age distribution of cases

Maggie Walters
June 22, 2017

Document Synopsis

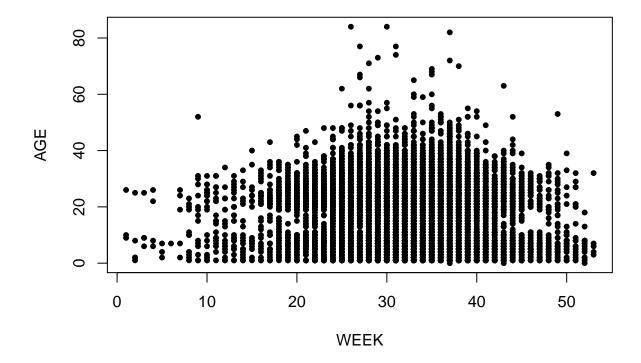
This document aims to visually and quantitatively describe the different age distribution of cases in the counties represented in spm.data.long.csv. Densities of cases for age groups will be calculated and then compared between counties.

The difference between counties will be used in order to explore possible predictive relationships between population size or urbanness of a county and it's density of cases in different age classes.

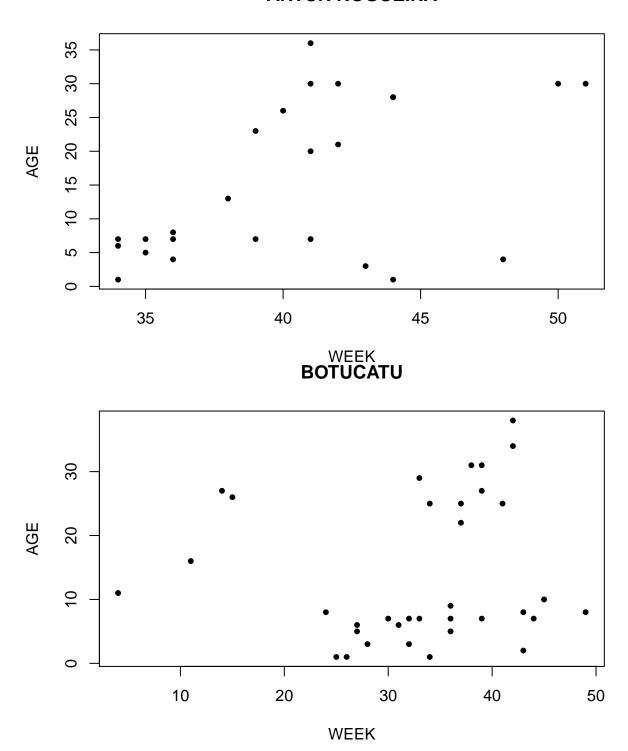
Age distribution scatter plots by county

Scatter plots will be created in order to visually identify the age groups which frequently have a low density of cases. Depending on results, this may be limited to counties with number of cases above a certain threshold.

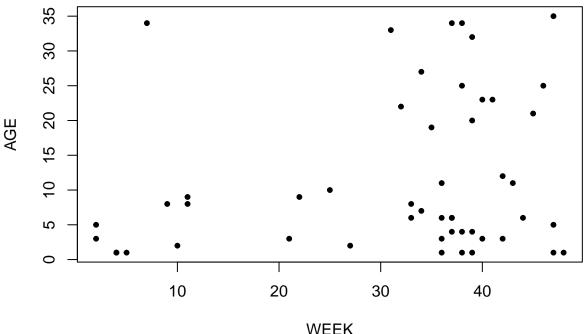
SAO PAULO



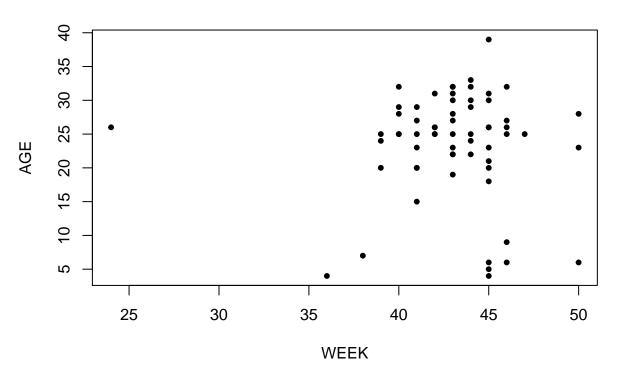
ARTUR NOGUEIRA



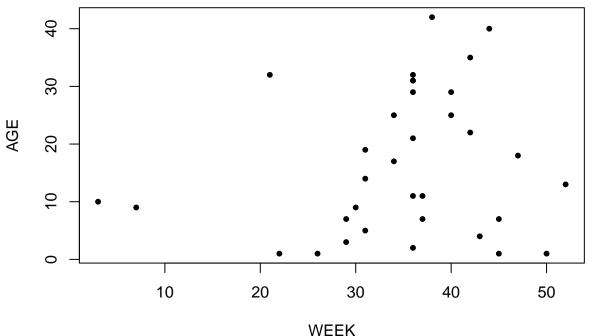
MOGI GUACU



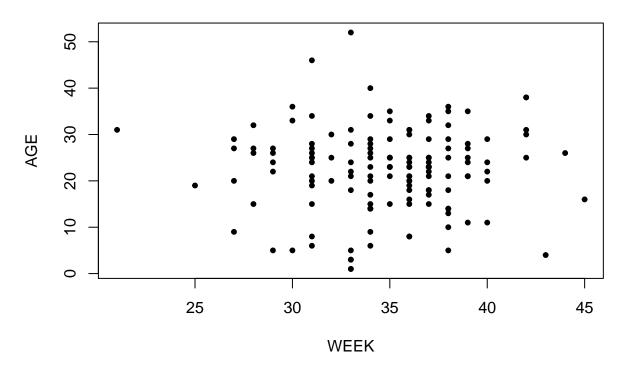
WEEK **ENGENHEIRO COELHO**



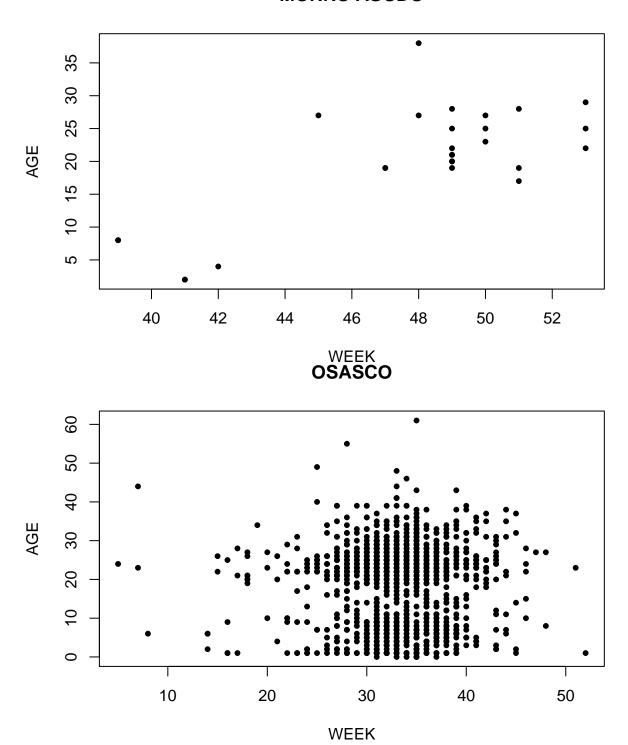
BAURU



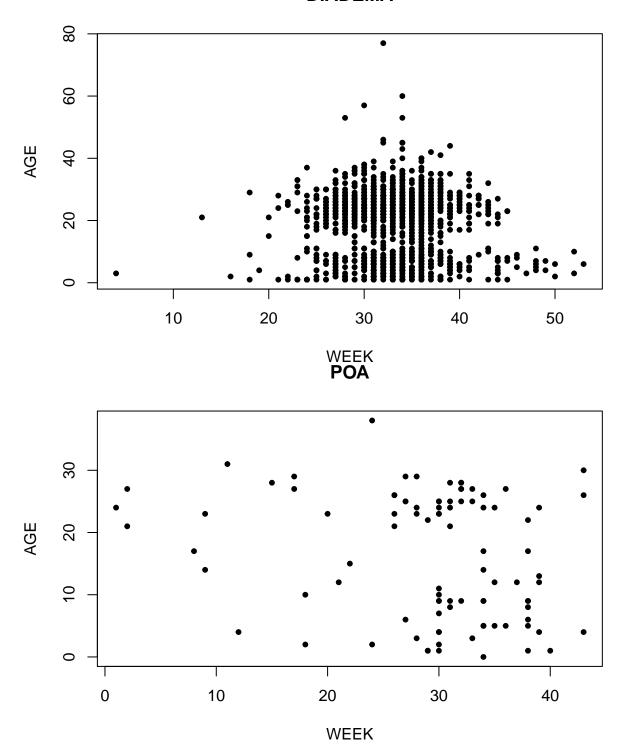




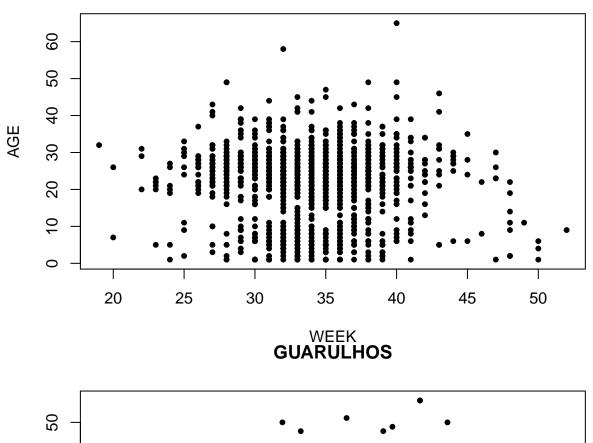
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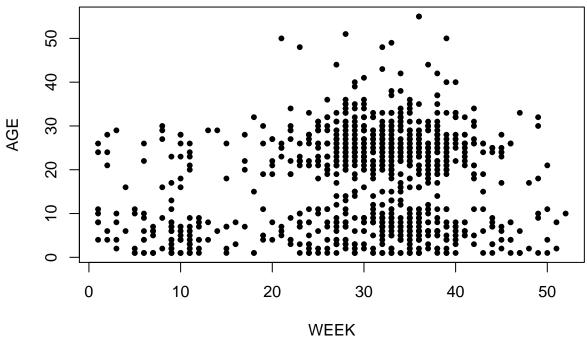


