

# What do we know about COVID-19 risk factors?

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#### Part 1 Screenshots:

- 1. Data Preprocessing
  - Eliminating Duplicates

```
# removing duplicate values by usung drop_duplicates
df_covid.drop_duplicates(['abstract', 'body_text'], inplace=True)
df_covid['abstract'].describe(include='all')

count     30184
unique     22480
top
freq     7677
Name: abstract, dtype: object
```

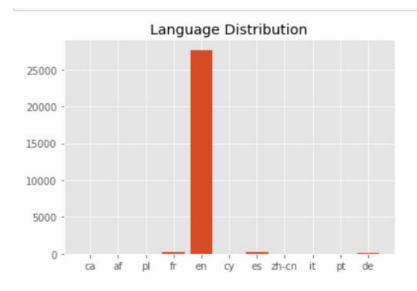
#Removing null values

Removing Null values

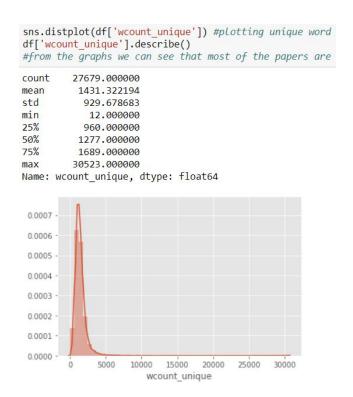
2. Using the Langdetect package to detect different languages.

```
#Printing python data structures by pretty print module
from pprint import pprint
#Langage disctionary to store different language codes
lang dictionary = {}
for lang in set(languages):
    lang_dictionary[lang] = languages.count(lang)
print("Total count: {}\n".format(len(languages)))
#Printing dictionary of words
pprint(lang_dictionary)
Total count: 28314
{'af': 1,
 'ca': 1,
 'cy': 1,
 'de': 48,
 'en': 27679,
 'es': 257,
 'fr': 297,
 'it': 13,
'pl': 2,
 'pt': 14,
 'zh-cn': 1}
```

Plotting different languages



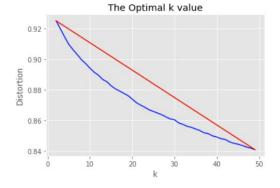
## 3. Plotting words counts



### 4. Plotting distortions and finding optimal K value

```
#Plotting the distortions
X_line = [K[0], K[-1]]
Y_line = [distortions[0], distortions[-1]]

# Plot the elbow
plt.plot(K, distortions, 'b-')
plt.plot(X_line, Y_line, 'r')
plt.xlabel('k')
plt.ylabel('Distortion')
plt.title('The Optimal k value')
plt.show()
```



#### 5. K- means algorithm

```
#K means algorithm with optimal k value, here optimal k value is between 18 to 25
# we will use k=20
k = 20 # defining K
kmeans = KMeans(n_clusters=k, random_state=42, n_jobs=-1) #kmeans model
y_pred = kmeans.fit_predict(x_reduced_pca)
df['y'] = y_pred
```

#### 6. t-SNE algorithm

```
#t-distributed stochastic neighbouring entities
#Dimensionality reduction, visualization for high dimensional dataset
#We use TSNE to reduce the dimensions of the data, bring down higher dimensions to 2D, i.e. x-y plane
from sklearn.manifold import TSNE

tsne_data = TSNE(verbose=1, perplexity=100, random_state=42) # computing tsne
X_embedded_data = tsne_data.fit_transform(X.toarray()) #transforming the data in array
```

#### 7. Plotting the data

```
: #Plotting data using matplot and seaborn modules

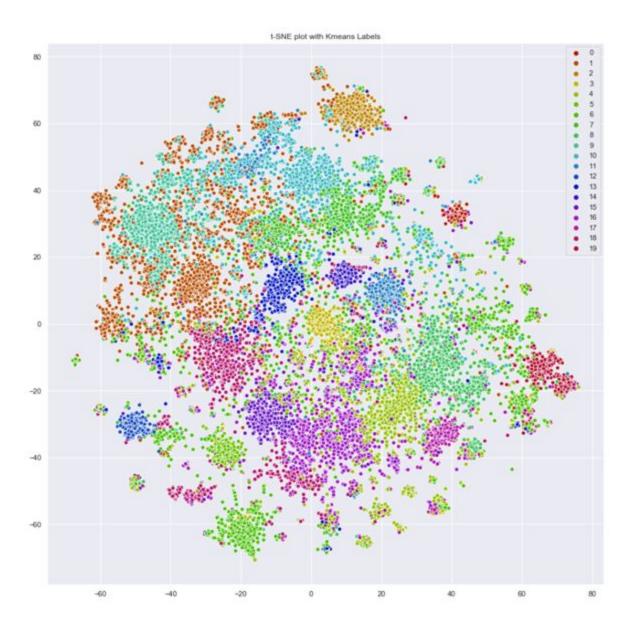
Xmatplotlib inline
from matplotlib import pyplot as plt
import seaborn as sns

# sns settings for the plot
sns.set(rc={'figure.figsize':(15, 15)})

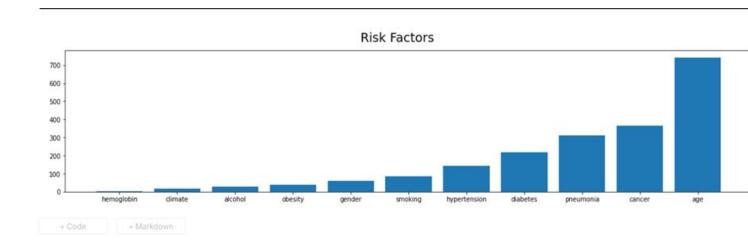
# different colors for the plot
palette = sns.hls_palette(20, l=.4, s=.9)

# plottin the plot with different colors and labels
sns.scatterplot(X_embedded_data[:,0], X_embedded_data[:,1], hue=y_pred, legend='full', palette=palette)
plt.title('t-SNE plot with Kmeans Labels')
#saving plot on the disk
plt.savefig("D:/ITM Sem 2/Data Mining/Project/Plots/cluster_tsne.png")
#displaying the plot
plt.show()
```

# 8. Clustered Data



Results obtained from spaCy pattern matching are a list of risk factors and a bar graph visualizing it.



Results of the top N papers based on query.

