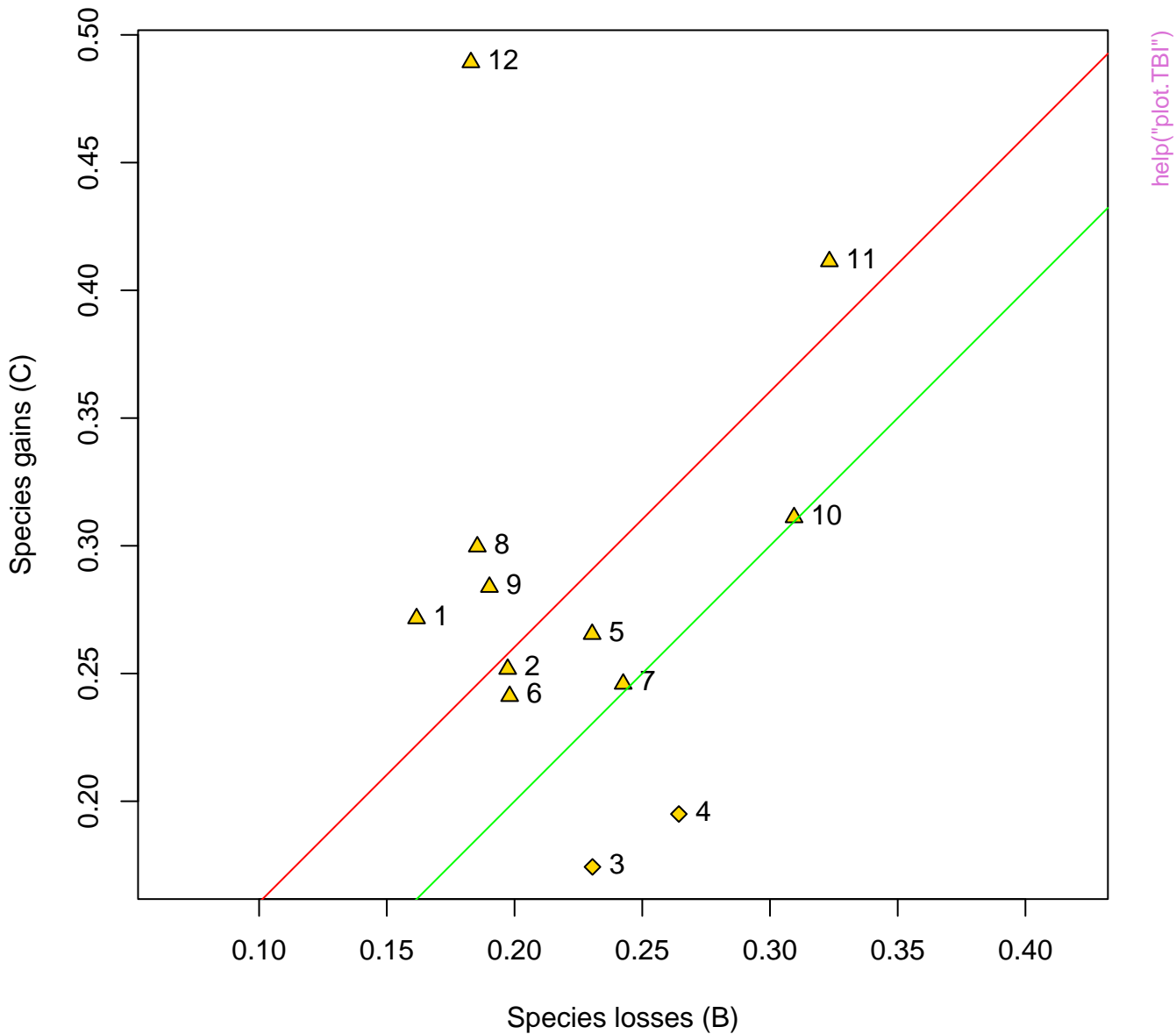