DIADEMA

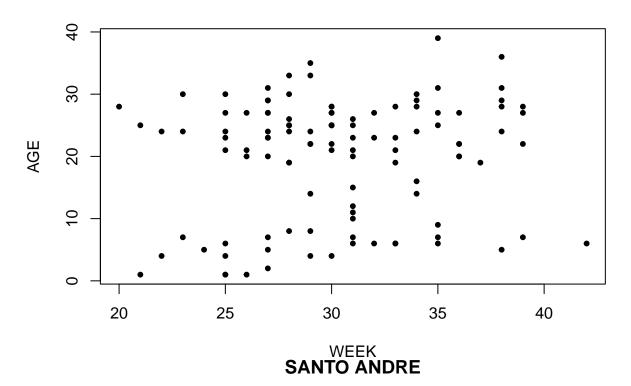


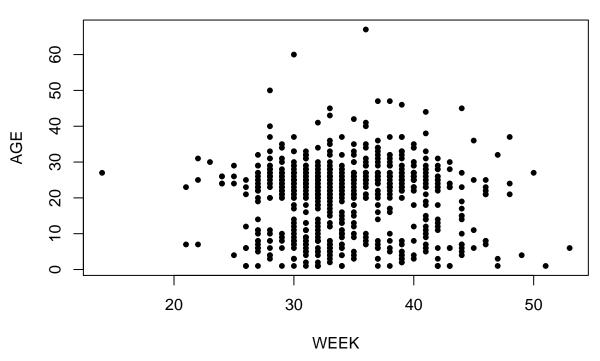
SAO BERNARDO DO CAMPO



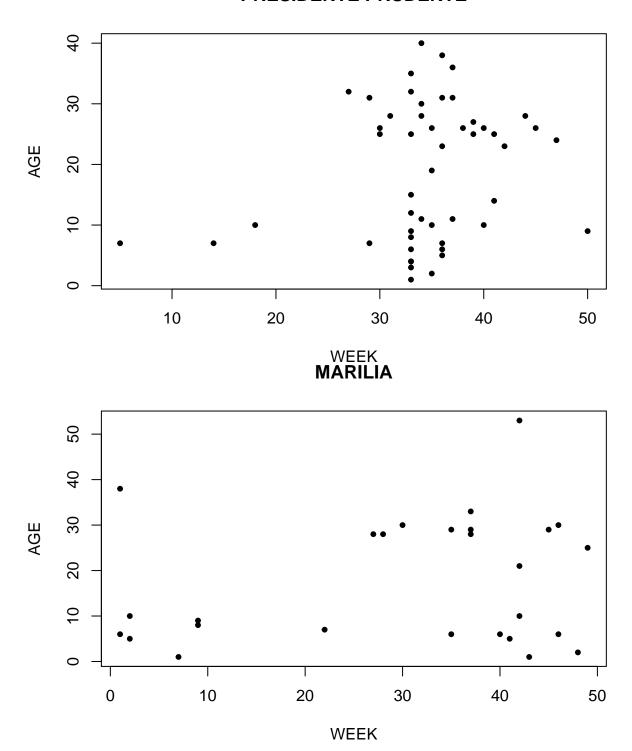


CAJAMAR

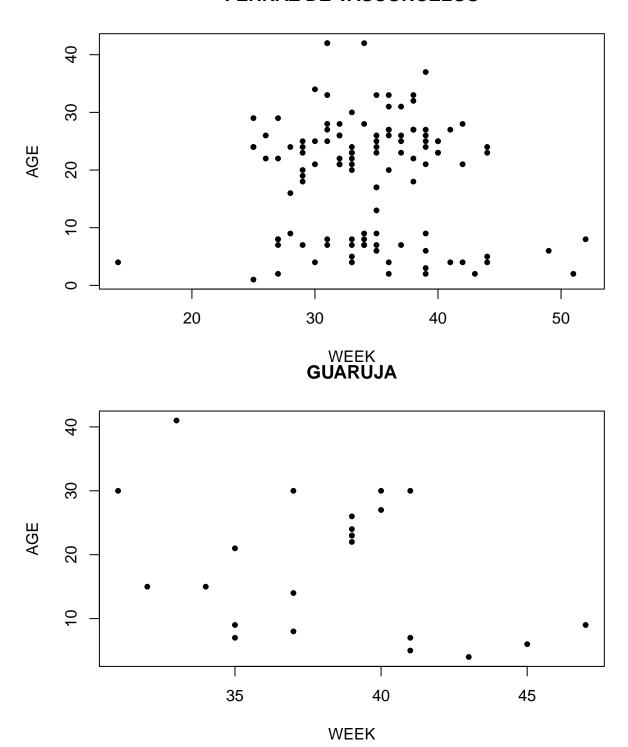




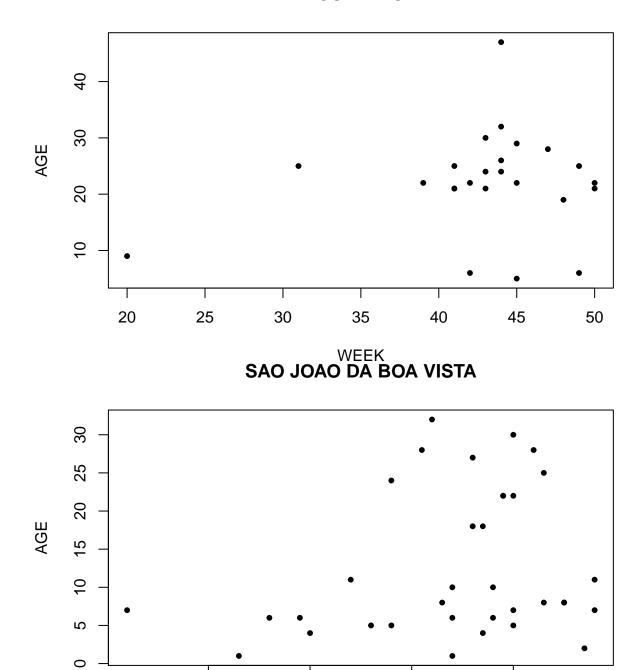
PRESIDENTE PRUDENTE



FERRAZ DE VASCONCELOS

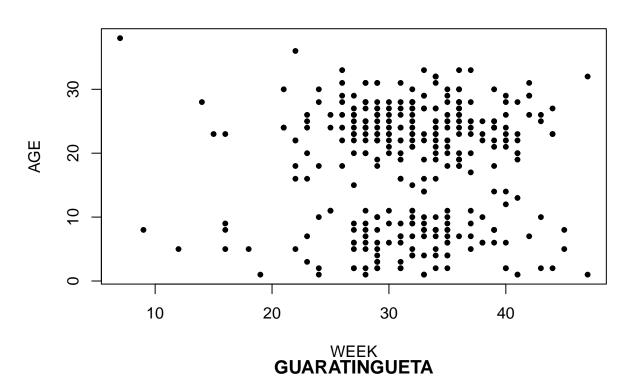


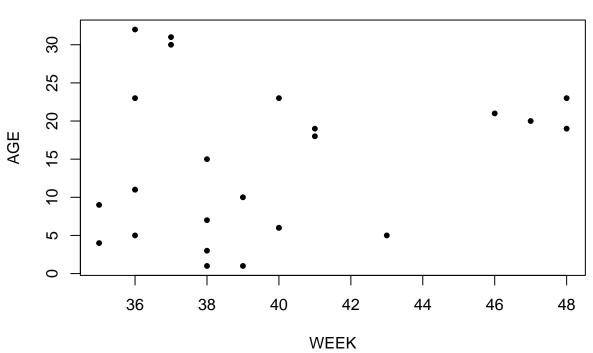
CUBATAO



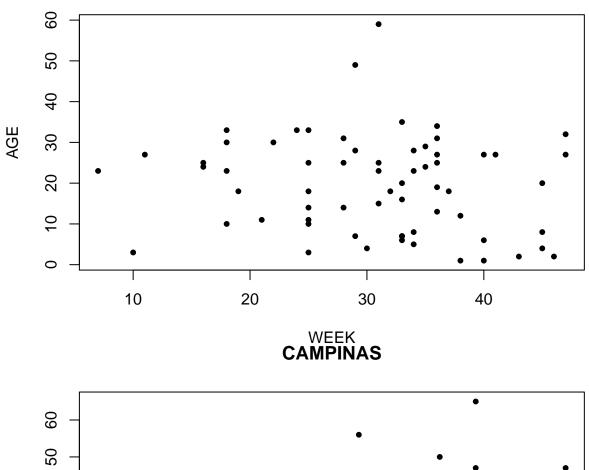
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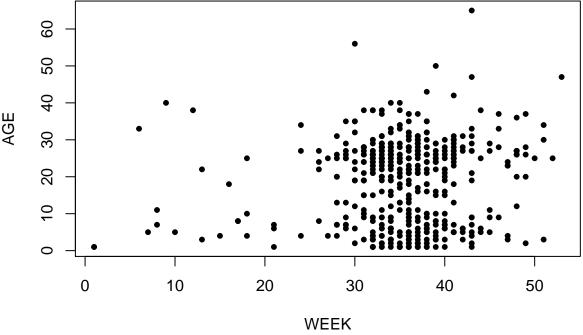
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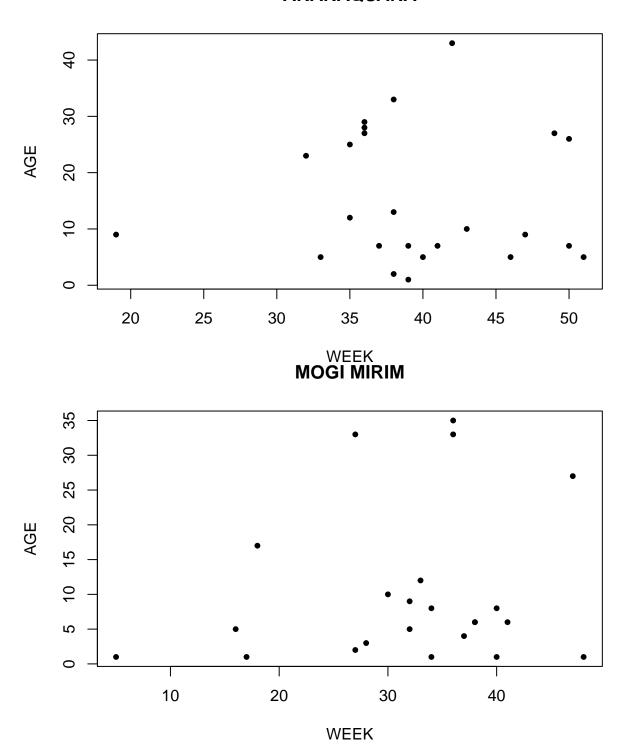


SANTOS

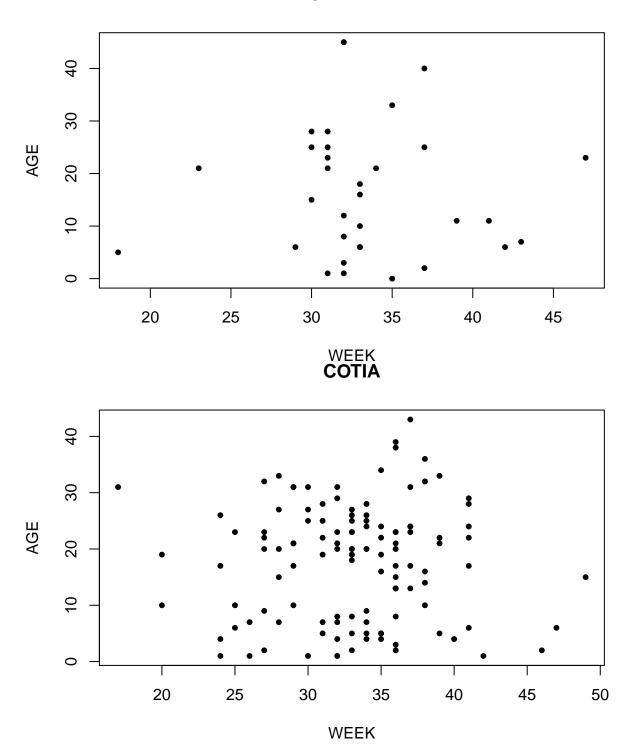




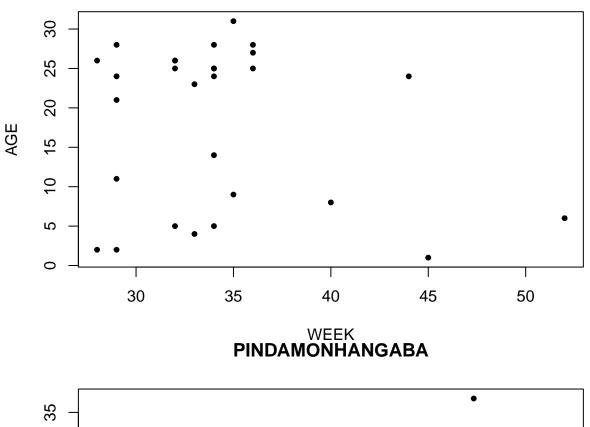
ARARAQUARA

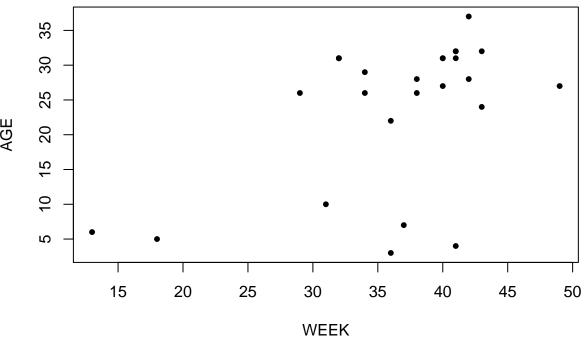


JANDIRA

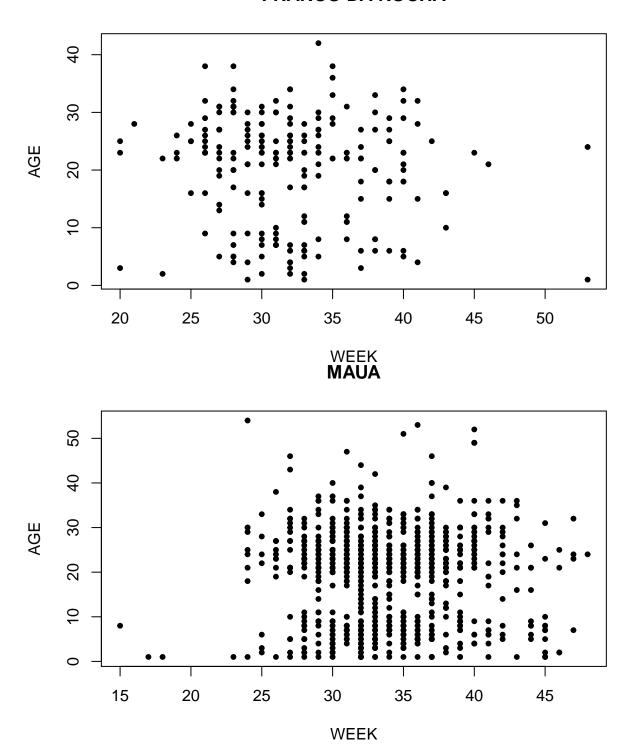


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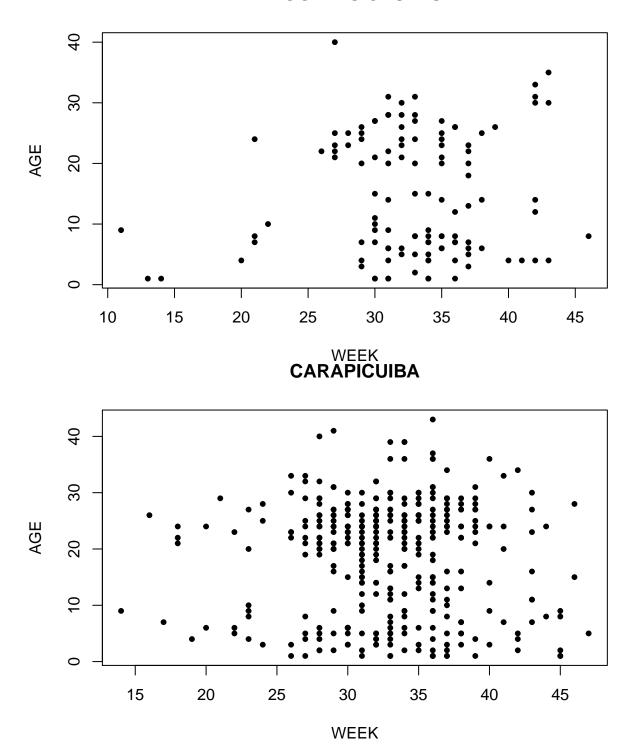




