

```
In [1]: import pandas as pd      # Loading data
df=pd.read_csv(r'https://raw.githubusercontent.com/iiscleap/Coswara-Data/master/combined_data.csv')
```

```
In [2]: df
```

```
Out[2]:
```

		id	a	covid_status	ep	g	I_c	I_l	I_s	rU	smoker	...	diarrhoea	pneumoni
0	iV3Db6t1T8b7c5HQY2TwxlhjbzD3	28		healthy	y	male	India	Anantapur	Andhra Pradesh	n	NaN	...	NaN	NaN
1	AxuYWBNOjFVLINCBqIW5aZmGCdu1	25		healthy	y	male	India	BENGALURU URBAN	Karnataka	n	True	...	NaN	NaN
2	C5elsssb9GSkaAgIfsHMH6fSh1	28		healthy	y	female	United States	Pittsburgh	Pennsylvania	n	NaN	...	NaN	NaN
3	YjbEAECMBIaZKyfQvWy5DDImUb2	26		healthy	y	male	India	Bangalore	Karnataka	n	NaN	...	NaN	NaN
4	aGOvk4jioVqlzCs1jHnzlw2UEy2	32		healthy	y	male	India	Nalanda	Bihar	n	NaN	...	NaN	NaN
...
1890	aH6o5IYN6vh40RCFhfYBYjkQVH13	33		healthy	y	male	India	24pgs	West Bengal	n	True	...	NaN	NaN
1891	hRMSJF9xD6h9CmQJzB3phXKNnTI2	42		recovered_full	y	male	India	Bangalore	Karnataka	n	NaN	...	NaN	NaN
1892	ZNTuppbDaCNybeDYGmyJeYr93s1	27		healthy	y	female	India	NaN	Karnataka	n	NaN	...	NaN	NaN
1893	J7R2UklD7jU0aOSw5amaS8zJVIm1	37		healthy	y	male	India	NaN	Karnataka	n	NaN	...	NaN	NaN
1894	8MdsvTBMWAUlnavE1pdmFQAZPF13	38		recovered_full	y	female	India	Bengaluru	Karnataka	n	NaN	...	NaN	NaN

1895 rows × 35 columns



```
In [3]: df.isnull().sum()      #checking for null
```

```
Out[3]: id                0
a                0
covid_status      0
ep                0
g                0
I_c              0
I_l             229
I_s              0
rU              680
smoker           1660
cold            1753
ht              1770
diabetes         1784
fever           1768
asthma          1822
um              1038
cough           1679
ihd             1884
bd              1825
st              1810
ftg             1803
mp              1815
loss_of_smell    1837
cld             1887
test_status      1413
diarrhoea        1886
pneumonia        1885
ctScan           1783
testType         1808
test_date        1808
vacc             1783
ctDate           1884
ctScore          1884
others_resp      1894
others_preexist  1887
dtype: int64
```

```
In [5]: df["rU"]=df["rU"].fillna("n")      #data - preprocessing
df["um"]=df["um"].fillna("n")
df=df.fillna(False)
```

In [6]:

df

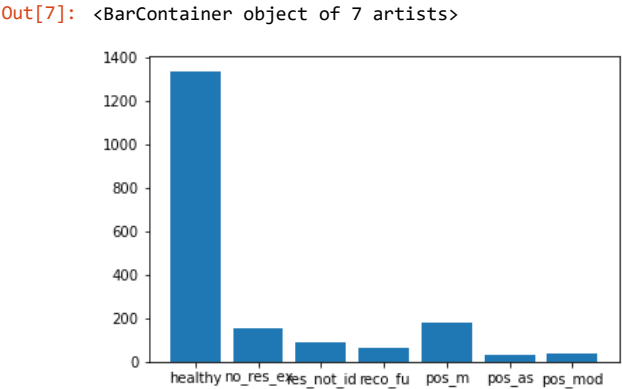
Out[6]:

