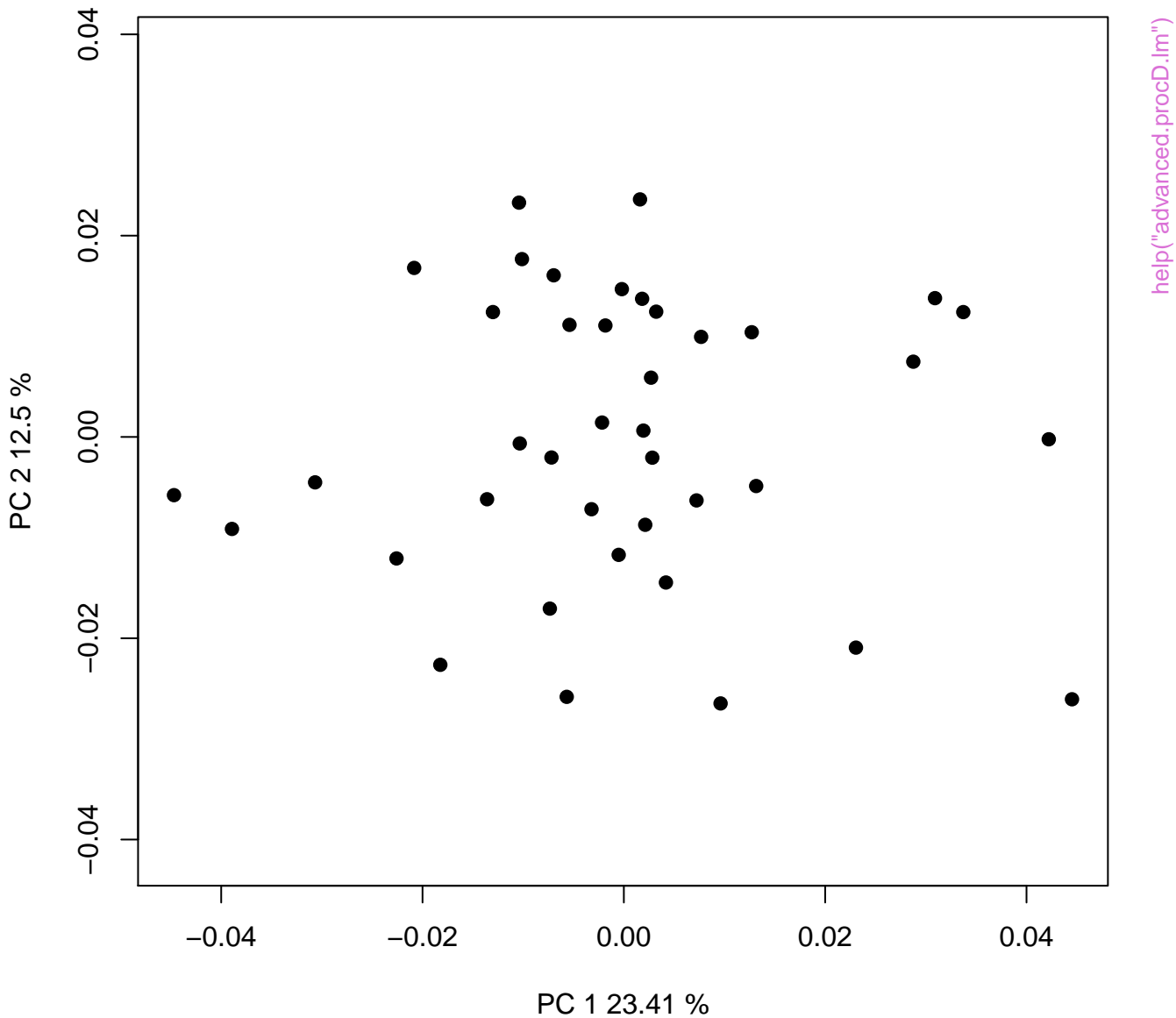
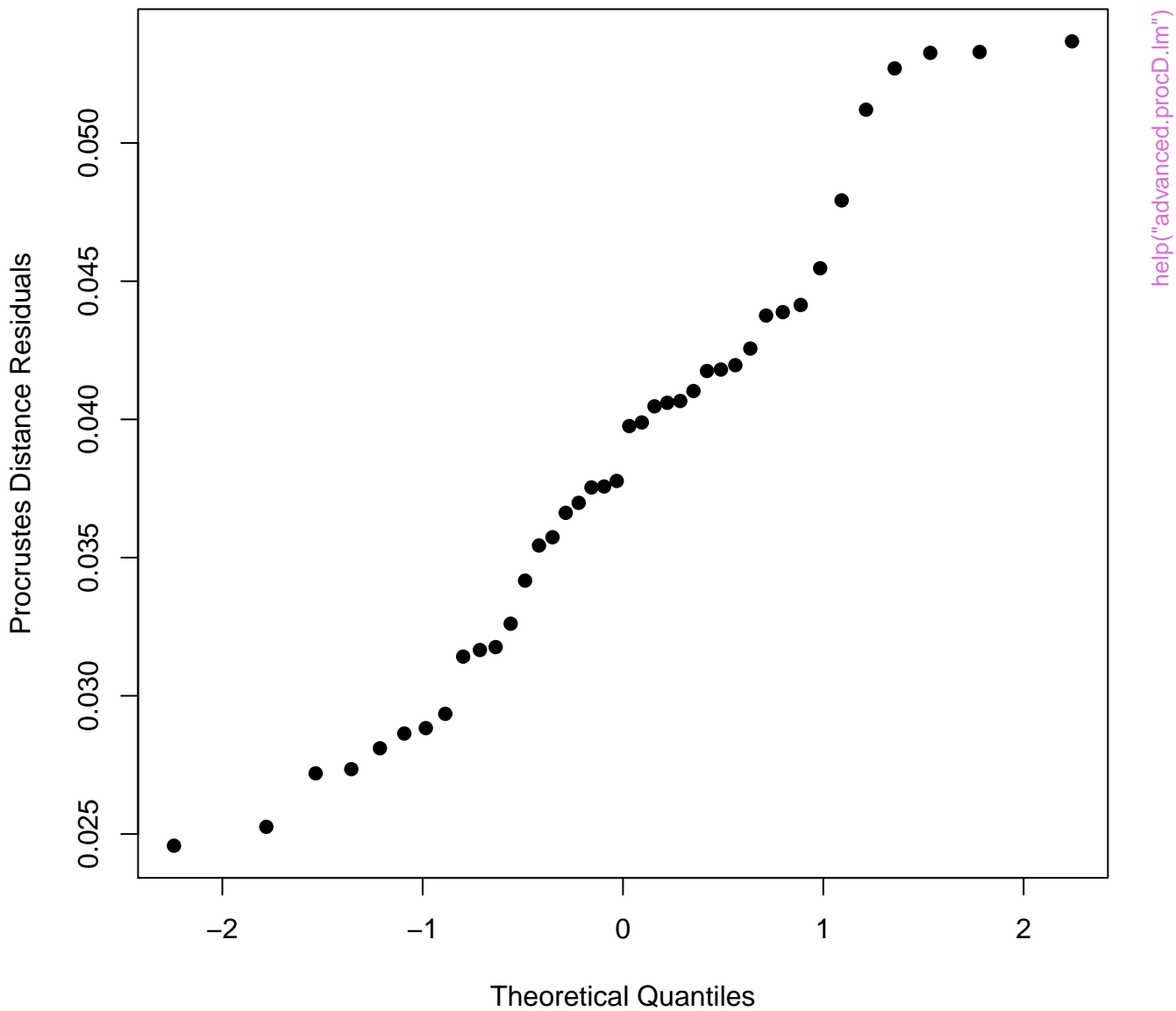


PCA Residuals

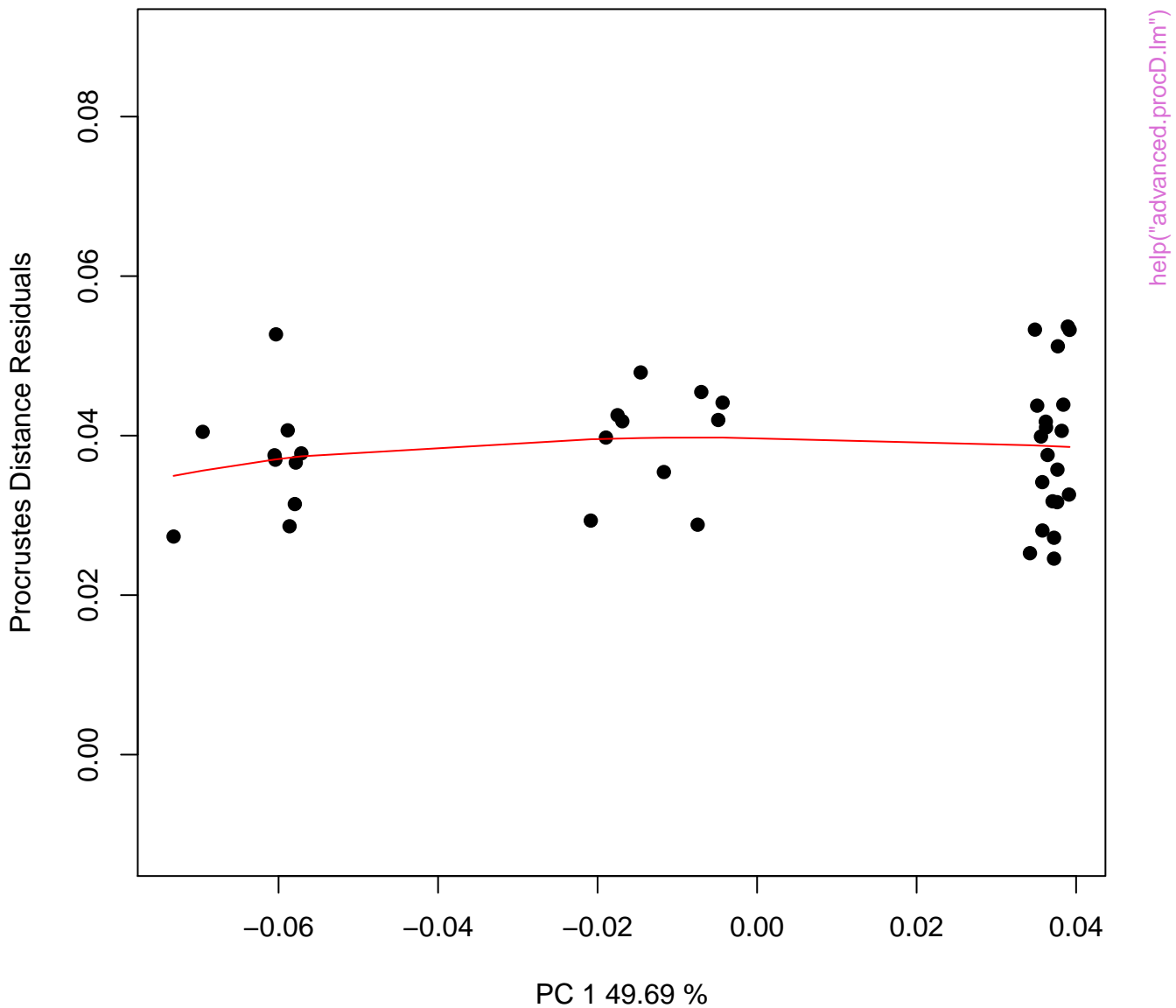


Q-Q plot

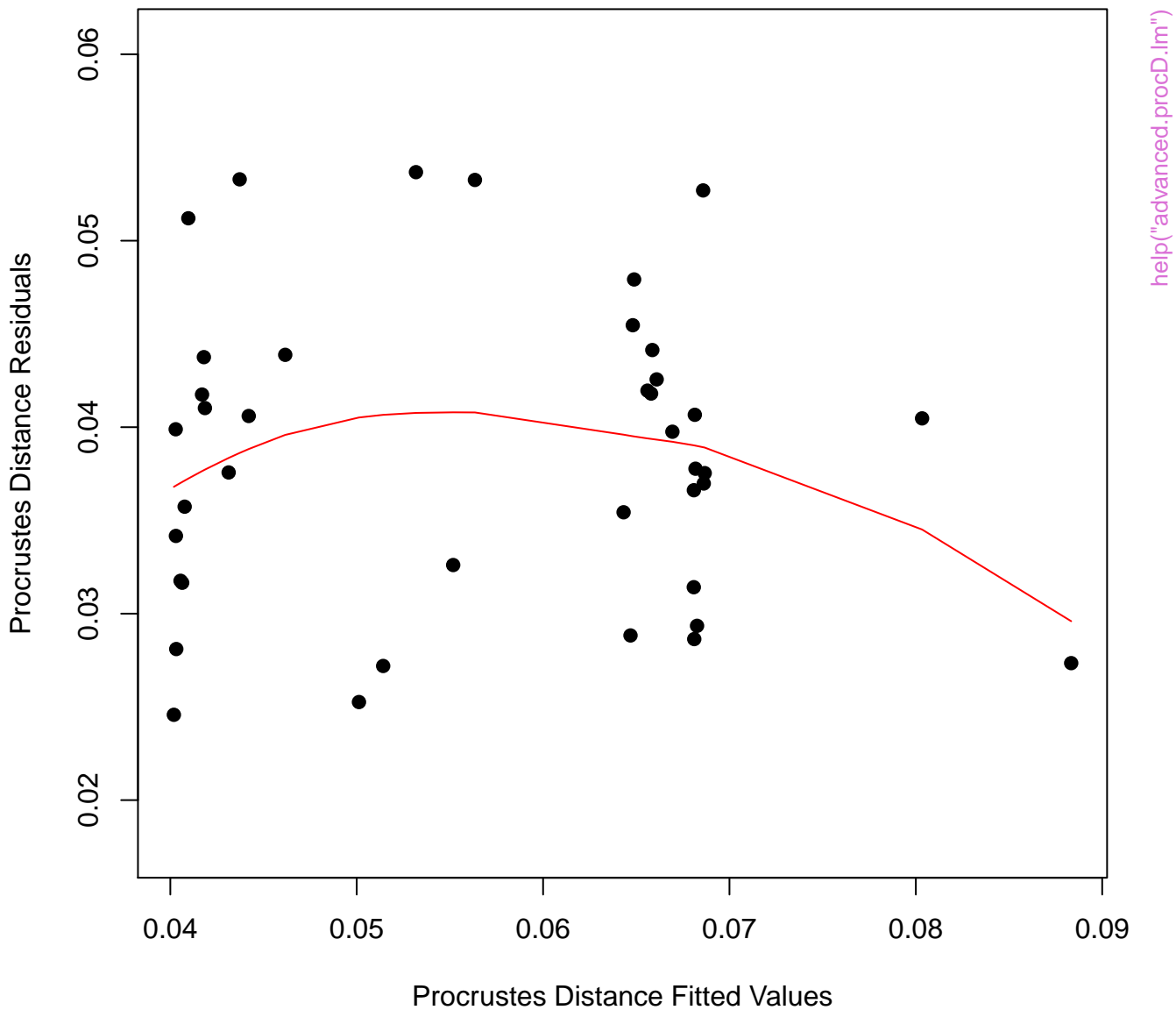


help("advanced.procD.lm")

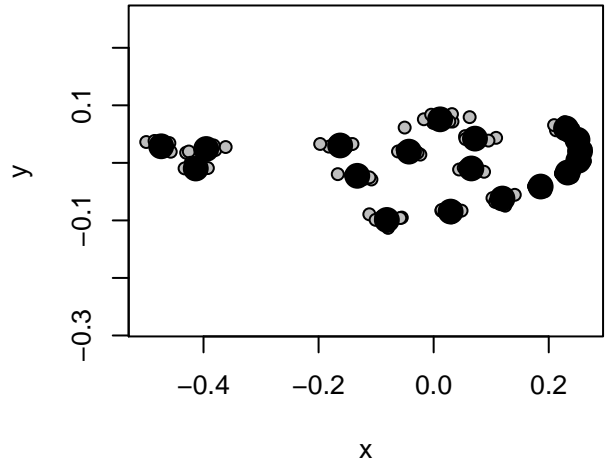
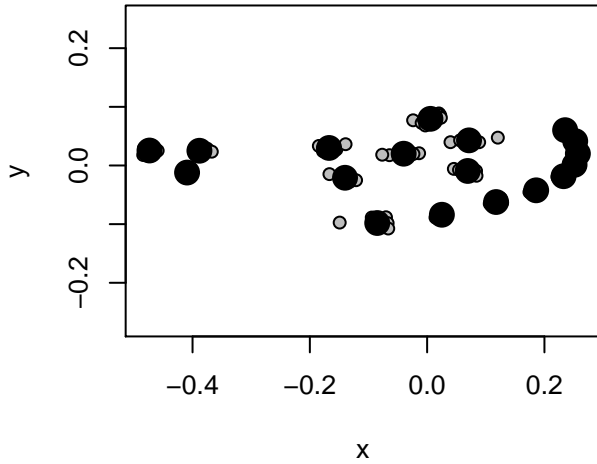
Residuals vs. PC 1 fitted



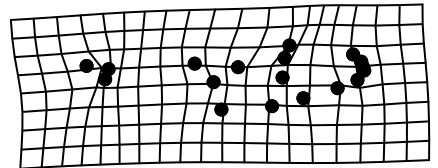
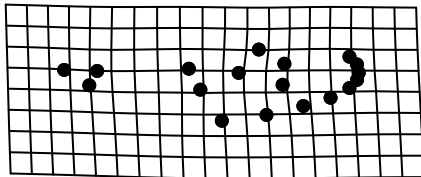
Residuals vs. Fitted



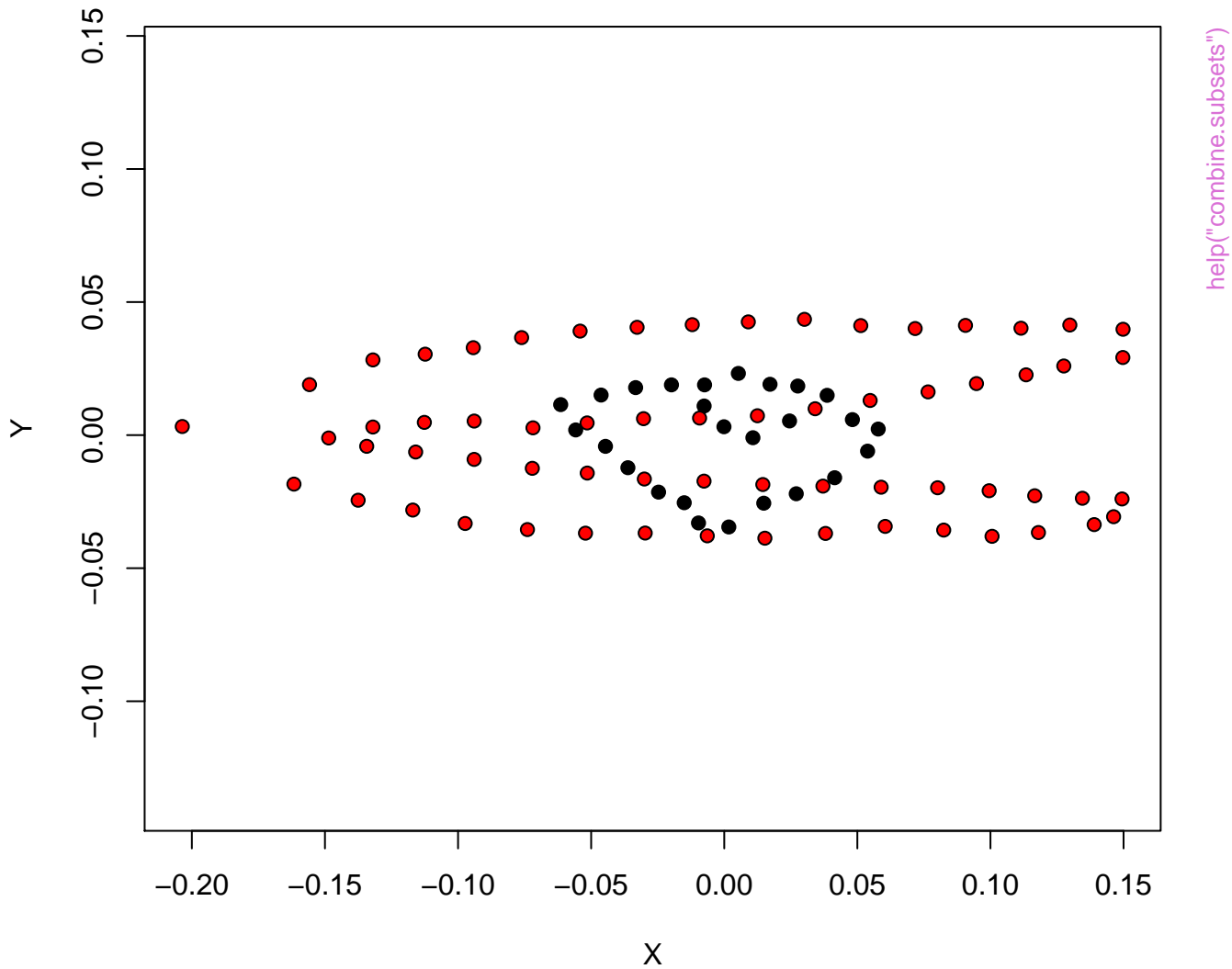
Symmetric Shape Component (left) and Asymmetric Shape Component (right)

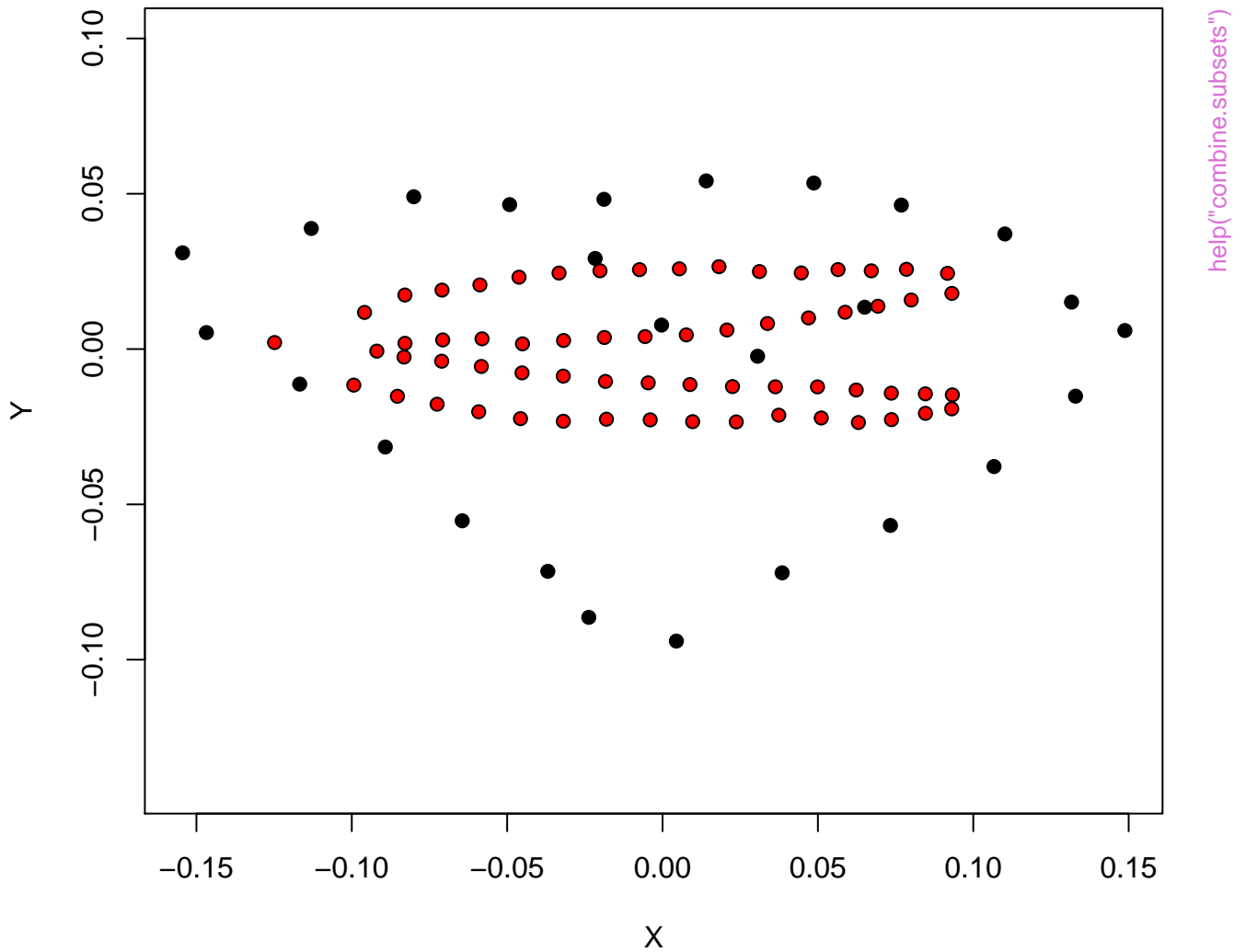


help("bilat.symmetry")

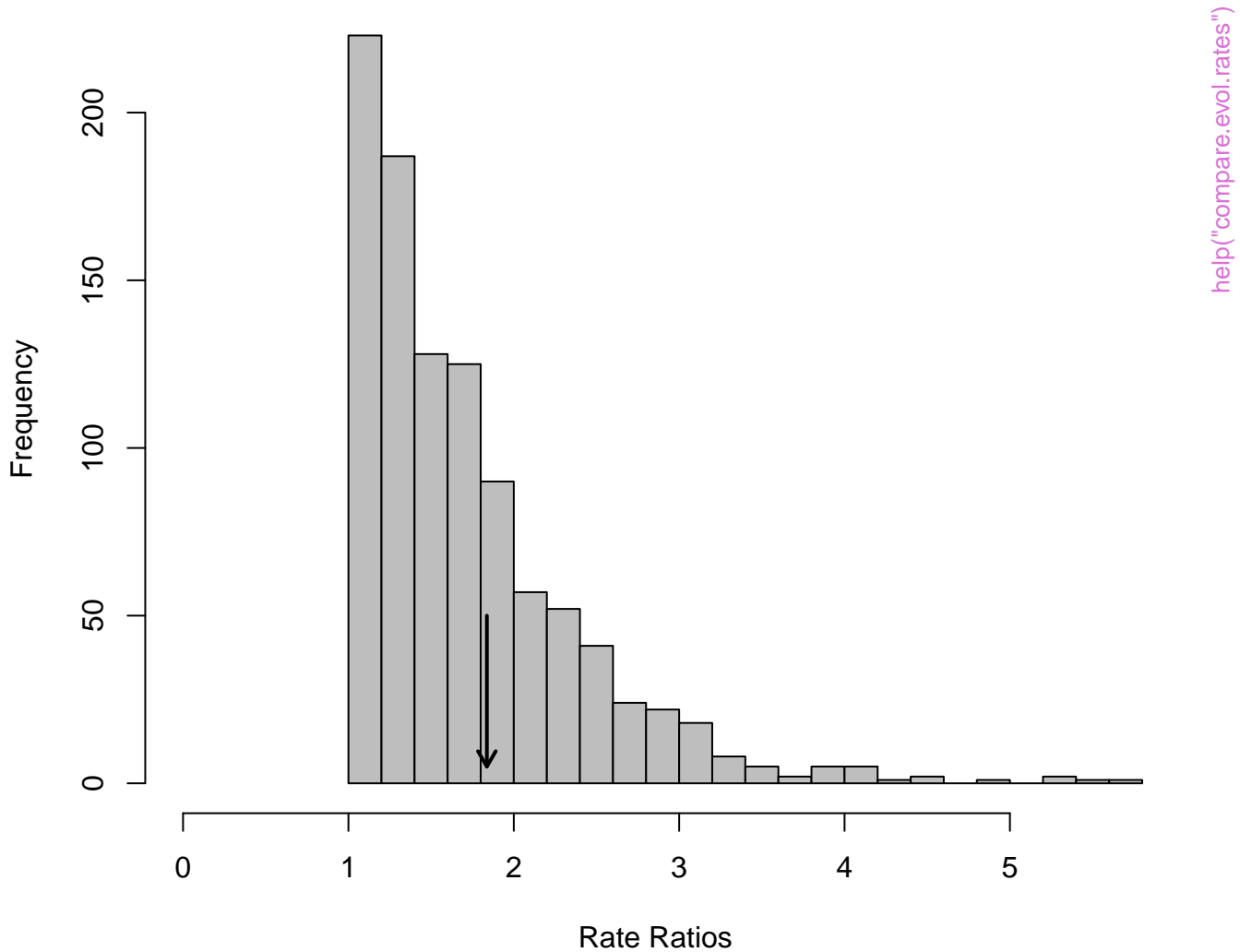


Mean directional (left) and fluctuating (right) asymmetry

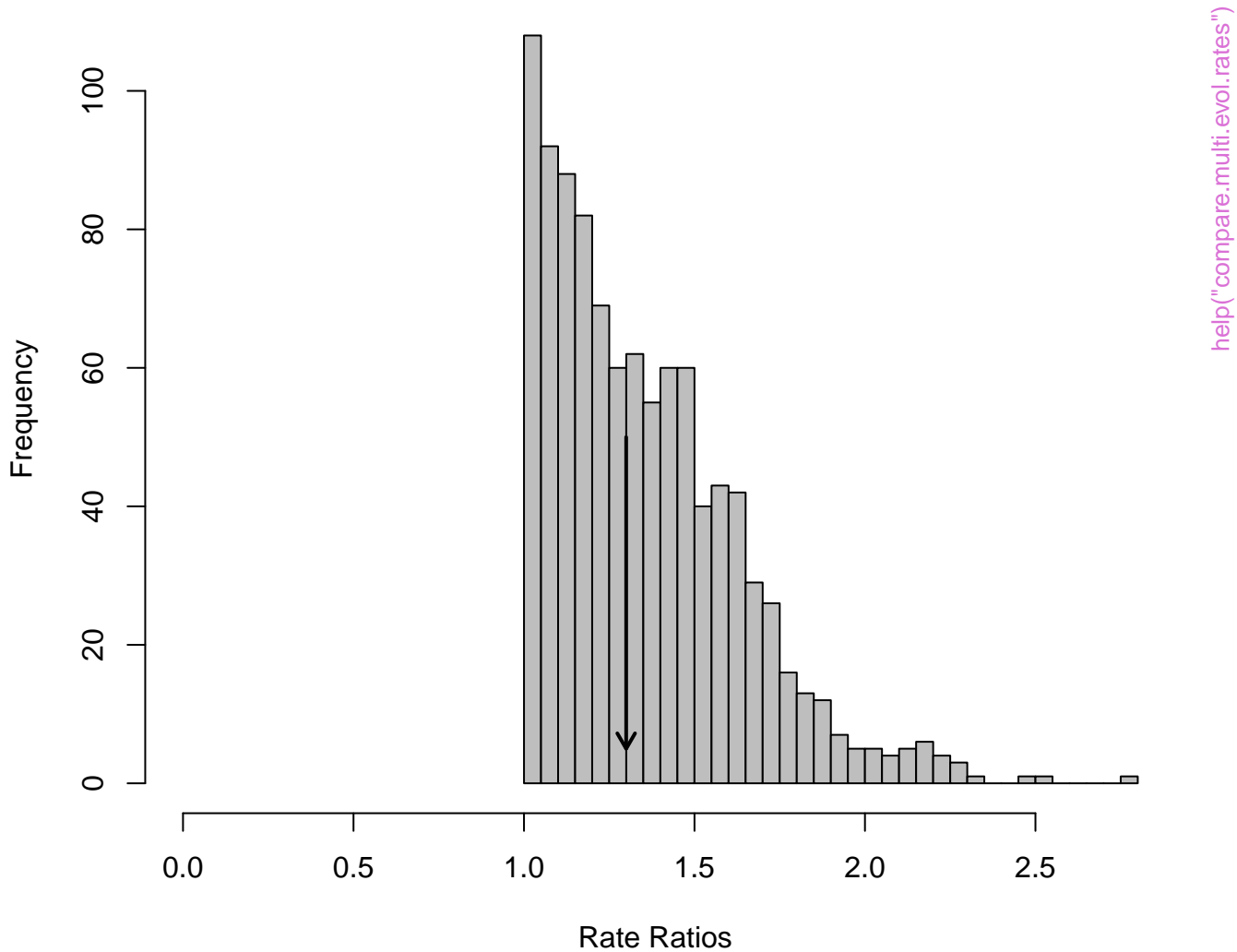




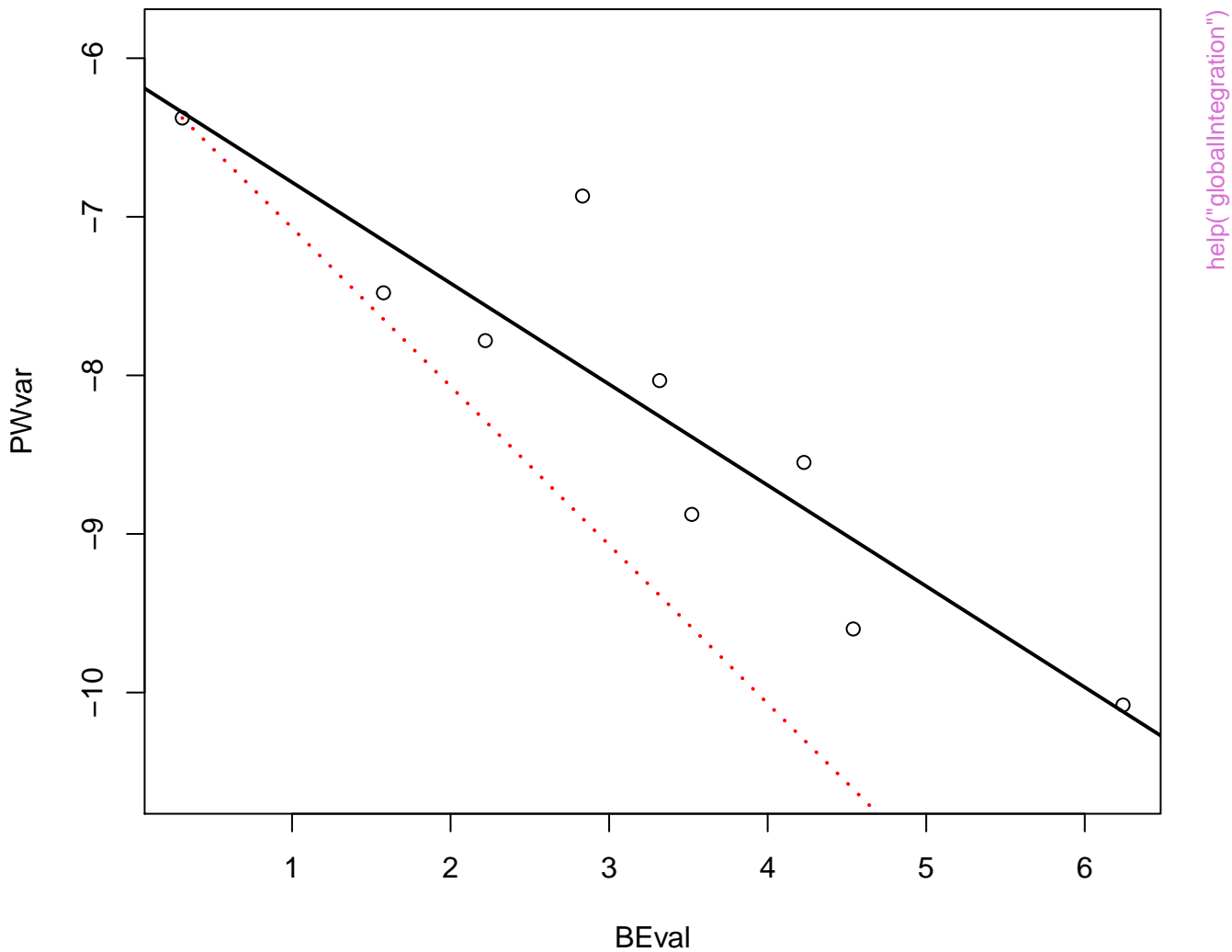
Observed Rate Ratio = 1.8372 ; P-value = 0.321



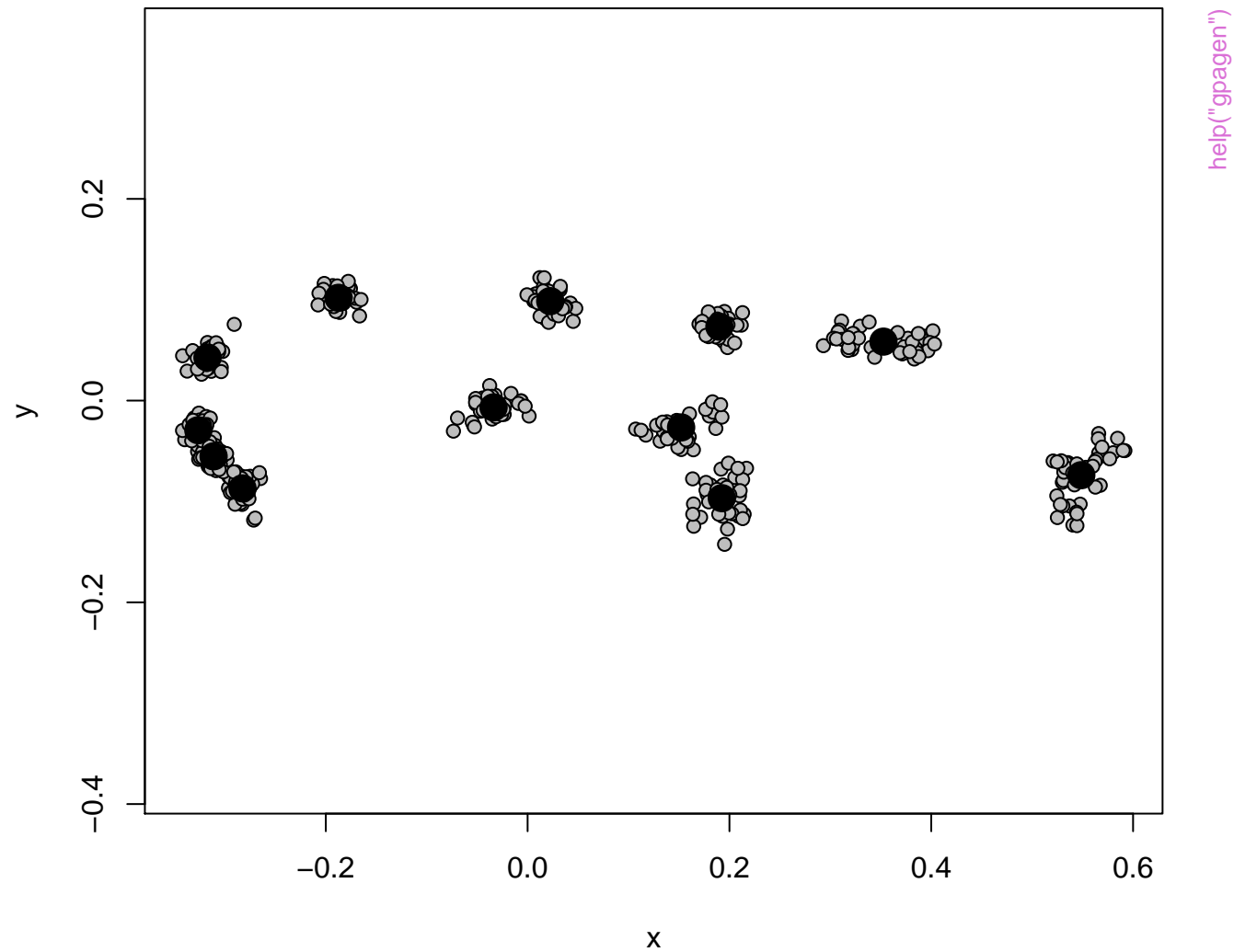
Observed Rate Ratio = 1.2997 ; P-value = 0.503

