

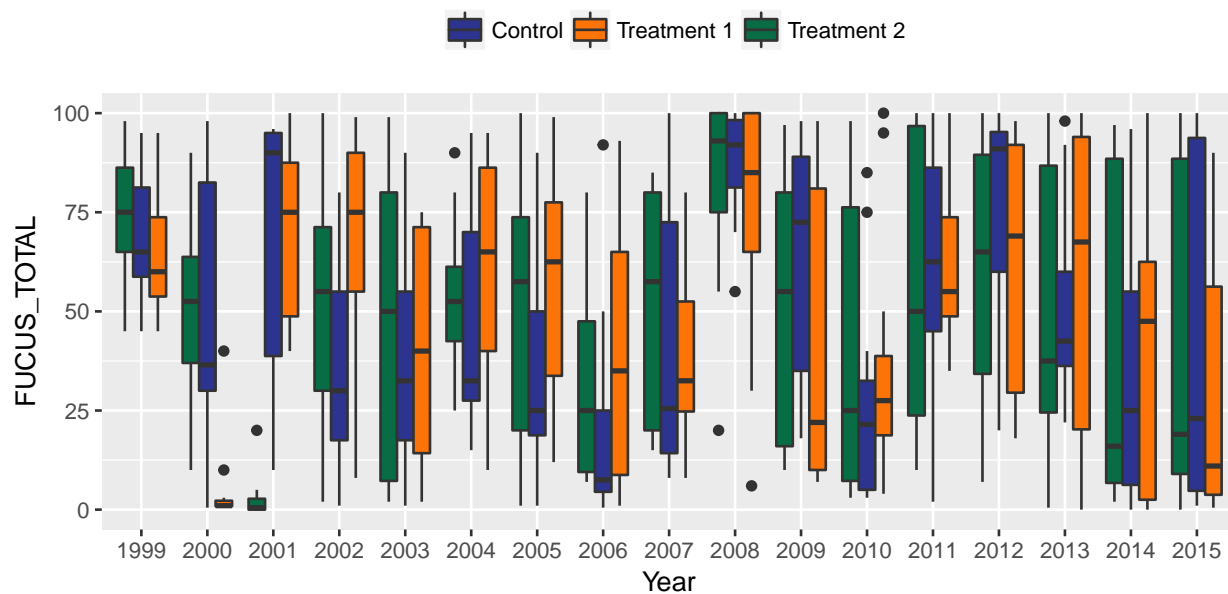
# Data Visualization

*Rachael E. Blake*

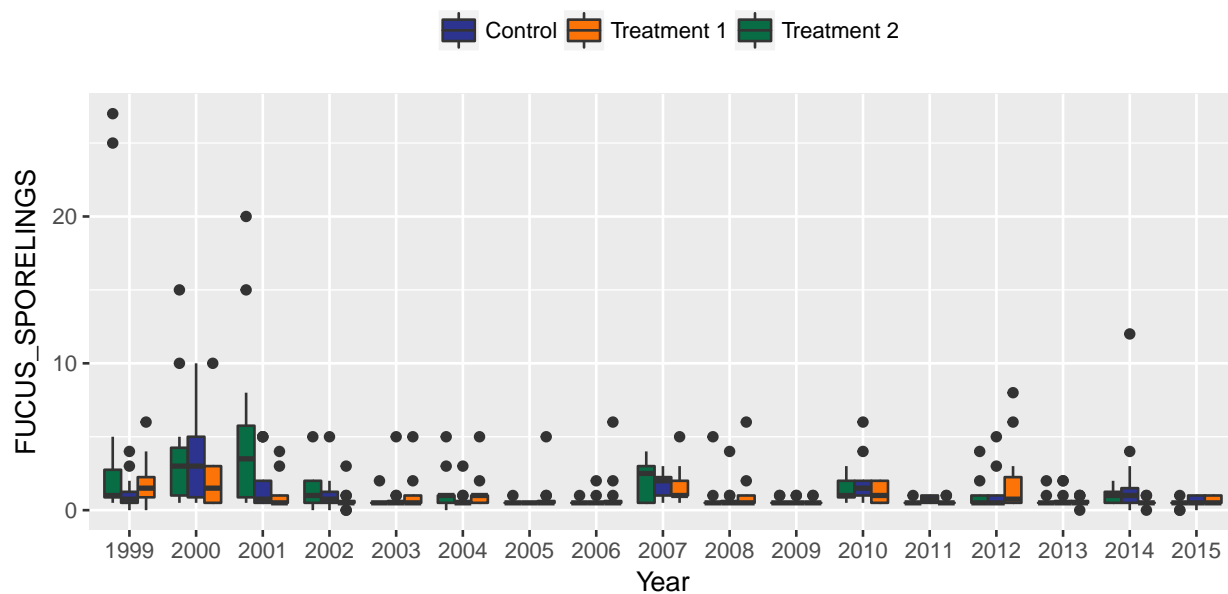
*May 1, 2017*

## Invertebrates: Percent Cover Data

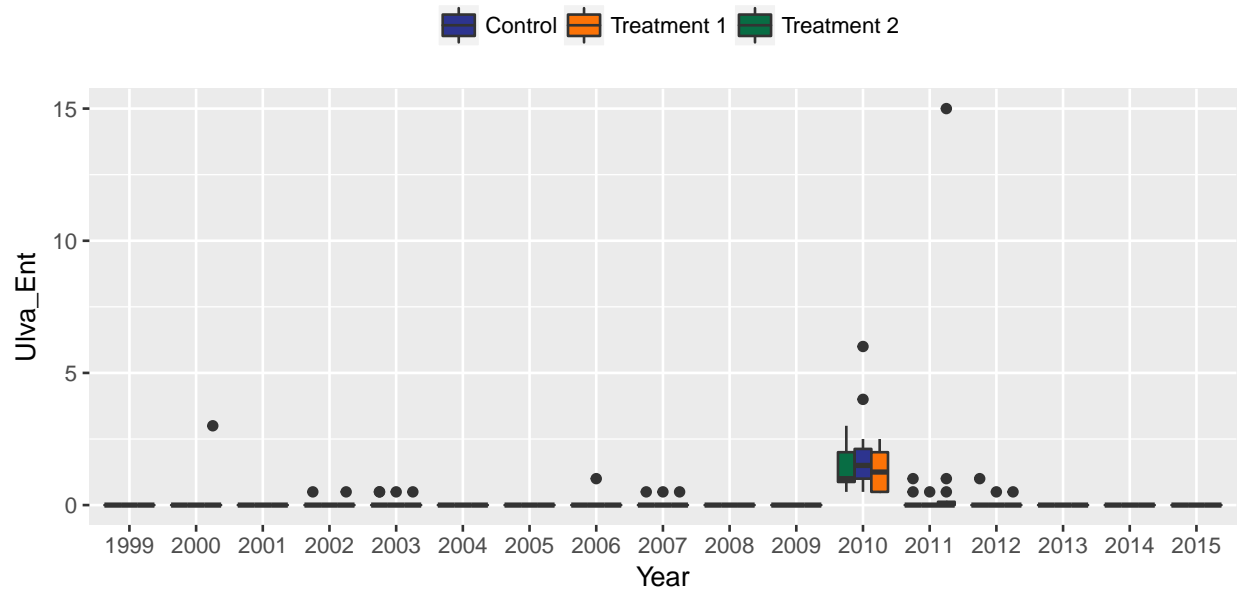
```
## [[1]]
```



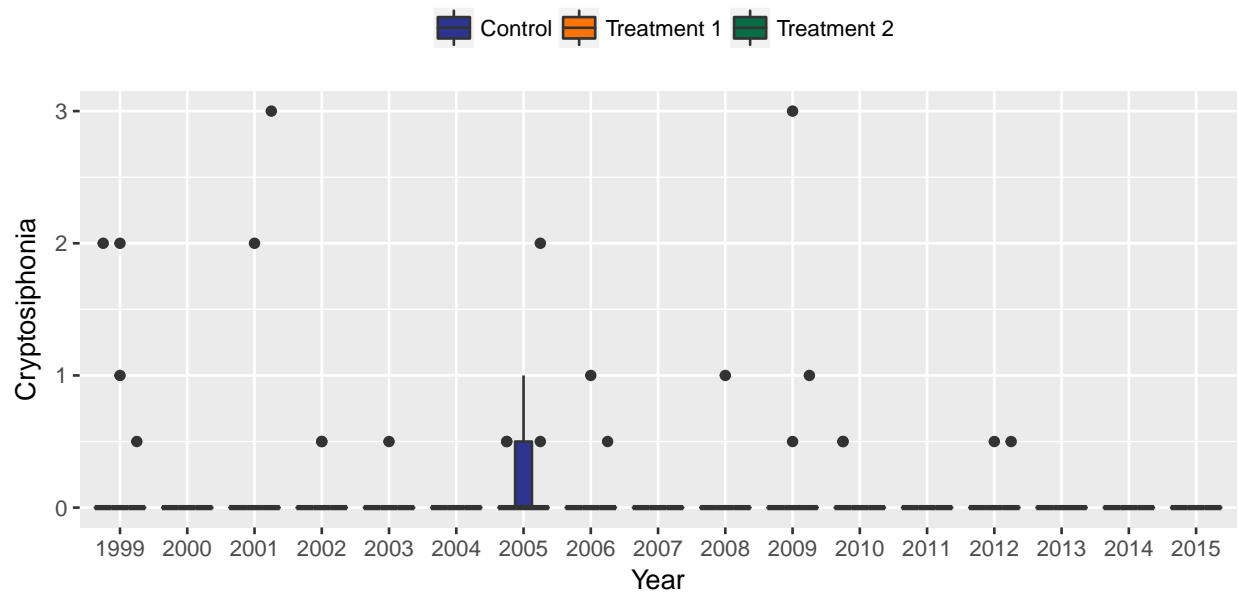
```
##  
## [[2]]
```



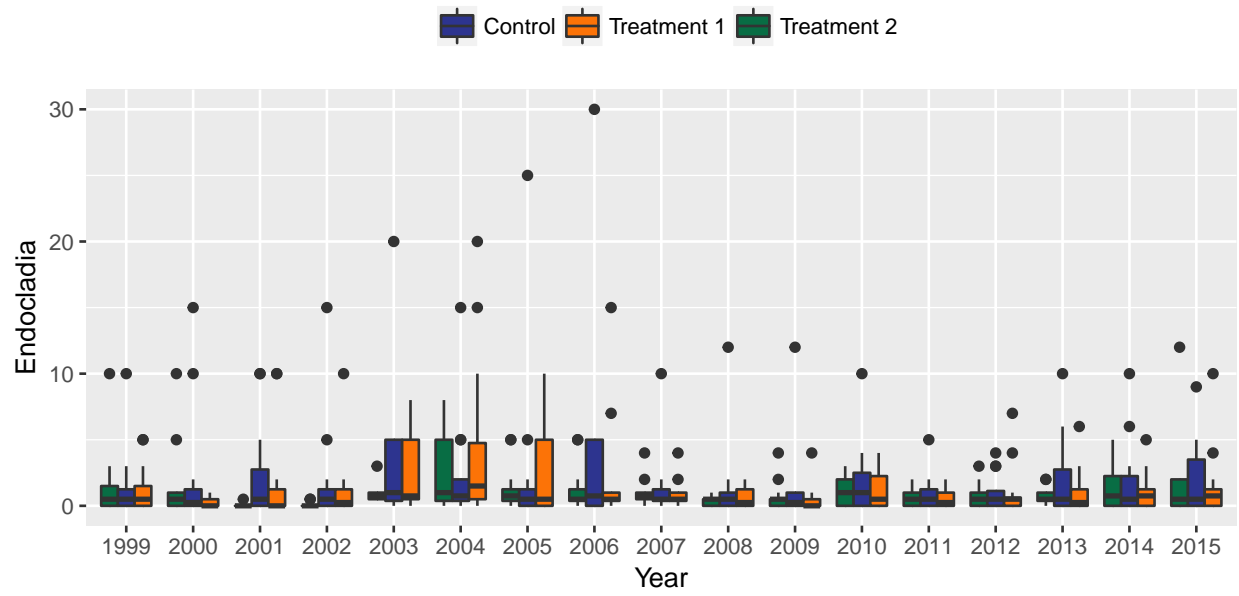
```
##  
## [[3]]
```



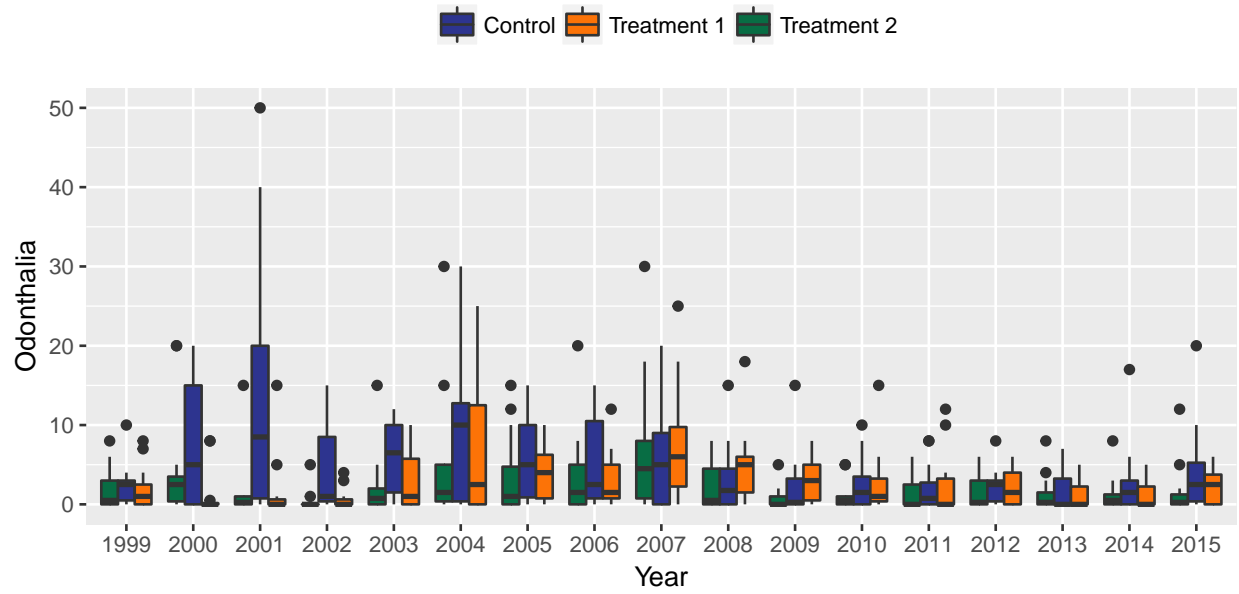
```
##  
## [[4]]
```



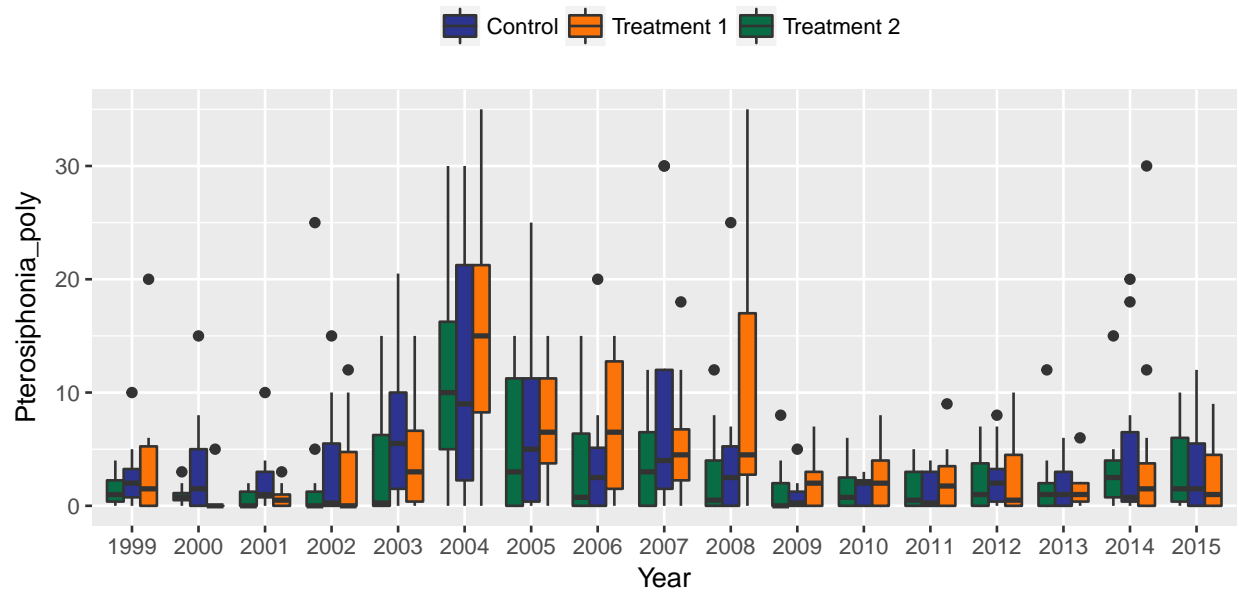
```