		id	a	covid_status	ep	g	I_c	I_l	I_s	rU	smoker	...	diarrhoea	pneumoni
0	iV3Db6t1T8b7c5HQY2TwxIhjbzD3	28		healthy	y	male	India	Anantapur	Andhra Pradesh	n	False	...	False	Fals
1	AxuYWBNOjFVLINCBqIW5aZmGCdu1	25		healthy	y	male	India	BENGALURU URBAN	Karnataka	n	True	...	False	Fals
2	C5elsssb9GSkaAgIfsHMHeR6fSh1	28		healthy	y	female	United States	Pittsburgh	Pennsylvania	n	False	...	False	Fals
3	YjbEAECMBlaZKyfqOvWy5DDImUb2	26		healthy	y	male	India	Bangalore	Karnataka	n	False	...	False	Fals
4	aGOvk4ji0cVqlzCs1jHnzlw2UEy2	32		healthy	y	male	India	Nalanda	Bihar	n	False	...	False	Fals
...
1890	aH6o5IYN6vh40RCFhyBYjkQVH13	33		healthy	y	male	India	24pgs	West Bengal	n	True	...	False	Fals
1891	hRMSJF9xD6h9CmQJzB3phXKNnTI2	42		recovered_full	y	male	India	Bangalore	Karnataka	n	False	...	False	Fals
1892	ZNTuppbDaCNybeDYGmyJeYr93s1	27		healthy	y	female	India	False	Karnataka	n	False	...	False	Fals
1893	J7R2UklD7jU0aOSw5amaS8zJVIm1	37		healthy	y	male	India	False	Karnataka	n	False	...	False	Fals
1894	8MdsvTBMWAUInavE1pdmFQAZPF13	38		recovered_full	y	female	India	Bengaluru	Karnataka	n	False	...	False	Fals

1895 rows × 35 columns

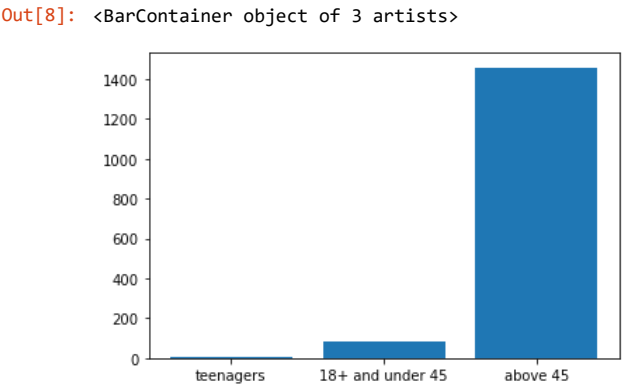
In [7]:

identified]],len(a[a=="recovered_full"]),len(a[a=="positive_mild"]),len(a[a=="positive_asymp"]),len(a[a=="positive_mo



In [8]:

d=df["a"]
t1=0
t2=0
t3=0
for i in d:
 if (i<=11 and i<=18):
 t1+=1
 elif (i<=19 and i<=45):
 t2+=1
 elif (i<=46 and i<=100):
 t3+=1
plt.bar(["teenagers", "18+ and under 45", "above 45"], [t1,t2,t3])



```
In [9]: from sklearn import preprocessing
l=preprocessing.LabelEncoder()
#a=df["covid_status"]
df["covid_status"]=l.fit_transform(df["covid_status"])

In [10]: df["rU"]=l.fit_transform(df["rU"])
df["um"]=l.fit_transform(df["um"])
df["g"]=l.fit_transform(df["g"])

In [11]: df["smoker"]=df["smoker"]*1

In [12]: df["cold"]=df["cold"].astype(int)
df
```