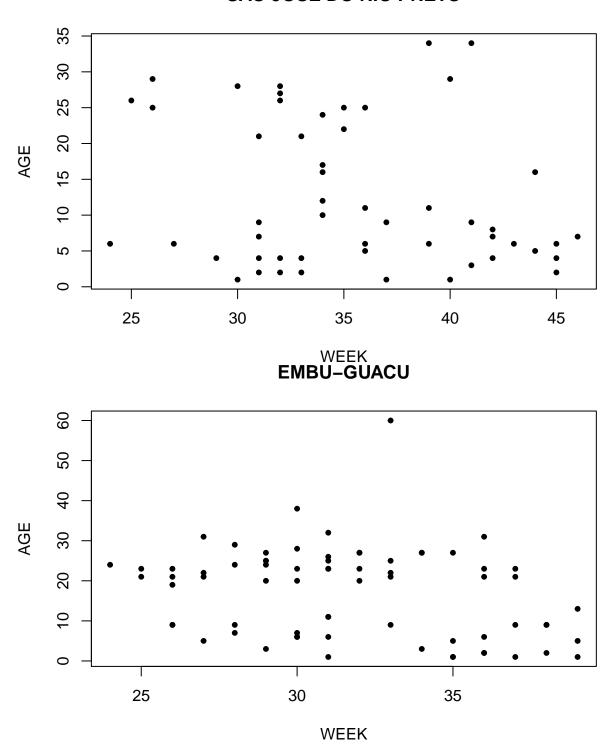
FRANCO DA ROCHA



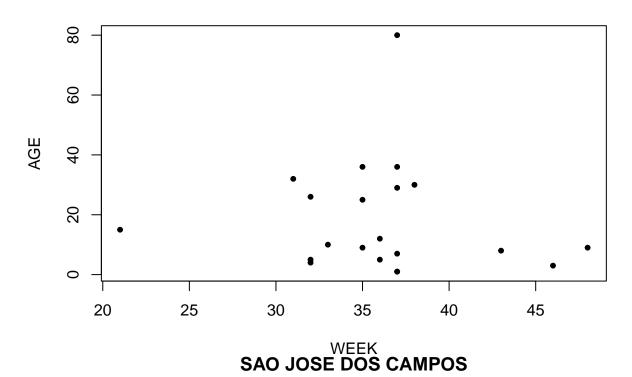
MOGI DAS CRUZES

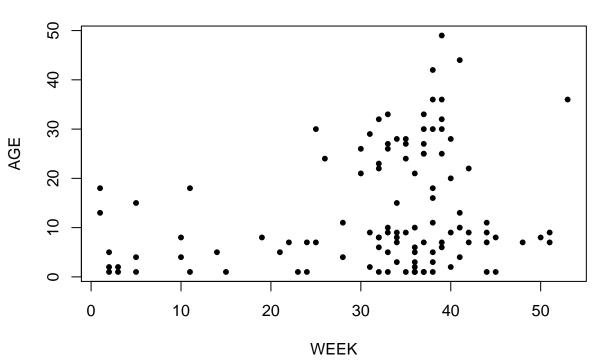


SAO JOSE DO RIO PRETO

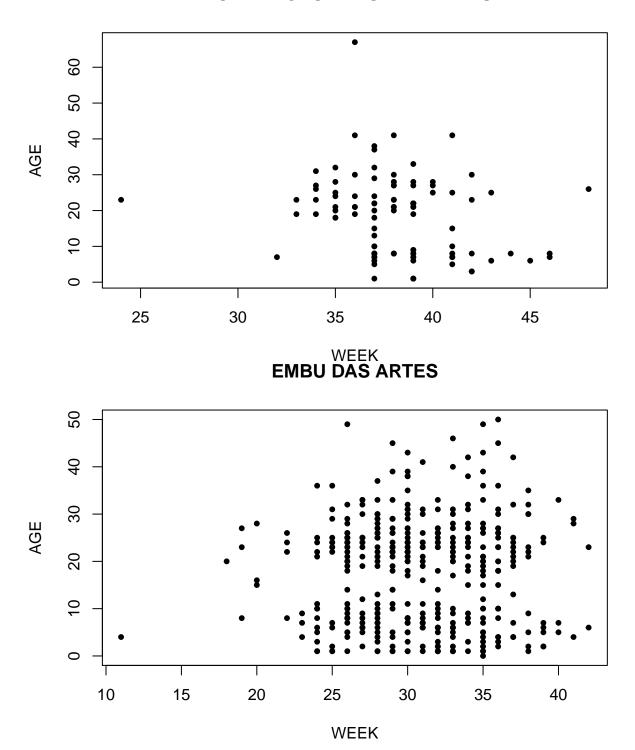


CATANDUVA

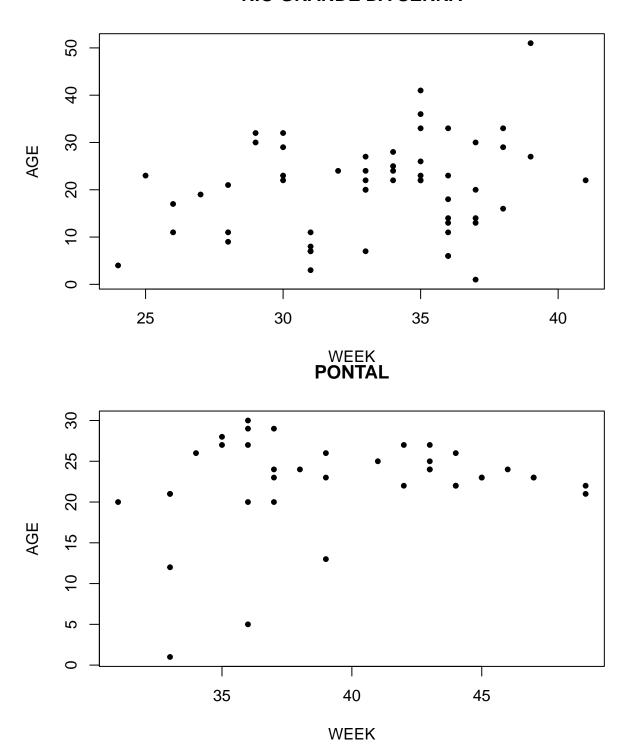




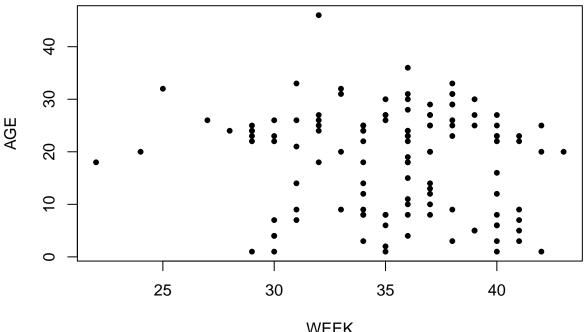
SANTA CRUZ DAS PALMEIRAS



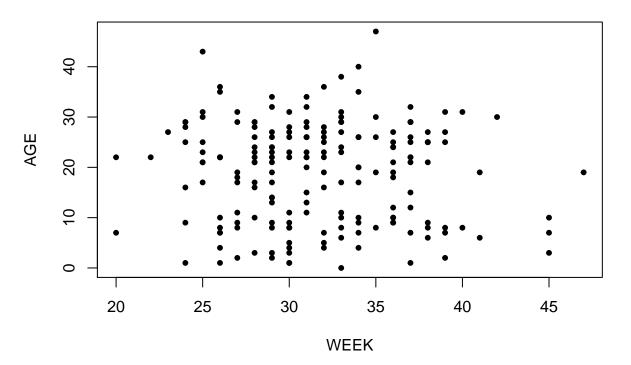
RIO GRANDE DA SERRA



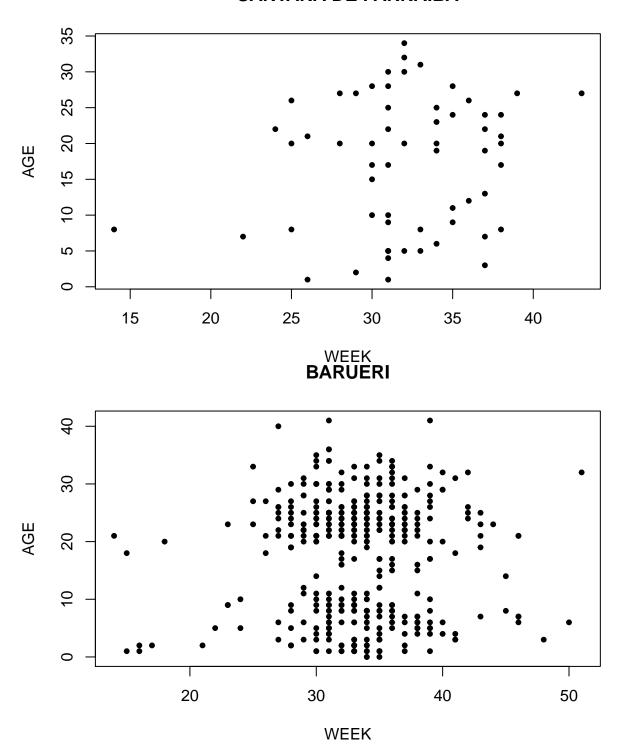
RIBEIRAO PIRES



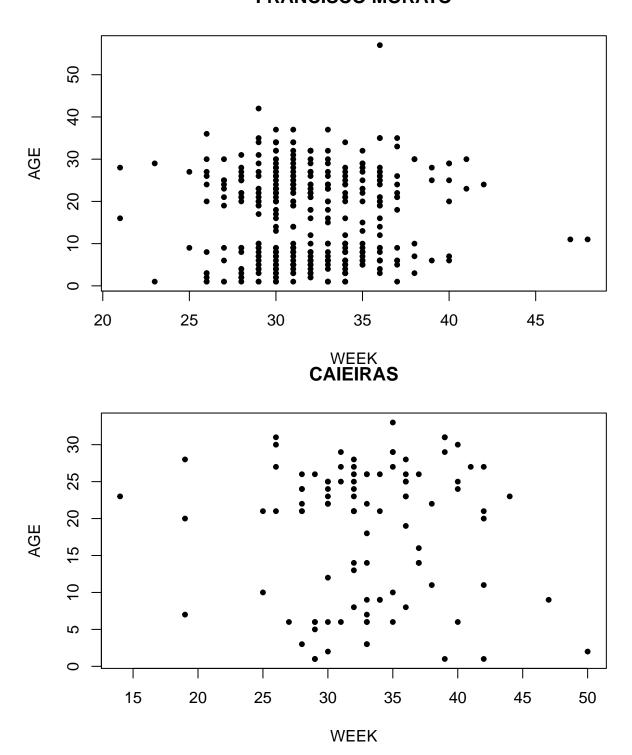




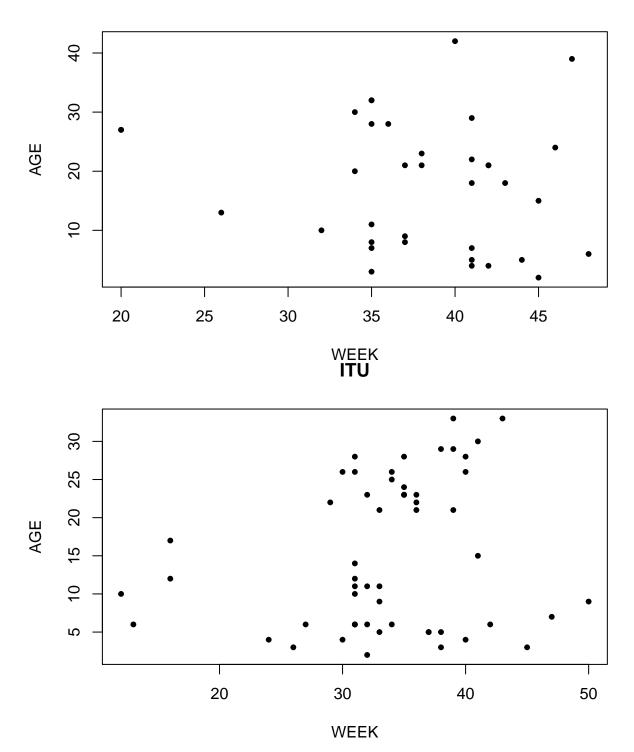
SANTANA DE PARNAIBA



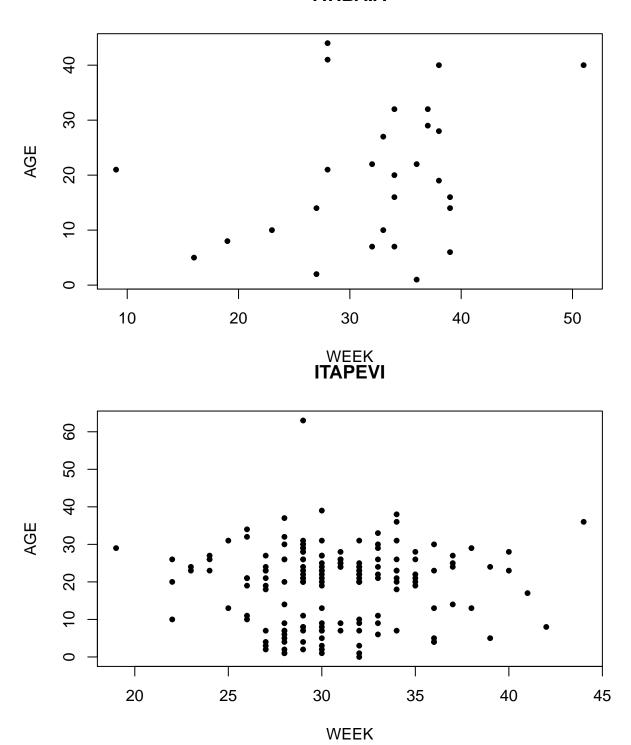
FRANCISCO MORATO



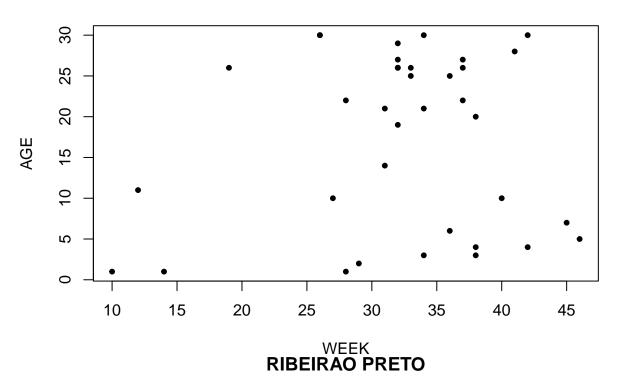


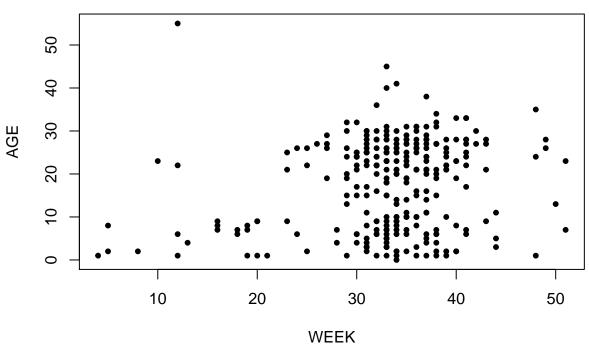


ATIBAIA

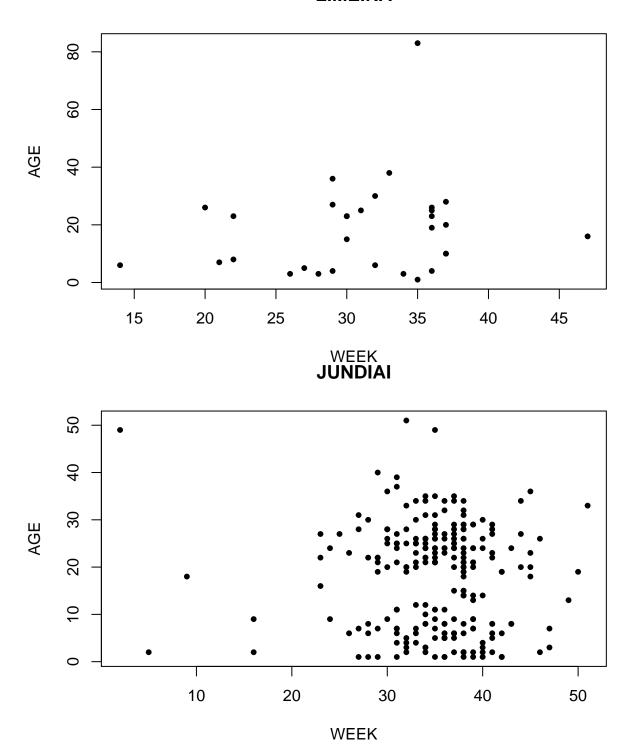


MAIRIPORA

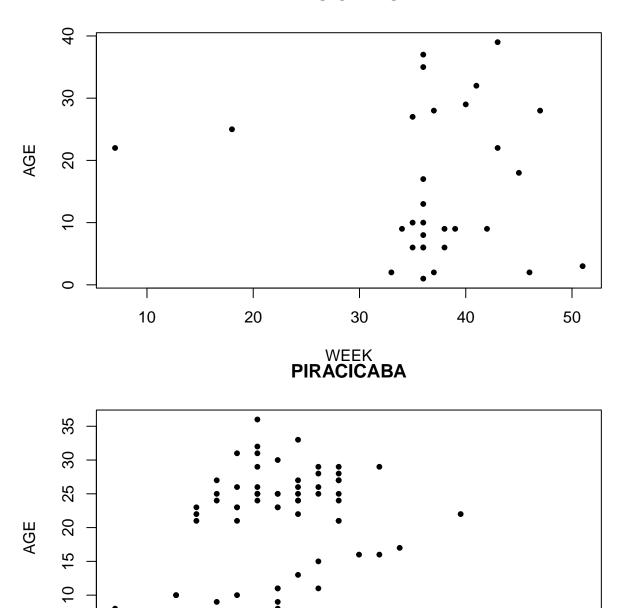




LIMEIRA

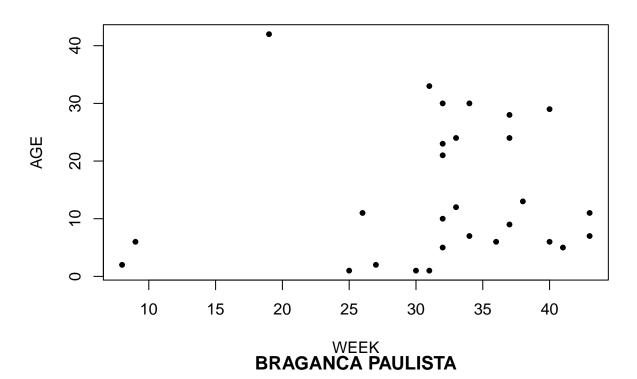


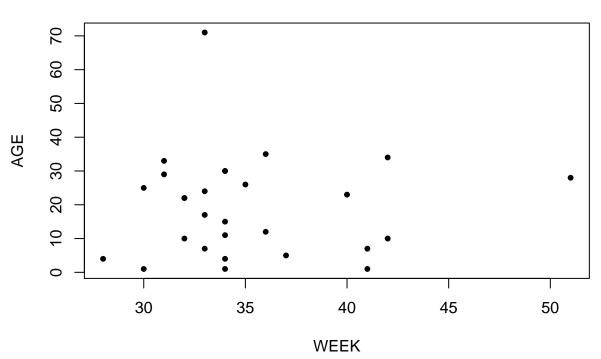
RIO CLARO



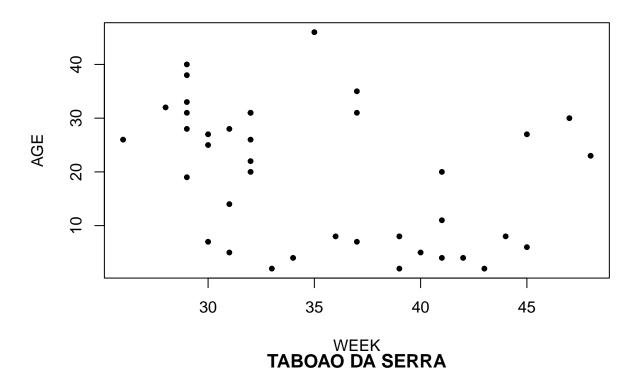
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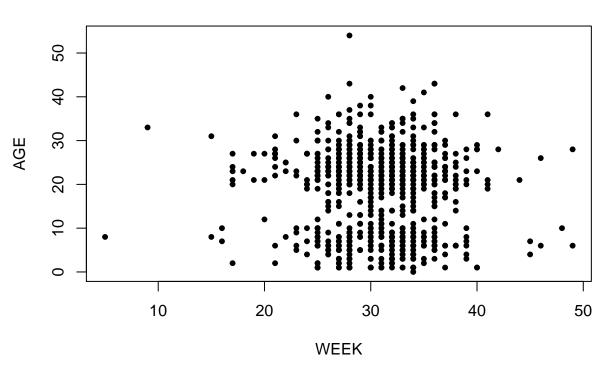
ARUJA



