

DEEP LEARNING - SE4050

ASSIGNMENT 02

B.Sc. (Hons) Degree in Information Technology

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Problem

Short Message Service (SMS) has become one of the most common and heavily used ways of communication in society today. According to the research conducted by Ghourabi, Mahmood, and Alzubi, mobile phone users around the world have sent more than 8.3 trillion SMS messages in the year 2017 [1]. Furthermore, the number of SMS messages sent monthly is more than 690 billion [1].

However, SMS spam has been widely spreading among most mobile phone users recently. Any undesired or unsolicited text message sent indiscriminately to your mobile phone, usually for commercial motives, is referred to as SMS spam [1, 2, 3]. According to a survey conducted in the research paper [1], more than 68% of mobile phone users around the globe are affected by SMS spam messages.

Most people face many difficulties in their day-to-day lives due to SMS spam messages. Spams annoy most people as their valuable time is wasted and their workflow is interrupted and disturbed when they have to go through unwanted messages [3]. Additionally, the important text messages can be missed due to the inbox being filled with unwanted spam SMS messages. Furthermore, these SMS spams waste network resources of the device [3]. Therefore, SMS spam can cause significant negative sociological and economical effects.

Furthermore, in many cases, SMS spams consist of malicious activities such as smishing (SMS + phishing) which is a cyber security threat for mobile users aimed at deceiving them via SMS spam messages that may include a link or malicious software [1]. Cyber-attackers are trying to steal users' secret and sensitive information such as credit card numbers, passwords, and bank account details [1] using these malicious SMS spam messages. Many individuals and even major organizations suffer huge financial losses due to the cyber security threats caused by SMS spam messages [1].

The filtration of SMS spam in smartphones is still not very robust compared to the filtration of email spam detection [1]. Identification of text spam messages is also proven to be a very hard and time-consuming task according to most research [1, 2, 3]. Most existing lexicon-based methodologies as well as traditional NLP (Natural Language Processing) and Machine Learning algorithms are not accurate and efficient enough. Therefore, addressing this real-world problem and coming up with a reliable technique to classify spam SMS messages from ham SMS messages can be very useful.

Dataset

Background

Link: http://archive.ics.uci.edu/ml/datasets/SMS+Spam+Collection

The dataset which we are going to use is the **SMS Spam Collection Dataset** which is hosted on UCI Machine Learning Repository [4, 5, 6]. The dataset has been published on the 22nd of June 2012 [4]. It contains 5,574 English SMS phone messages and each of these messages have already been labeled as either ham (legitimate) or spam [4, 6] as shown in Figure 1, Figure 2, and Figure 3. The main purpose of the SMS Spam Collection Dataset is to assist NLP and computational linguistics research focused on detecting mobile phone SMS text message spams [4].



Figure 1: Sample data instances of SMS Spam Collection Dataset

The dataset has been constructed using multiple free resources from the internet. Out of the total of 5,574 messages, 425 SMS messages were extracted manually from the Grumbletext website [7] which is a forum for mobile phone users to discuss SMS spam messages. Further, 3,375 SMS ham messages have been chosen randomly from the NUS SMS Corpus (NSC) [8] which consists of about 10,000 ham SMS messages collected mostly from Singaporean students

studying in the Department of Computer Science at the National University of Singapore [4]. Additionally, 450 SMS ham messages have been gathered from the Ph.D. thesis of Dr. Caroline Tagg [4]. Furthermore, 1,002 SMS ham messages and 322 spam messages have been accumulated from SMS Spam Corpus v.0.1 Big [4].

Figure 2: Concise summary of the dataframe

	sms_message	class
count	5572	5572
unique	5169	2
top	Sorry, I'll call later	ham
freq	30	4825

Figure 3: Descriptive statistics of the dataframe

Attributes

The SMS Spam Collection Dataset is composed of one text file consisting of only two columns.

1. SMS message text

This is the only input feature of the dataset. It contains the raw text of the mobile phone SMS message. The data type of this attribute is a string.

We have analyzed the text length distribution of each of these SMS messages in terms of the character count and plotted a graph as shown in Figure 4.

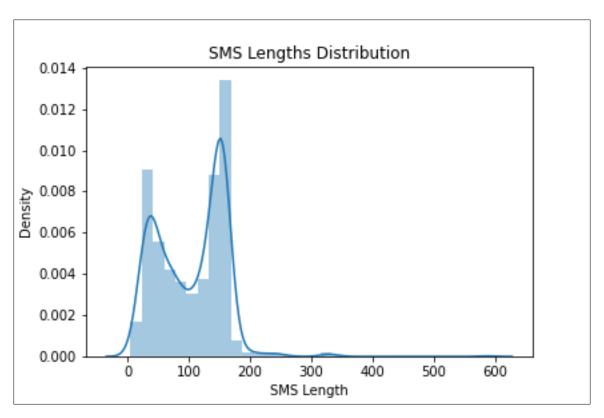


Figure 4: SMS lengths distribution plot

2. Label

This is the target attribute (class label) of the dataset. It has two distinct values: "ham" and "spam". The data type of this attribute is a string.

We have analyzed the data distribution of the target attribute of the dataset and plotted the distributions in a pie chart.

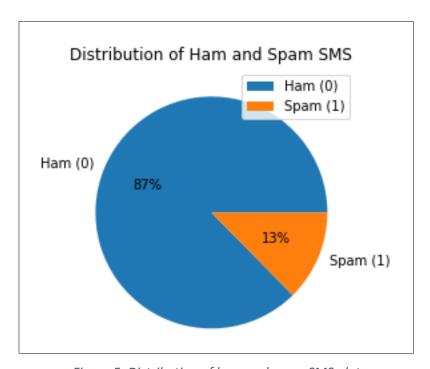


Figure 5: Distribution of ham and spam SMS plot

As shown in Figure 5, about 87% of SMS messages in the dataset are hams and the remaining 13% of SMS messages are spam. Therefore, the majority of the SMS messages in this dataset are labeled as ham as shown in Figure 6.

	class	ham	spam
sms_message	count	4825	747
	unique	4516	653
			Please call our customer service representative on FREEPHONE 0808 145 4742 between 9am-11pm as you have WON a guaranteed £1000 cash or £5000 prize!
	freq	30	4

Figure 6: Descriptive statistics for each class of the dataframe

We further generated the Word Cloud plots for ham and spam SMS messages separately to visualize the most used words in ham and spam SMS messages as shown in Figure 3 and Figure 4 respectively.



Figure 7: Ham SMS Word Cloud plot



Figure 8: Spam SMS Word Cloud plot

Methodology

Data Cleaning and Preprocessing

Real-world data sets contain several problems such as missing or null values, data inconsistency, incompleteness, and outliers of the dataset. Therefore, data preprocessing is a mandatory step before feeding the dataset into a deep learning model. Data preprocessing includes transforming row data into the machine-understandable and efficient format. Since the SMS Spam Collection dataset may also contain some of these issues, we had to do preprocessing tasks before using the dataset for model training purposes.

Initially, the dataset was downloaded and extracted.

```
# downloading UCI SMS Spam Collection dataset
!wget --no-check-certificate https://archive.ics.uci.edu/ml/machine-
learning-databases/00228/smsspamcollection.zip
# extracting the downloaded dataset
!unzip /content/smsspamcollection.zip
```

Then the dataset was imported to a Pandas dataframe.