##  
## [[5]]
```



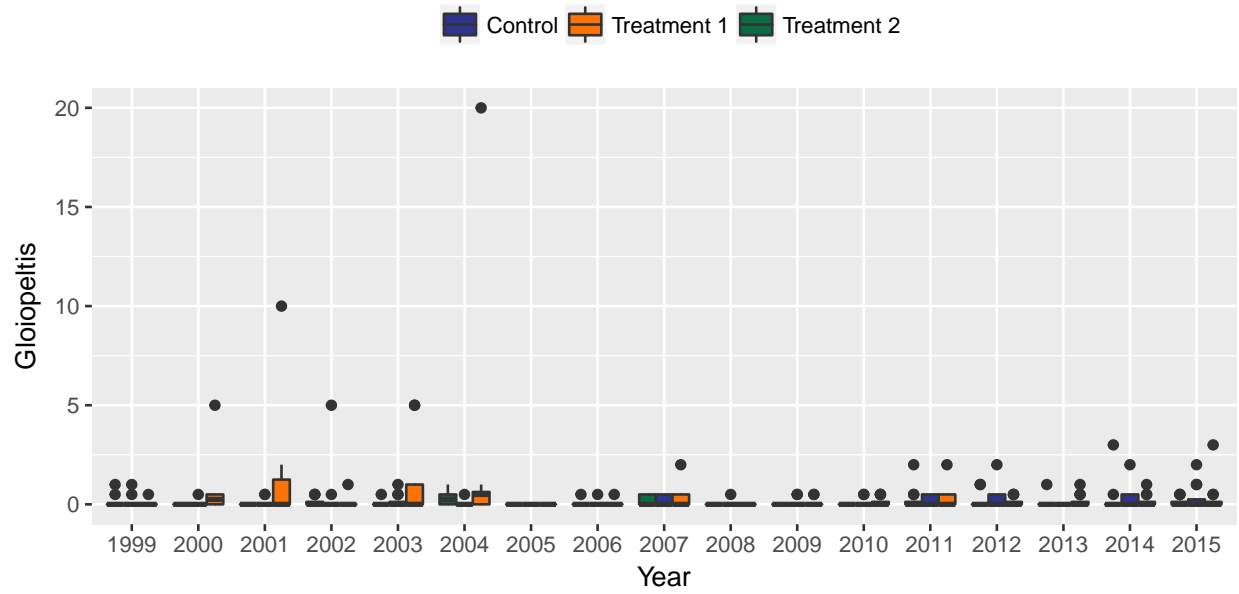
```
##
## [[6]]
```



```
##
## [[7]]
```



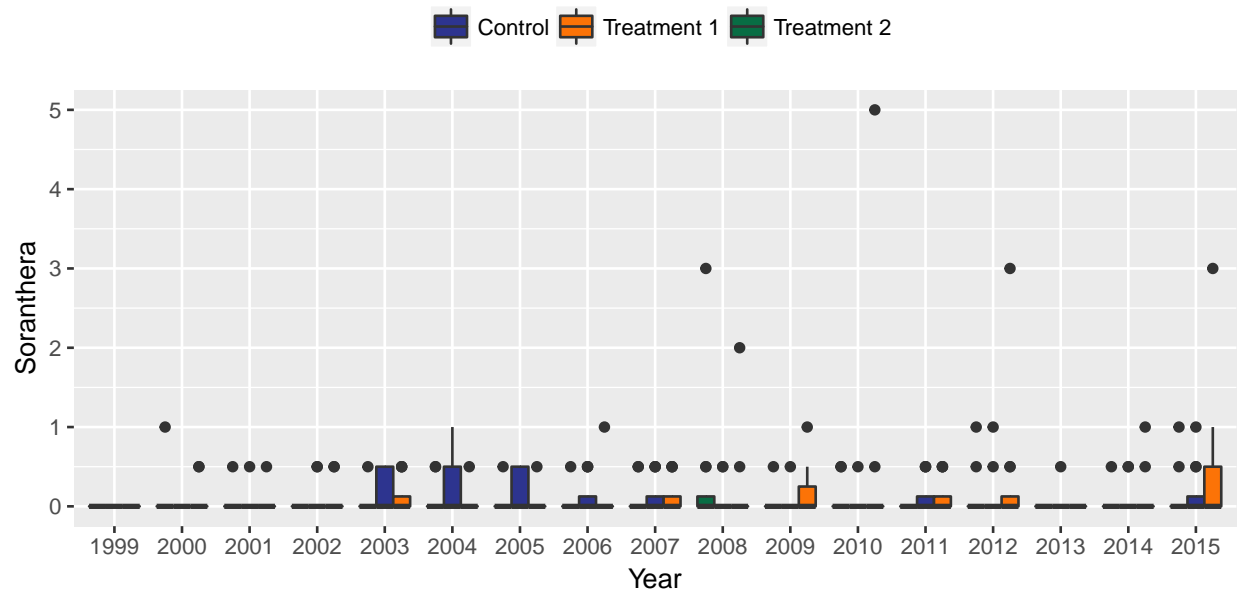
```
##
## [[8]]
```



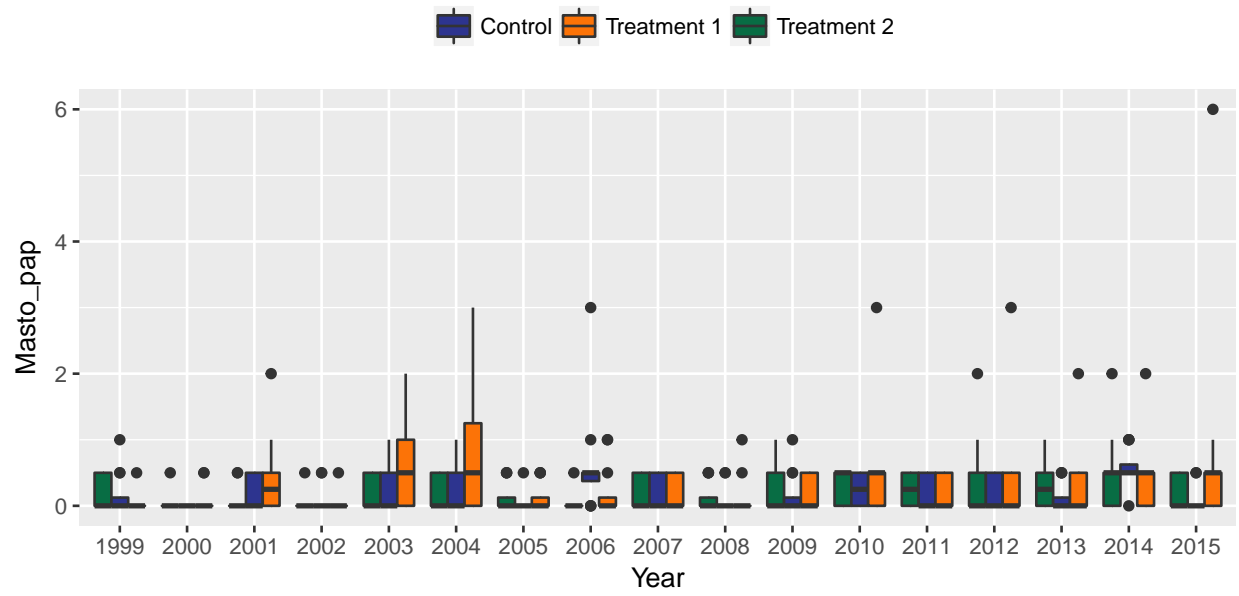
```
##
## [[9]]
```



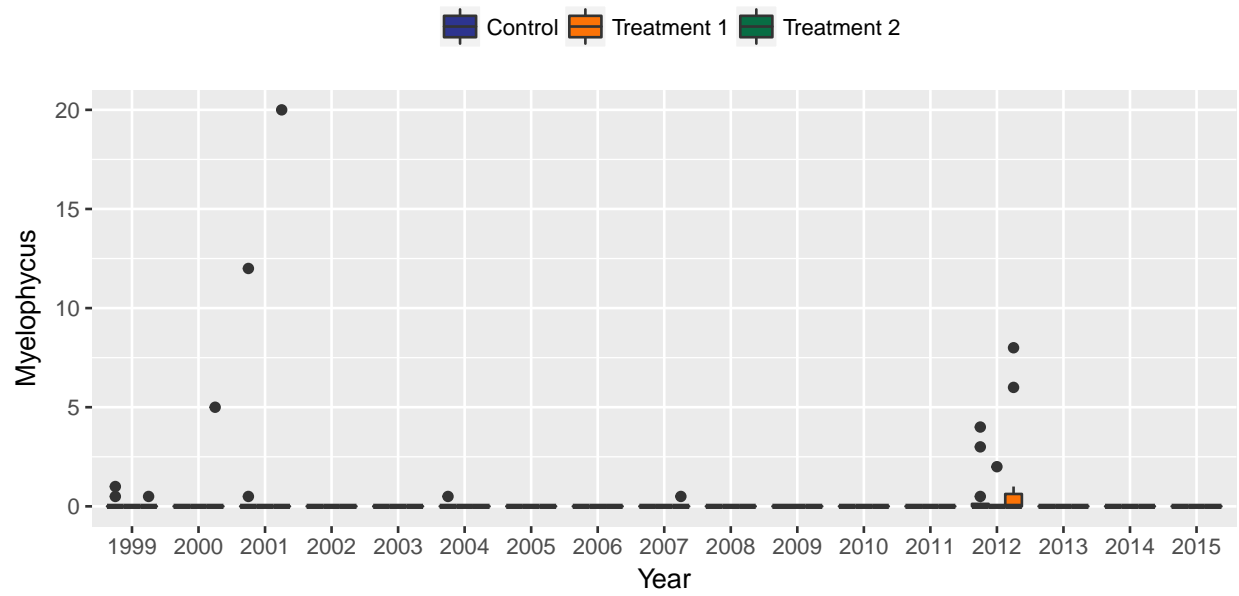
```
##
## [[10]]
```



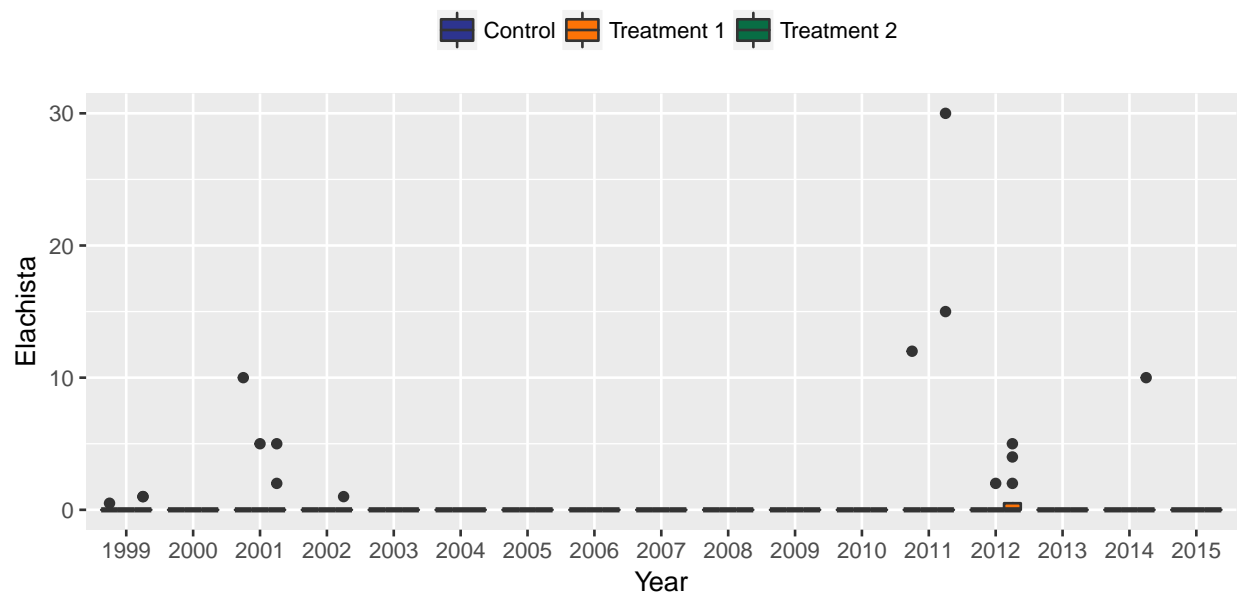
```
##
## [[11]]
```



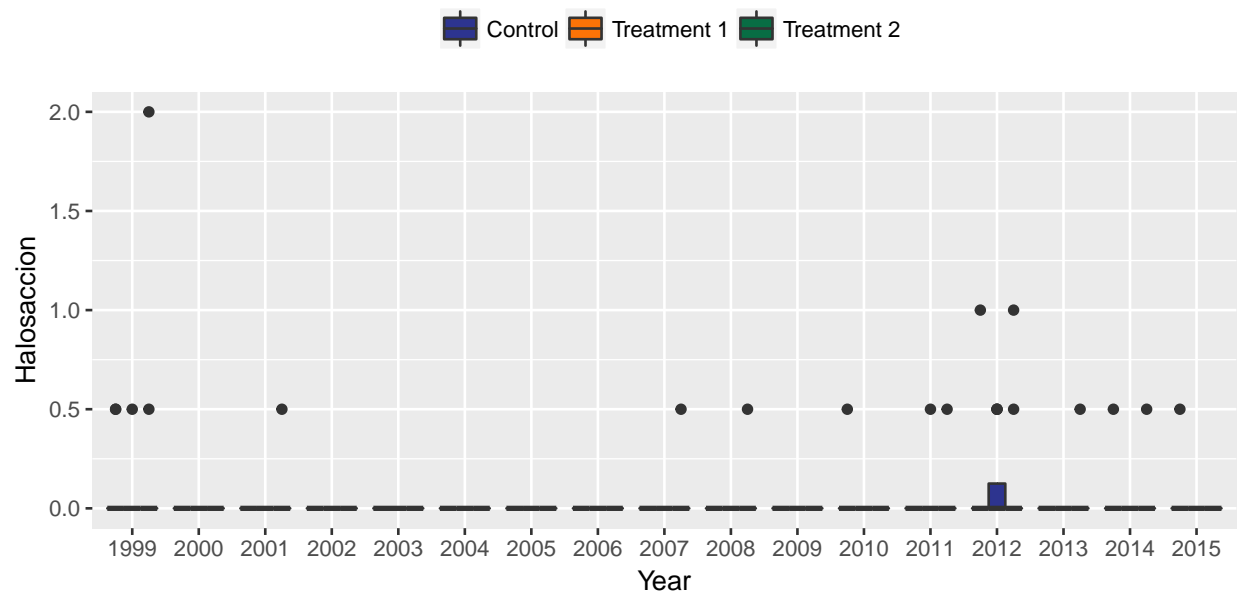
##  
## [[12]]



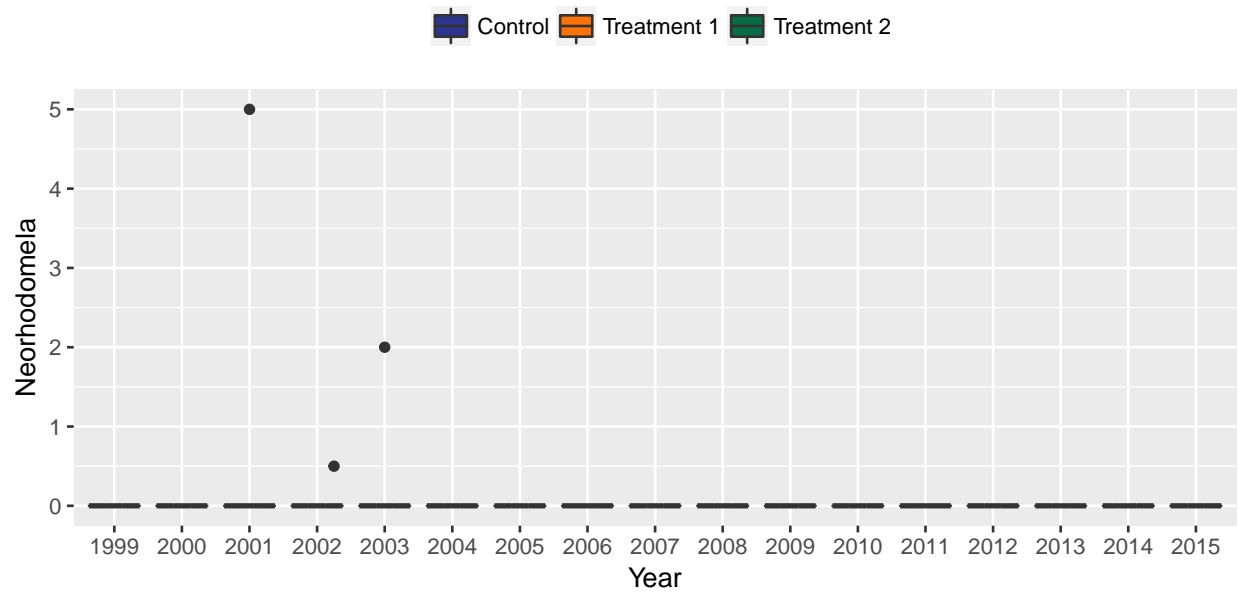
##  
## [[13]]



