#### CS 6375 - MACHINE LEARNING

### **Project Status Report**

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Our Project is on **predicting the occurrences of "total\_cases"** of Dengue based on "**DengAl: Predicting Disease Spread**" dataset which is in active driven data competitions. Here we have a set of weather information (precipitation, temperature, vegetation) from the two cities: San Juan (sj) and Iquitos (iq) with total cases of dengue by year and week of the year. We aim at making a complete analysis of the DengAl dataset to find the total number of Dengue affected cases in the given two cities with respect to set of climate variables as mentioned above.

#### The Dataset - DengAI: Predicting Disease Spread

The DengAI dataset is taken from an active DrivenData Competition and the link of which is given below:

https://www.drivendata.org/competitions/44/dengai-predicting-disease-spread/

This dataset has two cities: San Juan (sj) and Iquitos (iq). Since we assume that the spread of dengue may follow different patterns between the two different cities, we will divide the dataset, train separate models for each city, and then join our predictions finally.

Number of attributes = 24

Number of instances = 1456

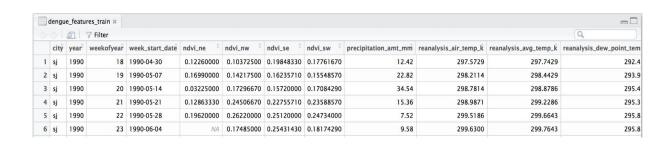
The attributes are the following:

- city we will divide all city = San Juan (sj) data into one dataset and
   all city = Iquitos (iq) data into other dataset
- 2. year

- 3. week\_of\_year
- 4. week\_start\_date
- 5. station\_max\_temp\_c Maximum temperature
- 6. station\_min\_temp\_c Minimum temperature
- 7. station\_avg\_temp\_c Average temperature
- 8. station\_precip\_mm Total precipitation
- station\_diur\_temp\_rng\_c Diurnal temperature
  - 10. precipitation\_amt\_mm Total precipitation
- 11. reanalysis\_sat\_precip\_amt\_mm Total precipitation
- 12. reanalysis\_dew\_point\_temp\_k Mean dew point temperature
  - 13. reanalysis\_air\_temp\_k Mean air temperature
- 14. reanalysis\_relative\_humidity\_percent Mean relative humidity
- 15. reanalysis\_specific\_humidity\_g\_per\_kg Mean specific humidity
- 16. reanalysis\_precip\_amt\_kg\_per\_m2 Total precipitation
- ${\bf 17.\ reanalysis\_max\_air\_temp\_k-Maximum\ air}$   ${\bf temperature}$
- ${\bf 18.\ reanalysis\_min\_air\_temp\_k-Minimum\ air}$   ${\bf temperature}$ 
  - 19. reanalysis\_avg\_temp\_k Average air temperature
  - 20. ndvi se Pixel southeast of city centroid
  - 21. ndvi sw Pixel southwest of city centroid
  - 22. ndvi\_ne Pixel northeast of city centroid
  - 23. ndvi\_nw Pixel northwest of city centroid
  - 24. reanalysis\_tdtr\_k Diurnal temperature range

Here is the snapshot of the data:

dengue\_features\_train <- read\_csv ("~/Downloads/dengue\_features\_train.csv ")
head(dengue\_features\_train)</pre>



station_precip_mm	station_min_temp_c	station_max_temp_c	station_diur_temp_rng_c	station_avg_temp_c	reanalysis_tdtr_k
16.0	20.0	29.4	6.900000	25.44286	2.628571
8.6	22.2	31.7	6.371429	26.71429	2.371429
41.4	22.8	32.2	6.485714	26.71429	2.300000
4.0	23.3	33.3	6.771429	27.47143	2.428571
5.8	23.9	35.0	9.371429	28.94286	3.014286
39.1	23.9	34.4	6.942857	28.11429	2.100000

dengue\_labels\_train <- read\_csv("~/Downloads/dengue\_labels\_train.csv")
head(dengue\_labels\_train)</pre>



