

Effects of urban green infrastructure on supply provision of ecosystem service: Temperature reduction

Irina Lerner

30/01/24

Contents

Aim and scope	1
Data	1
Exploratory Analysis	1
Response	1
Predictors	4

Aim and scope

CLEAR AIM: TO SHOW THE EFFECT OF VEGETATION CONFIGURATION IN MEAN TEMPERATURE.

Here we show results, flow and maybe missing steps

Data

DATA NEEDS URGENT CARE: - SEE IF ITS BETTER USING VI PROPORTIONS WITH OR WITHOUT NA. PROBLY WITH NA IF WE ARE CONSIDERING OTHER VARIABLES - PROP_WATER AND PROP_GRASS THAT ARE NOT CORRECTLY CALCULATED FROM THE MAPS

Exploratory Analysis

Response

We begin exploring the response data.

```
analyse_model(data, lm(T_mean ~ 1, data))
```

```

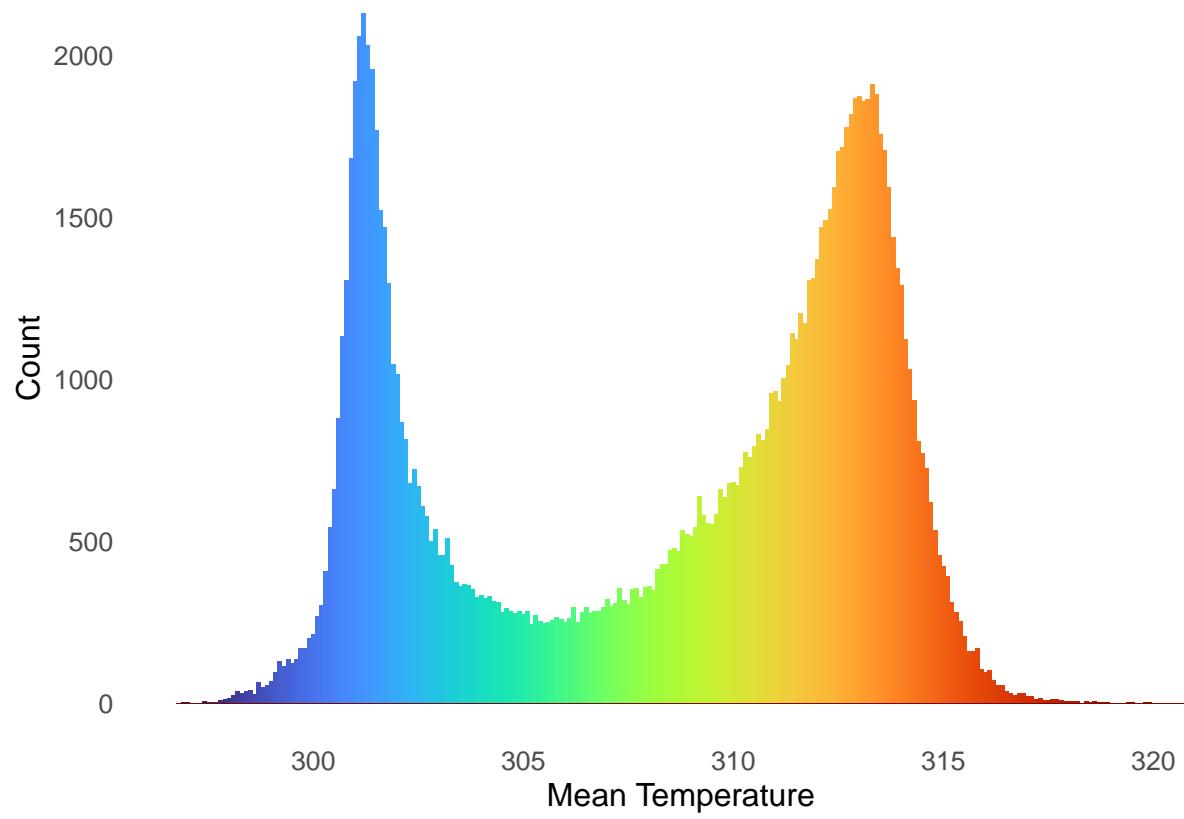
## [1] "Residuals not normally distributed:"
## [1] -0.2141189
##
## Moran I test under randomisation
##
## data: residuals(model)
## weights: listw
##
## Moran I statistic standard deviate = 736.58, p-value < 2.2e-16
## alternative hypothesis: greater
## sample estimates:
## Moran I statistic      Expectation      Variance
##         9.653109e-01     -6.748504e-06     1.717531e-06

## [1] 898590.2

ggplot(cdata, aes(x = T_mean)) +
  geom_histogram(aes(fill = as.factor(cut_width(T_mean, 0.1))), binwidth = 0.1, color = NA) +
  scale_fill_viridis_d(option = "turbo", guide = FALSE) +
  labs(title = "Distribution of T_mean by Bin Order", x = "Mean Temperature", y = "Count") +
  custom_theme

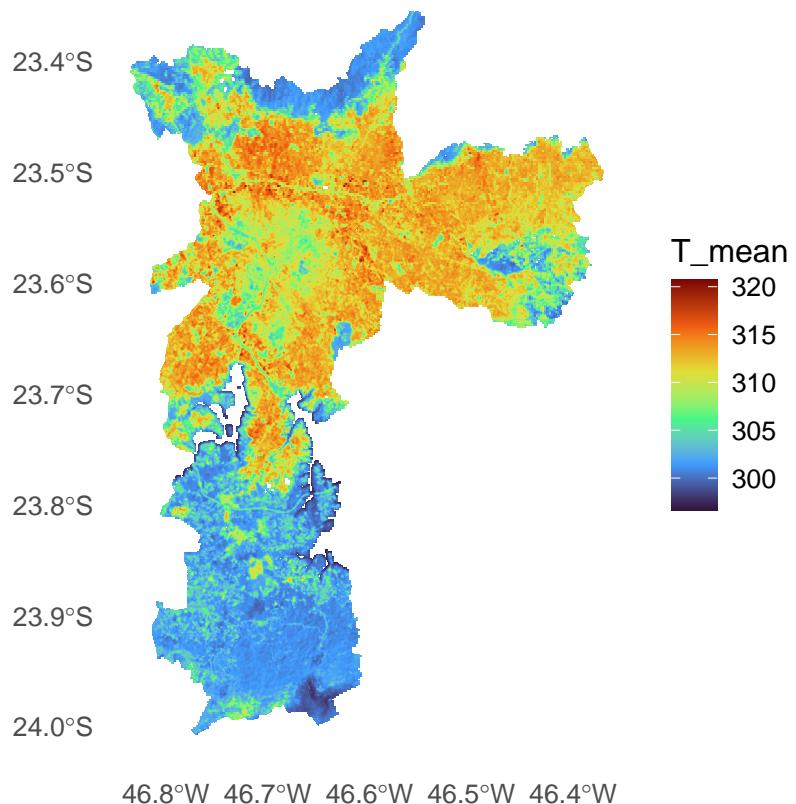
## Warning: The 'guide' argument in 'scale_()' cannot be 'FALSE'. This was deprecated in
## ggplot2 3.3.4.
## i Please use "none" instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.

