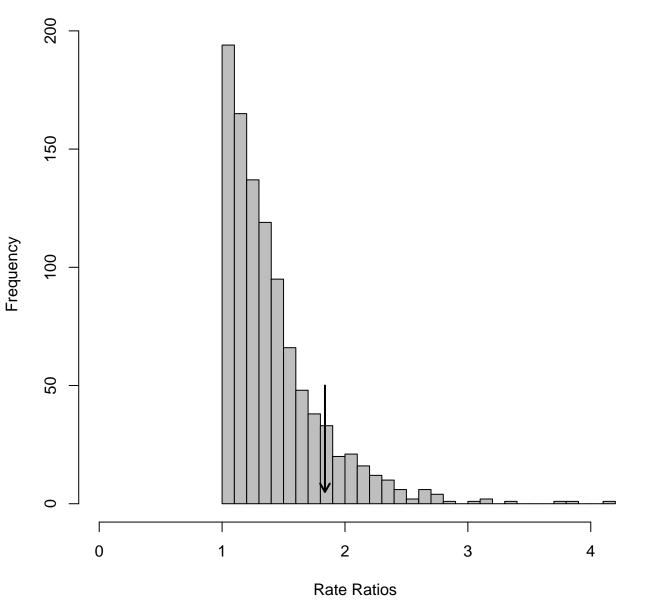
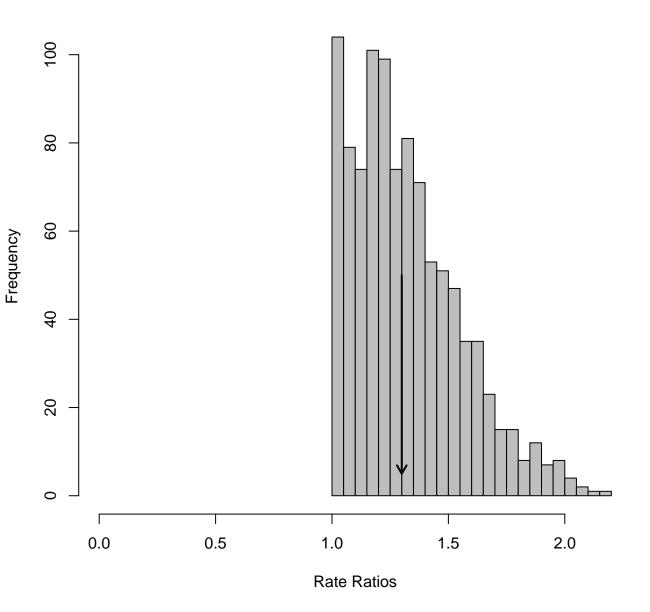
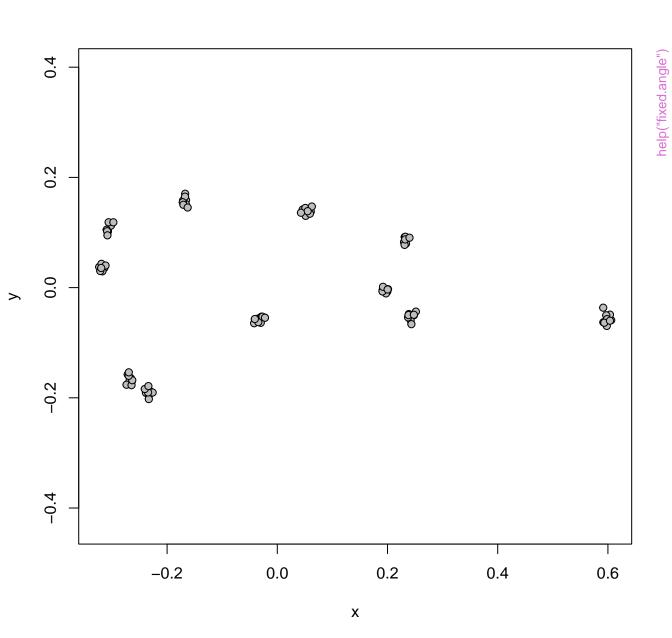
Observed Rate Ratio = 1.8372 ; P-value = 0.124

