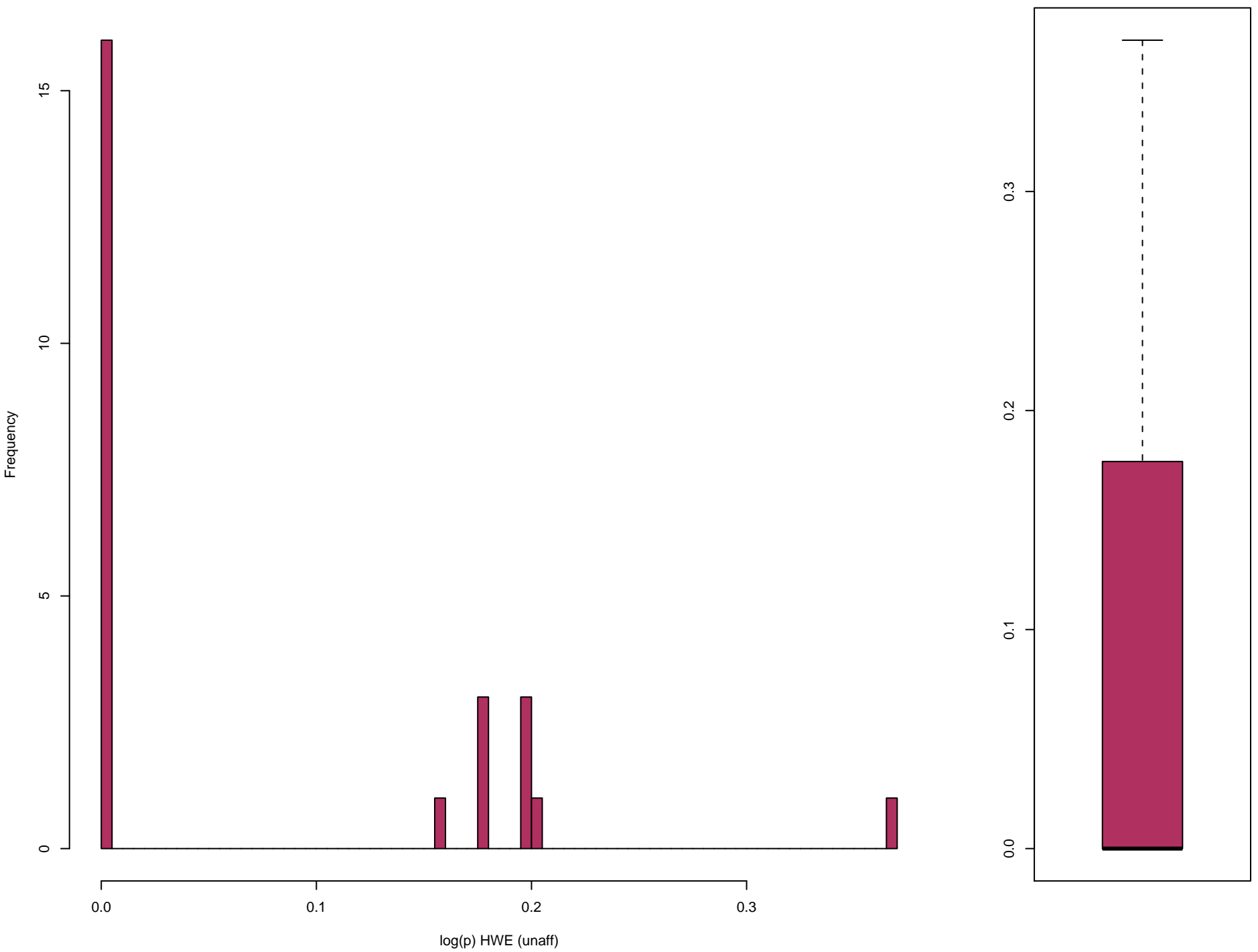