We are dividing dengue\_features\_train and dengue\_labels\_train into two datasets based on the city value by using the following commands:

```
sj_dengue_train_labels<-subset(dengue_labels_train,city=="sj")
iq_dengue_train_labels<-subset(dengue_labels_train,city=="iq")
sj_dengue_train_features<-subset(dengue_train_features,city=="sj")
iq_dengue_train_features<-subset(dengue_train_features,city=="iq")</pre>
```

#### <u>Techniques we planned to use</u>

We planned to apply the following techniques on the data to complete the required analysis –

- k-Nearest Neighbors (kNN)
- Random Forests
- Bagging
- Gradient Boosting

#### **Experimental Methodology**

We employ the following procedure in our project -

- 1. Pre-processing of the dataset
  - This step involves dealing with the NA values,
  - Scaling the required attributes,
  - Removing the uncorrelated attributes.
- 2. On the dataset
  - We perform each of the aforementioned techniques,
  - Also, vary the parameters and find the best set of parameters for the technique.
- 3. We evaluate the techniques using the following metrics
  - Accuracy
  - Precision
  - Recall

- F-measure
- 4. We plot the results that aid in comparing the performance of the classifiers.

## **Programming Language**

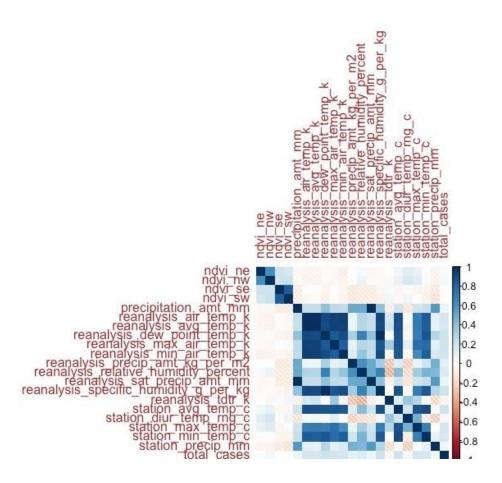
We plan to use **R** programming for the project.

## **Preliminary Results**

We now present the results of the work we've done so far.

- → Removing "week\_start\_date" attribute since it is not a feature for our model and removing it will not make any difference in the result.
- → We plotted the CORRPLOT which aids in identifying the correlation of the attribute with the class attribute (total\_cases).

The plot is as follows for each of the cities Iquitos (iq) and San Juan (sj) :-



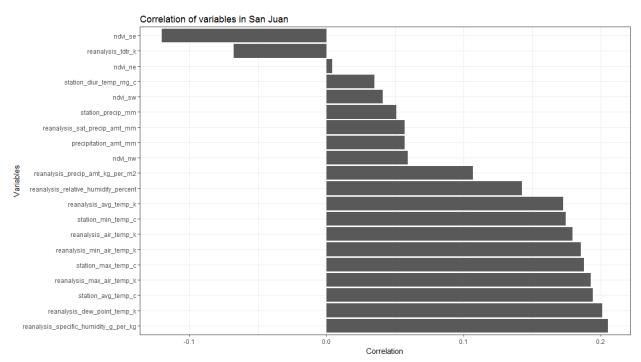
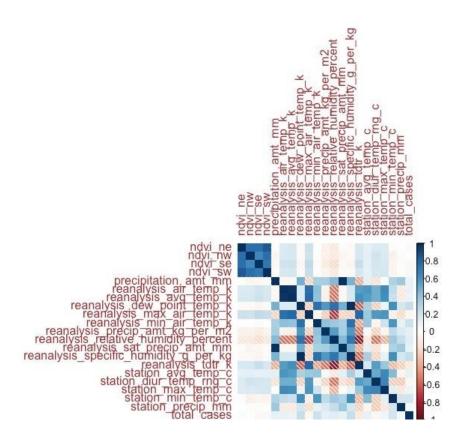


Fig 1. Correlation heat plot and bar plot of city San Juan (sj)