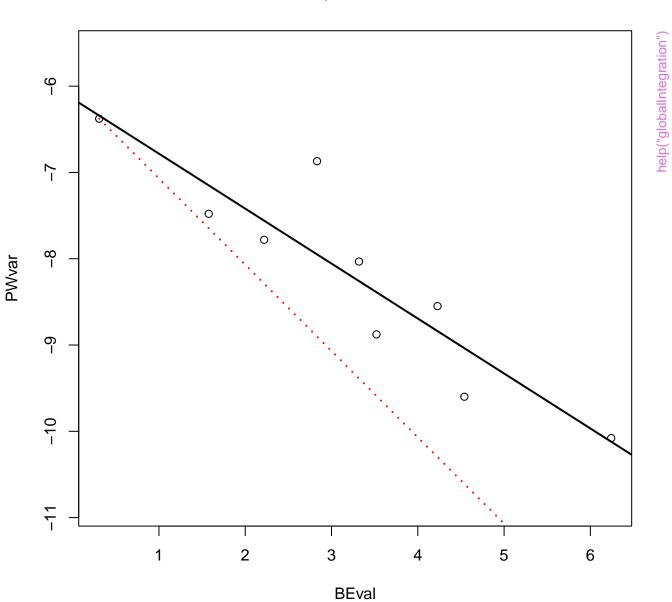




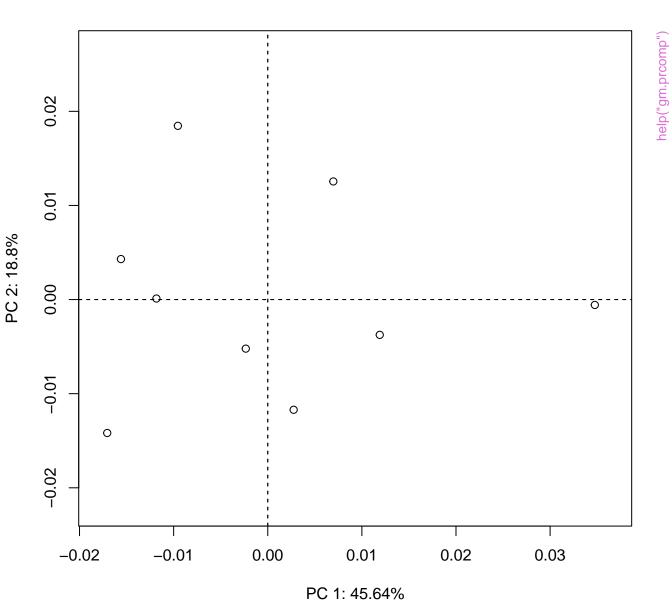
Χ



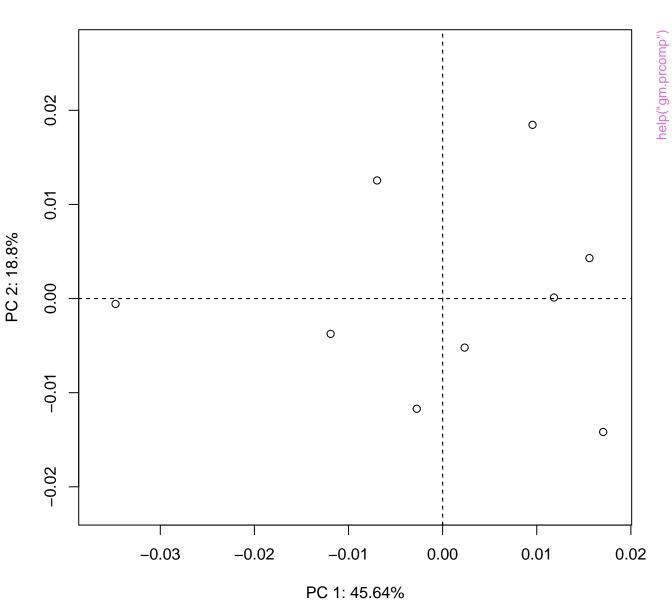
# $ObservedSlope_{black} = -0.6369198$



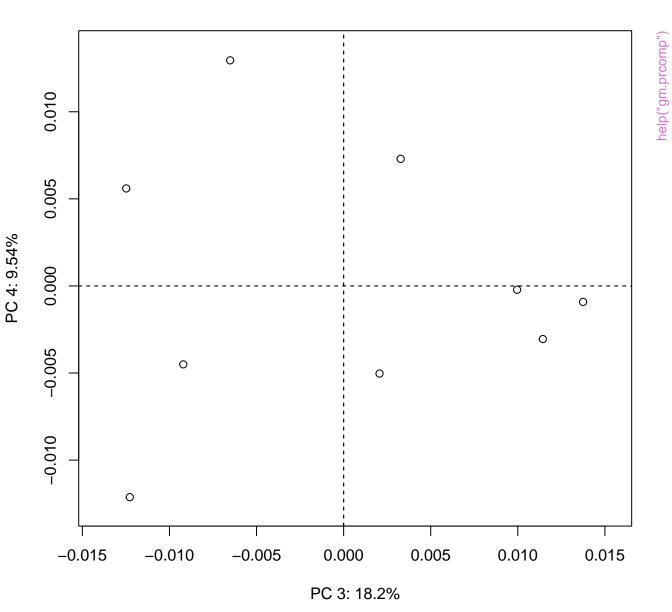




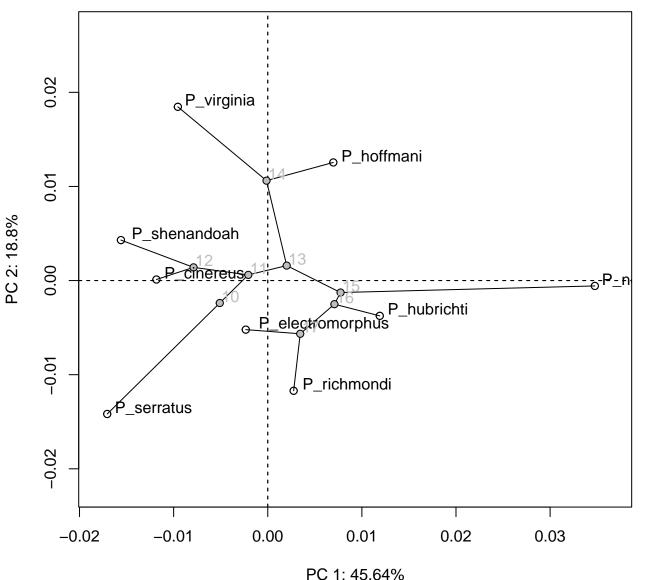




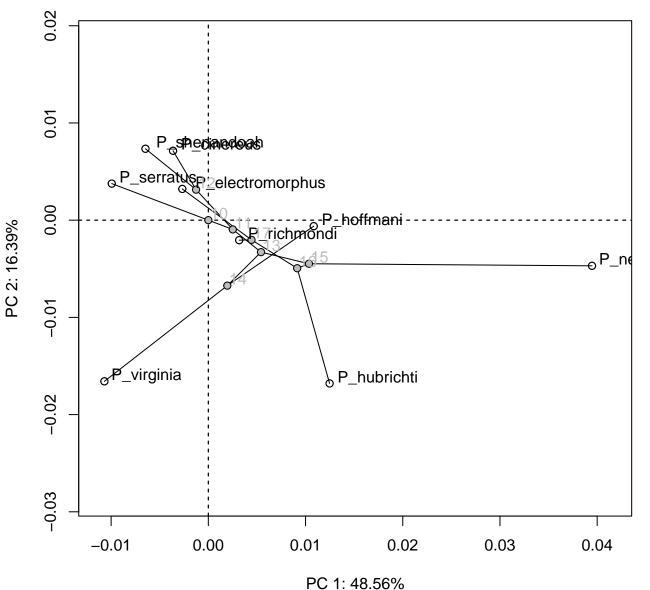




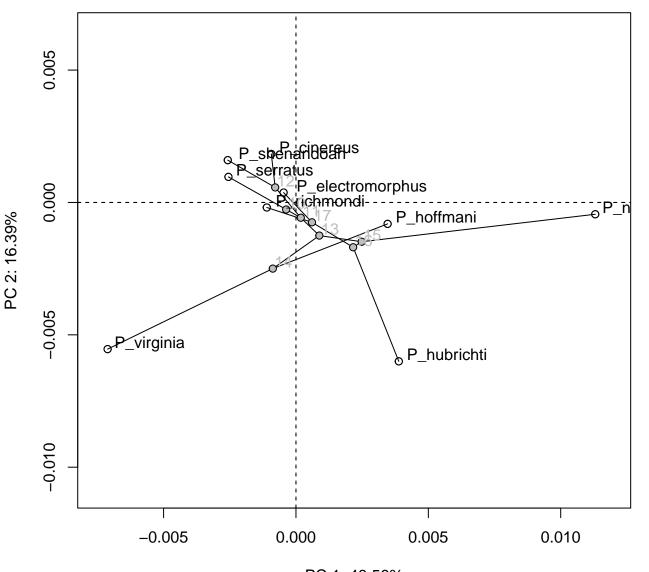
#### PCA.w.phylo



#### phylo PCA

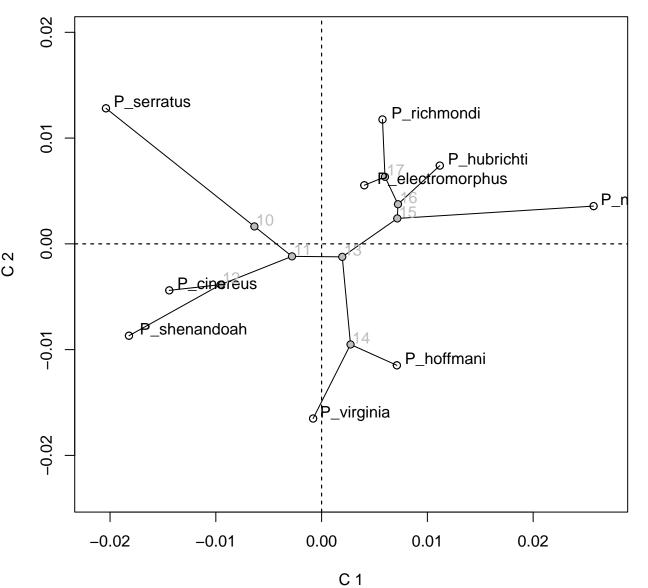


#### phylo PCA

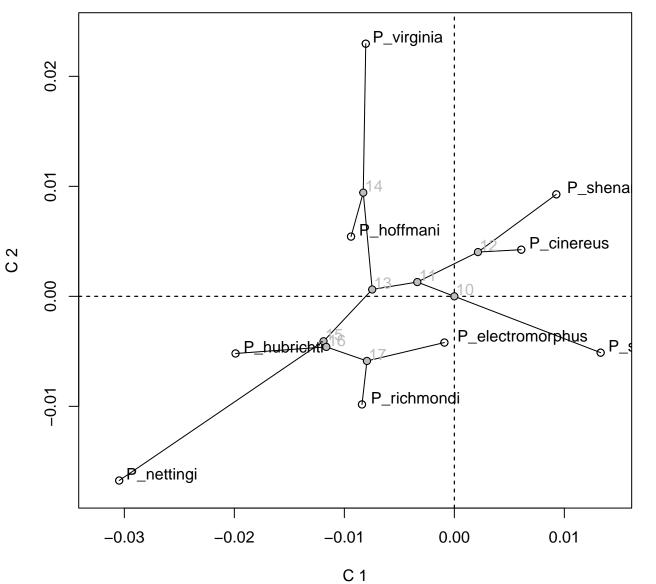


PC 1: 48.56%

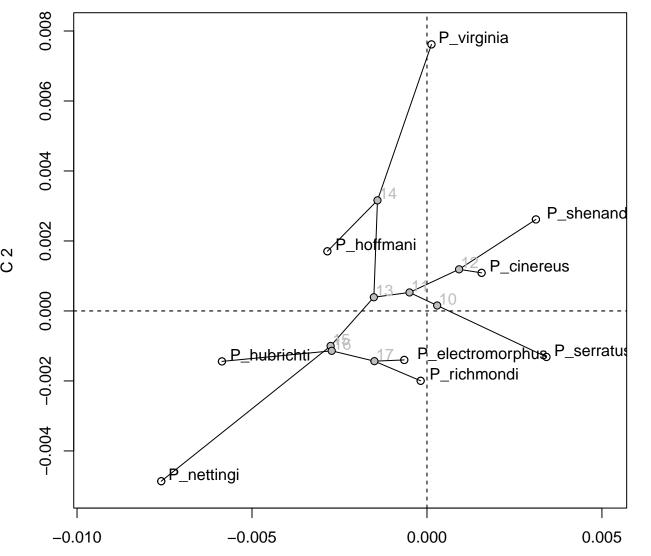