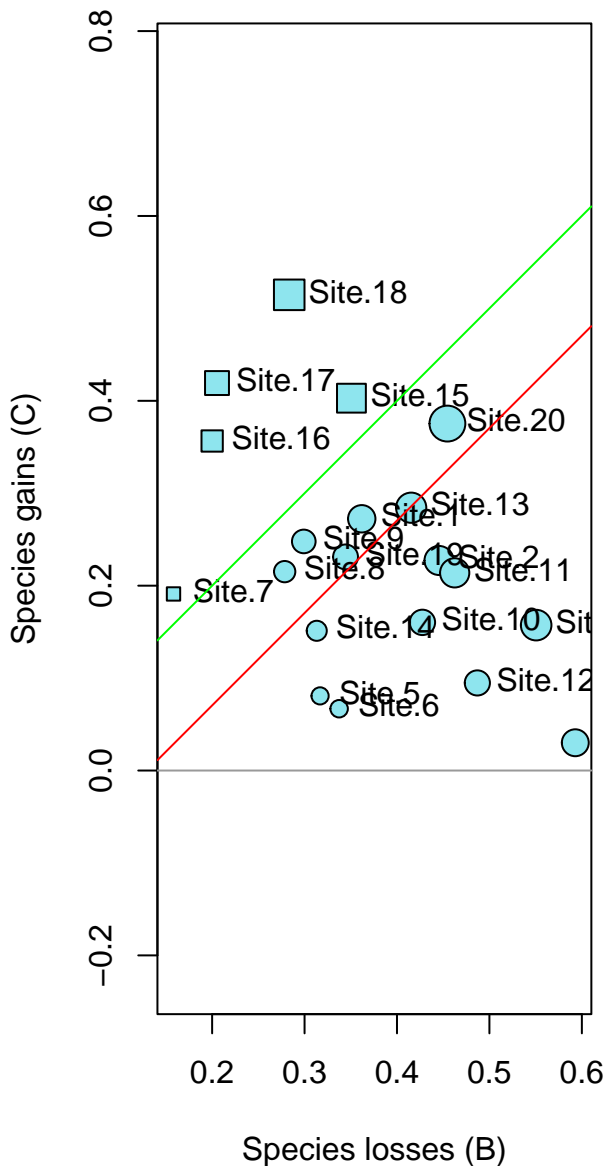