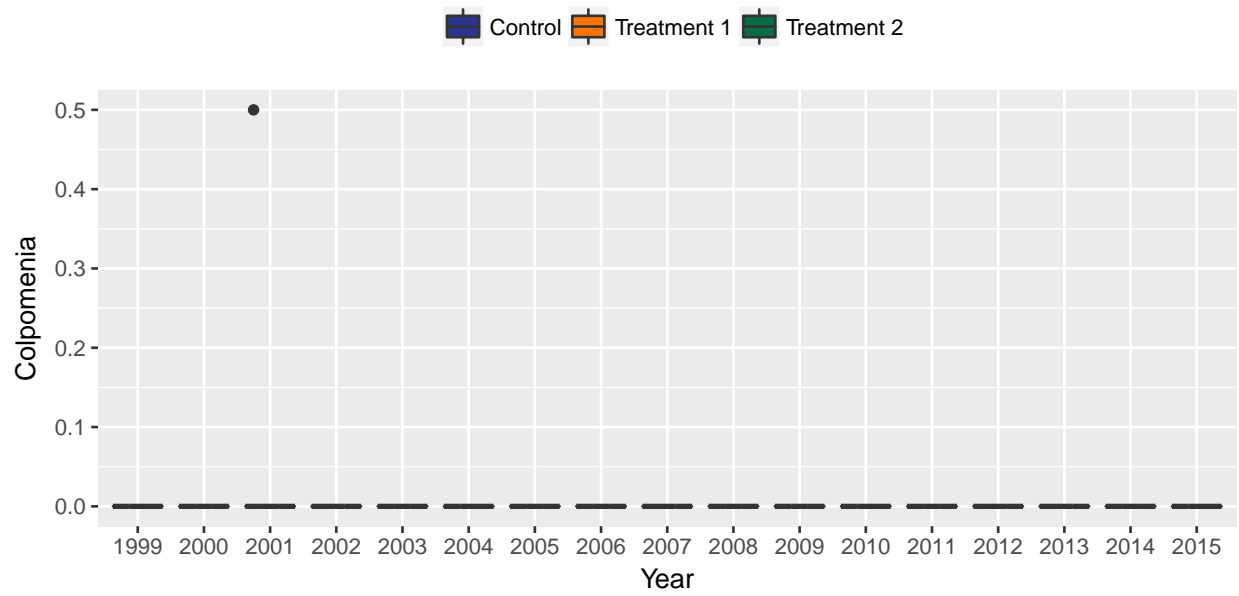
##  
## [[14]]