```



From the distribution of DT in this grid comparison, the values are not normally distributed, instead, they have two peaks.

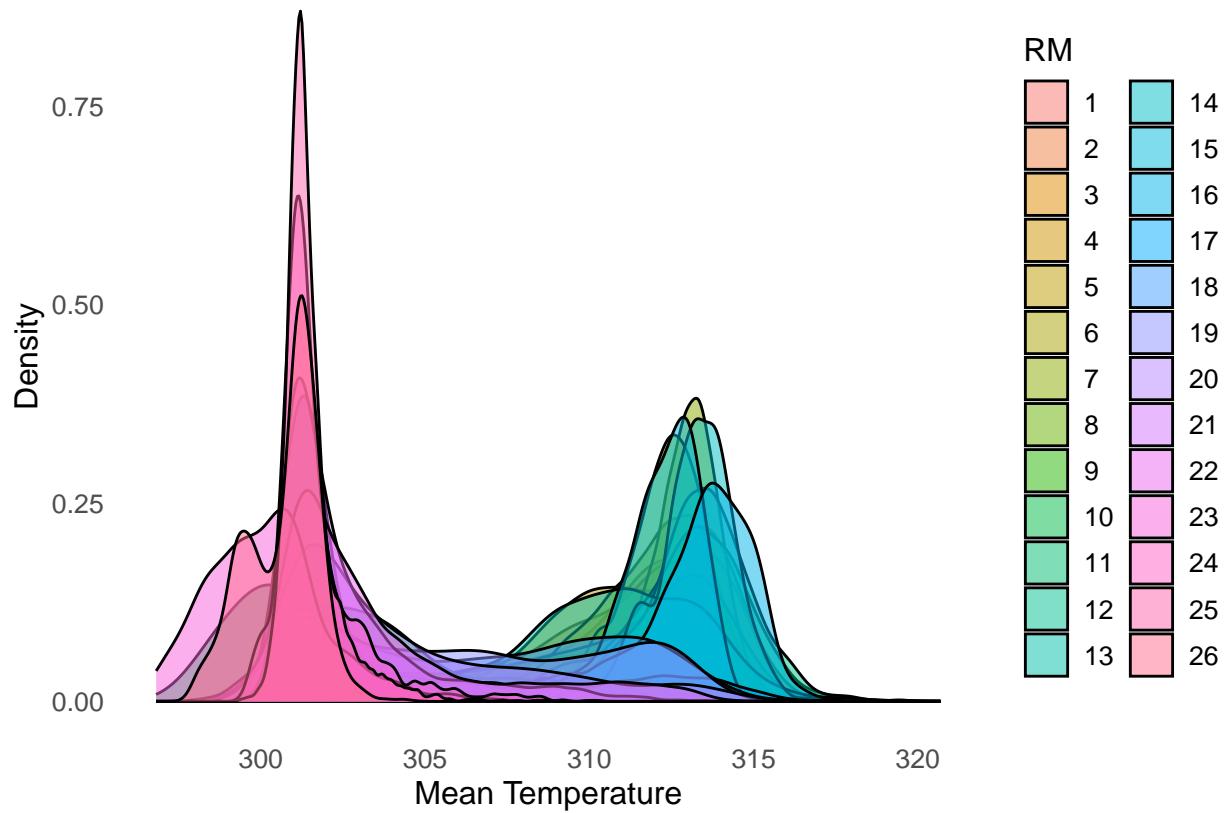
```
ggplot(data) +  
  geom_sf(aes(fill = T_mean), color = NA) + # Correct usage of color setting  
  scale_fill_viridis_c(option = "turbo") +  
  custom_theme
```