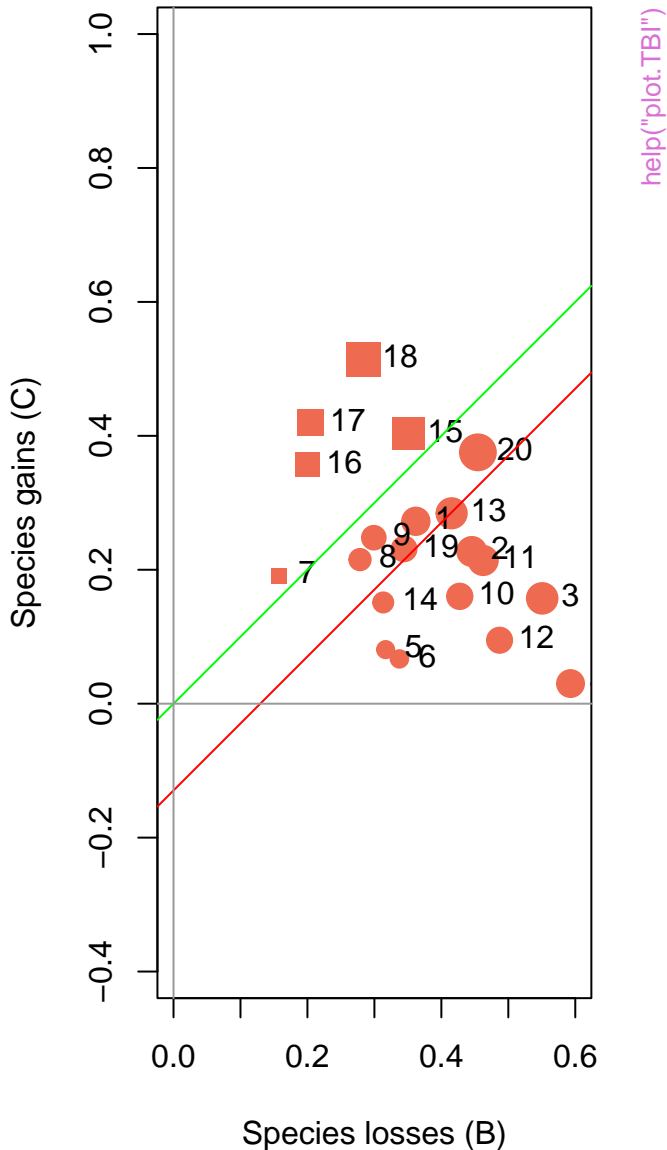
**B-C plot, Pyrifos, surveys 4 & 11**



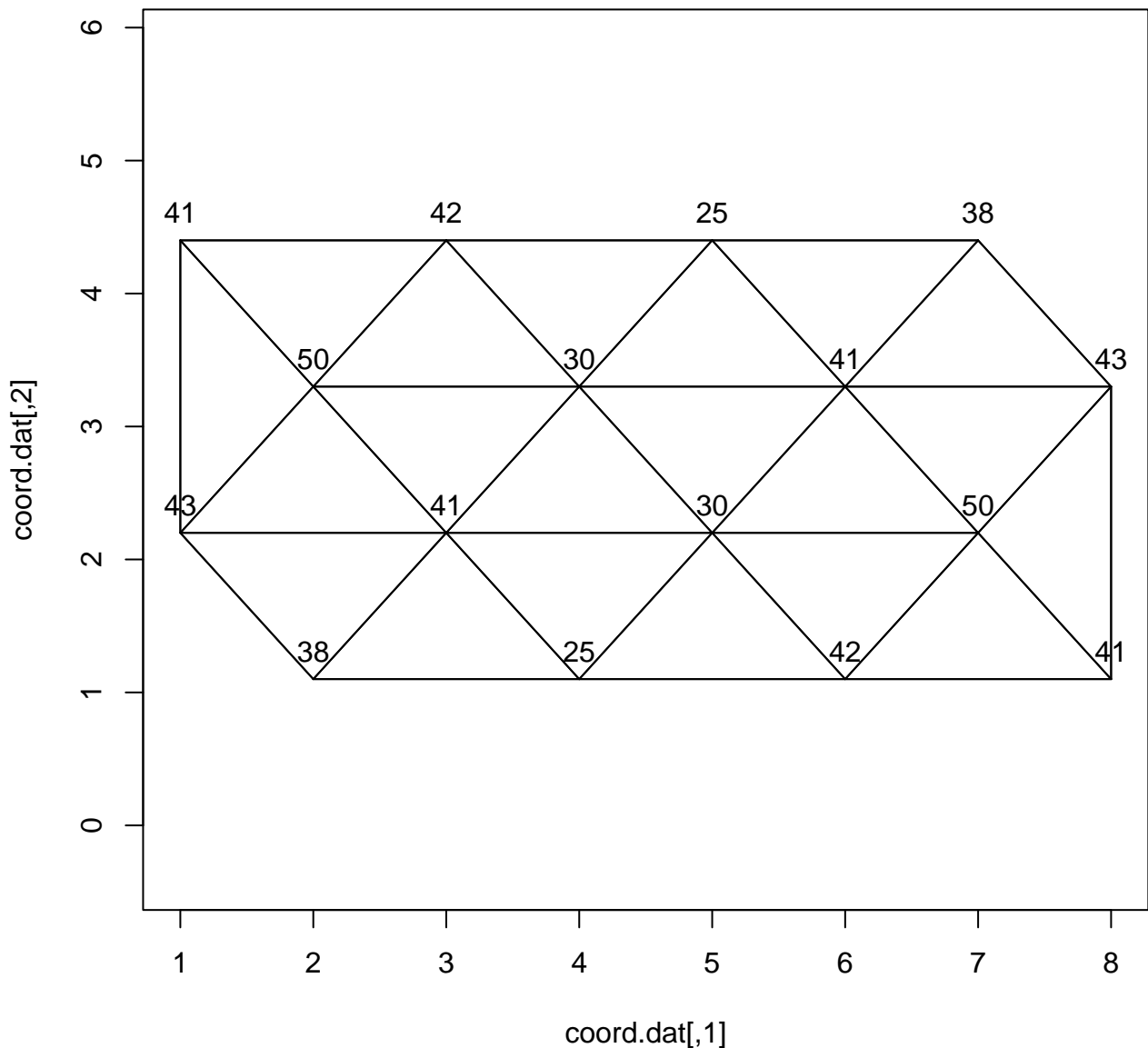
**B-C plot, Mite data**



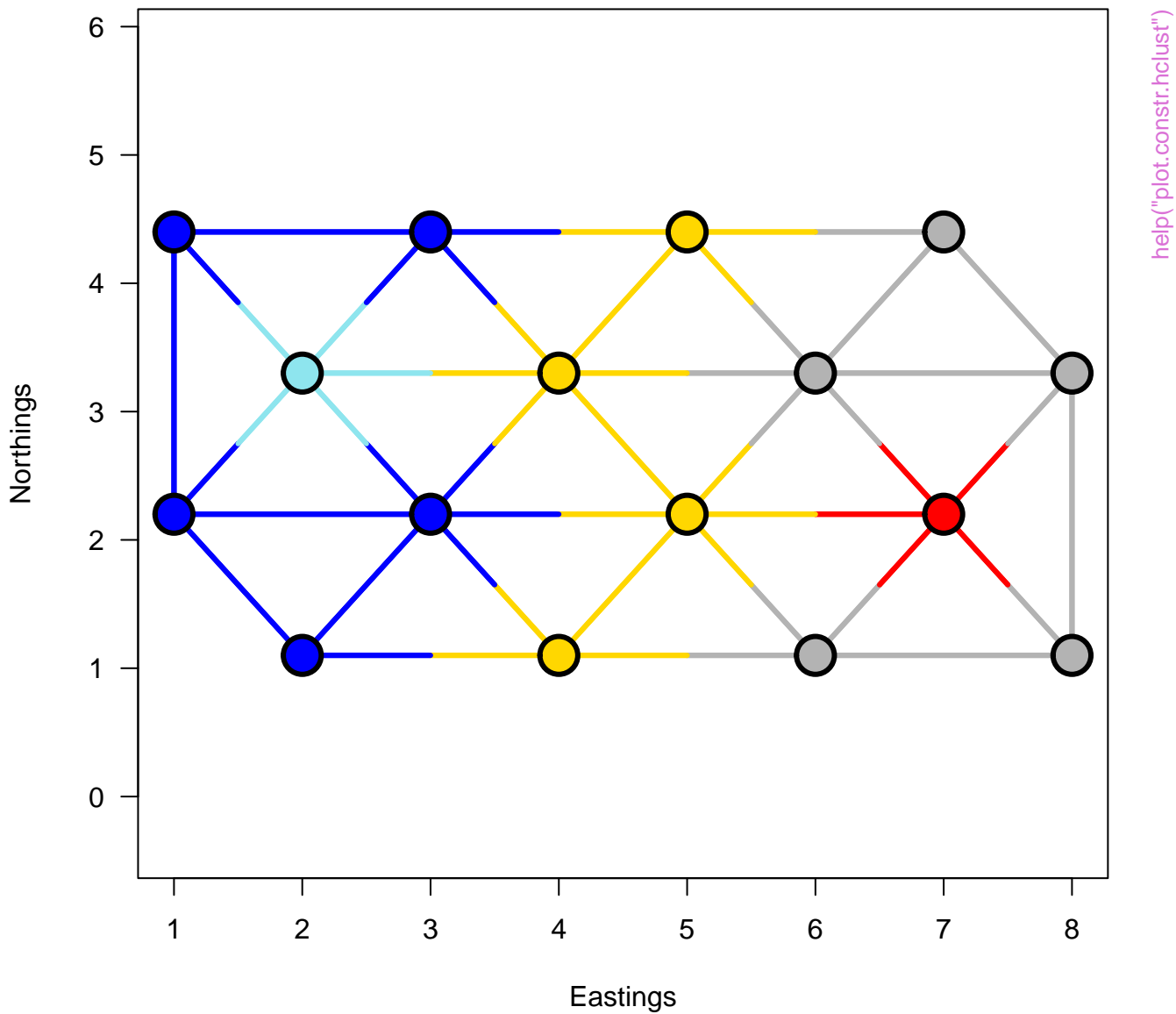
**B-C plot, Mite data**



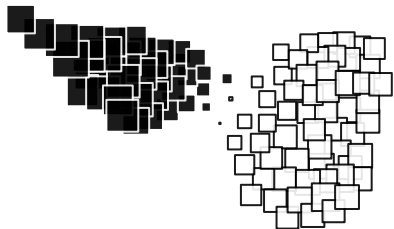