##  
## [[15]]

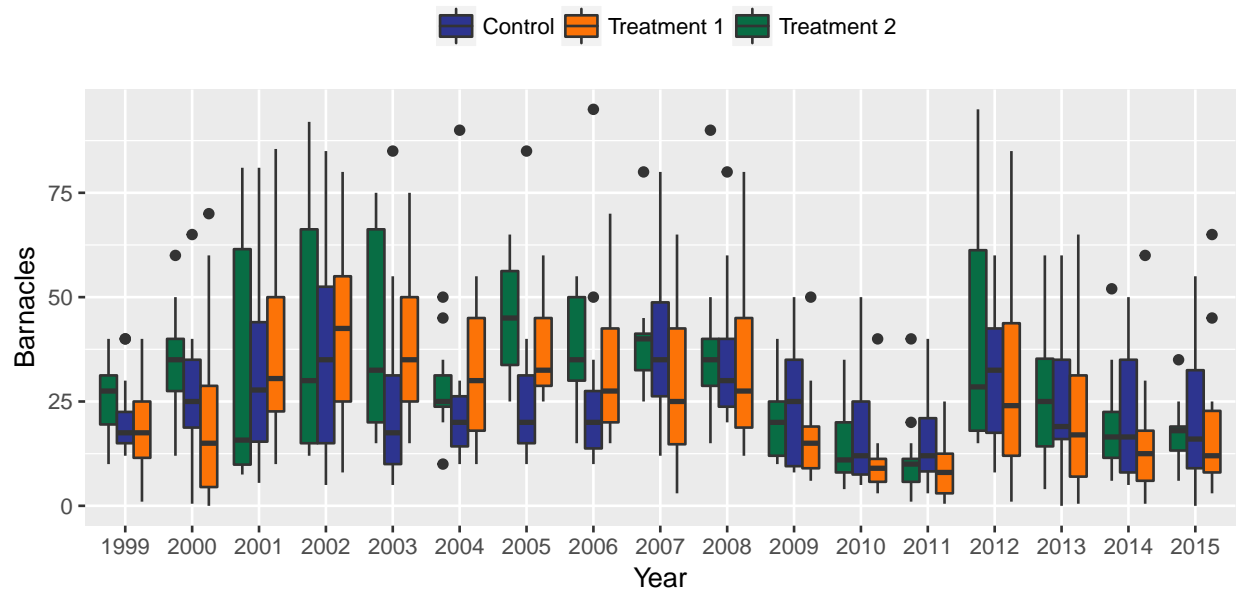


```
##
## [[16]]
```

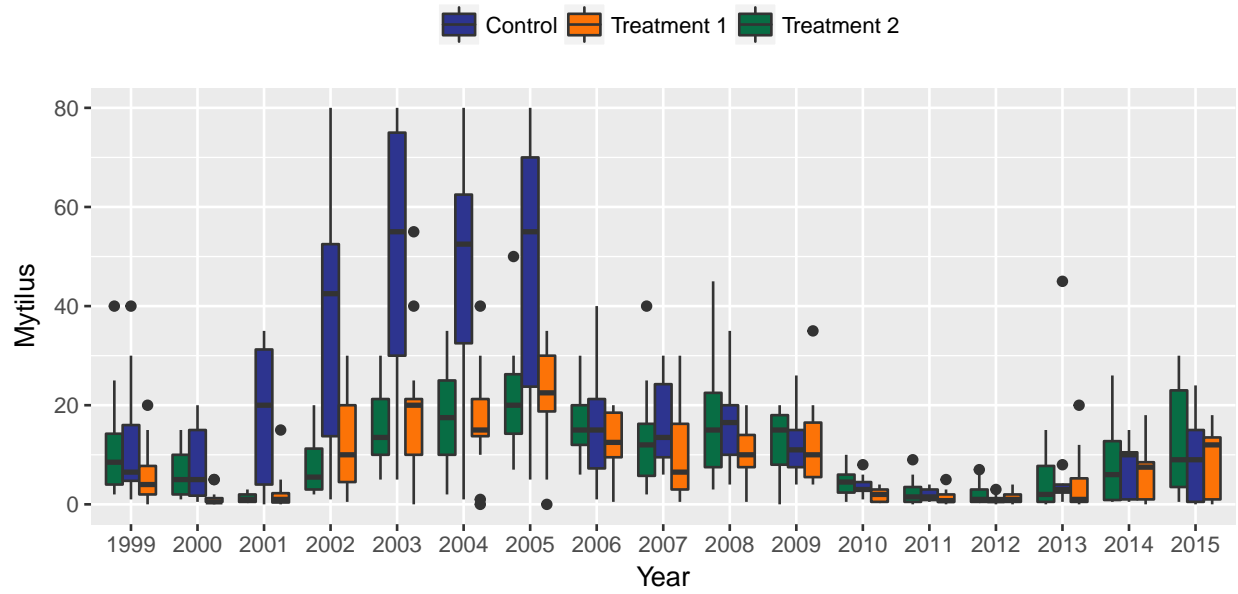


```
##
## [[17]]
```

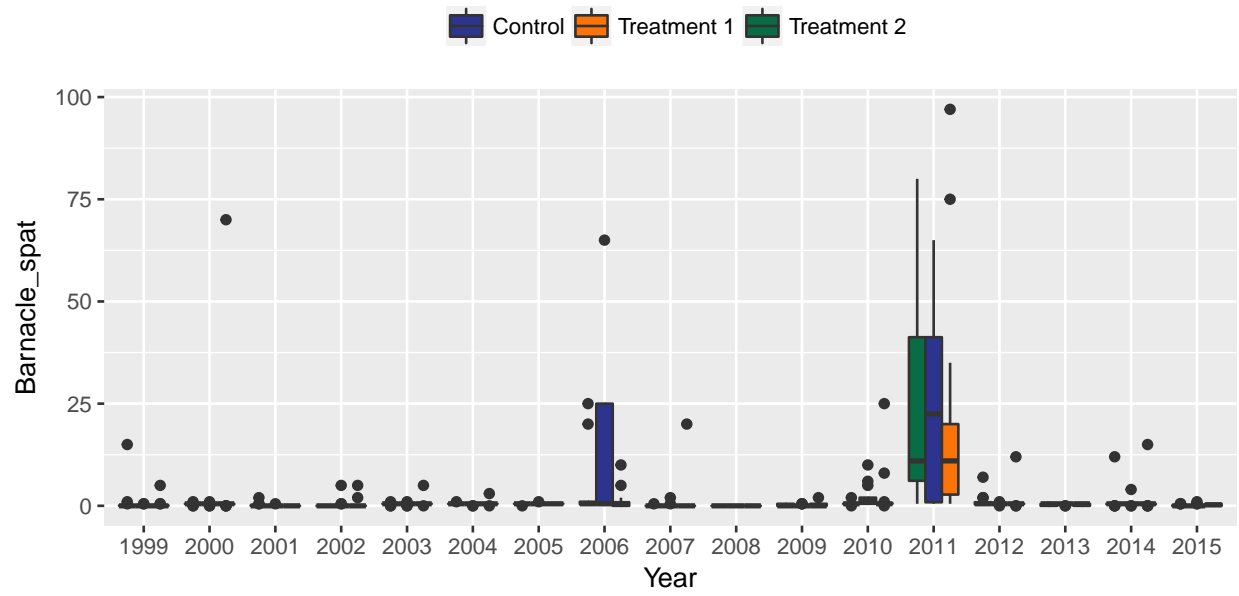




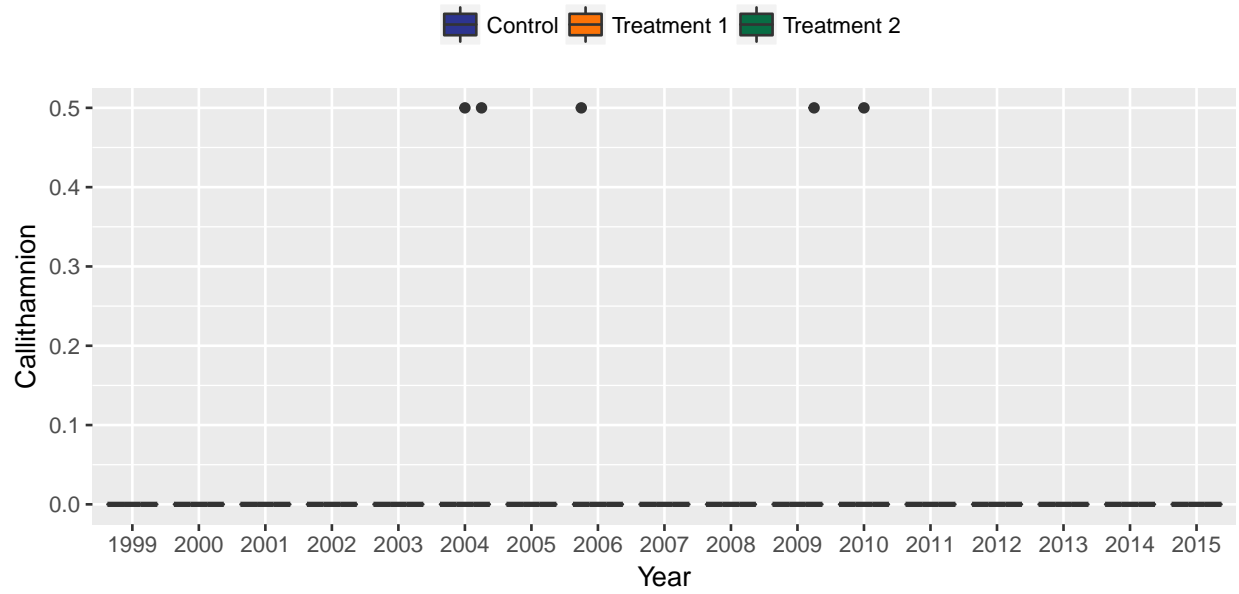
```
##
## [[18]]
```



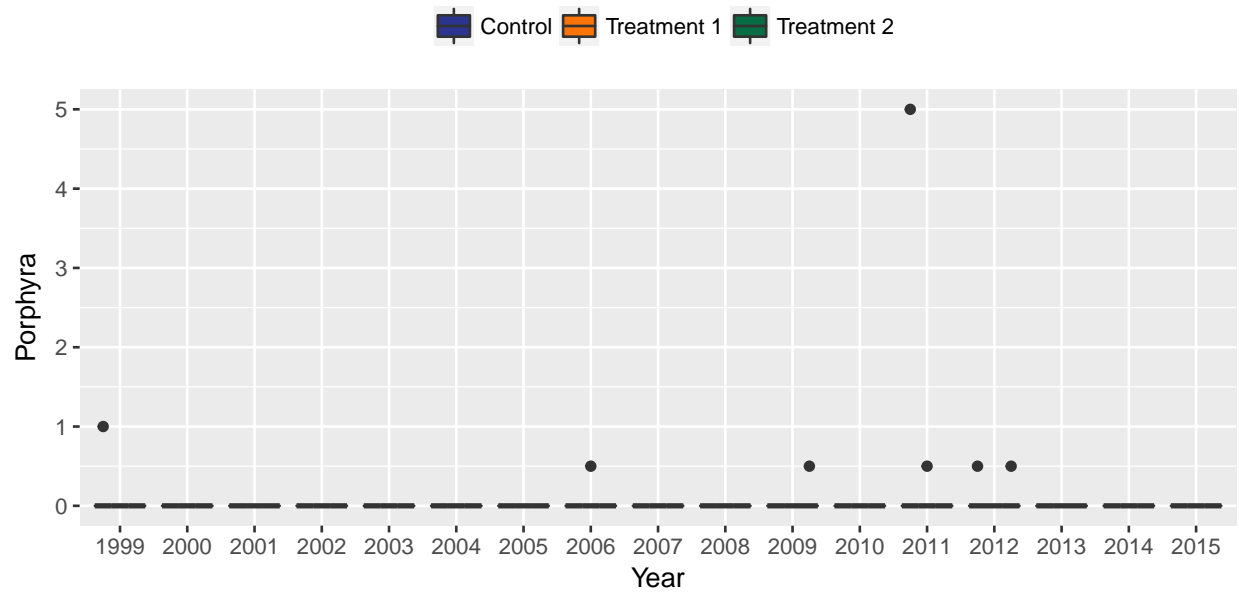
```
##
## [[19]]
```



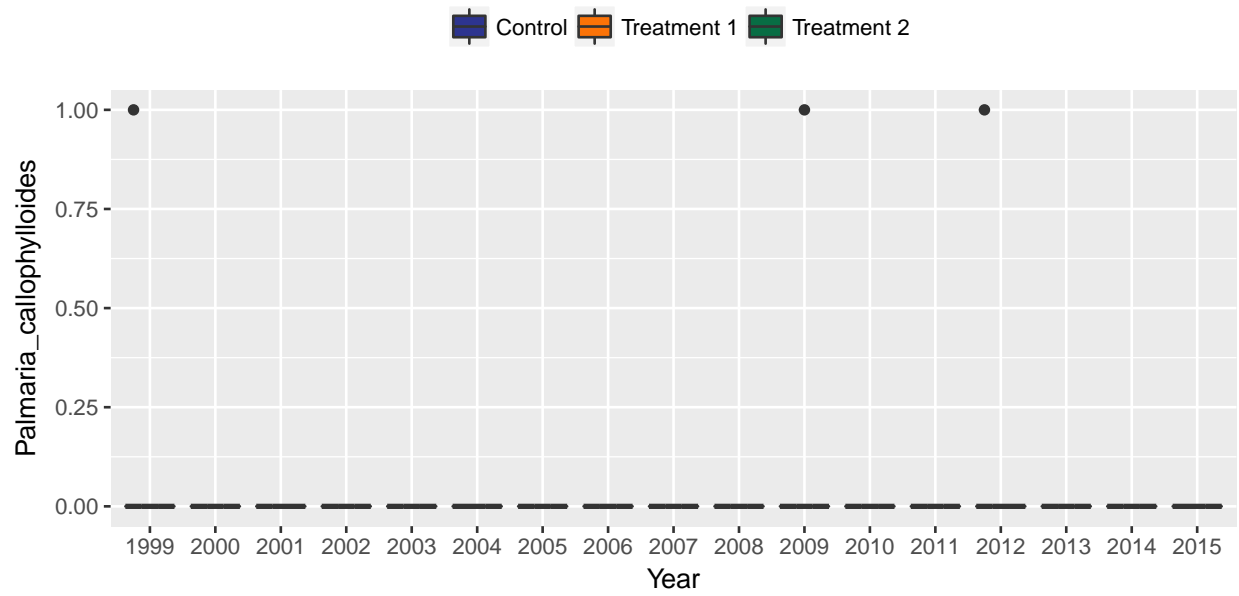
```
##
## [[20]]
```



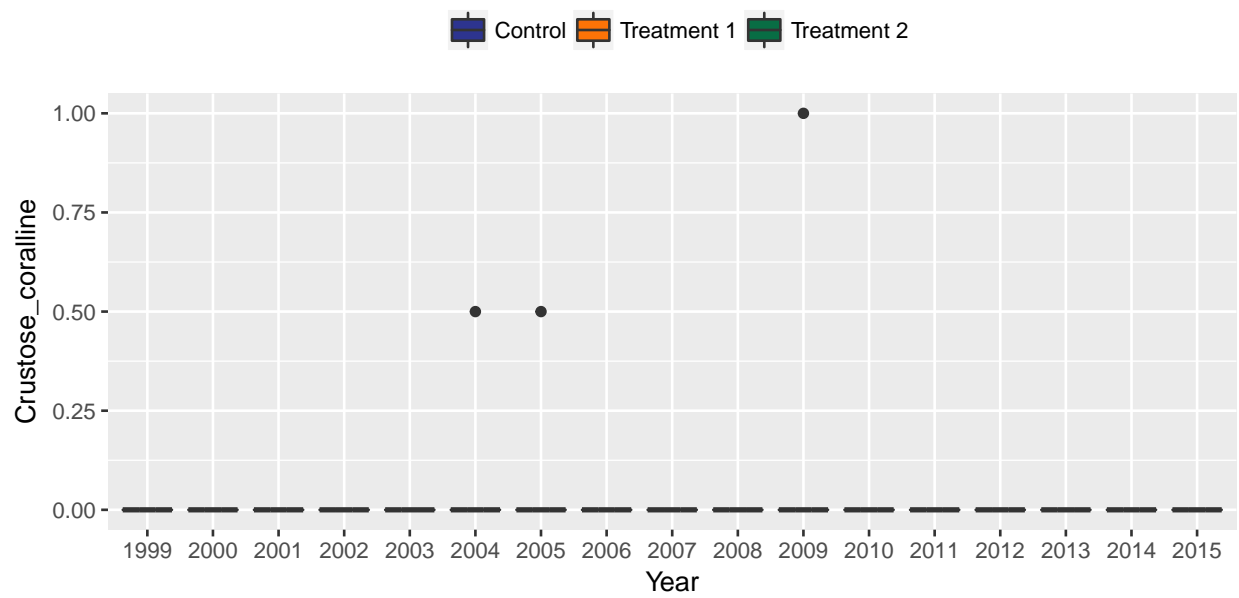
```
##
## [[21]]
```



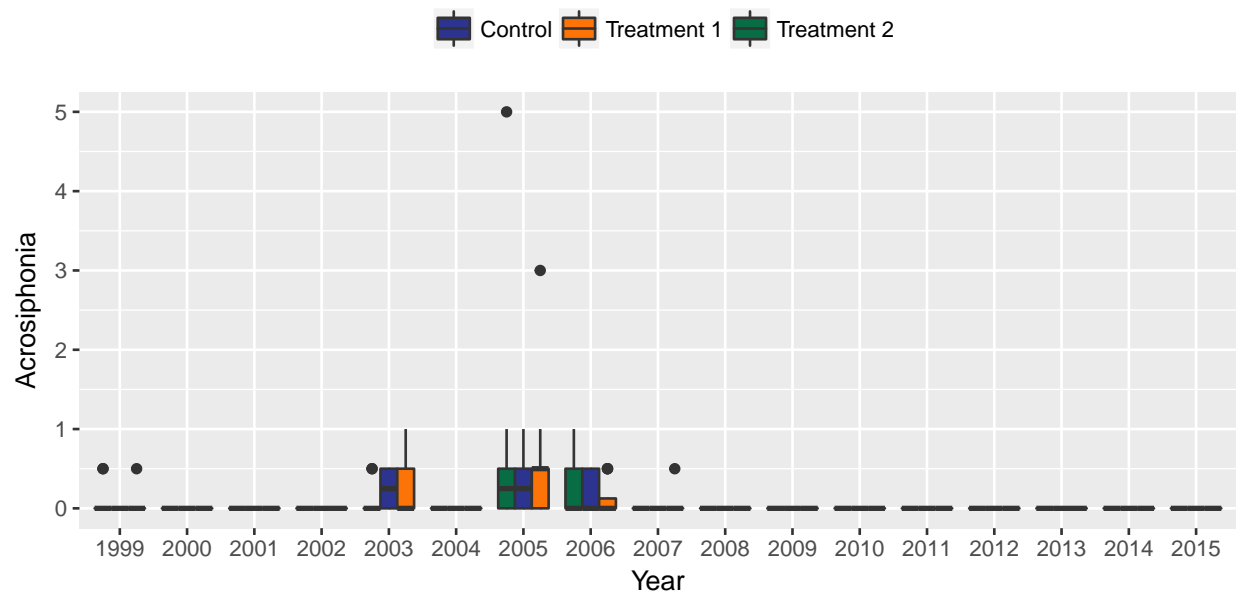
##  
## [[22]]



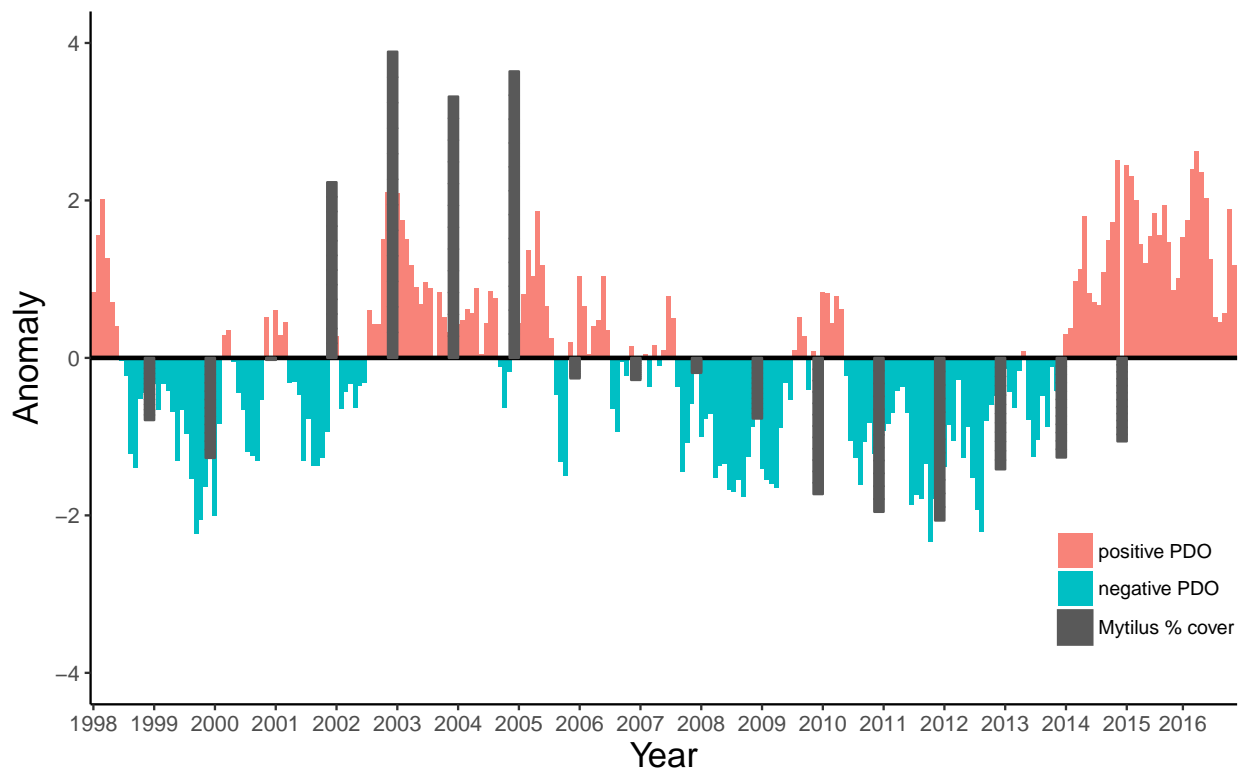
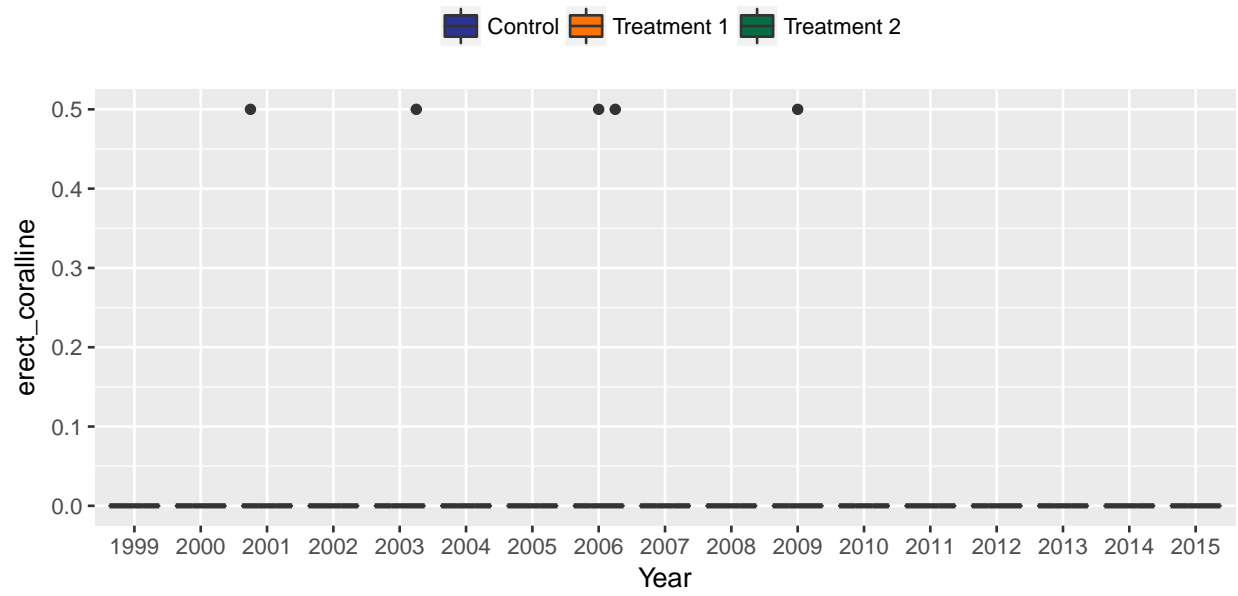
##  
## [[23]]



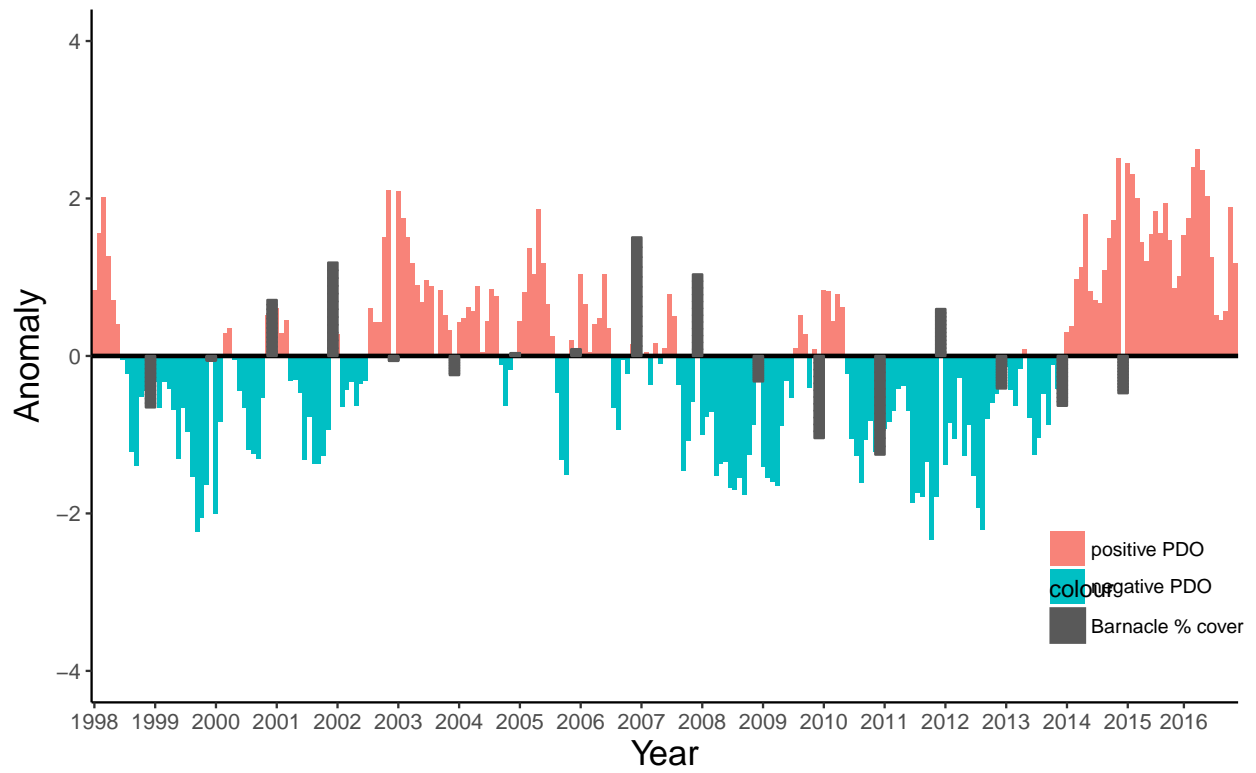
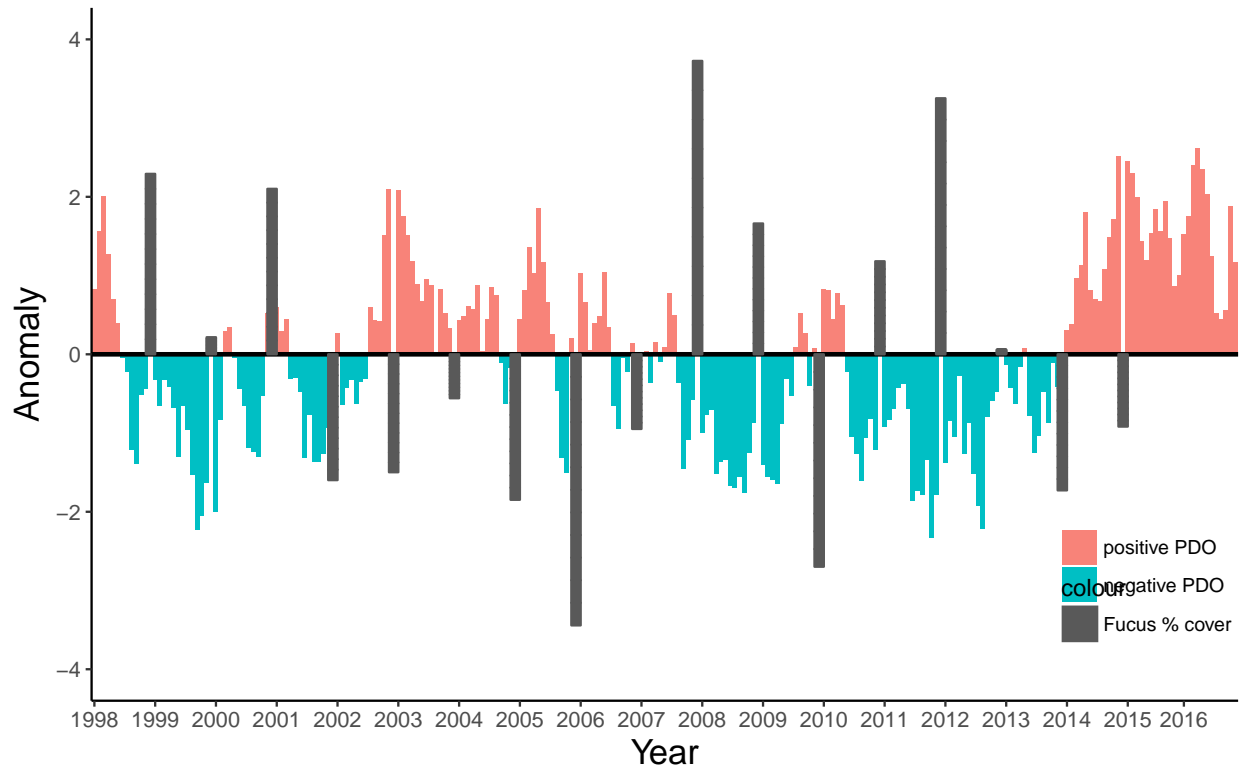
##  
## [[24]]