Predictors

This two peaks can be seen when we take the distribution of each micro climatic region

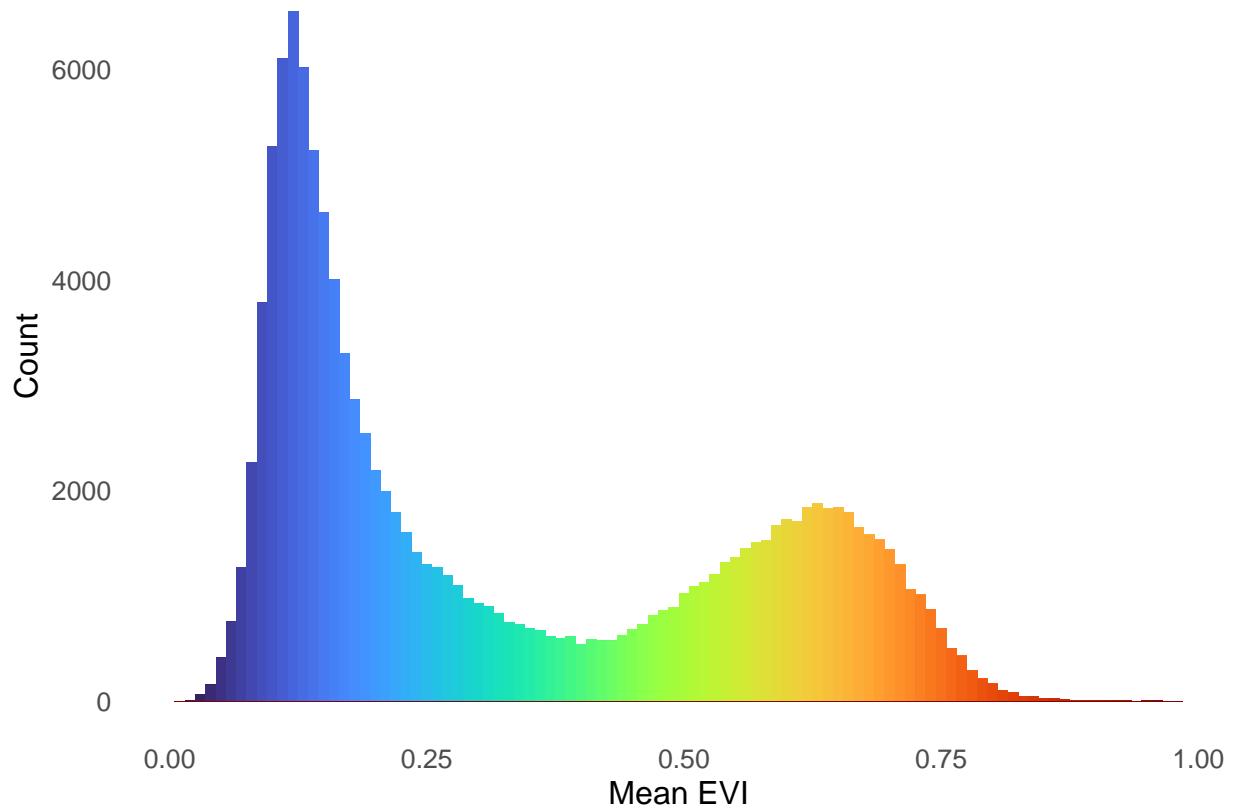
```
ggplot(cdata, aes(x = T_mean, fill = factor(RM))) +
  geom_density(alpha = 0.5) +
  labs(x = "Mean Temperature", y = "Density", fill = "RM") +
  custom_theme
```



we can either normalize by micro climatic region or consider it as a random effect in the distribution parameters.

