

Data Reading

```
Tobacco2<-read.table("D:/cscd477/assignment4/data/Youth_Tobacco_Survey__YTS__Data.csv",
header=TRUE, sep=";", quote="\")
```

Data cleaning:

```
TobaccoModified2<-Tobacco2[Tobacco2$Response=="Current" & Tobacco2$DisplayOrder==7,]
```

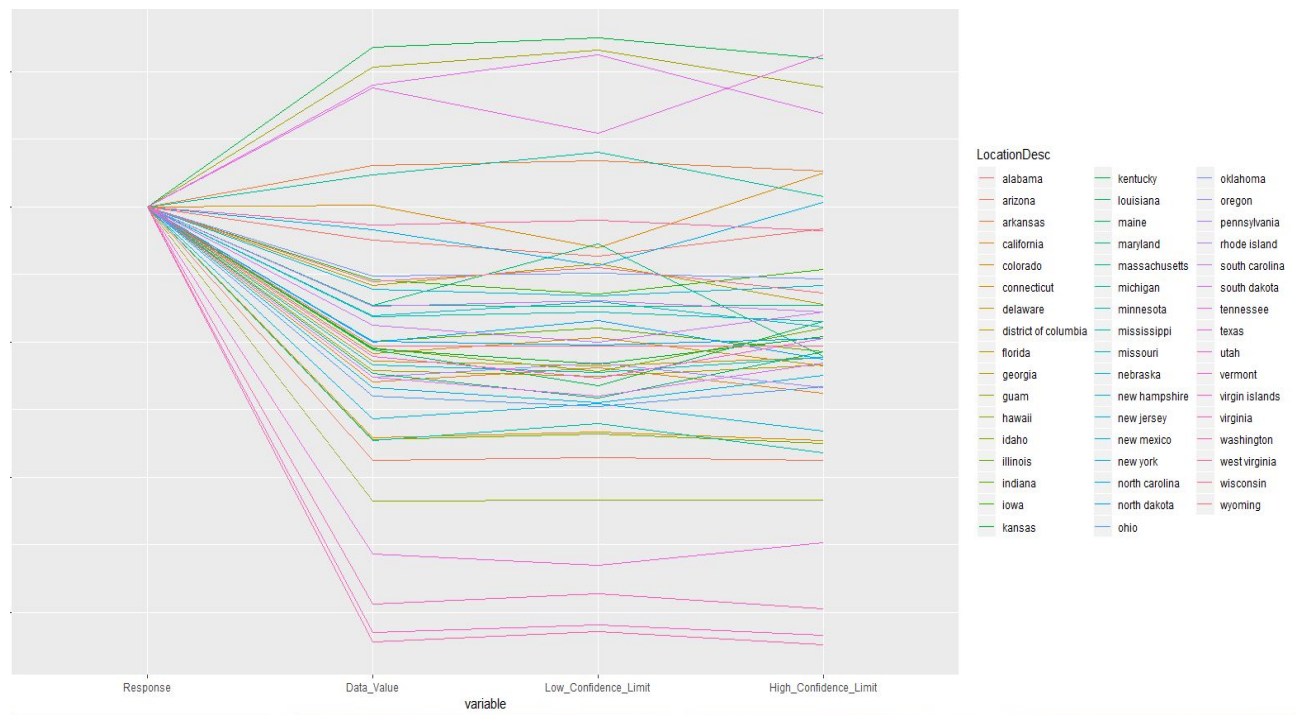
```
TobaccoModifiedFinal2<-TobaccoModified2[,c("LocationDesc","Response","Data_Value","Low_
Confidence_Limit","High_Confidence_Limit","DisplayOrder")]
```

```
TobaccoModifiedFinalAverage2<-aggregate(.$LocationDesc,data=TobaccoModifiedFinal2,FUN=
mean)
```

```
TobaccoModifiedFinalAverage2$LocationDesc<-tolower(TobaccoModifiedFinalAverage2$Locati
onDesc)
```