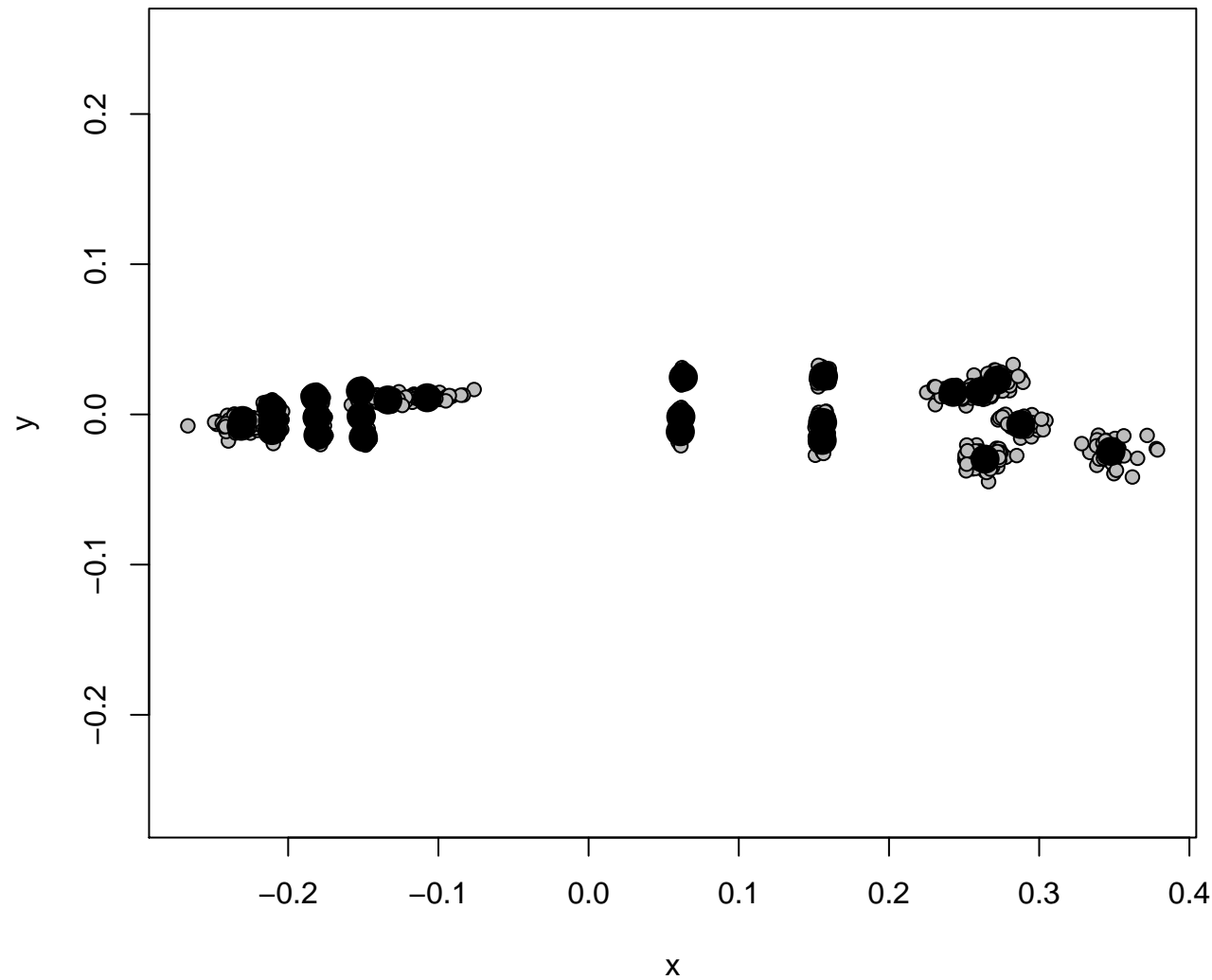


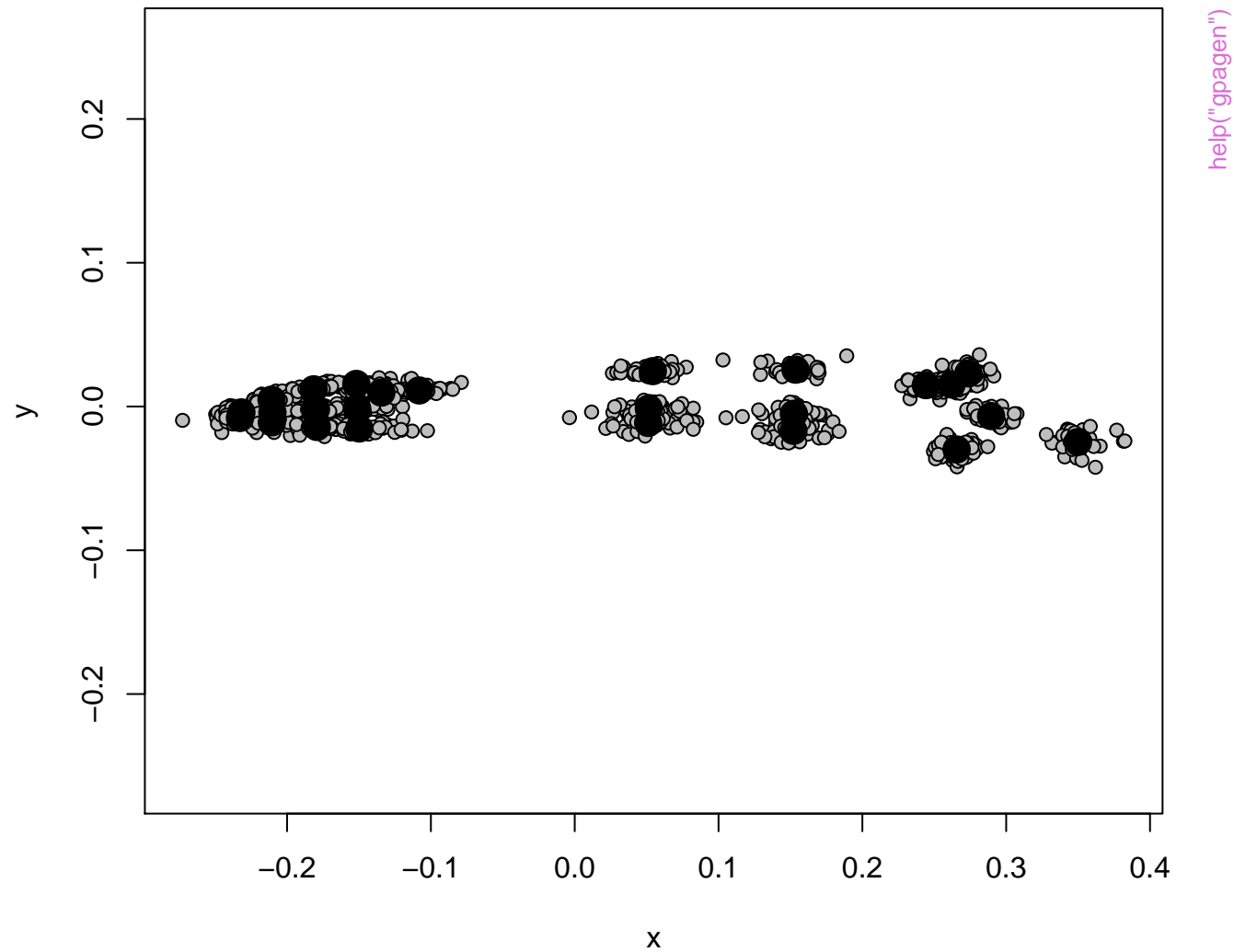
ObservedSlope_{black} = -0.6369197

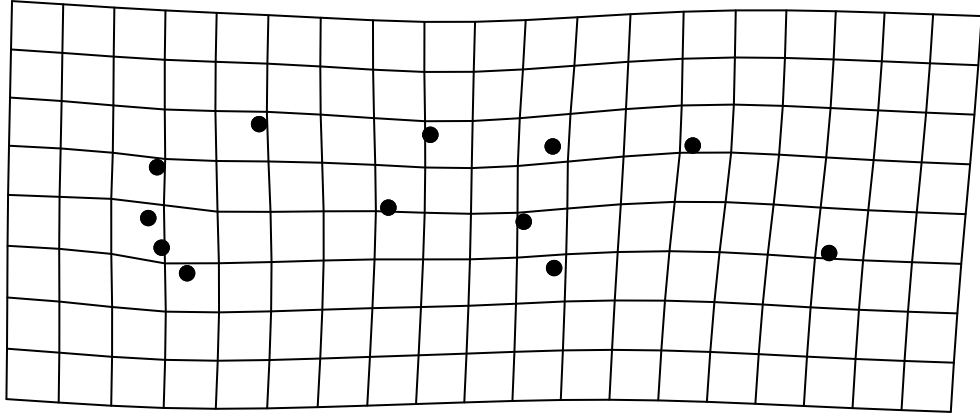


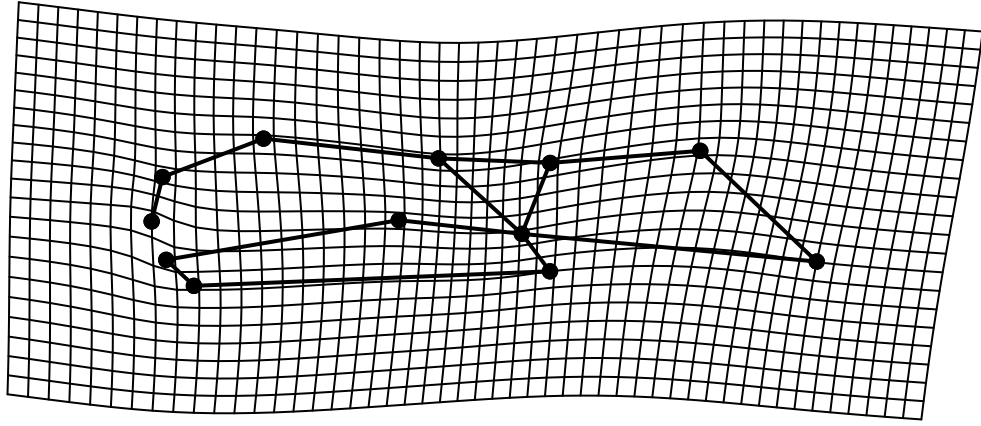
help("globalIntegration")

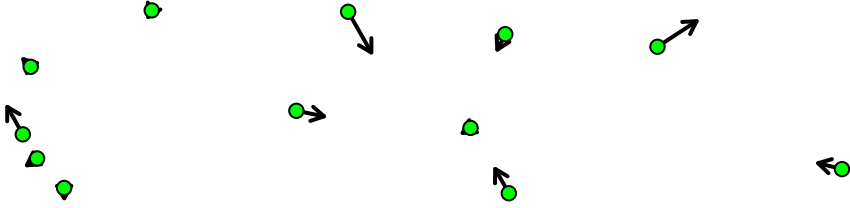


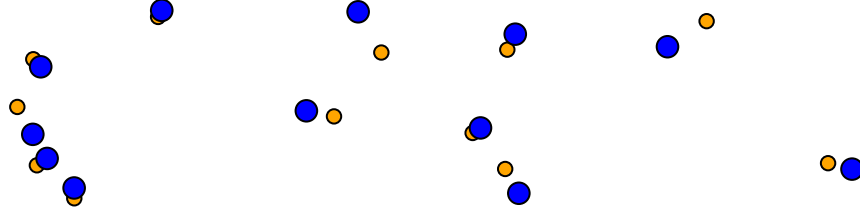


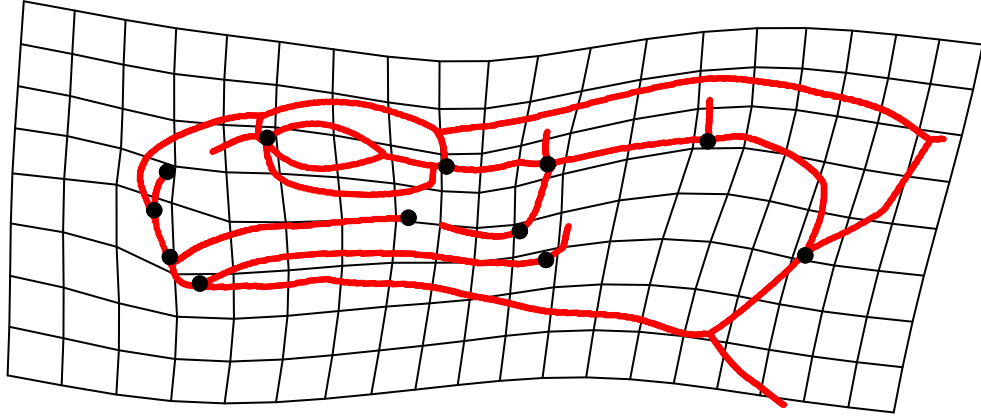


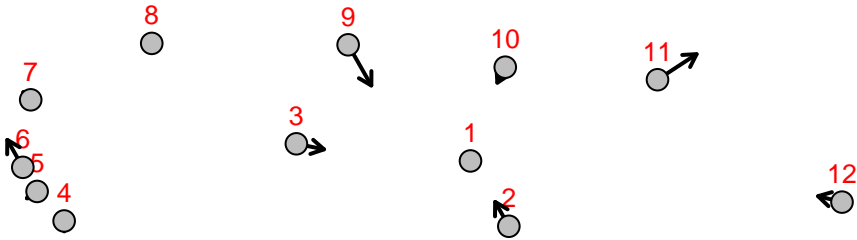




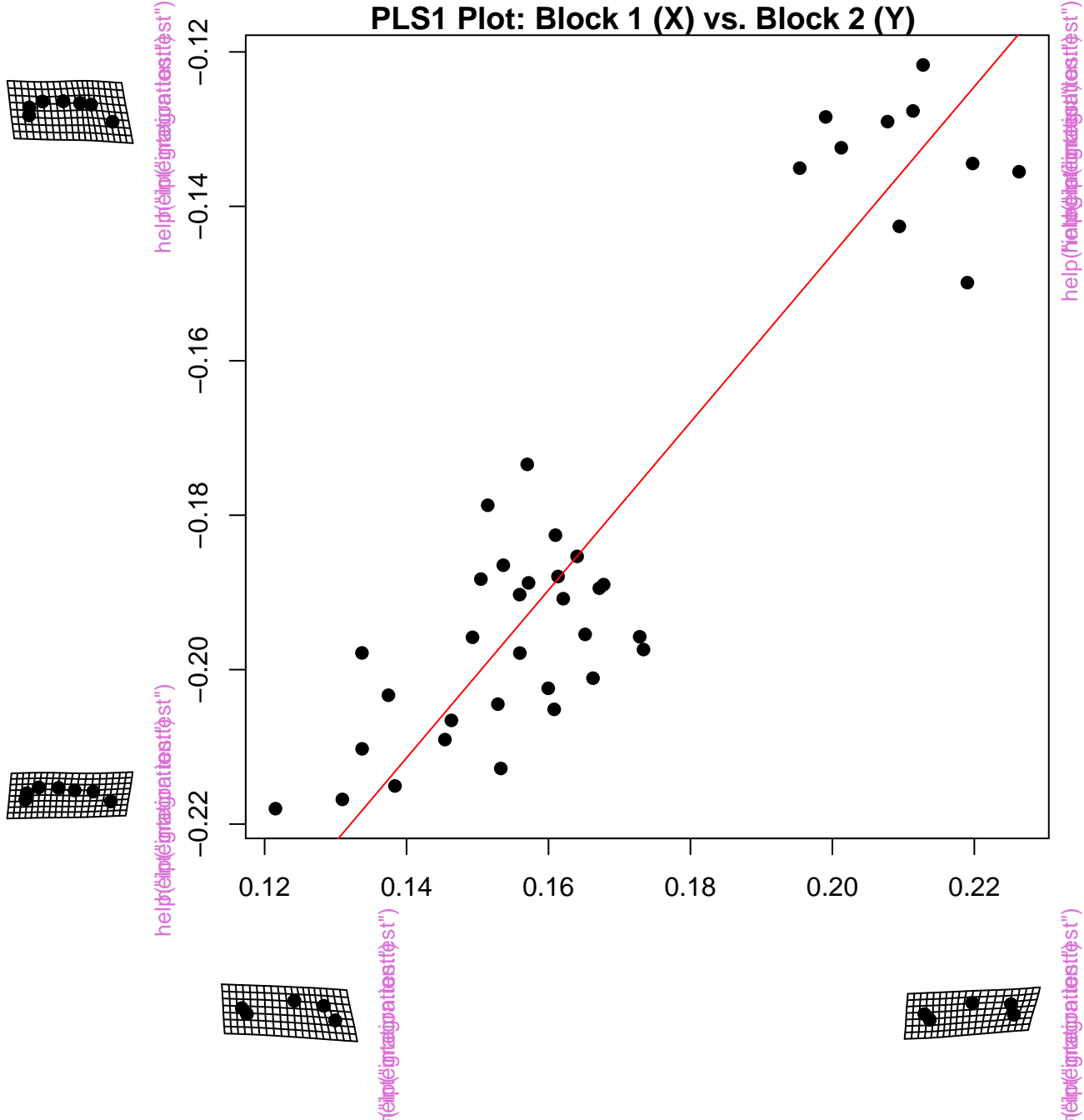




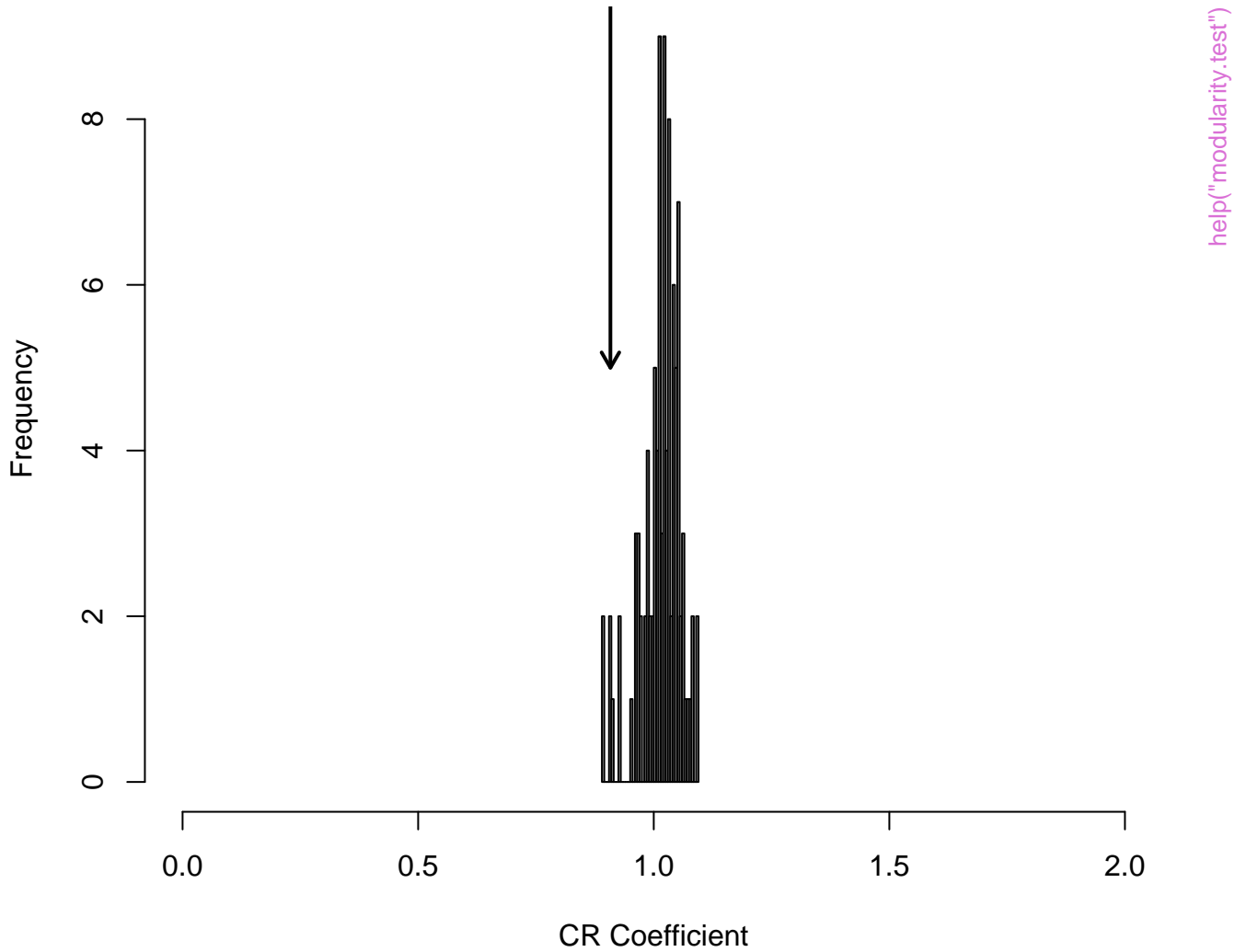




PLS1 Plot: Block 1 (X) vs. Block 2 (Y)

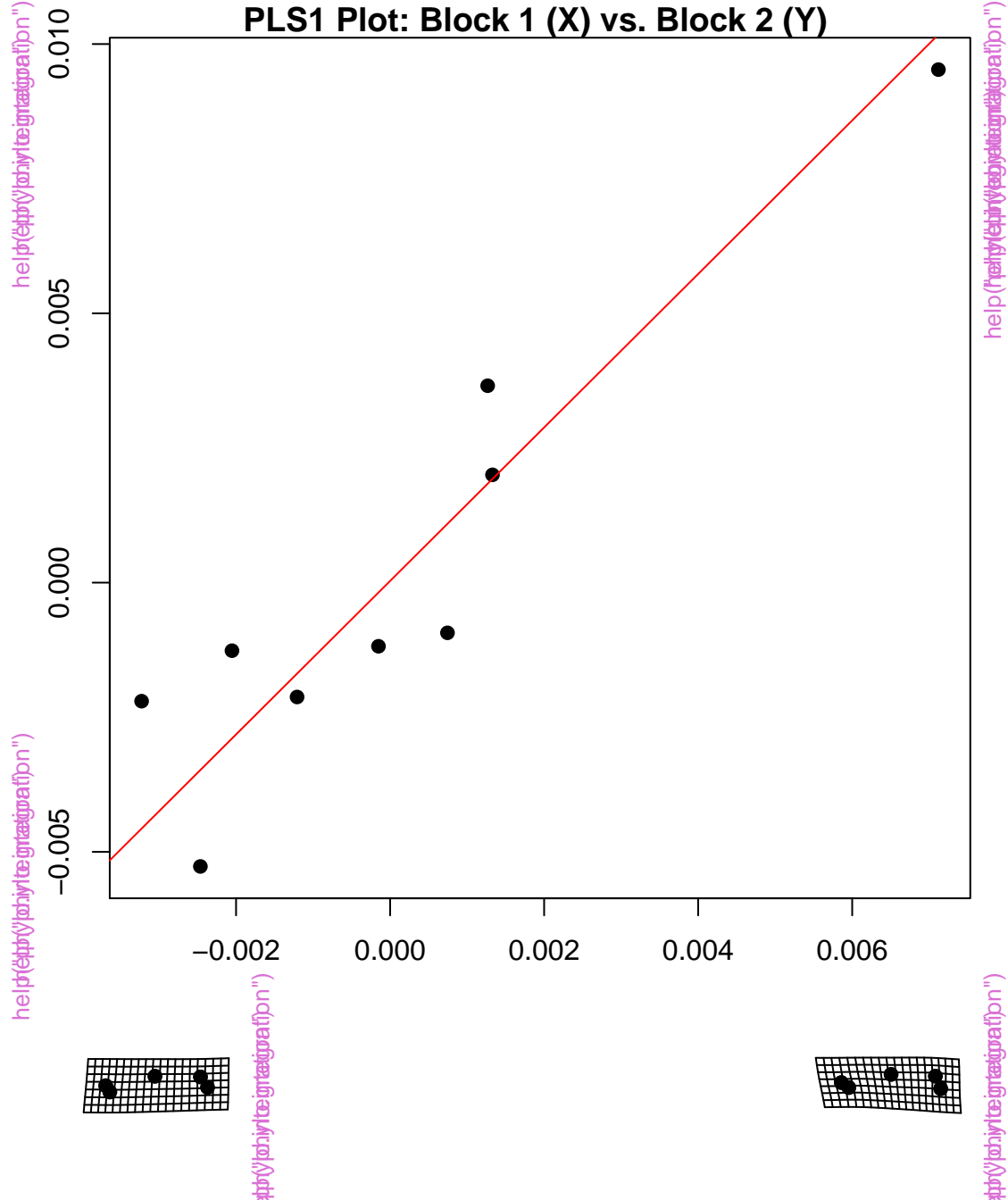


Observed CR = 0.908 ; P-value = 0.04

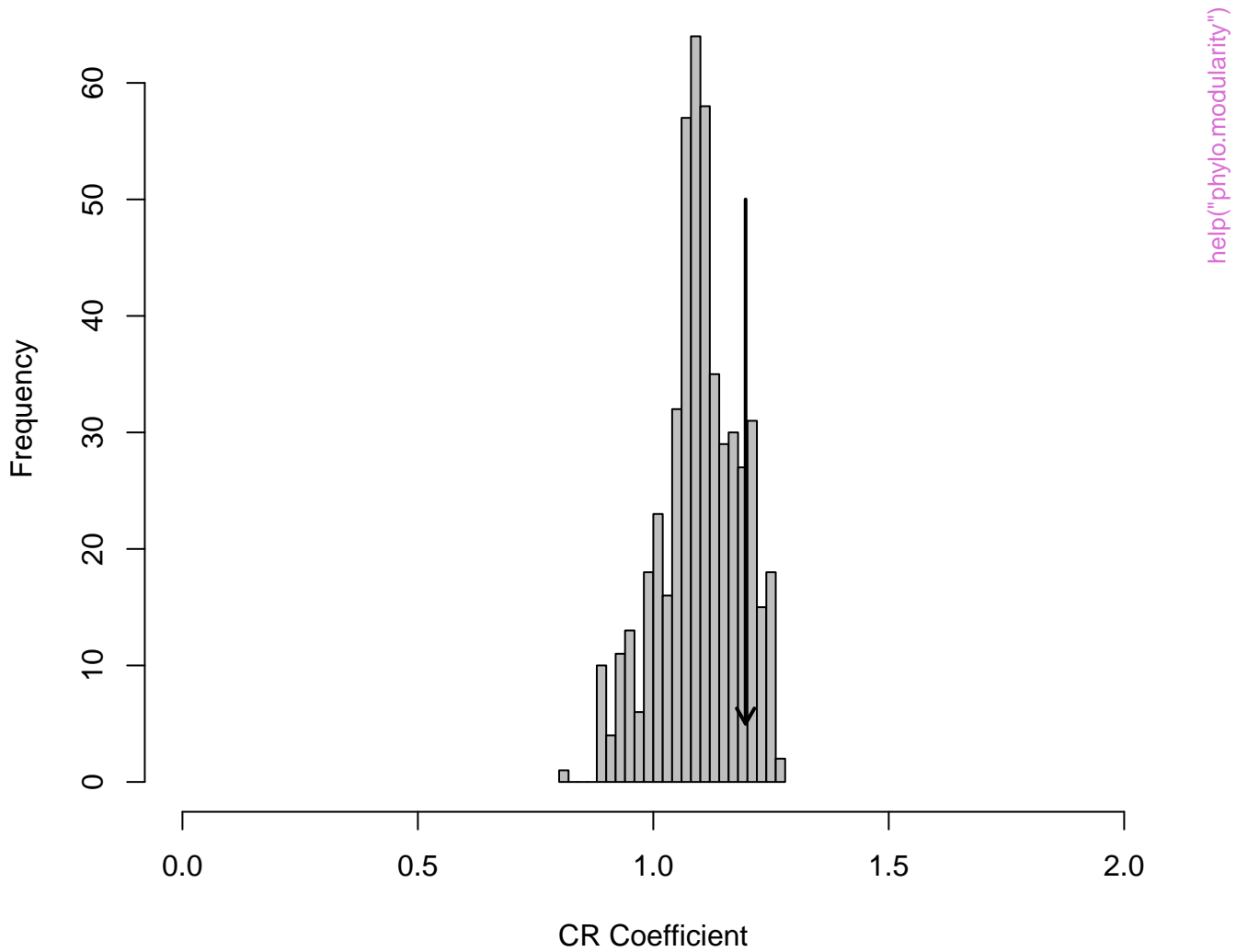


help("modularity.test")

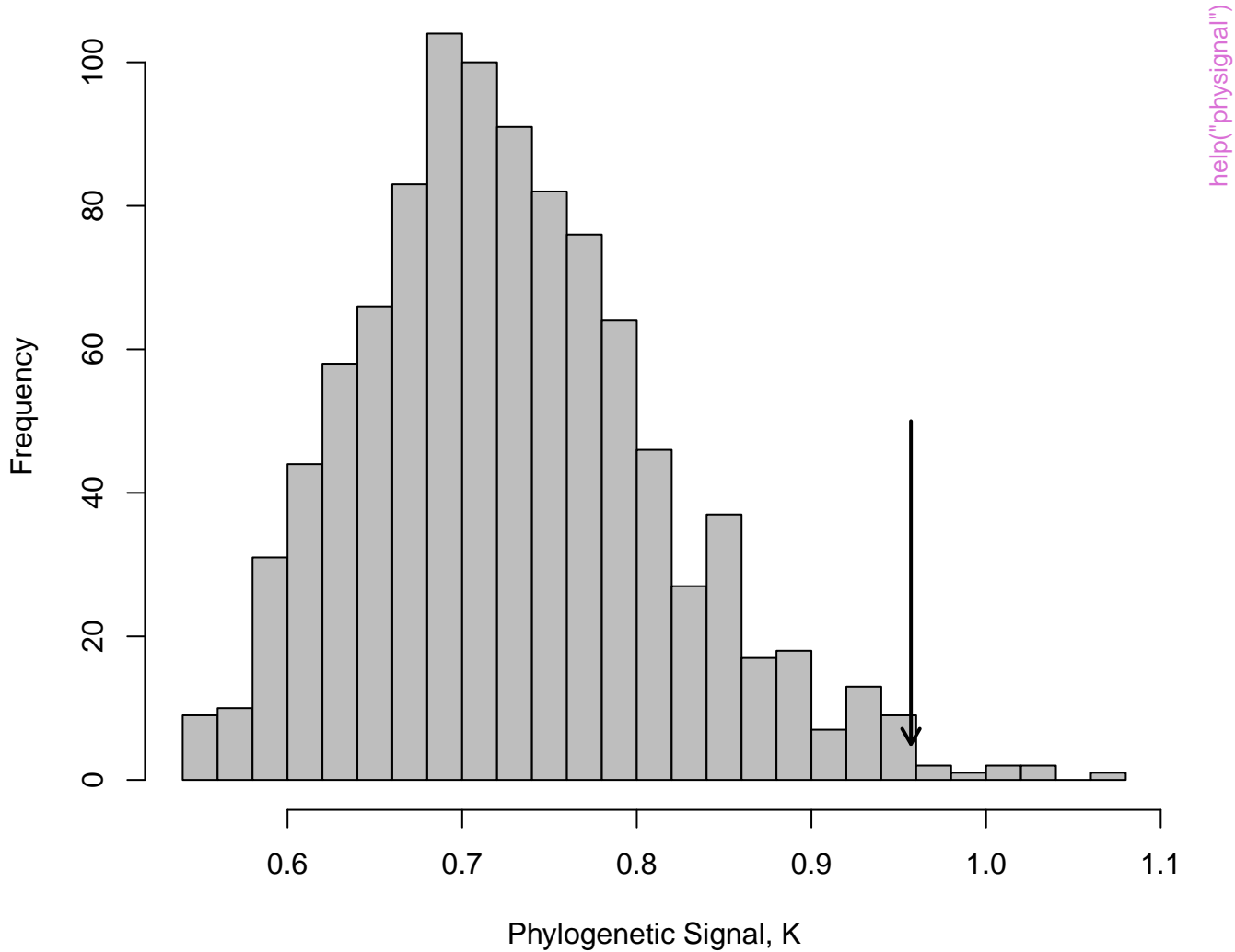
PLS1 Plot: Block 1 (X) vs. Block 2 (Y)