Out[12]:

		id	a	covid_status	ep	g	l_c	l_l	l_s	rU	smoker	...	diarrhoea	pneumonia	ct
0	iV3Db6t1T8b7c5HQY2TwxlhjbzD3	28		0	y	1	India	Anantapur	Andhra Pradesh	0	0	...	False	False	
1	AxuYWBNOjFVLINCBqIW5aZmGCdu1	25		0	y	1	India	BENGALURU URBAN	Karnataka	0	True	...	False	False	
2	C5elsssb9GSkaAgfshMHeR6fSh1	28		0	y	0	United States	Pittsburgh	Pennsylvania	0	0	...	False	False	
3	YjbEAECMBIaZKyfqOvWy5DDImUb2	26		0	y	1	India	Bangalore	Karnataka	0	0	...	False	False	
4	aGOvk4ji0cVqlzCs1jHnzlw2UEy2	32		0	y	1	India	Nalanda	Bihar	0	0	...	False	False	
...	
1890	aH6o5IYN6vh40RCFhfyBYjkQVH13	33		0	y	1	India	24pgs	West Bengal	0	True	...	False	False	
1891	hRMSJF9xD6h9CmQJzB3phXKNnTI2	42		5	y	1	India	Bangalore	Karnataka	0	0	...	False	False	
1892	ZNTuppbDaCNybeDYGmyJeYr93s1	27		0	y	0	India	False	Karnataka	0	0	...	False	False	
1893	J7R2Ukld7jU0aOSw5amaS8zJVIm1	37		0	y	1	India	False	Karnataka	0	0	...	False	False	
1894	8MdsVTBMWAUlnavE1pdmFQAZPF13	38		5	y	0	India	Bengaluru	Karnataka	0	0	...	False	False	

1895 rows × 35 columns

```
In [13]: #df["cold"]=df["cold"].astype(int)
df["ht"]=df["ht"].astype(int)
df["diabetes"]=df["diabetes"].astype(int)
df["fever"]=df["fever"].astype(int)
df["asthma"]=df["asthma"].astype(int)
df["cough"]=df["cough"].astype(int)
df["ihd"]=df["ihd"].astype(int)
df["bd"]=df["bd"].astype(int)
df["st"]=df["st"].astype(int)
df["ftg"]=df["ftg"].astype(int)
df["mp"]=l.fit_transform(df["mp"])
df["loss_of_smell"]=df["loss_of_smell"].astype(int)
df["cld"]=df["cld"].astype(int)
#df["test_status"]=df["test_status"].fillna("False")
df["diarrhoea"]=df["diarrhoea"].astype(int)
df["pneumonia"]=df["pneumonia"].astype(int)

In [14]: df["smoker"]=df["smoker"].replace("True",1)
#df["smoker"]=df["smoker"].astype(int)
```

```
In [15]: df
```

Out[15]:

		id	a	covid_status	ep	g	l_c	l_l	l_s	rU	smoker	...	diarrhoea	pneumonia	ct
0	iV3Db6t1T8b7c5HqY2TwXlhjbzD3	28		0	y	1	India	Anantapur	Andhra Pradesh	0	0	...	0	0	
1	AxuYWBN0jFVLINCBqIW5aZmGCdu1	25		0	y	1	India	BENGALURU URBAN	Karnataka	0	1	...	0	0	
2	C5elsssb9GSkaAgIfsHMHeR6fSh1	28		0	y	0	United States	Pittsburgh	Pennsylvania	0	0	...	0	0	
3	YjbEAECMBlaZKyfqOvWy5DDImUb2	26		0	y	1	India	Bangalore	Karnataka	0	0	...	0	0	
4	aGOvk4ji0cVqlzCs1jHnzlw2UEy2	32		0	y	1	India	Nalanda	Bihar	0	0	...	0	0	
...	
1890	aH6o5iYN6vh40RCFhfYBYjkQVH13	33		0	y	1	India	24pgs	West Bengal	0	1	...	0	0	
1891	hRMSJF9xD6h9CmQJzB3phXKNnTI2	42		5	y	1	India	Bangalore	Karnataka	0	0	...	0	0	
1892	ZNTuppbDaCNybeDYGmyJeYr93s1	27		0	y	0	India	False	Karnataka	0	0	...	0	0	
1893	J7R2UklD7jU0aOSw5amaS8zJVIm1	37		0	y	1	India	False	Karnataka	0	0	...	0	0	
1894	8MdsvTBMWAUInavE1pdmFQAZPF13	38		5	y	0	India	Bengaluru	Karnataka	0	0	...	0	0	