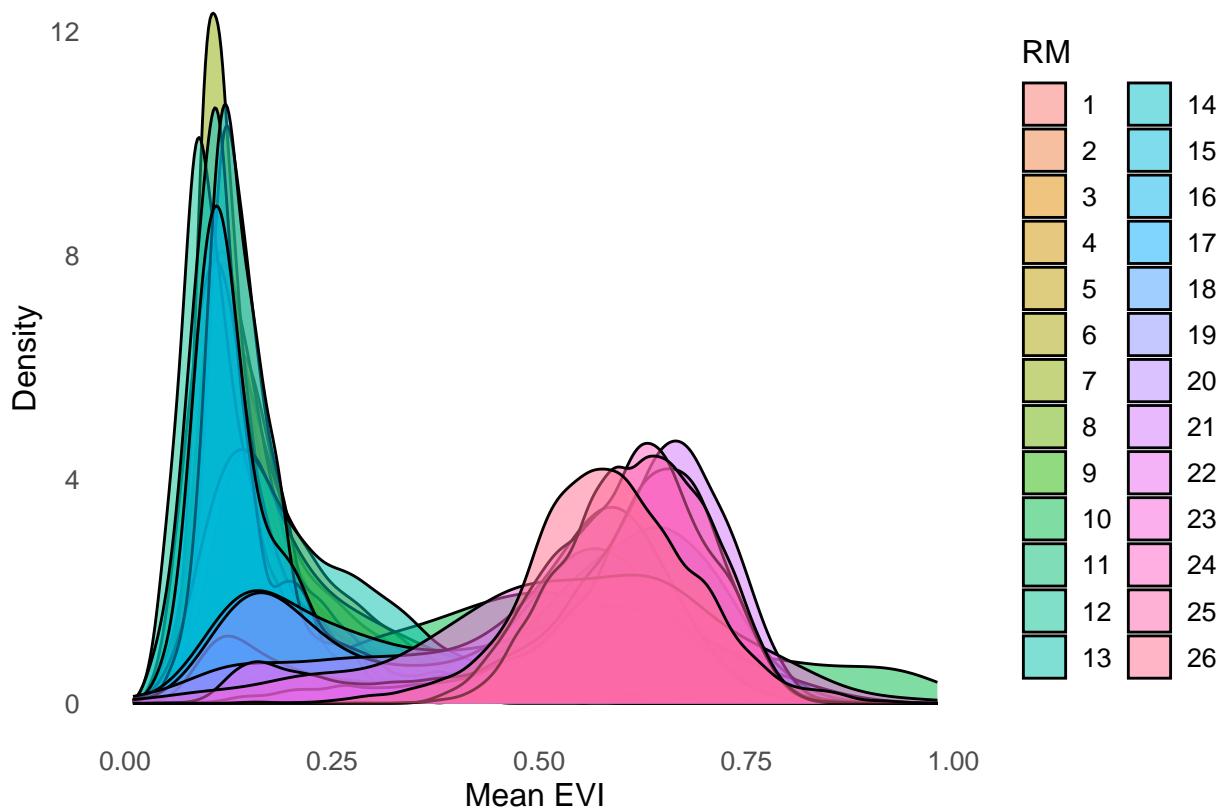
But looking at the distribution of São Paulo mean EVI values in the 100m² grid, we see again the peaks,

```
ggplot(cdata, aes(x = EVI_mean)) +
  geom_histogram(aes(fill = as.factor(cut_width(EVI_mean, 0.01))), binwidth = 0.01, color = NA) +
  scale_fill_viridis_d(option = "turbo", guide = FALSE) +
  labs(x = "Mean EVI", y = "Count") +
  custom_theme
```



Plotting the histogram divided by micro climatic region gives us the peaks again, indicating similar pattern found in the last image.

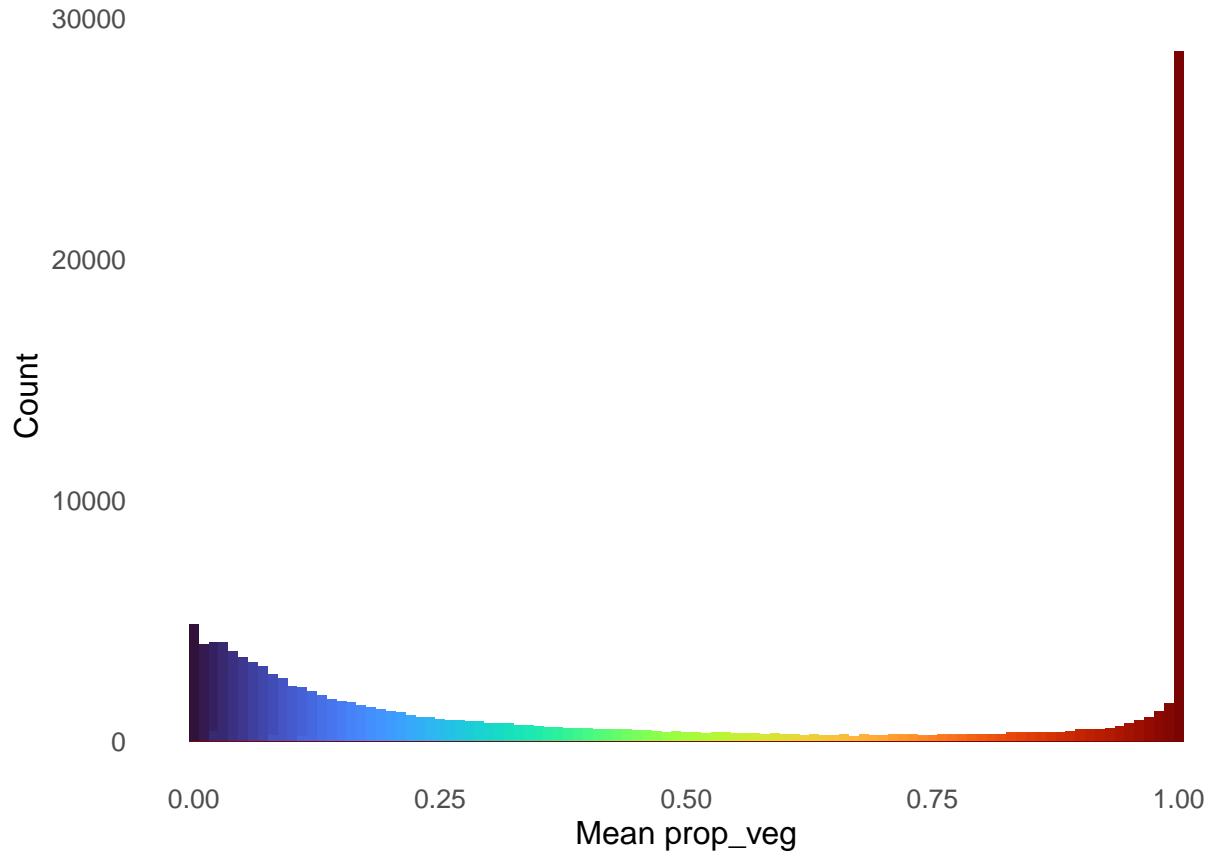
```
ggplot(cdata, aes(x = EVI_mean, fill = factor(RM))) +  
  geom_density(alpha = 0.5) +  
  labs(title = "Density Plot of mean EVI by Microclimatic region", x = "Mean EVI", y = "Density", fill = "RM") +  
  theme_minimal() +  
  theme(panel.grid.major = element_line(colour = "#D9E1F2"), panel.grid.minor = element_line(colour = "#D9E1F2"))
```