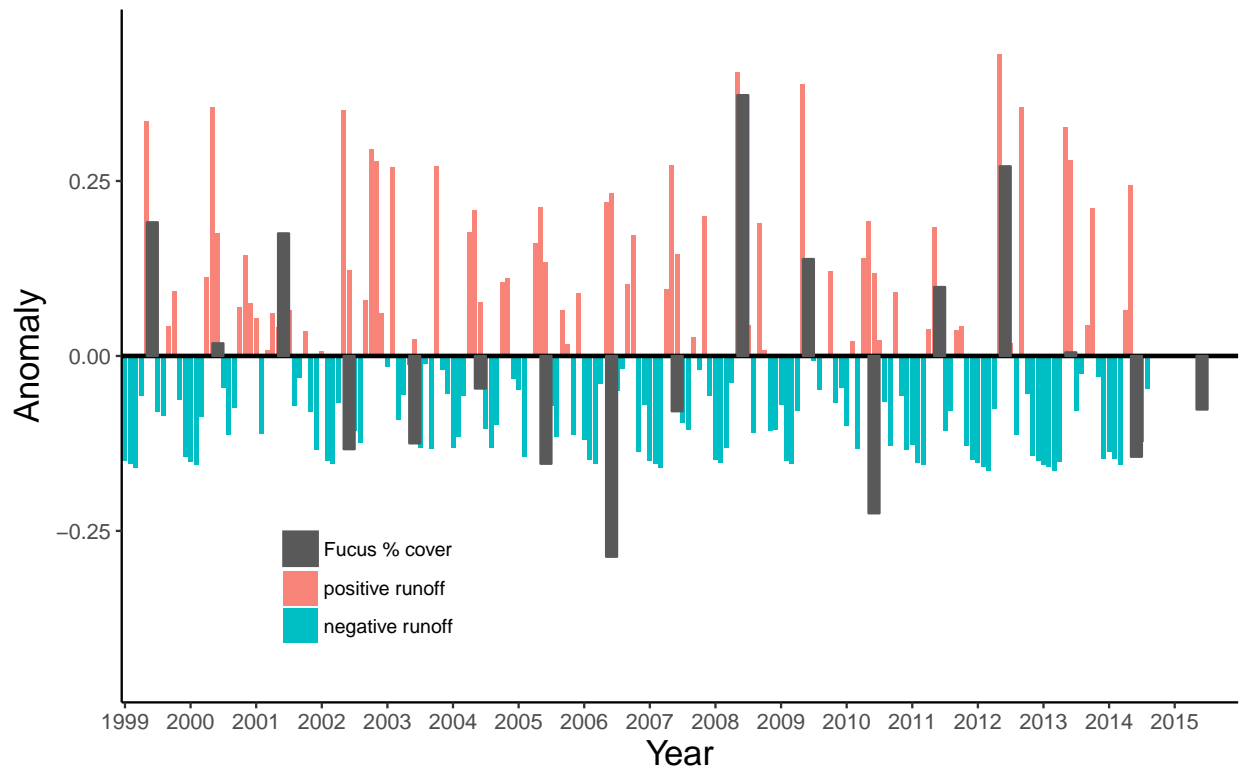
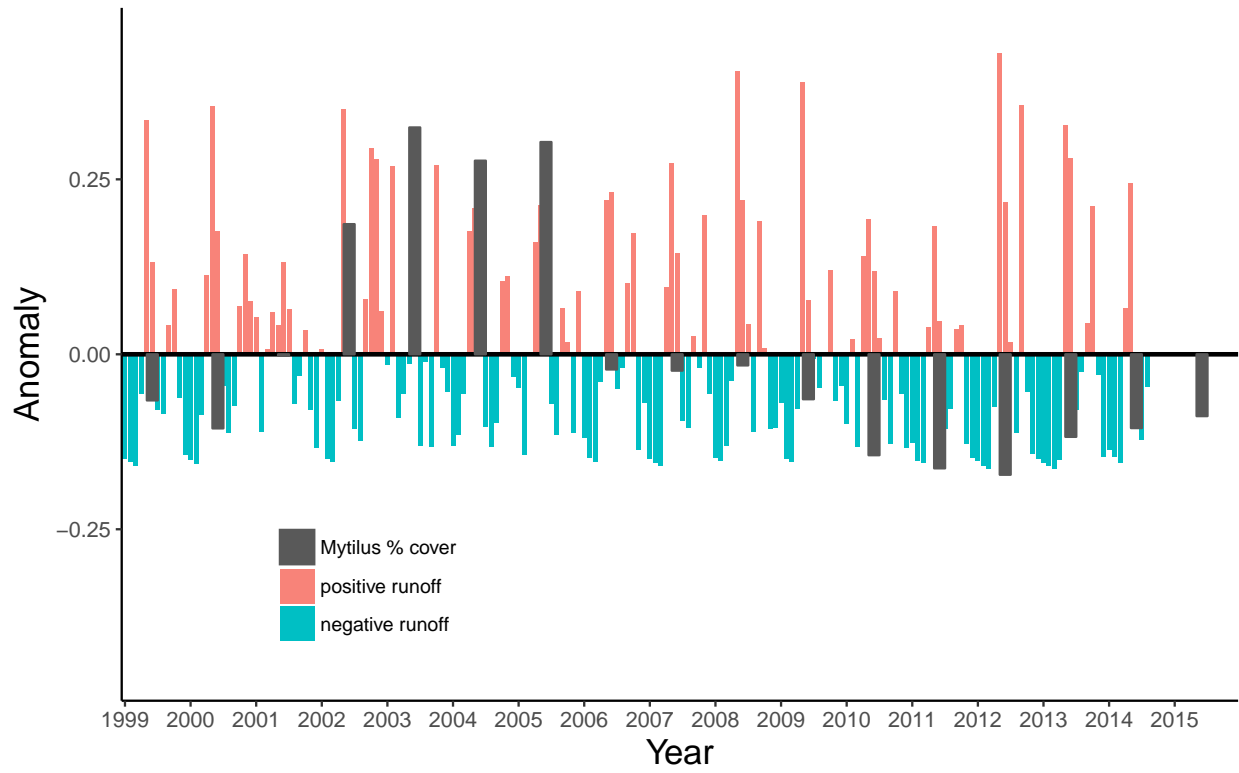


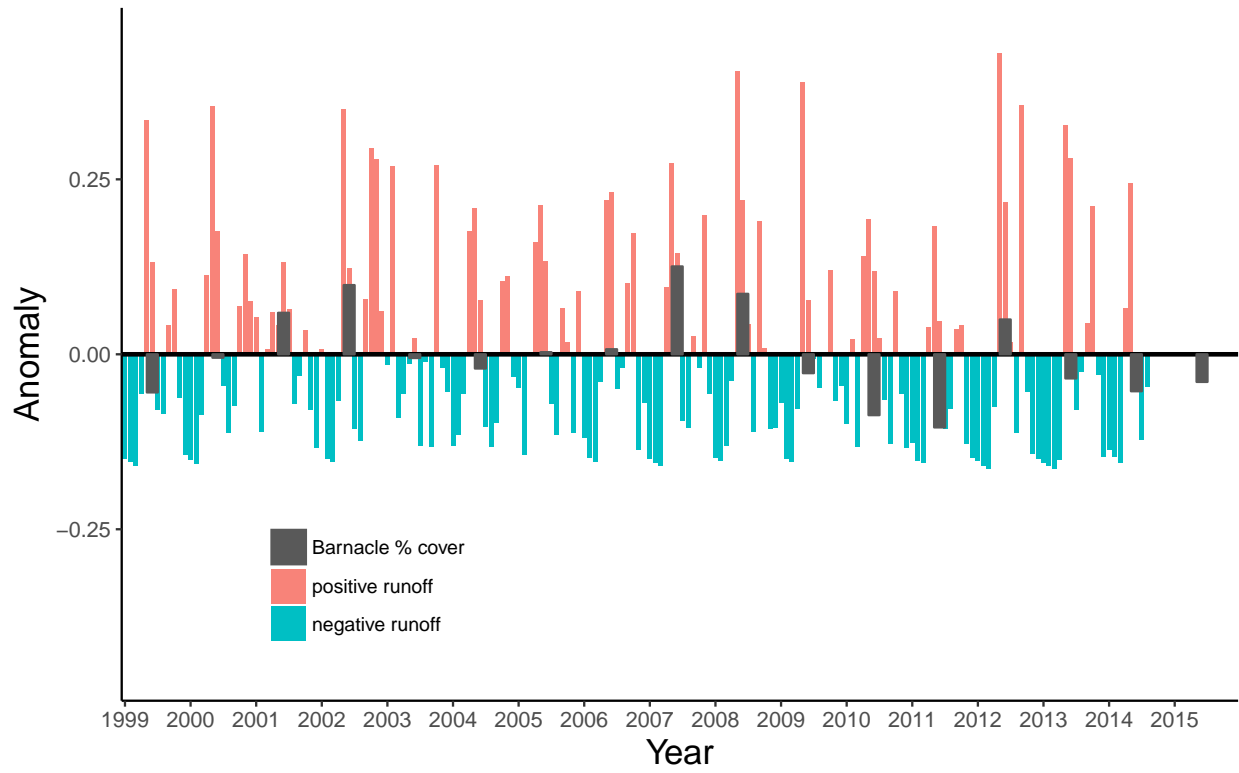
##  
## [[25]]