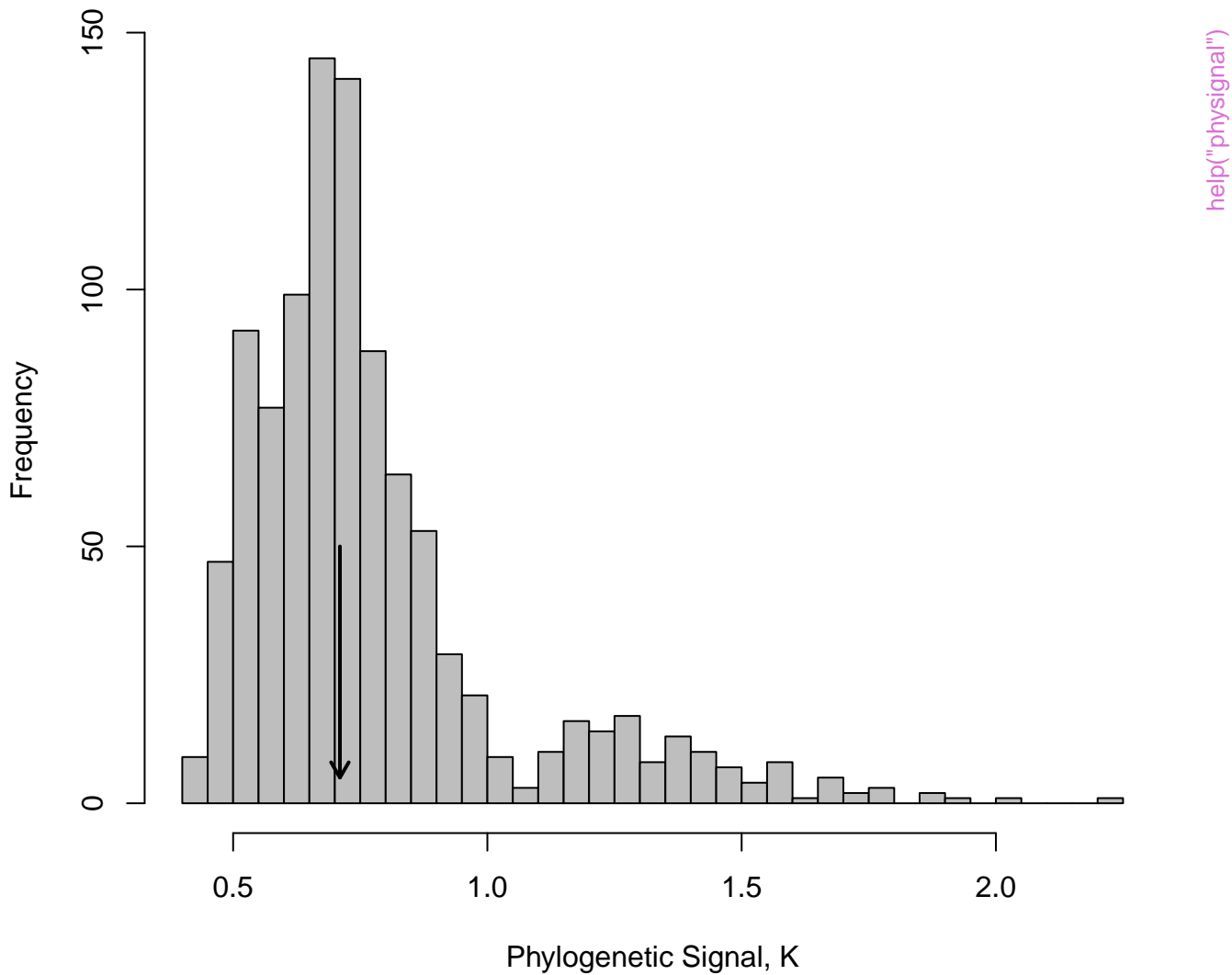
Observed CR = 1.196 ; P-value = 0.86

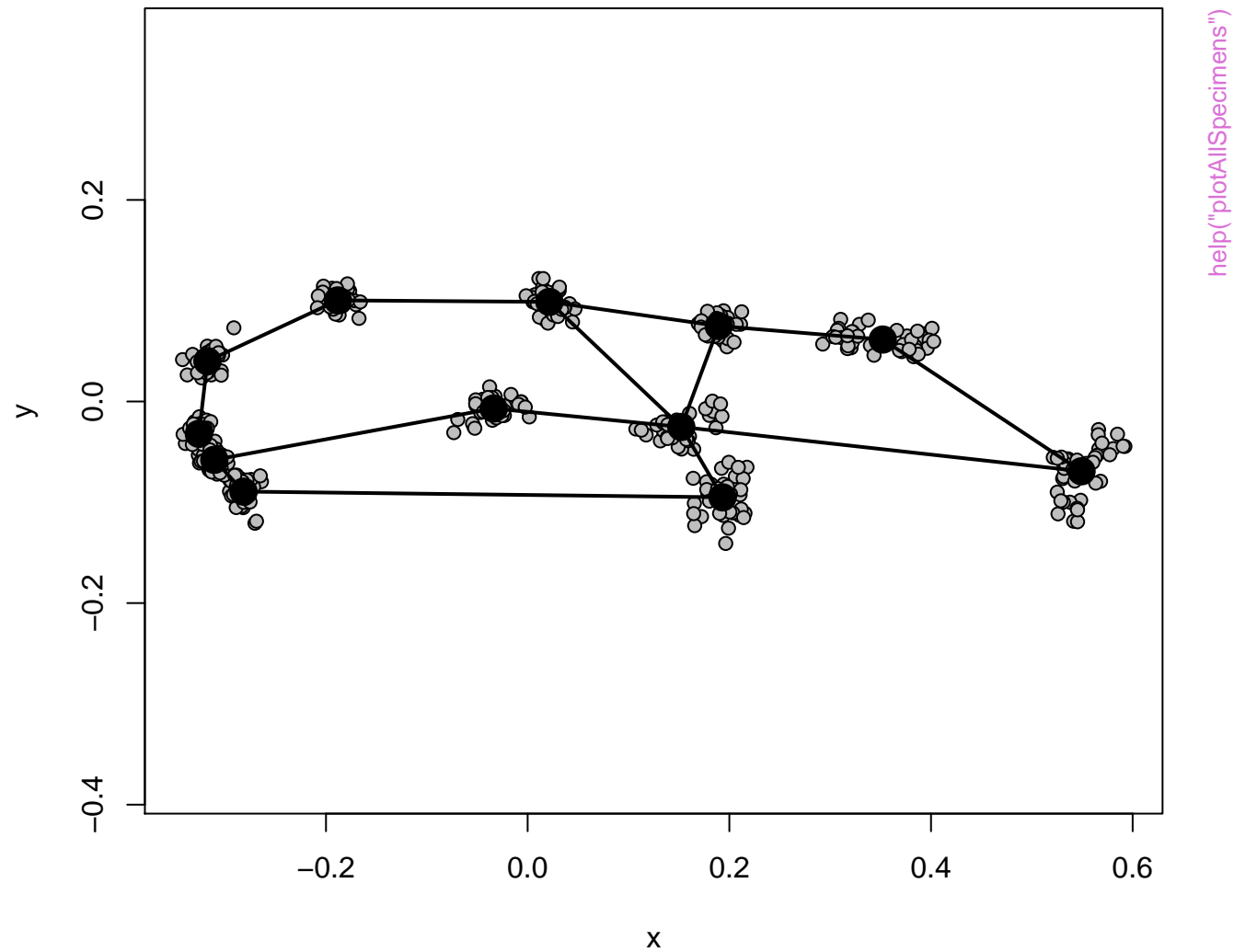


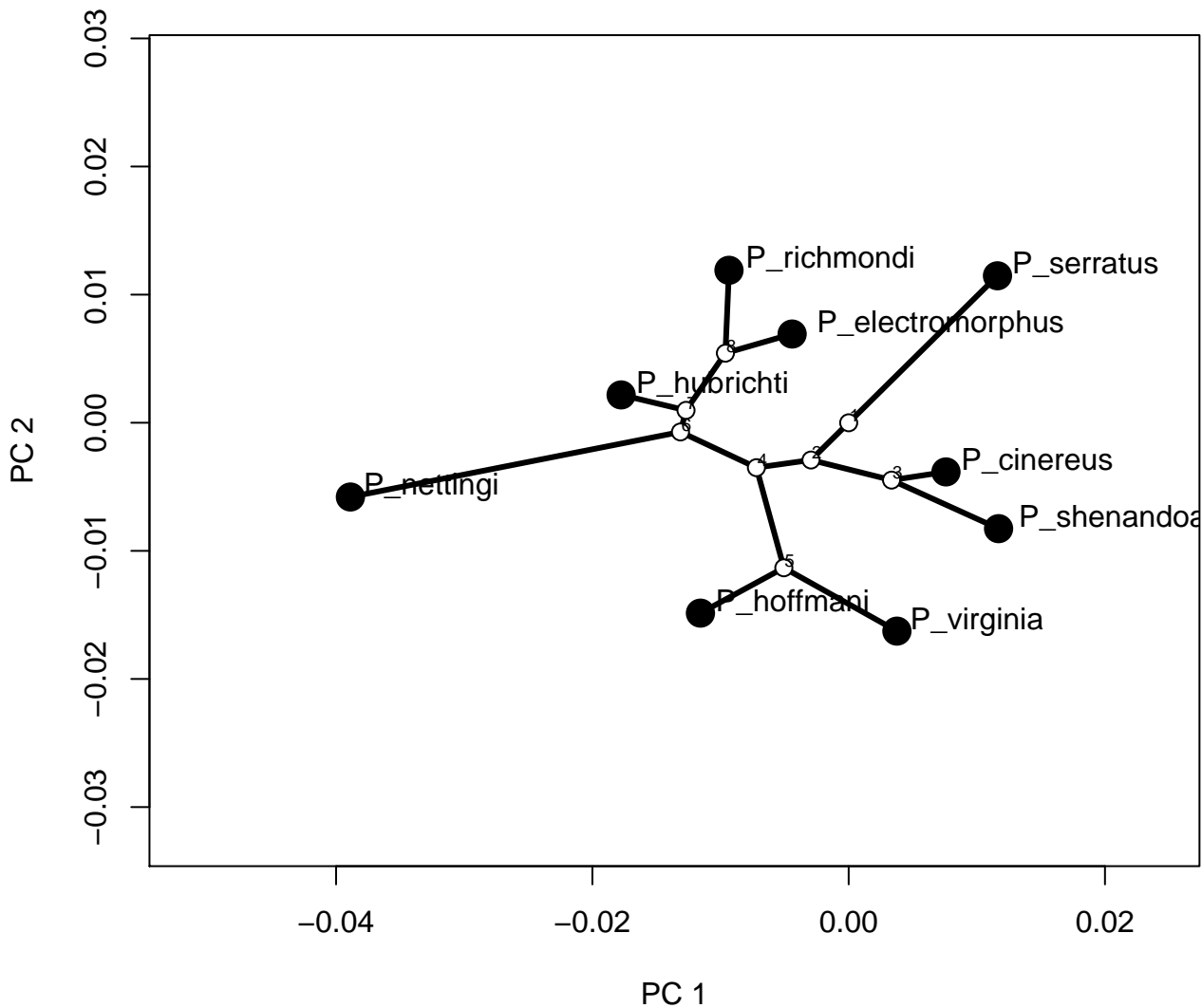
Observed K = 0.957 ; P-value = 0.011

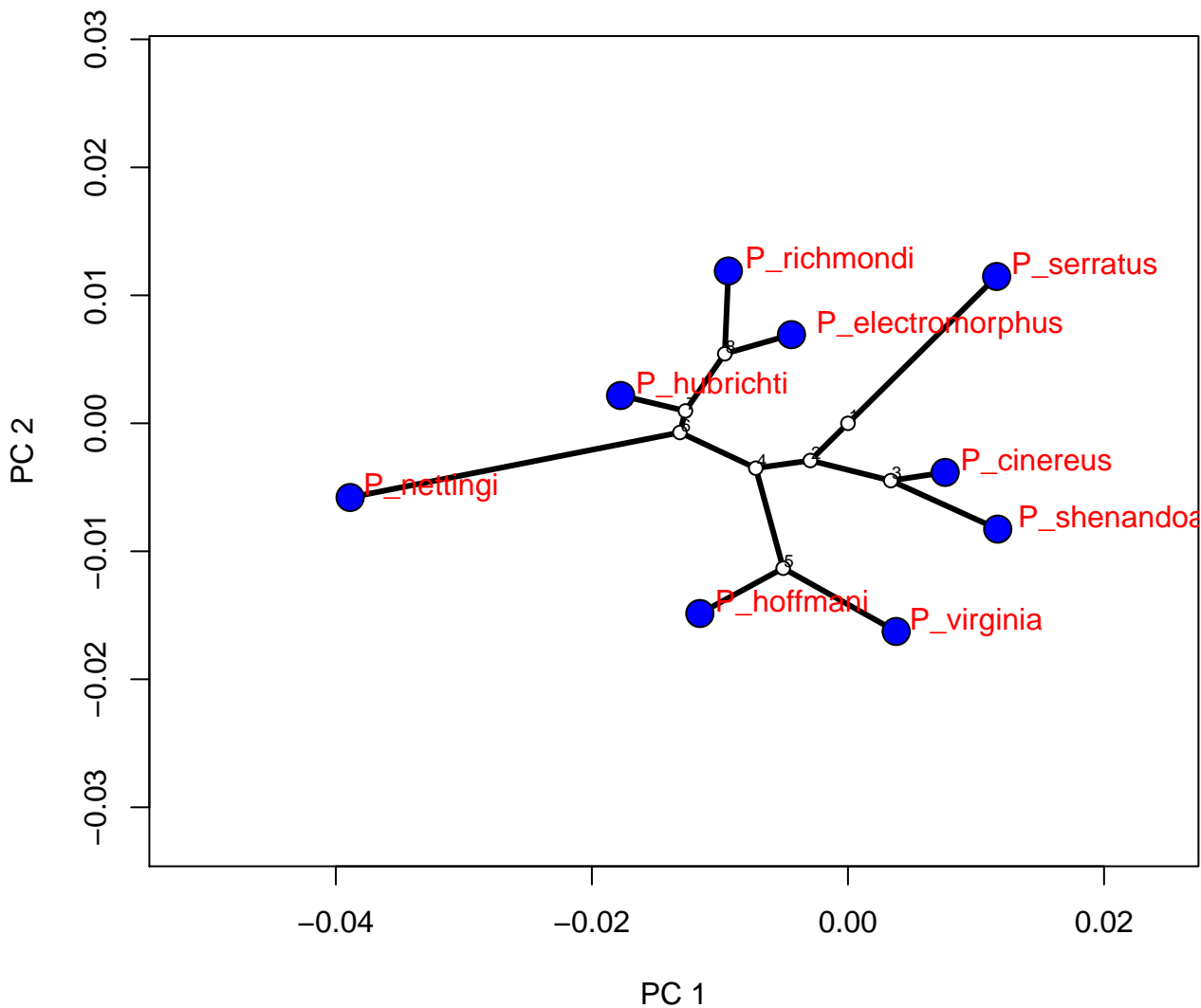


Observed K = 0.71 ; P-value = 0.497

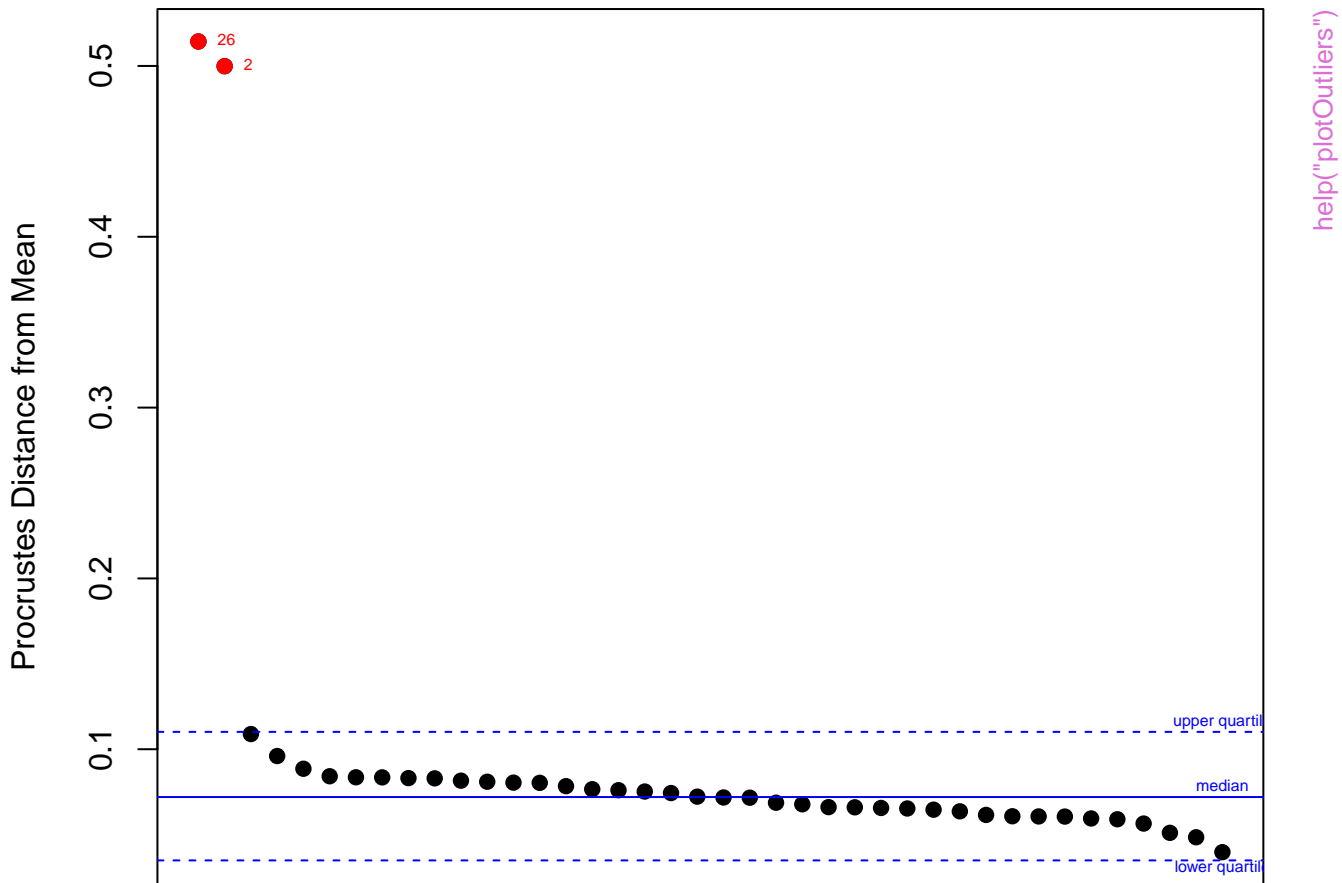






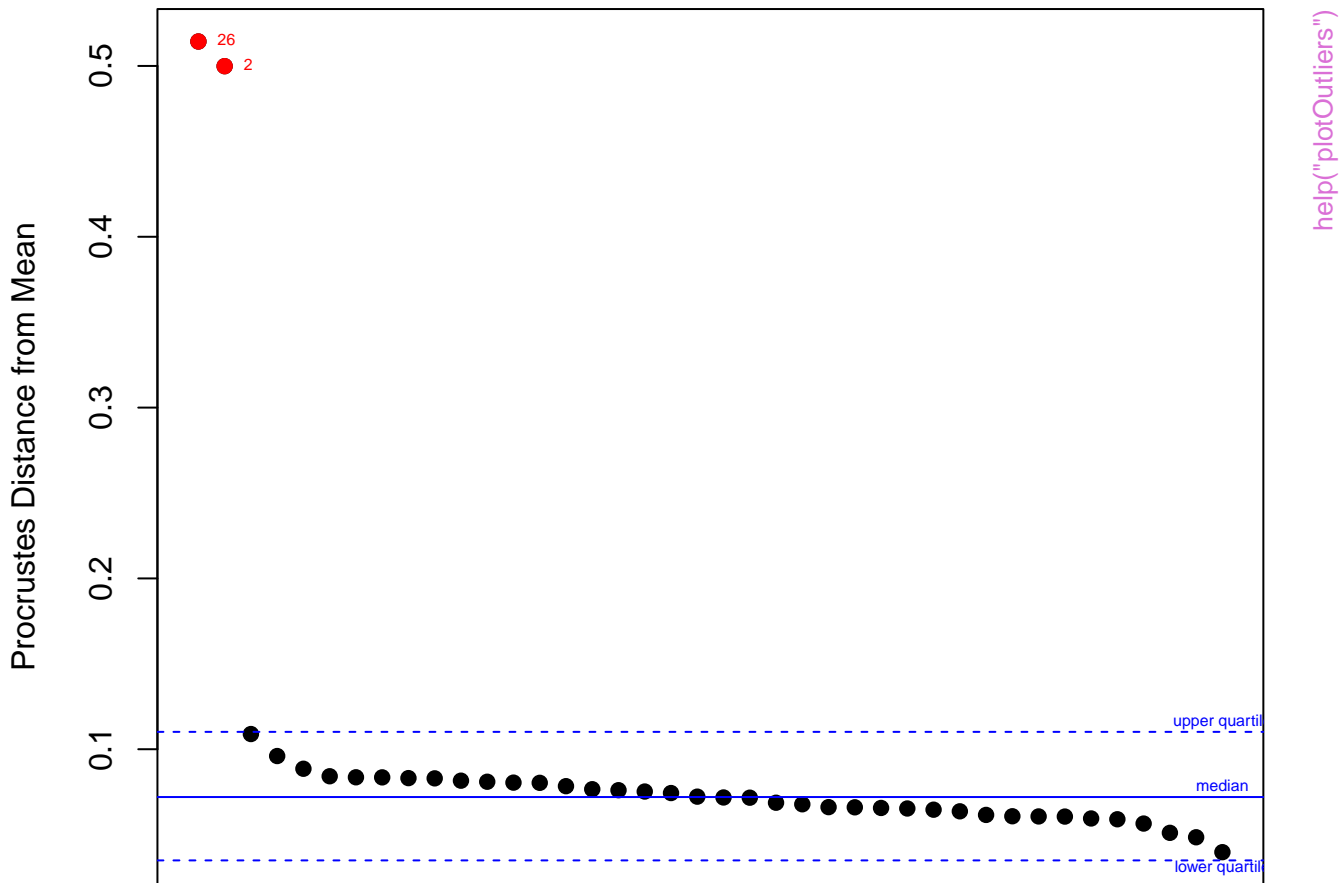


All Specimens



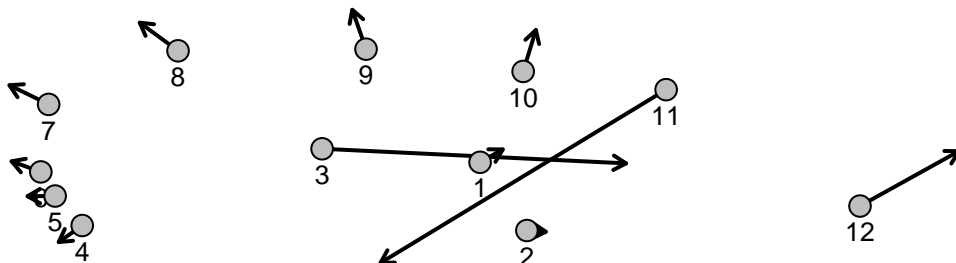
help("plotOutliers")

All Specimens

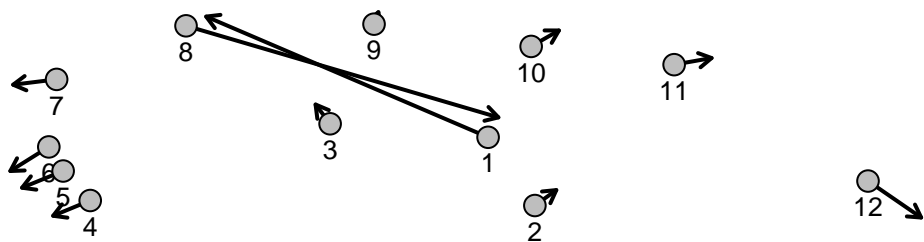


help("plotOutliers")

group: All Specimens, specimen: 26

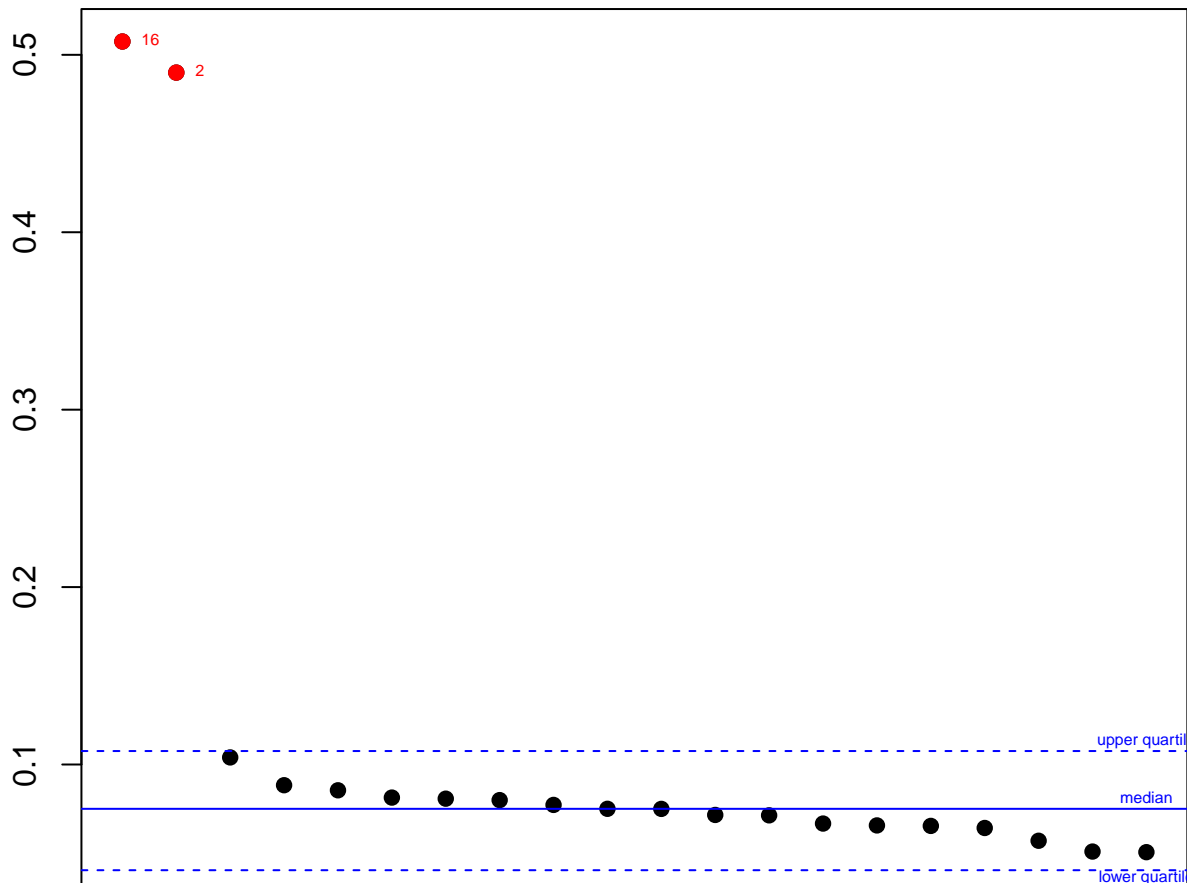


group: All Specimens, specimen: 2



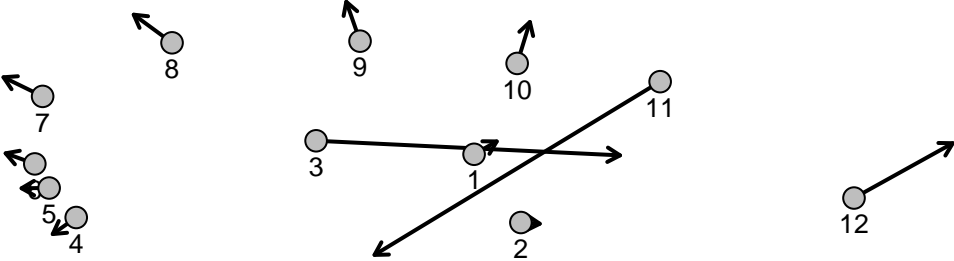
Jord

Procrustes Distance from Mean

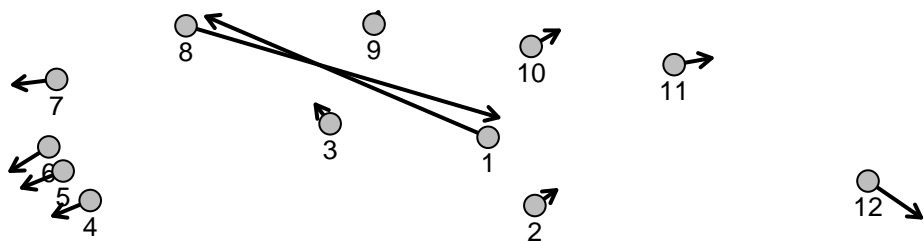


help("plotOutliers")

group: Jord, specimen: 16

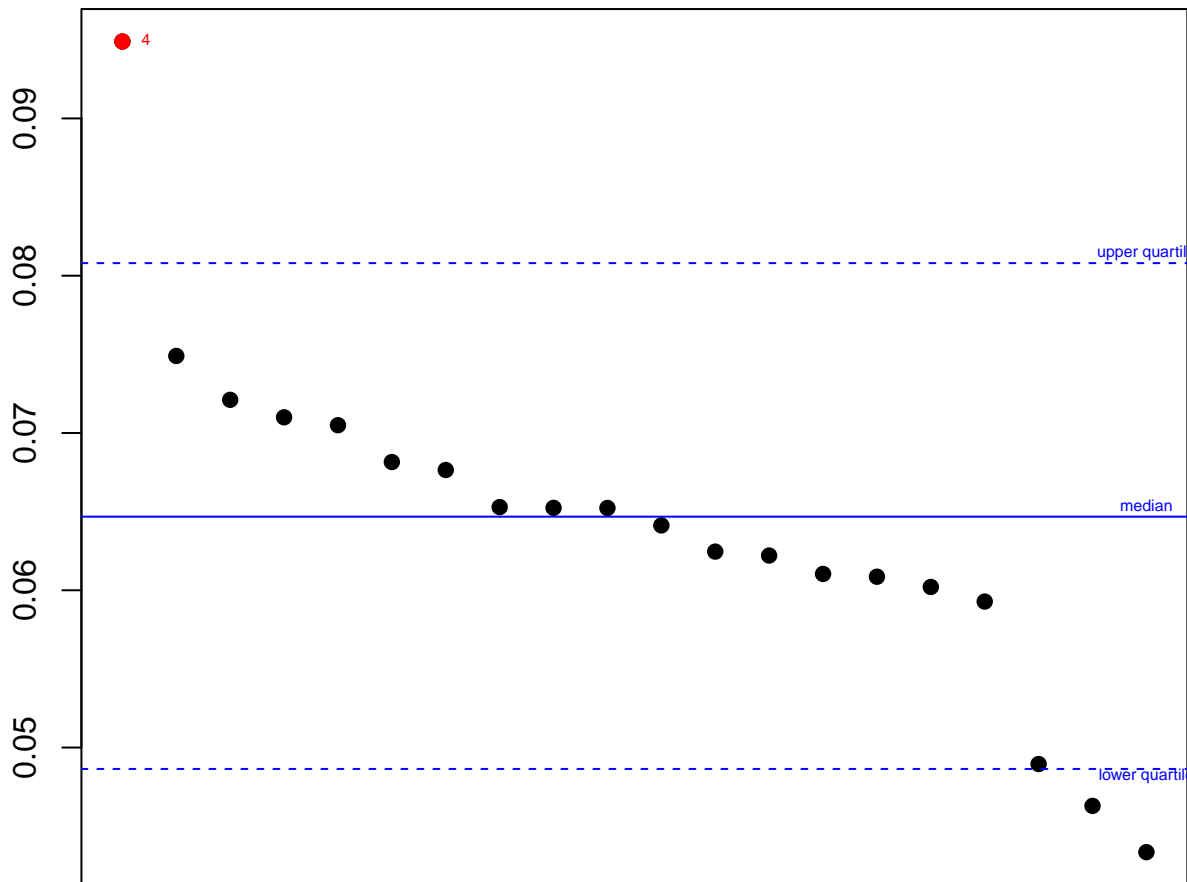


group: Jord, specimen: 2



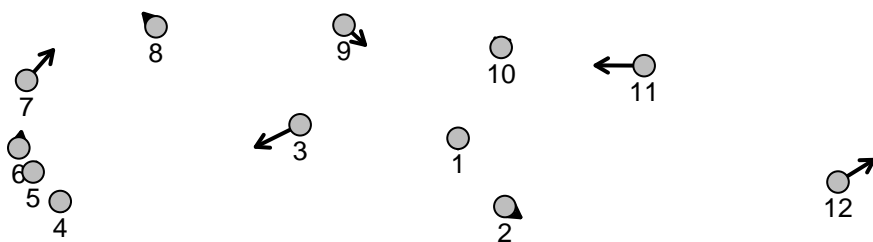
Teyah

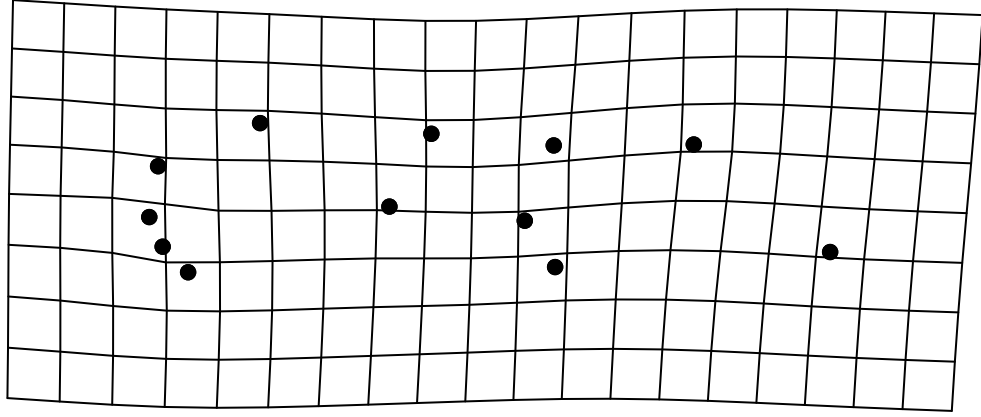
Procrustes Distance from Mean

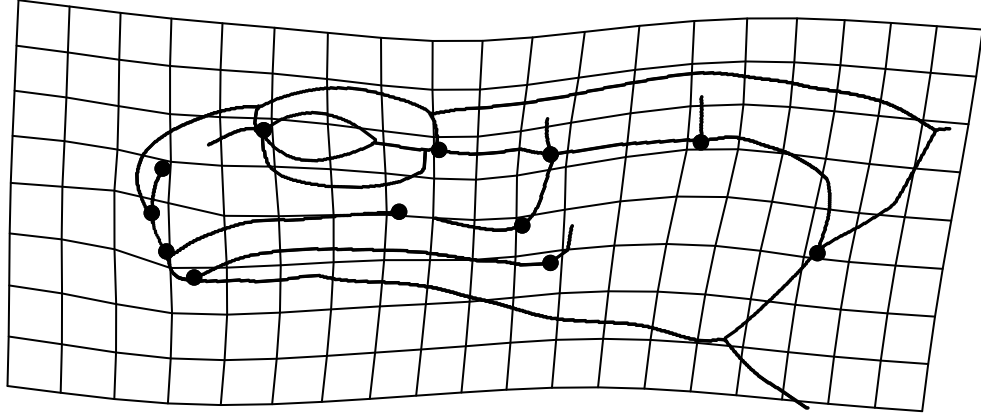


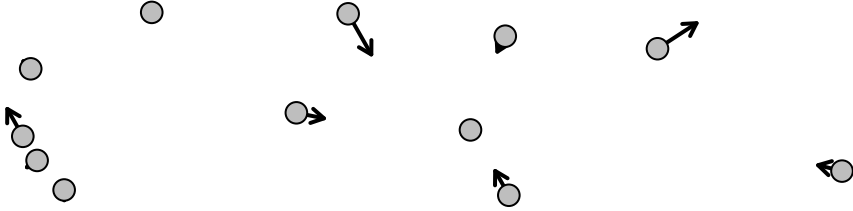
help("plotOutliers")