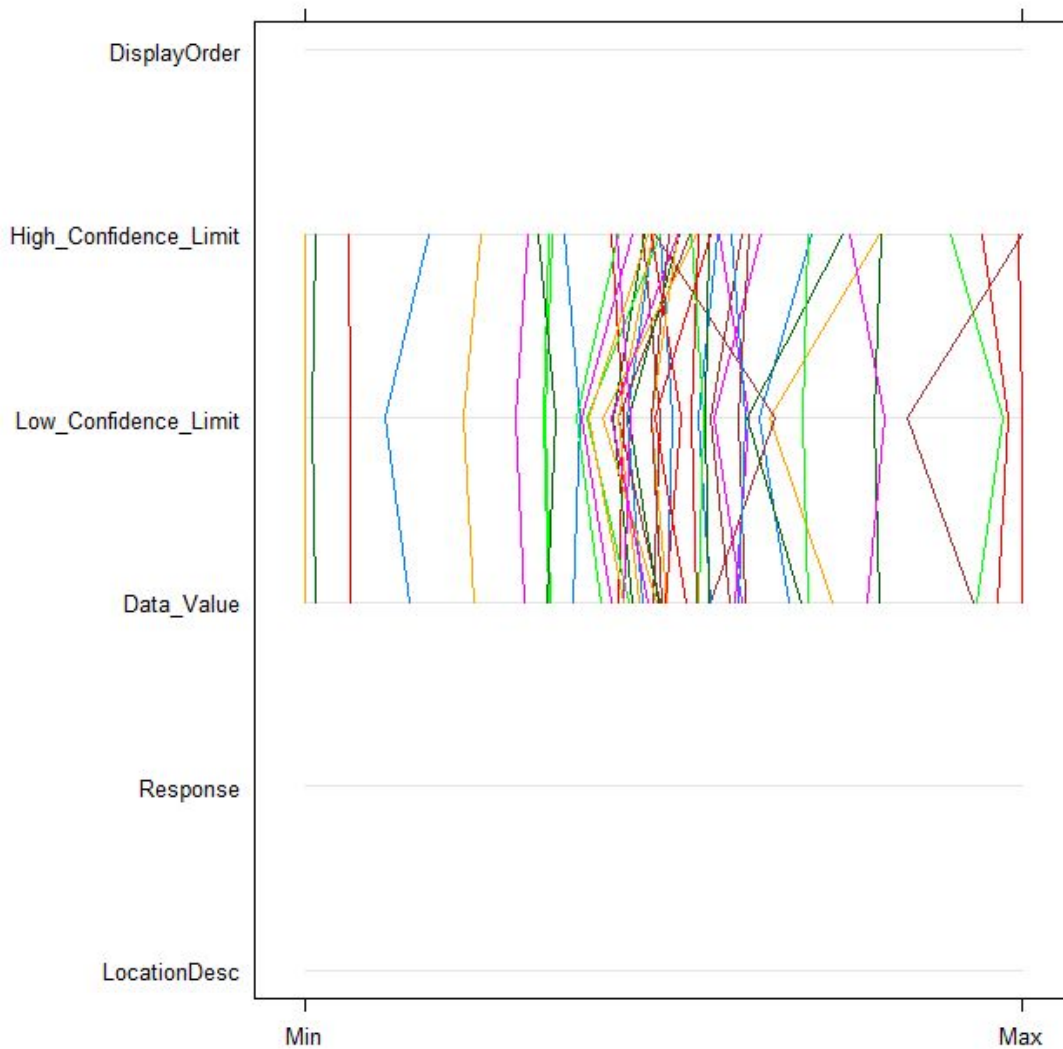
```
> TobaccoModifiedFinalAverage2[1:20,]
      LocationDesc Response Data_Value Low_Confidence_Limit High_Confidence_Limit DisplayOrder
1      alabama      2    15.959259      12.492593      19.425926              7
2      arizona      2     8.800000       6.613889      10.997222              7
3      arkansas      2    18.391667      15.275000      21.512500              7
4      california      2    11.333333       9.233333      13.441667              7
5      colorado      2    17.100000      12.750000      21.450000              7
6      connecticut      2     9.517647       7.341176      11.703922              7
7      delaware      2    12.264583      10.110417      14.414583              7
8 district of columbia      2    12.016667       9.300000      14.733333              7
9      florida      2    14.466667      12.266667      16.666667              7
10     georgia      2    11.717949       8.969231      14.469231              7
11      guam      2    21.566667      18.533333      24.550000              7
12     hawaii      2     7.471429       5.364286       9.566667              7
13     idaho      2     9.466667       7.283333      11.600000              7
14     illinois      2    12.461111       9.113889      15.813889              7
15     indiana      2    12.658974      10.389744      14.930769              7
16     iowa      2    14.666667      11.391667      17.938889              7
17     kansas      2    12.429167       9.350000      15.512500              7
18     kentucky      2    22.236364      18.878788      25.596970              7
19     louisiana      2    12.375000       8.713889      16.033333              7
20      maine      2    11.616667       8.333333      14.933333              7
> |
```

```
ggparcoord(TobaccoModifiedFinalAverage2, columns = 2:5, groupColumn= "LocationDesc",
title="Youth Tobacco Survey")
```