## **PaCA using OLS**

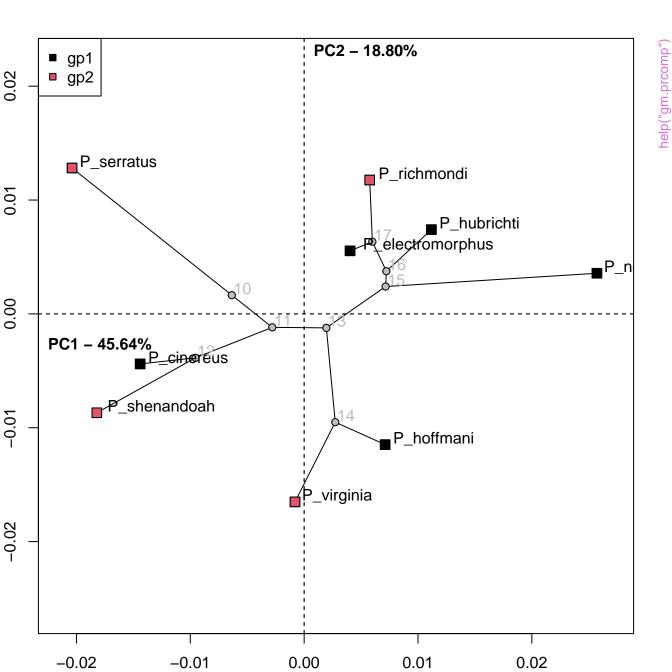


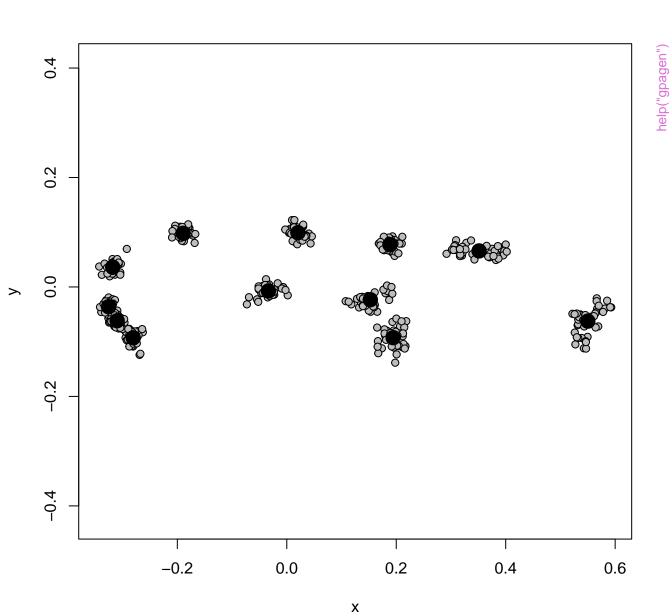
## **PaCA using GLS**

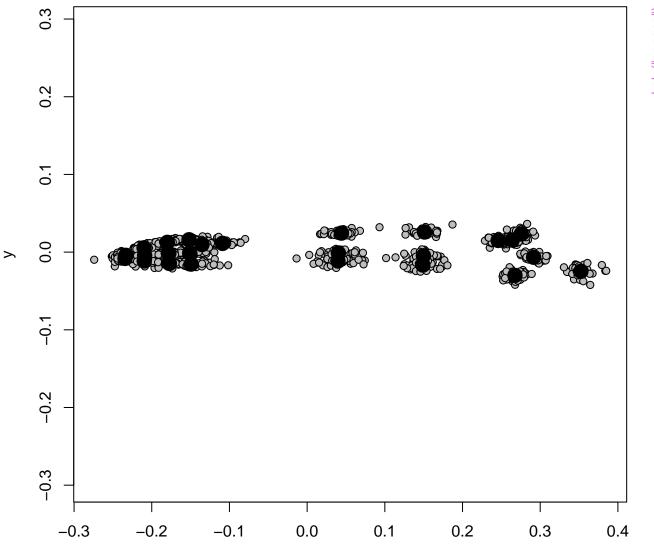


### PaCA using GLS and transformed projection

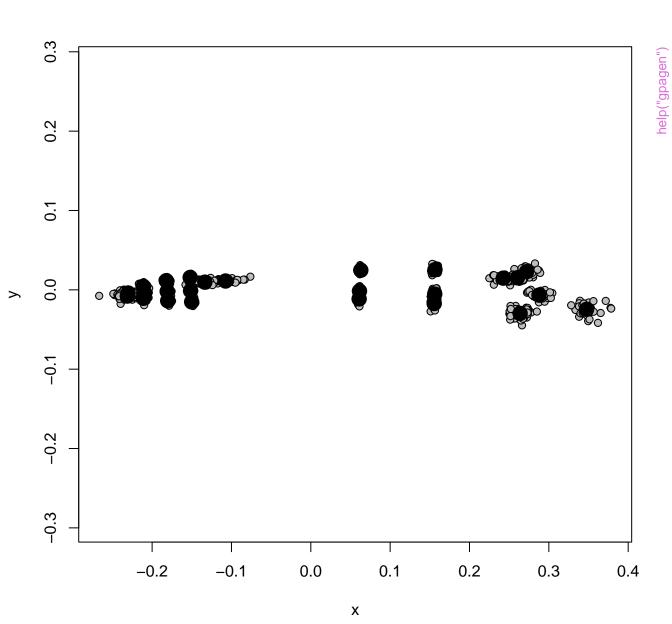


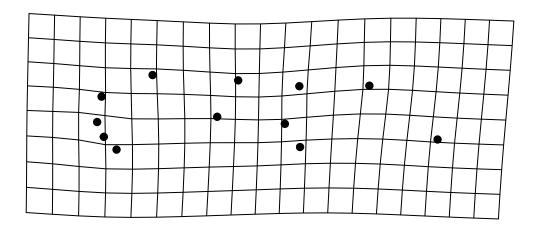


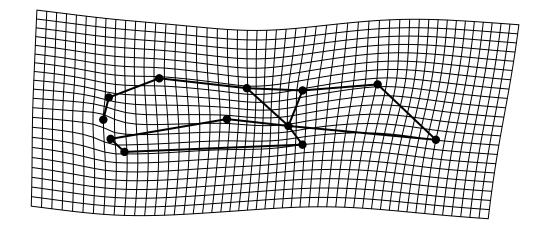


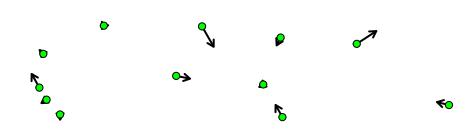


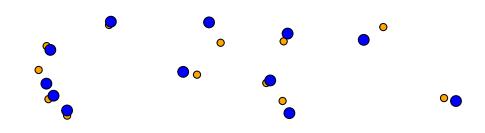
Χ

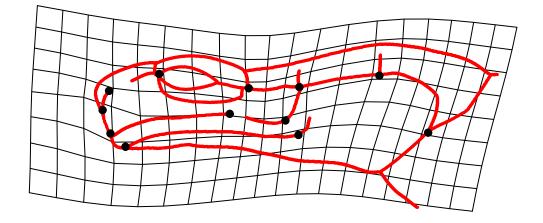


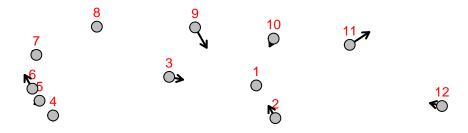




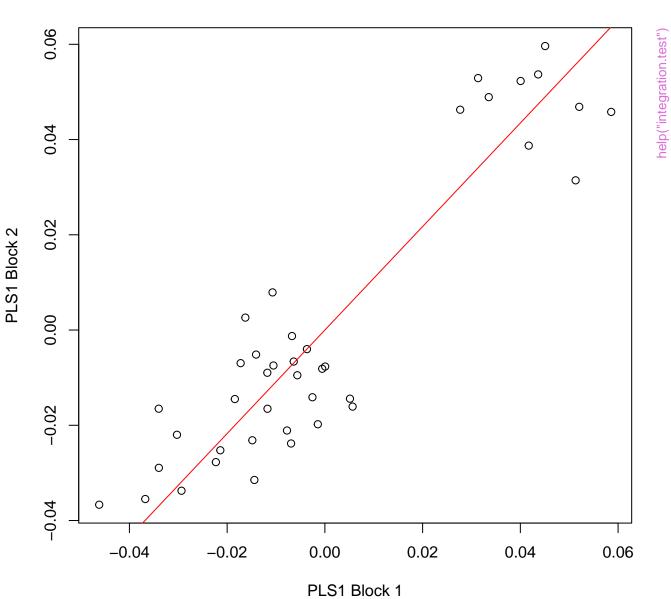


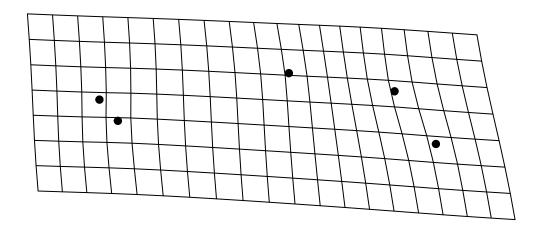