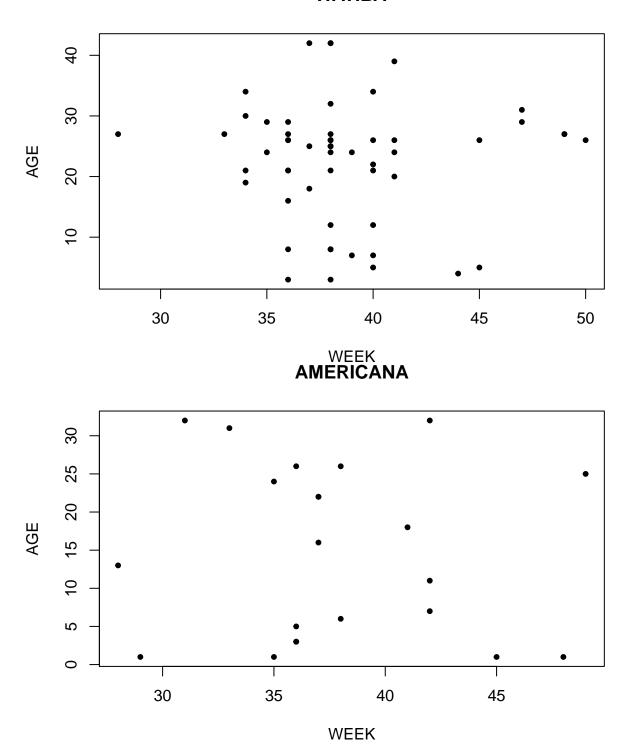


CAMPO LIMPO PAULISTA

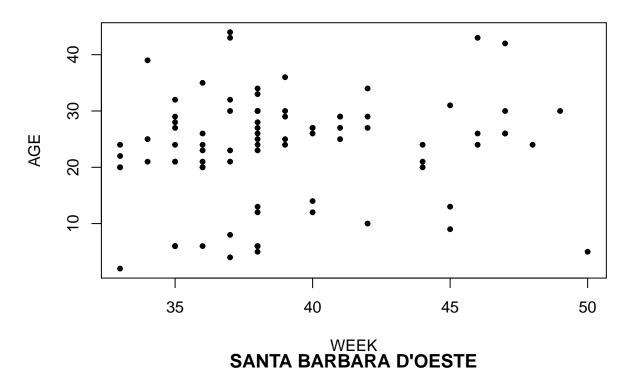


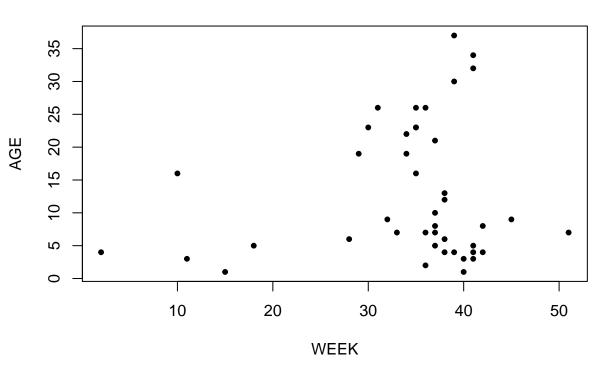


ITATIBA

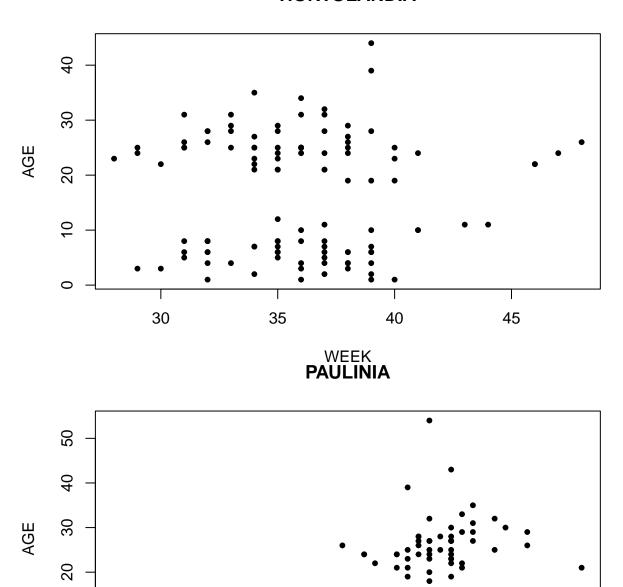


COSMOPOLIS



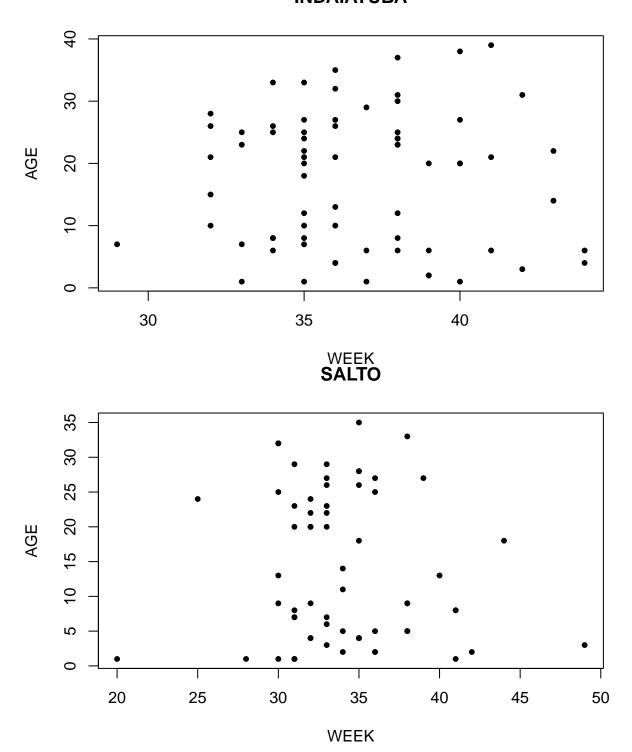


HORTOLANDIA

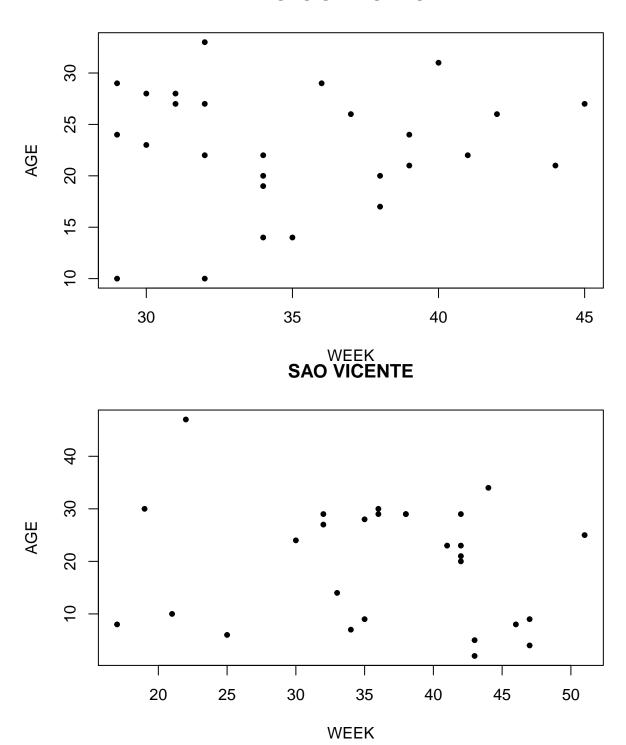


WEEK

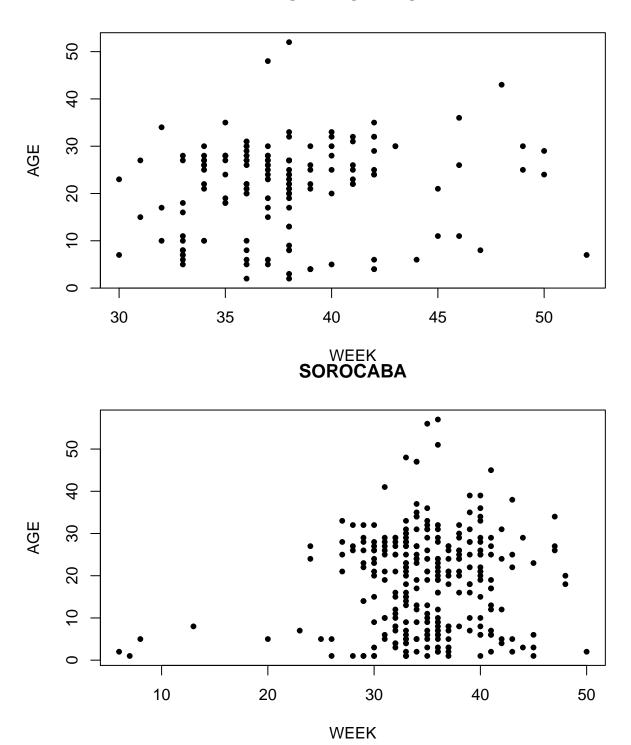
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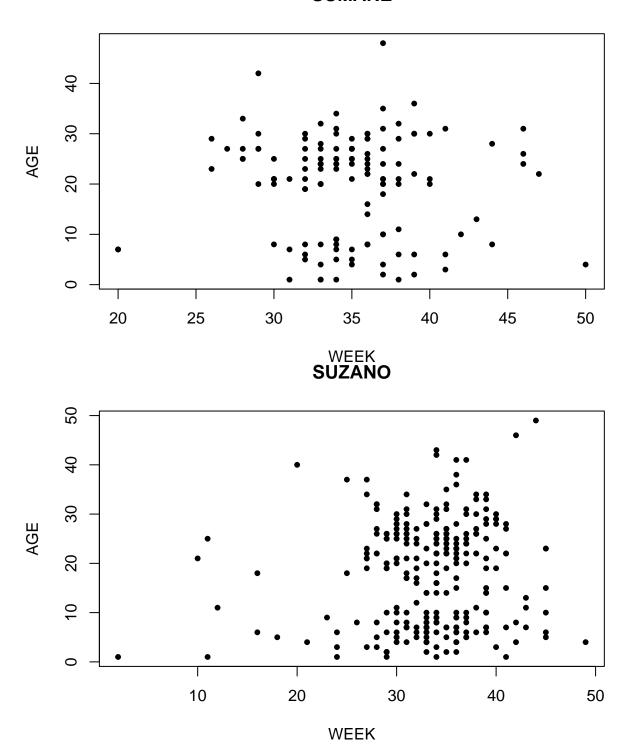
SAO SEBASTIAO



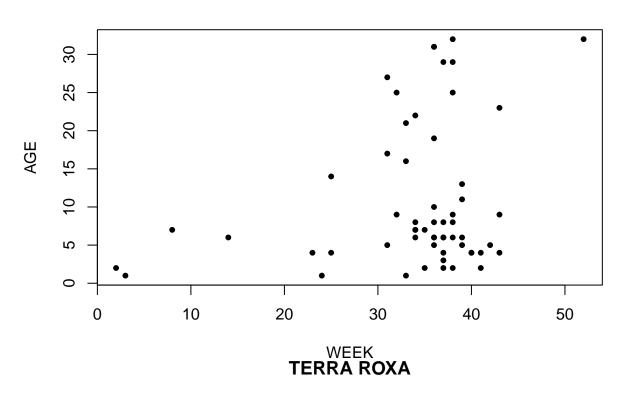
SERTAOZINHO

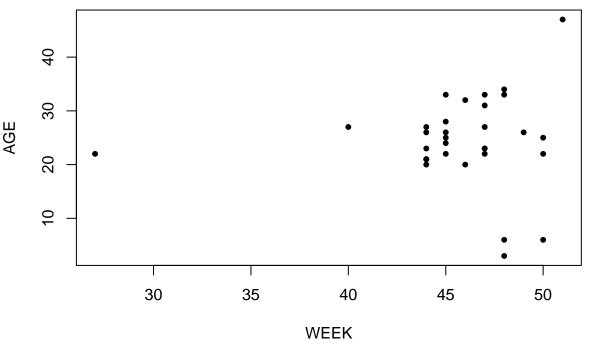


SUMARE

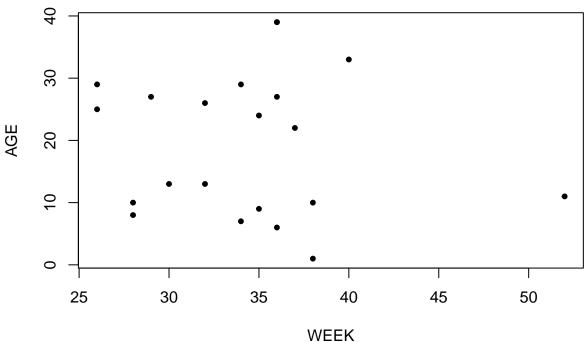


TAUBATE

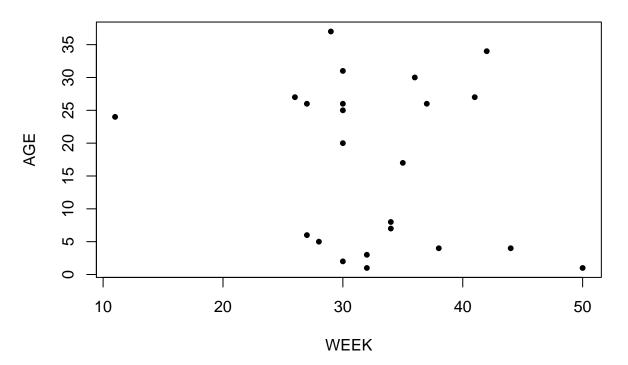




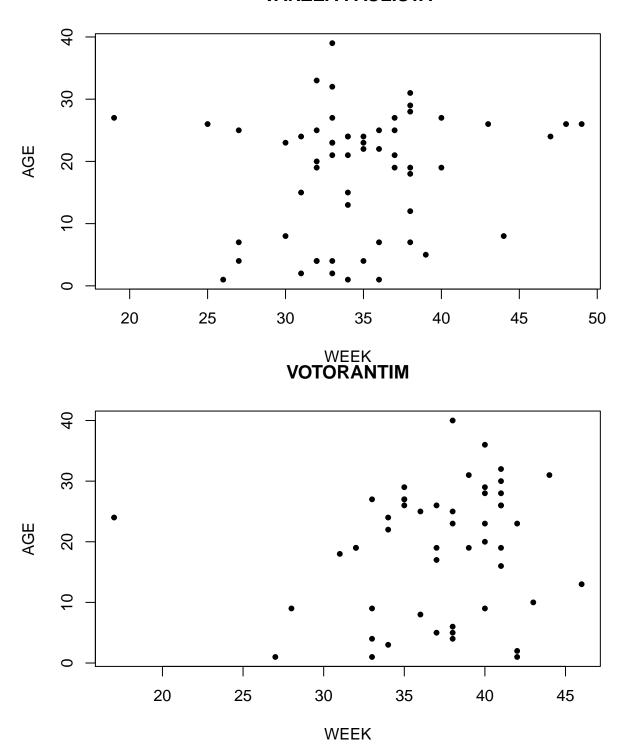
UBATUBA







VARZEA PAULISTA



Preliminary Analysis

Confidence intervals for age group densities

Created two_matrix, three_matrix, four_matrix, and five_matrix which contain the density of cases in age groups of the corresponding span for each county. Density was just calculated as the number of cases in that specific class divided by the number of cases in that county total.

```
#find appropriate age windows, look at age groups with ranges ranging from 2 to 5 years
density_age <- function(max.age, size){</pre>
  amount <- round(max.age / size)</pre>
  amount.vec <- rep(NA, amount)
  amount.vec <- as.data.frame(amount.vec)</pre>
  amount.vec[1,] <- length(which(subset_county$AGE <= size))</pre>
  for(i in 2:amount){
    before <- as.integer(sum(amount.vec[1: i - 1,]))</pre>
    amount.vec[i,] <- length(which(subset_county$AGE <= (size * i))) - before
  density.vec <- amount.vec / length(subset_county$AGE)</pre>
  return(density.vec)
#matrix for age group of 2's
two matrix <- matrix(rep(NA, round(84/2) * length(county vec)), ncol = length(county vec))
two_matrix <- as.data.frame(two_matrix)</pre>
colnames(two_matrix) <- county_vec</pre>
row.names(two_matrix) <- c("<2", "2-4", "4-6", "6-8", "8-10",
                             "10-12", "12-14", "14-16", "16-18", "18-20",
                                     "22-24", "24-26", "26-28", "28-30",
                            "30-32", "32-34", "34-36", "36-38", "38-40",
                            "40-42", "42-44", "44-46", "46-48", "48-50",
                            "50-52", "52-54", "54-56", "56-58", "58-60",
                            "60-62", "62-64", "64-66", "66-68", "68-70",
                            "70-72", "72-74", "74-76", "76-78", "78-80",
                            "80-82", "82-84")
for(j in 1:length(county vec)){
  subset county <- subset(data, data$COUNTY == county vec[j])</pre>
  max.age <- max(subset_county$AGE)</pre>
  density <- unlist(density_age(max.age, 2))</pre>
  length d <- length(density)</pre>
  density <- c(density, rep(NA, (42 - length_d)))
  two_matrix[,j] <- density</pre>
#matrix for age groups of 3's
three_matrix <- matrix(rep(NA, round(84/3) * length(county_vec)), ncol = length(county_vec))
three_matrix <- as.data.frame(three_matrix)</pre>
colnames(three_matrix) <- county_vec</pre>
rownames(three_matrix) <- c("<3", "3-6", "6-9",
                               "9-12", "12-15", "15-18",
                               "18-21", "21-24", "24-27",
                               "27-30", "30-33", "33-36",
                               "36-39", "39-42", "42-45",
                               "45-48", "48-51", "51-54",
```

```
"54-57", "57-60", "60-63",
                                "63-66", "66-69", "69-72",
                                "72-75", "75-78", "78-81",
                                "81-84")
for(j in 1:length(county_vec)){
  subset_county <- subset(data, data$COUNTY == county_vec[j])</pre>
  max.age <- max(subset_county$AGE)</pre>
  density <- unlist(density age(max.age, 3))</pre>
  length_d <- length(density)</pre>
  density <- c(density, rep(NA, (28 - length_d)))</pre>
  three_matrix[,j] <- density</pre>
}
#matrix for age groups of 4's
four_matrix <- matrix(rep(NA, 21 * length(county_vec)), ncol = length(county_vec))</pre>
four_matrix <- as.data.frame(four_matrix)</pre>
colnames(four_matrix) <- county_vec</pre>
row.names(four_matrix) <- c("<4", "4-8", "8-12", "12-16",
                              "16-20", "20-24", "24-28", "28-32",
                              "32-36", "36-40", "40-44", "44-48",
                              "48-52", "52-56", "56-60", "60-64",
                              "64-68", "68-72", "72-76", "76-80",
                              "80-84")
for(j in 1:length(county_vec)){
  subset_county <- subset(data, data$COUNTY == county_vec[j])</pre>
  max.age <- max(subset_county$AGE)</pre>
  density <- unlist(density_age(max.age, 4))</pre>
  length_d <- length(density)</pre>
  density <- c(density, rep(NA, (21 - length_d)))</pre>
  four_matrix[,j] <- density</pre>
}
#matrix for age groups of 5's
five_matrix <- matrix(rep(NA, round(84/5) * length(county_vec)), ncol = length(county_vec))
five matrix <- as.data.frame(five matrix)</pre>
colnames(five_matrix) <- county_vec</pre>
row.names(five_matrix) <- c("<5", "5-10", "10-15", "15-20",
                             "20-25", "25-30", "30-35", "35-40",
                             "40-45", "45-50", "50-55", "55-60",
                             "60-65", "65-70", "70-75", "75-80",
                             "80-85")
for(j in 1:length(county_vec)){
  subset_county <- subset(data, data$COUNTY == county_vec[j])</pre>
  max.age <- max(subset_county$AGE)</pre>
  density <- unlist(density_age(max.age, 5))</pre>
  length_d <- length(density)</pre>
  density <- c(density, rep(NA, (17 - length_d)))</pre>
  five_matrix[,j] <- density</pre>
}
```

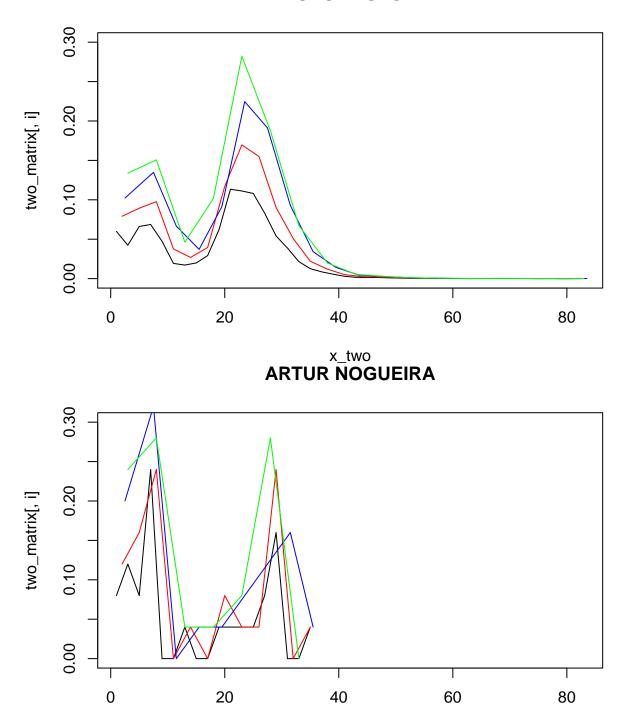
Preliminary Analysis

It appears that the size of the window does have a significant (colloquially) effect on the density in that close age classes have pretty different densities. Going to compare densities on a line graph.