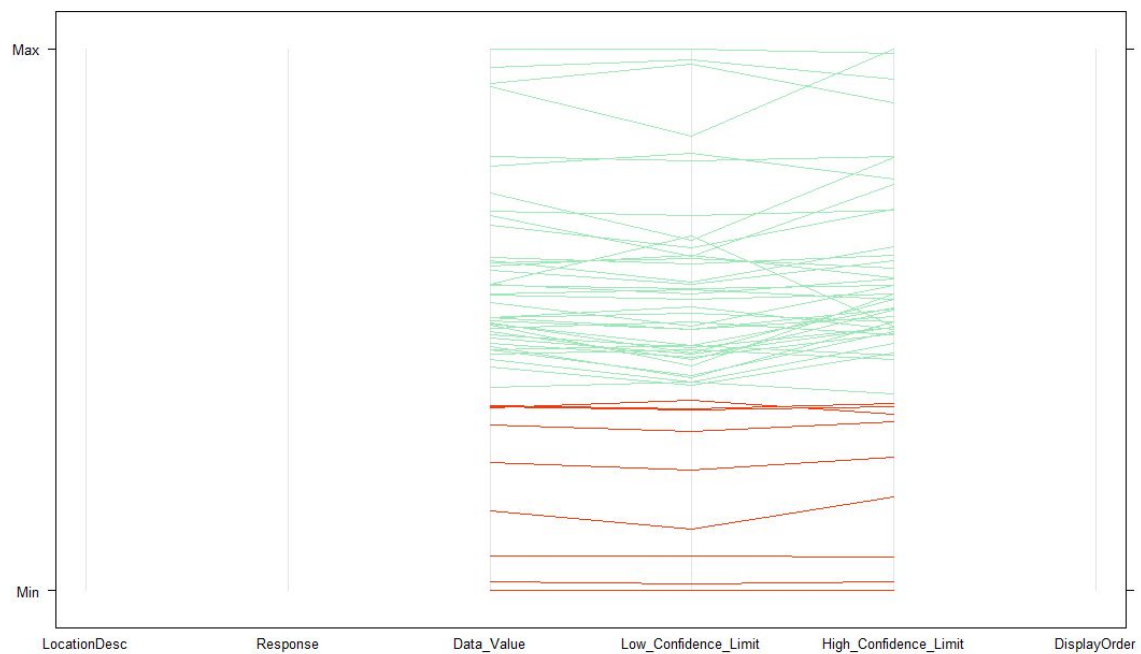


Making Clusters:

`parallelplot(TobaccoModifiedFinalAverage2)`



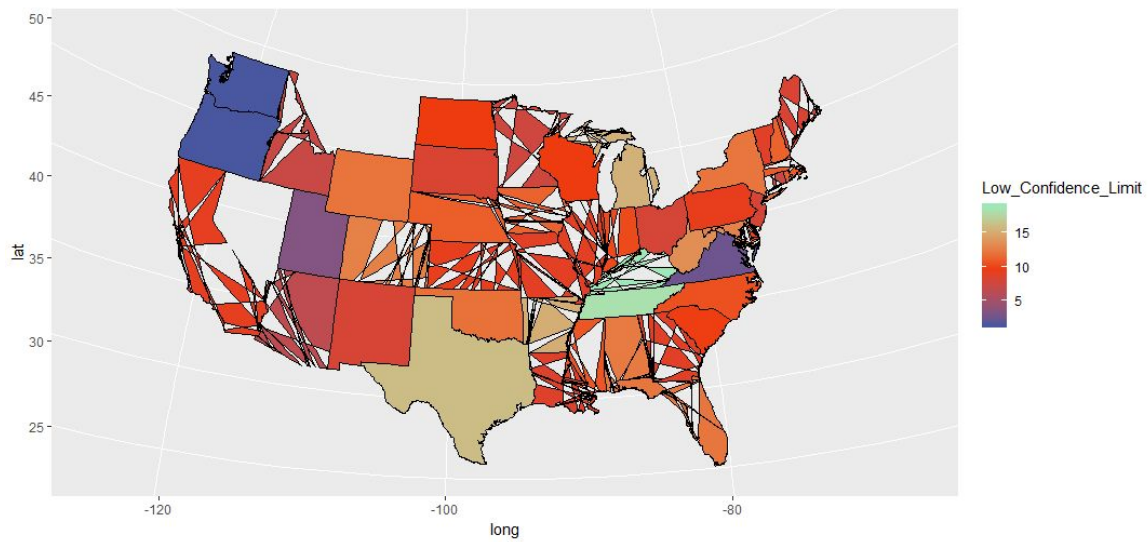
```
reading_colors<-c()
for (i in 1:length(TobaccoModifiedFinalAverage2$LocationDesc)){if
(TobaccoModifiedFinalAverage2$Data_Value[i] > 10){col<-"#9fe9b9"}else{col<-"#ef3b10" }
reading_colors<-c(reading_colors, col)}
parallelplot(TobaccoModifiedFinalAverage2, horizontal.axis=FALSE, col=reading_colors)
```



```
TobaccoAverageMap2<-merge(states_map,TobaccoModifiedFinalAverage2, by.x="region",
by.y="LocationDesc")
```

```
ggplot(TobaccoAverageMap2, aes(x=long, y=lat, group=group,
fill=Low_Confidence_Limit))+geom_polygon(colour="black")+coord_map("polyconic")
```

```
ggplot(TobaccoAverageMap2, aes(x=long, y=lat , group=group,
fill=Low_Confidence_Limit))+geom_polygon(colour="black")+scale_fill_gradient2(low="#2158aa",mid="#ef3b10",
high="#9fe9b9",midpoint=median(TobaccoModifiedFinalAverage2$Low_Confidence_Limit)) +
coord_map("polyconic")
```



Conclusion:

For this Tobacco Survey, we can make relationship with each variables by parallel plot. With parallel plot we can easily compare each state. Correlations can be observed as states are plotted on the chart. Each state corresponds to a line drawn through point on each axis corresponding to the value of the variable.

Yes, Clustering help in visualizing information. For example, In this information, if the Data_value is more than 10 then the color is greenish and if less than 10 then the color is redish.