
subplot(1,2,1),imshow(a),title('Original');
subplot(1,2,2),imshow(b),title('Gray Slicing');
```



Original



Gray Slicing



**Aim: Basic Transformation** 

# Practical No. 04

```
A) Log Transformation:

Code:

a=imread('D:\klaroline.jpeg');

[m,n]=size(a);

for i=1:m

for j=1:n

x=a(i,j);

b(i,j)=20*log(1+double(x));

end

end

subplot(1,2,1),imshow(a),title('Original');

subplot(1,2,2),imshow(b),title('Log Transform');
```



Original

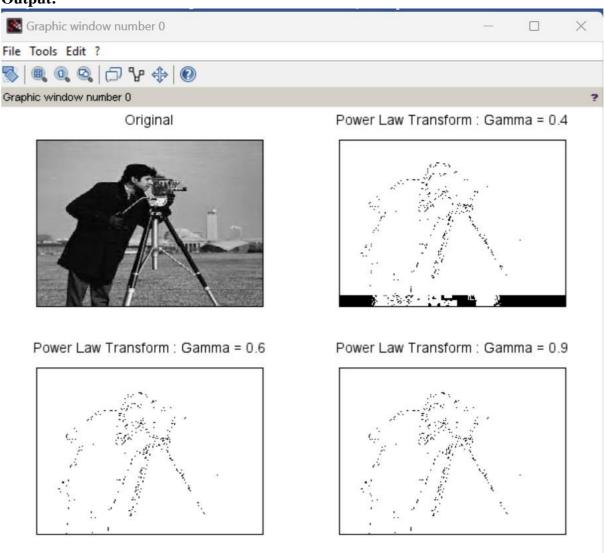


Log Transform



# B) Power Law Transformation: code:

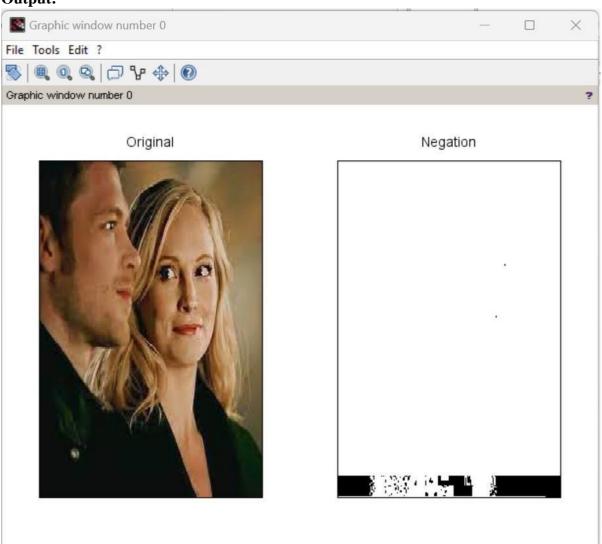
```
 \begin{array}{l} a=\underline{imread}('D:\langle cameraman.jpeg');\\ [m,n]=\underline{size}(a);\\ for \ i=1:m\\ for \ j=1:n\\ x=\underline{double}(a(i,j));\\ b(i,j)=20*(x^0.4);\\ c(i,j)=20*(x^0.6);\\ d(i,j)=20*(x^0.9);\\ end\\ end\\ \underline{subplot}(2,2,1),\underline{imshow}(a),\underline{title}('Original');\\ \underline{subplot}(2,2,2),\underline{imshow}(b),\underline{title}('Power \ Law\ Transform:\ Gamma=0.4');\\ \underline{subplot}(2,2,3),\underline{imshow}(c),\underline{title}('Power \ Law\ Transform:\ Gamma=0.6');\\ \underline{subplot}(2,2,4),\underline{imshow}(d),\underline{title}('Power \ Law\ Transform:\ Gamma=0.9');\\ \underline{subplot}(2,2,4),\underline{title}('Power \ Law\ Transform:\ Gamma=0.9');\\ \underline{subplot}(2,2,4),\underline{title}(2,2,4),\underline{title}(2,2,4),\underline{title}(2,2,4),\underline{title}(2,2,4),\underline{title}(2,2,4),\underline{title}(2,2,4),\underline{
```



# C) Negation code

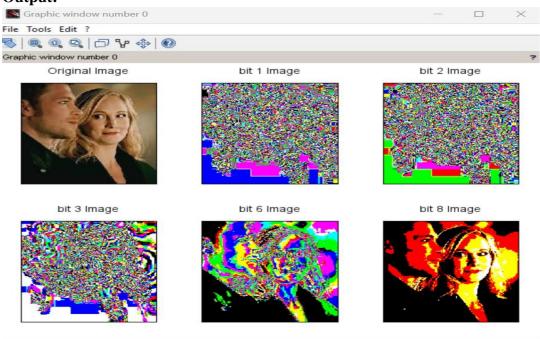
# code:

```
a=imread('D:\klaroline.jpeg');
[m,n]=size(a);
for i=1:m
for j=1:n
b(i,j)=255 - a(i,j);
end
end
subplot(1,2,1),imshow(a),title('Original');
subplot(1,2,2),imshow(b),title('Negation');
```



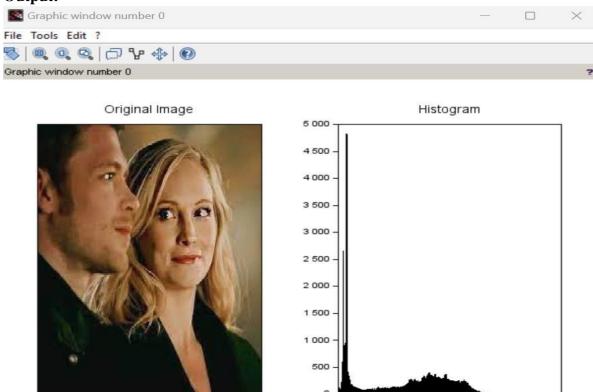
# D) Piecewise linear transformations code:

```
clc;
clear all;
a=<u>imread('D:\klaroline.jpeg');</u>
b=double(a);
subplot(2,3,1);
imshow(a);
title('Original Image');
f1=bitget(b,1);
subplot(2,3,2);
imshow(f1);
title('bit 1 Image');
f2=bitget(b,2);
subplot(2,3,3);
imshow(f2);
title('bit 2 Image');
f3=bitget(b,4);
\underline{\text{subplot}}(2,3,4);
imshow(f3);
title('bit 3 Image');
f4 = \underline{bitget(b,6)};
\underline{\text{subplot}}(2,3,5);
imshow(f4);
title('bit 6 Image');
f5=bitget(b,8);
subplot(2,3,6);
imshow(f5);
title('bit 8 Image');
```



# Aim: A) Write a program to plot a Histogram for Colour and Grayscale Images. Code:

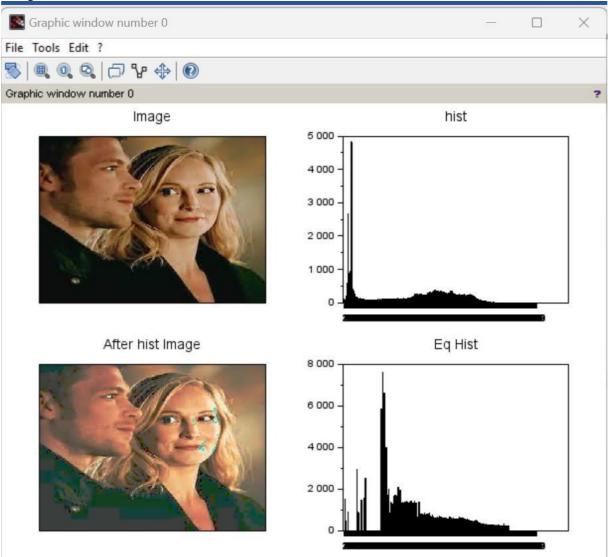
```
a = \underline{imread}('D:\klaroline.jpeg');
a = double(a);
[row col] = size(a);
h = zeros(1,300);
for n = 1:1:row
for m = 1:1:col
if a(n,m) == 0
a(n,m) = 1;
end
end
end
for n = 1:1:row
for m = 1:1:col
t = a(n,m);
h(t) = h(t) + 1;
end
end
subplot(1,2,1),imshow(uint8(a)); title('Original Image');
subplot(1,2,2),bar(h),title('Histogram');
```



# B) Write a program to apply histogram equalization.

### **Code:**

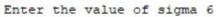
```
a = imread('D:\klaroline.jpeg');
a = double(a);
big = 256;
[row col d] = size(a);
c = row*col;
h = zeros(1,300);
z = zeros(1,300);
for e = 1:1:d
for n = 1:1:row
for m = 1:1:col
if a(n,m,e) == 0
a(n,m,e) = 1;
end
end
end
end
for n = 1:1:row
for m = 1:1:col
t = a(n,m);
h(t) = h(t) + 1;
end
end
pdf = h/c;
cdf(1) = pdf(1);
for x = 2:1:big
cdf(x) = pdf(x) + cdf(x-1);
end
new = round (cdf*big);
new = new + 1;
for r = 1:1:d
for p = 1:1:row
for q = 1:1:col
temp = a(p,q,r);
b(p,q,r) = new(temp);
t = b(p,q,r);
z(t) = z(t) + 1;
end
end
end
b = b-1;
subplot(2,2,1); imshow(uint8(a)); title('Image');
subplot(2,2,2); bar(h); title('hist');
subplot(2,2,3); imshow(uint8(b)); title('After hist Image');
subplot(2,2,4); bar(z); title('Eq Hist');
```



# Aim: Write a program to apply Gaussian filter on an image. Code:

```
m=input('Enter the Size ');
s=input('Enter the value of sigma');
sum1=0;
a=m/2;
p=0;q=0;
r=1;
t=1;
w=floor(a);
for i=-w:w
for j=-w:w
p=i*i;
q=j*j;
g(r,t) = \exp(-(p+q)/(2*s*s));
sum1=sum(sum(g(r,t)+sum1));
t=t+1;
end
t=1;
r=r+1;
end
for r=1:m
for t=1:m
h(r,t)=g(r,t)/sum1;
t=t+1;
end
t=1:
r=r+1;
end
im=imread('D:\cameraman.jpeg');
p=double(im);
s1=0;
[M N] = size(p);
for x=0:M-m
for y=0:N-m
for s=1:m
for z=1:m
s1=(h(s,z)*(p(x+s,y+z)))+s1;
end
end
N_{img(x+1,y+1)=s1};
s1=0;
end
end
subplot(1,2,1),imshow(uint8(im)),title('Original Image');
subplot(1,2,2),imshow(uint8(N_img)),title('Image After Gaussian Filter');
```

Output: Enter the Size 20





Aim: 1) Write a program to apply following morphological operations on the image.

# A. Opening Code:

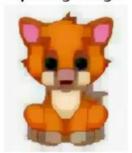
```
img=imread('cameraman.tif');
se1 = strel('square',11);
im2 = imerode(img,se1);
im3 = imdilate(im2,se1);
subplot(1,2,1),imshow(img),title('orignal image');
subplot(1,2,2),imshow(im3),title('opening image');
```

### **Output:**

### orignal image



opening image



# **B.** Closing Code:

```
aa=imread('cameraman.tif');
se1=strel('square',11);
IM2=imdilate(aa,se1);
IM3=imerode(IM2,se1);
subplot(1,2,1),imshow(aa),title('Original Image');
subplot(1,2,2),imshow(IM3),title('Closed Image');
```

Original Image



**Closed Image** 



# **C.Morphological Gradient**

### Code:

img=imread('cameraman.tif');
se1=strel('square',12);
im1=imdilate(img,se1);
im2=imerode(im1,se1);
g=im1-im2;
subplot(2,2,1),imshow(img),title('Orignal Image');
subplot(2,2,2),imshow(im1),title('Dilation Image');
subplot(2,2,3),imshow(im2),title('Erotion Image');
subplot(2,2,4),imshow(g),title('Gradient Image');

# **Output:**

# Orignal Image



# Dilation Image



# Erotion Image



**Gradient Image** 



# **D.Top-hat transformation** Code:

i=imread('cameraman.tif');
se1=strel('square',22);
im1=imerode(i,se1);
im2=imdilate(im1,se1);
h=i-im2;
subplot(2,2,1),imshow(i),title('Orignal Image');
subplot(2,2,2),imshow(im1),title('Erotion Image');
subplot(2,2,3),imshow(im2),title('Dilation Image');
subplot(2,2,4),imshow(h),title('Top Hat Transformation Image');

# **Output:**

# Orignal Image



# **Erotion Image**



# Dilation ImageTop Hat Transformation Ima





# Aim: 2) Write a program for boundary detection.

# **Code:**

clear all; clc; aa=imread('moon.jpeg'); se1=strel('square',11); subplot(2,1,1),imshow(aa); m1=imerode(aa,se1); m2=aa-m1; title('orignal image'); subplot(2,1,2),imshow(m2); title('edge detection');

# **Output:**

# orignal image



# edge detection



# Aim: A) Write a program to show RGB planes

### **Code:**