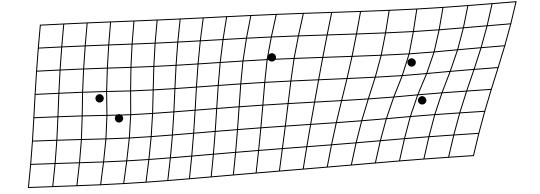


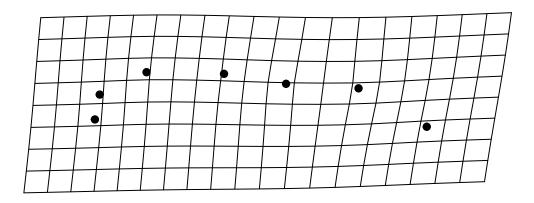


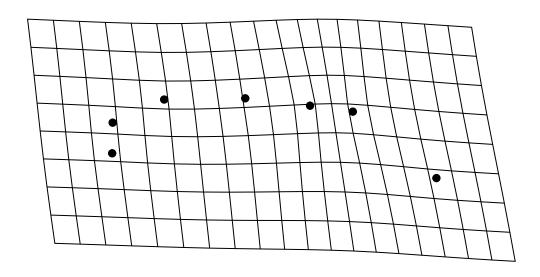
PLS1 Plot: Block 1 (X) vs. Block 2 (Y)



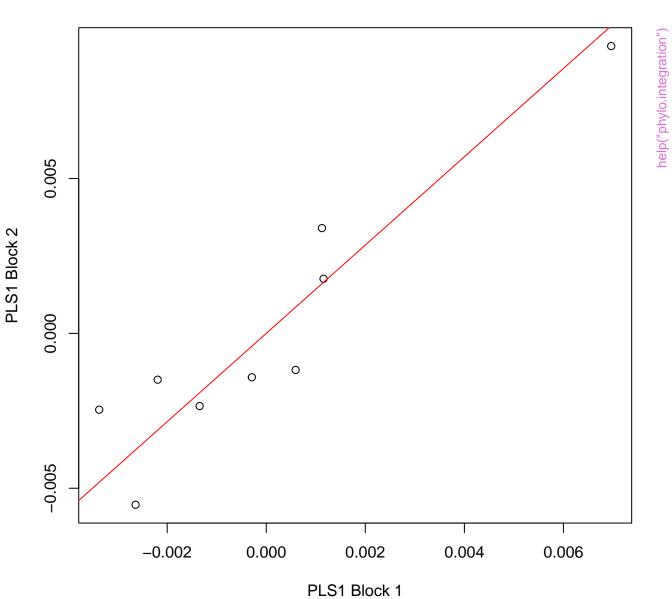




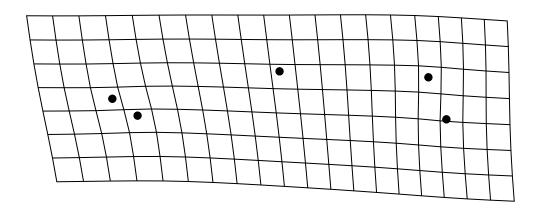


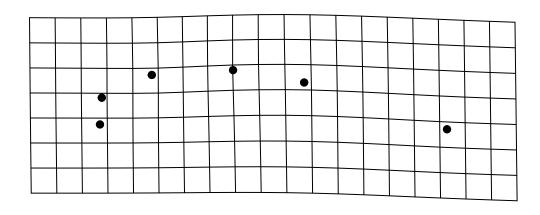


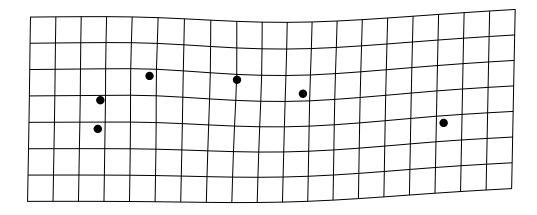
PLS1 Plot: Block 1 (X) vs. Block 2 (Y)

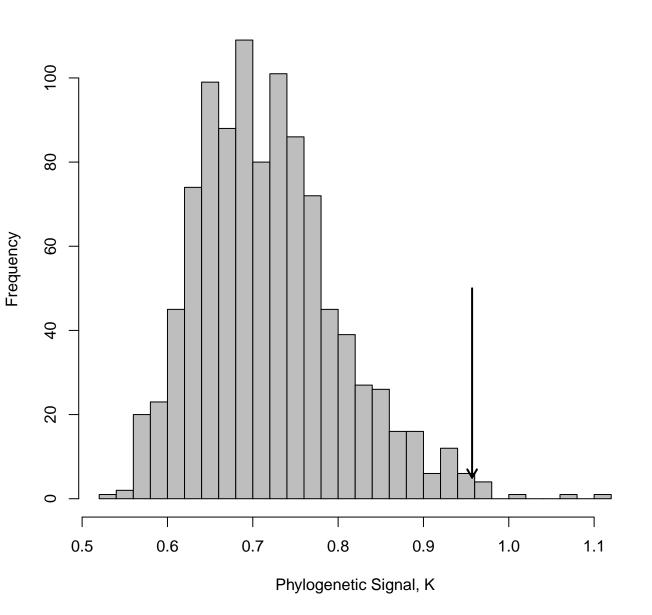


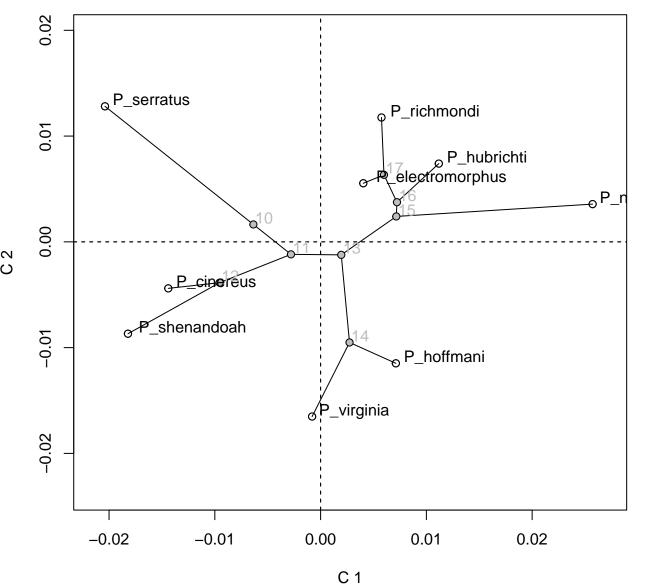
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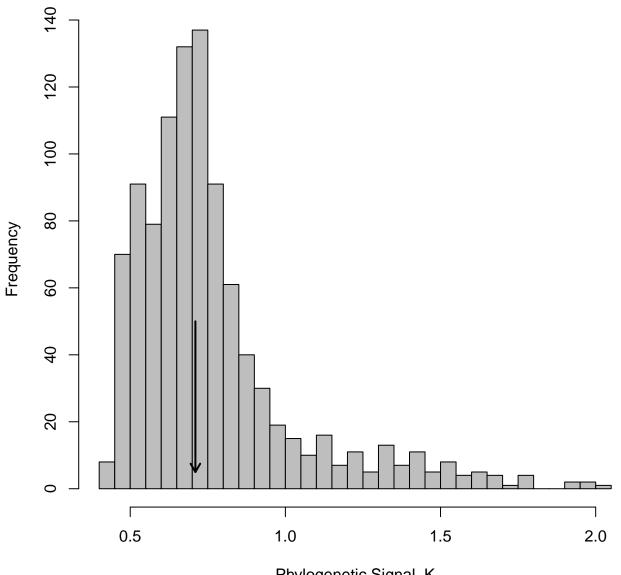