## Delaunay triangulation



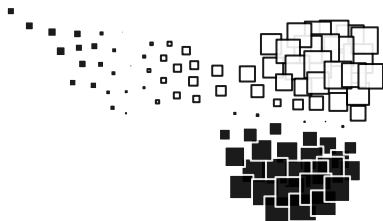
```
help("plot.constr.hclust")
```



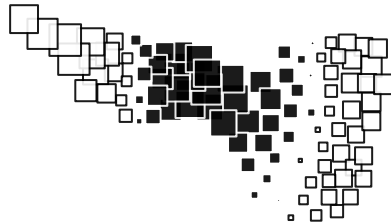
MEM1



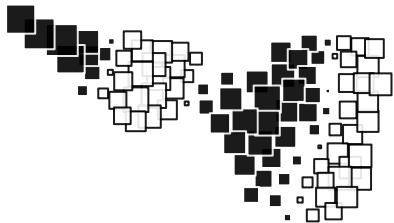
MEM2



MEM3



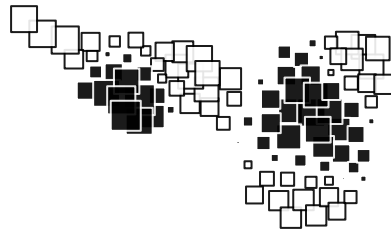
MEM4