```
original=<u>imread('D:\klaroline.jpeg');</u>
im_red=original(:,:,1);
im_green=original(:,:,1);
im_blue=original(:,:,3);
subplot(2,2,1),imshow(original),title('Original');
subplot(2,2,2),imshow(im_red),title('Red');
subplot(2,2,3),imshow(im_green),title('Green');
subplot(2,2,4),imshow(im_blue),title('Blue');
```

# **Output:**



### Original



Red



Green



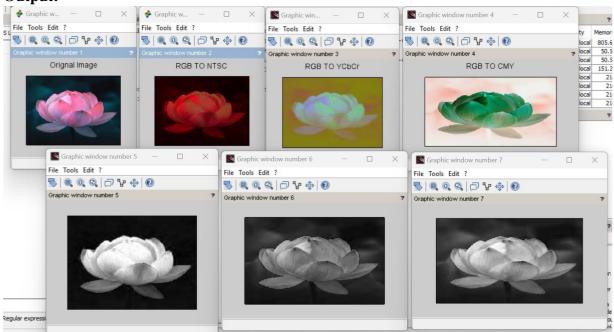
Blue



```
Aim: B) WAP to convert
RGB to NTSC
RGB to YCbCr
RGB to CMY
Code:
clc:
clear all;
close all;
a = imread('D:\lotus.jpeg');
figure(1),imshow(a);
title('Orignal Image');
k=rgb2ntsc(a);
figure(2),imshow(k);
title('RGB TO NTSC');
l=rgb2ycbcr(a);
figure(3), imshow(1);
title('RGB TO YCbCr');
m=imcomplement(a);
figure(4), imshow(m);
title('RGB TO CMY');
imr=a(:,:,1);
img=a(:,:,2);
imb=a(:,:,3);
figure(5), imshow(imr);
figure(6), imshow(img);
figure(7), imshow(imb);
I=(imr+img+imb)/3;
[m,n]=size(imr);
for c=1:m
for d=1:n
min1=min(imr(c,d),img(c,d));
min2=min(min1,imb(c,d));
S(c,d) = 1-(3/(imr(c,d)+img(c,d)+imb(c,d)))*min2;
end
end
for c=1:m
for d=1:n
temp= (0.5*(imr(c,d)-img(c,d))+(imr(c,d)-img(c,d))
imb(c,d))/sqrt(double(imr(c,d)*imr(c,d)+(imr(c,d)-imb(c,d))*(img(c,d)-imb(c,d))));
H(c,d)=acos(double(temp));
end
end
for c=1:m
for d=1:n
finali(c,d,1)=I(c,d);
finali(c,d,2)=S(c,d);
finali(c,d,3)=H(c,d);
end
end
```

# **Image Processing**

figure(8),imshow(finali);
title('Final image');



# Aim: Write a program to achieve Pseudo coloring. Code:

```
a=imread('D:\lotus.jpeg');
[1,m,n]=size(a);
for i=1:1
for i=1:m
for k=1:n
if a(i,j) >= 0 & a(i,j) < 50
b(i,j,1)=a(i,j,1)+50;
b(i,j,2)=a(i,j,1)+100;
b(i,j,3)=a(i,j,1)+10;
end
if a(i,j) > = 50 \& a(i,j) < 100
b(i,j,1)=a(i,j,1)+35;
b(i,j,2)=a(i,j,1)+128;
b(i,j,3)=a(i,j,1)+10;
end
if a(i,j) > = 100 \& a(i,j) < 150
b(i,j,1)=a(i,j,1)+152;
b(i,j,2)=a(i,j,1)+130;
b(i,j,3)=a(i,j,1)+15;
end
if a(i,j) > = 150 \& a(i,j) < 200
b(i,j,1)=a(i,j,1)+50;
b(i,j,2)=a(i,j,1)+140;
b(i,j,3)=a(i,j,1)+25;
end
if a(i,j) > = 200 \& a(i,j) < 256
b(i,j,1)=a(i,j,1)+120;
b(i,j,2)=a(i,j,1)+160;
b(i,j,3)=a(i,j,1)+45;
end
end
end
end
subplot(1,2,1),imshow(a),title('Original');
subplot(1,2,2),imshow(b),title('Pseudo Image');
```