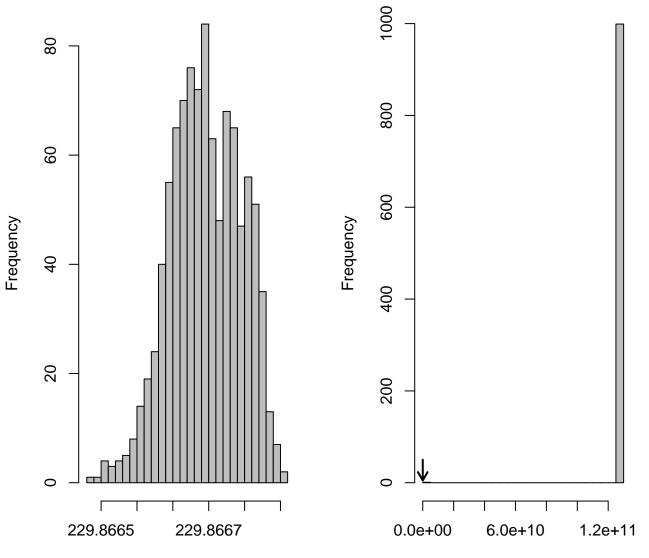




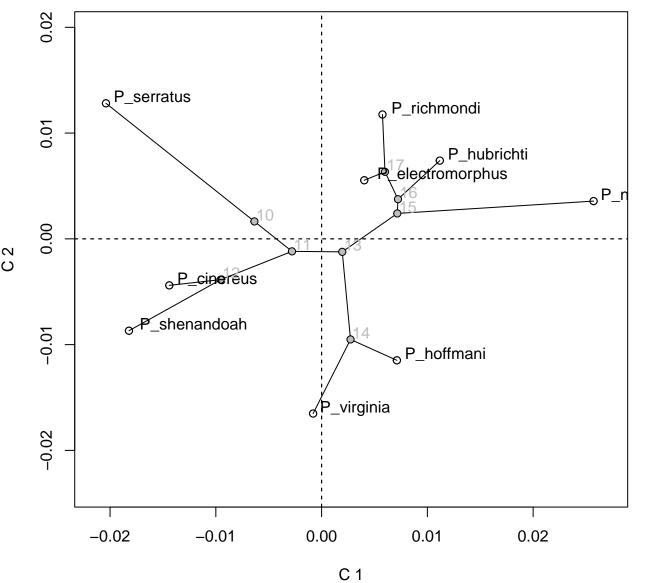


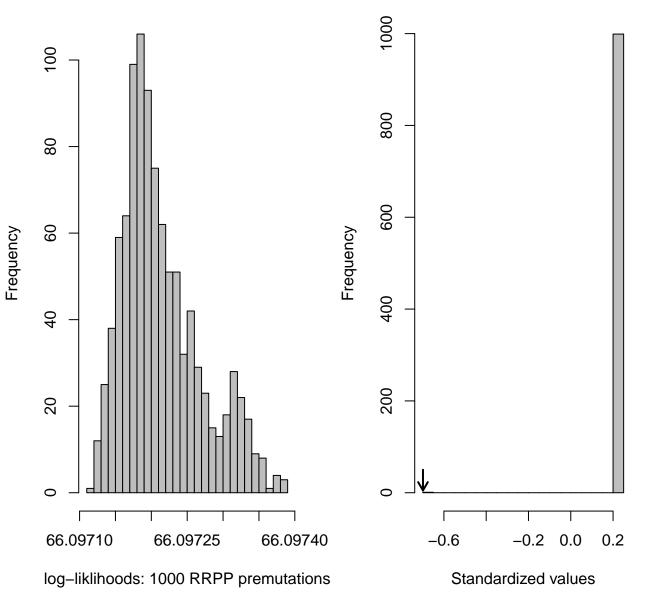
Phylogenetic Signal, K

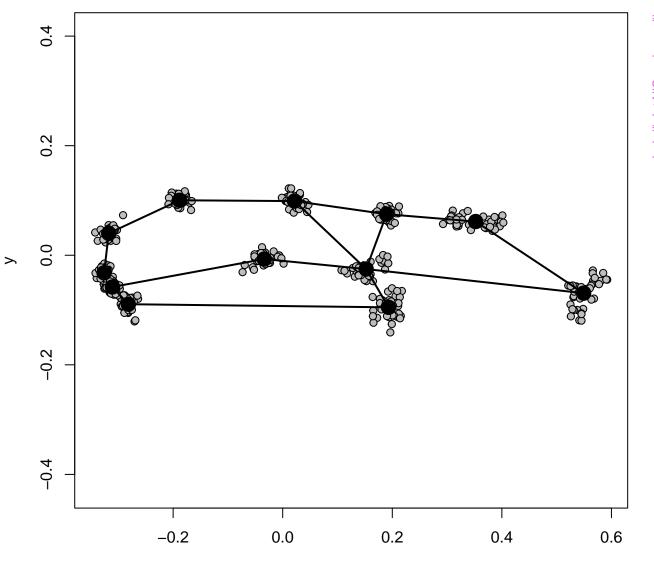
Standardized values



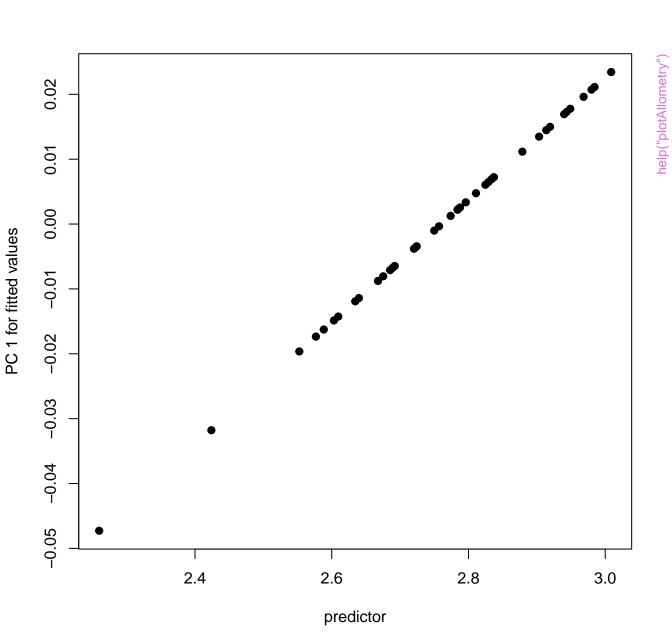
log-liklihoods: 1000 RRPP premutations

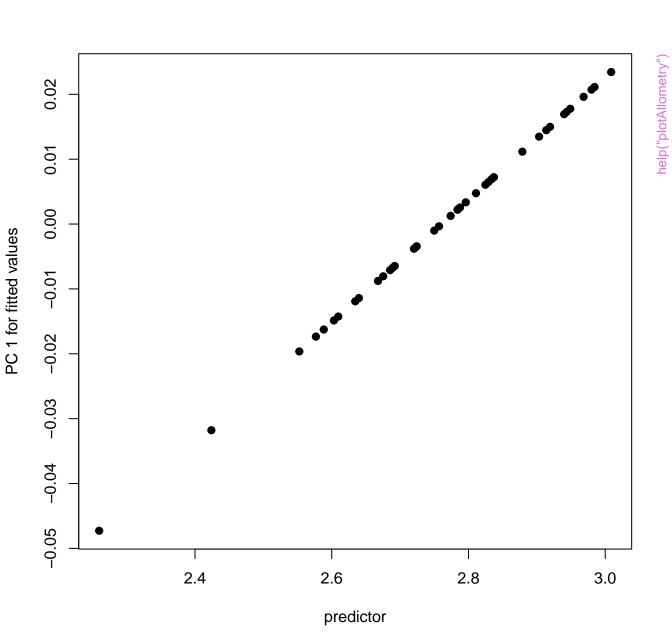


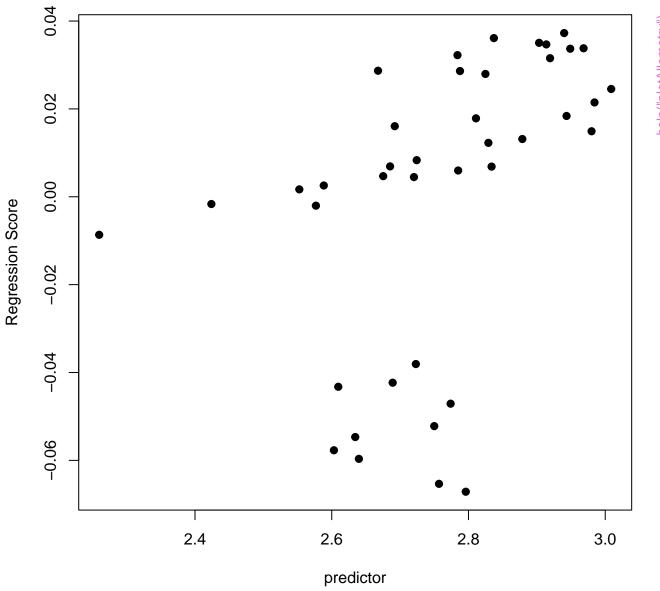


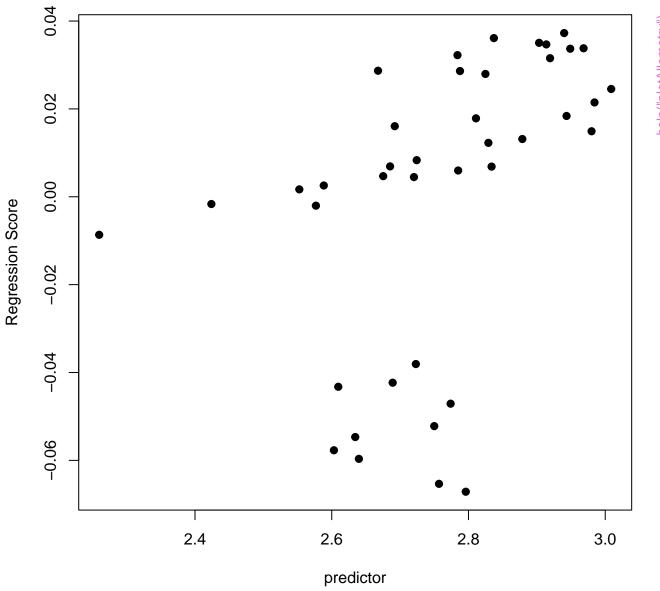