Then the data was analyzed to check whether there are any missing or null values existed within the dataset. If any null values existed within the dataset, there are usually two ways to deal with them. The first method is to delete the particular rows which contain multiple null values. The second method is to replace the missing values with mean, median, or most frequent values. Although our analysis proved that the SMS Spam Collection dataset did not contain any missing or null values as shown in Figure 9, we implemented the null or missing values removal step programmatically.

```
sms_spam_dataframe =
sms_spam_dataframe[sms_spam_dataframe.notna().all(axis=1)]
```

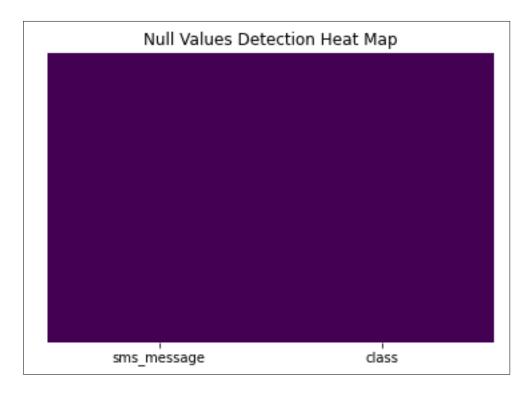


Figure 9: Null values detection heat map

Furthermore, the dataset was checked to identify duplicate rows.

```
# detecting duplicate rows exist in the dataframe before cleaning
duplicated_records = sms_spam_dataframe[sms_spam_dataframe.duplicated()]

# checking the number of duplicate rows exist in the dataframe
# before cleaning
sms_spam_dataframe.duplicated().sum()
```

Since there were 403 duplicates in the data frame, we implemented the duplicate row removal step.

```
# removing the duplicate rows from the dataframe if exist
sms_spam_dataframe = sms_spam_dataframe.drop_duplicates()
```

When we analyzed the data distribution of ham and spam SMS messages, it clearly shows that the data is imbalanced between the two classes as shown in Figure 10.

```
# printing count of values in each class of the dataframe
cleaned_sms_spam_dataframe['class'].value_counts()
ham     4516
spam     653
```

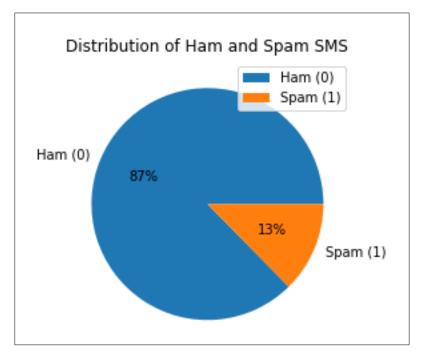


Figure 10: Distribution of ham and spam SMS plot before downsampling

To address this problem of imbalanced data, we decided to apply the downsampling technique which is a process where you randomly delete some of the observations from the majority class (ham) so that the numbers in majority and minority (spam) classes are matched. The dataset is separated into two dataframes based on the class label, and the majority class is downsampled.

```
# extracting the data instances with class label 'spam'
spam_dataframe =
cleaned_sms_spam_dataframe[cleaned_sms_spam_dataframe['class'] == 'spam']
# extracting the data instances with class label 'ham'
ham_dataframe =
cleaned_sms_spam_dataframe[cleaned_sms_spam_dataframe['class'] == 'ham']
```

After applying downsampling and merging the two dataframes, there were the same amount of 653 messages for each class.

```
# merging the two dataframes (spam + downsampled ham dataframes)
merged_dataframe = pd.concat([downsampled_ham_dataframe, spam_dataframe])
# printing count of values in each class of the merged dataframe
merged_dataframe['class'].value_counts()

spam 653
ham 653
```

As shown in Figure 11, now the dataframe is balanced with 50% of data instances for each class.

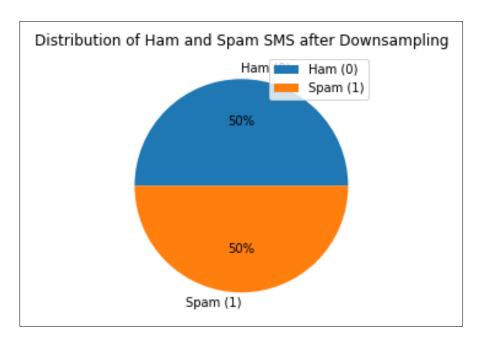


Figure 11: Distribution of ham and spam SMS plot after downsampling

Then we inserted a new column to the dataframe containing the length of the SMS message texts in terms of the number of characters.

```
# inserting a new column called 'length' to the merged dataframe
# the column contains the number of characters of the sms_message text
merged_dataframe['length'] = merged_dataframe['sms_message'].apply(len)
```

Then the text labels "ham" and "spam" were converted to numeric values 0 and 1 respectively as shown in Table 1.

```
# inserting a new column called 'label' to the merged dataframe
# if class is 'ham' label = 0
# if class is 'spam' label = 1
merged_dataframe['label'] = merged_dataframe['class'].map({'ham': 0, 'spam': 1})
```

Table 1: Numeric value for textual class labels

Class	Label
Ham	0
Spam	1

Then the dataset was split into train and test sets. The training set was 80% of the entire dataset while the test set was 20% of the entire dataset. Therefore, out of 1306 total records, 1044 records were selected as training set and 262 records were selected as testing set.

Table 2: Row count after train test split

Total dataset rows	1306
Train dataset rows	1044
Test dataset rows	262

References

- [1] A. Ghourabi, M. A. Mahmood, and Q. M. Alzubi, "A Hybrid CNN-LSTM Model for SMS Spam Detection in Arabic and English Messages," Future Internet, vol. 12, no. 9, p. 156, Sep. 2020, doi: 10.3390/fi12090156.
- [2] J. Ma, Y. Zhang, J. Liu, K. Yu, and X. Wang, "Intelligent SMS Spam Filtering Using Topic Model," 2016 International Conference on Intelligent Networking and Collaborative Systems (INCoS), 2016, pp. 380-383, doi: 10.1109/INCoS.2016.47.
- [3] M. Popovac, M. Karanovic, S. Sladojevic, M. Arsenovic, and A. Anderla, "Convolutional Neural Network Based SMS Spam Detection," 2018 26th Telecommunications Forum (TELFOR), 2018, pp. 1-4, doi: 10.1109/TELFOR.2018.8611916.
- [4] Archive.ics.uci.edu. 2012. UCI Machine Learning Repository: SMS Spam Collection Data Set. [online] Available at: http://archive.ics.uci.edu/ml/datasets/SMS+Spam+Collection.
- [5] "SMS Spam Collection Dataset", Kaggle.com, 2016. [online]. Available: https://www.kaggle.com/uciml/sms-spam-collection-dataset.
- [6] T. A. Almeida, J. M. G. Hidalgo, and A. Yamakami, "Contributions to the study of SMS spam filtering," presented at the 11th ACM symposium, 2011. doi: 10.1145/2034691.2034742.
- [7] Grumbletext.co.uk, 2021. [Online]. Available: http://www.grumbletext.co.uk/.
- [8] Comp.nus.edu.sg, 2021. [Online]. Available: https://www.comp.nus.edu.sg/.
- [9] a
- [10] a
- [11] a
- [12] a
- [13] a
- [14] a
- [15] a
- [16] a
- [17] a
- [18] a

Appendixes

#88 md

SMS Spam Detection with Deep Learning

Short Message Service (SMS) is being heavily used as a way of communication. However, SMS spams have targeted most mobile phone users recently. In some cases, SMS spams contain malicious activities such as smishing. Smishing (SMS + Phishing) is a cyber-security threat for mobile users aimed at deceiving them via SMS spam messages that may include a link or malicious software or both. The attackers attempt to steal users' secret and sensitive information, like credit card numbers, bank account details, and passwords.