group: Teyah, specimen: 4

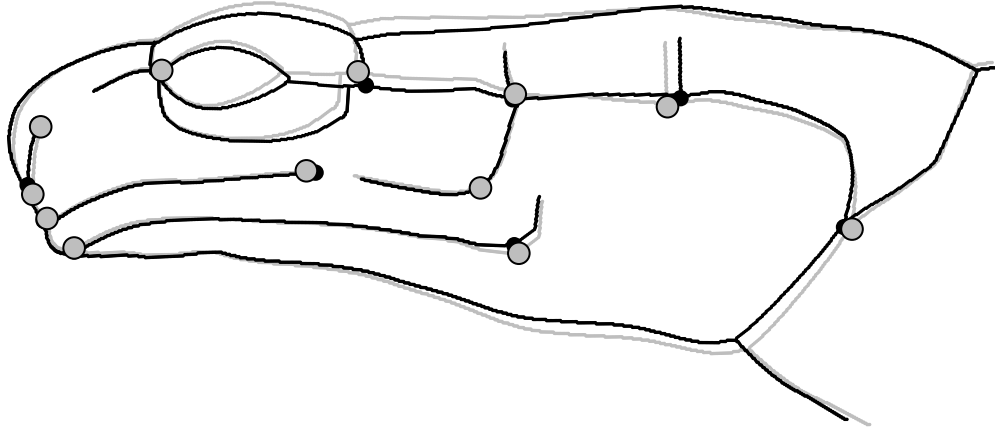


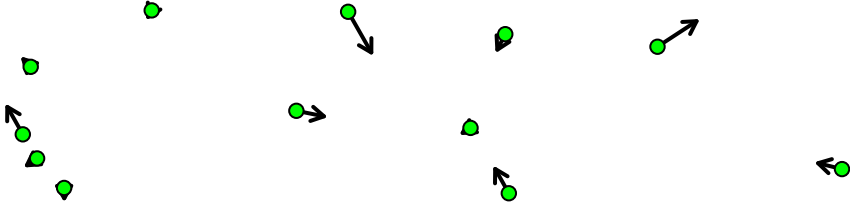


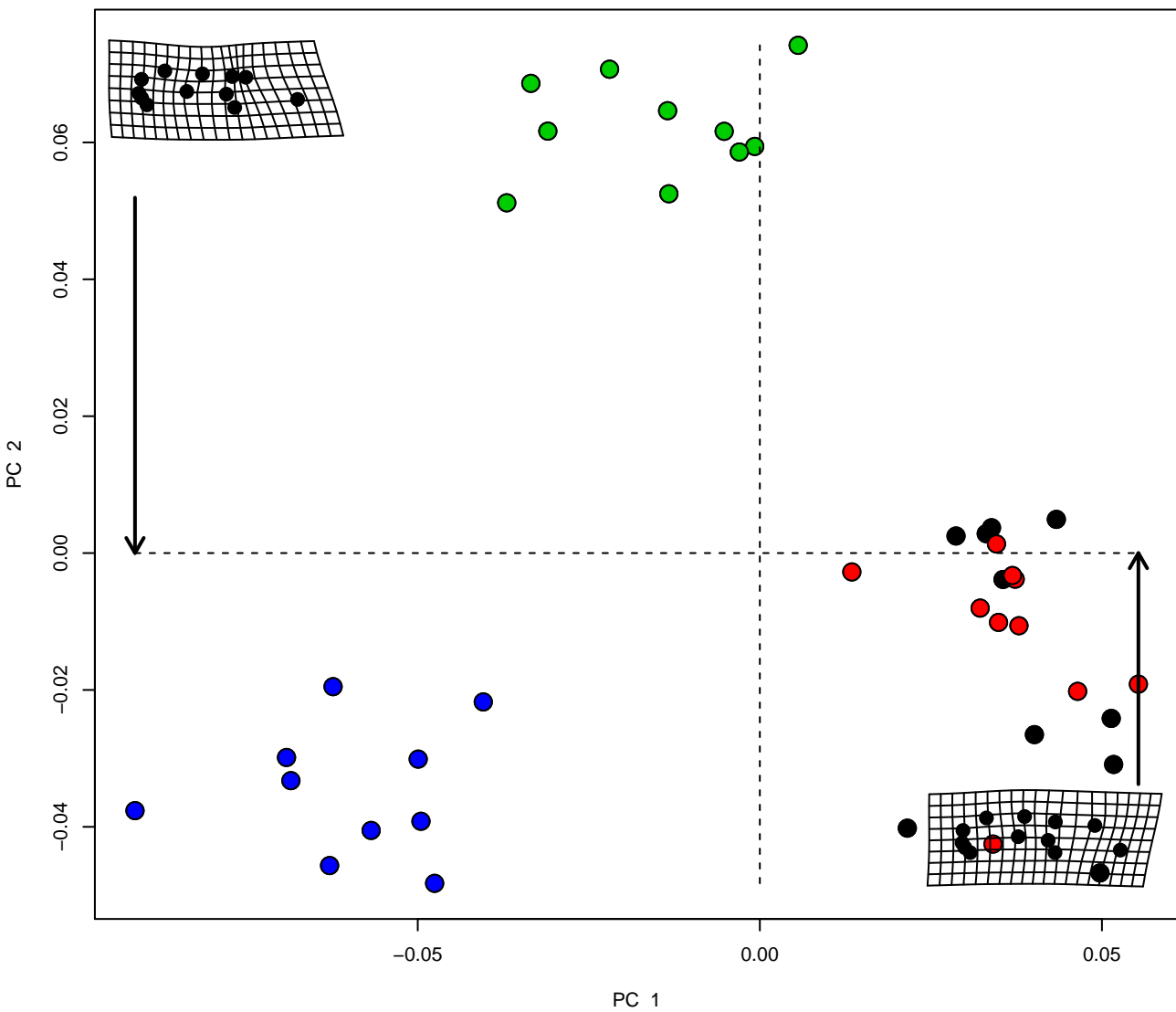


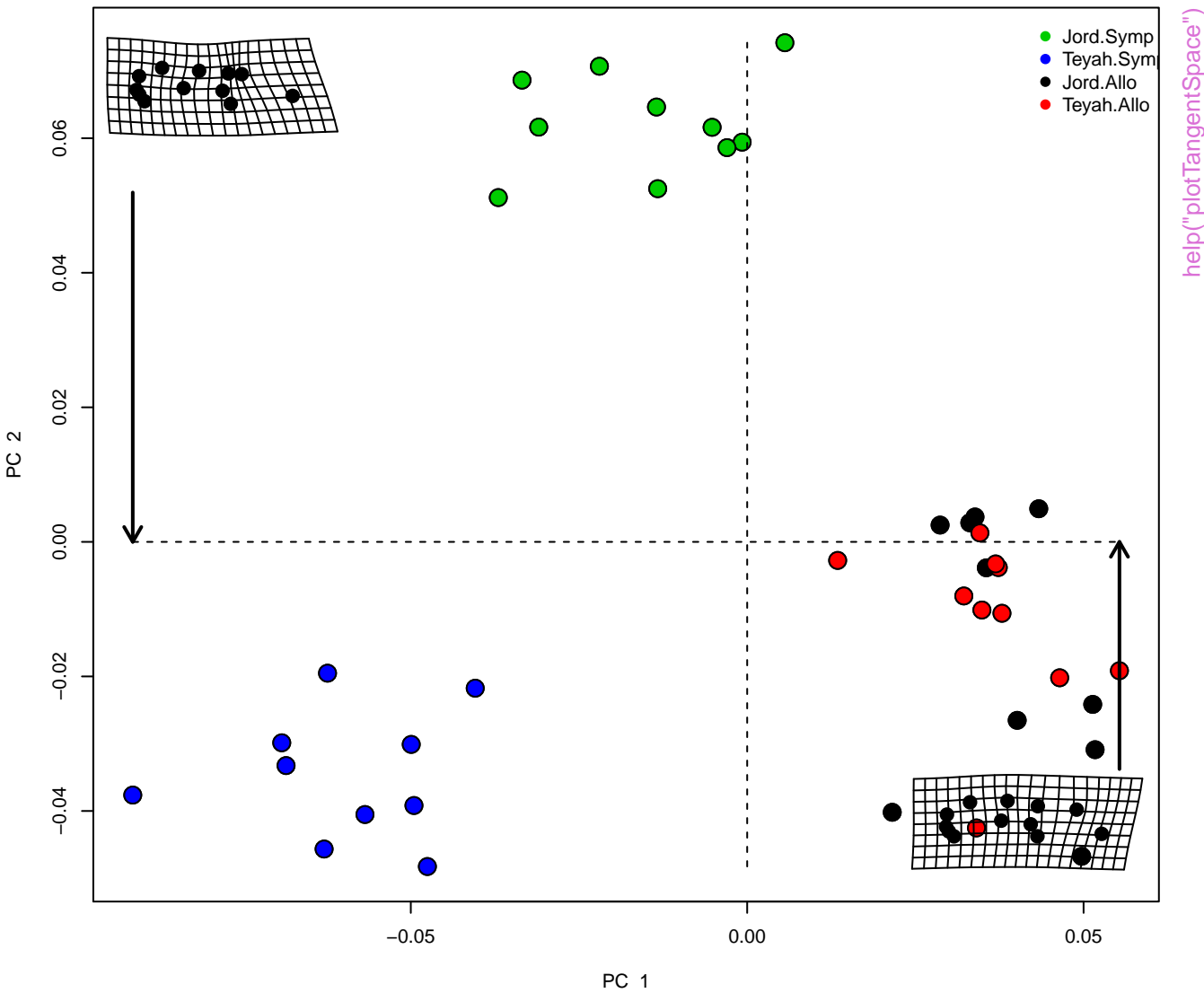


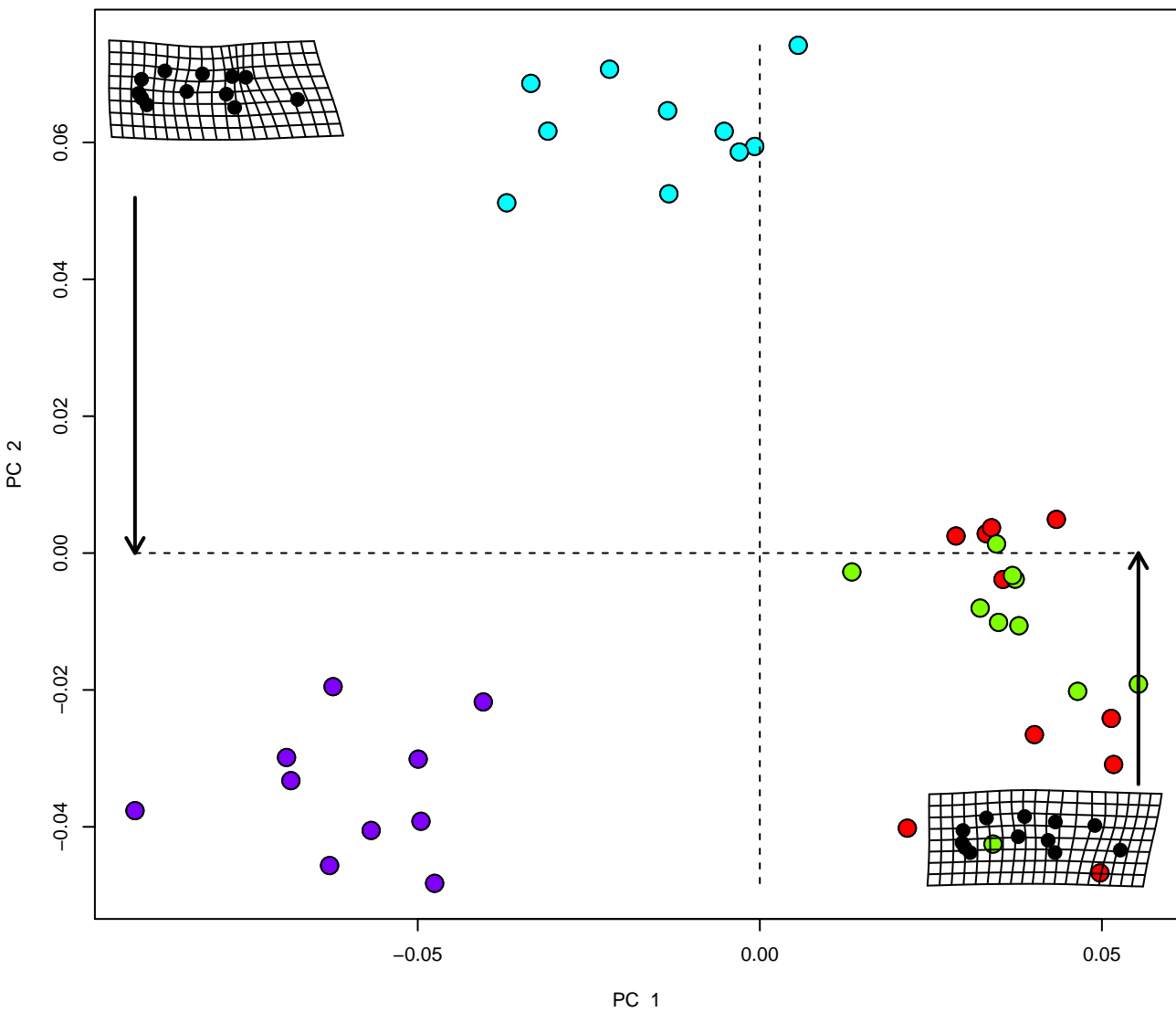
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help("plotRefToTarget")
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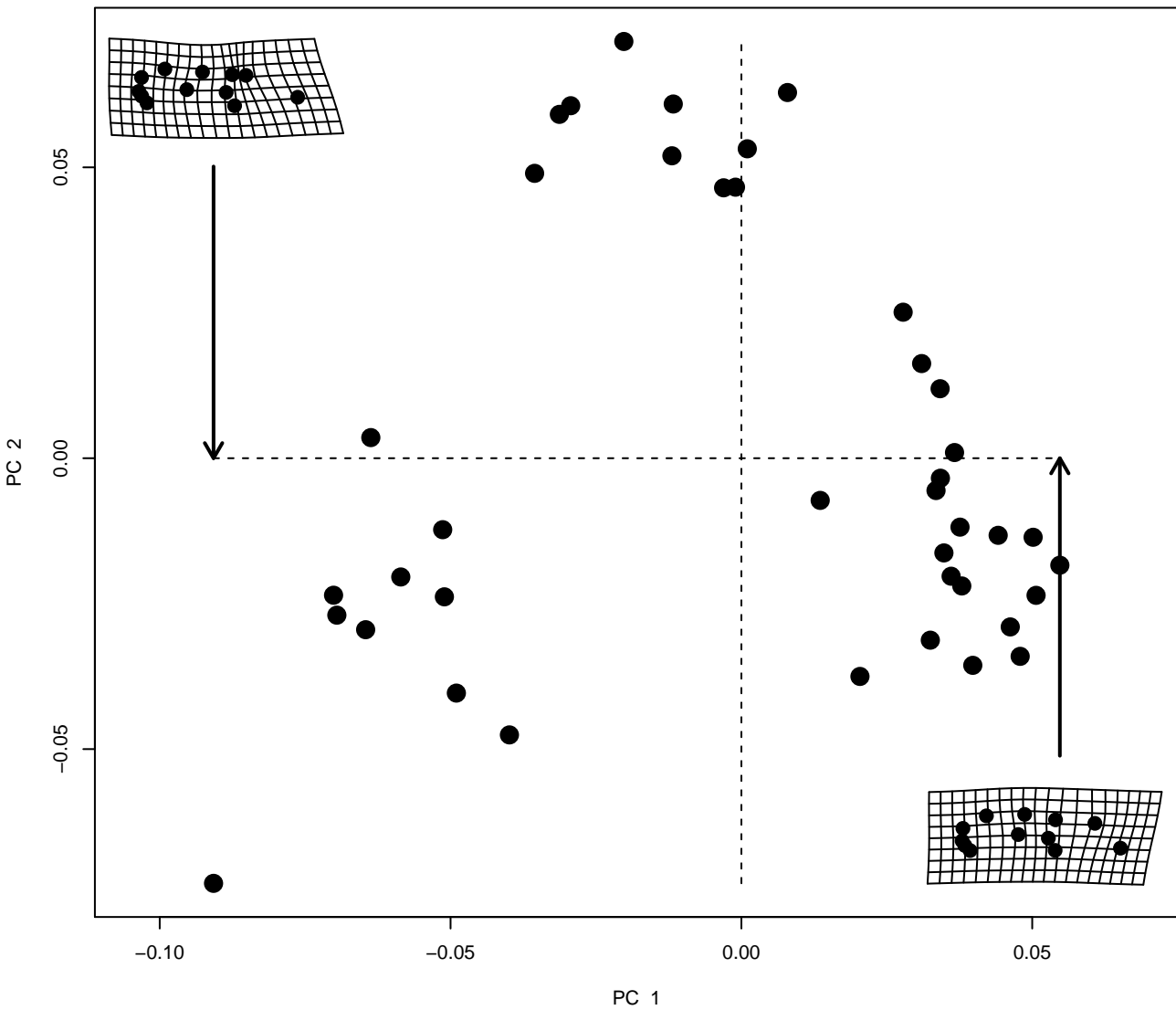



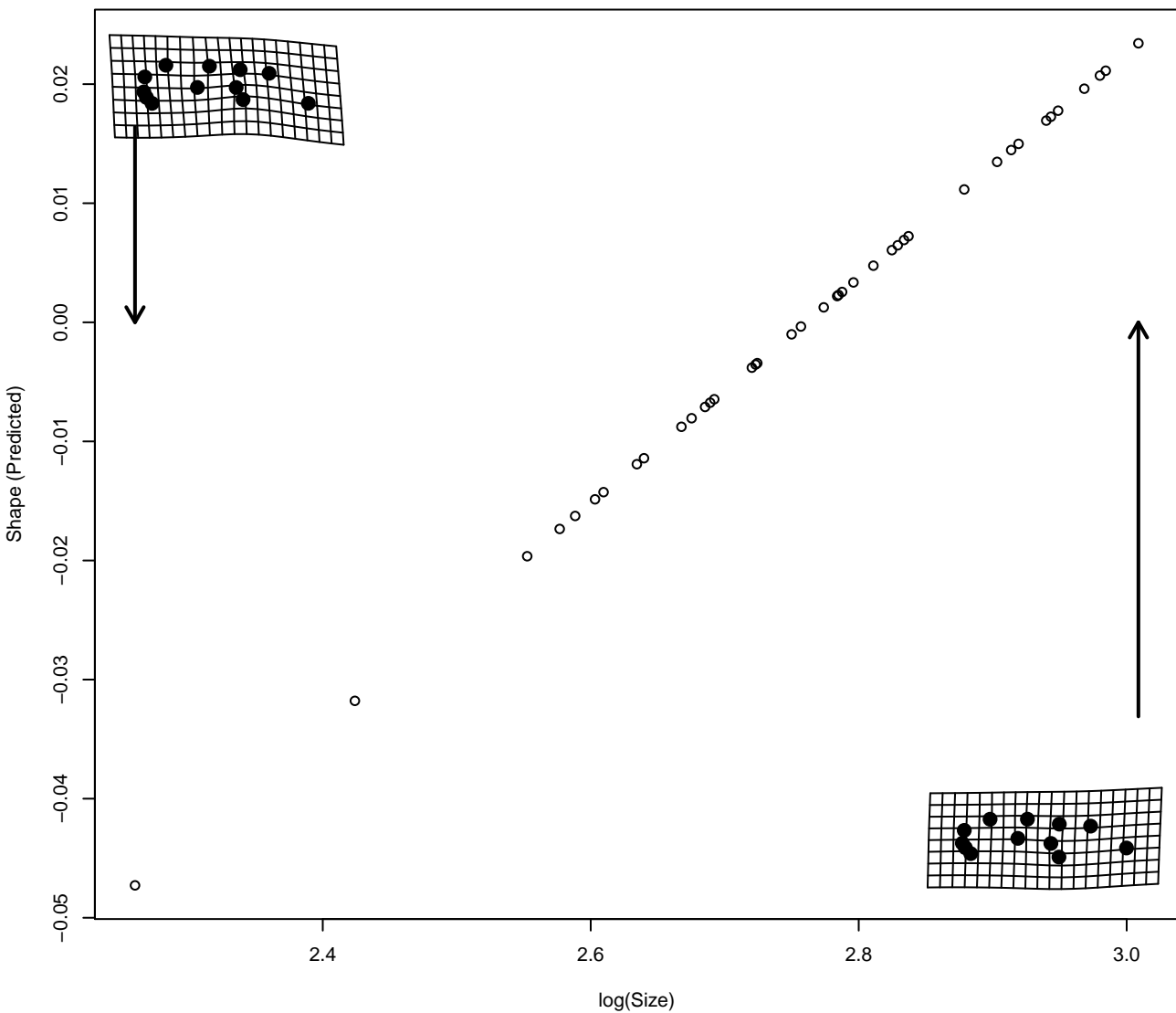


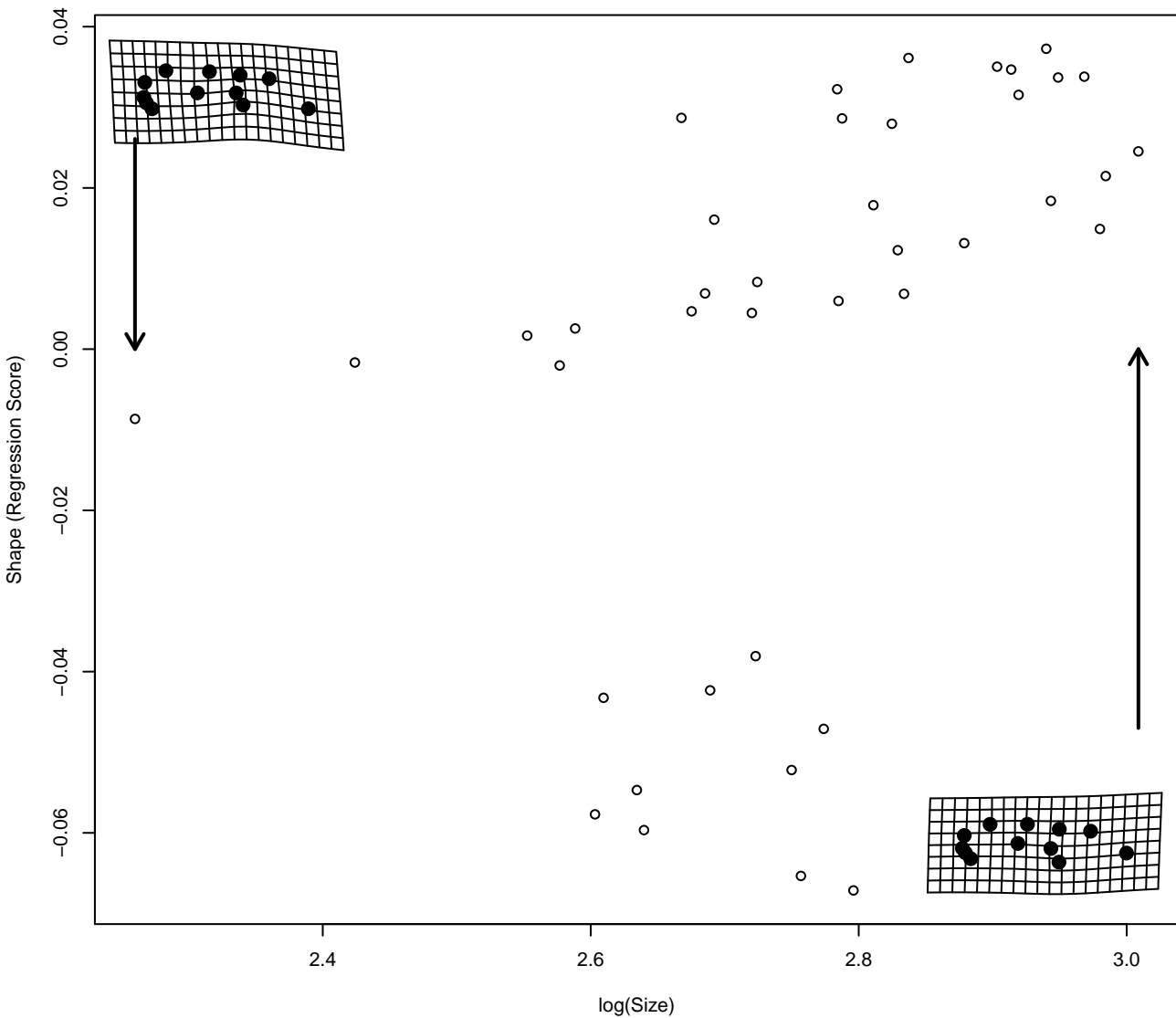


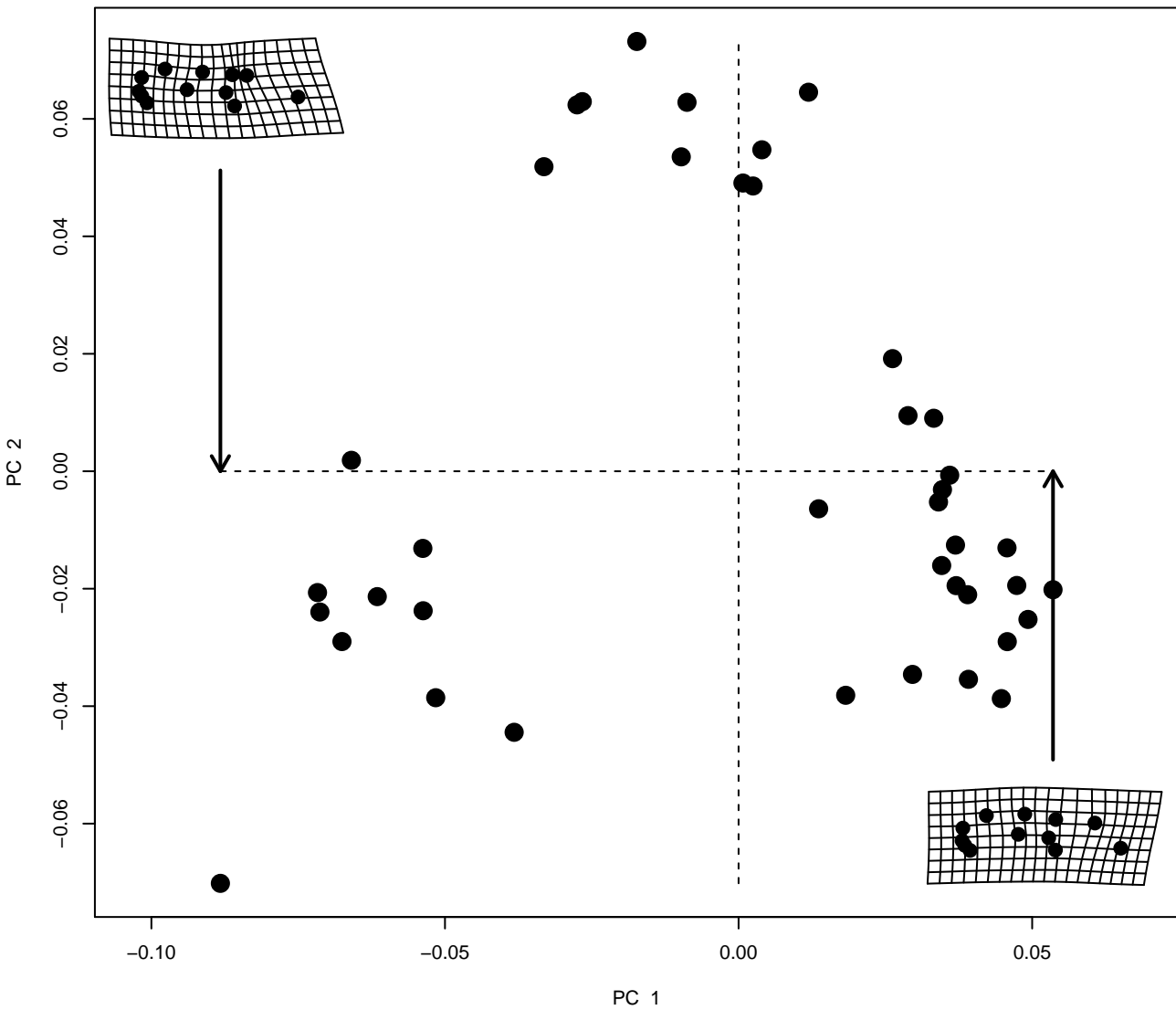


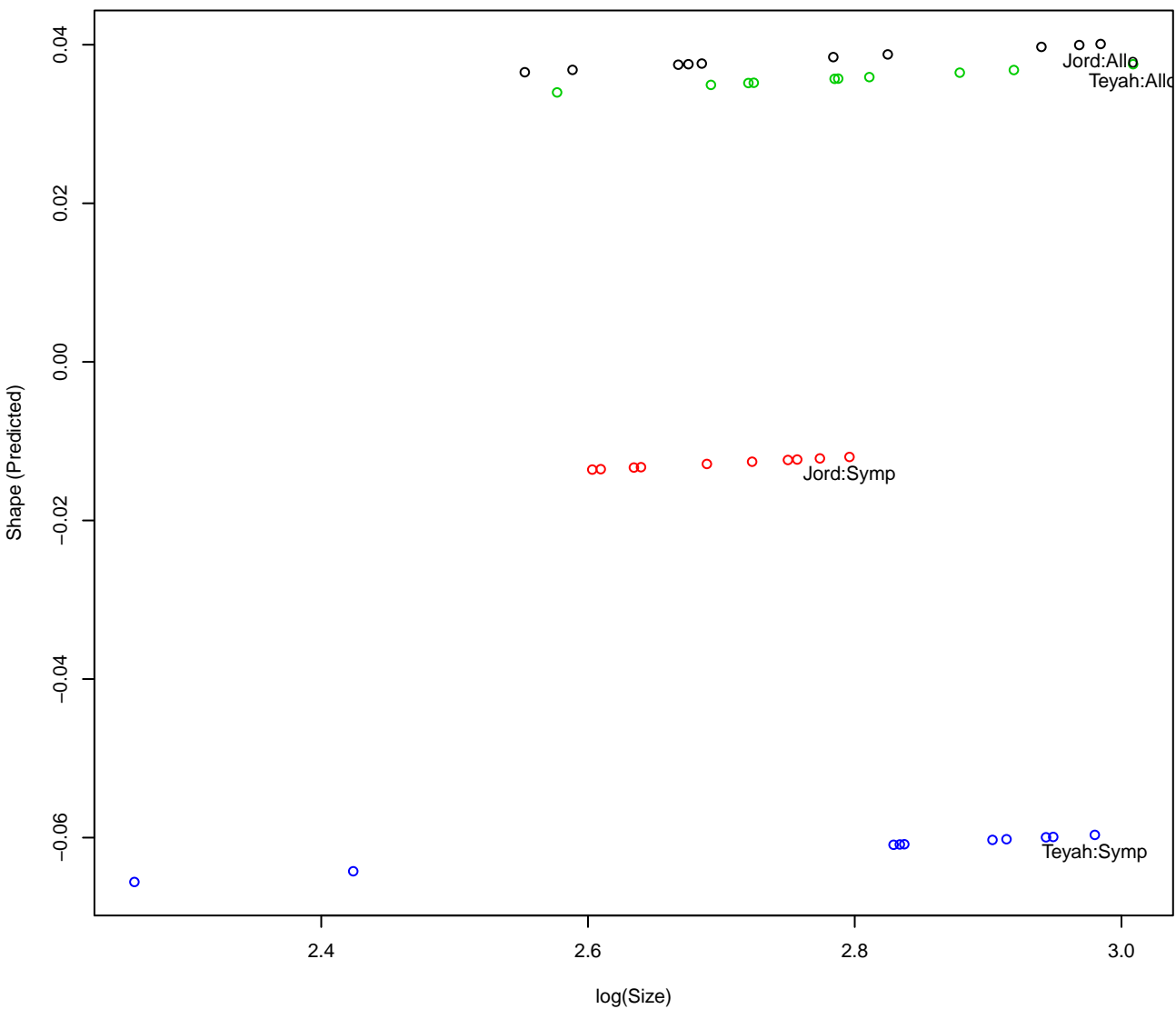


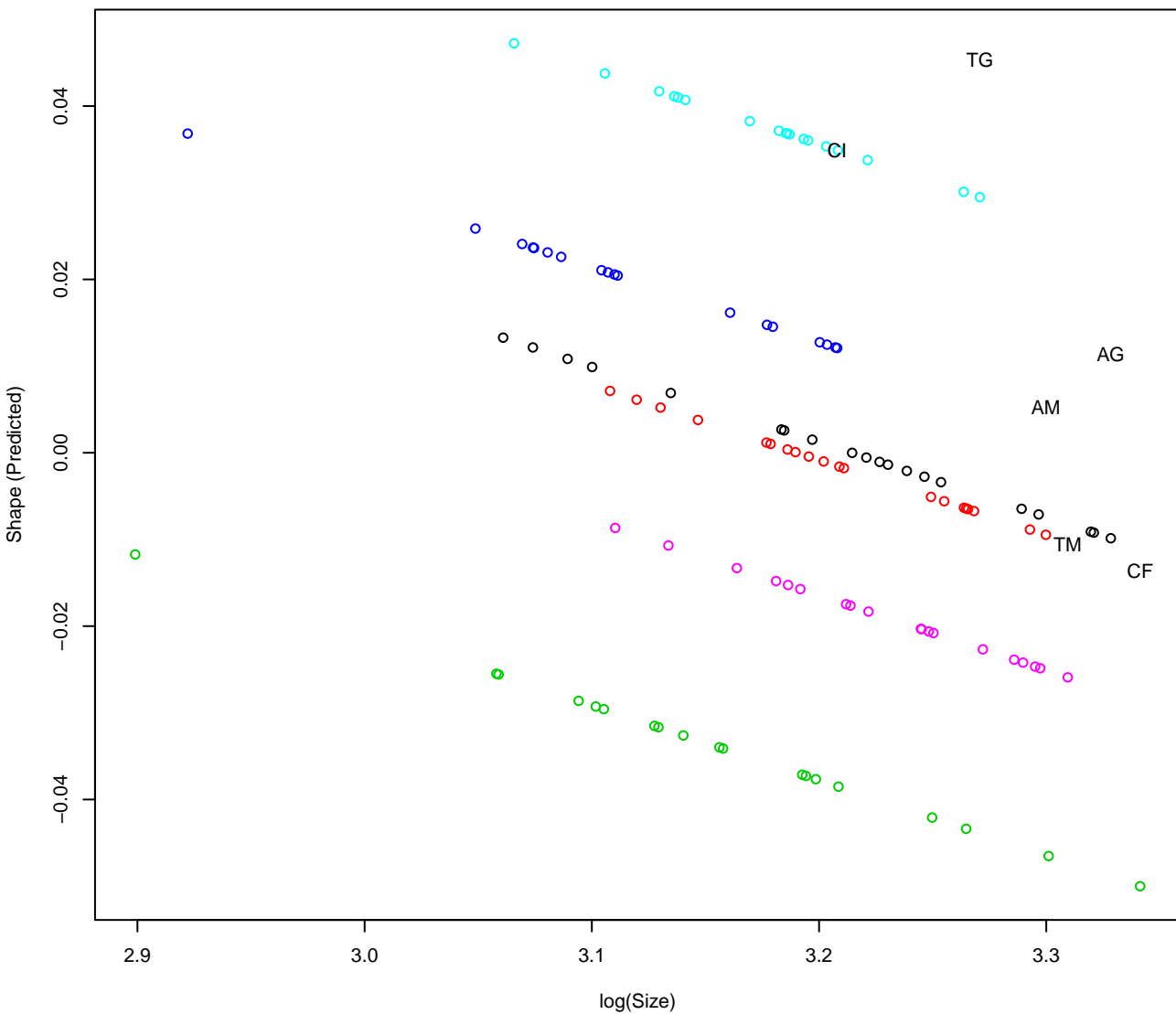




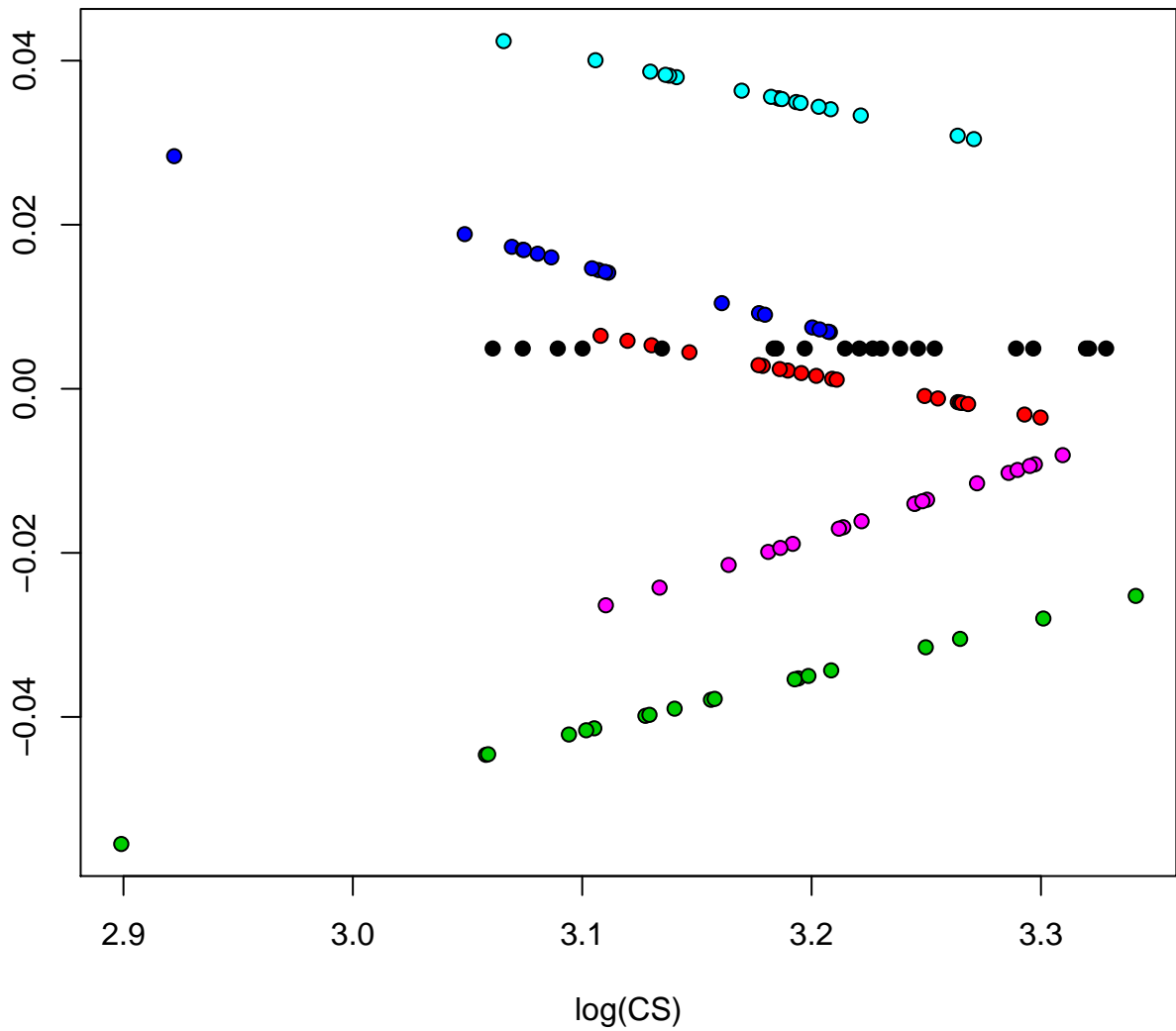






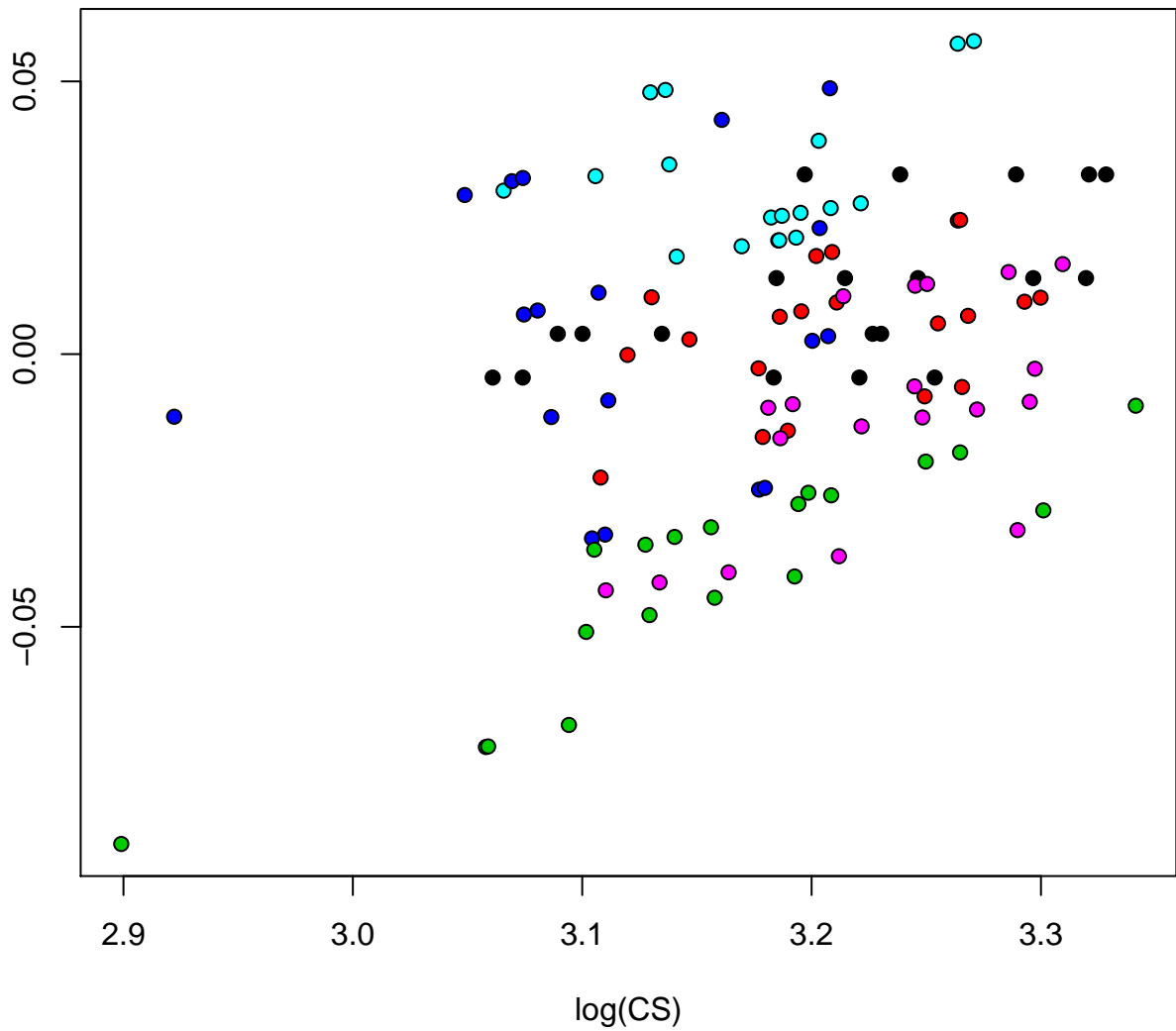


PC 1 for fitted values



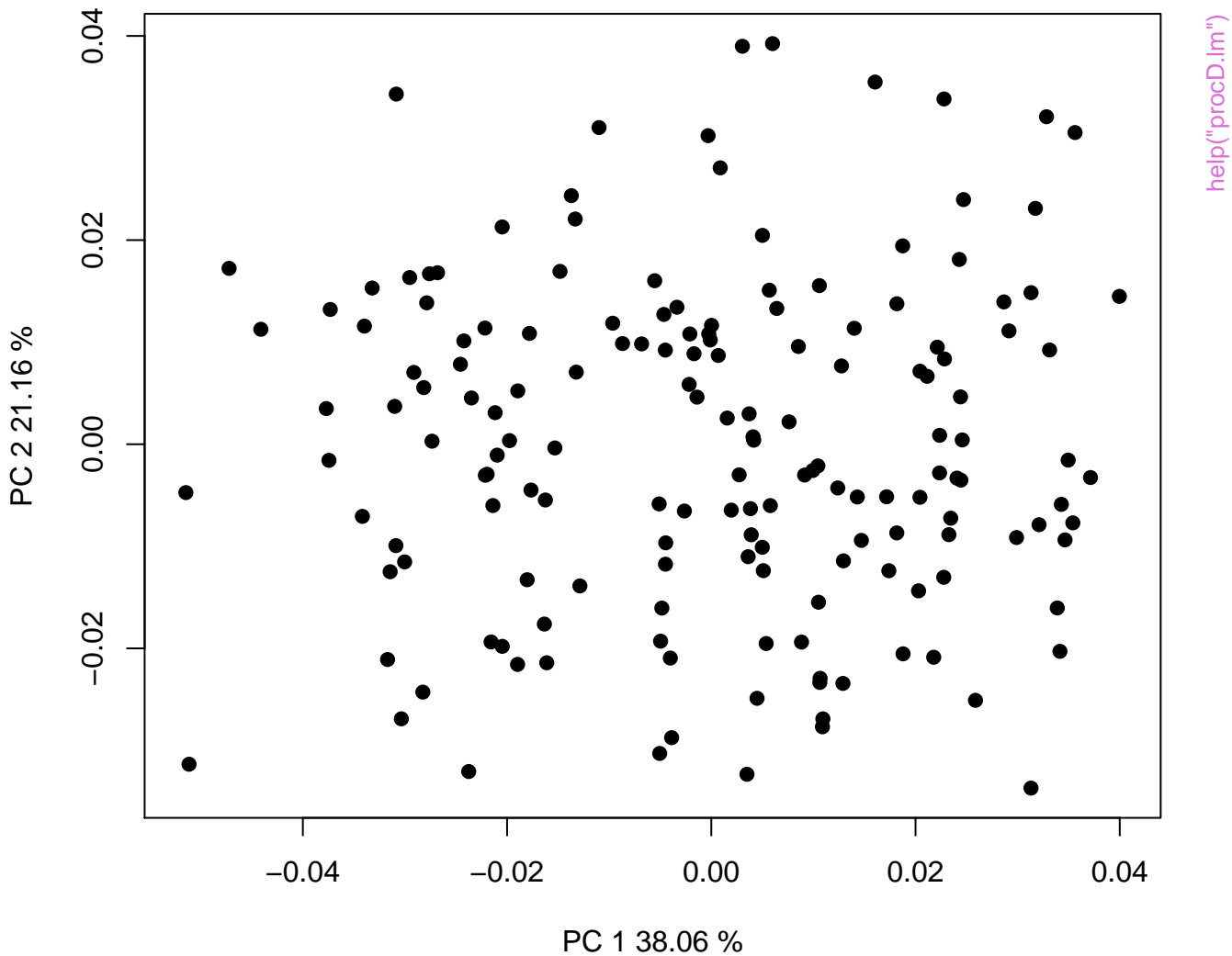
help("procD.allometry")