This implies that micro climatic regions are way different in vegetation and that might be what is responsible for this situation. But we can visually conclude that the number of places with low temperature peaks higher than the number of places with high EVI.

When we look at proportion of vegetation distribution we see what might lie behind this effect. While the high temperature and low vegetation peaks kind of match, the high EVI doesn't correspond to the peak, but we see that São Paulo has more fully vegetated patches than anything else.

```
ggplot(cdata, aes(x = prop_veg)) +
  geom_histogram(aes(fill = as.factor(cut_width(prop_veg, 0.01))), binwidth = 0.01, color = NA) +
  scale_fill_viridis_d(option = "turbo", guide = FALSE) +
  labs(title = "Distribution of prop_veg by Bin Order", x = "Mean prop_veg", y = "Count") +
  custom_theme
```

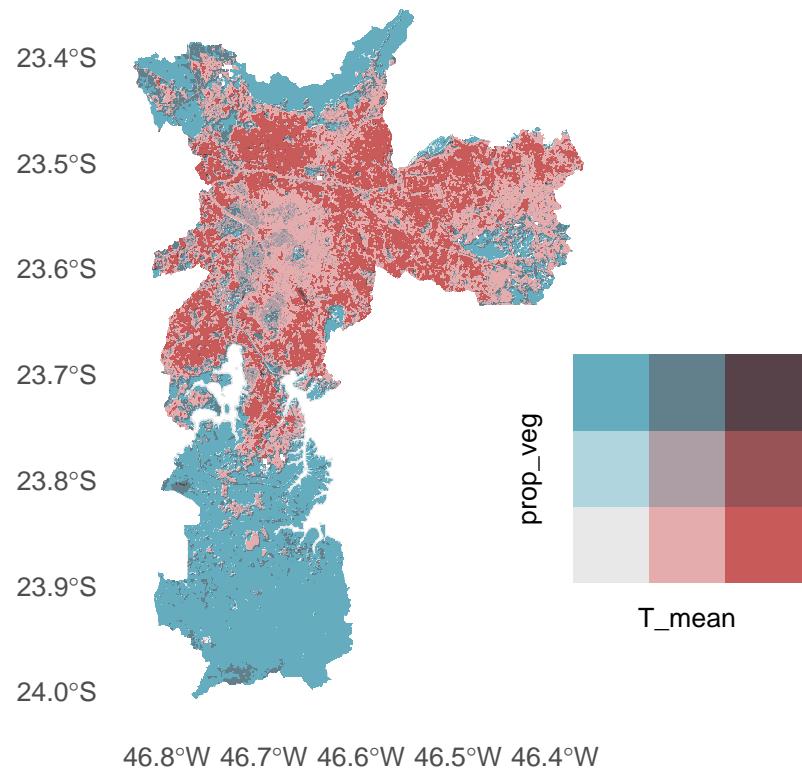


Some hypothesis can be drawn from that. In particular, that the effect of vegetation in temperature is not linear, but increases with the increase in vegetation. Also, that we might be seeing an spatial effect of vegetation, where it not only reduces local temperature but can cool down neighboring regions. Lets explore those hypothesis.

Let's see the maps overlaying

```
plot_bi_map(data, "T_mean", "prop_veg")
```

```
##  
##   1-1    1-2    1-3    2-1    2-2    2-3    3-1    3-2    3-3  
##  2487   2455  46675  35988  13712  12059  32919   1686    201
```