The filtration of SMS spams in smartphones is still not very robust compared to the filtration of email spams. The state-of-the-art methodologies based on Deep Learning can be utilized for solving this binary classification problem of SMS spam detection. We have used the two deep neural network architectures namely Long Short-Term Memory (LSTM) and DenseNet (Densely Connected Convolutional Neural Network (CNN)) for this purpose.

```
# 응 응
```

```
# magic function that renders the figure in a notebook instead of
# displaying a dump of the figure object
# sets the backend of matplotlib to the 'inline' backend
# with this backend, the output of plotting commands is displayed
# inline within frontends like the Jupyter notebook, directly below
# the code cell that produced it
# the resulting plots will then also be stored in the notebook document
%matplotlib inline
# 응 응
# creating a new directory named plots
!mkdir plots
# creating a new directory named models
!mkdir models
# creating a new directory named processed datasets
!mkdir processed datasets
# 응 응
# importing warnings library to handle exceptions, errors, and warning
# of the program
import warnings
```

```
# ignoring potential warnings of the program
warnings.filterwarnings('ignore')
# % %
# importing pandas library to perform data manipulation and analysis
import pandas as pd
# configuring the pandas dataframes to show all columns
pd.options.display.max columns = None
# configuring the pandas dataframes to increase the maximum column width
pd.options.display.max colwidth = 150
# % %
# downloading UCI SMS Spam Collection dataset
!wget --no-check-certificate https://archive.ics.uci.edu/ml/machine-
learning-databases/00228/smsspamcollection.zip
# 응 응
# extracting the downloaded dataset
!unzip /content/smsspamcollection.zip
# 응 응
# listing files and directories
#%% md
# SMS Spam Collection Dataset
The SMS Spam Collection Dataset was downloaded from UCI datasets. It
contains 5,574 SMS phone messages. The data were collected for the purpose
of mobile phone SMS text message spam research and have already been
labeled as either spam or ham.
Link to the dataset -
http://archive.ics.uci.edu/ml/datasets/SMS+Spam+Collection
# 응 응
# importing SMSSpamCollection dataset to a pandas dataframe
sms spam dataframe = pd.read csv('/content/SMSSpamCollection',
                                 sep='\t',
                                 header=None,
                                 names=['class', 'sms_message'])
sms spam dataframe
#88 md
```

