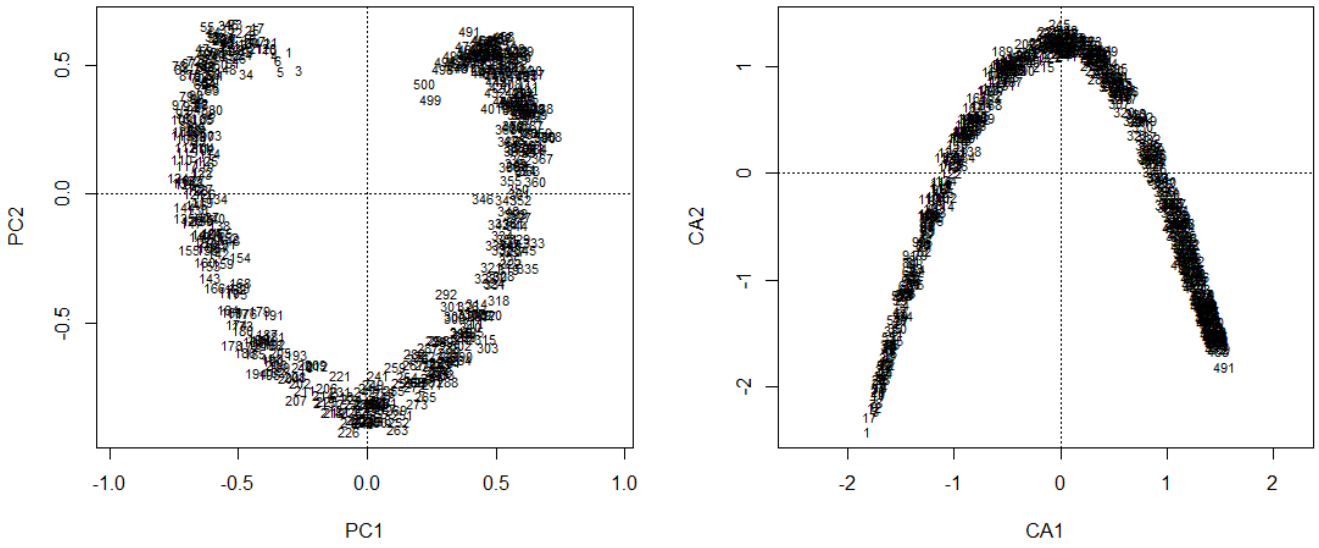
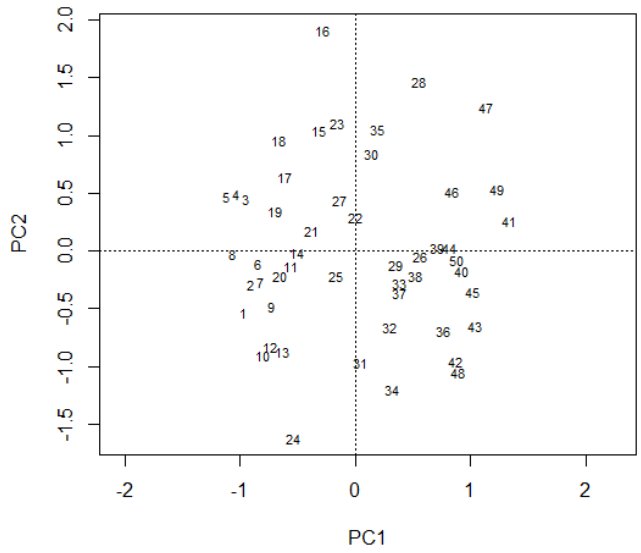
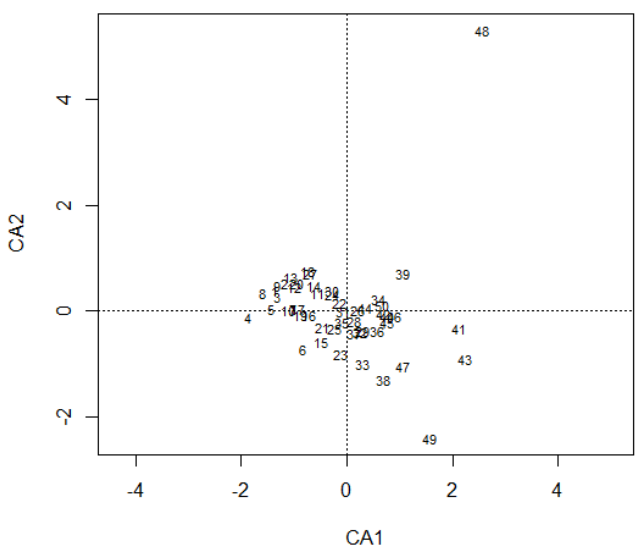
Řešení cvičení 1

```
library (vegan)
simul1.spe <- read.delim
('http://www.davidzeleny.net/anadat-r/data-download/simul1-spe.txt', row.names = 1)
PCA <- rda (simul1.spe)
CA <- cca (simul1.spe)
par (mfrow = c(1,2))
ordiplot (PCA)
ordiplot (CA)
```




```
PCA.short <- rda (simul1.spe[1:50,])
CA.short <- cca (simul1.spe [1:50,])
par (mfrow = c(1,2))
ordiplot (PCA.short)
ordiplot (CA.short)
```





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https://anadat-r.davidzeleny.net/ - **Analysis of community ecology data in R**

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