PC 1 for fitted values

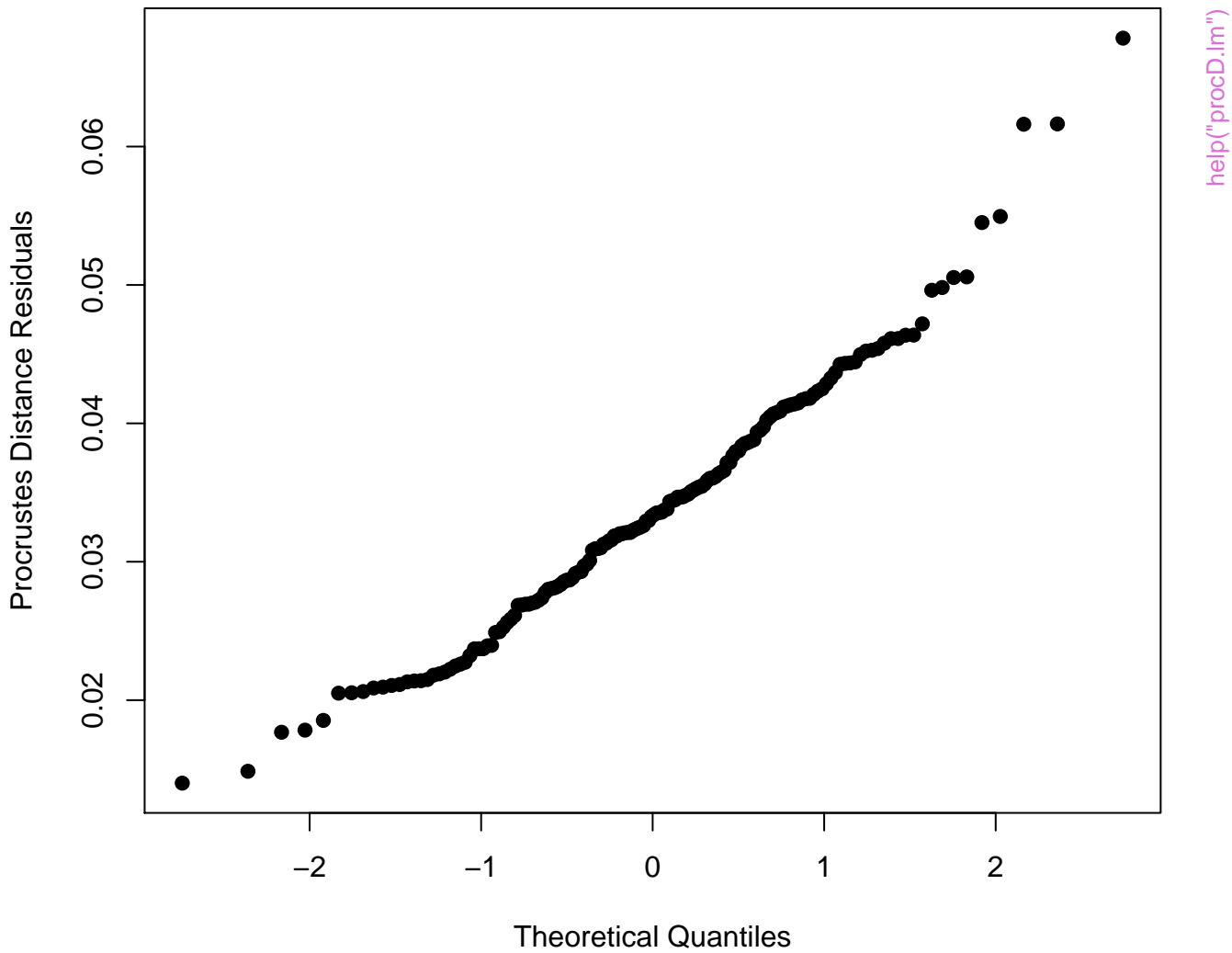


help("procD.allometry")

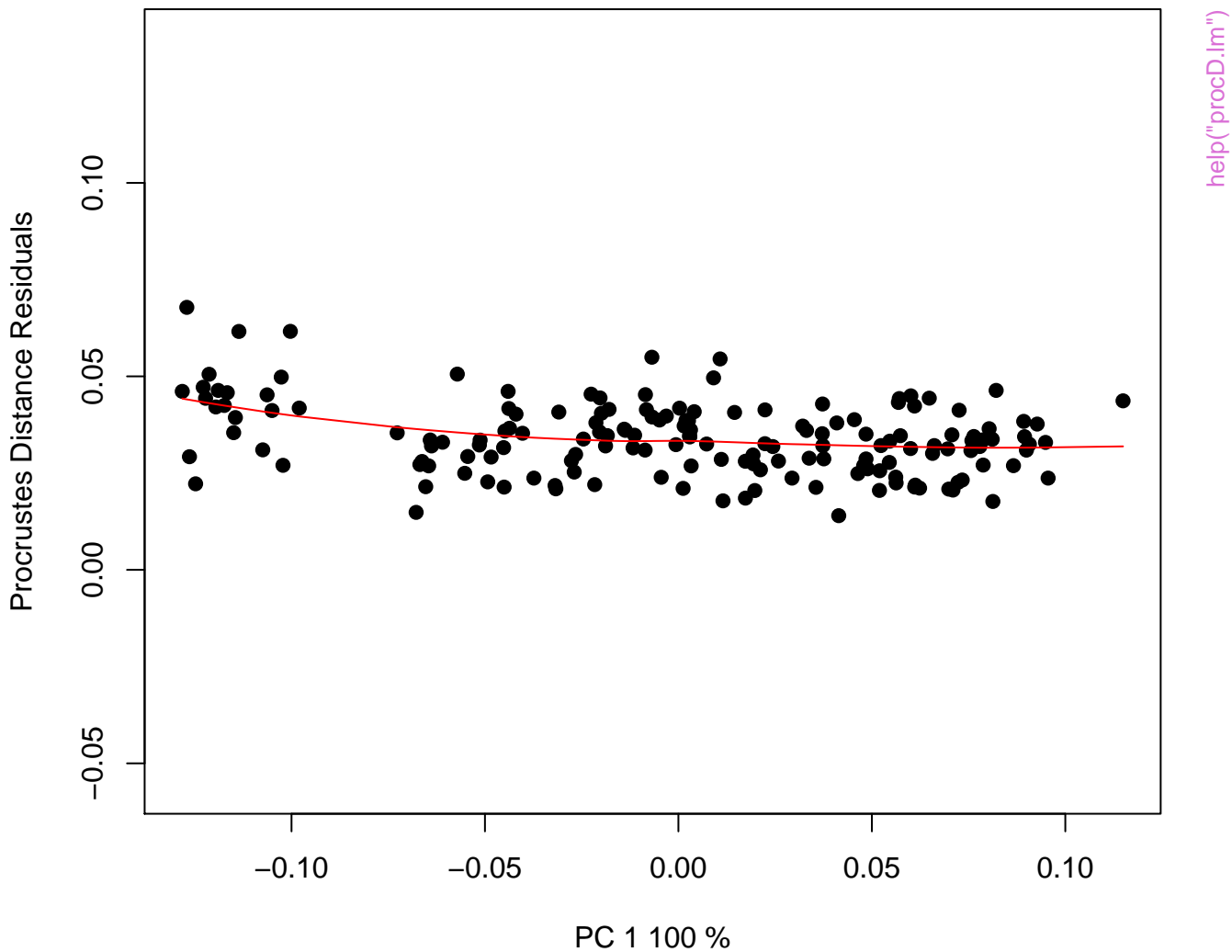
PCA Residuals



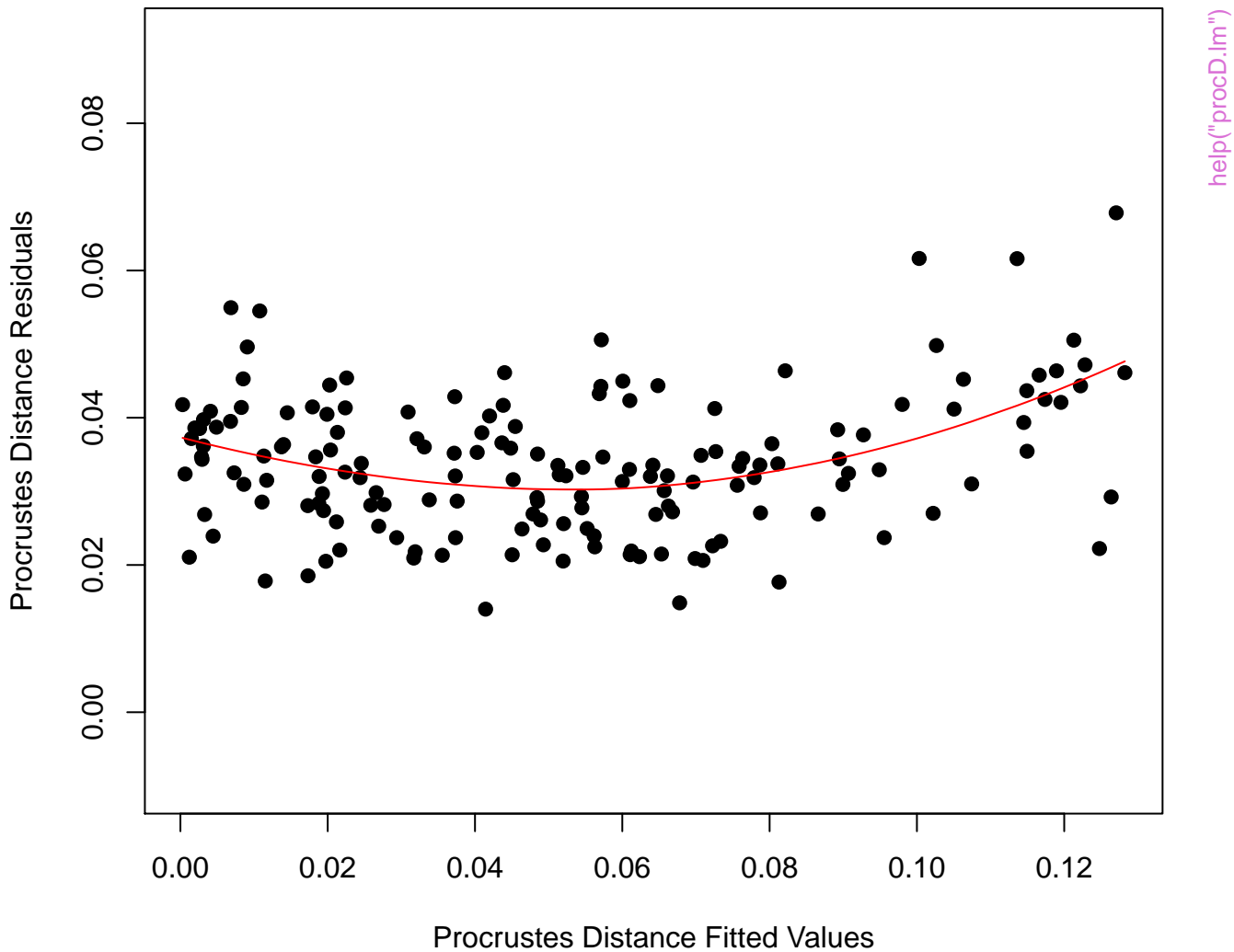
Q-Q plot



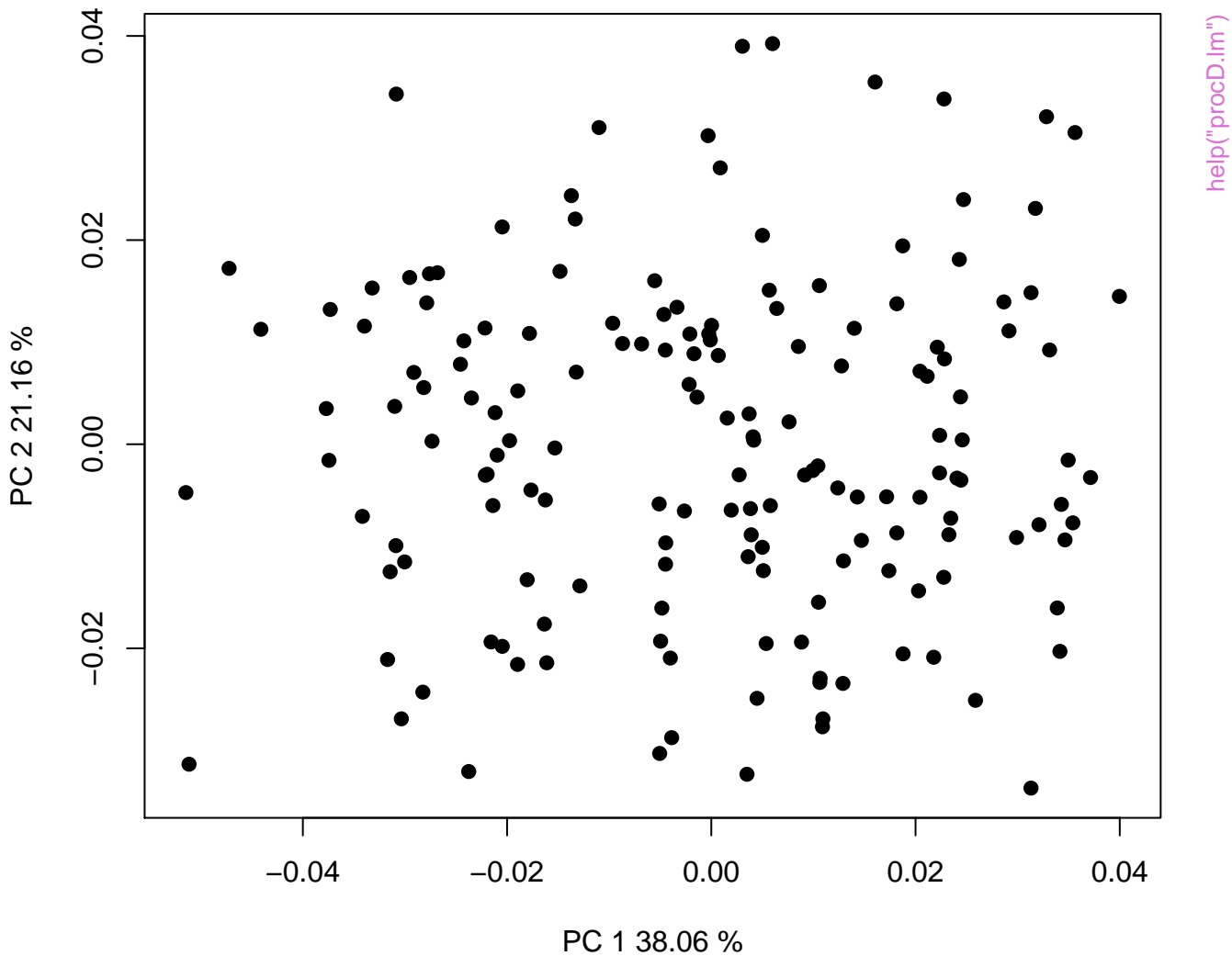
Residuals vs. PC 1 fitted



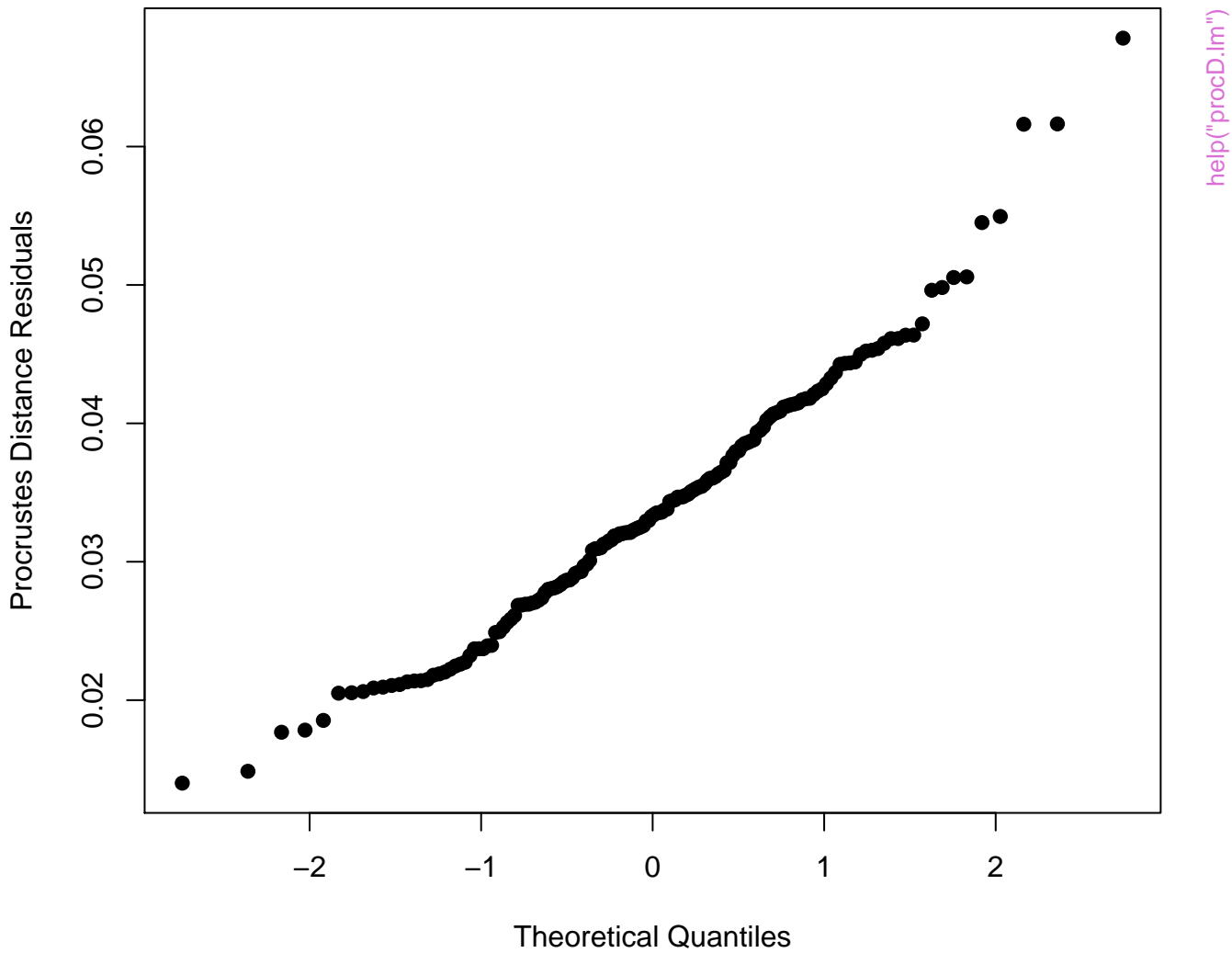
Residuals vs. Fitted



PCA Residuals

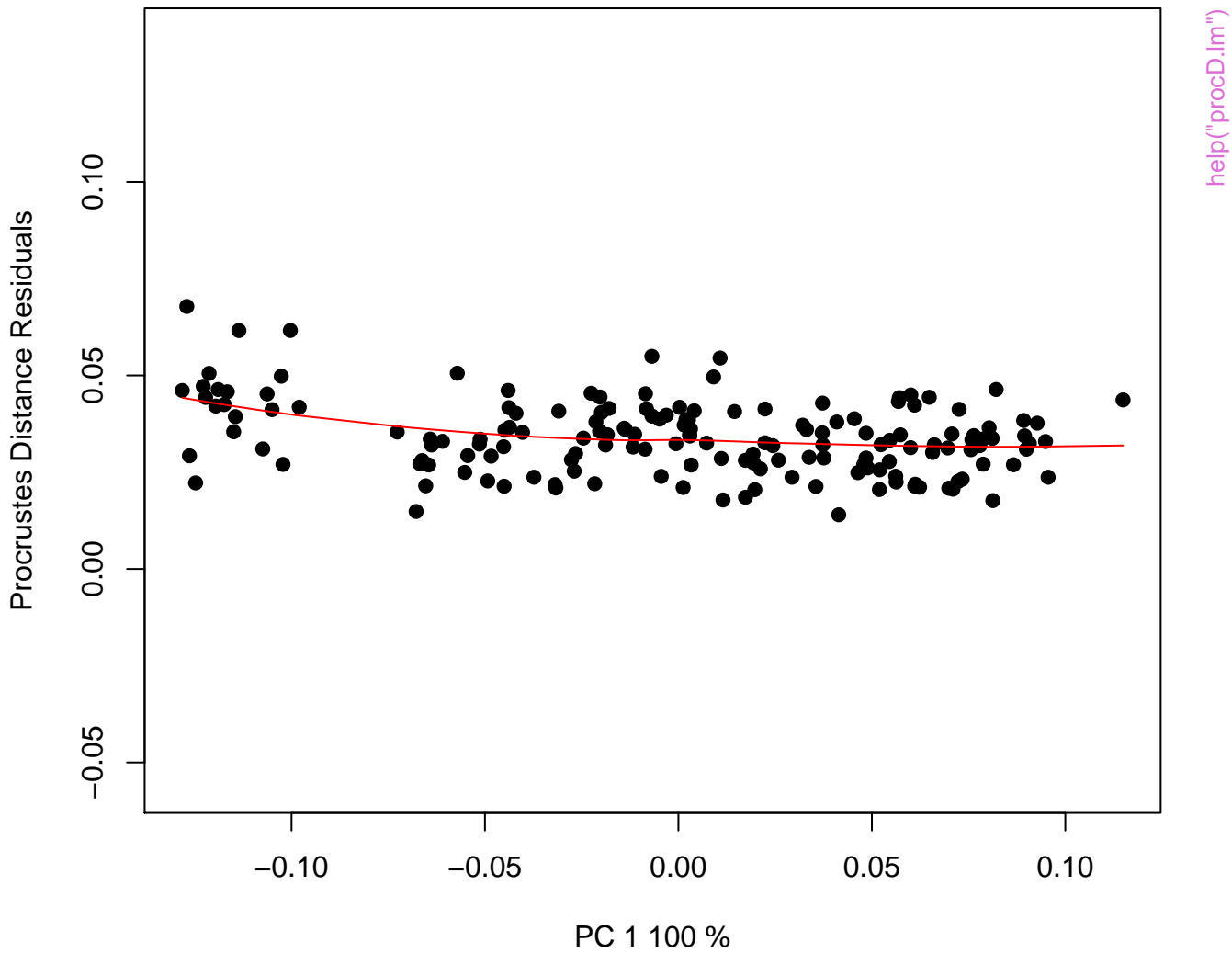


Q-Q plot

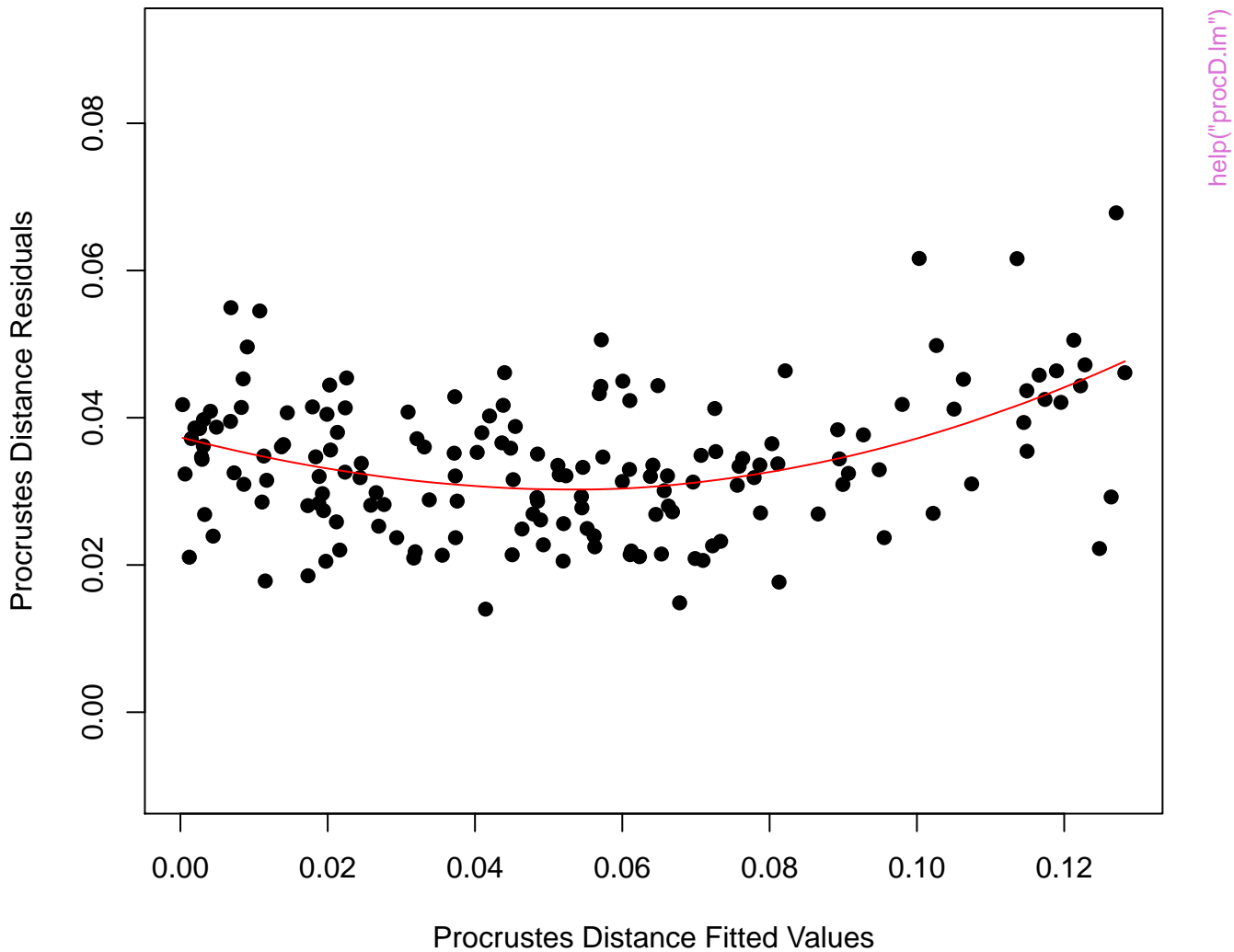


help("procD.lm")

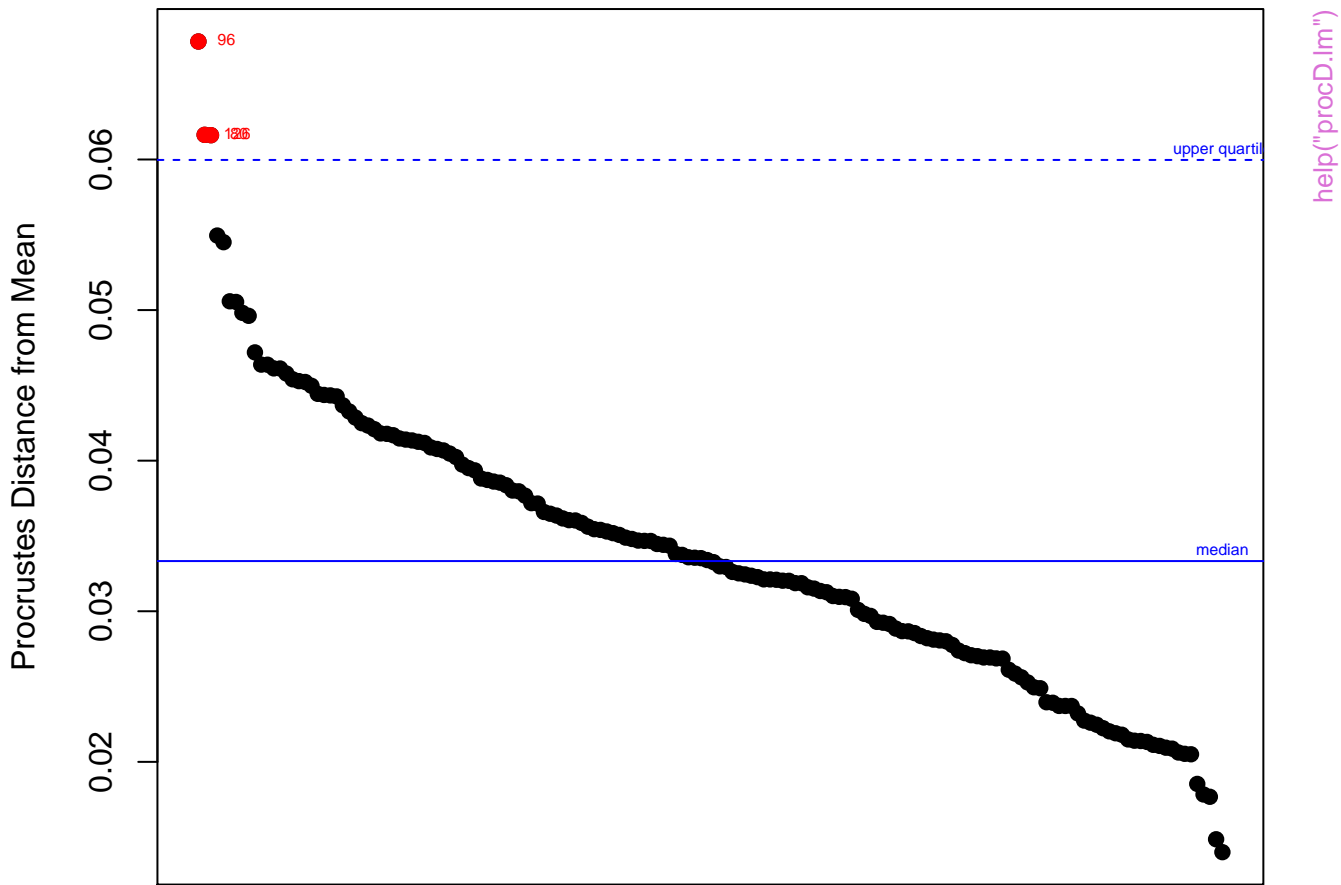
Residuals vs. PC 1 fitted



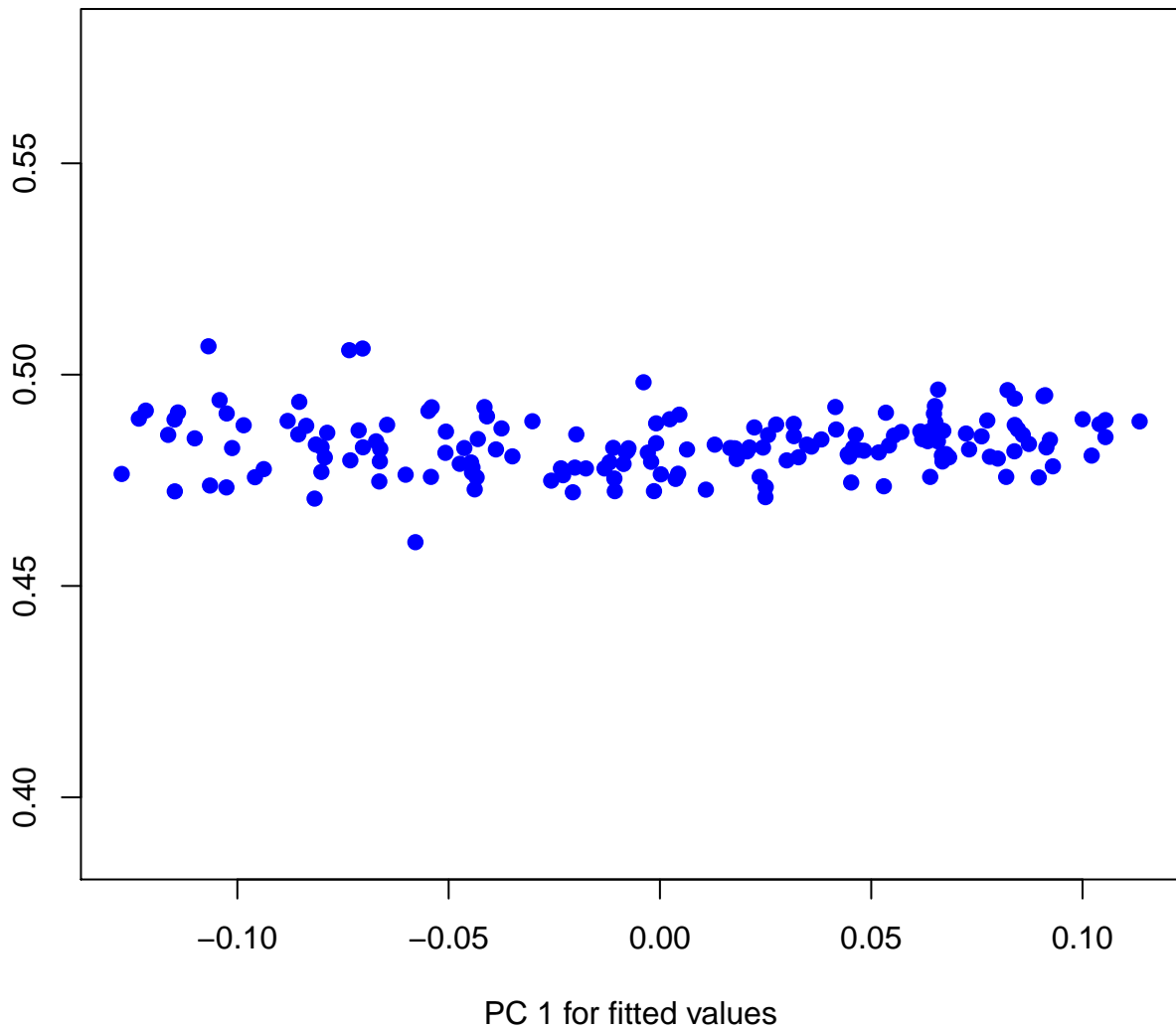
Residuals vs. Fitted



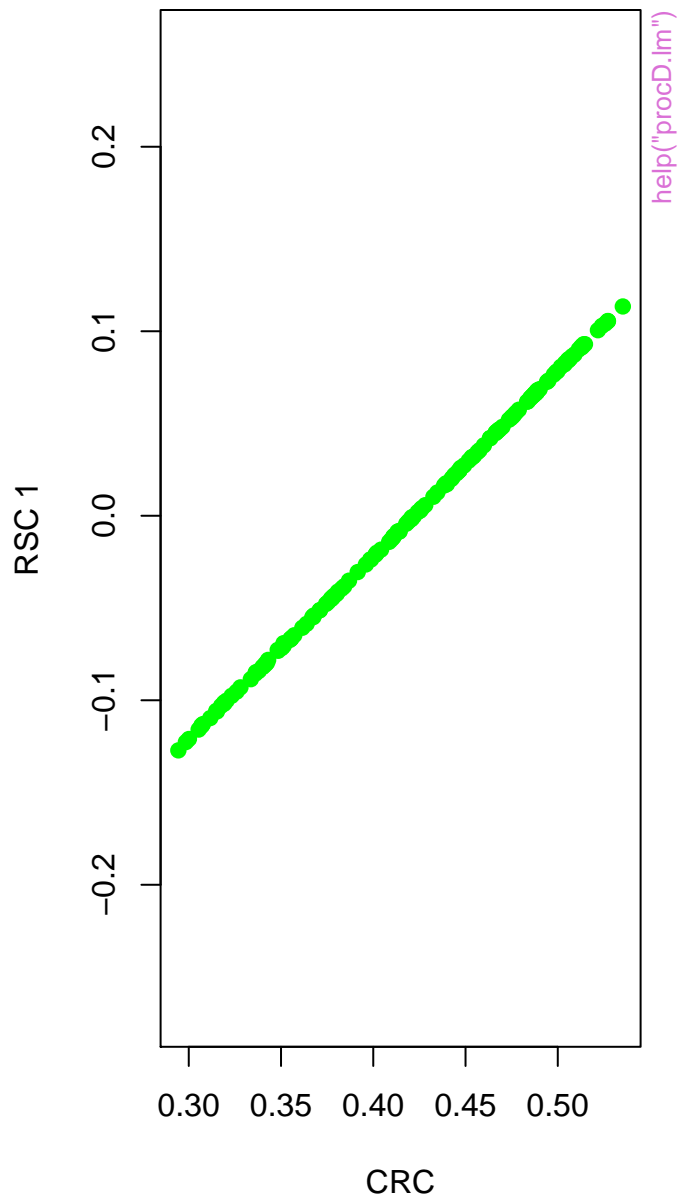
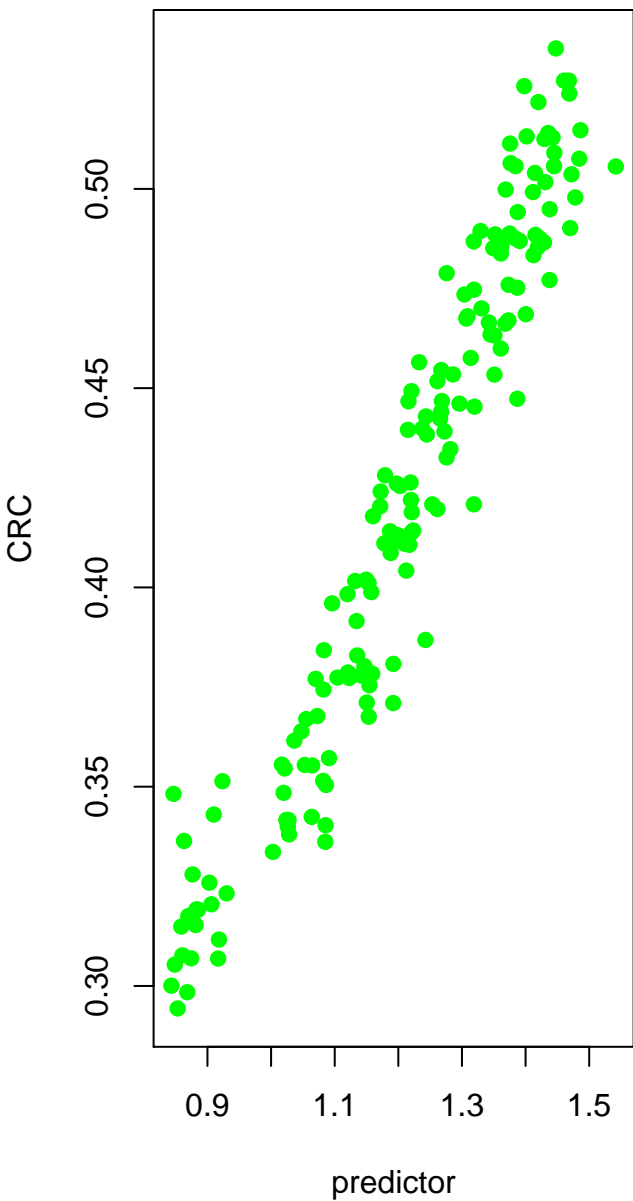
All Specimens