## Warning: Removed 2 rows containing missing values (geom\_col).

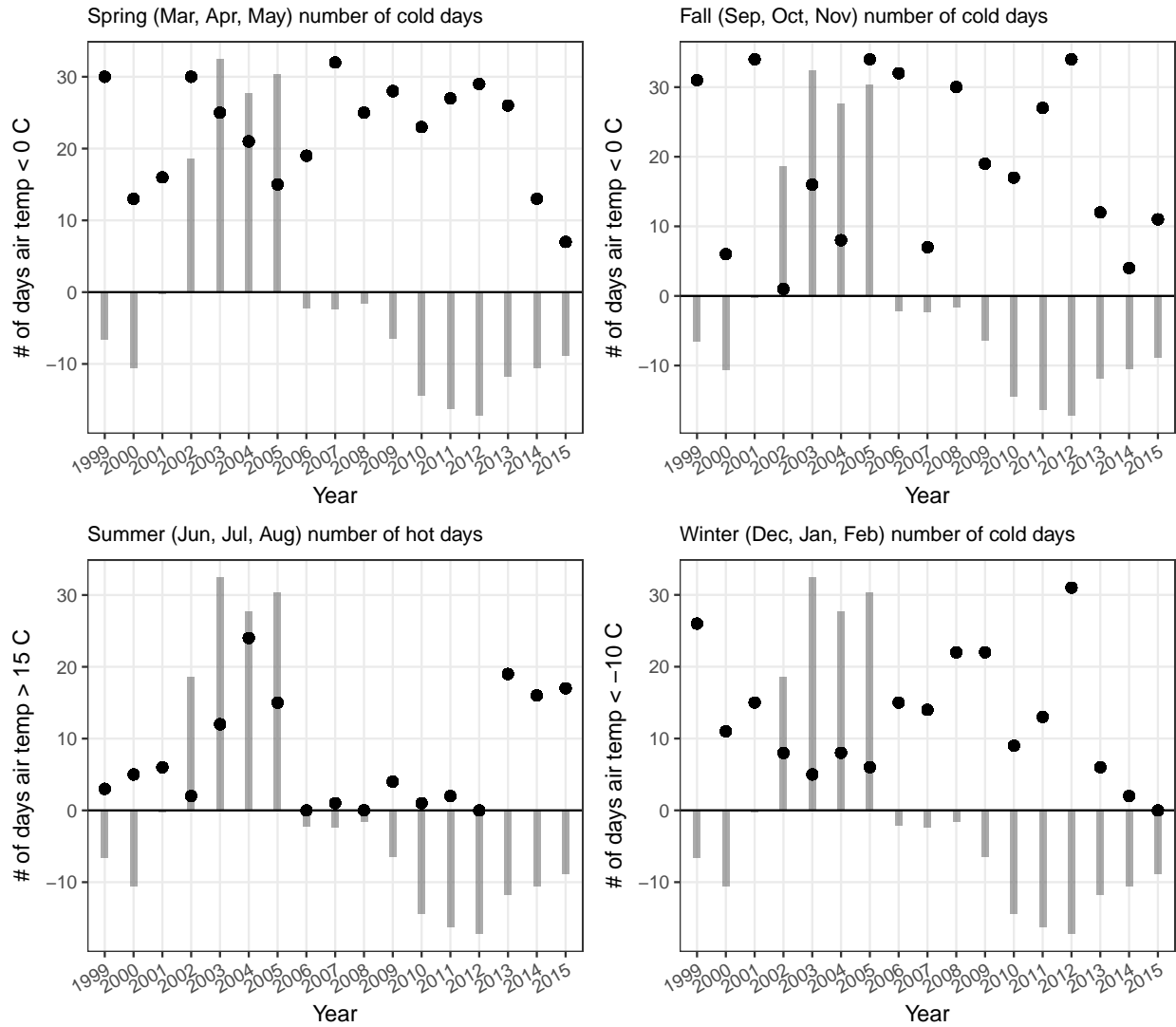




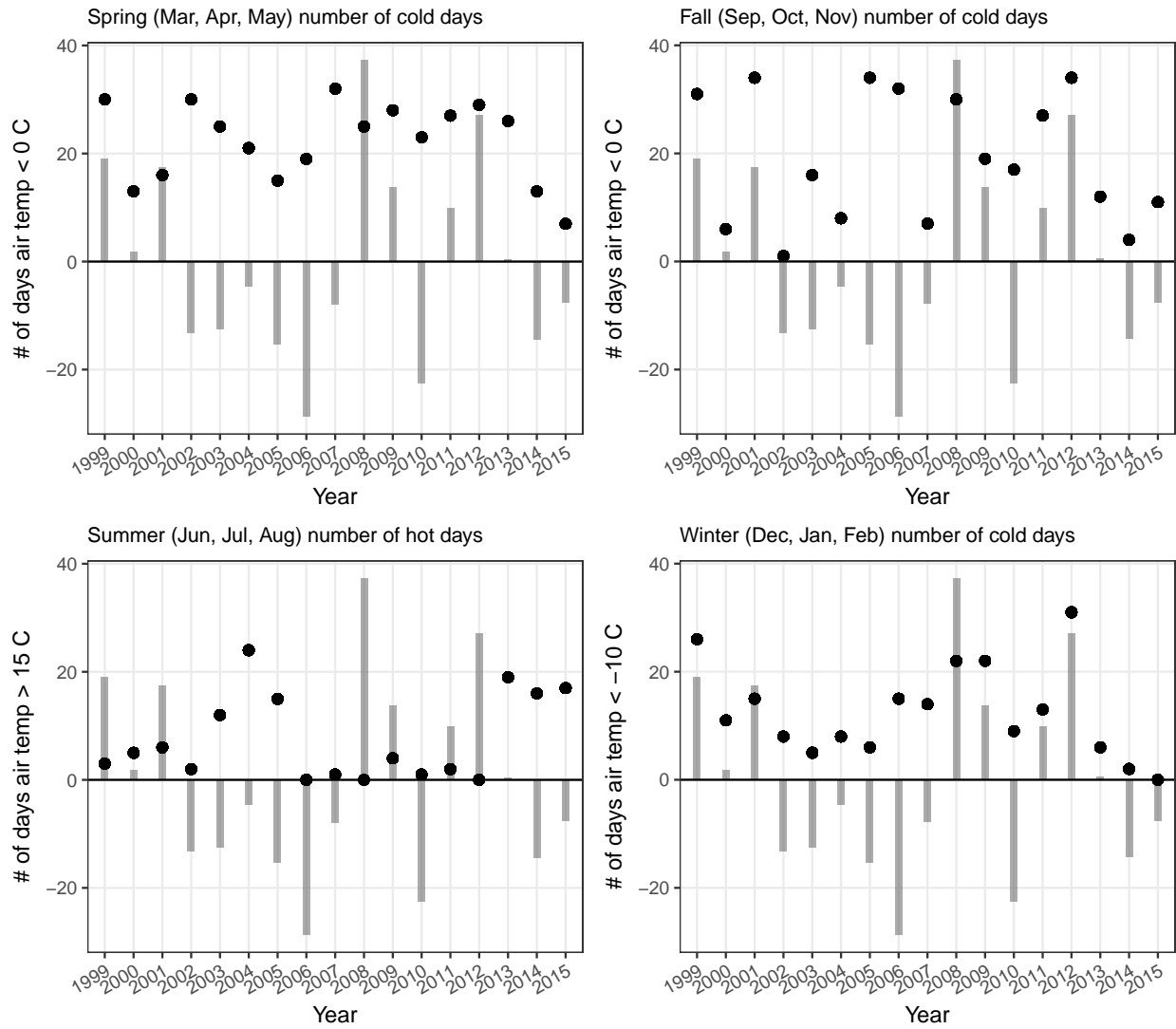




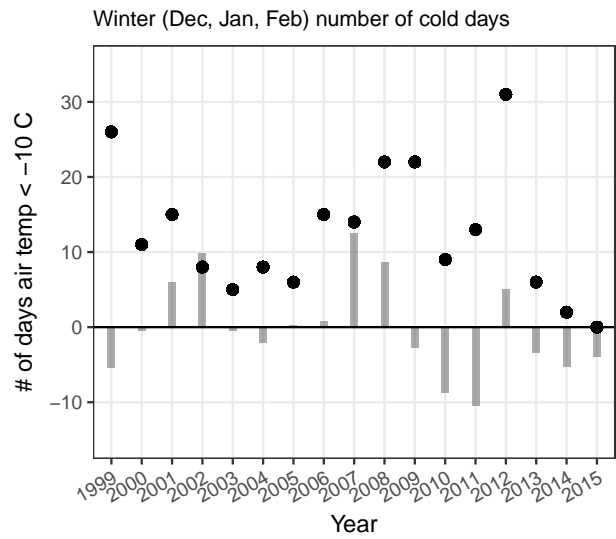
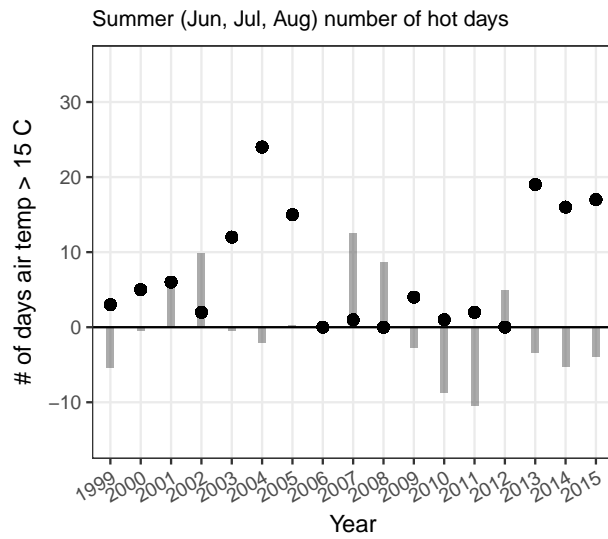
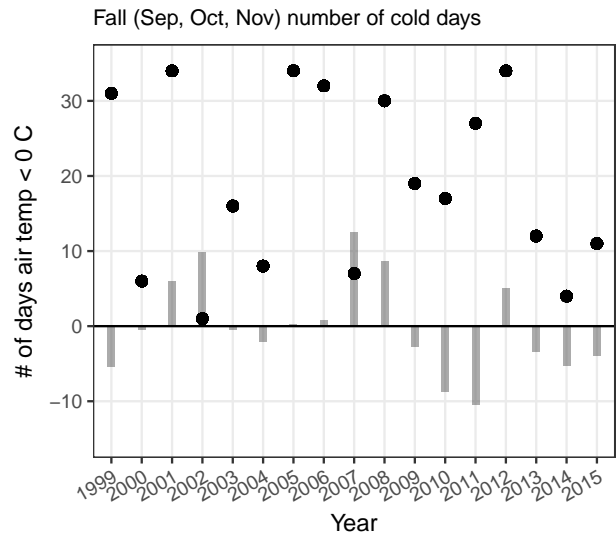
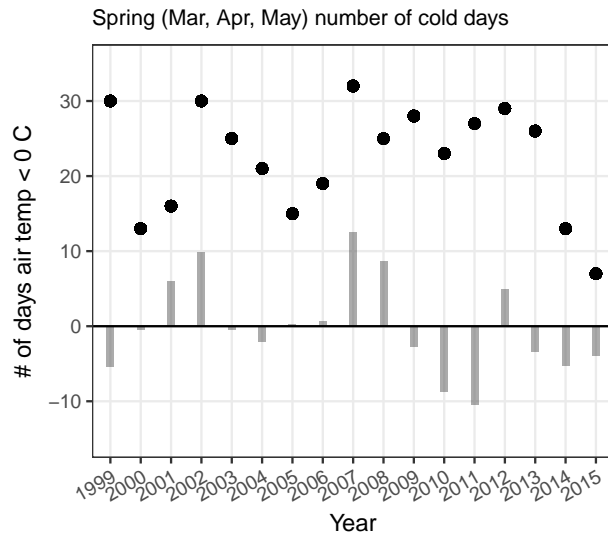
## Mytilus

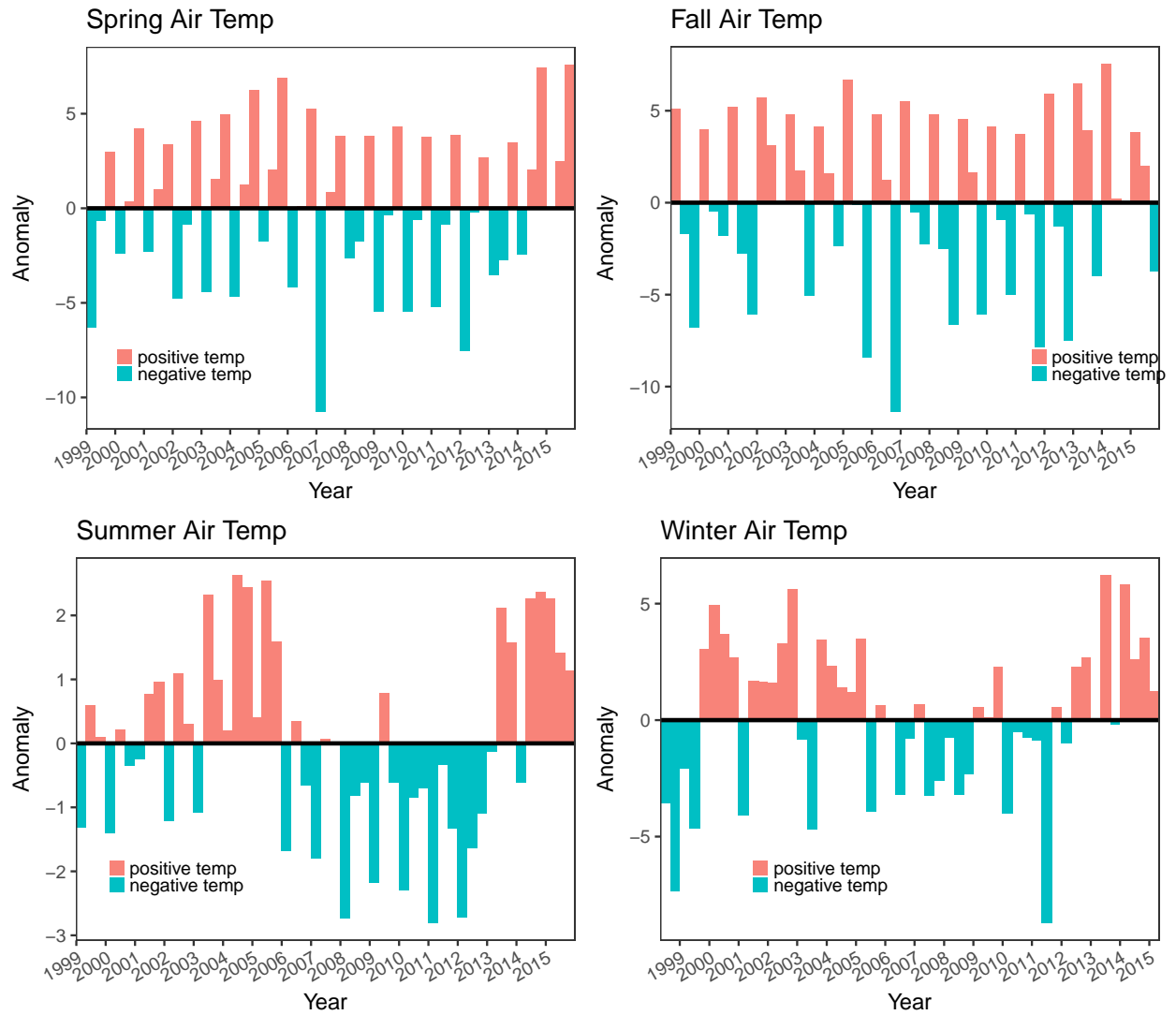


## Fucus

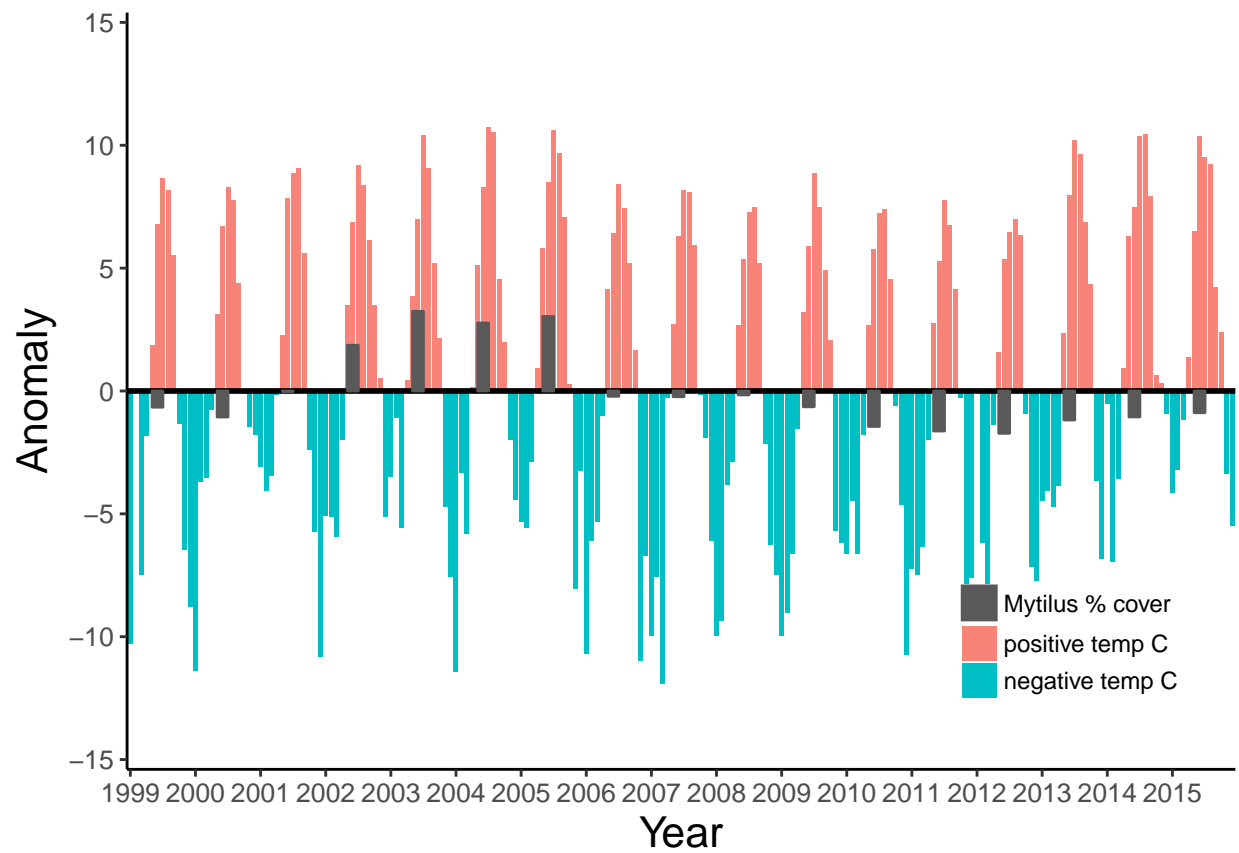


## Barnacles

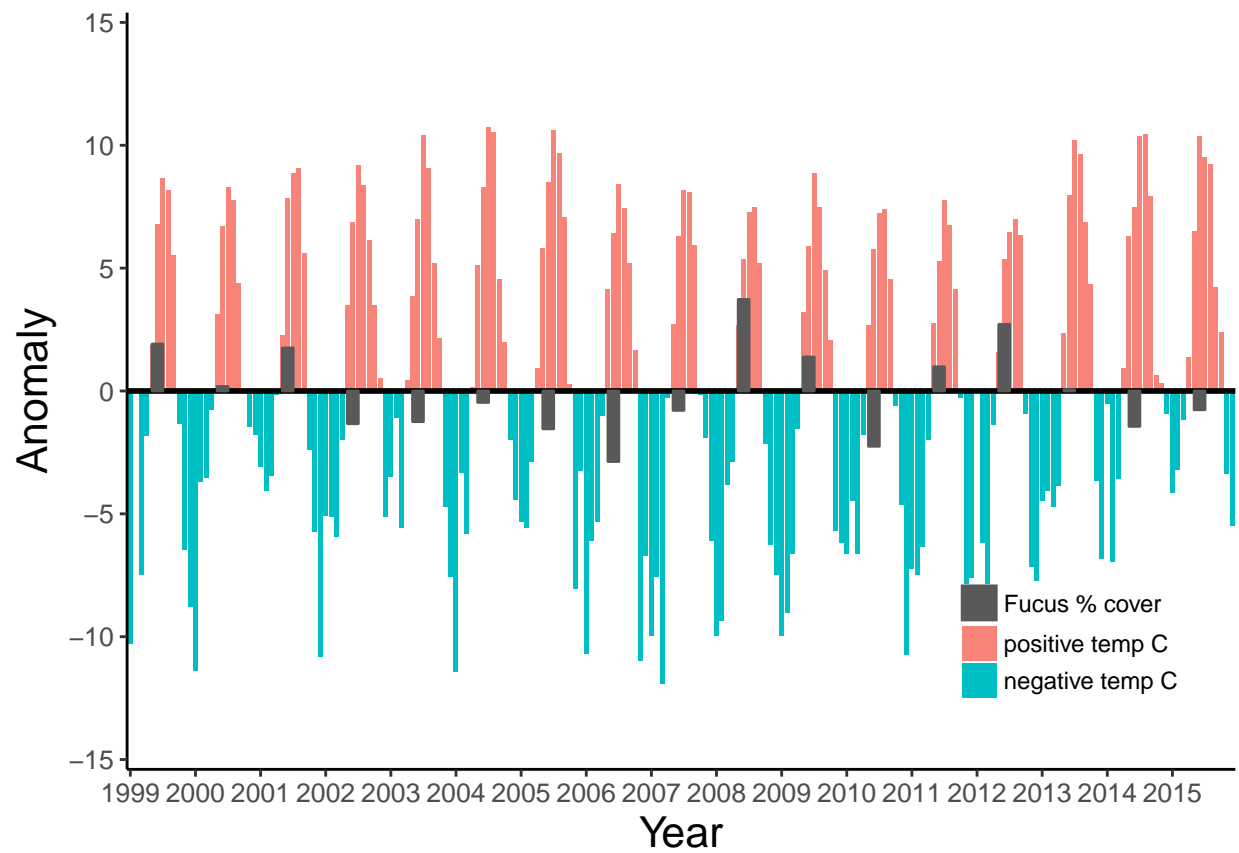




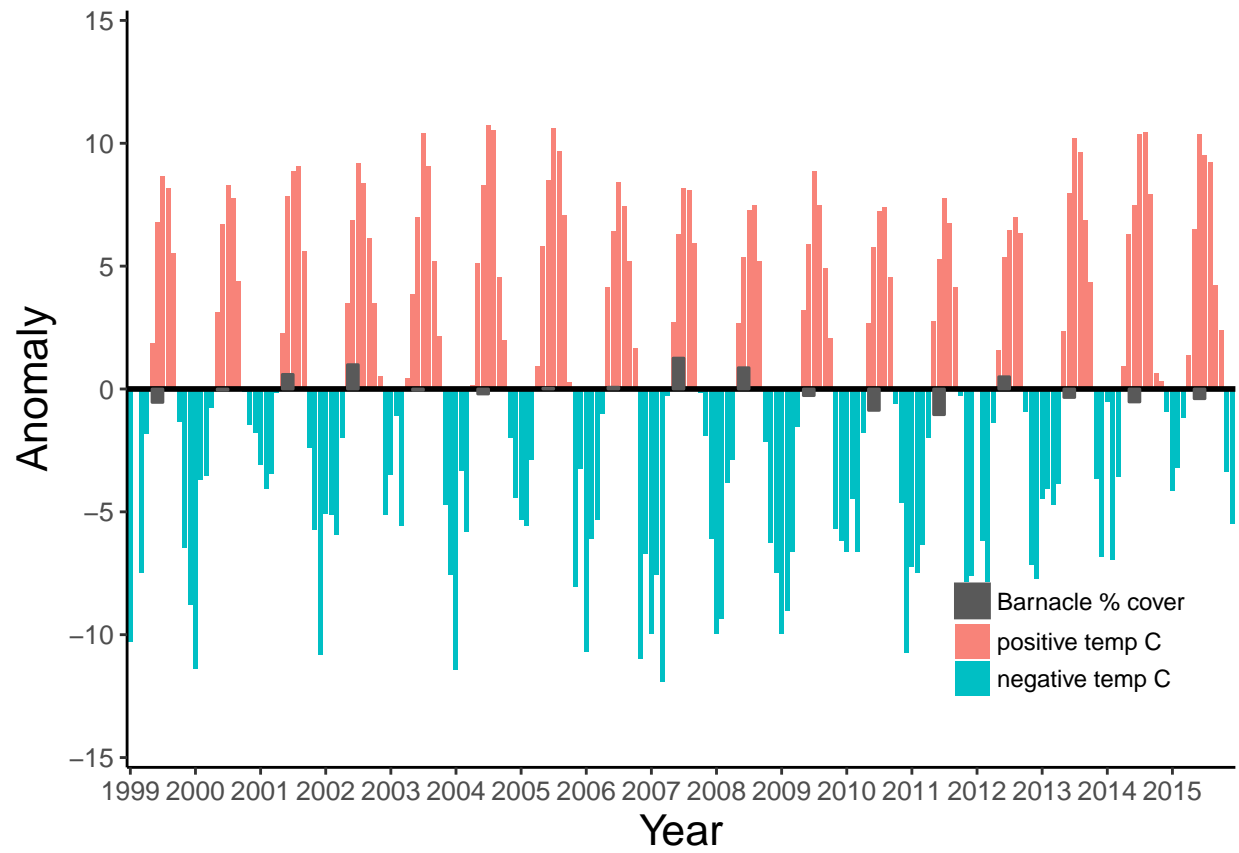
## Warning: Removed 2 rows containing missing values (position\_stack).