Dataset Analysis and Data Preprocessing

```
# 응 응
# printing the columns of the dataframe
sms spam dataframe.columns
# % %
# changing the order of the dataframe columns for better visualization
sms spam dataframe = sms spam dataframe[['sms message', 'class']]
sms spam dataframe
# % %
# displaying the dimensionality of the dataframe
sms spam dataframe.shape
# 응 응
# printing a concise summary of the dataframe
# information such as index, data type, columns, non-null values,
# and memory usage
sms spam dataframe.info()
# 응 응
# generating descriptive statistics of the dataframe
sms spam dataframe.describe()
# 응 응
# generating descriptive statistics for each class of the dataframe
# T property is used to transpose index and columns of the dataframe
sms spam dataframe.groupby('class').describe().T
# 응 응
# checking for missing or null values in the dataframe
dataframe null =
sms spam dataframe[sms spam dataframe.isnull().any(axis=1)]
dataframe null
# % %
# printing the number of rows with any missing or null values
# in the dataframe
dataframe null.shape[0]
# % %
# removing the missing or null values from the dataframe if exist
sms spam dataframe =
sms spam dataframe[sms spam dataframe.notna().all(axis=1)]
```

```
# printing the count of null values in class and sms message
# columns of the dataframe
sms spam dataframe[['class', 'sms message']].isnull().sum()
# 응 응
# importing pyplot from matplotlib library to create interactive
# visualizations
import matplotlib.pyplot as plt
# importing seaborn library which is built on top of matplotlib to
# create statistical graphics
import seaborn as sns
# plotting the heatmap for missing or null values in the dataframe
sns.heatmap(sms spam dataframe.isnull(),
            yticklabels=False,
            cbar=False,
            cmap='viridis')
plt.title('Null Values Detection Heat Map')
plt.savefig('plots/null detection_heat_map.png',
            facecolor='white')
plt.show()
# 응 응
# importing missingno library
# used to understand the distribution of missing values through
# informative visualizations
# visualizations can be in the form of heat maps or bar charts
# used to observe where the missing values have occurred
# used to check the correlation of the columns containing the missing
# with the target column
import missingno as msno
# plotting a matrix visualization of the nullity of the dataframe
fig = msno.matrix(sms spam dataframe)
fig copy = fig.get figure()
fig copy.savefig('plots/msno matrix.png',
                 bbox inches='tight')
fiq
# % %
# plotting a seaborn heatmap visualization of nullity correlation
# in the dataframe
fig = msno.heatmap(sms spam dataframe)
fig copy = fig.get figure()
fig copy.savefig('plots/msno_heatmap.png',
                 bbox inches='tight')
fiq
# % %
```

```
# detecting duplicate rows exist in the dataframe before cleaning
duplicated records = sms spam dataframe[sms spam dataframe.duplicated()]
duplicated records
# 응 응
# checking the number of duplicate rows exist in the dataframe
# before cleaning
sms spam dataframe.duplicated().sum()
# % %
# removing the duplicate rows from the dataframe if exist
sms spam dataframe = sms spam dataframe.drop duplicates()
sms spam dataframe
# % %
# checking the number of duplicate rows exist in the dataframe
# after cleaning
sms spam dataframe.duplicated().sum()
# % %
# displaying the dimensionality of the dataframe
sms spam dataframe.shape
# 응 응
# printing a concise summary of the dataframe
# information such as index, data type, columns, non-null values,
# and memory usage
sms spam dataframe.info()
# % %
# generating descriptive statistics of the dataframe
sms spam dataframe.describe()
# % %
# generating descriptive statistics for each class of the dataframe
# T property is used to transpose index and columns of the dataframe
sms spam dataframe.groupby('class').describe().T
# % %
# saving cleaned dataset to a csv file
file name = 'processed datasets/cleaned dataset.csv'
sms spam dataframe.to csv(file name,
                          encoding='utf-8',
                          index=False)
# loading dataset from the saved csv file to a pandas dataframe
```

```
cleaned sms spam dataframe = pd.read csv(file name)
cleaned sms spam dataframe
# 응 응
# importing set of stopwords from wordcloud library
from wordcloud import STOPWORDS
stopwords = set(STOPWORDS)
# printing number of stopwords defined in wordcloud library
len(stopwords)
# 응 응
# importing random library
# used for generating random numbers
import random
# printing 10 random values of stopwords set
for i, val in enumerate(random.sample(stopwords, 10)):
   print(val)
# 응 응
# importing WordCloud object for generating and drawing
# wordclouds from wordcloud library
from wordcloud import WordCloud
# defining a function to return the wordcloud for a given text
def plot wordcloud(text):
    wordcloud = WordCloud(width=600,
                          height=300,
                          background color='black',
                          stopwords=stopwords,
                          \max font size=50,
                          colormap='Oranges').generate(text)
    return wordcloud
# 응 응
# extracting the data instances with class label 'ham'
ham dataframe =
cleaned sms spam dataframe[cleaned sms spam dataframe['class'] == 'ham']
ham dataframe
# % %
# creating numpy list to visualize using wordcloud
ham sms message text = '
'.join(ham dataframe['sms message'].to numpy().tolist())
# generating wordcloud for ham sms messages
```

```
ham sms wordcloud = plot wordcloud(ham sms message text)
plt.figure(figsize=(16, 10))
plt.imshow(ham sms wordcloud,
           interpolation='bilinear')
plt.axis('off')
plt.title('Ham SMS Wordcloud')
plt.savefig('plots/ham wordcloud.png',
            facecolor='white')
plt.show()
# 응 응
# extracting the data instances with class label 'spam'
spam dataframe =
cleaned sms spam dataframe[cleaned sms spam dataframe['class'] == 'spam']
spam dataframe
# % %
# creating numpy list to visualize using wordcloud
spam sms message text = '
'.join(spam dataframe['sms message'].to numpy().tolist())
# generating wordcloud for spam sms messages
spam sms wordcloud = plot wordcloud(spam sms message text)
plt.figure(figsize=(16, 10))
plt.imshow(spam sms wordcloud,
           interpolation='bilinear')
plt.axis('off')
plt.title('Spam SMS Wordcloud')
plt.savefig('plots/spam wordcloud.png',
            facecolor='white')
plt.show()
# 응 응
# printing count of values in each class of the dataframe
cleaned sms spam dataframe['class'].value counts()
# % %
# plotting the distribution of target values
fig = plt.figure()
lbl = ['Ham (0)', 'Spam (1)']
pct = '%1.0f%%'
ax = cleaned sms spam dataframe['class'].value counts().plot(kind='pie',
                                                               labels=lbl,
                                                               autopct=pct)
ax.yaxis.set visible(False)
plt.title('Distribution of Ham and Spam SMS')
plt.legend()
fig.savefig('plots/ham spam pie chart.png',
            facecolor='white')
plt.show()
```

```
# 응 응
# downsampling is a process where you randomly delete some of the
# observations from the majority class so that the numbers in majority
# and minority classes are matched
# after downsampling the ham messages (majority class), there are now
# 653 messages in each class
downsampled ham dataframe = ham dataframe.sample(n=len(spam dataframe),
                                                  random state=44)
downsampled ham dataframe
# 응 응
# printing the dimensions of spam and downsampled ham dataframes
print('Spam dataframe shape:', spam dataframe.shape)
print('Ham dataframe shape:', downsampled ham dataframe.shape)
# % %
# merging the two dataframes (spam + downsampled ham dataframes)
merged dataframe = pd.concat([downsampled ham dataframe, spam dataframe])
merged dataframe = merged dataframe.reset index(drop=True)
merged dataframe
# 응 응
# printing count of values in each class of the merged dataframe
merged dataframe['class'].value counts()
# 응 응
# plotting the distribution of target values after downsampling
fig = plt.figure()
lbl = ['Ham (0)', 'Spam (1)']
pct = '%1.0f%%'
ax = merged dataframe['class'].value counts().plot(kind='pie',
                                                    labels=lbl,
                                                    autopct=pct)
ax.yaxis.set visible(False)
plt.title('Distribution of Ham and Spam SMS after Downsampling')
plt.legend()
fig.savefig('plots/ham spam pie chart after downsampling.png',
            facecolor='white')
plt.show()
# % %
# inserting a new column called 'label' to the merged dataframe
# if class is 'ham' label = 0
# if class is 'spam' label = 1
merged dataframe['label'] = merged dataframe['class'].map({'ham': 0,
'spam': 1})
merged dataframe
```

```
# 응 응
# inserting a new column called 'length' to the merged dataframe
# the column contains the number of characters of the sms message text
merged dataframe['length'] = merged dataframe['sms message'].apply(len)
merged dataframe
# % %
# displaying the first 5 rows of the dataframe
merged dataframe.head()
# 응 응
# displaying the last 5 rows of the dataframe
merged dataframe.tail()
# % %
# displaying the dimensionality of the dataframe
merged dataframe.shape