Comparison of windows

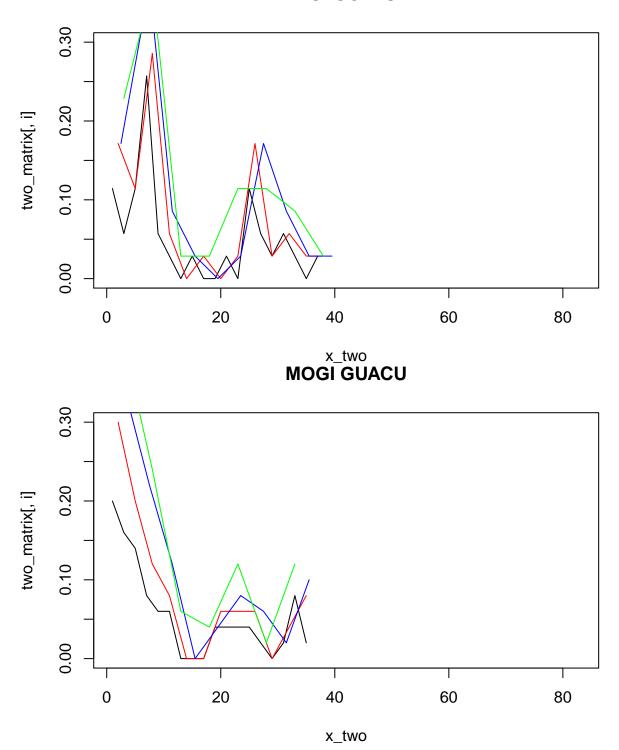
```
#x axis for twos
x_{two} \leftarrow rep(NA, 42)
x_two[1] <- 1
for(i in 2:42){
  x_add \leftarrow seq(1:41)
  x_two[i] \leftarrow i + x_add[i - 1]
#x axis for threes
x_{three} \leftarrow rep(NA, 28)
x_{three}[1] \leftarrow 2
for(i in 2:28){
  x_add \leftarrow seq(1:28)
  x_{three[i]} \leftarrow 2 * i + x_{add[i]} -1
#x axis for fours
x_{four} \leftarrow rep(NA, 21)
x_four[1] <- 2.5
for(i in 2:21){
  x_add <- seq(1:21)
  x_{four}[i] \leftarrow 3 * i + x_{add}[i] + 0.5 -1
}
#x axis for fives
x_{five} \leftarrow rep(NA, 17)
x_five[1] \leftarrow 3
for(i in 2:17){
  x_add \leftarrow seq(1:17)
  x_{five}[i] \leftarrow 4 * i + x_{add}[i] -2
for(i in 1:length(county_vec)){
  subset_county <- subset(data, data$COUNTY == county_vec[i])</pre>
  if(length(subset_county$AGE) > 20){
    {plot(x_two, two_matrix[,i], type = "l", ylim = c(0, 0.3), main = county_vec[i])}
     lines(x_three, three_matrix[,i], col = "red")
      lines(x_four, four_matrix[,i], col = "blue")
      lines(x_five, five_matrix[,i], col = "green")}
}
```