## Warning: Removed 2 rows containing missing values (position\_stack).

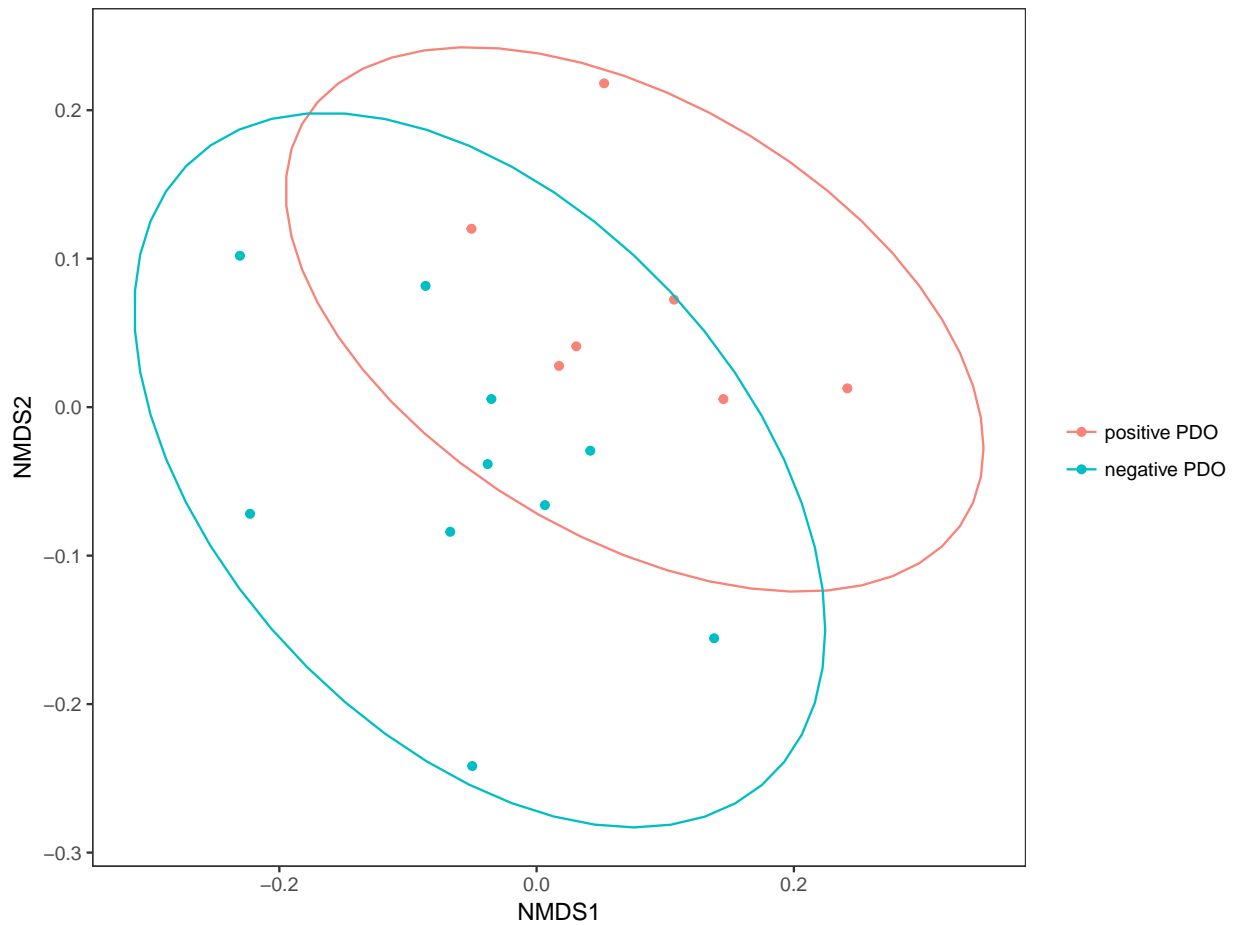


## Warning: Removed 2 rows containing missing values (position\_stack).



# NMDS Plots

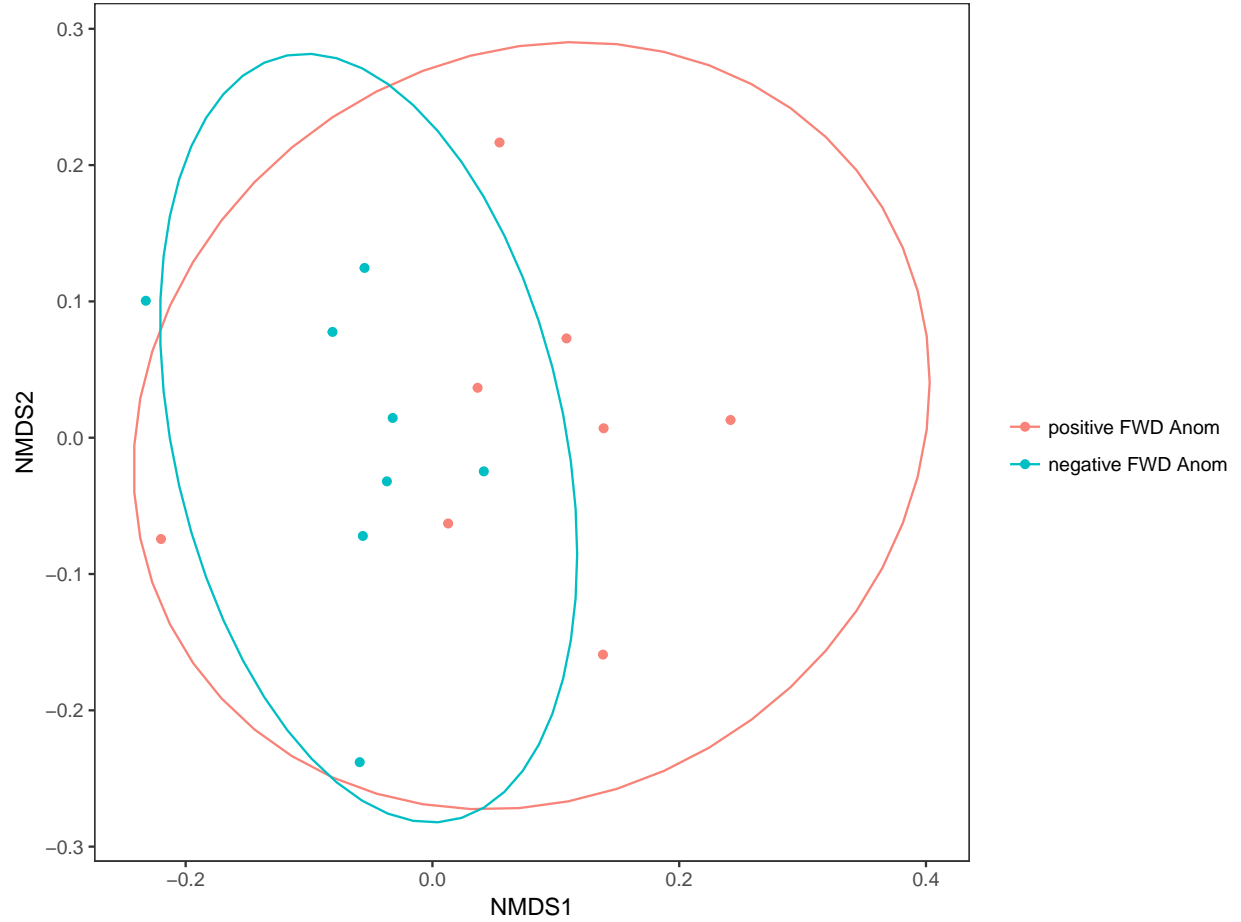
PDO



```
##
## Call:
## vegan::adonis(formula = sp_percov ~ PDO_anul_mn, data = pdo_treats,      permutations = 1000, method
##
## Permutation: free
## Number of permutations: 1000
##
## Terms added sequentially (first to last)
##
##              Df SumsOfSqs  MeanSqs F.Model    R2  Pr(>F)
## PDO_anul_mn   1   0.18366 0.183660   5.655 0.27378 0.000999 ***
## Residuals    15   0.48716 0.032477         0.72622
## Total        16   0.67082         1.00000
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

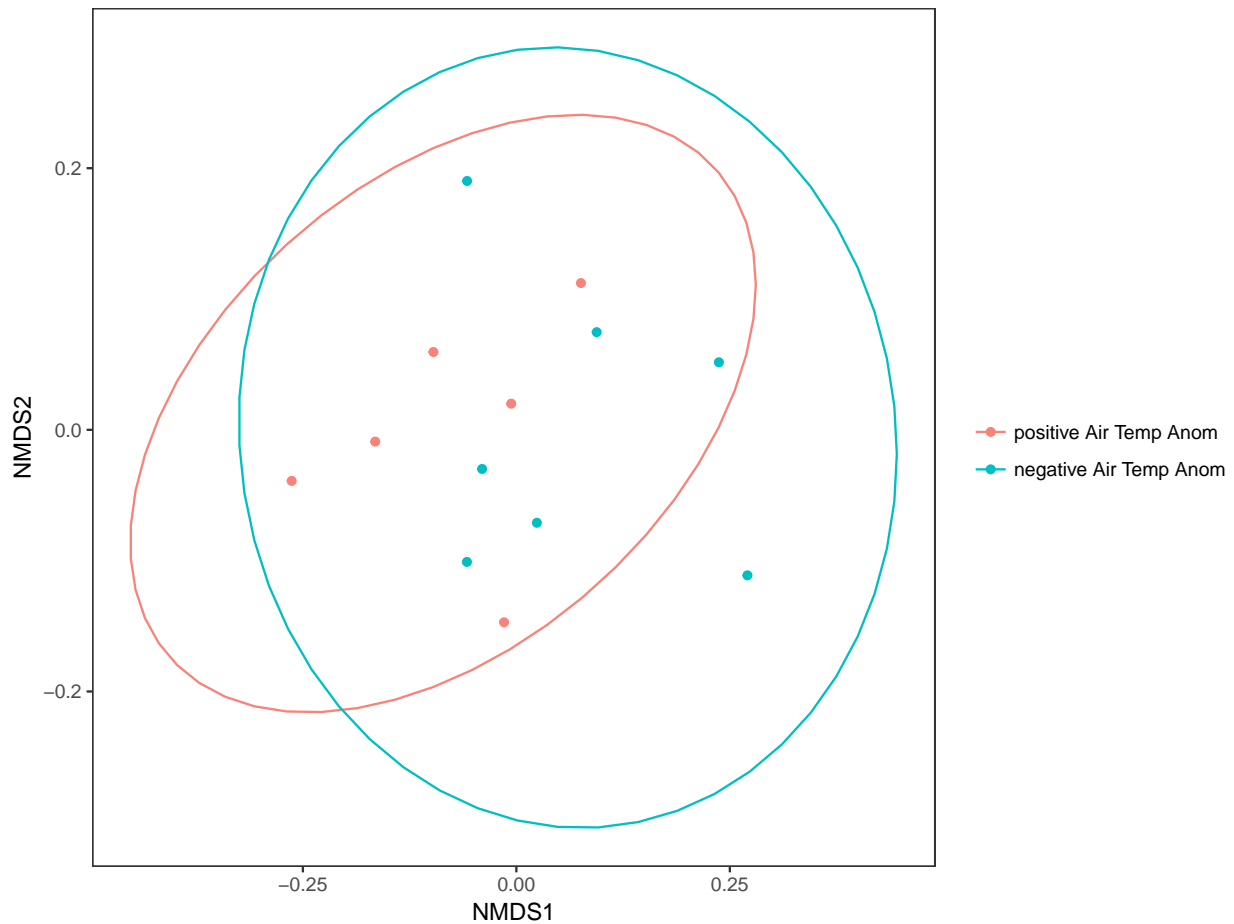


## Freshwater



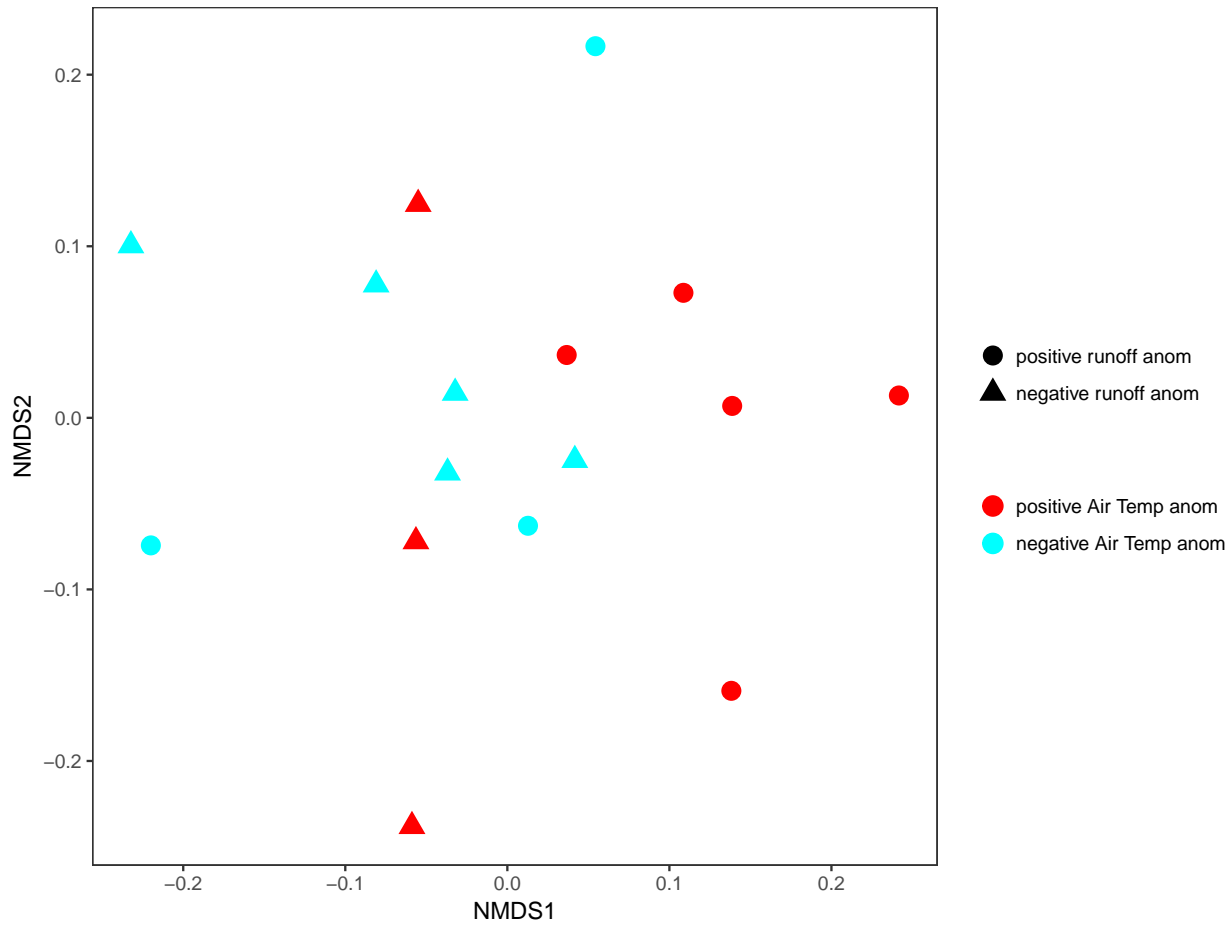
```
##
## Call:
## vegan::adonis(formula = sp_percov2 ~ mean_yearly_discharge_m3s1,      data = fresh_treats, permutati
##
## Permutation: free
## Number of permutations: 1000
##
## Terms added sequentially (first to last)
##
##              Df SumsOfSqs  MeanSqs F.Model    R2  Pr(>F)
## mean_yearly_discharge_m3s1  1   0.10794 0.107936  2.7347 0.16342 0.06094 .
## Residuals                14   0.55256 0.039469           0.83658
## Total                    15   0.66050           1.00000
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## Air Temp