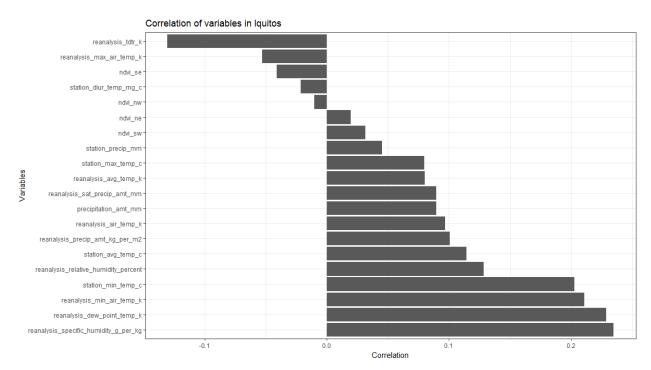


Fig.2 Correlation heat plot and bar plot of city Iquitos

#### **Observations**

- 1. We see that many of the temperature data are strongly correlated.
- 2. We noted that total\_cases has weak correlations with the other attributes and vegetation\_index attributes also has weak correlations with other attributes.
- 3. We also see that correlation strengths differ for each city and reanalysis\_specific\_humidity\_g\_per\_kg and reanalysis\_dew\_point\_temp\_k are the most strongly correlated with total\_cases. As we know that mosquitos thrive wet climates, we can infer why they are strongly correlated.
- 4. As minimum temperatures, maximum temperatures, and average temperatures rise, the total\_cases of dengue fever tend to rise as well.
- 5. We also note that the precipitation measurements bear little to no correlation to total\_cases, despite strong correlations to the humidity measurements.

Based on the correlation observations above, these are the attributes which have the strong correlations with the total cases in each city:

- reanalysis\_specific\_humidity\_g\_per\_kg
- reanalysis\_dew\_point\_temp\_k
- station\_avg\_temp\_c
- station\_min\_temp\_c

#### R\_CODE:

```
#Read the dataset

dengue_labels_train <- read_csv("~/Downloads/dengue_labels_train.csv")

dengue_features_train <- read_csv("~/Downloads/dengue_features_train.csv")

View(dengue_labels_train)

View(dengue_features_train)

#Loading data by City

sj_dengue_train_labels<-subset(dengue_labels_train,city=="sj")

iq_dengue_train_labels<-subset(dengue_labels_train,city=="iq")
```

```
sj_dengue_train_features<-subset(dengue_features_train,city=="sj")
iq_dengue_train_features<-subset(dengue_features_train,city=="iq")</pre>
#Merging features and labels
merged\_sj\_train\_features\_instances <-
merge(sj_dengue_train_features,sj_dengue_train_labels,by=c('city','year','weekofyear'))
merged_iq_train_features_instances <-
merge(iq_dengue_train_features,iq_dengue_train_labels,by=c('city','year','weekofyear'))
#Pre-processing
When the data set is loaded in R, the null values are replaced by NA.
We have used "gam" package in which NAs are replaced by the mean of the non-missing entries.
library(gam)
merged_sj_train_features_instances <- na.gam.replace(merged_sj_train_features_instances)
merged_iq_train_features_instances <- na.gam.replace(merged_iq_train_features_instances)
View(merged_sj_train_features_instances)
View(merged_iq_train_features_instances)
# Removing 'week_start_date' column
As it doesn't impact the result significantly.
We have used "dplyr" package is used to remove unnecessary columns.
merged_sj_train_features_instances <- dplyr::select(merged_sj_train_features_instances, -week_start_date)
merged_iq_train_features_instances <- dplyr::select(merged_iq_train_features_instances, -week_start_date)</pre>
View(merged_sj_train_features_instances)
View(merged_iq_train_features_instances)
#Finding the correlation plot
library(corrplot)
sj_corrplot<-cor(merged_sj_train_features_instances[,4:24])
```

corrplot(sj\_corrplot,type = 'full', tl.col = 'brown', method="shade")

iq\_corrplot<-cor(merged\_iq\_train\_features\_instances[,4:24])
corrplot(iq\_corrplot,type = 'full', tl.col = 'brown', method="shade")</pre>