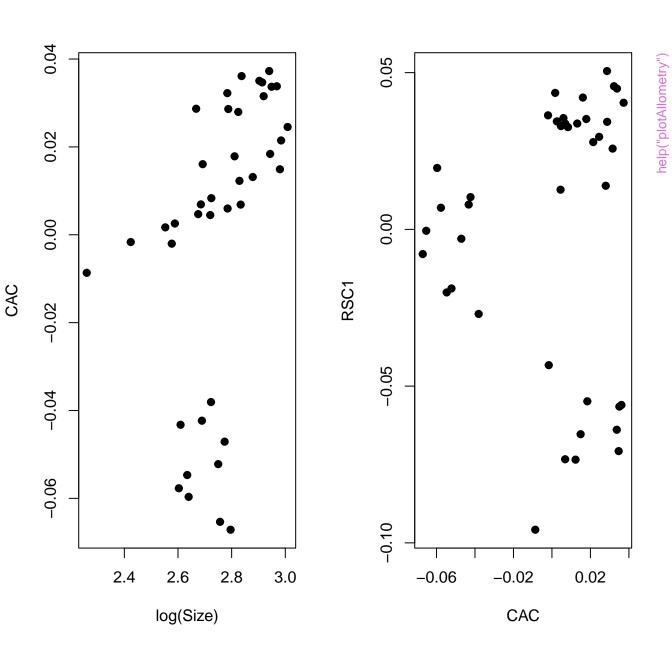
Χ

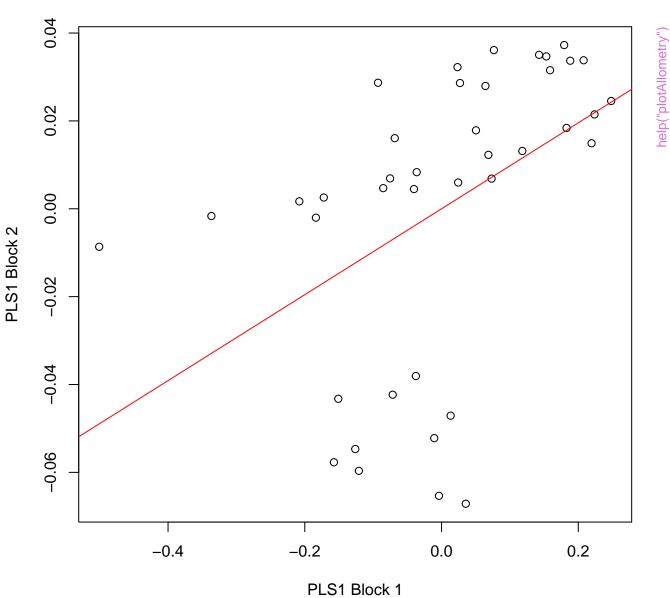


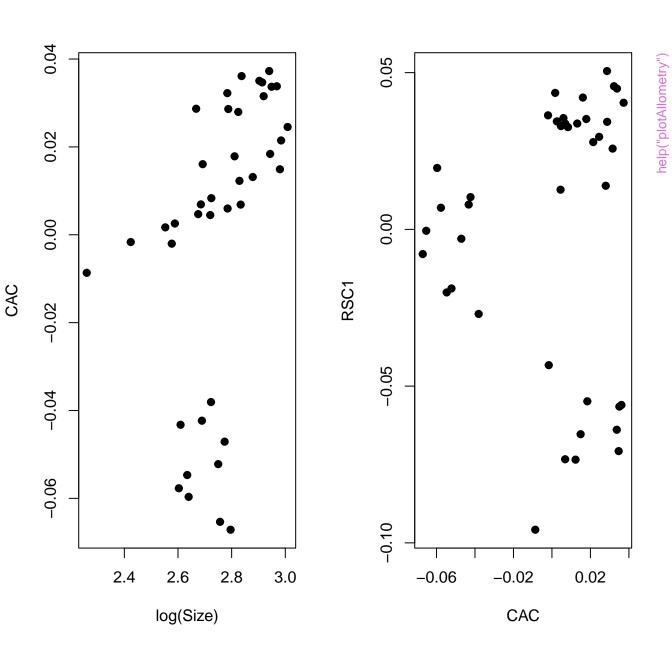


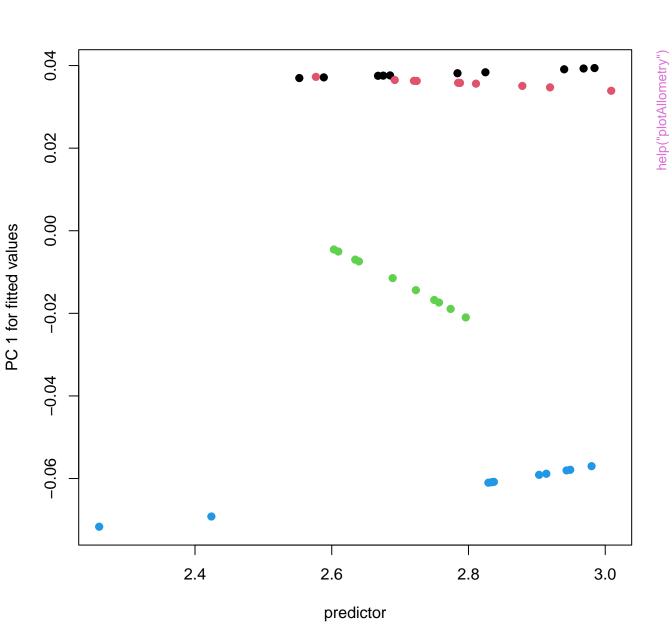




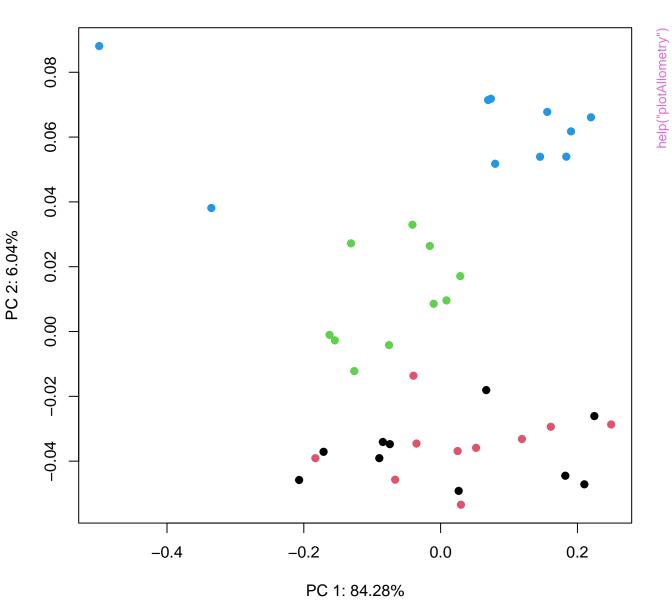




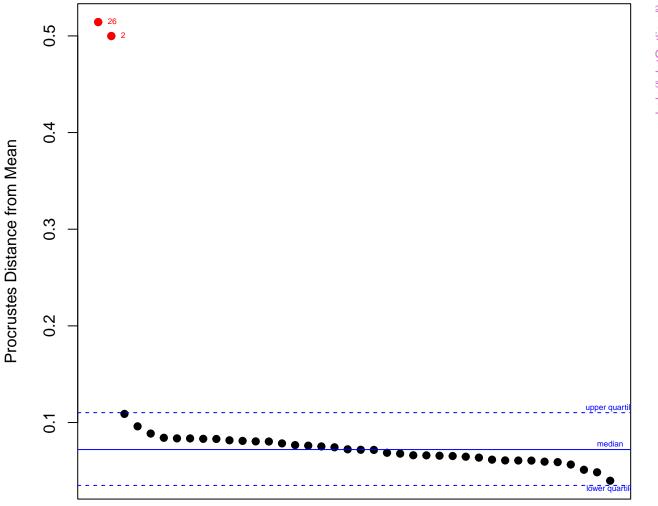




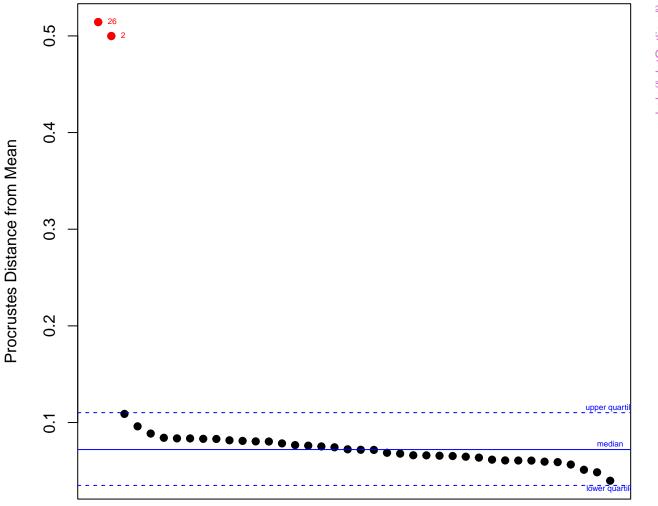
### Size-Shape PC plot



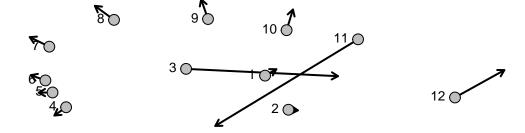
### **All Specimens**



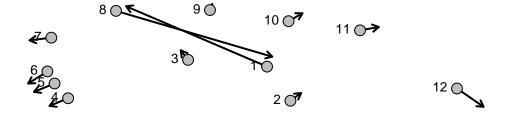