SAO PAULO

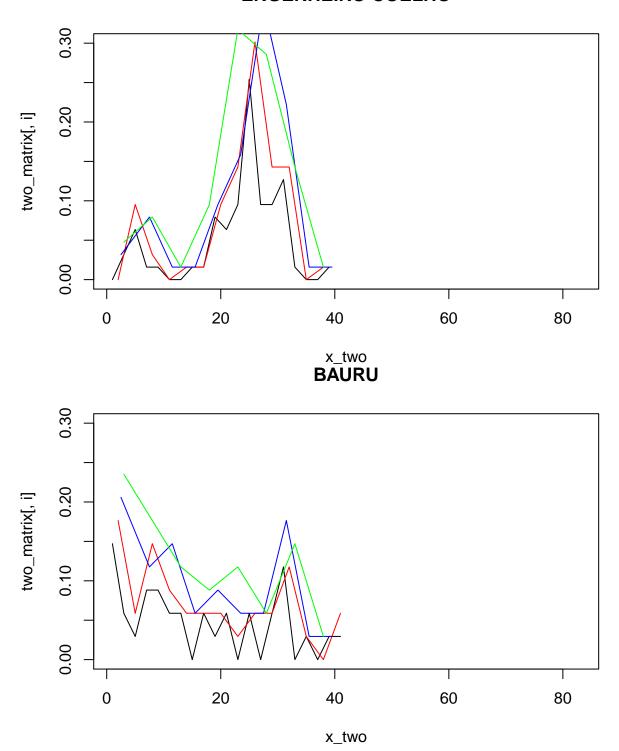


x_two

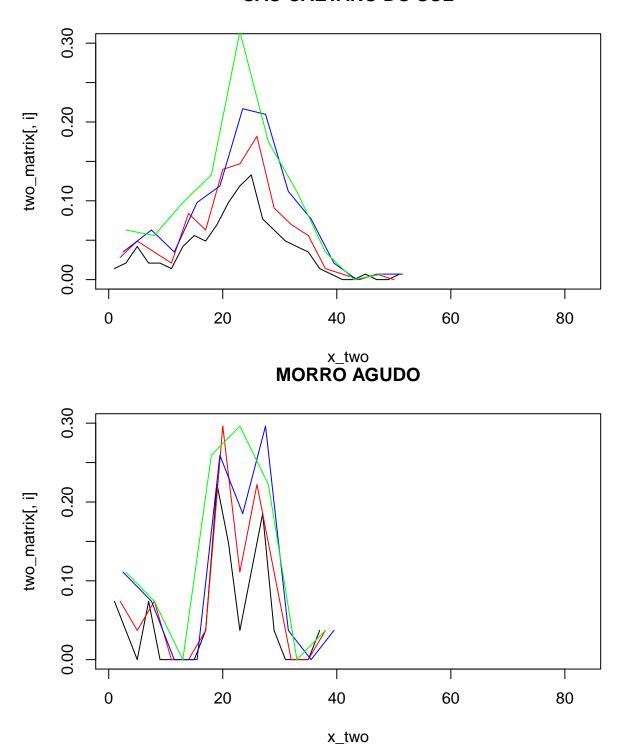
BOTUCATU



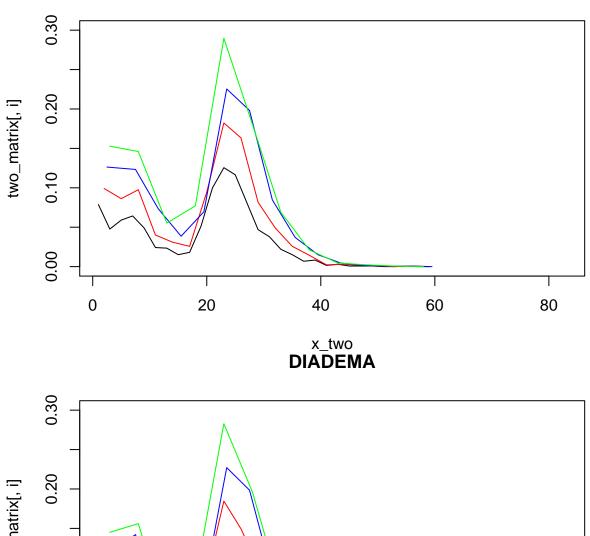
ENGENHEIRO COELHO

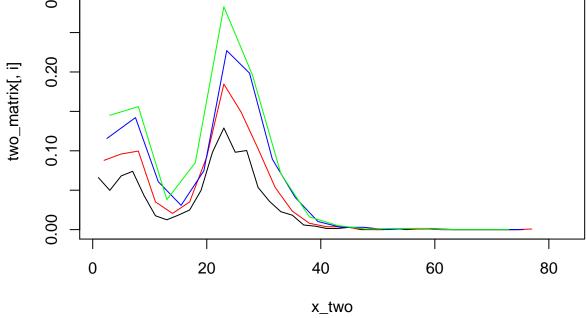


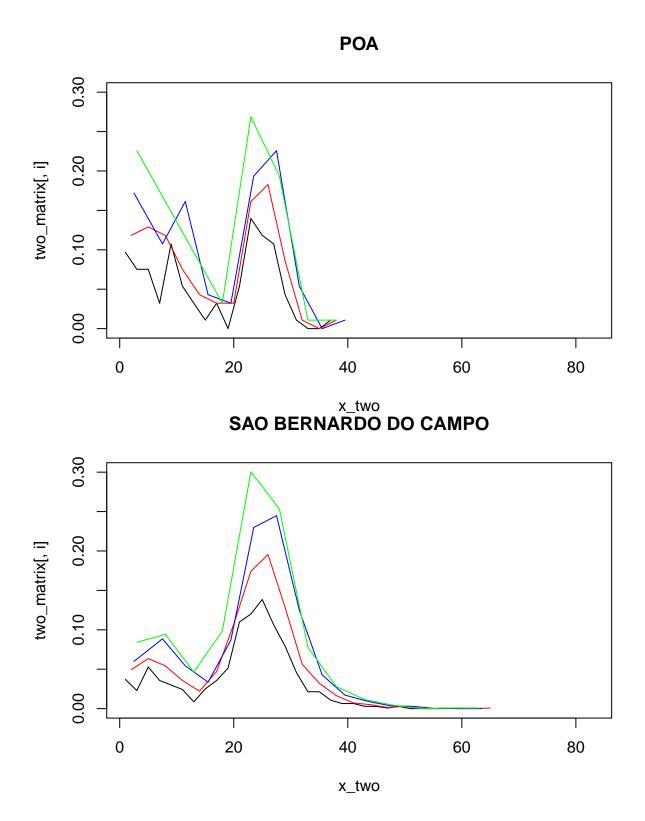
SAO CAETANO DO SUL



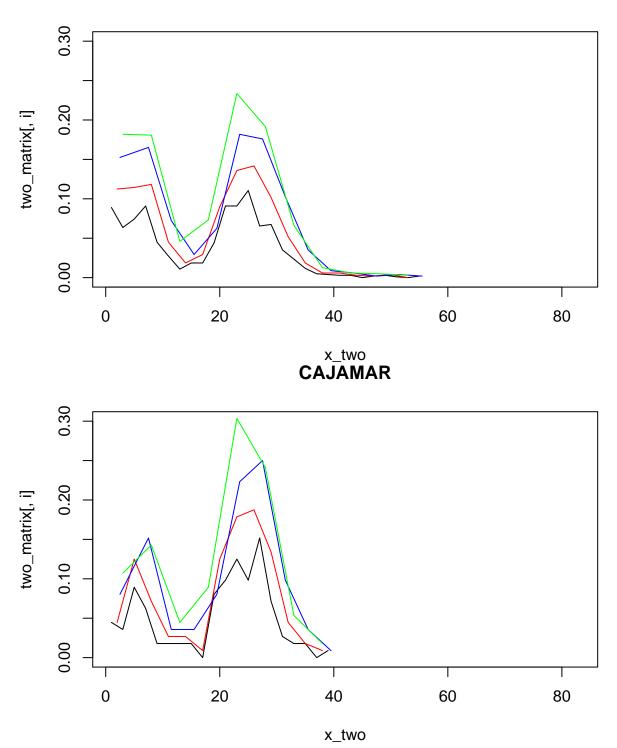




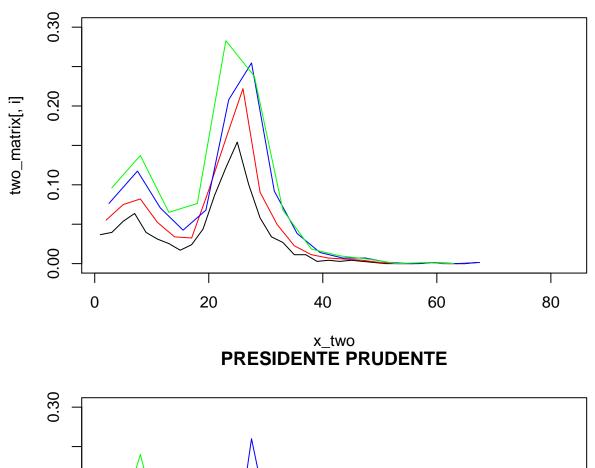


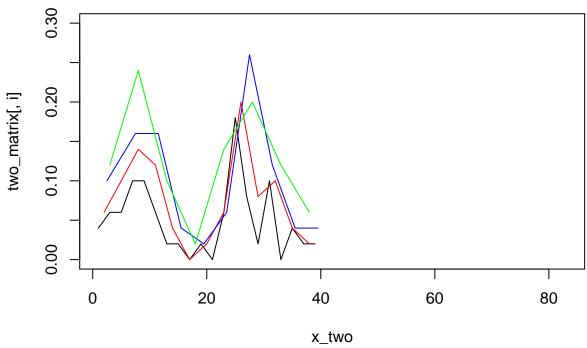


GUARULHOS

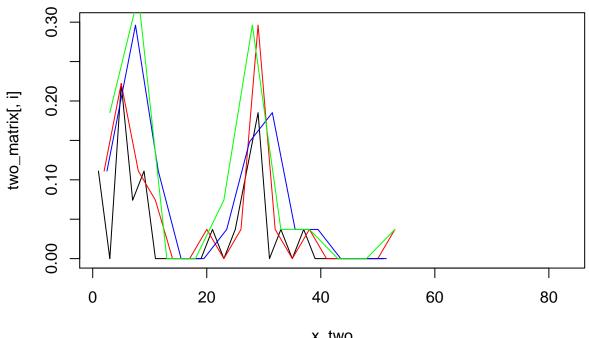


SANTO ANDRE

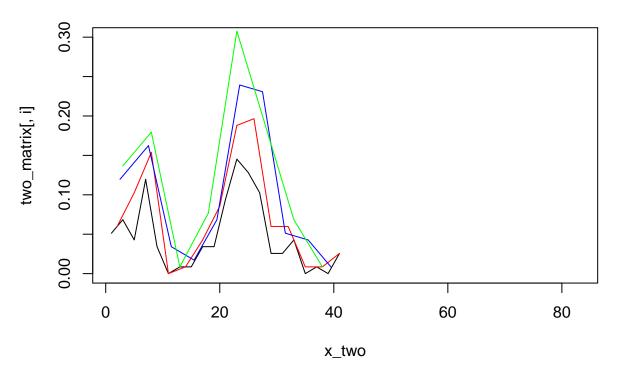




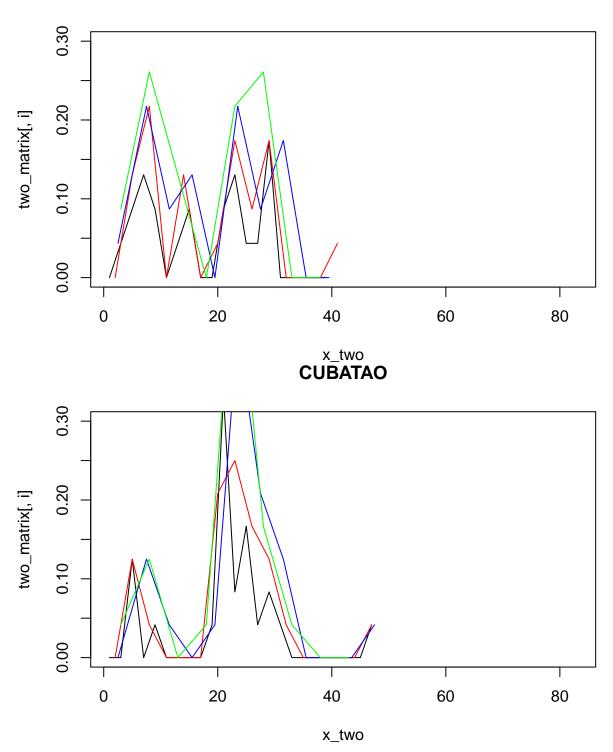
MARILIA



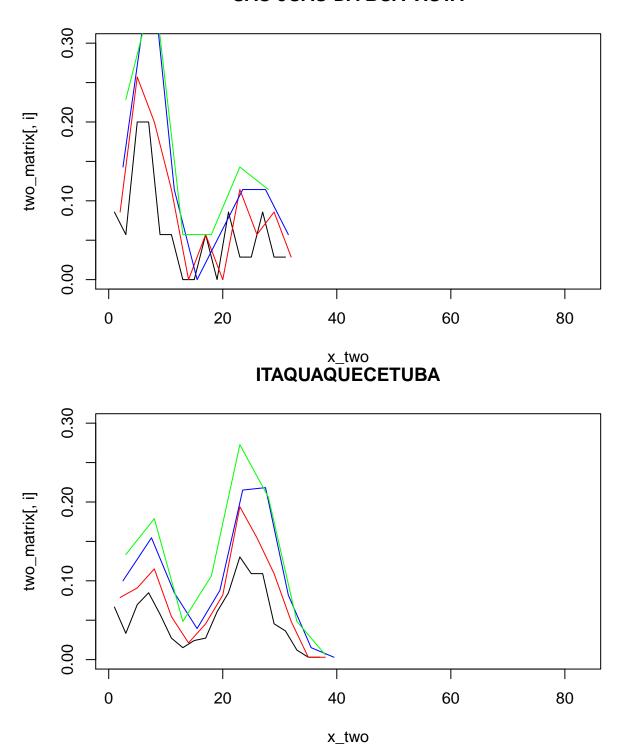
FERRAZ DE VASCONCELOS



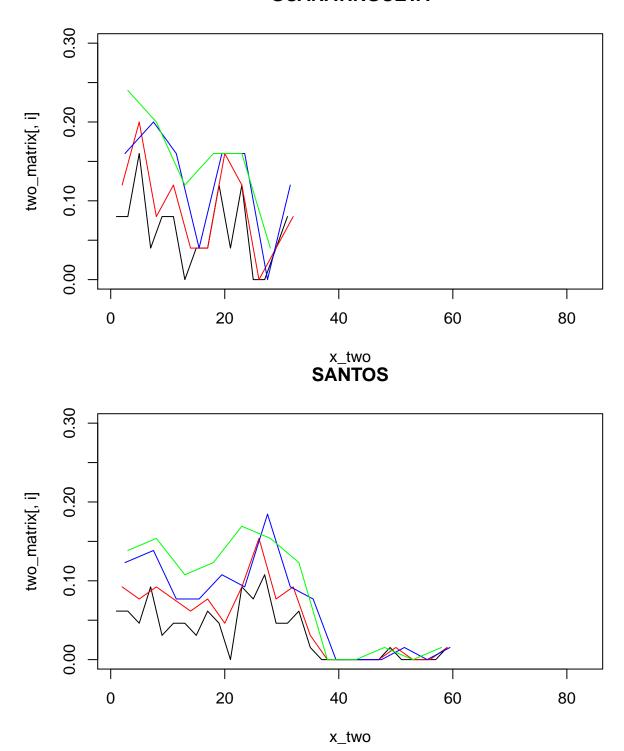




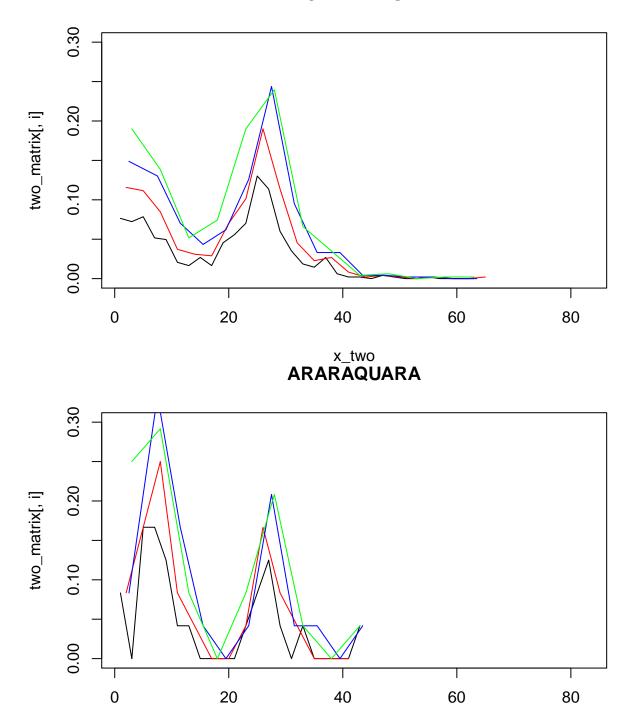
SAO JOAO DA BOA VISTA



GUARATINGUETA

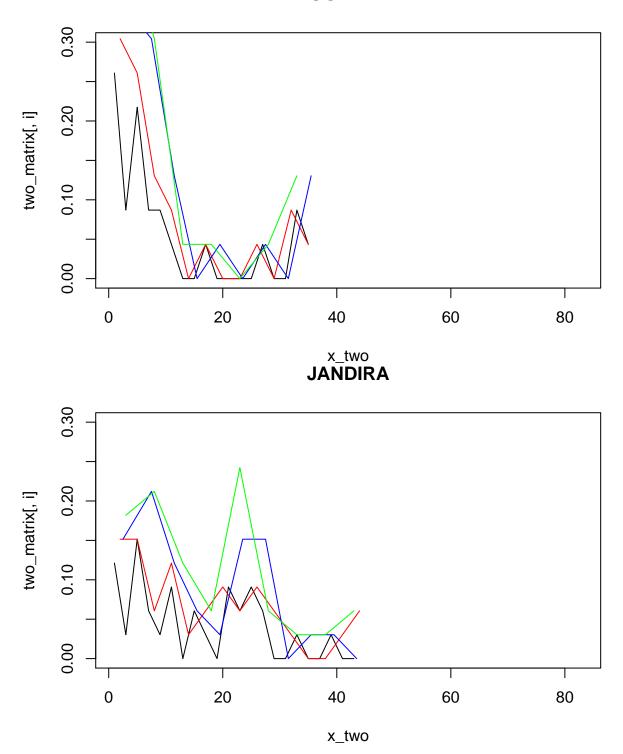


CAMPINAS

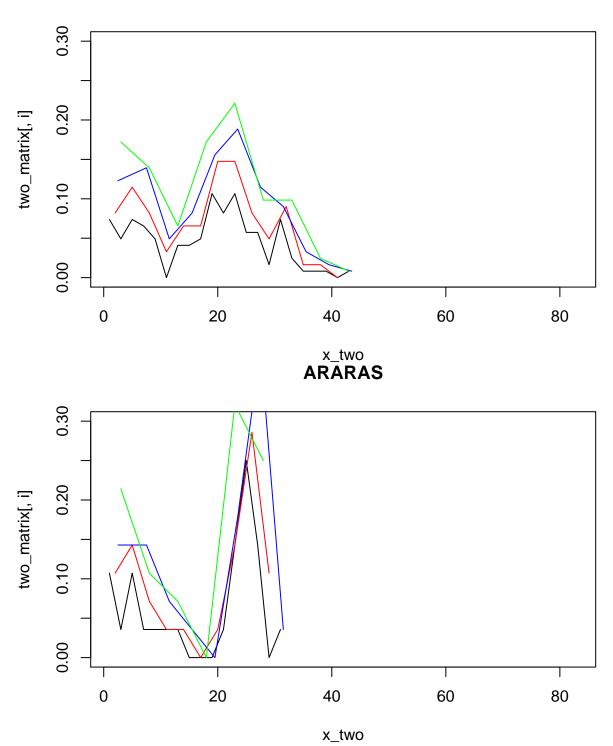


x_two

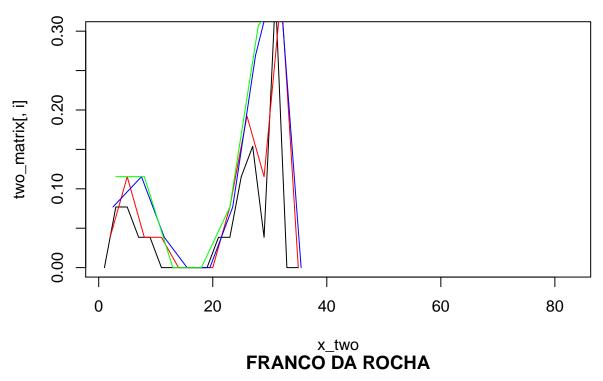
MOGI MIRIM

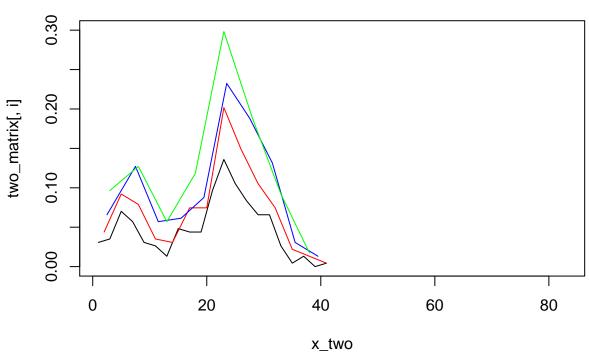




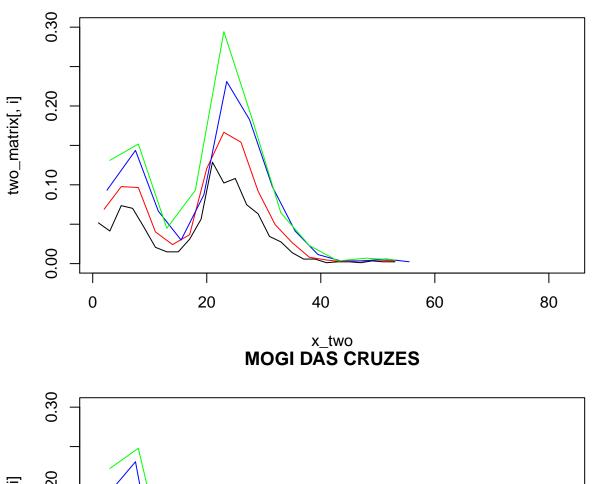


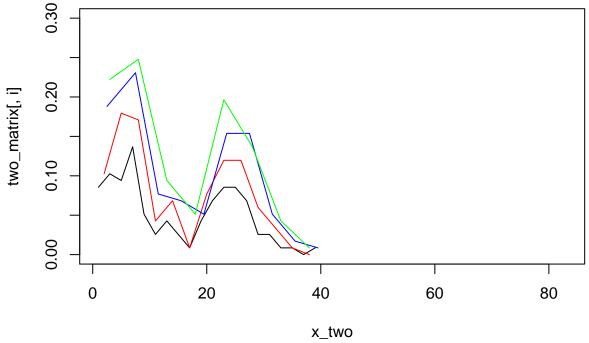
PINDAMONHANGABA



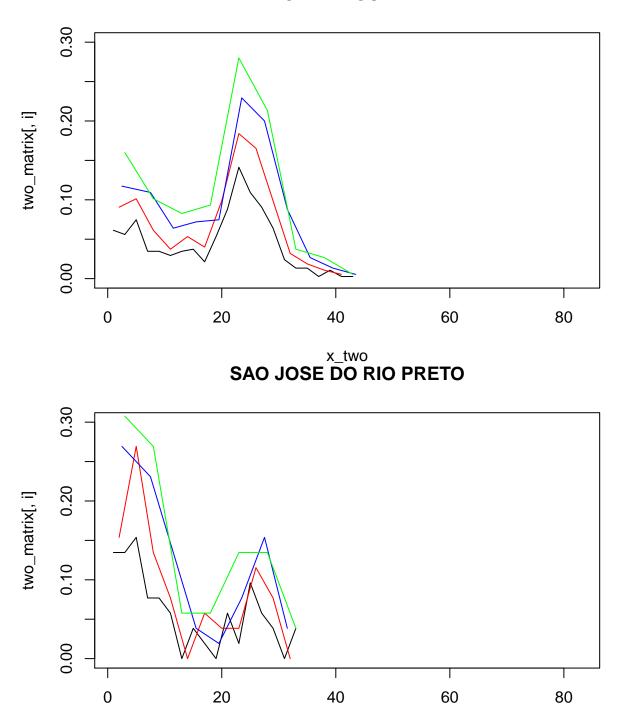






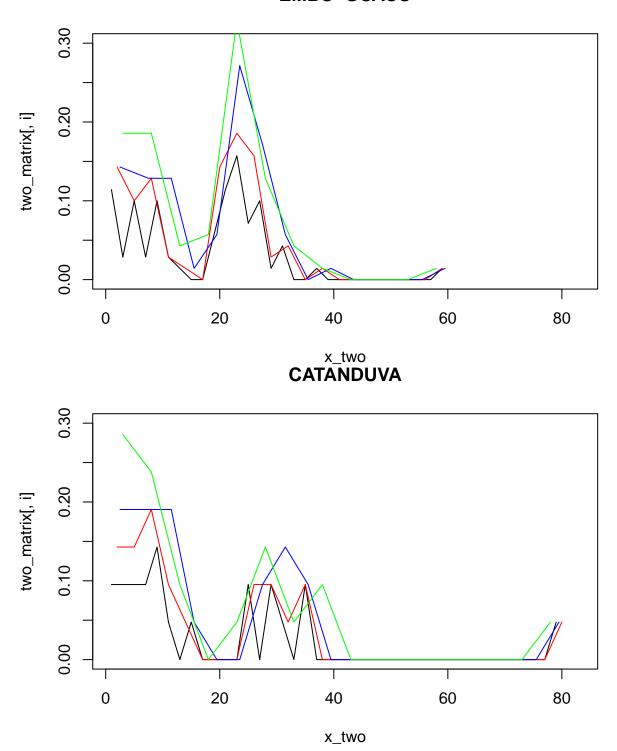


CARAPICUIBA

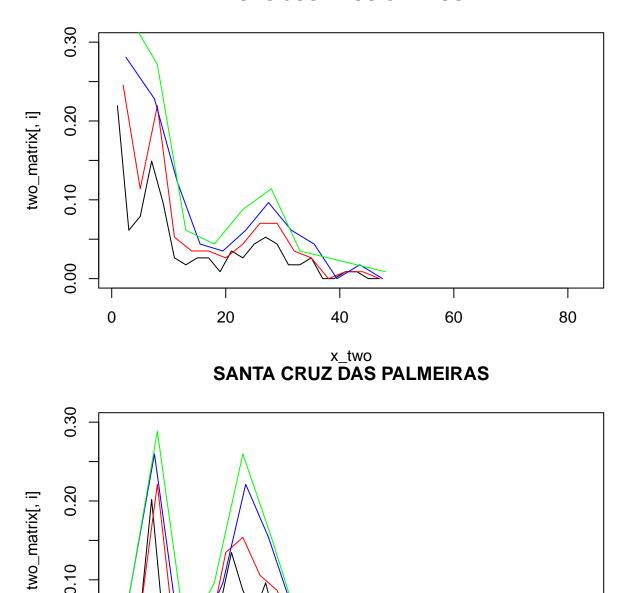


x_two

EMBU-GUACU



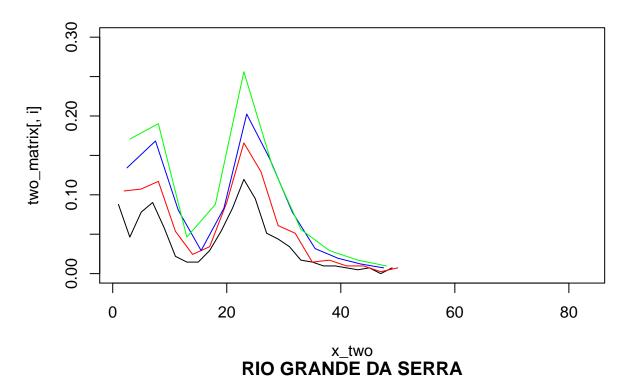
SAO JOSE DOS CAMPOS

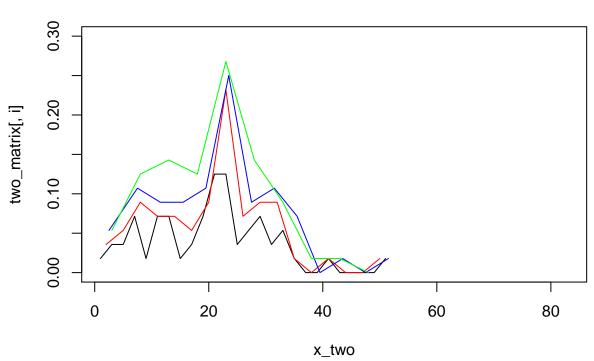


x_two

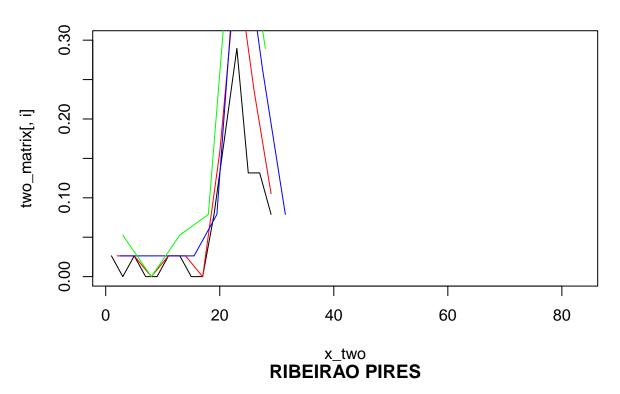
0.00

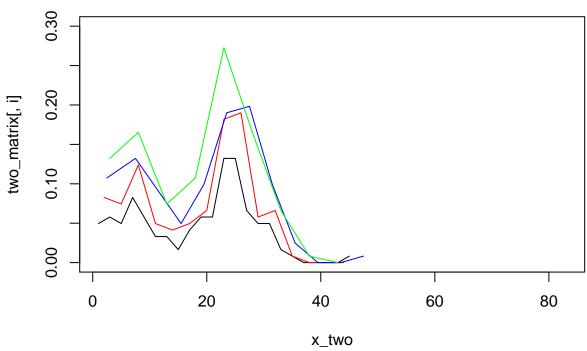
EMBU DAS ARTES



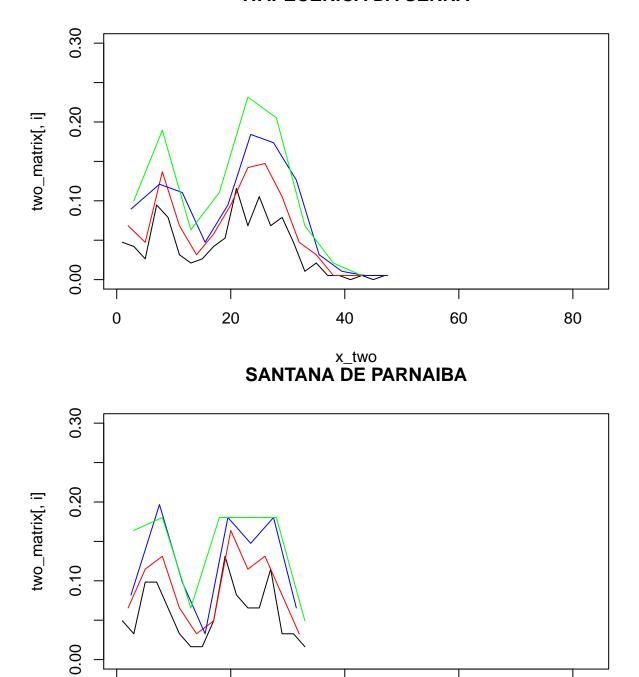


PONTAL



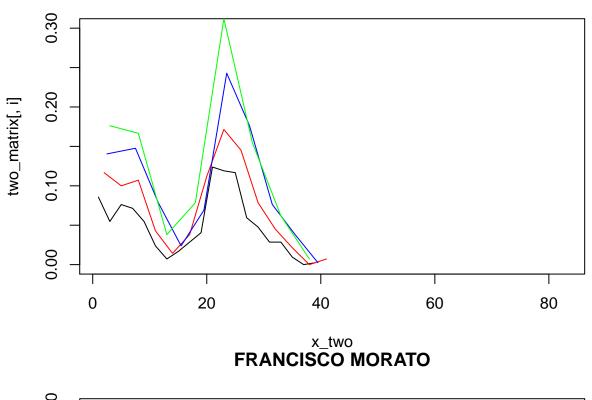


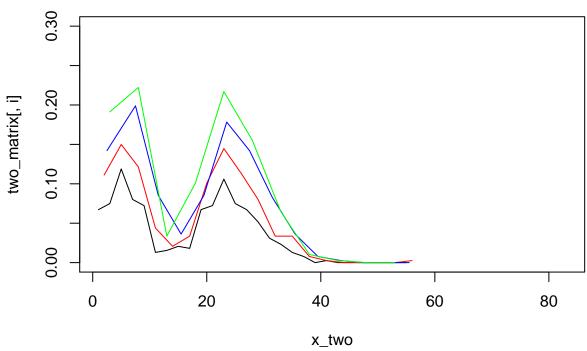
ITAPECERICA DA SERRA



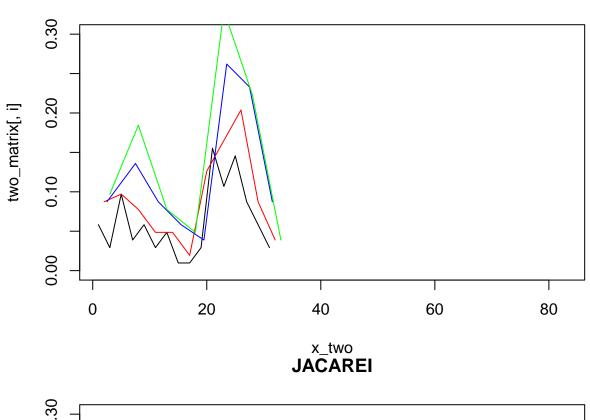
x_two

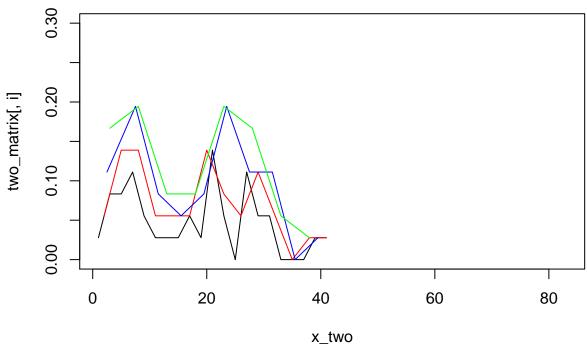
BARUERI



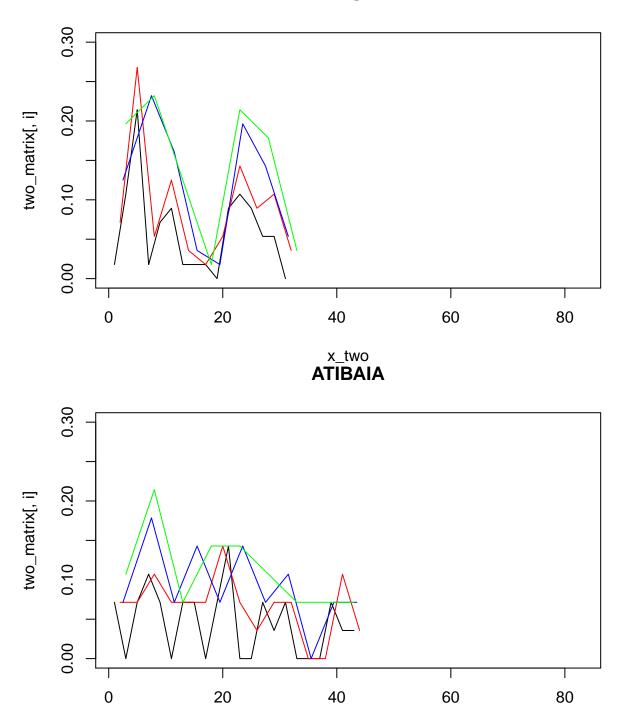


CAIEIRAS



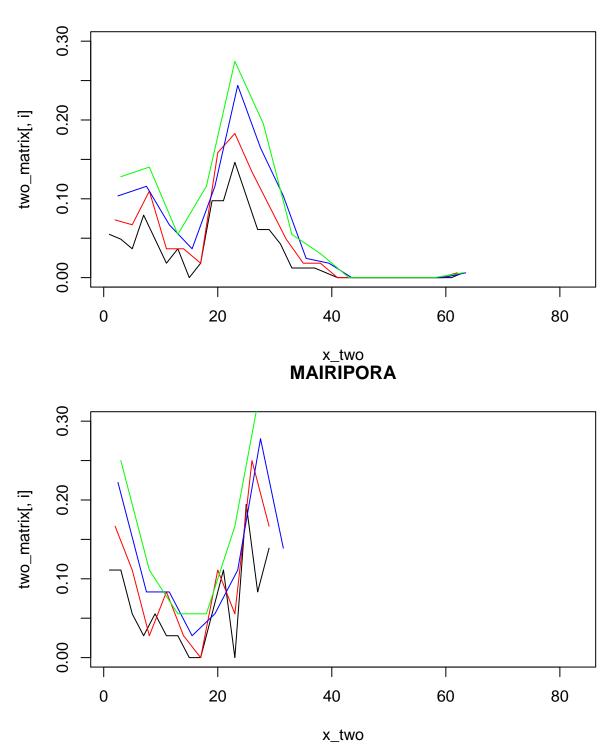




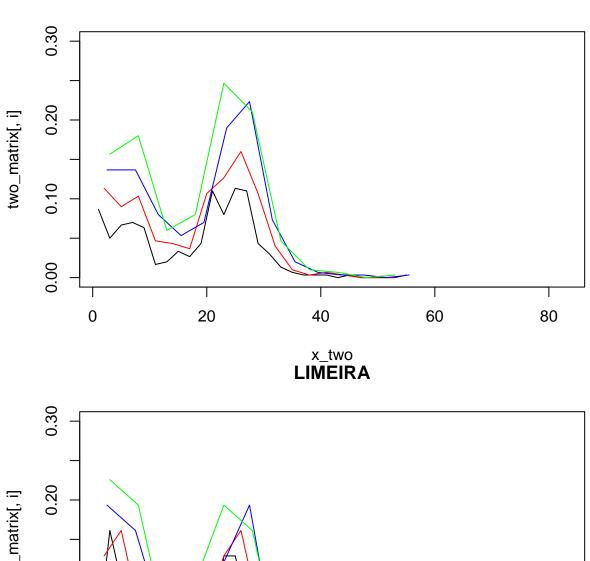


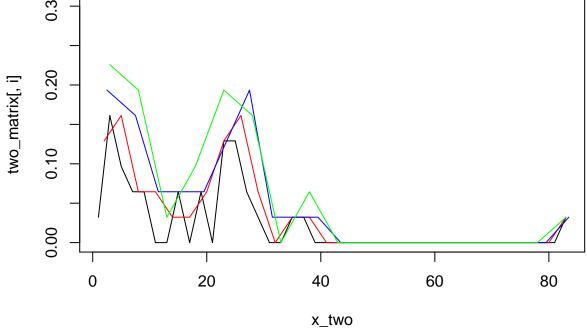
x_two



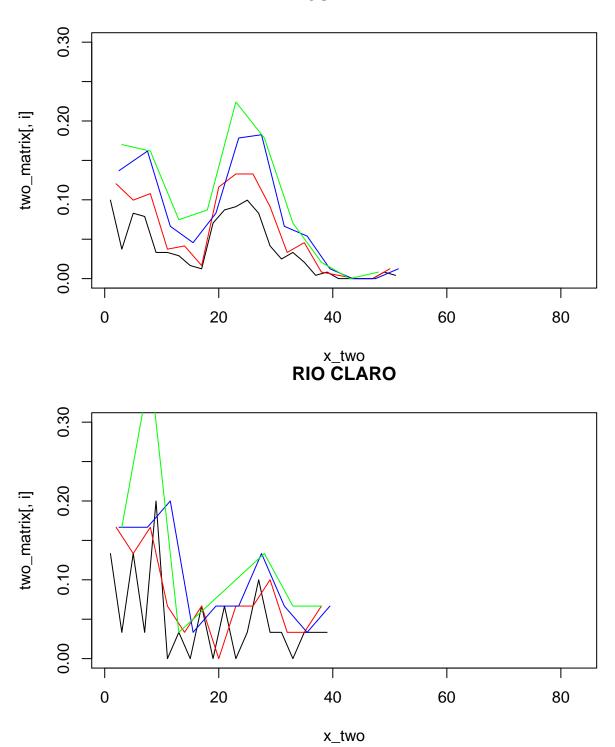


RIBEIRAO PRETO