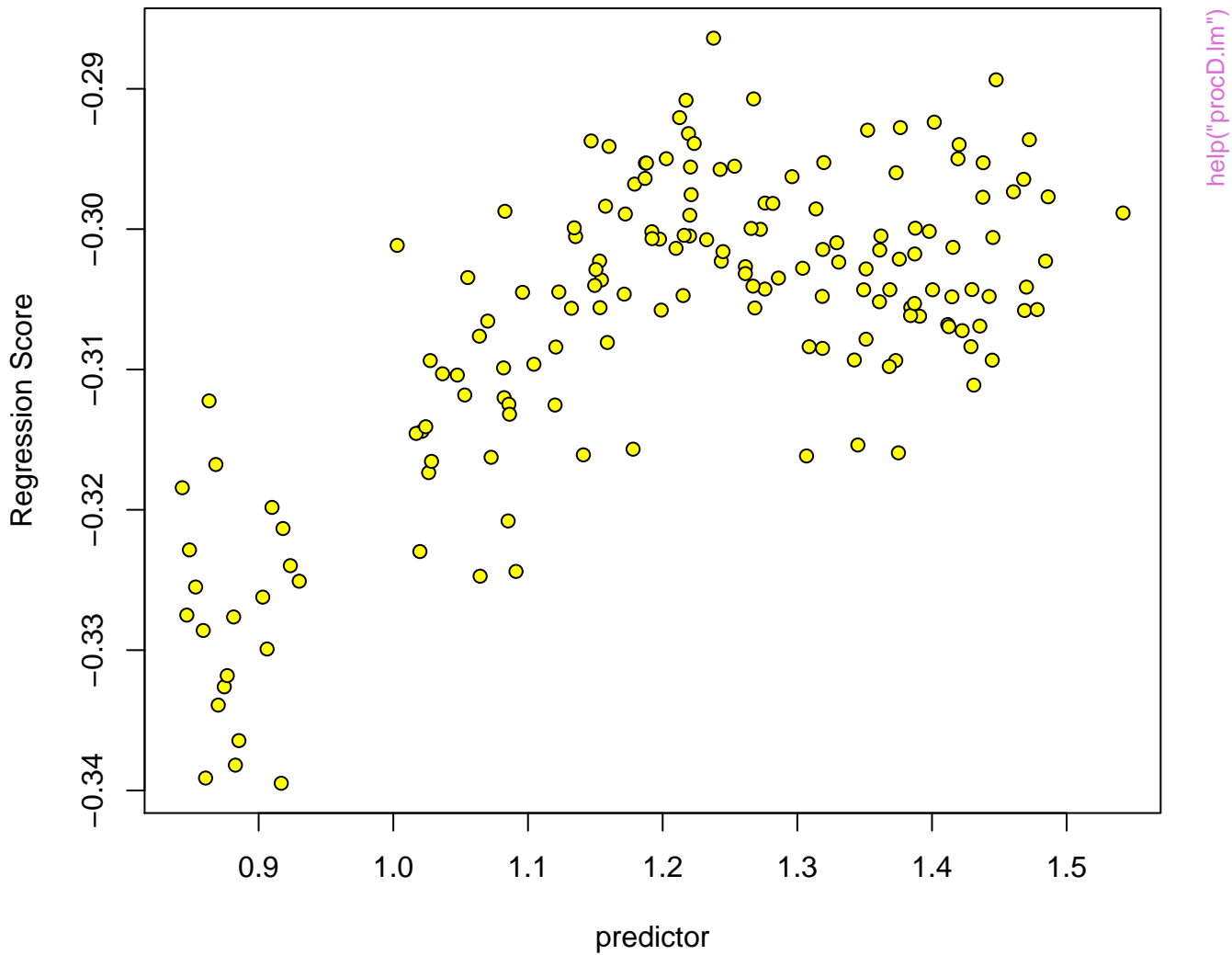


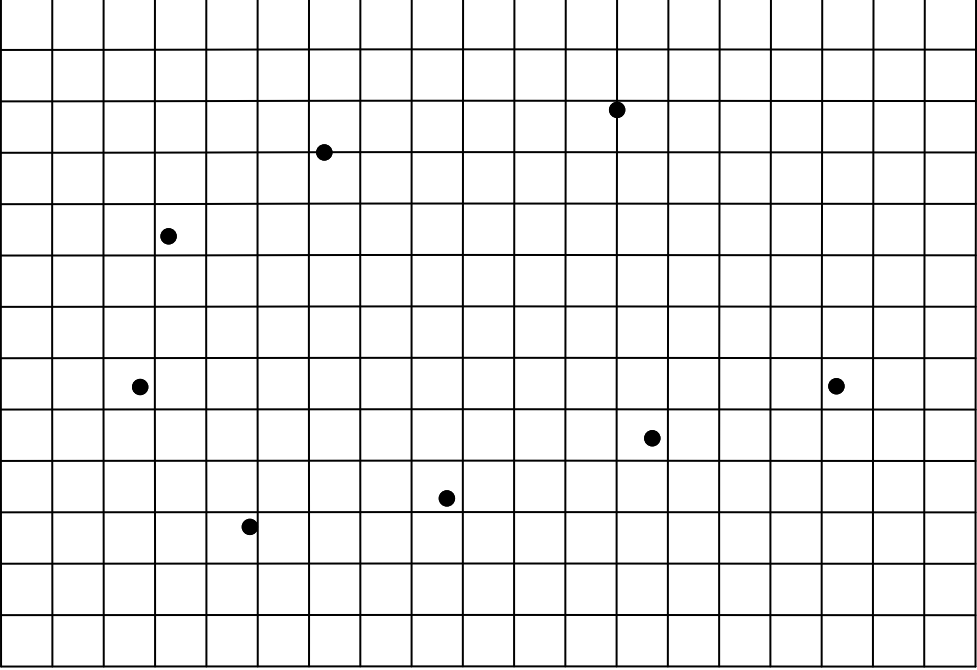
PC 2 for fitted values

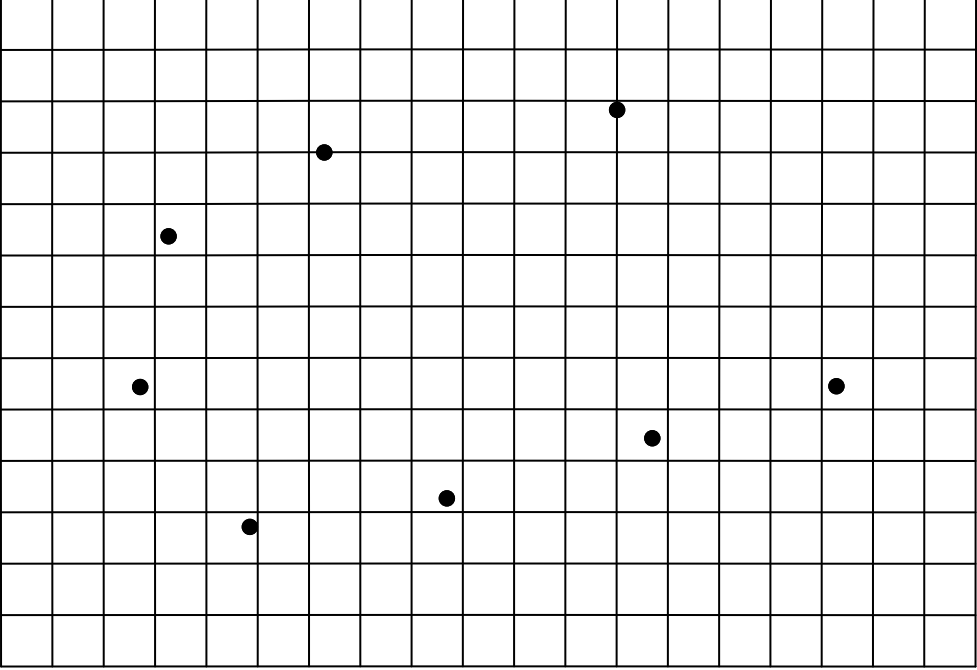


help("procD.lm")

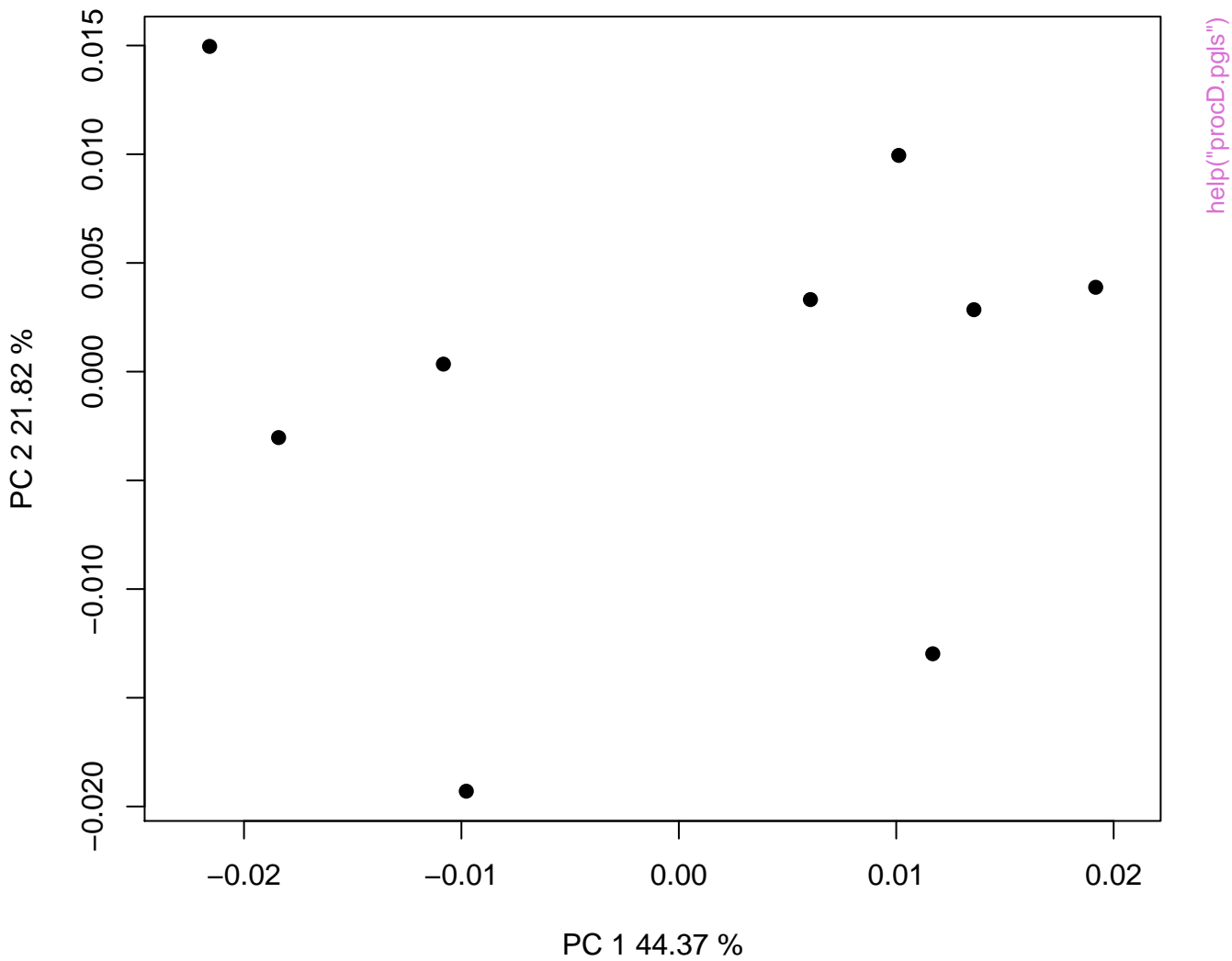




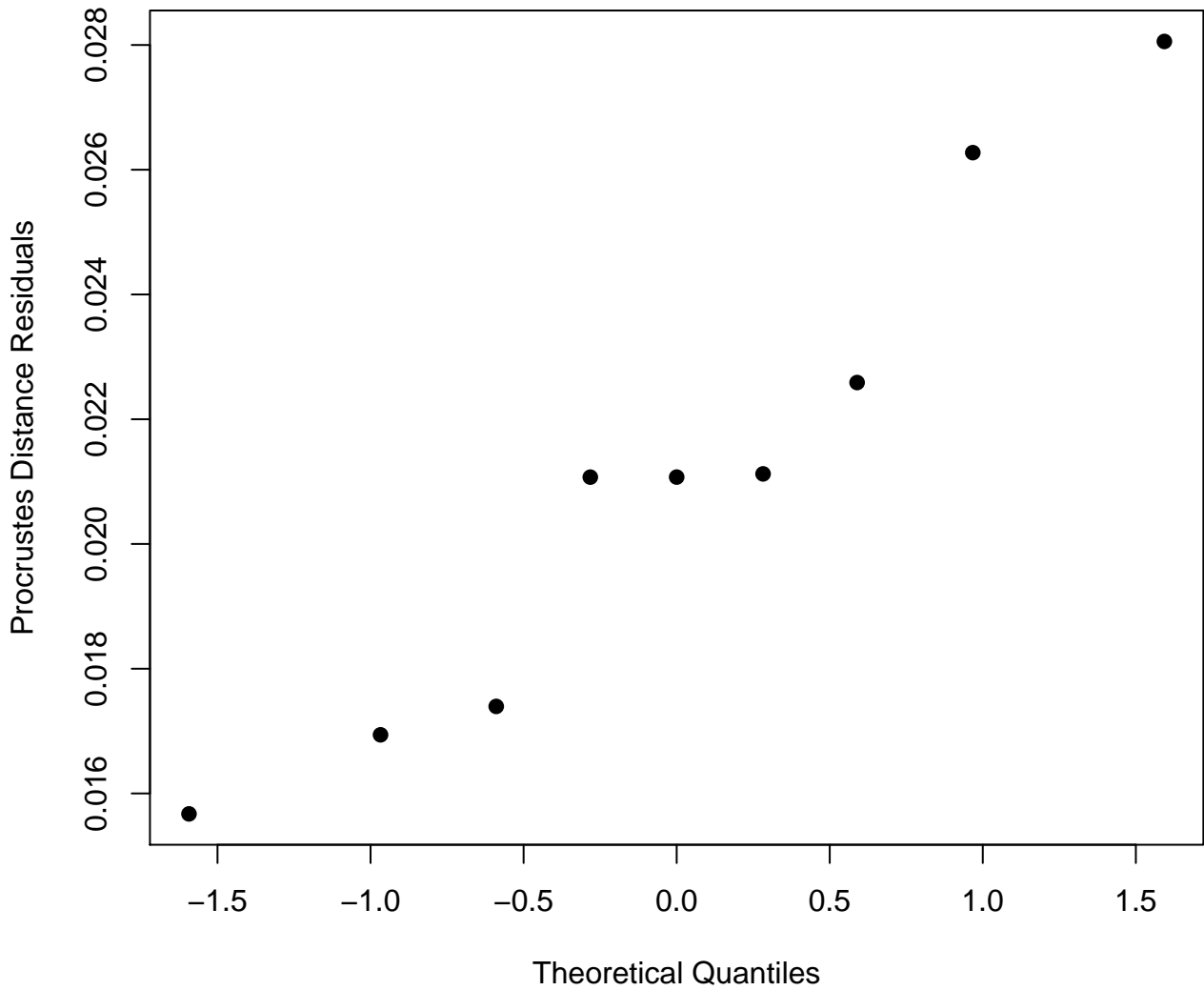




PCA Residuals

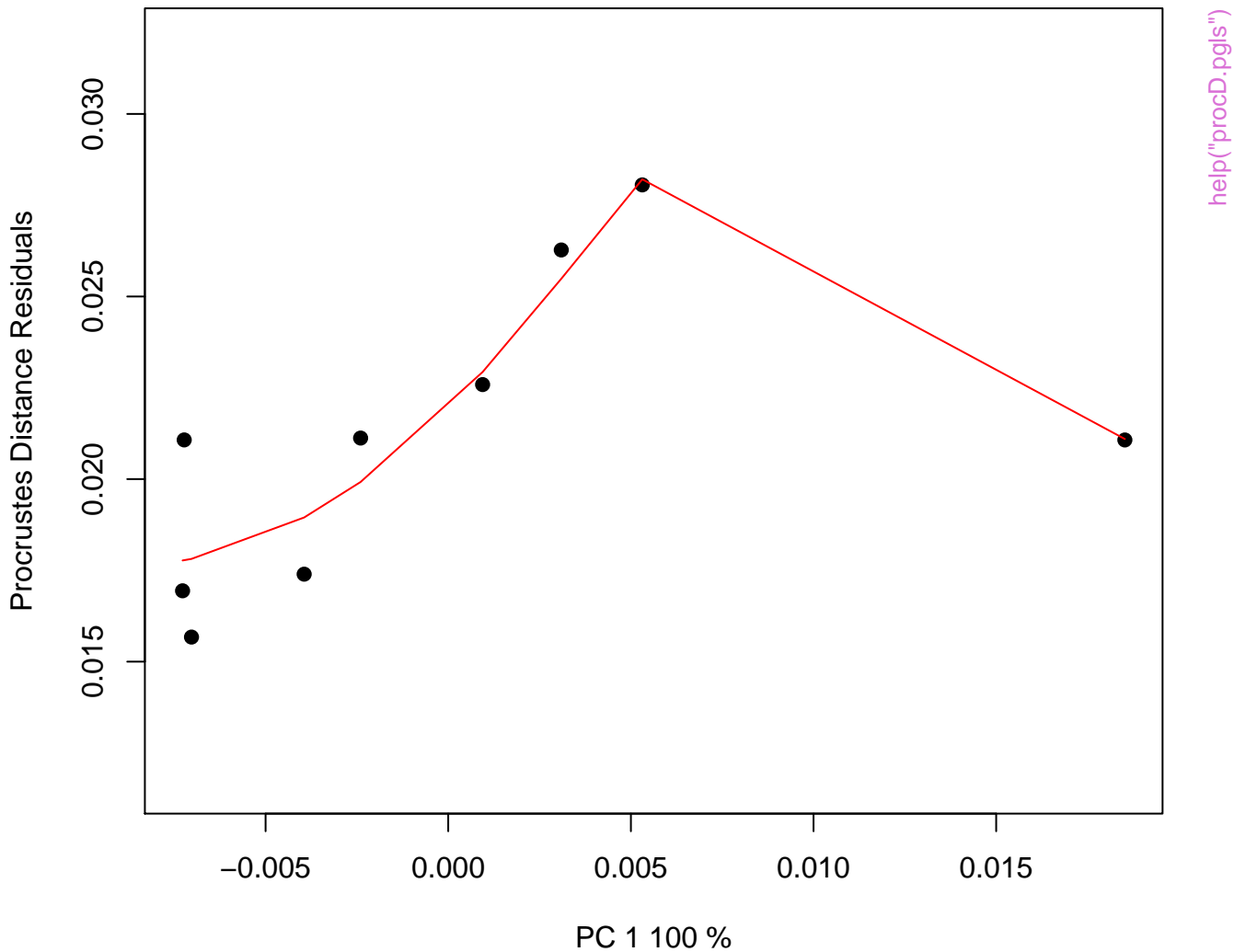


Q-Q plot



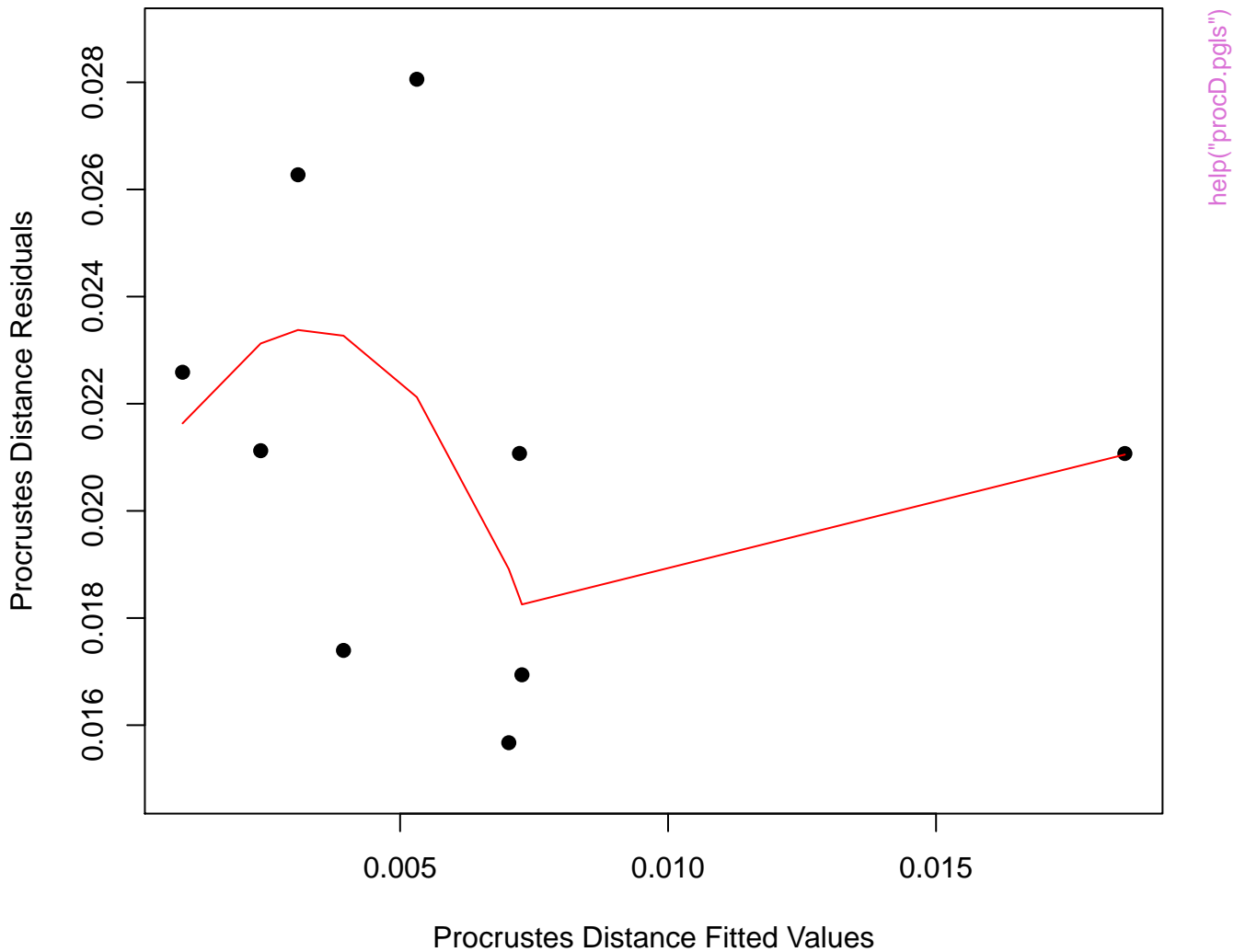
[help\("procD.pgls"\)](#)

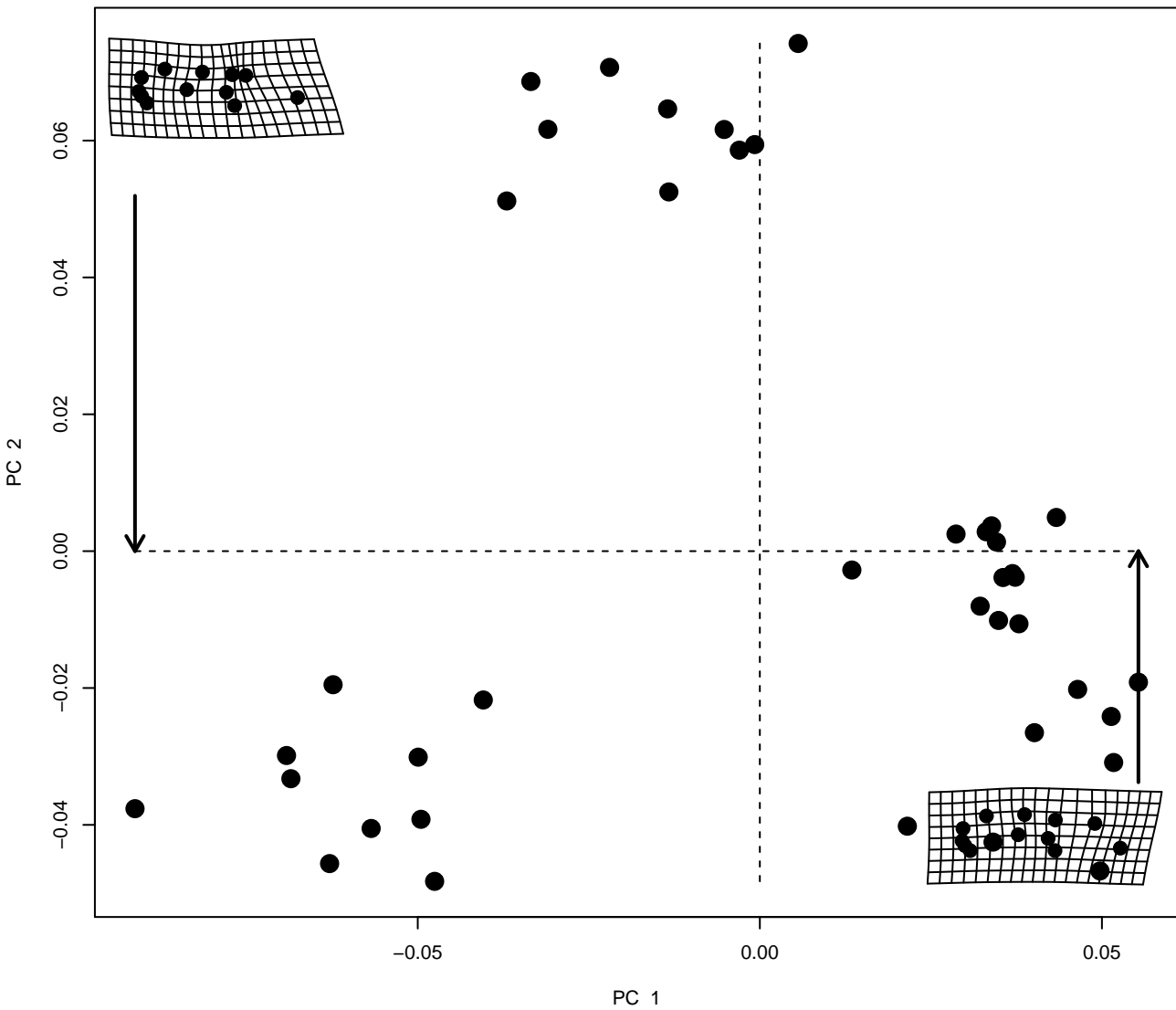
Residuals vs. PC 1 fitted

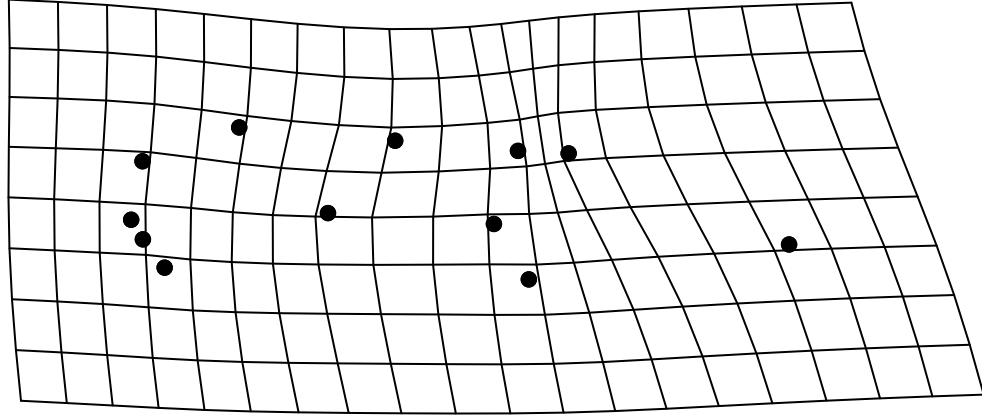


help("procD.pgls")