### **All Specimens**



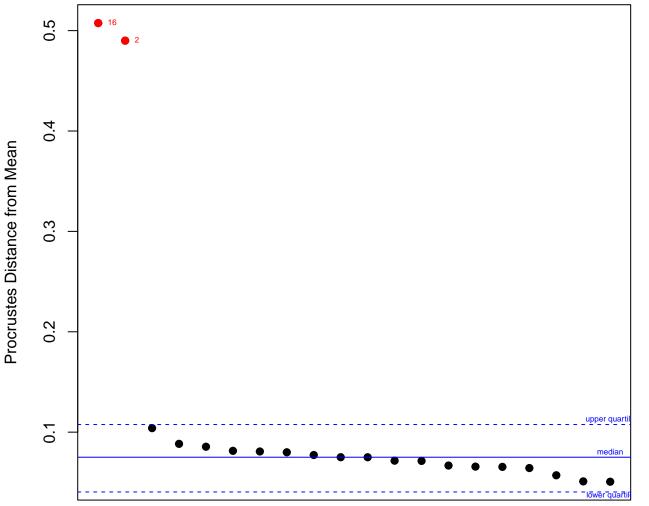
# group: All Specimens, specimen: 26



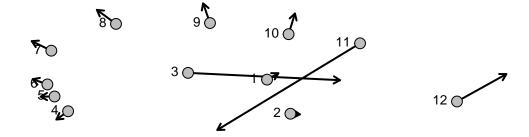
### group: All Specimens, specimen: 2



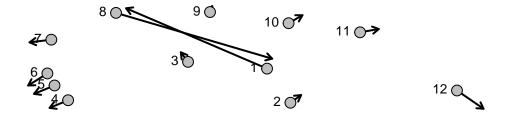
# Jord



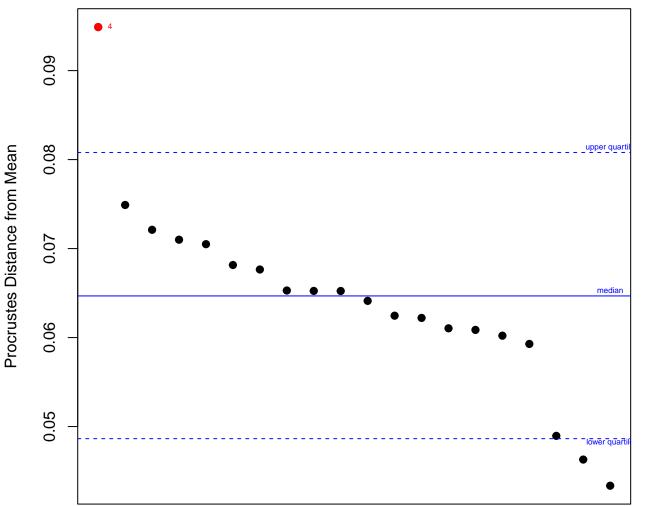
# group: Jord, specimen: 16



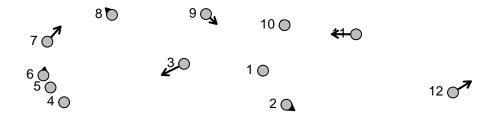
### group: Jord, specimen: 2

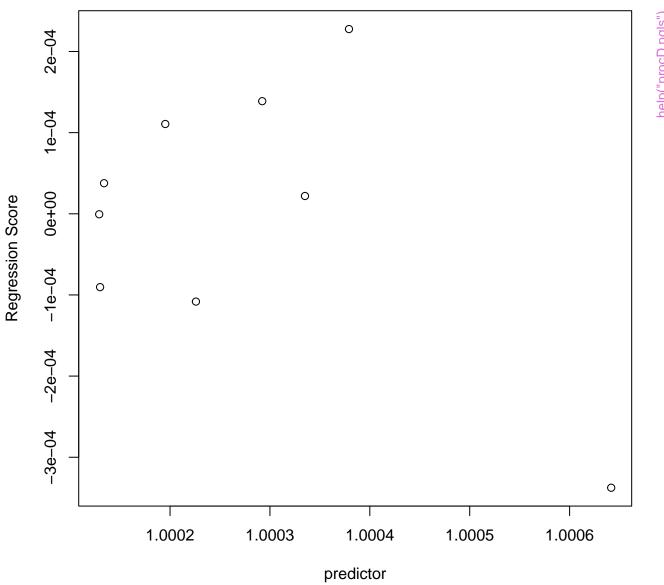