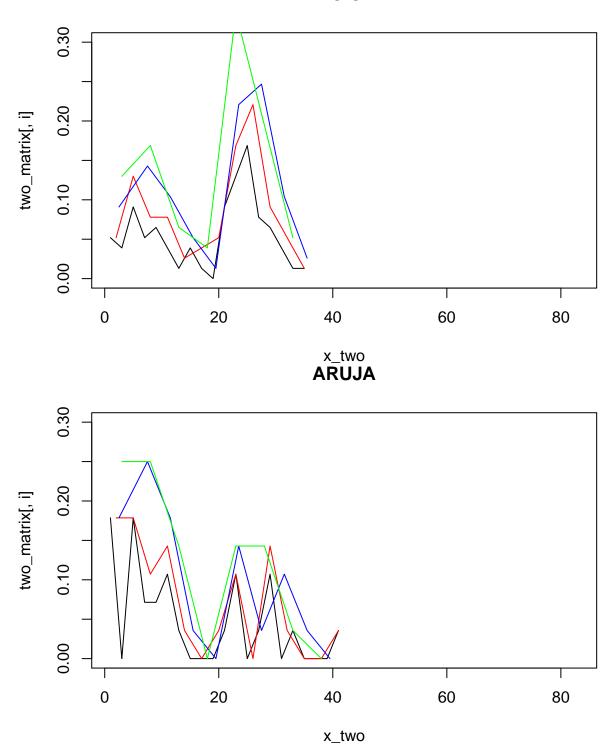




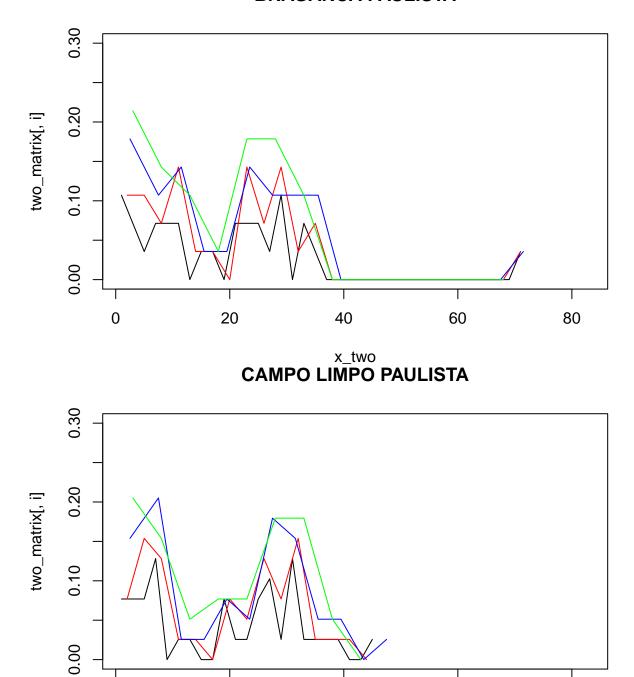
JUNDIAI



PIRACICABA

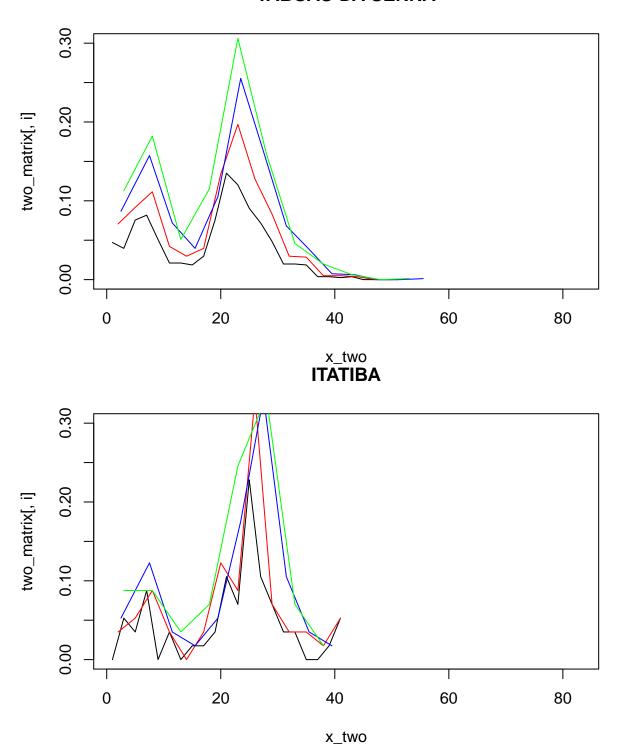


BRAGANCA PAULISTA

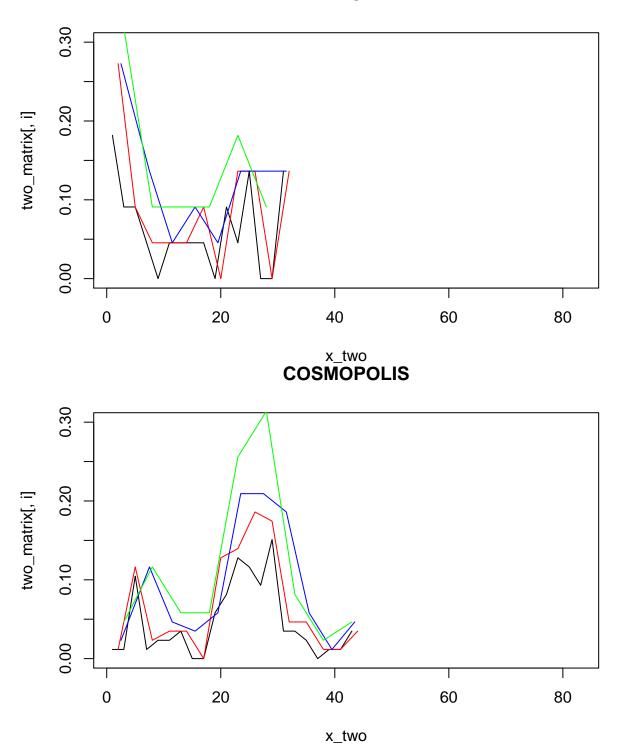


x_two

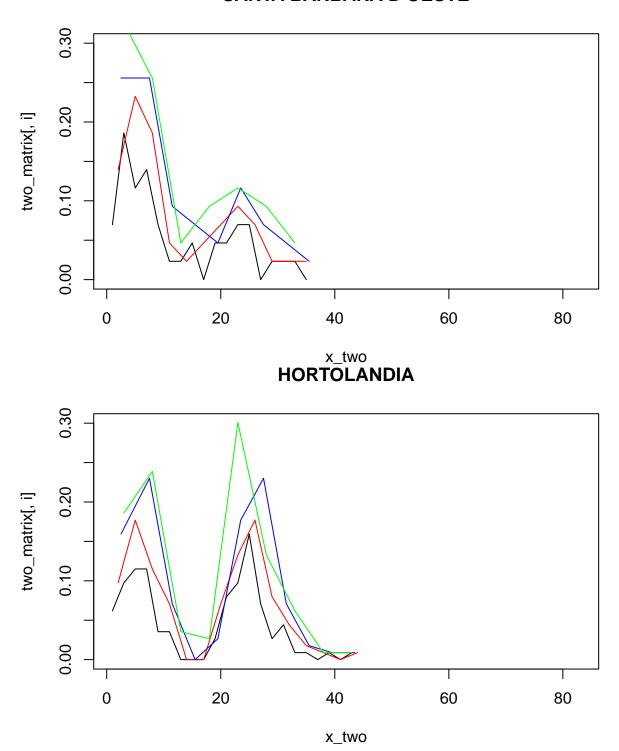
TABOAO DA SERRA



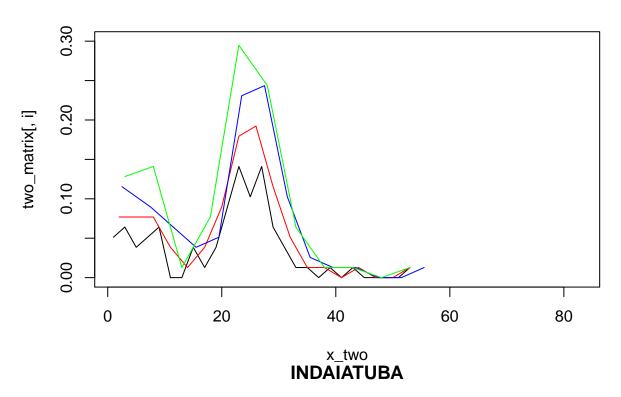
AMERICANA

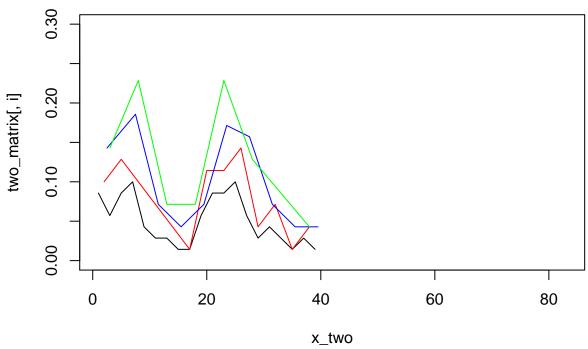


SANTA BARBARA D'OESTE

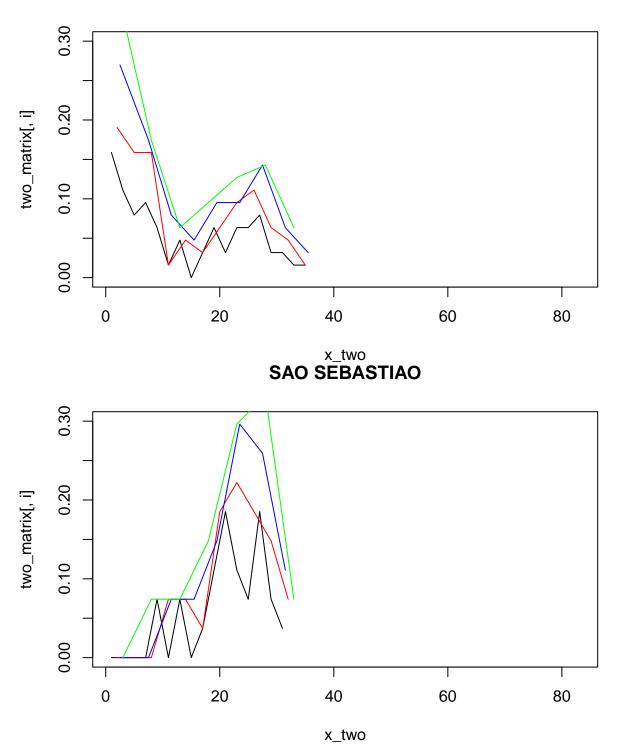


PAULINIA

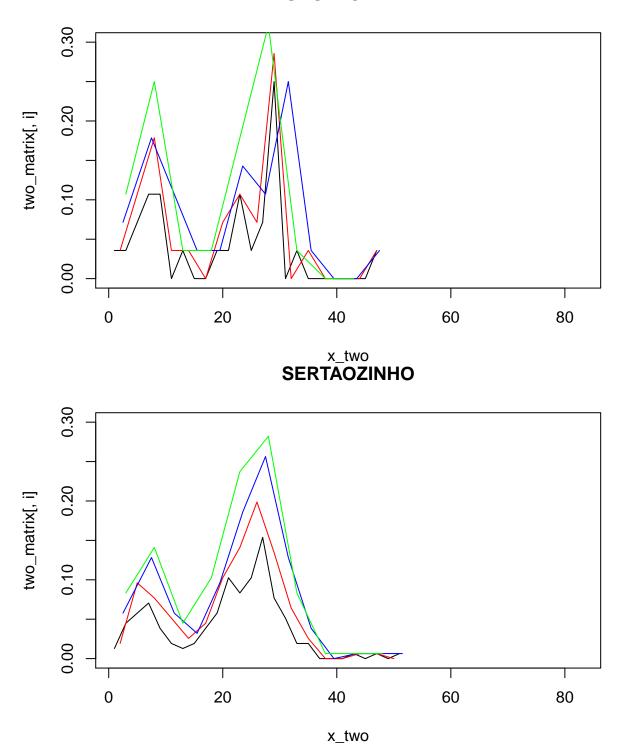




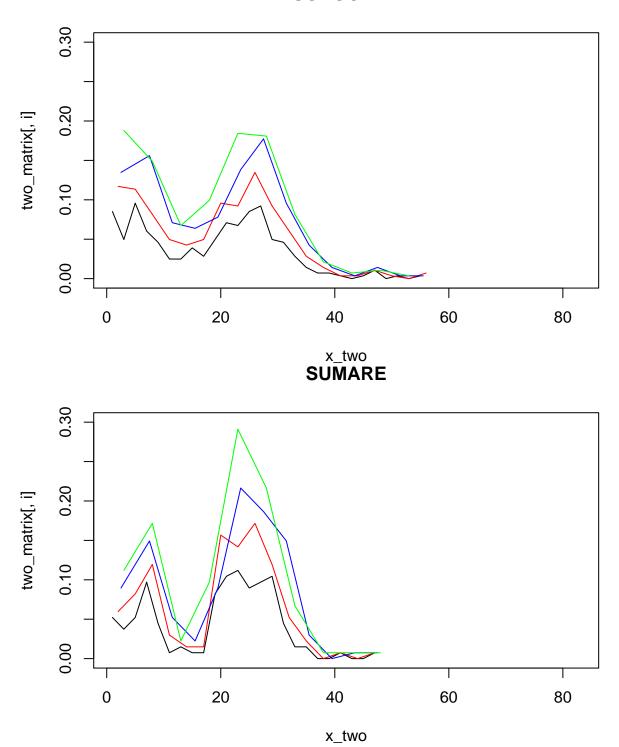




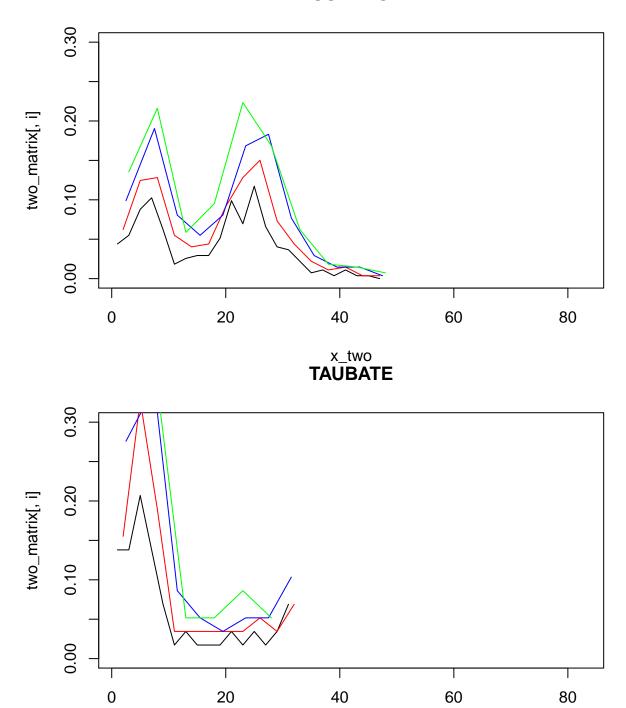
SAO VICENTE



SOROCABA

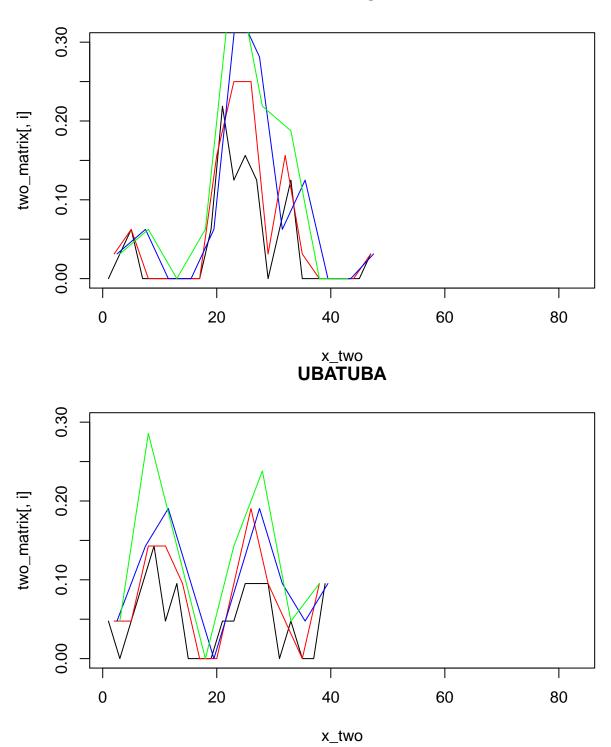


SUZANO

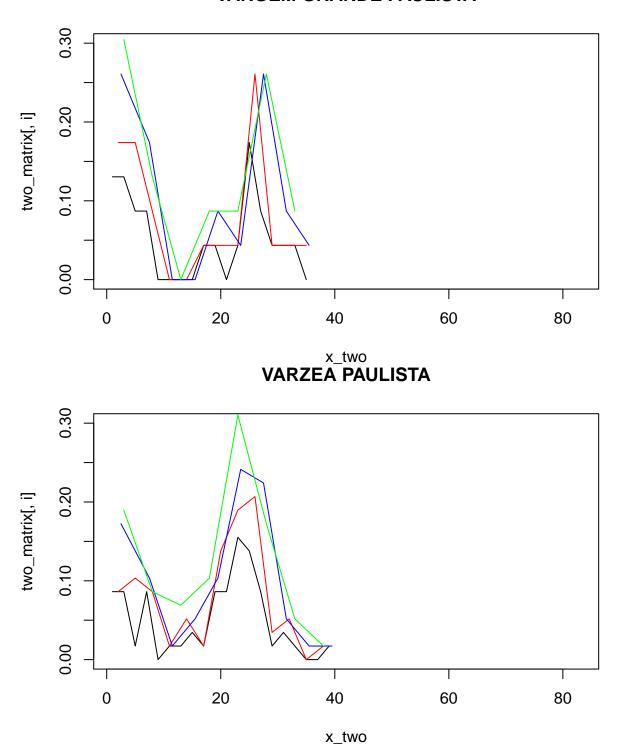


x_two

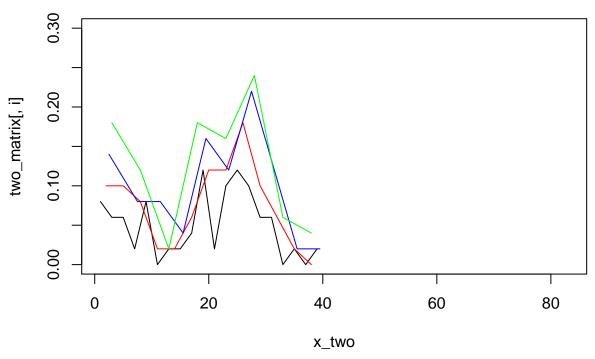
TERRA ROXA



VARGEM GRANDE PAULISTA



VOTORANTIM



```
#two
mean_two <- rep(NA, 42)</pre>
sd_two <- rep(NA, 42)</pre>
for(i in 1:42){
  mean_two[i] <- rowMeans(two_matrix[i,], na.rm = TRUE)</pre>
  sd_two[i] <- apply(two_matrix, 1, sd, na.rm = TRUE)[i]</pre>
}
#three
mean_three <- rep(NA, 28)
sd_three <- rep(NA, 28)
for(i in 1:28){
  mean_three[i] <- rowMeans(three_matrix[i,], na.rm = TRUE)</pre>
  sd_three[i] <- apply(three_matrix, 1, sd, na.rm = TRUE)[i]</pre>
}
#four
mean_four <- rep(NA, 21)</pre>
sd_four <- rep(NA, 21)</pre>
for(i in 1:21){
  mean_four[i] <- rowMeans(four_matrix[i,], na.rm = TRUE)</pre>
  sd_four[i] <- apply(four_matrix, 1, sd, na.rm = TRUE)[i]</pre>
}
#five
mean_five <- rep(NA, 17)</pre>
sd_five <- rep(NA, 17)</pre>
for(i in 1:17){
  mean_five[i] <- rowMeans(five_matrix[i,], na.rm = TRUE)</pre>
```

```
sd_five[i] <- apply(five_matrix, 1, sd, na.rm = TRUE)[i]</pre>
}
\#RMSP = 1
urban <- subset(data, data$RMSP == "1")</pre>
urban_counties <- as.character(unique(urban$COUNTY))</pre>
which.urban <- rep(NA, length(urban counties))
for(i in 1:length(urban counties)){
which.urban[i] <- which(county_vec[] == urban_counties[i])</pre>
}
two_urban_mat <- two_matrix[which.urban]</pre>
three_urban_mat <- three_matrix[which.urban]</pre>
