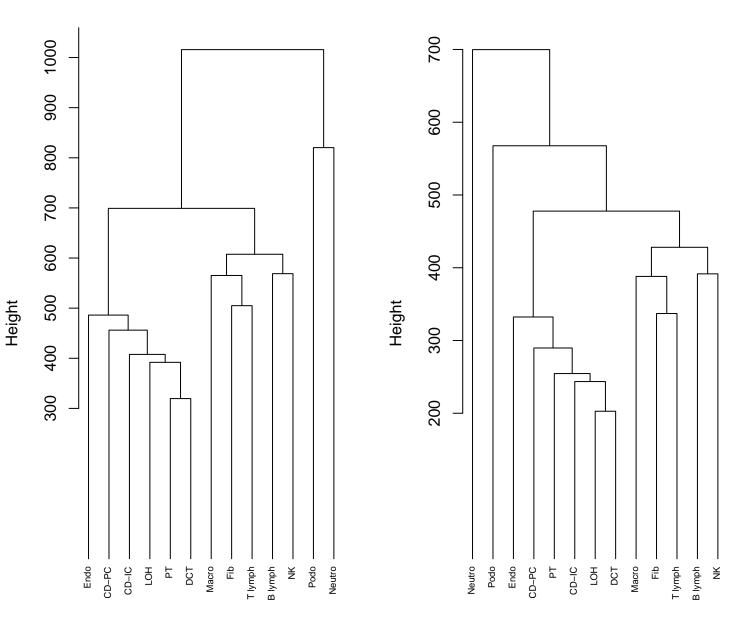


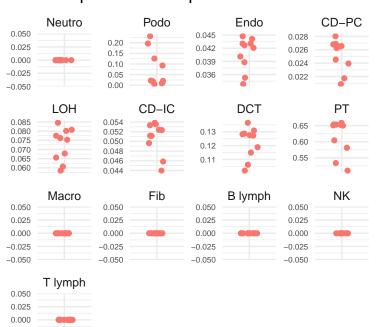
Cluster log(Mean of RA)



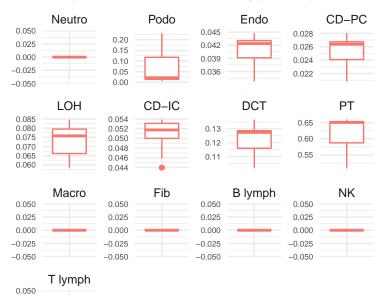
d1 hclust (*, "complete")

d2 hclust (*, "complete")

Jitter plot of Est Proportions



Boxplot of Estimated Cell Type Proportions



0.025

0.000

-0.025

-0.050

Heatmap of Estimated Cell Type Proportions

-0.025

-0.050