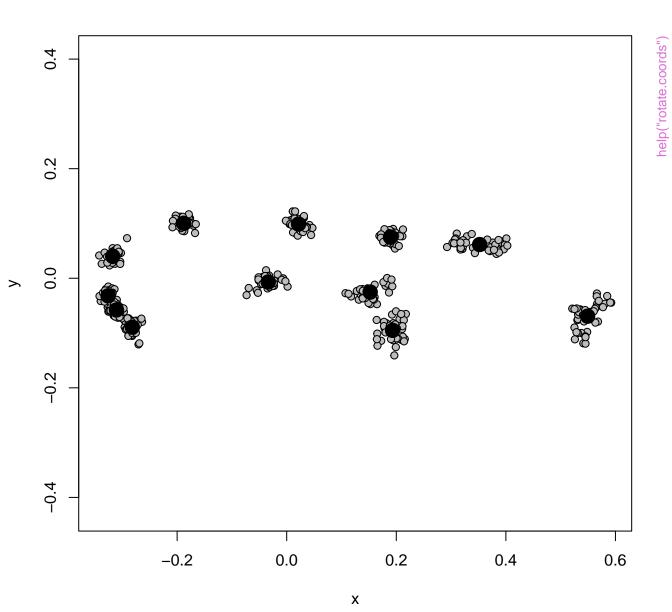
Teyah

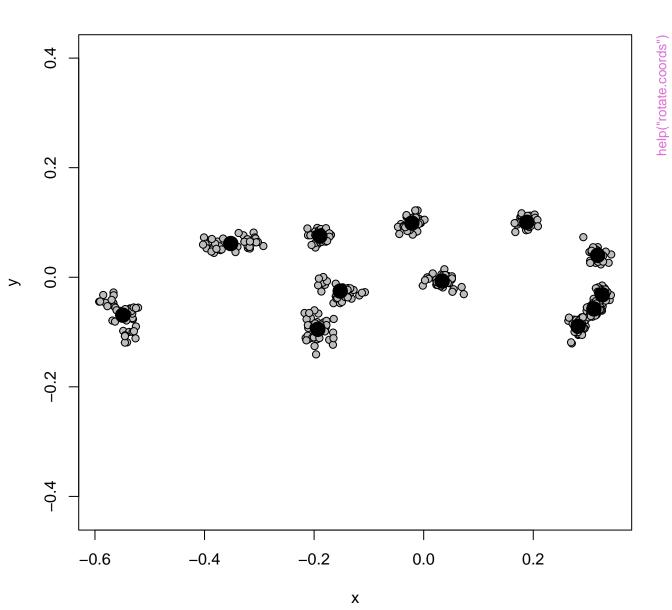


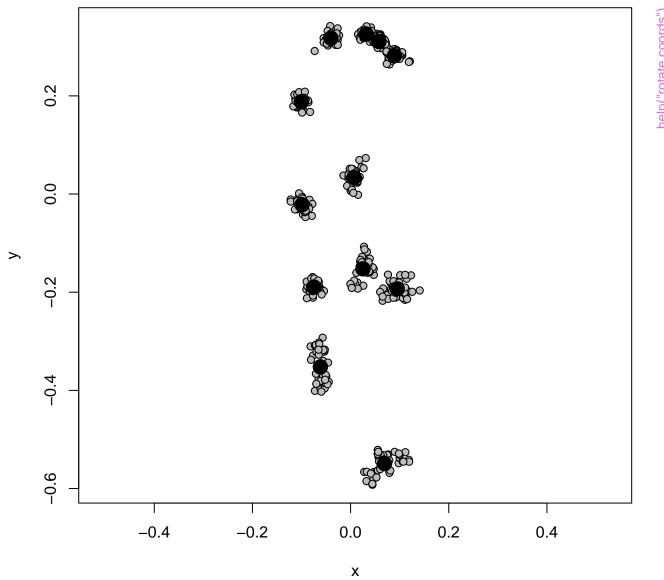
# group: Teyah, specimen: 4

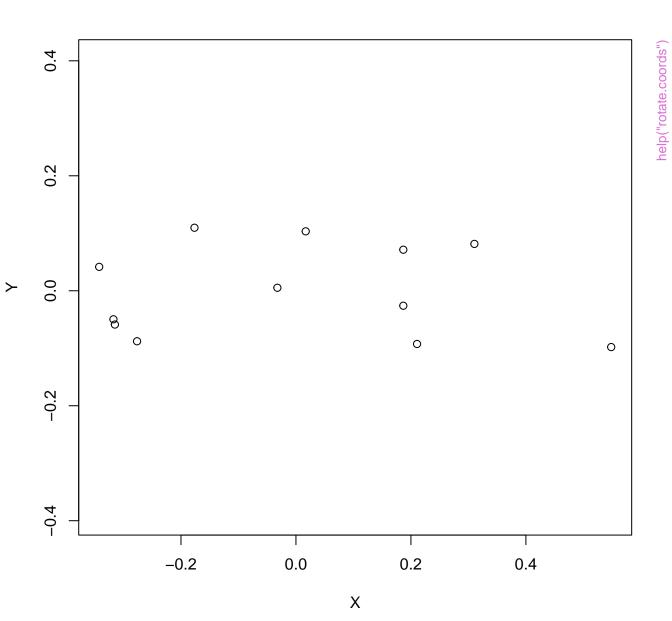


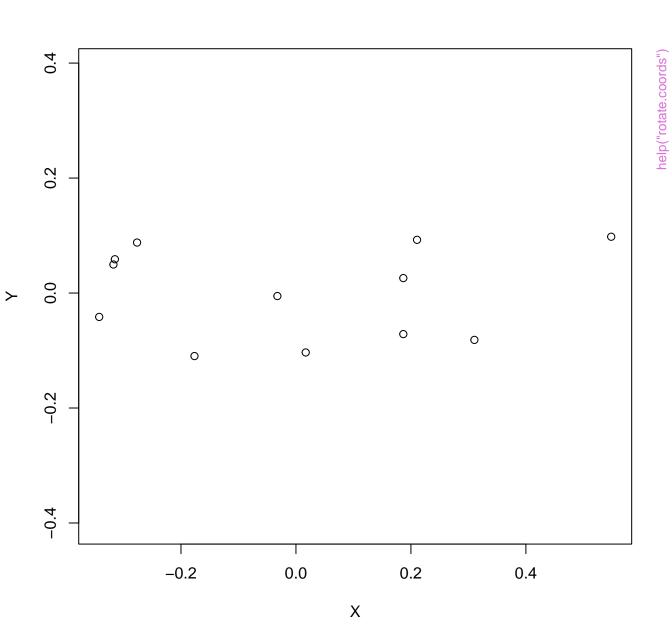


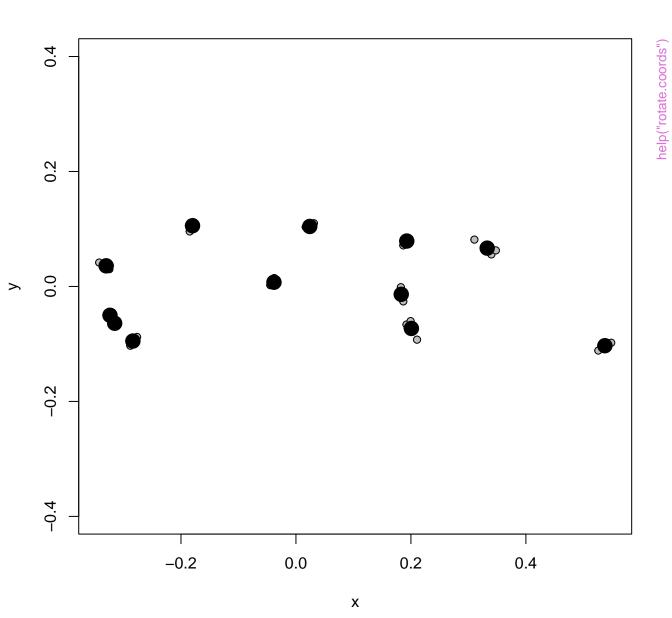




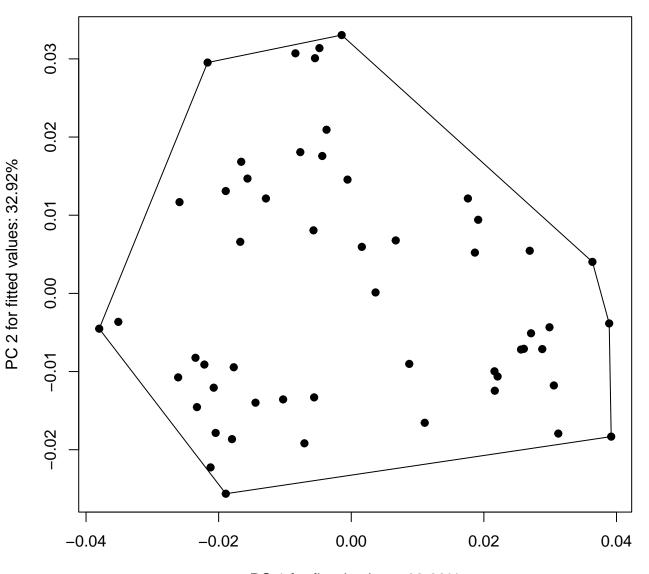




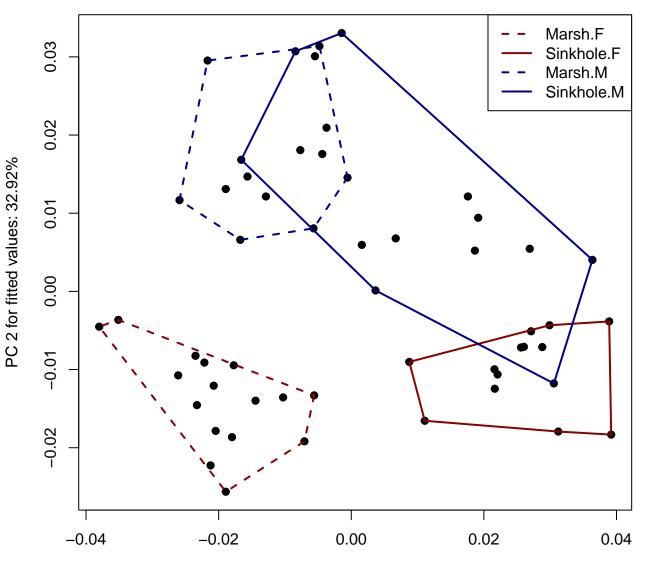




Χ



PC 1 for fitted values: 62.66%



PC 1 for fitted values: 62.66%