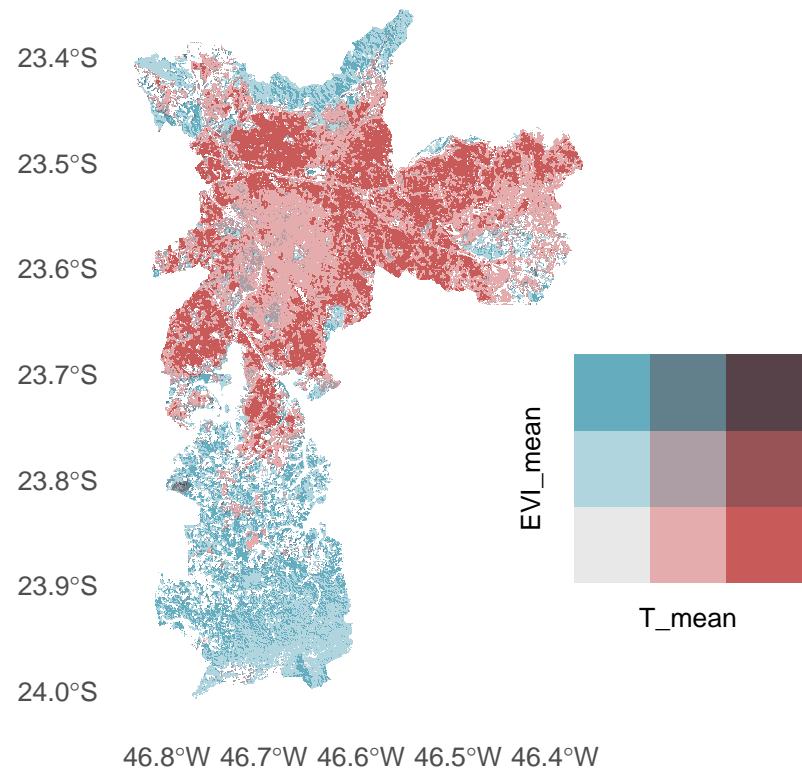


Combinations of low proportion of vegetation and low mean temperature seem to be around water, forest borders, and in a specific region in the center of the city. In the reserves, we see low temperature and high vegetation. In the urban part of the city we see low vegetation with high temperature as expected.

We can also check the biplots for other variables

```
plot_bi_map(cdata, "T_mean", "EVI_mean")
```

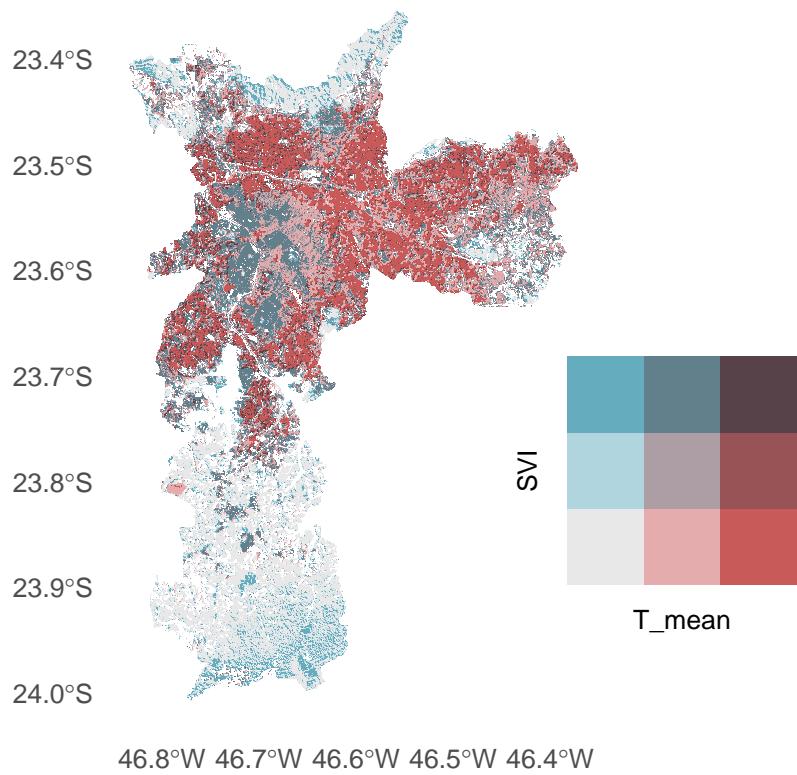
```
## 
##   1-1   1-2   1-3   2-1   2-2   2-3   3-1   3-2   3-3
##   883  25412 13569 40217  9598  1076 32299   428    18
```



Lets check for our response variables

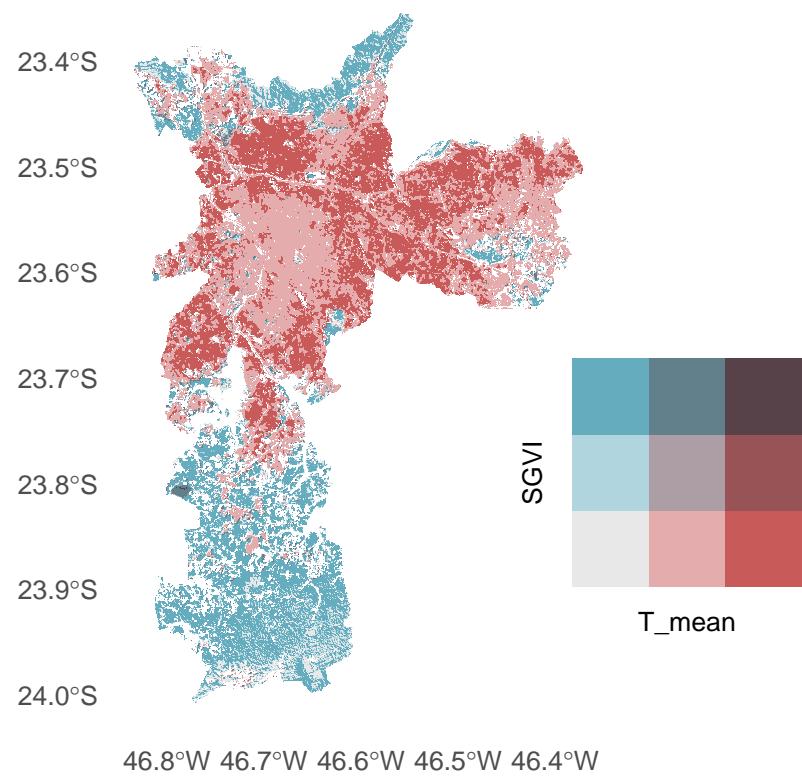
```
plot_bi_map(cdata, "T_mean", "SVI")
```

```
##  
##   1-1   1-2   1-3   2-1   2-2   2-3   3-1   3-2   3-3  
## 27288  6417  6159 19564 13044 18283 22131  6723  3891
```