# 응 응
# printing a concise summary of the dataframe
# information such as index, data type, columns, non-null values,
# and memory usage
merged dataframe.info()
# 응 응
# generating descriptive statistics of the dataframe
merged dataframe.describe().round(2)
# 응 응
# generating descriptive statistics for each class of the dataframe
# T property is used to transpose index and columns of the dataframe
merged dataframe.groupby('label').describe().T
# 응 응
# generating descriptive sms text length statistics by label types
merged dataframe.groupby('label')['length'].describe().round(2)
# 응 응
# plotting a univariate distribution of observations for sms lengths
sns.distplot(merged dataframe['length'].values)
plt.title('SMS Lengths Distribution')
plt.xlabel('SMS Length')
plt.savefig('plots/sms length.png',
            facecolor='white')
```

```
plt.show()
# 응 응
# saving merged dataset to a csv file
file name = 'processed datasets/merged dataset.csv'
merged dataframe.to csv(file name,
                        encoding='utf-8',
                        index=False)
# loading dataset from the saved csv file to a pandas dataframe
merged dataframe = pd.read csv(file name)
merged dataframe
# 응 응
# assigning attributes (features) to X
X = merged dataframe['sms message']
# 응 응
# assigning label (target) to y
y = merged dataframe['label'].values
У
# % %
# importing train test split from scikit-learn library
from sklearn.model selection import train test split
# splitting data into random train and test subsets
# train set - 80%, test set - 20%
X train, X test, y train, y test = train test split(X,
                                                     У,
                                                     test size=0.2,
                                                     random state=443)
# printing the dimension of train features dataframe
print('Shape of train features dataframe:', X train.shape)
# printing the dimension of train target dataframe
print('Shape of train target dataframe:', y train.shape)
# printing the dimension of test features dataframe
print('Shape of test features dataframe:', X test.shape)
# printing the dimension of test target dataframe
print('Shape of test target dataframe:', y test.shape)
# 응 응
# displaying train features dataframe
X train
```

```
# 응 응
# displaying train target dataframe
y train
# % %
# displaying test features dataframe
X test
# 응 응
# displaying test target dataframe
y test
# % %
# defining pre-processing hyperparameters
# oov token defines the out of vocabulary token
# oov token will be added to word index in the corpus which is used to
# build the model
# this is used to replace out of vocabulary words (words that are not
# in our corpus) during text to sequence calls.
oov token = '<00V>'
# vocabulary size indicates the maximum number of unique words to
# tokenize and load in training and testing data
vocabulary size = 500
# 응 응
# importing Tokenizer from keras library
# tensorflow is a free and open-source software library for machine
# learning used across a range of machine learning related tasks
# focus on training and inference of deep neural networks
# keras is a high-level api of tensorflow
# keras.preprocessing.text provides keras data preprocessing utils
# to pre-process datasets with textual data before they are fed to the
# machine learning model
from tensorflow.keras.preprocessing.text import Tokenizer
# Tokenizer allows to vectorize a text corpus, by turning each text into
# either a sequence of integers (each integer being the index of a token
# in a dictionary) or into a vector where the coefficient for each token
# could be binary, based on word count, based on tf-idf
tokenizer = Tokenizer(num words=vocabulary size,
                      char level=False,
                      oov token=oov token)
# updating internal vocabulary based on a list of text required before
# using texts to sequences
tokenizer.fit on texts(X train)
```

```
# 응 응
# getting the word index
word index = tokenizer.word index
word index
# 응 응
# printing length of the word index
len(word index)
# % %
# transforming each text in train data to a sequence of integers
X train sequences = tokenizer.texts to sequences(X train)
# printing the first sequence
X train sequences[0]
# 응 응
# getting lengths of each generated sequences of integers
# in train data
x train length of sequence = [len(sequence) for sequence in
X train sequences]
# printing the length of the first sequence
x train length of sequence[0]
# 응 응
# importing numpy library
# used to perform fast mathematical operations over python arrays
# and lists
import numpy as np
# printing maximum length of a sequence in the train data
np.max(x train length of sequence)
# 응 응
# plotting a univariate distribution of observations for
# sequence lengths of train data
sns.distplot(x train length of sequence)
plt.title('Train Data Sequence Lengths Distribution')
plt.xlabel('Sequence Length')
plt.savefig('plots/train sequence length.png',
            facecolor='white')
plt.show()
# 응 응
# transforming each text in test data to a sequence of integers
```

```
X_test_sequences = tokenizer.texts to sequences(X test)
# printing the first sequence
X test sequences[0]
# 응 응
# getting lengths of each generated sequences of integers
# in test data
x test length of sequence = [len(sequence) for sequence in
X test sequences]
# printing the length of the first sequence
x test length of sequence[0]
# % %
# printing maximum length of a sequence in the test data
np.max(x test length of sequence)
# 응 응
# plotting a univariate distribution of observations for sequence
# lengths of test data
sns.distplot(x test length of sequence)
plt.title('Test Data Sequence Lengths Distribution')
plt.xlabel('Sequence Length')
plt.savefig('plots/test sequence length.png',
            facecolor='white')
plt.show()
# % %
# defining pre-processing hyperparameters
# maximum length indicates the maximum number of words considered
# in a text
maximum length = 50
# truncating type indicates removal of values from sequences larger
# than maxlen, either at the beginning ('pre') or at the end ('post')
# of the sequences
truncating type = 'post'
# padding type indicates pad either before ('pre') or after ('post')
# each sequence
padding type = 'post'
# 응 응
# importing utilities for preprocessing sequence data from
# keras library
from tensorflow.keras.preprocessing.sequence import pad sequences
```

```
# padding on train data
# pad sequences pads sequences to the same length
# padding='post' to pad after each sequence
X train padded = pad sequences(X train sequences,
                               maxlen=maximum length,
                               padding=padding type,
                               truncating=truncating type)
# printing the first padded sequence
X train padded[0]
# 응 응
# getting lengths of each padded sequences of integers in train
x train length of padded sequence = [len(sequence) for sequence in
X train padded]
# printing the length of the first padded sequence
x train length of padded sequence[0]
# 응 응
# printing maximum length of a padded sequence in the train data
np.max(x train length of padded sequence)
# % %
# printing the dimension of padded training dataframe
X train padded.shape
# 응 응
# plotting a univariate distribution of observations for sequence
# lengths of train data after padding
sns.distplot(x train length of padded sequence)
plt.title('Train Data Padded Sequence Lengths Distribution')
plt.xlabel('Sequence Length')
plt.savefig('plots/train padded sequence length.png',
            facecolor='white')
plt.show()
# 응 응
# padding on test data
# pad sequences pads sequences to the same length
# padding='post' to pad after each sequence
X test padded = pad sequences(X test sequences,
                              maxlen=maximum length,
                              padding=padding type,
                              truncating=truncating type)
# printing the first padded sequence
X test padded[0]
```

```
# 응 응
# getting lengths of each padded sequences of integers in test
x test length of padded sequence = [len(sequence) for sequence in
X test padded]
# printing the length of the first padded sequence
x test length of padded sequence[0]
# 응 응
# printing maximum length of a padded sequence in the test data
np.max(x test length of padded sequence)
# % %
# printing the dimension of padded test dataframe
X test padded.shape
# 응 응
# plotting a univariate distribution of observations for
# sequence lengths of test data after padding
sns.distplot(x test length of padded sequence)
plt.title('Test Data Padded Sequence Lengths Distribution')
plt.xlabel('Sequence Length')
plt.savefig('plots/test padded sequence length.png',
            facecolor='white')
plt.show()
#88 md
# LSTM Model
Long Short Term Memory (LSTM) is a special kind of Recurrent Neural
Network (RNN). LSTM models are explicitly designed to avoid the long-term
dependency problem by remembering information for long periods of time.
# % %
# LSTM network architecture hyperparameters
# SpatialDropout1D is used to dropout the embedding layer which helps
# to drop entire 1D feature maps instead of individual elements
# dropout rate indicates the fraction of the units to drop for the
# linear transformation of the inputs
dropout rate = 0.2
# n lstm indicates the number of nodes in the hidden layers within the
# LSTM cell
```