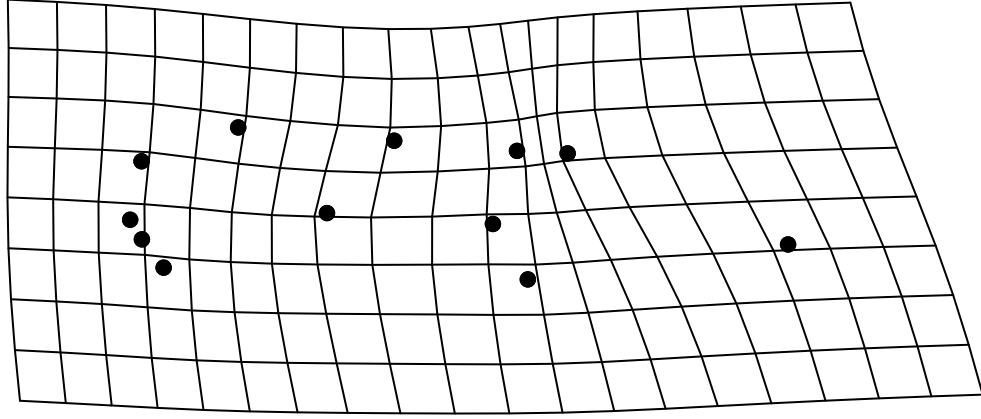
Residuals vs. Fitted



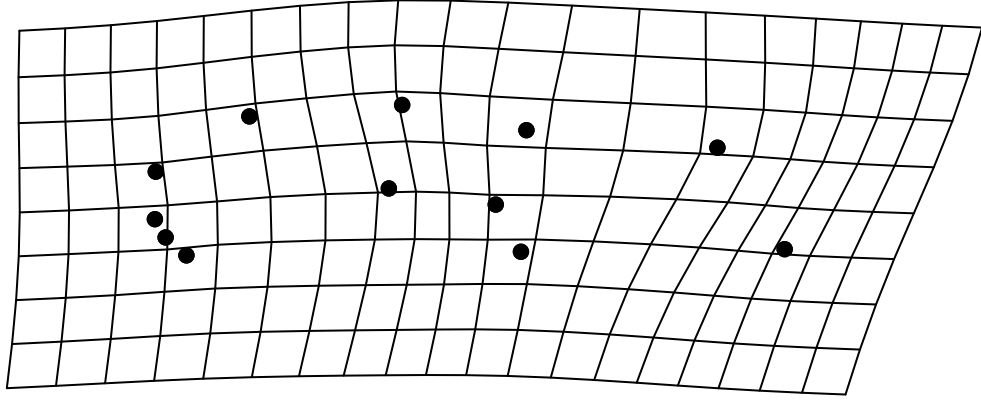


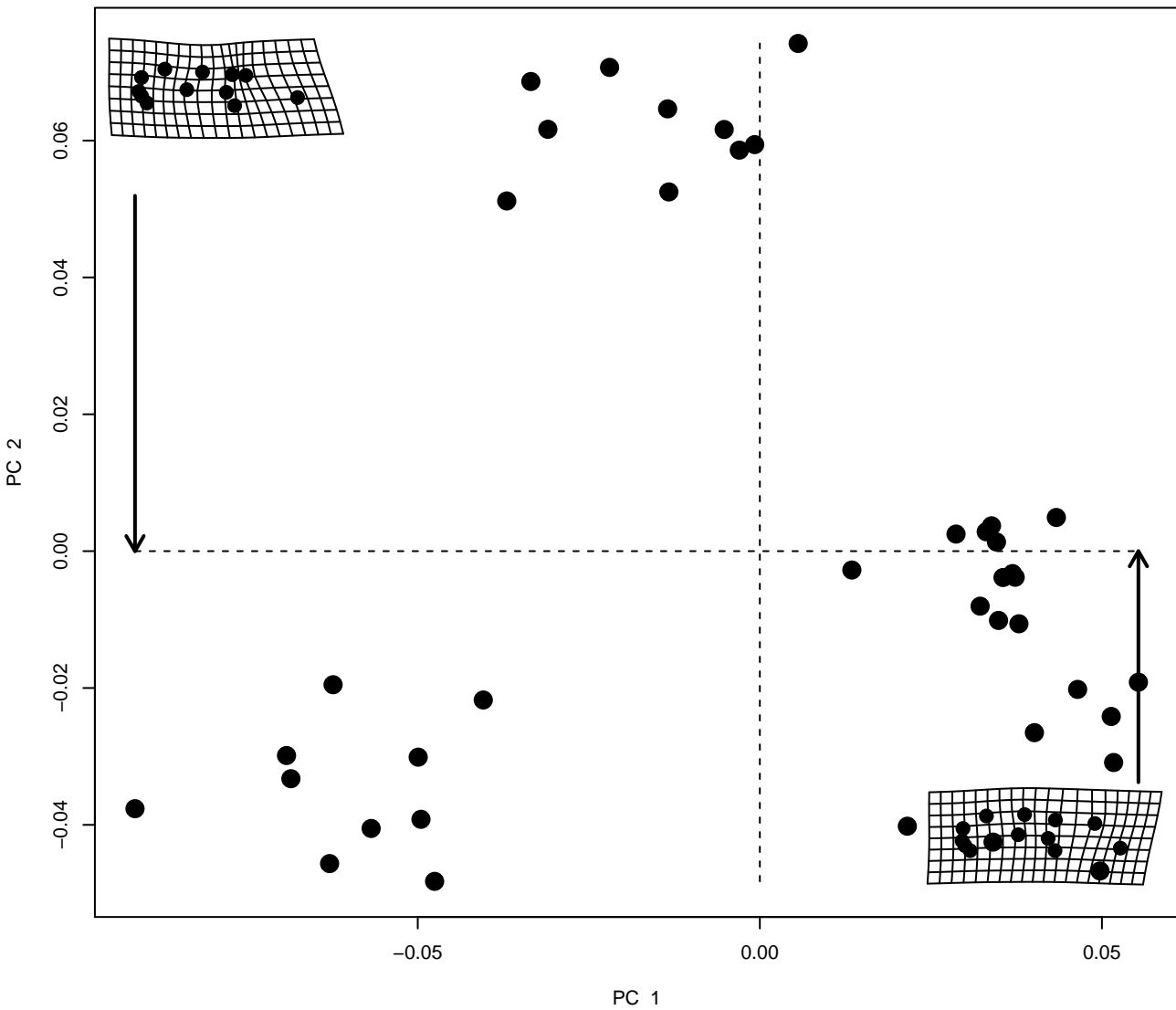


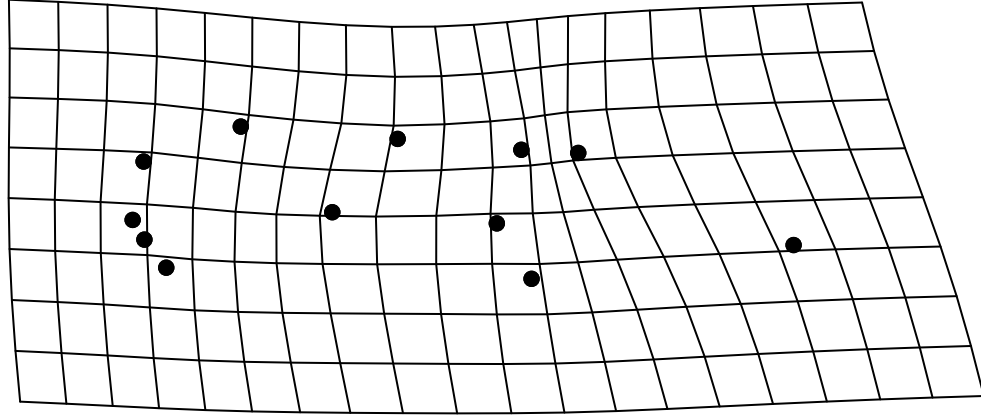
`help("shape.predictor")`



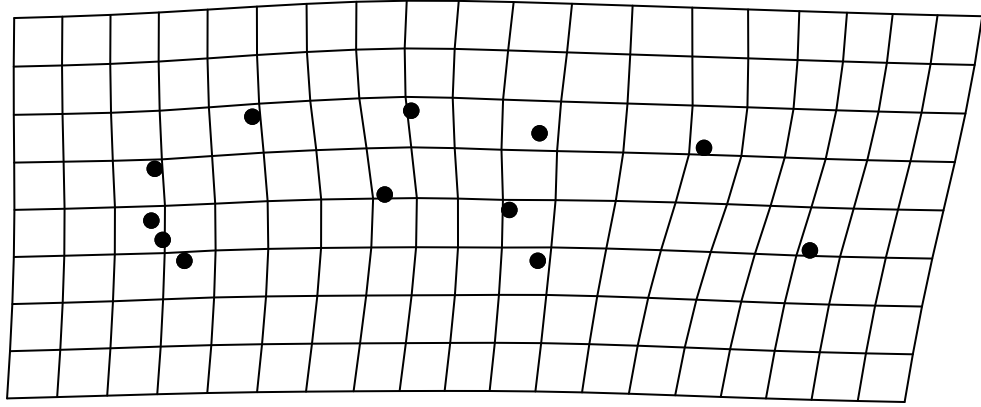
`help("shape.predictor")`



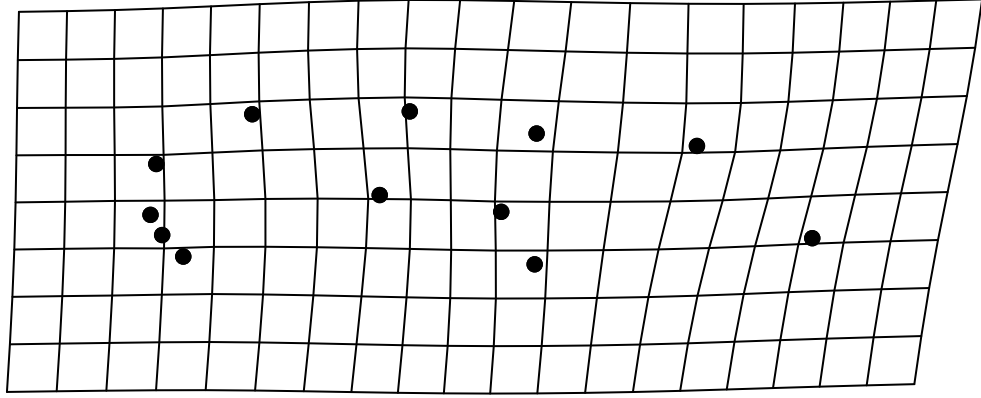


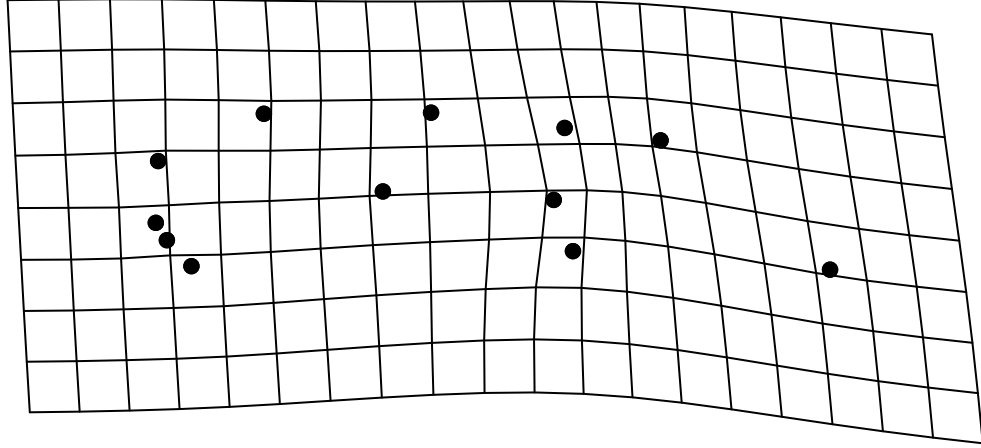


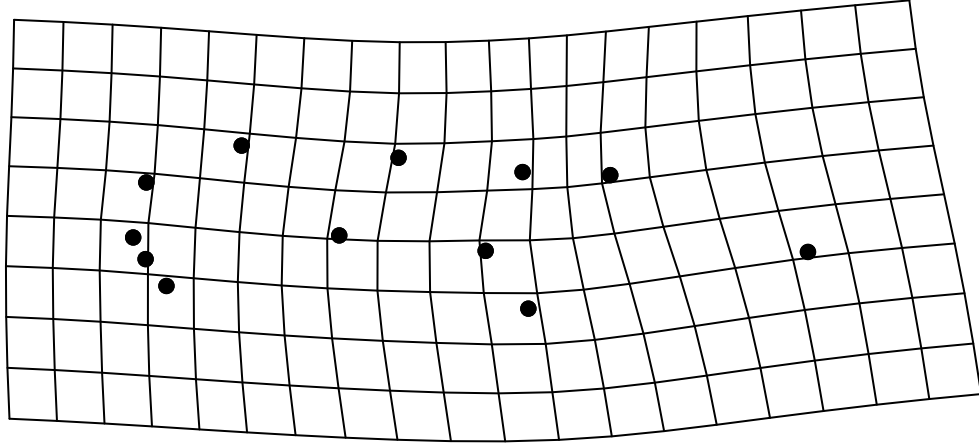
help("shape.predictor")



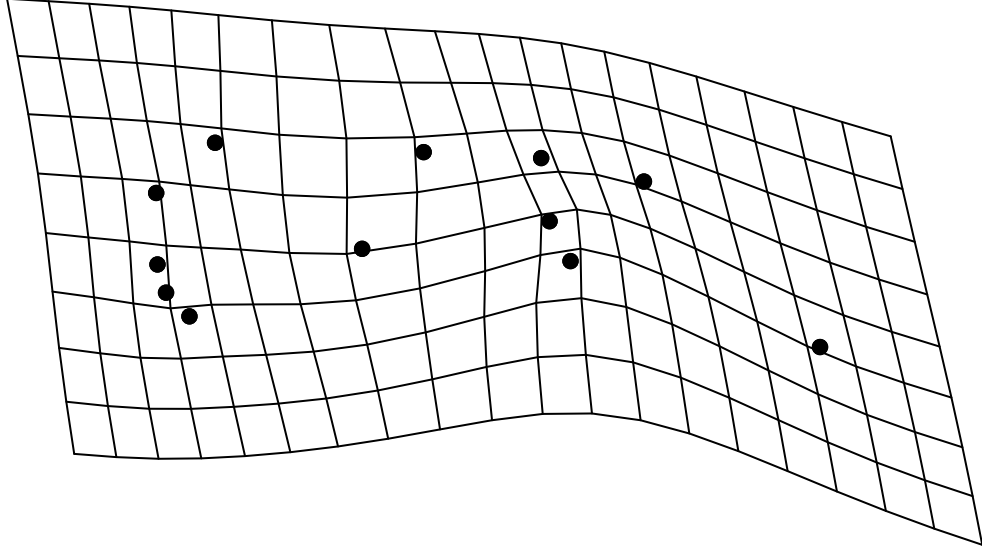
`help("shape.predictor")`

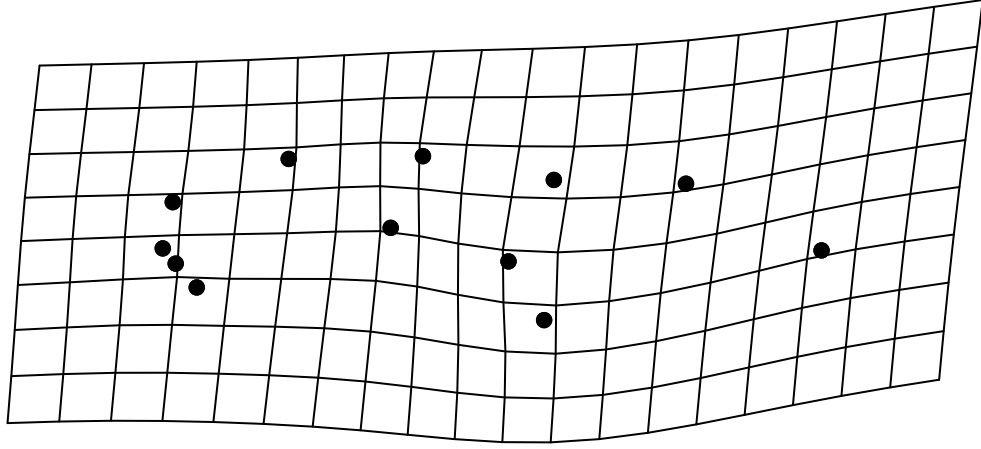




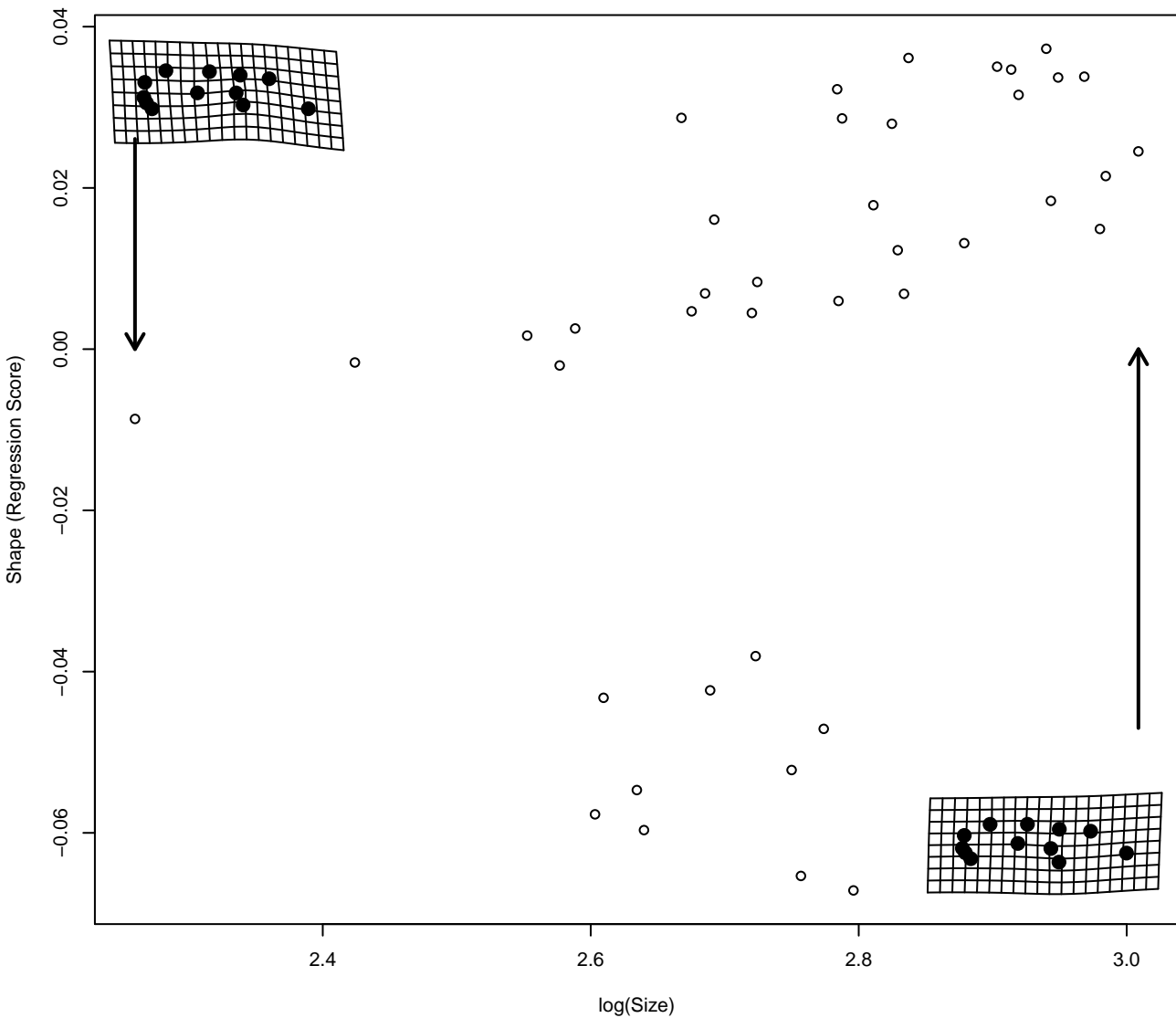


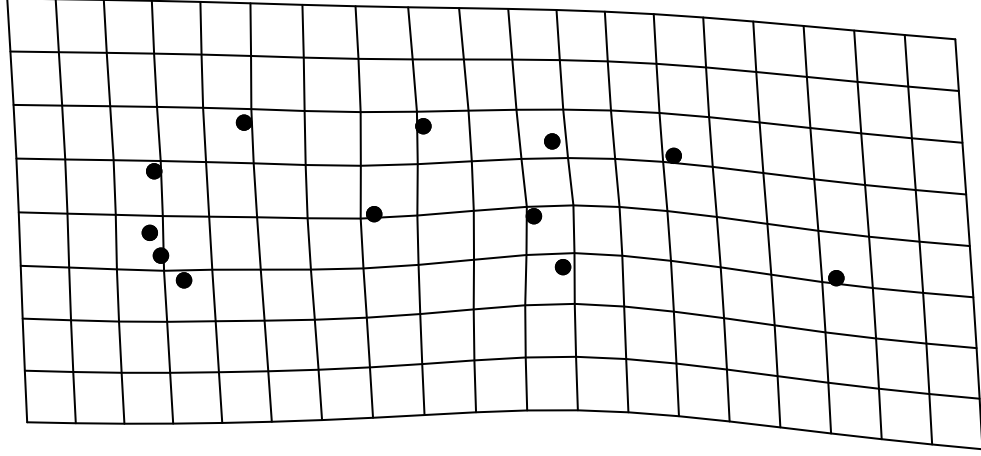
`help("shape.predictor")`



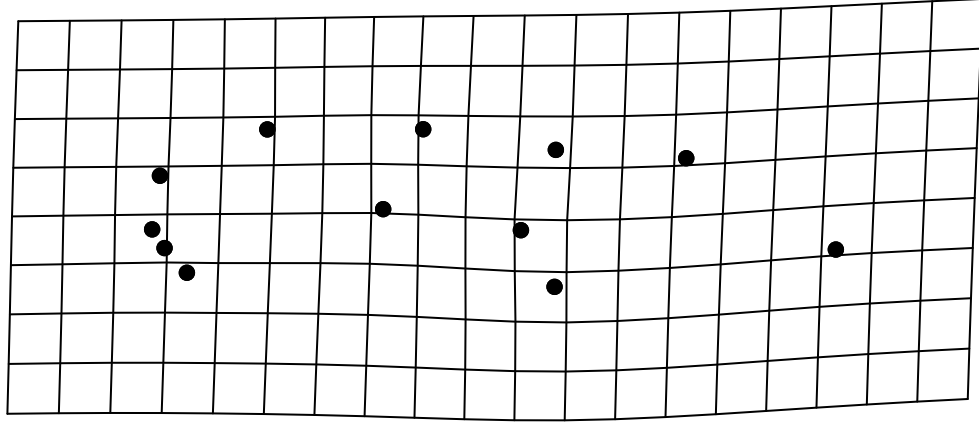


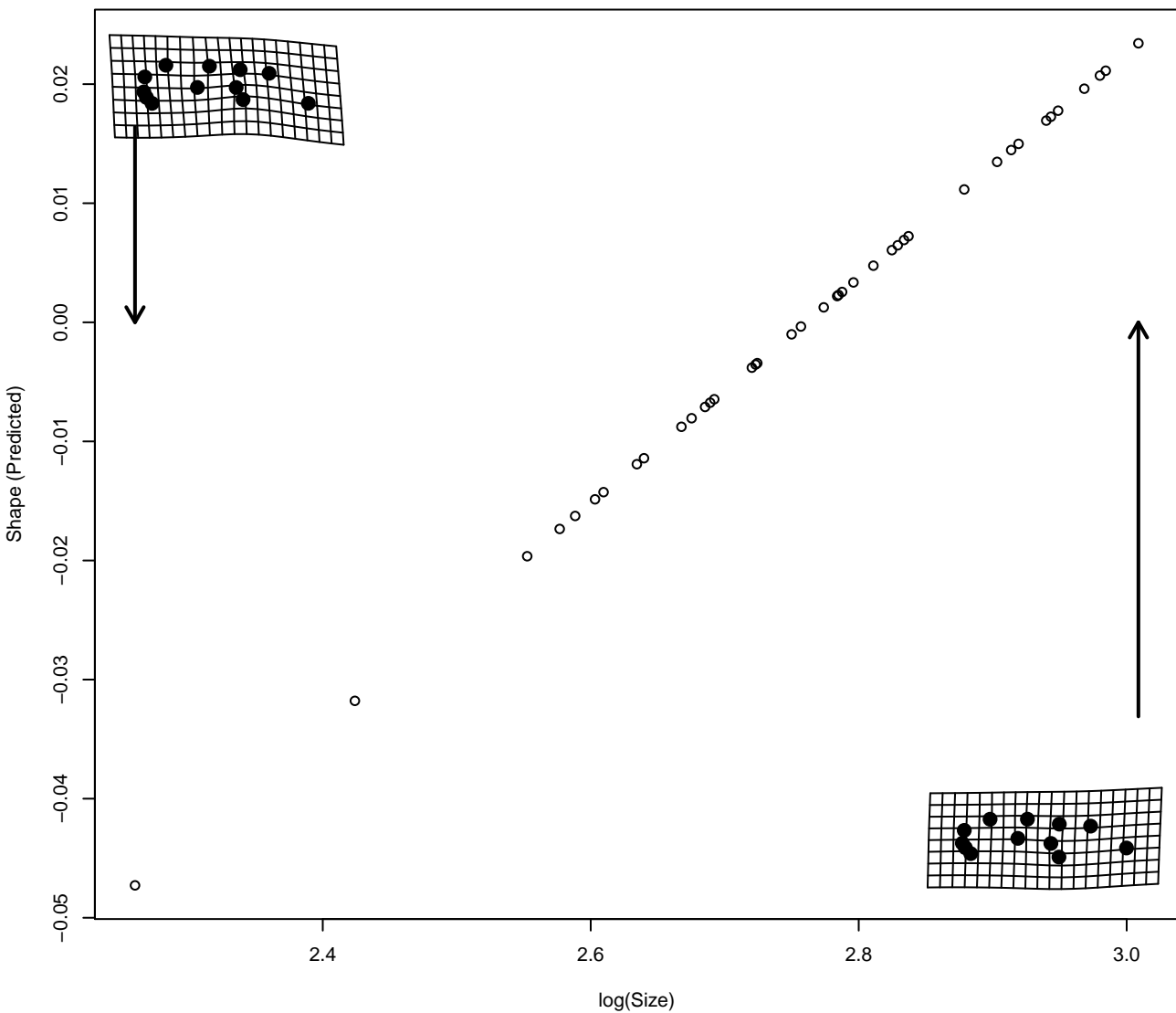
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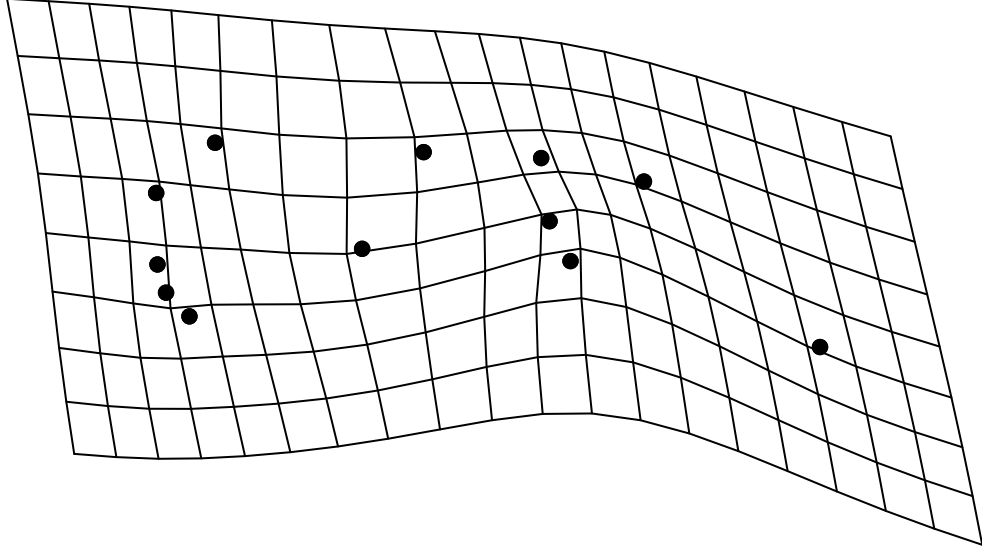


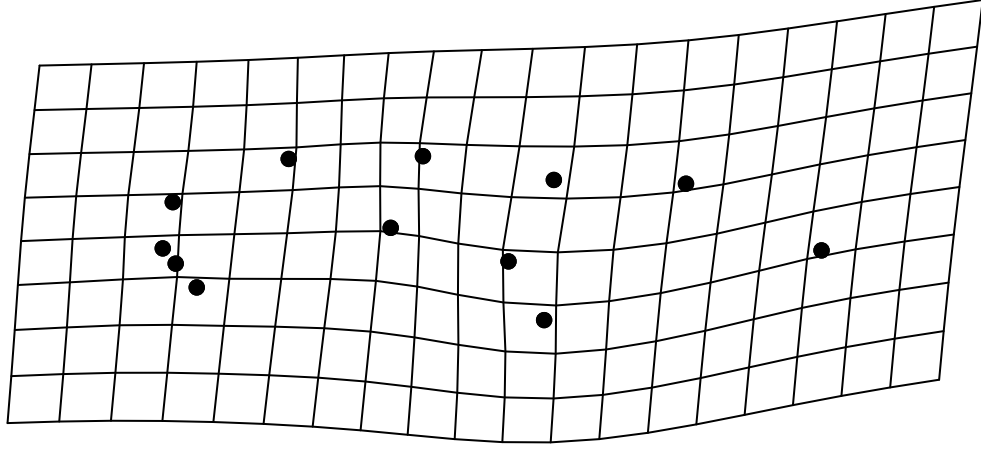


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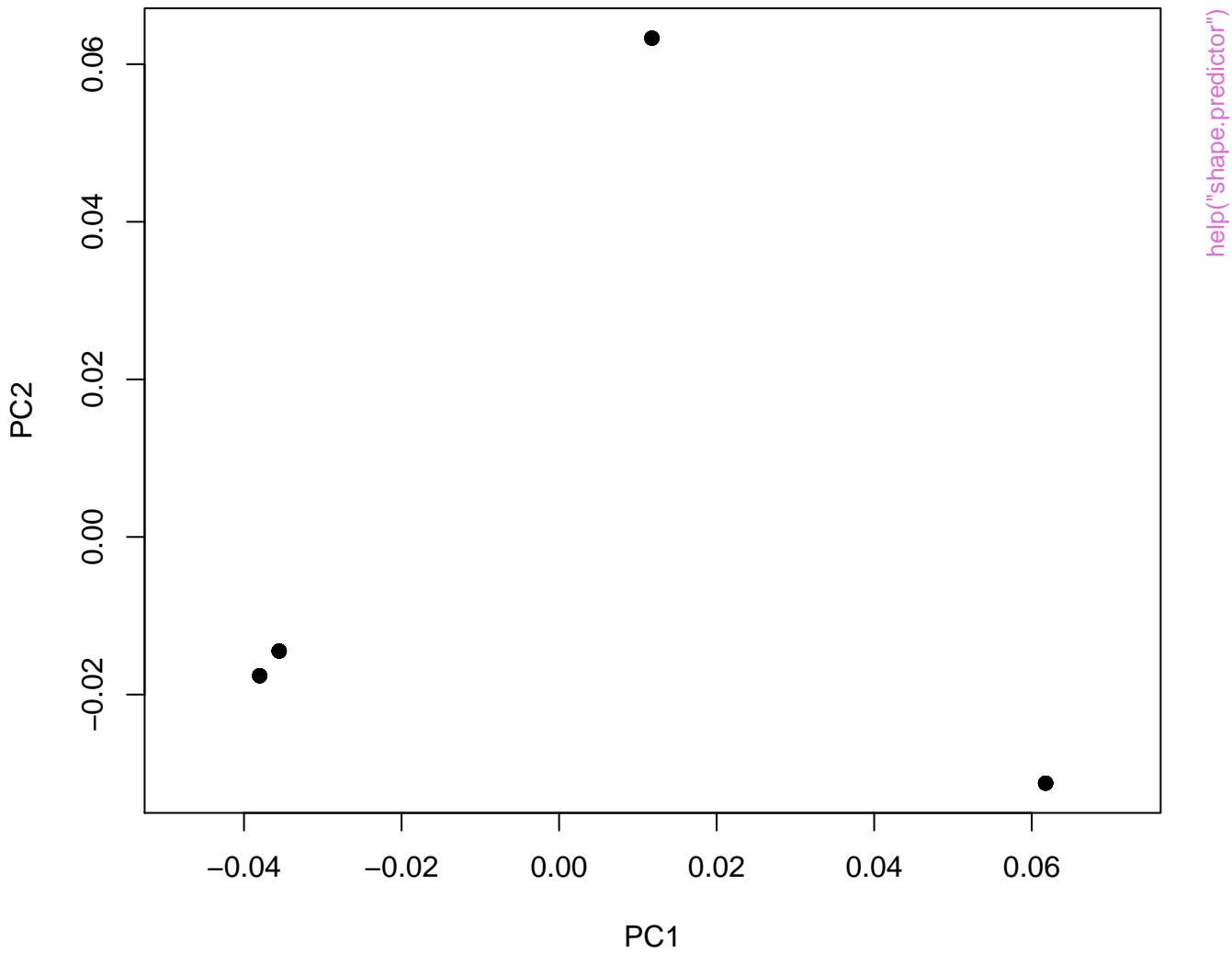


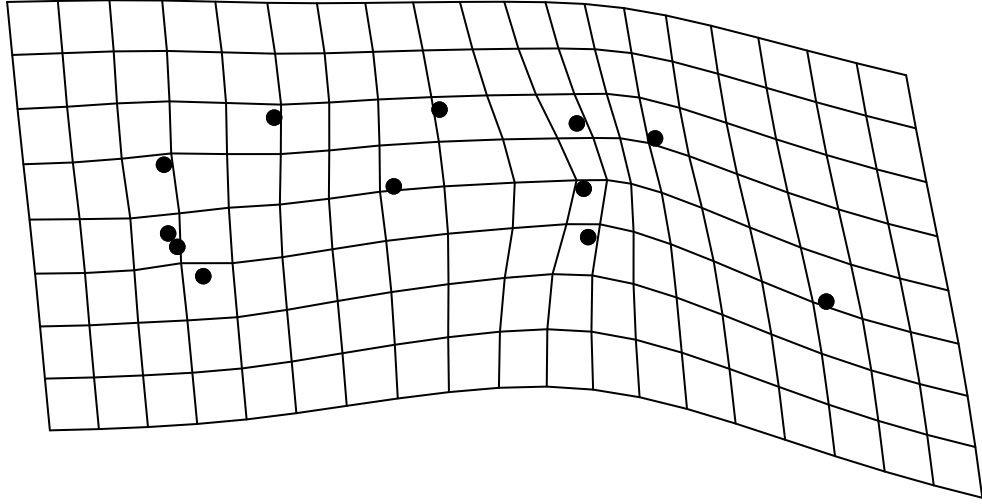




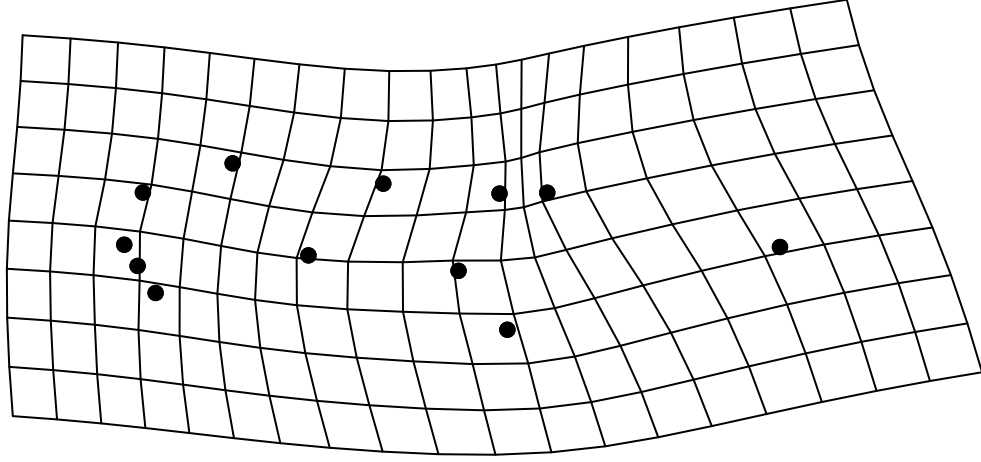


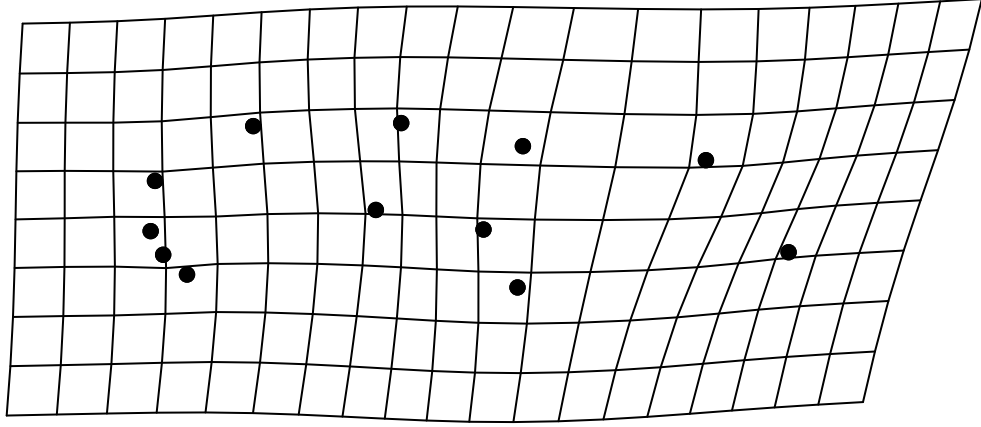
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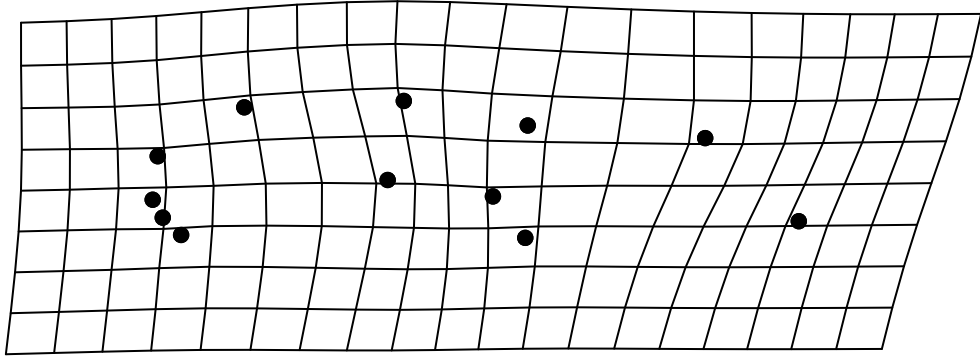


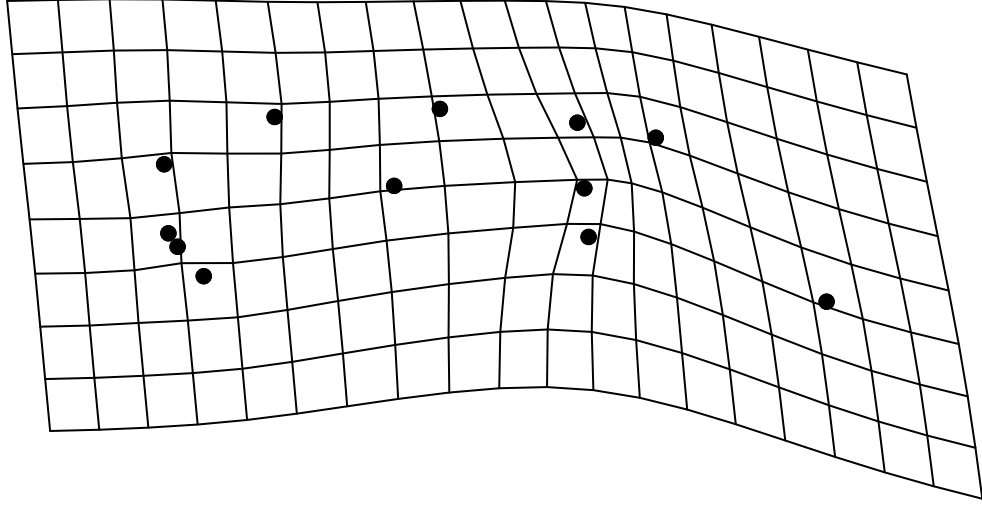


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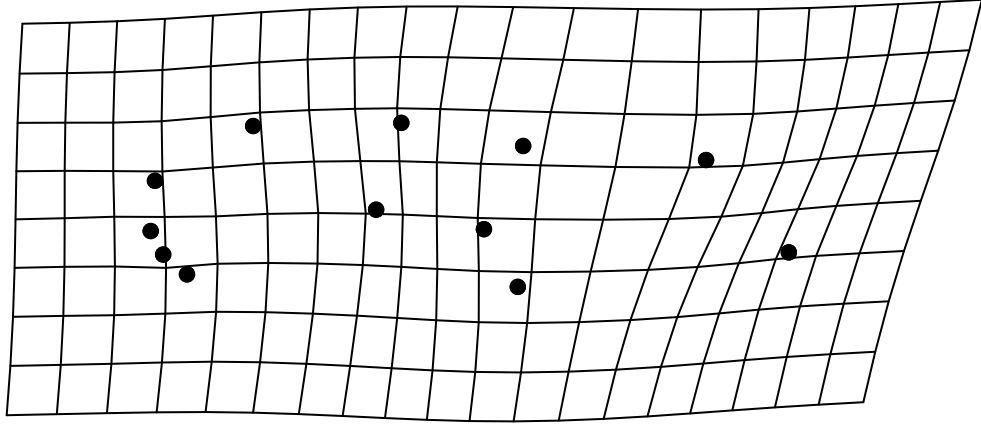




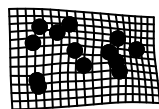




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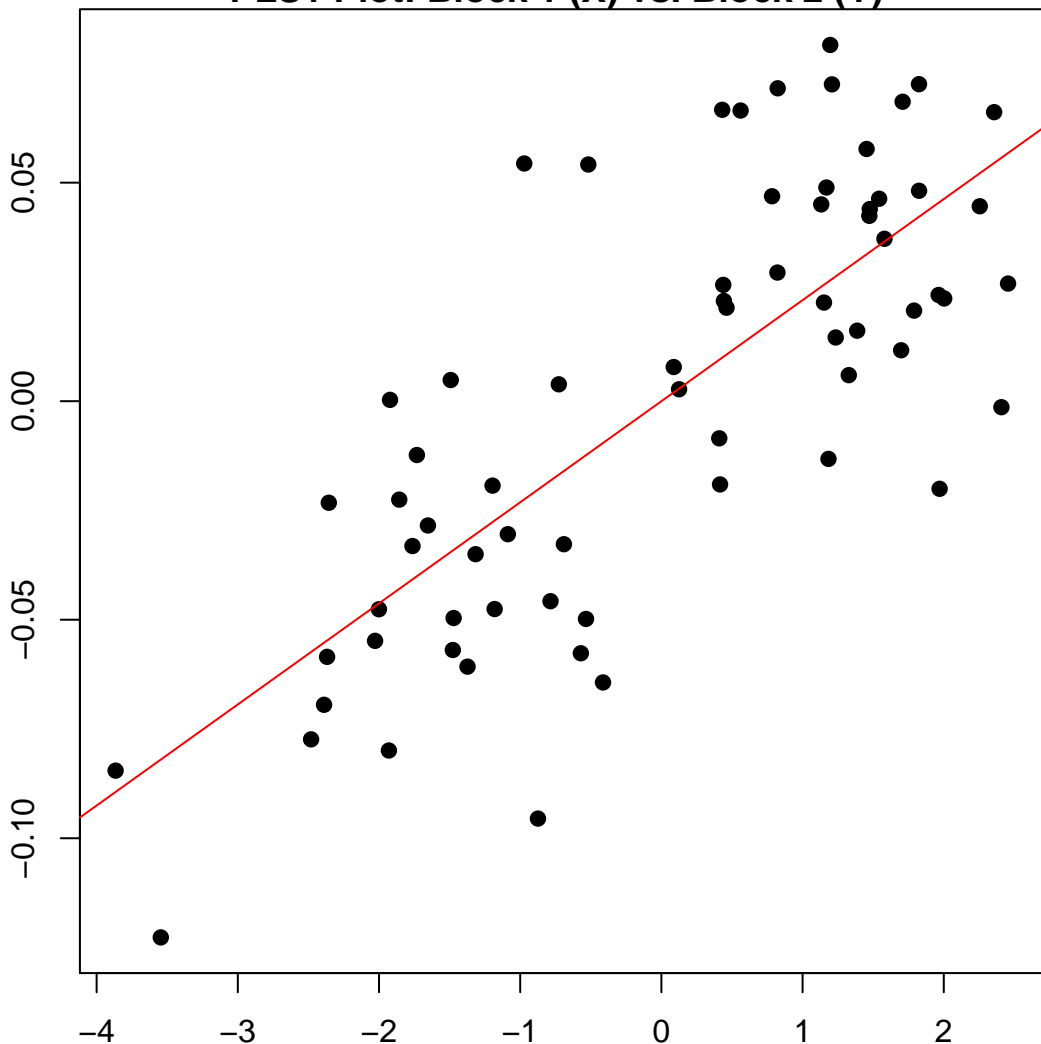
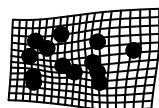