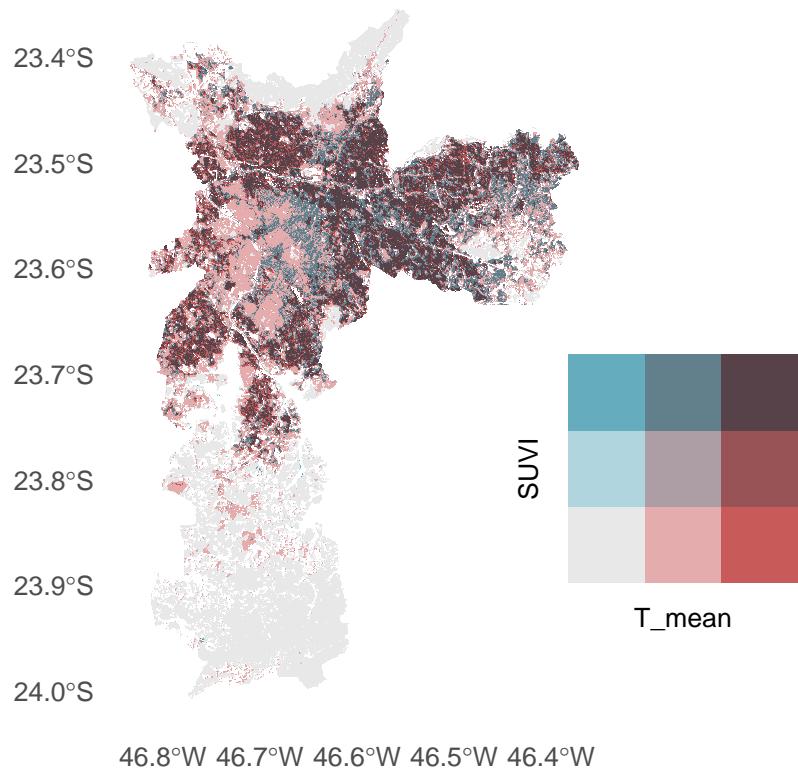
```
plot_bi_map(cdata, "T_mean", "SGVI")
```

```
##  
##   1-1    1-2    1-3    2-1    2-2    2-3    3-1    3-2    3-3  
##  7521   6462  25881  46602   2010   2279  32651     64     30
```



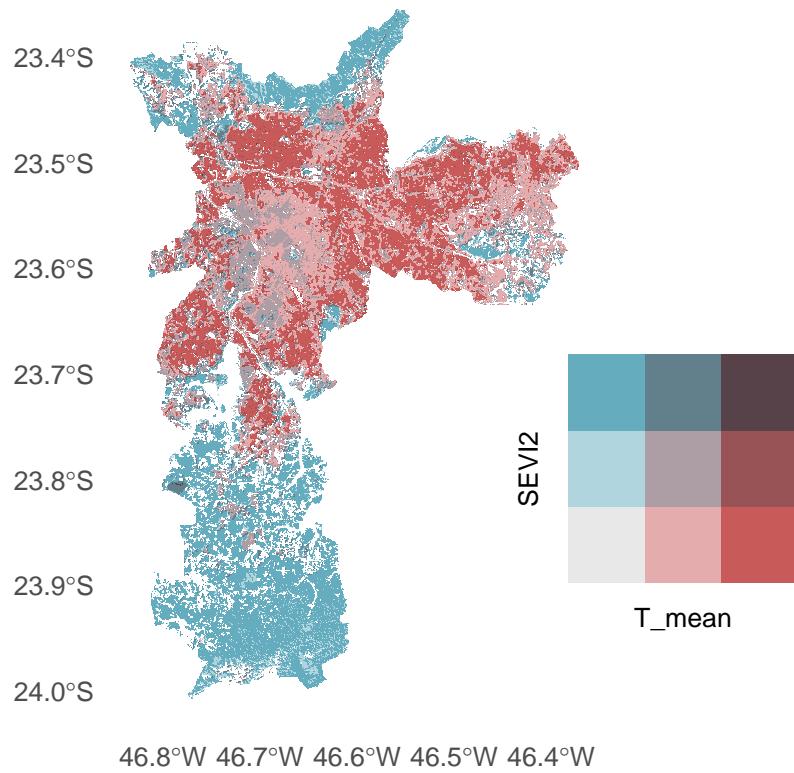
```
plot_bi_map(cdata, "T_mean", "SGVI")
```

```
##  
##   1-1    1-2    1-3    2-1    2-2    2-3    3-1    3-2    3-3  
## 39630    153     81 26020   9853 15018   4870   6736 21139
```



```
plot_bi_map(cdata, "T_mean", "SEVI2")
```