four_urban_mat <- three_matrix[which.urban]</pre>
five_urban_mat <- five_matrix[which.urban]</pre>
\#RMSP = 0
rural <- subset(data, data$RMSP == "0")</pre>
rural_counties <- as.character(unique(rural$COUNTY))</pre>
which.rural <- rep(NA, length(rural_counties))</pre>
for(i in 1:length(rural_counties)){
  which.rural[i] <- which(county_vec[] == rural_counties[i])</pre>
two_rural_mat <- two_matrix[which.rural]</pre>
two_rural <- matrix(rep(NA, 3 * 42 * length(rural_counties)), ncol = 3)</pre>
colnames(two_rural) <- c("COUNTY", "CLASS", "DENSITY")</pre>
two_rural <- as.data.frame(two_rural)</pre>
#fill counties
two_rural[,1] <- rep(rural_counties, 42)</pre>
#fill age class
two_rural[seq(1:length(rural_counties)),2] <- x_two[1]</pre>
for(i in 1:length(x_two)){
  x <- length(rural_counties) * i + seq(1:length(rural_counties))</pre>
  two_rural[x,2] <- x_two[i+1]</pre>
}
#fill density
for(i in 1:length(rural_counties)){
  two_rural[i,3] <- two_rural_mat[1,i]</pre>
for(i in 1:length(rural_counties)){
  for(j in 2:length(x_two)){
    two_rural[(i + 348 * (j - 1) ),3] <- two_rural_mat[j,i]</pre>
  }
  }
two rural$CLASS <- as.factor(two rural$CLASS)</pre>
two_rural$COUNTY <- as.factor(two_rural$COUNTY)</pre>
three_rural_mat <- three_matrix[which.rural]</pre>
four_rural_mat <- three_matrix[which.rural]</pre>
```

five_rural_mat <- five_matrix[which.rural]</pre>

Preliminary analysis

Created a method (via for loops) to make a matrix which is compatible with ANOVAs.

Up next:

- Look for correlation or association between density and class on both a rural and urban level.
 - This may be best to do with the five_rural_mat and five_urban_mat because they have fewer factor levels. Starting with urban may also be more promising because it will have fewer factors for county.
- Look into whether the moving windows/data smoothing can be used for the windows of age groups. Try to understand how this works.
- Plot row means with row sd error bars for the different windows.