PLS1 Plot: Block 1 (X) vs. Block 2 (Y)

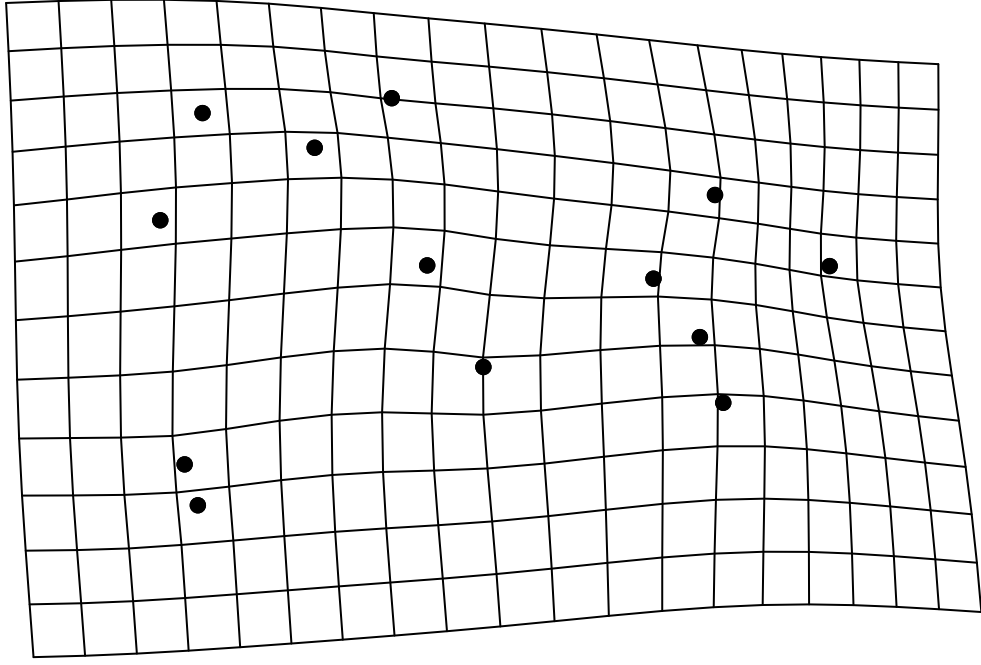


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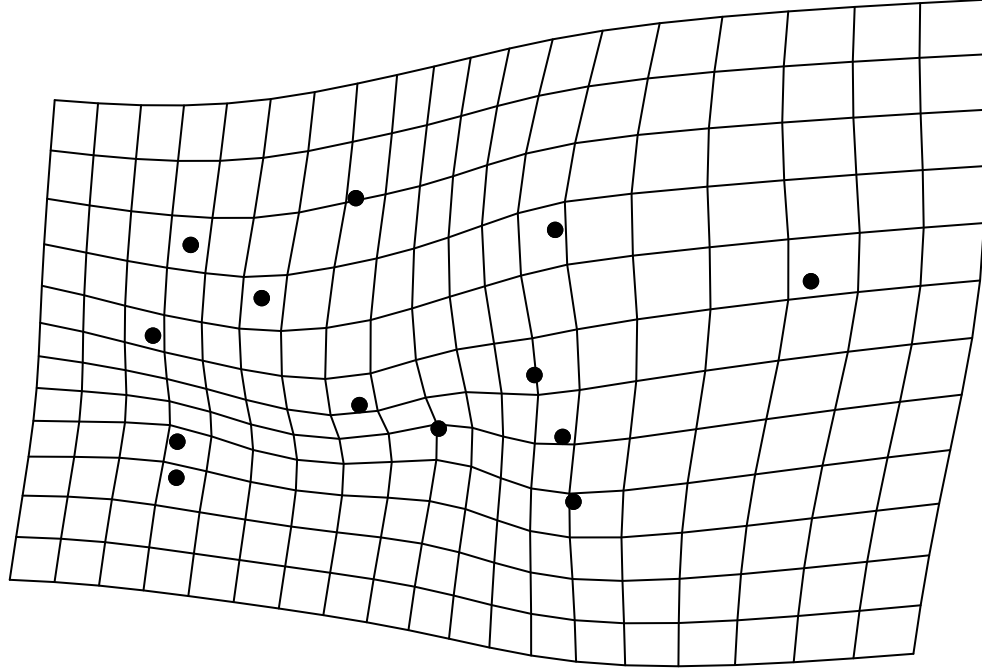
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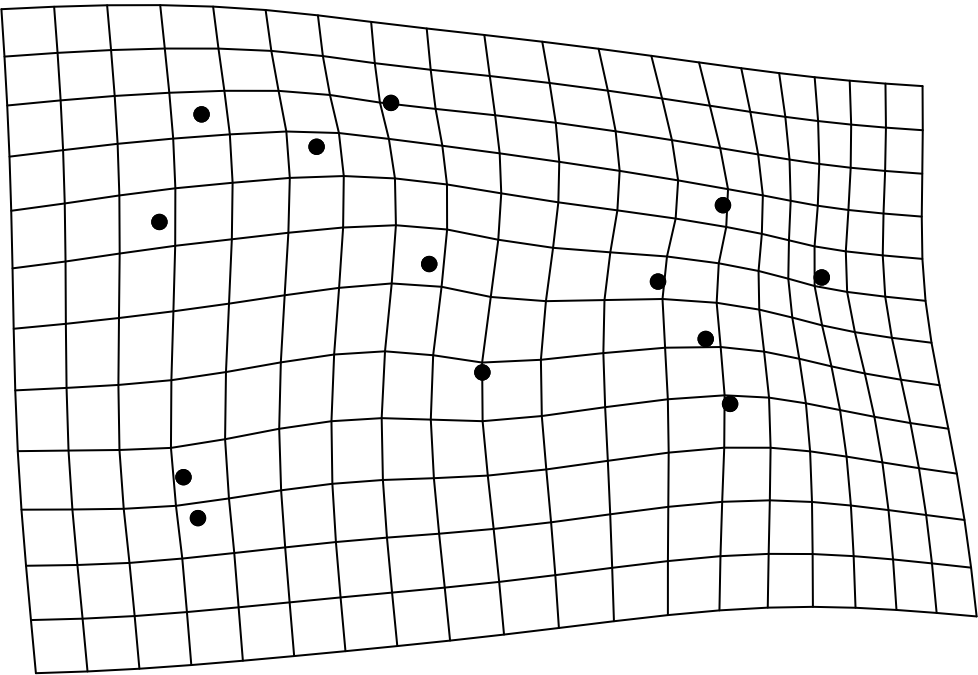
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help("shape.predictor")

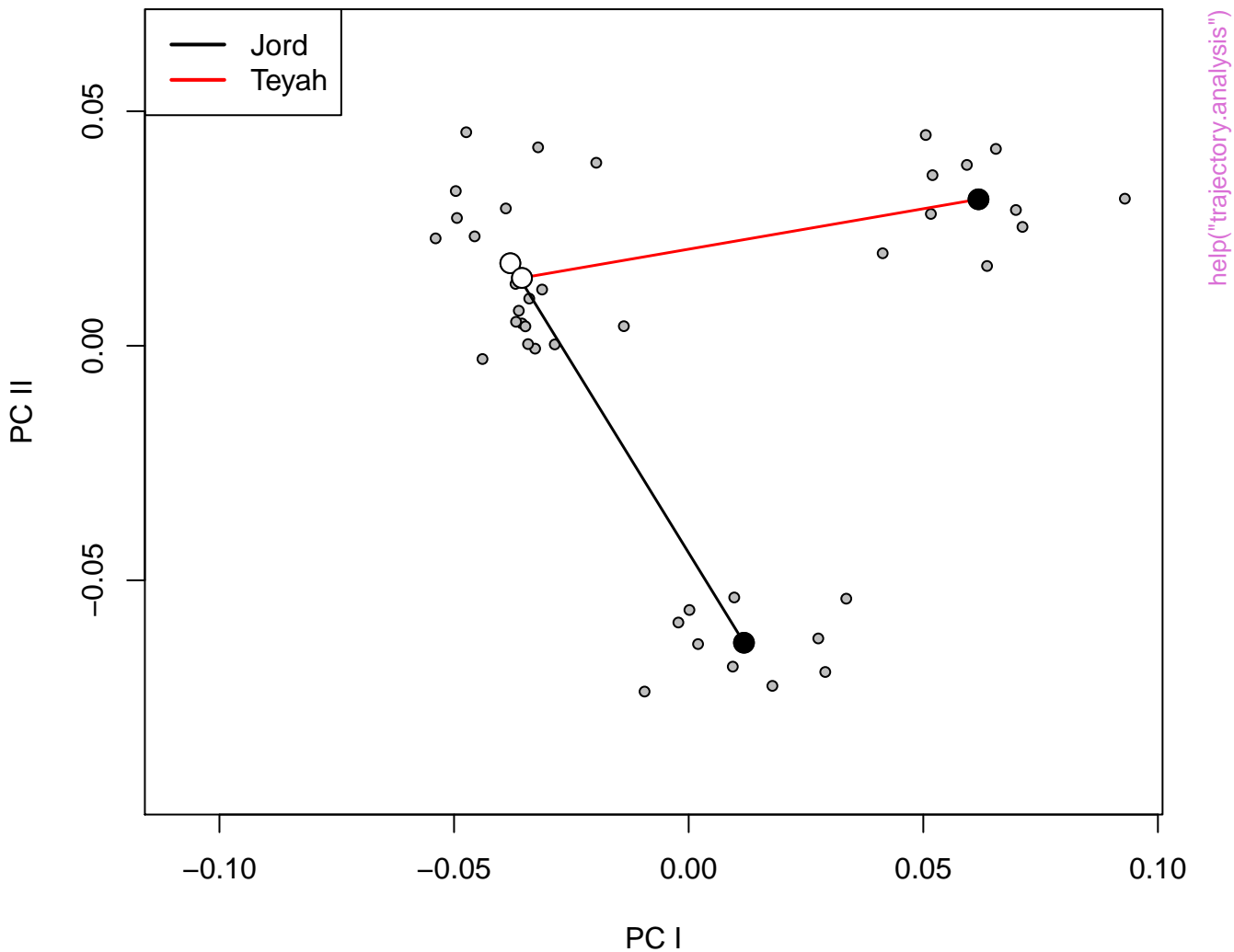


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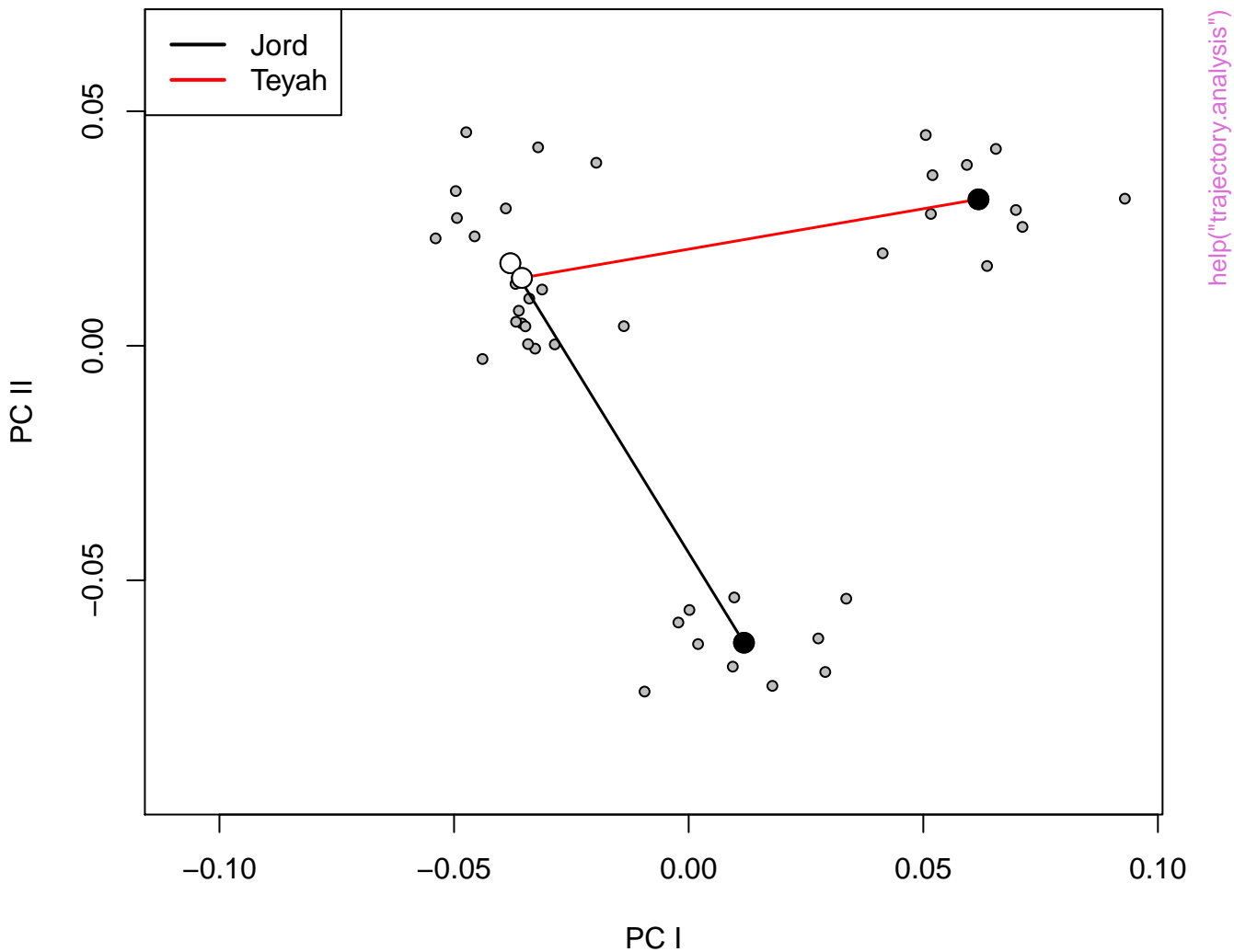


help("shape.predictor")

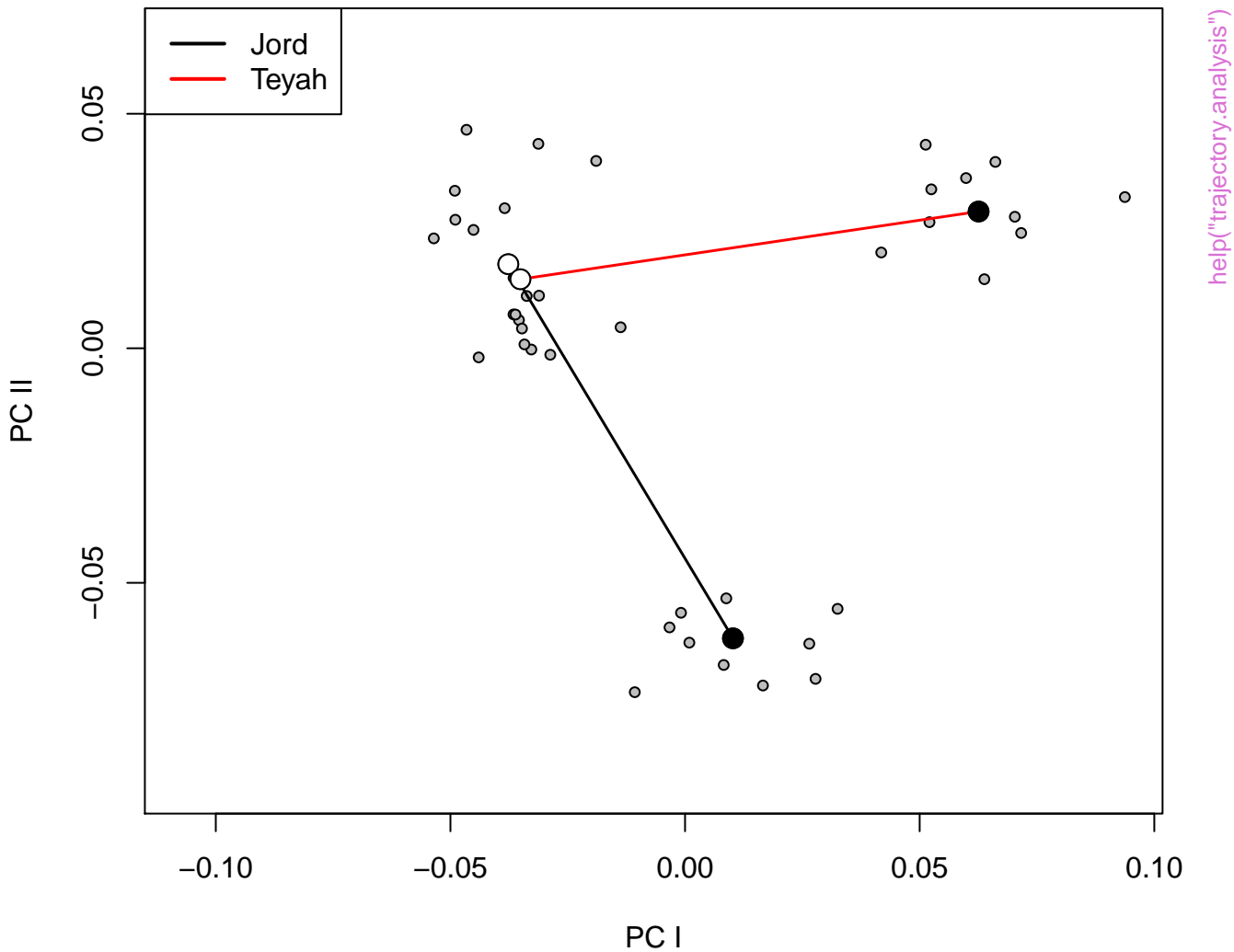
Two Dimensional View of Phenotypic Trajectories



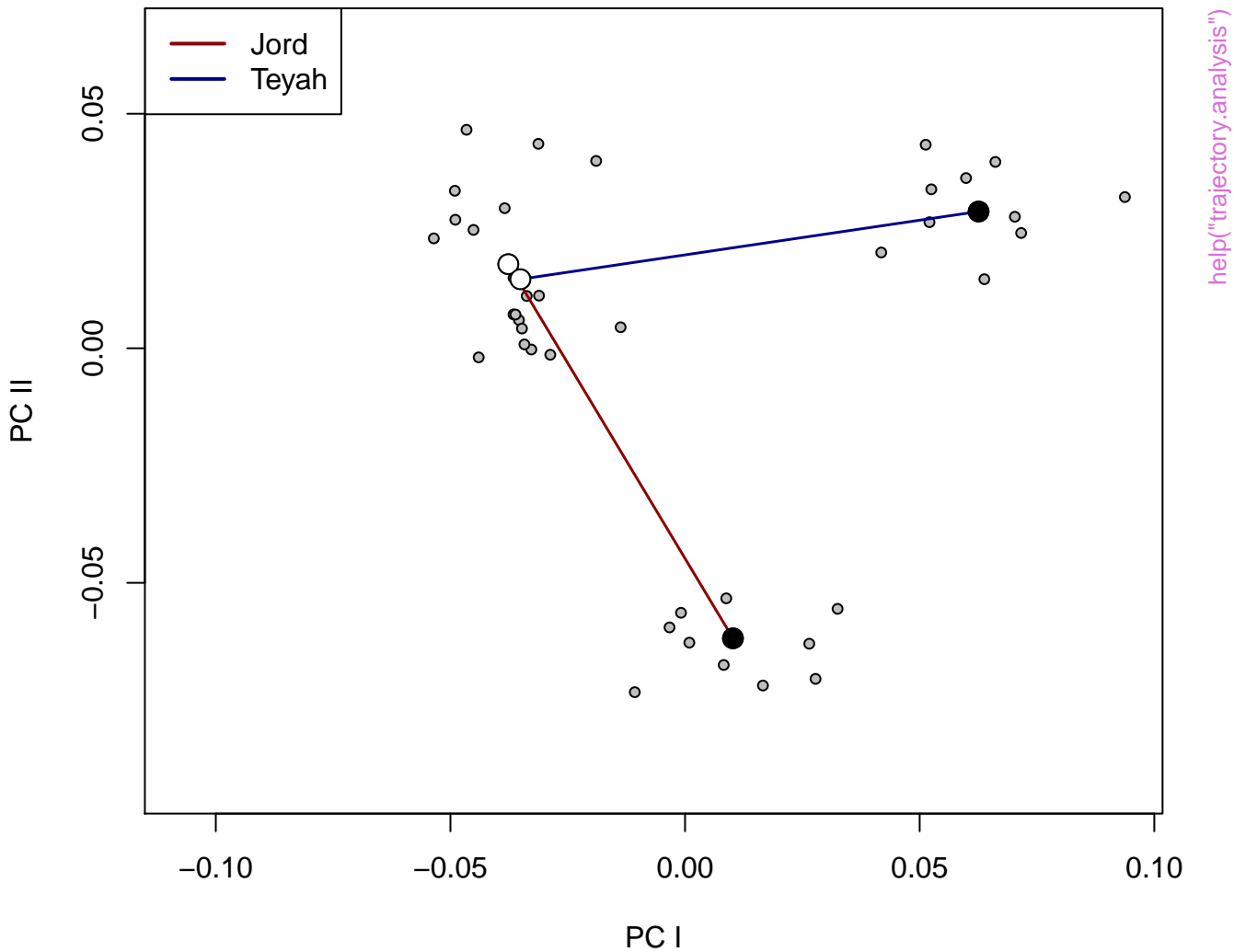
Two Dimensional View of Phenotypic Trajectories



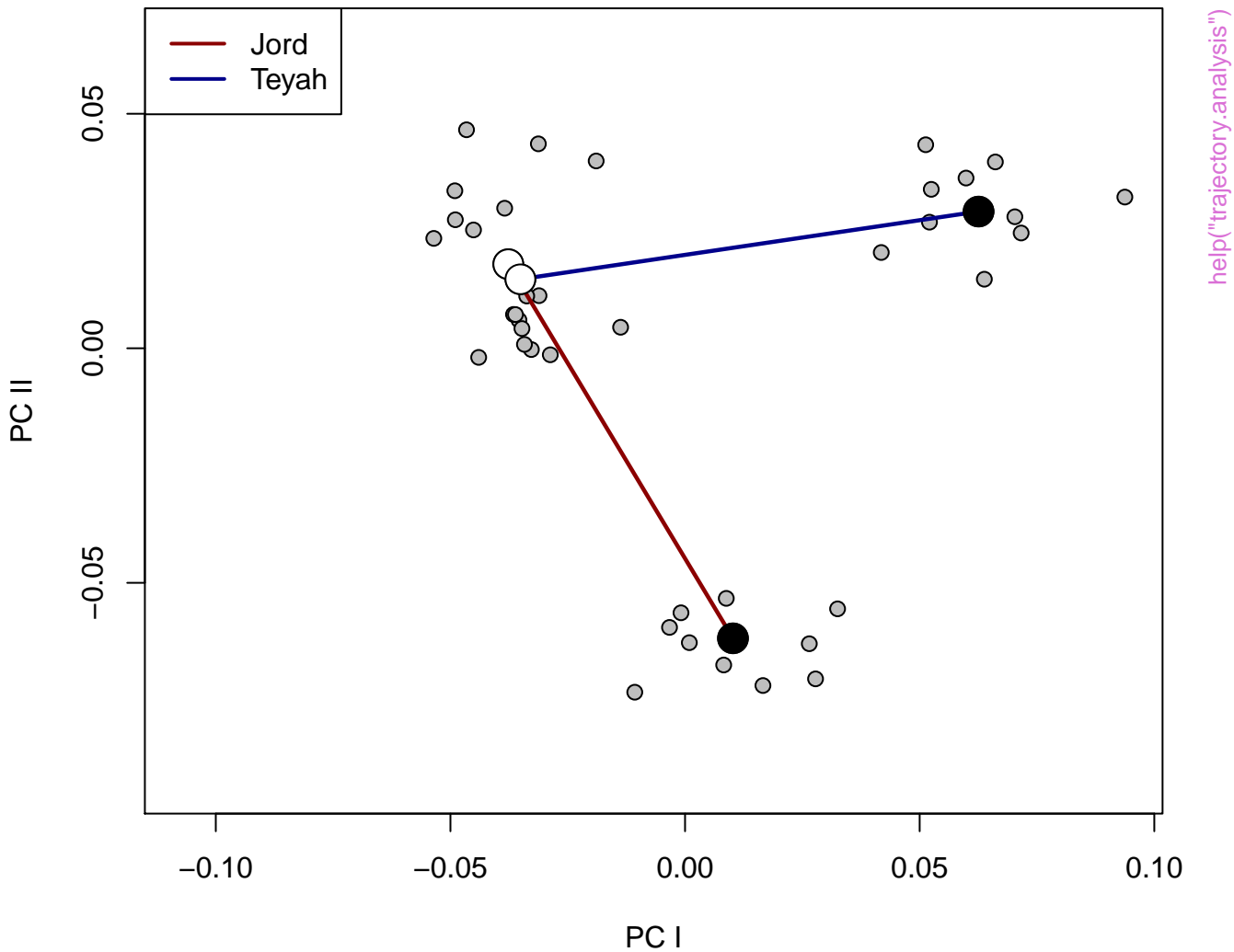
Two Dimensional View of Phenotypic Trajectories



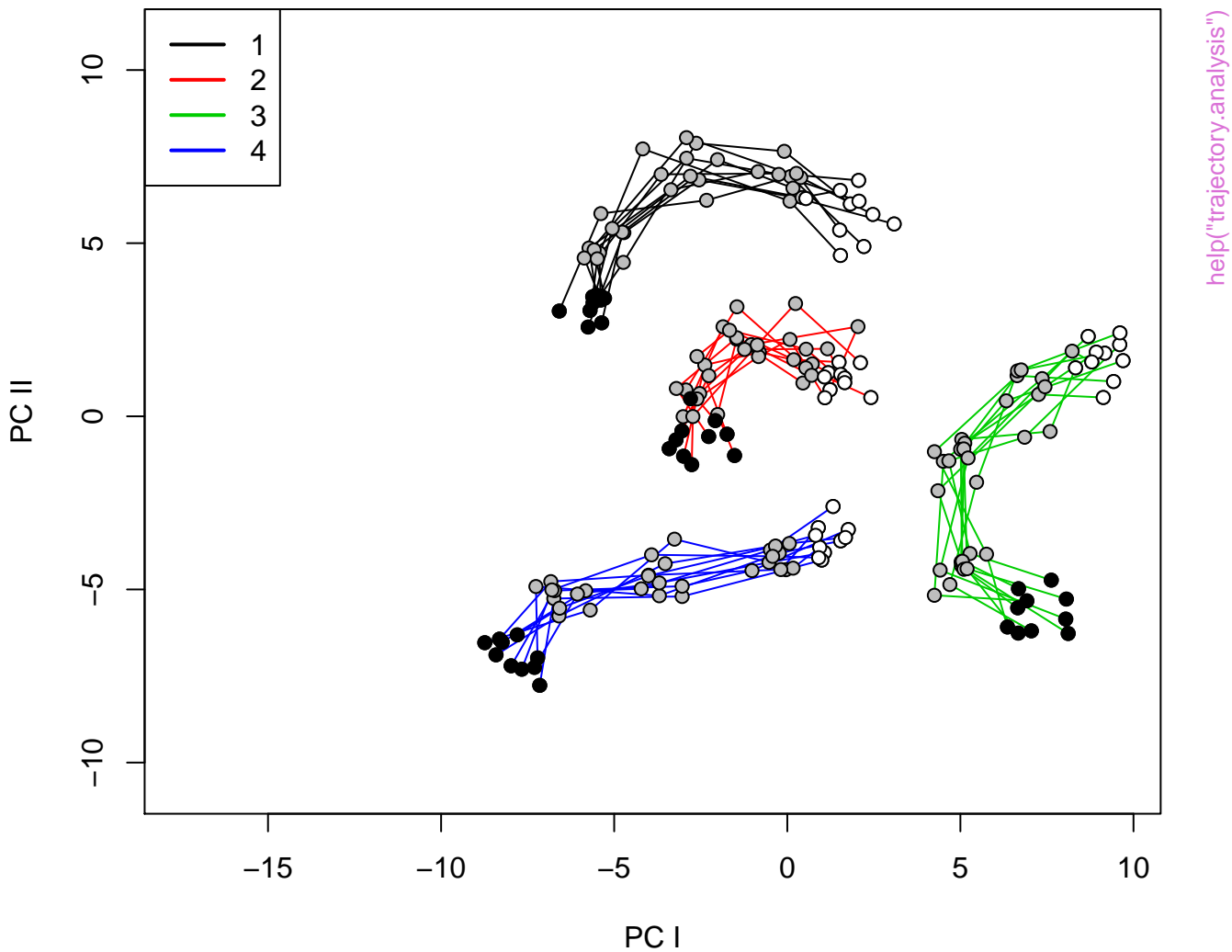
Two Dimensional View of Phenotypic Trajectories



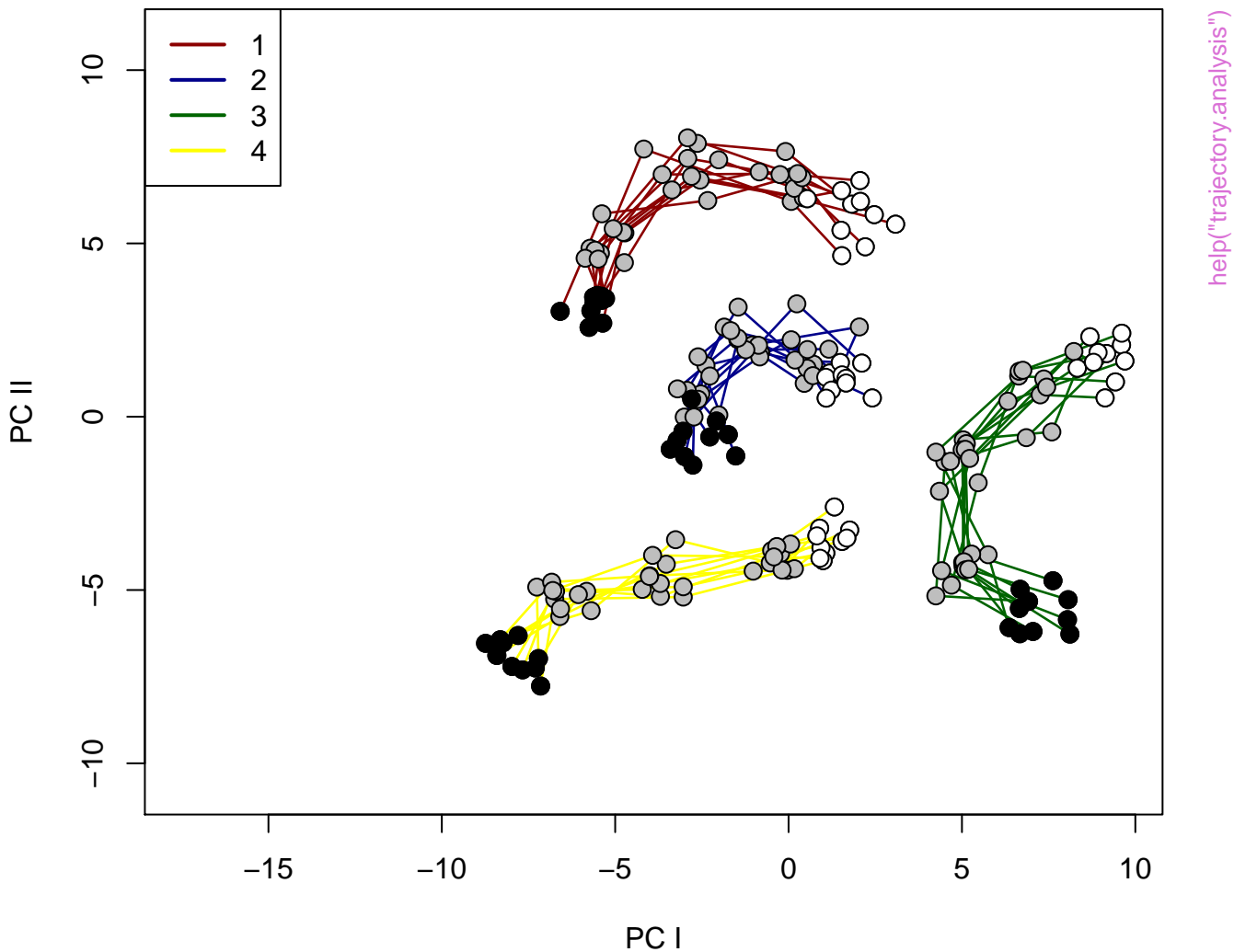
Two Dimensional View of Phenotypic Trajectories



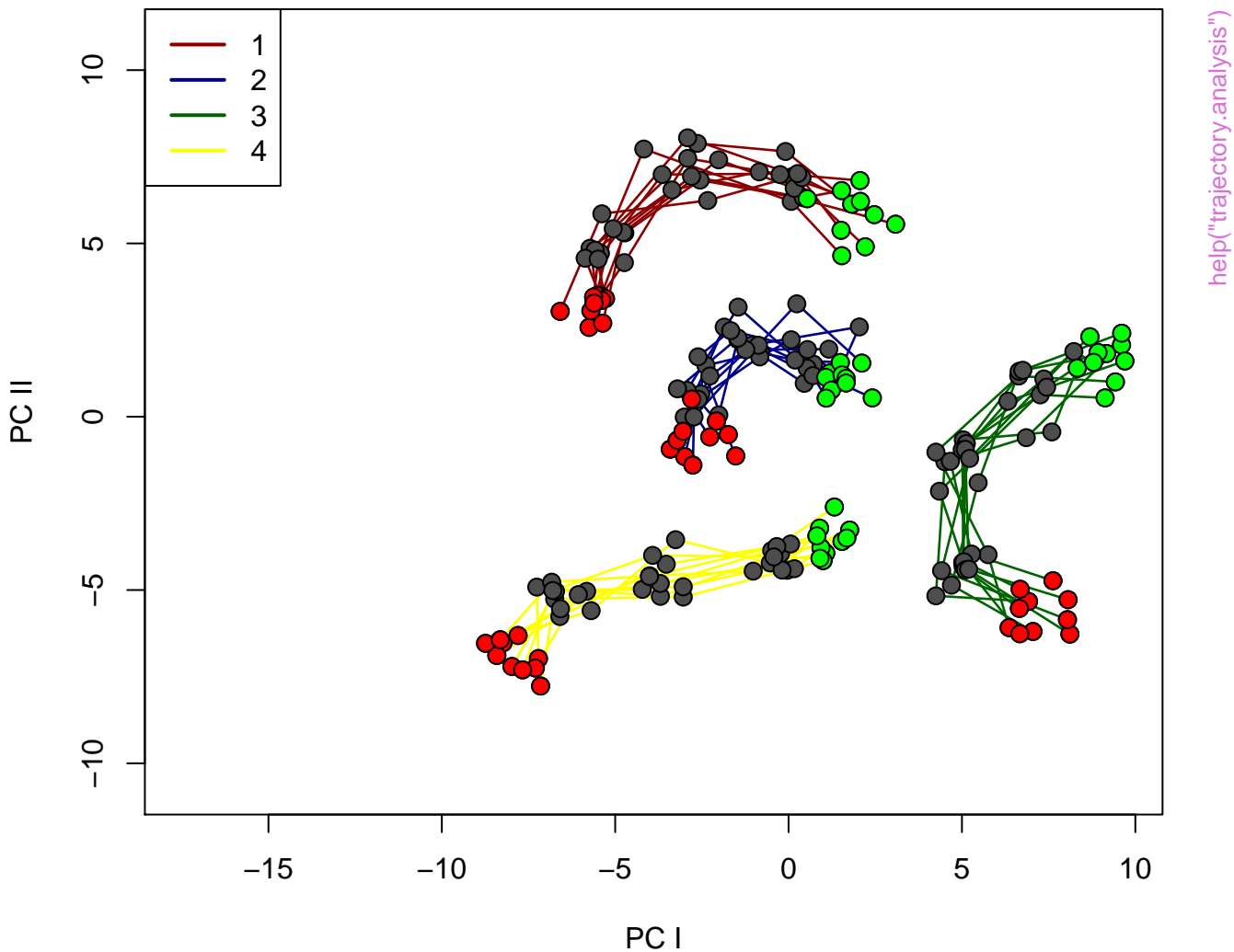
Two Dimensional View of Phenotypic Trajectories



Two Dimensional View of Phenotypic Trajectories



Two Dimensional View of Phenotypic Trajectories



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