1895 rows × 35 columns



```
In [16]: df.isnull().sum()
```

Out[16]:

id	0
a	0
covid_status	0
ep	0
g	0
l_c	0
l_l	0
l_s	0
rU	0
smoker	0
cold	0
ht	0
diabetes	0
fever	0
asthma	0
um	0
cough	0
ihd	0
bd	0
st	0
ftg	0
mp	0
loss_of_smell	0
cld	0
test_status	0
diarrhoea	0
pneumonia	0
ctScan	0
testType	0
test_date	0
vacc	0
ctDate	0
ctScore	0
others_resp	0
others_preexist	0
dtype: int64	

```
In [17]: df
```

Out[17]:

		id	a	covid_status	ep	g	L_c	L_l	L_s	rU	smoker	...	diarrhoea	pneumonia	ct
0	iV3Db6t1T8b7c5HQY2TwxlhjbzD3	28		0	y	1	India	Anantapur	Andhra Pradesh	0	0	...	0	0	
1	AxuYWBNOjFVLINCBqIW5aZmGCdu1	25		0	y	1	India	BENGALURU URBAN	Karnataka	0	1	...	0	0	
2	C5elsssb9GSkaAgfshMHeR6fSh1	28		0	y	0	United States	Pittsburgh	Pennsylvania	0	0	...	0	0	
3	YjbEAECMBlaZKyfqOvWy5DDImUb2	26		0	y	1	India	Bangalore	Karnataka	0	0	...	0	0	
4	aGOvk4ji0cVqlzCs1jHnzlw2UEy2	32		0	y	1	India	Nalanda	Bihar	0	0	...	0	0	
...	
1890	aH6o5iYN6vh40RCFhyBYjkQVH13	33		0	y	1	India	24pgs	West Bengal	0	1	...	0	0	
1891	hRMSJF9xD6h9CmQJzB3phXKNnTI2	42		5	y	1	India	Bangalore	Karnataka	0	0	...	0	0	
1892	ZNTuppbDaCNybeDYGmyjJeYr93s1	27		0	y	0	India	False	Karnataka	0	0	...	0	0	
1893	J7R2Ukl7jU0aOSw5amaS8zJVIm1	37		0	y	1	India	False	Karnataka	0	0	...	0	0	
1894	8MdsvtBMWauInavE1pdmFQAZPF13	38		5	y	0	India	Bengaluru	Karnataka	0	0	...	0	0	

1895 rows × 35 columns



```
In [18]: df["covid_status"]
```

Out[18]:

0	0
1	0
2	0
3	0
4	0
...	...
1890	0
1891	5
1892	0
1893	0
1894	5

Name: covid_status, Length: 1895, dtype: int32

In [19]:

df.corr()

#finding correlation for target column to extraxt the features

Out[19]:

	a	covid_status	g	rU	cold	ht	diabetes	fever	asthma	um	...	bd
a	1.000000	0.070818	-0.004444	0.007648	-0.004834	0.329970	0.323958	0.070876	-0.002322	0.075909	...	0.141344
covid_status	0.070818	1.000000	-0.074612	0.037475	0.234706	0.106719	0.089614	0.310393	0.366153	0.247199	...	0.260441
g	-0.004444	-0.074612	1.000000	0.024445	-0.004895	0.011394	0.048830	0.004336	-0.003138	-0.091514	...	0.016974
rU	0.007648	0.037475	0.024445	1.000000	0.059672	-0.019958	0.011304	0.042883	-0.017802	0.098737	...	0.033719
cold	-0.004834	0.234706	-0.004895	0.059672	1.000000	0.021263	0.031429	0.364572	-0.025726	0.123828	...	0.156791
ht	0.329970	0.106719	0.011394	-0.019958	0.021263	1.000000	0.268686	0.056307	0.035180	0.010695	...	0.094485
diabetes	0.323958	0.089614	0.048830	0.011304	0.031429	0.268686	1.000000	0.067949	0.043484	0.029102	...	0.141777
fever	0.070876	0.310393	0.004336	0.042883	0.364572	0.056307	0.067949	1.000000	-0.009785	0.191502	...	0.361498
asthma	-0.002322	0.366153	-0.003138	-0.017802	-0.025726	0.035180	0.043484	-0.009785	1.000000	-0.016493	...	0.033486
um	0.075909	0.247199	-0.091514	0.098737	0.123828	0.010695	0.029102	0.191502	-0.016493	1.000000	...	0.159796
cough	0.101326	0.326308	-0.023673	0.098252	0.396169	0.038480	0.037815	0.448407	-0.020025	0.201046	...	0.273108
ihd	0.131344	0.058897	0.014676	0.028663	0.031020	0.119621	0.188007	0.062860	0.056894	-0.000380	...	0.169181
bd	0.141344	0.260441	0.016974	0.033719	0.156791	0.094485	0.141777	0.361498	0.033486	0.159796	...	1.000000
st	-0.011312	0.194681	-0.001744	0.054822	0.228823	-0.037047	0.011086	0.247791	-0.003635	0.089384	...	0.160314
ftg	0.093844	0.271610	-0.041149	0.064611	0.271435	0.078456	0.027301	0.391139	-0.019699	0.165408	...	0.294222
mp	0.035811	0.234686	-0.057022	0.027134	0.259210	0.018216	0.037035	0.332047	-0.014751	0.168709	...	0.237159
loss_of_smell	-0.001018	0.209661	-0.024261	0.080044	0.205421	0.002149	0.033954	0.283174	-0.019648	0.059721	...	0.176356
cld	0.072513	0.100924	0.057541	-0.011151	0.012382	0.015486	0.053077	0.015098	0.071548	-0.020769	...	0.073551
diarrhoea	0.019490	0.055659	0.024279	-0.011831	0.096951	-0.018358	0.048140	0.165658	0.065938	0.057500	...	0.108555
pneumonia	0.071803	0.120242	-0.005415	0.031301	0.034599	0.039331	0.074882	0.155247	0.098958	0.077398	...	0.178818
others_resp	0.039826	0.027341	0.013831	-0.003935	-0.006540	-0.006106	-0.005732	0.085733	-0.004599	-0.007329	...	0.117326
others_preexist	0.105169	0.100924	-0.052545	-0.011151	0.074208	0.146645	0.122396	0.047647	0.029257	0.091680	...	0.030400

22 rows × 22 columns

In [20]:

test=df["covid_status"]#train and test data

train=df[["fever","asthma","cough","ftg","bd","cold","um","mp","loss_of_smell","pneumonia"]]

In [21]:

from sklearn.model_selection import train_test_split#train and test data split

X_train, X_test, y_train, y_test = train_test_split(train, test, test_size = .3, random_state = 0)

In [22]:

import tensorflow#importing libraries

from tensorflow import keras

from keras.models import Sequential

from keras.layers import Dense

In [23]:

df

Out[23]:

	id	a	covid_status	ep	g	L_c	L_I	L_s	rU	smoker	...	diarrhoea	pneumonia	ct
0	iV3Db6t1T8b7c5HQY2TwxlhjbzD3	28		0	y	1	India	Anantapur	Andhra Pradesh	0	0	...	0	0
1	AxuYWBNOjFVLINCBqIW5aZmGCdu1	25		0	y	1	India	BENGALURU URBAN	Karnataka	0	1	...	0	0
2	C5elsssb9GSkaAgIfsHMHeR6fSh1	28		0	y	0	United States	Pittsburgh	Pennsylvania	0	0	...	0	0
3	YjbEAECMBlaZKyfQvWy5DDImUb2	26		0	y	1	India	Bangalore	Karnataka	0	0	...	0	0
4	aGOvk4jii0cVqlzCs1jHnzlw2UEy2	32		0	y	1	India	Nalanda	Bihar	0	0	...	0	0
...
1890	aH6o5IYN6vh40RCFhfYBYjkQVH13	33		0	y	1	India	24pgs	West Bengal	0	1	...	0	0
1891	hRMSJF9xD6h9CmQJzB3phXKNnTI2	42		5	y	1	India	Bangalore	Karnataka	0	0	...	0	0
1892	ZNTuppbDaCNybeDYGmyJeYr93s1	27		0	y	0	India	False	Karnataka	0	0	...	0	0
1893	J7R2UklD7jU0aOSw5amaS8zJVIm1	37		0	y	1	India	False	Karnataka	0	0	...	0	0
1894	8MdsVBMWAUInavE1pdmFQAZPF13	38		5	y	0	India	Bengaluru	Karnataka	0	0	...	0	0

1895 rows × 35 columns

```
In [24]: from sklearn.preprocessing import StandardScaler           #standardizing data
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)
```

```
In [44]: from sklearn.neural_network import MLPClassifier           #training MLP classifier
clf = MLPClassifier(solver='sgd', alpha=1e-5,hidden_layer_sizes=(100, 100,10), random_state=1)
clf.fit(X_train, y_train)
```

C:\Users\sahit\anaconda3\lib\site-packages\sklearn\neural_network_multilayer_perceptron.py:582: ConvergenceWarning: Stochastic Optimizer: Maximum iterations (200) reached and the optimization hasn't converged yet.
warnings.warn(

```
Out[44]: MLPClassifier(alpha=1e-05, hidden_layer_sizes=(100, 100, 10), random_state=1,
solver='sgd')
```

```
In [45]: a=clf.predict(X_test)
```

```
In [46]: from sklearn.metrics import accuracy_score,confusion_matrix           #accuracy of MLP classifier
print(accuracy_score(y_test,a))
```

```
0.7557117750439367
```

```
In [25]: from keras.models import Sequential           #importing libraries for cnn and lstm
import keras
from keras.models import Sequential
from keras.layers import Dense, Flatten, Conv1D, Dropout
from keras import utils
from keras.layers.convolutional import MaxPooling1D
from keras.utils import np_utils

from keras.utils.np_utils import to_categorical
```

```
In [26]: X_train.shape
```

```
Out[26]: (1326, 10)
```

```
In [27]: X_test.shape
```

```
Out[27]: (569, 10)
```

```
In [28]: _train=y_train-1           #reshapes of X_train and y_train, y_test and x_
_test=y_test-1
_train=keras.utils.np_utils.to_categorical(y_train,7)
_test=keras.utils.np_utils.to_categorical(y_test,7)
_train=X_train.reshape(1326,10,1)
_test=X_test.reshape(569,10,1)
```

```
In [29]: model = Sequential()                                     #cnn model
model.add(Conv1D(64,3,input_shape=(10,1), activation='relu'))
#model.add(Dense(128, activation='relu'))
model.add(Dropout(0.1))
model.add(MaxPooling1D(pool_size=1))
model.add(Flatten())
model.add(Dense(64, activation='relu'))
model.add(Dense(7, activation='softmax'))

model.compile(loss=keras.losses.categorical_crossentropy,
              optimizer=keras.optimizers.Adam(),
              metrics=['accuracy'])
model.summary()
```

Model: "sequential"