no of nodes = 20

```
# embedding dimension indicates the dimension of the state space used for
# reconstruction
embedding dimension = 16
# no of epochs indicates the number of complete passes through the
# training dataset
no of epochs = 30
# vocabulary size indicates the maximum number of unique words to
# tokenize and load in training and testing data
vocabulary size = 500
# % %
# importing Sequential class from keras
# Sequential groups a linear stack of layers into a keras Model
from tensorflow.keras.models import Sequential
# importing Embedding class from keras layers api package
# turning positive integers (indexes) into dense vectors of fixed size
# this layer can only be used as the first layer in a model
from tensorflow.keras.layers import Embedding
# importing LSTM class from keras layers api package
# LSTM - Long Short-Term Memory layer
from tensorflow.keras.layers import LSTM
# importing Dense class from keras layers api package
# Dense class is a regular densely-connected neural network layer
# Dense implements the operation:
# output = activation(dot(input, kernel) + bias)
# activation is the element-wise activation function passed as
# the activation argument
# kernel is a weights matrix created by the layer
# bias is a bias vector created by the layer
# (only applicable if use bias is True)
from tensorflow.keras.layers import Dense
# 응 응
# LSTM model architecture
lstm model = Sequential()
1stm model.add(Embedding(vocabulary size,
                         embedding dimension,
                         input length=maximum length))
# return sequences=True ensures that the LSTM cell returns all of the
# outputs from the unrolled LSTM cell through time
# if this argument is not used, the LSTM cell will simply provide the
# output of the LSTM cell from the previous step
lstm model.add(LSTM(no of nodes,
                    dropout=dropout rate,
```

```
return sequences=True))
1stm model.add(LSTM(no of nodes,
                    dropout=dropout rate,
                    return sequences=True))
# sigmoid is a non-linear and easy to work with activation function
# that takes a value as input and outputs another value between 0 and 1
lstm model.add(Dense(1,
                     activation='sigmoid'))
# 응 응
# compiling the model
# configuring the model for training
lstm model.compile(loss='binary crossentropy',
                   optimizer='adam',
                   metrics=['accuracy'])
# 응 응
# printing a string summary of the network
lstm model.summary()
# 응 응
# importing EarlyStopping class from callbacks module
# in keras library
# callbacks module includes utilities called at certain
# points during model training
# used to stop training when a monitored metric has
# stopped improving
from tensorflow.keras.callbacks import EarlyStopping
# monitoring the validation loss and if the validation loss is not
# improved after three epochs, then the model training is stopped
# it helps to avoid overfitting problem and indicates when to stop
# training before the deep learning model begins overfitting
early stopping = EarlyStopping(monitor='val loss',
                               patience=3)
# 응 응
# training the LSTM model
history = lstm model.fit(X train padded,
                         y train,
                         epochs=no of epochs,
                         validation data=(X test padded, y test),
                         callbacks=[early stopping],
                         verbose=2)
# % %
# visualizing the history results by reading as a dataframe
```

```
metrics lstm = pd.DataFrame(history.history)
metrics 1stm
# 응 응
# renaming the column names of the dataframe
metrics lstm.rename(columns={'loss': 'Training Loss',
                              'accuracy': 'Training Accuracy',
                              'val loss': 'Validation Loss',
                              'val accuracy': 'Validation Accuracy'},
                    inplace=True)
metrics 1stm
# 응 응
# plotting the training and validation loss by number of epochs for
# the LSTM model
metrics_lstm[['Training_Loss', 'Validation Loss']].plot()
plt.title('LSTM Model - Training and Validation Loss vs. Epochs')
plt.xlabel('Number of Epochs')
plt.ylabel('Loss')
plt.legend(['Training_Loss', 'Validation_Loss'])
plt.savefig('plots/lstm loss vs epochs.png',
            facecolor='white')
plt.show()
# 응 응
# plotting the training and validation accuracy by number of epochs for
# the LSTM model
metrics lstm[['Training Accuracy', 'Validation Accuracy']].plot()
plt.title('LSTM Model - Training and Validation Accuracy vs. Epochs')
plt.xlabel('Number of Epochs')
plt.ylabel('Accuracy')
plt.legend(['Training Accuracy', 'Validation Accuracy'])
plt.savefig('plots/lstm_accuracy_vs_epochs.png',
            facecolor='white')
plt.show()
# 응 응
# saving the trained LSTM model as an h5 file
# h5 is a file format to store structured data
# keras saves deep learning models in this format as it can easily store
# the weights and model configuration in a single file
lstm path = 'models/lstm model.h5'
lstm model.save(lstm path)
1stm model
# 응 응
# importing load model function from keras to load a saved keras
# deep learning model
from tensorflow.keras.models import load model
```

```
# loading the saved LSTM model
loaded lstm model = load model(lstm path)
loaded 1stm model
#88 md
# LSTM Model Evaluation
# % %
# evaluating the LSTM model performance on test data
# validation loss = 0.21674150228500366
# validation accuracy = 0.930610716342926
loaded 1stm model.evaluate(X test padded,
                            y test)
# 응 응
# predicting labels of X test data values on the basis of the
# trained model
y pred lstm = [1 \text{ if } x[0][0] > 0.5 \text{ else } 0 \text{ for } x \text{ in}
loaded lstm model.predict(X test padded)]
# printing the length of the predictions list
len(y pred lstm)
# 응 응
# printing the first 25 elements of the predictions list
y pred lstm[:25]
# 응 응
# importing mean squared error from scikit-learn library
from sklearn.metrics import mean squared error
# mean squared error (MSE)
print('MSE :', mean squared error(y test,
                                    y pred lstm))
# root mean squared error (RMSE)
# square root of the average of squared differences between predicted
# and actual value of variable
print('RMSE:', mean squared_error(y_test,
                                    y pred lstm,
                                    squared=False))
# % %
# importing mean absolute error from scikit-learn library
from sklearn.metrics import mean absolute error
# mean absolute error (MAE)
```

```
print('MAE:', mean absolute error(y test,
                                  y pred lstm))
# 응 응
# importing accuracy score from scikit-learn library
from sklearn.metrics import accuracy score
# accuracy
# ratio of the number of correct predictions to the total number of
# input samples
print('Accuracy:', accuracy score(y test,
                                  y pred lstm))
# 응 응
# importing precision recall fscore support from scikit-learn library
from sklearn.metrics import precision recall fscore support
print('\t\tPrecision \t\tRecall \t\tF-Measure \tSupport')
# computing precision, recall, f-measure and support for each class
# with average='micro'
print('average=micro
                        -', precision recall fscore support(y test,
                                                             y pred lstm,
average='micro'))
# computing precision, recall, f-measure and support for each class
# with average='macro'
print('average=macro
                       -', precision recall fscore support (y test,
                                                             y pred lstm,
average='macro'))
# computing precision, recall, f-measure and support for each class
# with average='weighted'
print('average=weighted -', precision recall fscore support(y test,
                                                             y pred lstm,
average='weighted'))
# 응 응
# importing classification report from scikit-learn library
# used to measure the quality of predictions from a
# classification algorithm
from sklearn.metrics import classification report
# report shows the main classification metrics precision, recall and
# f1-score on a per-class basis
print(classification report(y test,
                            y pred lstm))
```

```
# importing confusion matrix from scikit-learn library
from sklearn.metrics import confusion matrix
# confusion matrix is a summarized table used to assess the performance
# of a classification model
# number of correct and incorrect predictions are summarized with their
# count according to each class
print(confusion_matrix(y_test,
                       y pred lstm))
# 응 응
# importing plot confusion matrix from scikit-learn library
# plotting the confusion matrix
cm = confusion matrix(y test,
                      y pred lstm)
sns.heatmap(cm,
            annot=True,
            cbar=False,
            fmt='g')
plt.title('LSTM Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('True')
plt.savefig('plots/lstm confusion matrix.png',
            facecolor='white')
plt.show()
# 응 응
# plotting scatter plot to visualize overlapping of predicted and
# test target data points
plt.scatter(range(len(y pred lstm)),
            y pred lstm,
            color='red')
plt.scatter(range(len(y test)),
            y test,
            color='green')
plt.title('LSTM - Predicted label vs. Actual label')
plt.xlabel('SMS Messages')
plt.ylabel('Label')
plt.savefig('plots/lstm predicted vs real.png',
            facecolor='white')
plt.show()
#88 md
```