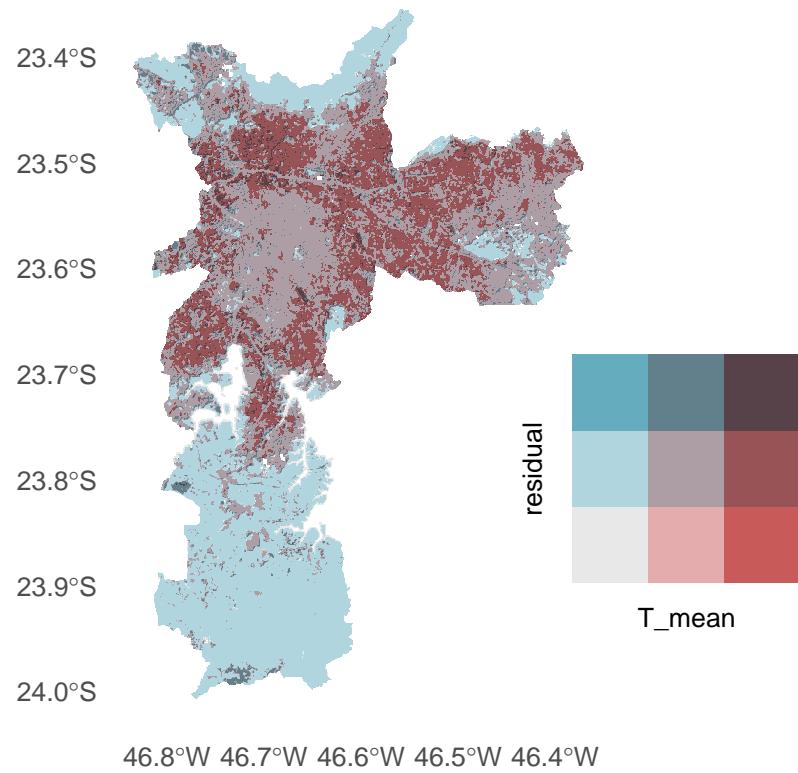
```
##  
##   1-1    1-2    1-3    2-1    2-2    2-3    3-1    3-2    3-3  
##   267    6129   33468  26938  19253  4700   29045  3596   104
```



Those maps show us important features about the homogeneity of the process that is happening in the city. Let's check the residuals of a linear model:

```
linear <- lm(T_mean ~ prop_veg, data)
data$residual <- resid(linear)
plot_bi_map(data, "T_mean", "residual")

## 
##   1-1   1-2   2-1   2-2   2-3   3-2   3-3
##  2871 48746   252 54775   6732 30974   3832
```



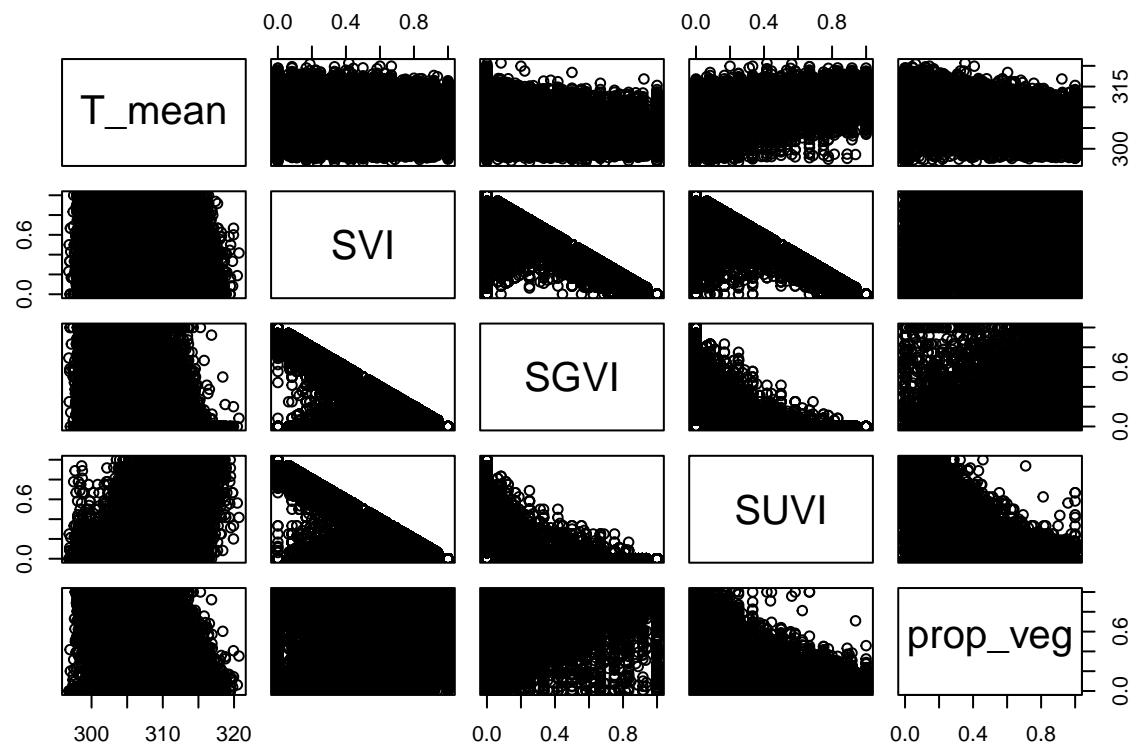
The variables are highly colinear, as we can see in pairwise correlation

```

predictors <- c("T_mean",
                 "#"SBmean",
                 "SVI",
                 "SGVI",
                 "SUVI",
                 "prop_veg")

pairs(as.data.frame(cdata)[, predictors])

```



we can see high correlation among predictor variables. Lets check the collective explained variance