Layer (type)	Output Shape	Param #
conv1d (Conv1D)	(None, 8, 64)	256
dropout (Dropout)	(None, 8, 64)	0
max_pooling1d (MaxPooling1D)	(None, 8, 64)	0
flatten (Flatten)	(None, 512)	0
dense (Dense)	(None, 64)	32832
dense_1 (Dense)	(None, 7)	455
Total params: 33,543		
Trainable params: 33,543		
Non-trainable params: 0		

```
In [30]: batch_size = 240                                     #training and evaluating cnn model
epochs = 200
history= model.fit(X_train, y_train,
                  batch_size=batch_size,
                  epochs=epochs,
                  verbose=1,validation_split=0.3,shuffle=False)
[loss,score]=model.evaluate(X_test,y_test,verbose=0)
```

```
4/4 [=====] - 0s 12ms/step - loss: 0.7094 - accuracy: 0.7993 - val_loss: 0.8318 - val_acc
uracy: 0.7638
Epoch 35/200
4/4 [=====] - 0s 12ms/step - loss: 0.7093 - accuracy: 0.8017 - val_loss: 0.8335 - val_acc
uracy: 0.7638
Epoch 36/200
4/4 [=====] - 0s 13ms/step - loss: 0.7068 - accuracy: 0.7987 - val_loss: 0.8338 - val_acc
uracy: 0.7613
Epoch 37/200
4/4 [=====] - 0s 12ms/step - loss: 0.7096 - accuracy: 0.8004 - val_loss: 0.8326 - val_acc
uracy: 0.7663
Epoch 38/200
4/4 [=====] - 0s 12ms/step - loss: 0.7058 - accuracy: 0.8027 - val_loss: 0.8339 - val_acc
uracy: 0.7663
Epoch 39/200
4/4 [=====] - 0s 12ms/step - loss: 0.7019 - accuracy: 0.8036 - val_loss: 0.8383 - val_acc
uracy: 0.7613
Epoch 40/200
4/4 [=====] - 0s 12ms/step - loss: 0.7029 - accuracy: 0.8050 - val_loss: 0.8417 - val_acc
uracy: 0.7588
```

```
In [32]: print(score*100)                                     #accuracy of cnn model
```

75.92267394065857


```
In [33]: from keras.layers import LSTM           #defining LSTM, training and evaluationg lstm model
model1 = Sequential()
model1.add(LSTM(64, input_shape=(10,1)))
model1.add(Dropout(0.1))
model1.add(Dense(7))
model1.compile(loss=keras.losses.categorical_crossentropy, optimizer='adam',metrics=['accuracy'])
lstm=model1.fit(X_train, y_train, epochs=200, batch_size=240, verbose=1,validation_split=0.3,shuffle=False)
l1,acc1=model1.evaluate(X_test,y_test,verbose=0)
```

```
Epoch 1/200
4/4 [=====] - 5s 185ms/step - loss: 4.3221 - accuracy: 0.1304 - val_loss: 10.7632 - val_a
ccuracy: 0.6256
Epoch 2/200
4/4 [=====] - 0s 19ms/step - loss: 10.9397 - accuracy: 0.5949 - val_loss: 10.8766 - val_a
ccuracy: 0.6256
Epoch 3/200
4/4 [=====] - 0s 19ms/step - loss: 11.0258 - accuracy: 0.6259 - val_loss: 10.8325 - val_a
ccuracy: 0.6256
Epoch 4/200
4/4 [=====] - 0s 18ms/step - loss: 11.0166 - accuracy: 0.6271 - val_loss: 10.8259 - val_a
ccuracy: 0.6256
Epoch 5/200
4/4 [=====] - 0s 22ms/step - loss: 10.9539 - accuracy: 0.6267 - val_loss: 10.6906 - val_a
ccuracy: 0.6256
Epoch 6/200
4/4 [=====] - 0s 21ms/step - loss: 10.9548 - accuracy: 0.6209 - val_loss: 10.6733 - val_a
ccuracy: 0.6256
Epoch 7/200
4/4 [=====] - 0s 21ms/step - loss: 10.9548 - accuracy: 0.6209 - val_loss: 10.6733 - val_a
ccuracy: 0.6256
```

```
In [34]: print(acc1*100)           # accuracy of lstm model
```

```
73.6379623413086
```

```
In [ ]:
```