Densely Connected CNN (DenseNet) Model

응 응

A DenseNet is a type of Convolutional Neural Network (CNN) that utilises dense connections between layers, where all layers with matching feature-map sizes are connected directly with each other. With the dense

connections, higher accuracy is achieved with fewer parameters compared to a traditional CNN.

```
# % %
# Densely Connected CNN (DenseNet) architecture hyperparameters
# SpatialDropout1D is used to dropout the embedding layer which helps
# to drop entire 1D feature maps instead of individual elements
# dropout rate indicates the fraction of the units to drop for the
# linear transformation of the inputs
dropout rate = 0.2
# embedding dimension indicates the dimension of the state space used for
# reconstruction
embedding dimension = 16
# no of epochs indicates the number of complete passes through the
# training dataset
no of epochs = 30
# vocabulary size indicates the maximum number of unique words to
# tokenize and load in training and testing data
vocabulary size = 500
# 응 응
# importing Sequential class from keras
# Sequential groups a linear stack of layers into a keras Model
from tensorflow.keras.models import Sequential
# importing Embedding class from keras layers api package
# turning positive integers (indexes) into dense vectors of fixed size
# this layer can only be used as the first layer in a model
from tensorflow.keras.layers import Embedding
# importing GlobalAveragePooling1D class from keras layers api package
# used to perform global average pooling operation for temporal data
from tensorflow.keras.layers import GlobalAveragePooling1D
# importing Dropout class from keras layers api package
# used to apply Dropout to the input
# Dropout is one of the most effective and most commonly used
# regularization techniques for neural networks
# Dropout, applied to a layer, consists of randomly 'dropping out'
# (set to zero) a number of output features of the layer during training
from tensorflow.keras.layers import Dropout
# importing Dense class from keras layers api package
# Dense class is a regular densely-connected neural network layer
# Dense implements the operation:
# output = activation(dot(input, kernel) + bias)
# activation is the element-wise activation function passed as
# the activation argument
```

```
# kernel is a weights matrix created by the layer
# bias is a bias vector created by the layer
# (only applicable if use bias is True)
from tensorflow.keras.layers import Dense
# 응 응
# Densely Connected CNN (DenseNet) model architecture
densenet cnn model = Sequential()
densenet cnn model.add(Embedding(vocabulary size,
                                 embedding dimension,
                                 input length=maximum length))
densenet cnn model.add(GlobalAveragePooling1D())
# Rectified Linear Activation Function (ReLU) is a piecewise linear
# function that will output the input directly if it is positive,
# otherwise, it will output zero
densenet cnn model.add(Dense(24,
                             activation='relu'))
densenet cnn model.add(Dropout(dropout rate))
# sigmoid is a non-linear and easy to work with activation function
# that takes a value as input and outputs another value between 0 and 1
densenet cnn model.add(Dense(1,
                             activation='sigmoid'))
# 응 응
# compiling the model
# configuring the model for training
densenet cnn model.compile(loss='binary crossentropy',
                           optimizer='adam',
                           metrics=['accuracy'])
# 응 응
# printing a string summary of the network
densenet cnn model.summary()
# 응 응
# monitoring the validation loss and if the validation loss is not
# improved after three epochs, then the model training is stopped
# it helps to avoid overfitting problem and indicates when to stop
# training before the deep learning model begins overfitting
early stopping = EarlyStopping(monitor='val_loss',
                               patience=3)
# % %
```

```
# training the DenseNet CNN model
history = densenet cnn model.fit(X train padded,
                                  y train,
                                  epochs=no of epochs,
                                  validation data=(X test padded, y test),
                                  callbacks=[early stopping],
                                  verbose=2)
# % %
# visualizing the history results by reading as a dataframe
densenet cnn metrics = pd.DataFrame(history.history)
densenet cnn metrics
# 응 응
# renaming the column names of the dataframe
densenet cnn metrics.rename(columns={'loss': 'Training Loss',
                                      'accuracy': 'Training Accuracy',
                                      'val loss': 'Validation Loss',
                                      'val accuracy':
'Validation Accuracy' },
                             inplace=True)
densenet cnn metrics
# 응 응
# plotting the training and validation loss by number of epochs for
# the DenseNet CNN model
densenet cnn metrics[['Training_Loss', 'Validation_Loss']].plot()
plt.title('DenseNet CNN Model - Training and Validation Loss vs. Epochs')
plt.xlabel('Number of Epochs')
plt.ylabel('Loss')
plt.legend(['Training_Loss', 'Validation_Loss'])
plt.savefig('plots/densenet cnn loss vs epochs.png',
            facecolor='white')
plt.show()
# 응 응
# plotting the training and validation accuracy by number of epochs
# for the DenseNet CNN model
densenet cnn metrics[['Training Accuracy', 'Validation Accuracy']].plot()
plt.title('DenseNet CNN Model - Training and Validation Accuracy vs.
Epochs')
plt.xlabel('Number of Epochs')
plt.ylabel('Accuracy')
plt.legend(['Training Accuracy', 'Validation Accuracy'])
plt.savefig('plots/densenet cnn accuracy vs epochs.png',
            facecolor='white')
plt.show()
# % %
```

```
# saving the trained DenseNet CNN model as an h5 file
# h5 is a file format to store structured data
# keras saves deep learning models in this format as it can easily store
# the weights and model configuration in a single file
densenet cnn path = 'models/densenet cnn model.h5'
densenet cnn model.save (densenet cnn path)
densenet cnn model
# 응 응
# loading the saved DenseNet CNN model
loaded densenet cnn model = load model(densenet cnn path)
loaded densenet cnn model
#88 md
# Densely Connected CNN (DenseNet) Model Evaluation
# evaluating the DenseNet CNN model performance on test data
# validation loss = 0.11119994521141052
# validation accuracy = 0.9732824563980103
loaded densenet cnn model.evaluate(X test padded,
                                    y test)
# % %
# predicting labels of X test data values on the basis of the
# trained model
y pred densenet cnn = [1 \text{ if } x[0] > 0.5 \text{ else } 0 \text{ for } x \text{ in }
loaded densenet cnn model.predict(X test padded)]
# printing the length of the predictions list
len(y pred densenet cnn)
# 응 응
# printing the first 25 elements of the predictions list
y pred densenet cnn[:25]
# 응 응
# mean squared error (MSE)
print('MSE :', mean squared_error(y_test,
                                   y pred densenet cnn))
# root mean squared error (RMSE)
# square root of the average of squared differences between predicted
# and actual value of variable
print('RMSE:', mean squared error(y test,
                                   y pred densenet cnn,
                                   squared=False))
```

```
# 응 응
# mean absolute error (MAE)
print('MAE:', mean absolute error(y test,
                                  y pred densenet cnn))
# 응 응
# accuracy
# ratio of the number of correct predictions to the total number of
# input samples
print('Accuracy:', accuracy score(y test,
                                  y pred densenet cnn))
# 응 응
print('\t\tPrecision \t\tRecall \t\tF-Measure \tSupport')
# computing precision, recall, f-measure and support for each class
# with average='micro'
                     -', precision recall fscore support (y test,
print('average=micro
y pred densenet cnn,
average='micro'))
# computing precision, recall, f-measure and support for each class
# with average='macro'
print('average=macro
                       -', precision recall fscore support (y test,
y pred densenet cnn,
average='macro'))
# computing precision, recall, f-measure and support for each class
# with average='weighted'
print('average=weighted -', precision_recall_fscore_support(y_test,
y pred densenet cnn,
average='weighted'))
# 응 응
# report shows the main classification metrics precision, recall and
# f1-score on a per-class basis
print(classification report(y test,
                            y_pred_densenet_cnn))
# 응 응
# confusion matrix is a summarized table used to assess the performance
# of a classification model
# number of correct and incorrect predictions are summarized with their
```

```
# count according to each class
print(confusion matrix(y_test,
                       y pred densenet cnn))
# % %
# plotting the confusion matrix
cm = confusion matrix(y_test,
                      y pred densenet cnn)
sns.heatmap(cm,
            annot=True,
            cbar=False,
            fmt='g')
plt.title('DenseNet CNN Confusion Matrix')
plt.xlabel('Predicted')
plt.ylabel('True')
plt.savefig('plots/densenet cnn confusion matrix.png',