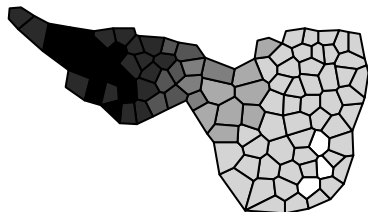
MEM5



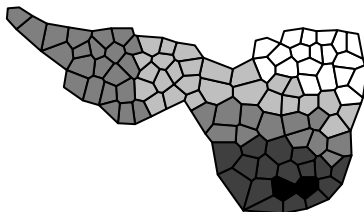
MEM6



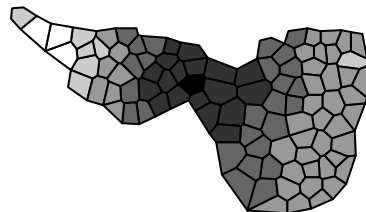
MEM1



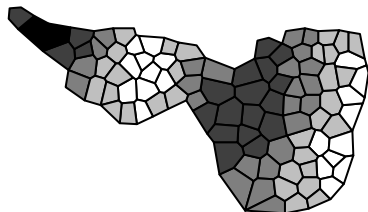
MEM2



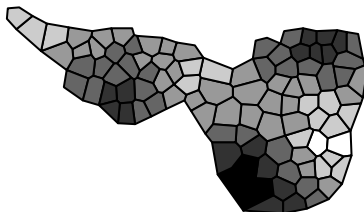
MEM3



MEM4



MEM5



MEM6