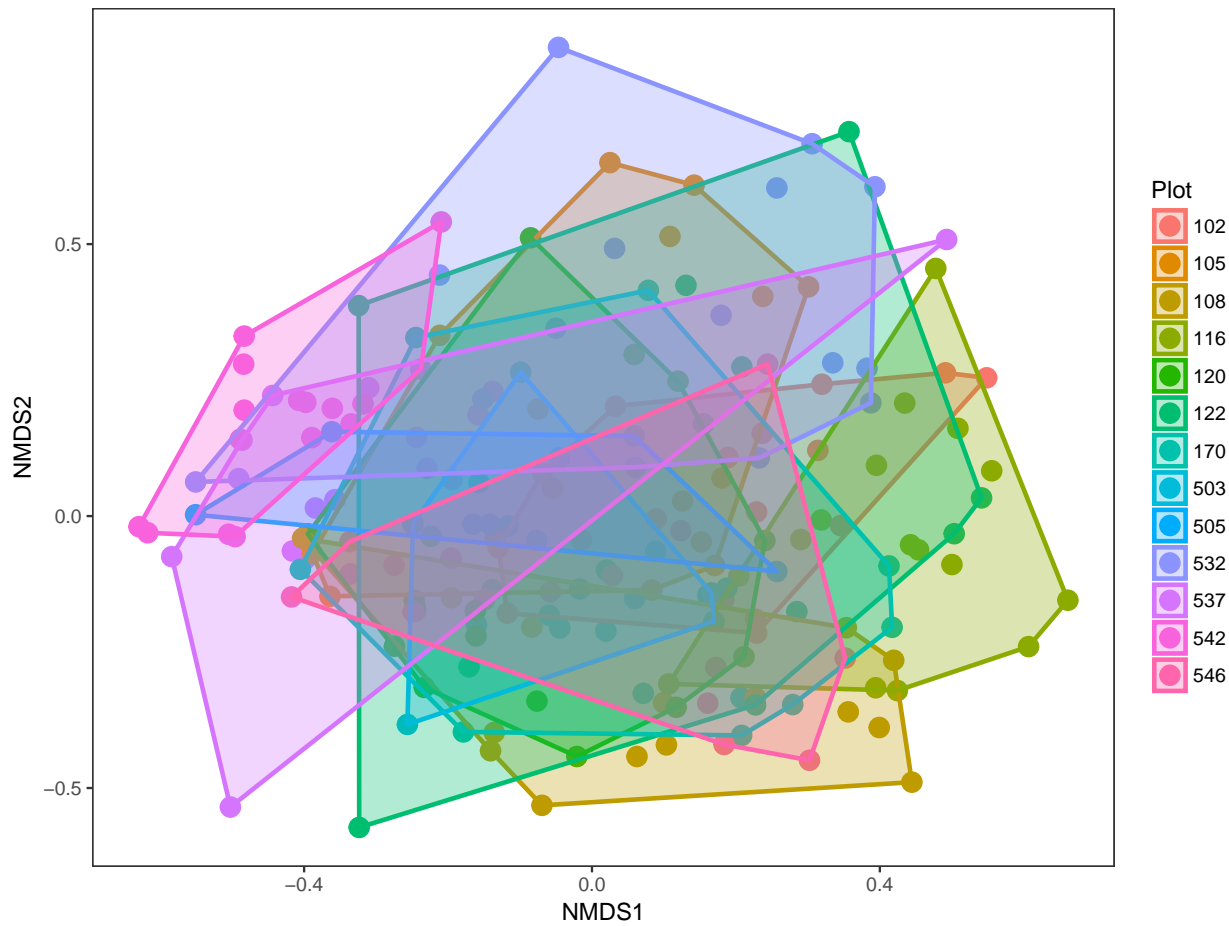
```
##
## Call:
## vegan::adonis(formula = sp_percov3 ~ ATemp_YearMn, data = atemp_treats,      permutations = 1000, me
##
## Permutation: free
## Number of permutations: 1000
##
## Terms added sequentially (first to last)
##
##              Df SumsOfSqs  MeanSqs F.Model    R2  Pr(>F)
## ATemp_YearMn  1   0.12079 0.120791  2.9682 0.2125 0.03297 *
## Residuals    11   0.44764 0.040695         0.7875
## Total        12   0.56843          1.0000
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

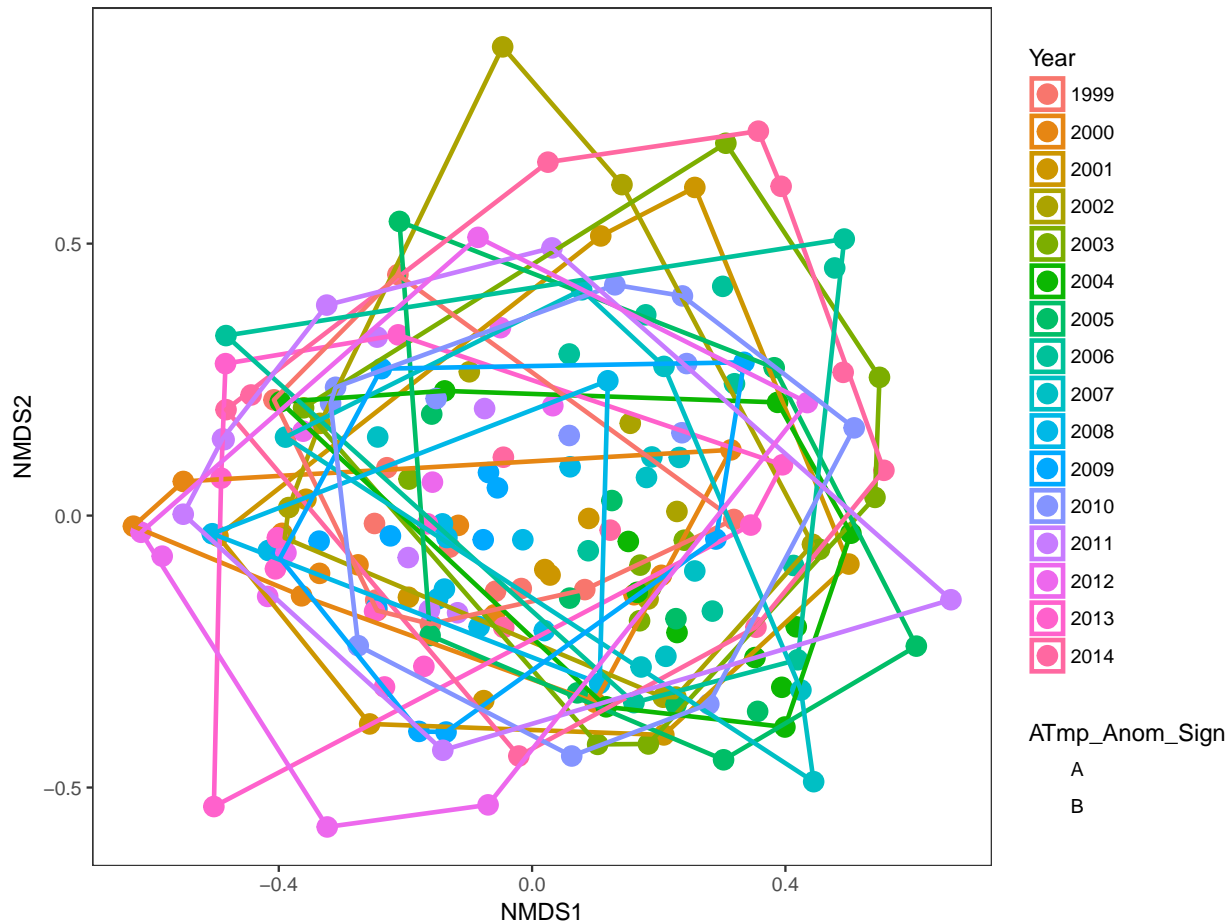
All env. variables



```
##
## Call:
## vegan::adonis(formula = sp_percov4 ~ ATemp_YearMn + mn_yr_discharge,      data = all_treats, permuta
##
## Permutation: free
## Number of permutations: 1000
##
## Terms added sequentially (first to last)
##
##              Df SumsOfSqs  MeanSqs F.Model    R2  Pr(>F)
## ATemp_YearMn   1  0.15102 0.151023  4.5478 0.22865 0.004995 **
## mn_yr_discharge 1  0.07778 0.077779  2.3422 0.11776 0.090909 .
## Residuals     13  0.43170 0.033207          0.65359
## Total         15  0.66050              1.00000
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

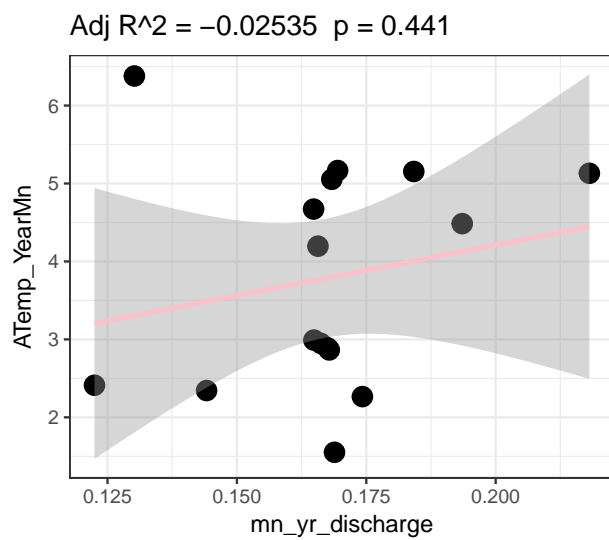
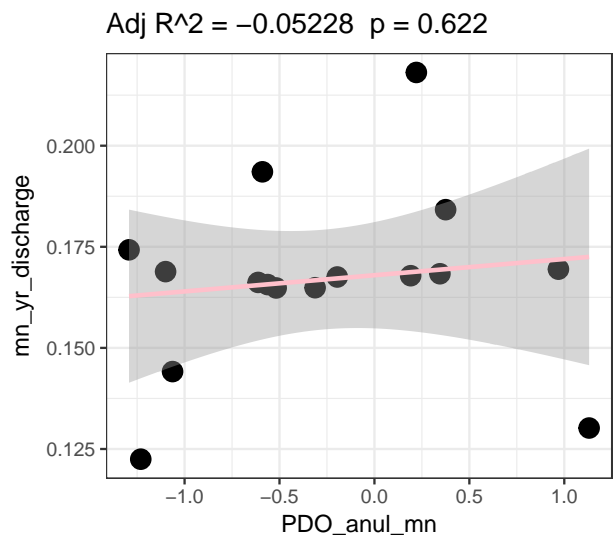
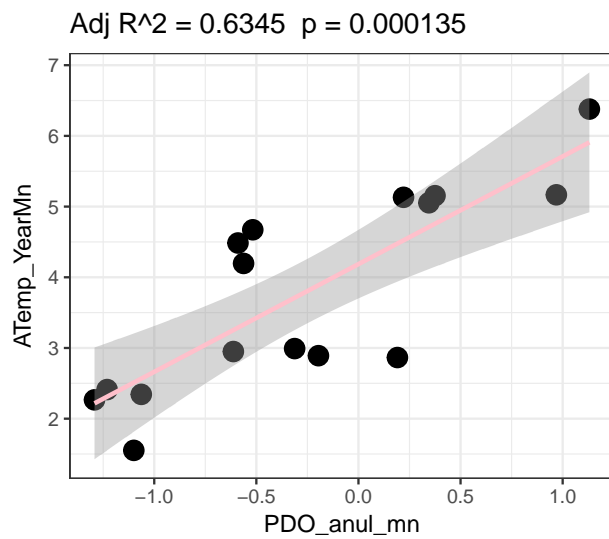
Biological NMDS



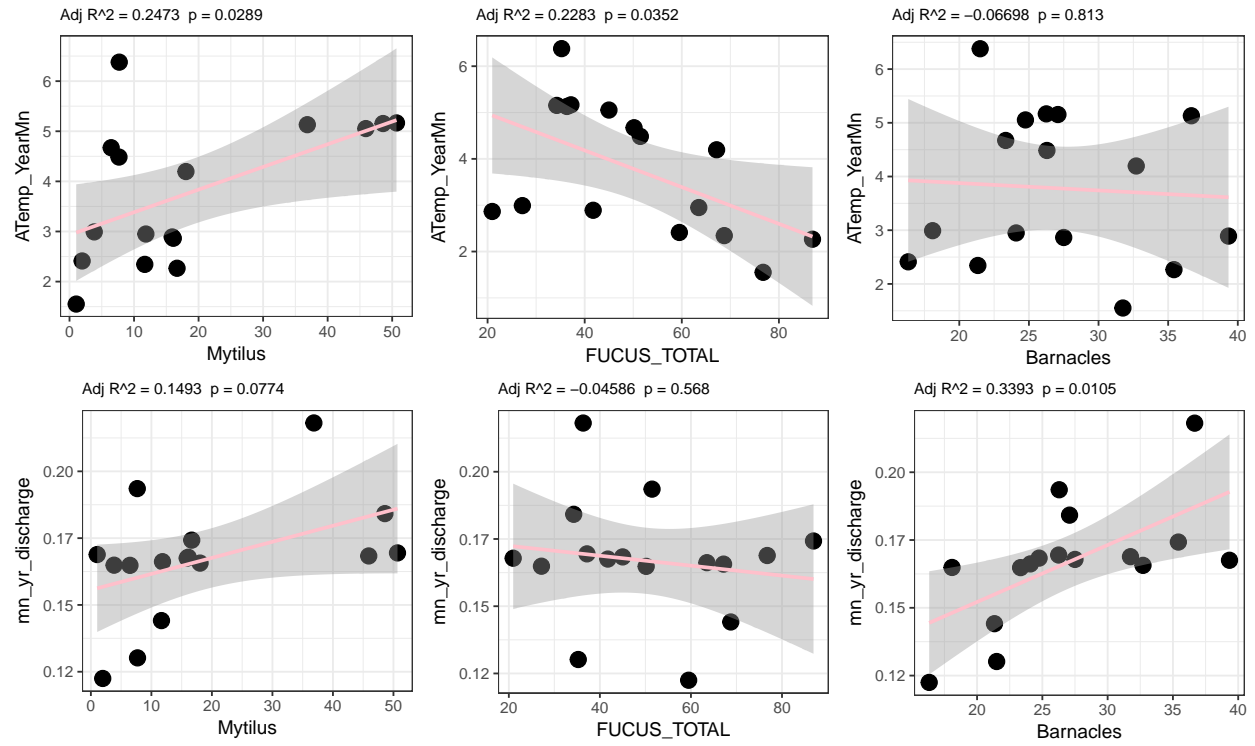


```
##
## Call:
## vegan::adonis(formula = sp_percov5 ~ ATemp_YearMn * mn_yr_discharge, data = all_treats2, permut
##
## Permutation: free
## Number of permutations: 1000
##
## Terms added sequentially (first to last)
##
##              Df SumsOfSqs MeanSqs F.Model    R2
## ATemp_YearMn    1   1.4209  1.42087 12.3037 0.05888
## mn_yr_discharge  1   0.6750  0.67500   5.8450 0.02797
## ATemp_YearMn:mn_yr_discharge  1   0.3240  0.32396   2.8053 0.01343
## Residuals     188  21.7109  0.11548         0.89972
## Total         191  24.1308         1.00000
##
##              Pr(>F)
## ATemp_YearMn    0.000999 ***
## mn_yr_discharge 0.000999 ***
## ATemp_YearMn:mn_yr_discharge 0.033966 *
## Residuals
## Total
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## Correlations between env. variables



## Correlations between other variables



## MVABUND Results