            facecolor='white')
plt.show()
# 응 응
# plotting scatter plot to visualize overlapping of predicted and
# test target data points
plt.scatter(range(len(y pred densenet cnn)),
            y pred densenet cnn,
            color='red')
plt.scatter(range(len(y test)),
            y test,
            color='green')
plt.title('DenseNet CNN - Predicted label vs. Actual label')
plt.xlabel('SMS Messages')
plt.ylabel('Label')
plt.savefig('plots/densenet cnn predicted vs real.png',
            facecolor='white')
plt.show()
#%% md
# Making Predictions with Real-World Examples
# 응 응
# defining pre-processing hyperparameters
# maximum length indicates the maximum number of words considered
# in a text
maximum length = 50
# truncating type indicates removal of values from sequences larger
# than maxlen, either at the beginning ('pre') or at the end ('post')
# of the sequences
```

```
truncating type = 'post'
# padding_type indicates pad either before ('pre') or after ('post')
# each sequence
padding type = 'post'
# % %
# defining a function to preprocess the sms message text to feed to the
# trained deep learning models
# the input text is transformed to a sequence of integers
# then the sequence is padded to the same length
# padding='post' to pad after each sequence
# the function returns the padded sequence
def preprocess text(sms messages):
    sequence = tokenizer.texts to sequences(sms messages)
    padded sequence = pad sequences(sequence,
                                    maxlen=maximum length,
                                    padding=padding type,
                                    truncating=truncating type)
    return padded sequence
# 응 응
# defining a set of real-world samples for ham and spam sms messages
sms messages = [
    'IMPORTANT - You could be entitled up to £3,160 in compensation from
mis-sold PPI on a credit card or loan. Please reply PPI for info or STOP
to opt out.',
    'Hello, Janith! Did you go to the school yesterday? If you did, can
you please send me the notes of all the subjects?',
    'Congratulations ur awarded 500 of CD vouchers or 125 gift guaranteed
& Free entry 2 100 wkly draw txt MUSIC to 87066.',
    'A loan for £950 is approved for you if you receive this SMS. 1 min
verification & cash in 1 hr at www.abc.co.uk to opt out reply stop',
    'If he started searching, he will get job in few days. He has great
potential and talent.',
    'One chance ONLY! Had your mobile 11mths+? You are entitled to update
to the latest colour camera mobile for FREE! Call The Mobile Update Co
FREE on 08002986906.',
    'Valentines Day Special! Win over 1000 USD in cash in our quiz and
take your partner on the trip of a lifetime! Send GO to 83600 now. 150
p/msg rcvd.',
    'Now I am better. Made up for Friday and stuffed myself like a pig
yesterday. Now I feel bad.',
    'I got another job! The one at the hospital, doing data analysis or
something, starts on Monday! Not sure when my thesis will finish.'
]
# 응 응
# invoking the preprocess text function to preprocess and get the
# padded sequences of the set of real-world samples for ham and
# spam sms messages
```

```
padded sequences = preprocess text(sms messages)
padded sequences
# 응 응
# making prediction for the given set of real-world sms messages
# using the trained LSTM model
print('LSTM Model Predictions',
      end='\n\n')
lstm prediction = []
for index, sms message in enumerate(sms messages):
    prediction = loaded lstm model.predict(padded sequences)[index][0][0]
    lstm prediction.append(prediction)
    if prediction > 0.5:
        print('SPAM -', prediction, '-', sms_message)
    else:
       print('HAM -', prediction, '-', sms message)
# 응 응
# making prediction for the given set of real-world sms messages
# using the trained DenseNet CNN model
print('DenseNet CNN Model Predictions',
      end='\n\n'
densenet cnn prediction = []
for index, sms message in enumerate(sms messages):
    prediction =
loaded densenet cnn model.predict(padded sequences)[index][0]
    densenet cnn prediction.append(prediction)
    if prediction > 0.5:
        print('SPAM -', prediction, '-', sms_message)
    else:
        print('HAM -', prediction, '-', sms message)
#88 md
# Comparison of Deep Learning Models
# 응 응
# plotting line graphs of the predicted values for LSTM and
# DenseNet CNN deep learning models to compare the separation
# of classes by each model
plt.plot(list(range(len(sms messages))),
         1stm prediction,
         label='LSTM',
         color='blue',
         marker='o')
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plt.plot(list(range(len(sms messages))),
         densenet cnn prediction,
         label='CNN',
         color='red',
         marker='o')
plt.plot(list(range(len(sms messages))),
         [0.5 for x in range(len(sms messages))],
         color='black',
         linestyle='dashed')
plt.title('Comparison of Predicted Values by LSTM and DenseNet Models')
plt.xlabel('SMS Message ID')
plt.ylabel('Predicted Value')
plt.legend(loc='upper right',
           bbox to anchor=(1, 1)
plt.savefig('plots/prediction values comparison.png',
            facecolor='white')
plt.show()
# % %
# importing recall score from scikit-learn library
from sklearn.metrics import recall score
# importing precision score from scikit-learn library
from sklearn.metrics import precision score
# importing f1 score from scikit-learn library
from sklearn.metrics import f1 score
accuracy = {}
recall = {}
precision = {}
f1 = {}
rmse = {}
mae = {}
y pred dict = {
    'LSTM': y pred lstm,
    'CNN': y pred densenet cnn
}
for y pred in y pred dict:
    accuracy[y_pred] = accuracy_score(y_test,
                                       y pred dict[y pred])
    recall[y pred] = recall score(y test,
                                   y pred dict[y pred],
                                   average='weighted')
    precision[y pred] = precision score(y test,
                                         y pred dict[y pred],
                                         average='weighted')
    f1[y pred] = f1 score(y test,
                           y pred dict[y pred],
                           average='weighted')
    rmse[y pred] = mean squared error(y test,
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y pred dict[y pred],
                                       squared=False)
    mae[y pred] = mean absolute error(y test,
                                       y pred dict[y pred])
# ratio of the number of correct predictions to the total number
# of input samples
print('Accuracy:', accuracy)
# recall is the ratio tp / (tp + fn)
# tp is the number of true positives
# fn the number of false negatives
print('Recall:', recall)
# precision is the ratio tp / (tp + fp)
# tp is the number of true positives
# fp the number of false positives
print('Precision:', precision)
# F1 score is also known as balanced F-score or F-measure
# F1 score can be interpreted as a weighted average of the
# precision and recall
\# F1 = 2 * (precision * recall) / (precision + recall)
print('F1 Score:', f1)
# root mean squared error (RMSE)
# square root of the average of squared differences between predicted
# and actual value of variable
print('RMSE:', rmse)
# mean absolute error (MAE)
print('MAE:', mae)
# 응 응
# sorting the accuracy scores of two deep learning models in
# descending order
sorted(accuracy.items(),
       key=lambda kv: kv[1],
       reverse=True)
# 응 응
# plotting the accuracy comparison bar chart for the two
# deep learning models
fig, ax = plt.subplots(figsize=(5, 5))
plt.bar(y pred dict.keys(),
        accuracy.values(),
        color='rg')
plt.title('Accuracy Comparison')
plt.xlabel('Algorithm')
plt.ylabel('Accuracy')
plt.savefig('plots/accuracy comparison.png',
            facecolor='white')
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plt.show()
# 응 응
# defining a function to plot a bar chart with multiple bars
def bar plot(ax,
             data,
             colors=None,
             total width=0.8,
             single width=1,
             legend=True):
    if colors is None:
        colors = plt.rcParams['axes.prop cycle'].by key()['color']
    n bars = len(data)
    bar width = total width / n bars
    bars = []
    for i, (name, values) in enumerate(data.items()):
        x_offset = (i - n_bars / 2) * bar_width + bar_width / 2
        for x, y in enumerate(values):
            bar = ax.bar(x + x offset,
                         width=bar width * single width,
                          color=colors[i % len(colors)])
        bars.append(bar[0])
    if legend:
        ax.legend(bars,
                  data.keys())
# 응 응
# plotting the algorithm comparison chart for all evaluation metrics
data = \{\}
for key in y pred dict.keys():
    data[key] = [accuracy[key],
                 recall[key],
                 precision[key],
                 f1[key],
                 rmse[key],
                 mae[key]]
fig, ax = plt.subplots(figsize=(7, 5))
bar plot(ax,
         data,
         total width=0.9,
         single width=0.9)
plt.title('Algorithm Comparison')
plt.xlabel('Metrics')
plt.ylabel('Scores')
plt.xticks(range(len(data[key])),
           ['Accuracy', 'Recall', 'Precision', 'F1-Score', 'RMSE', 'MAE'])
```