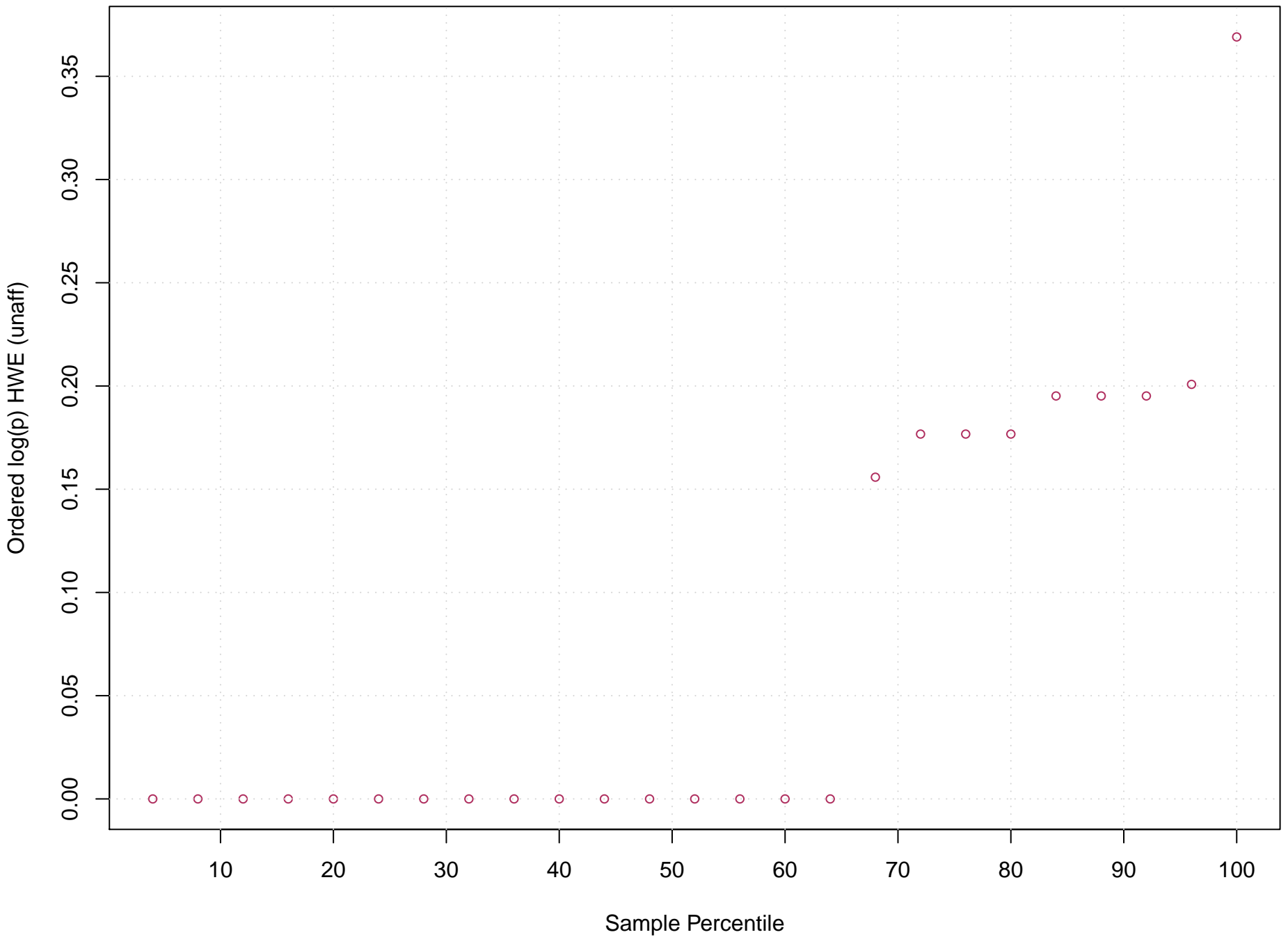


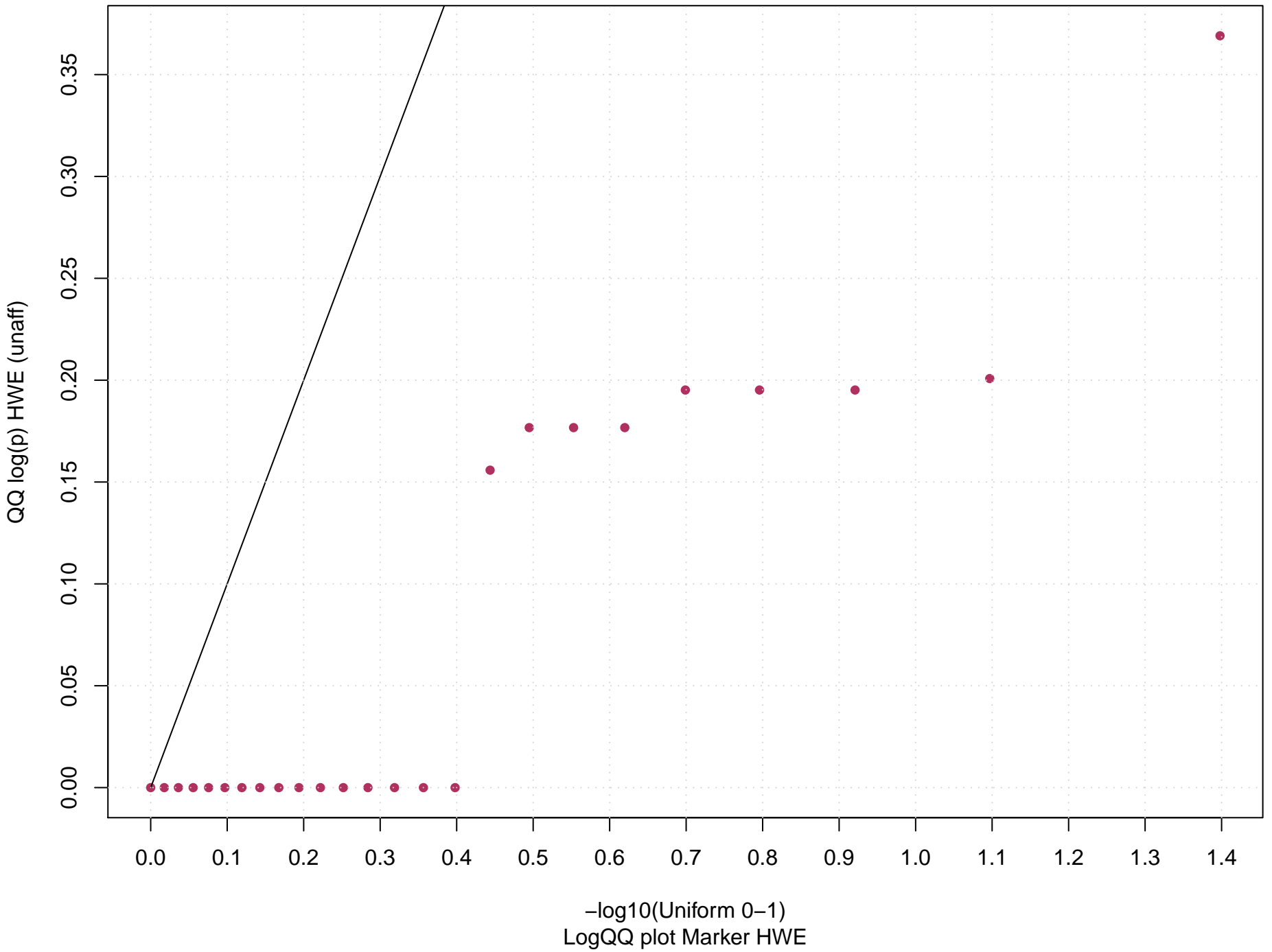
# QC for tinywga – Marker HWE



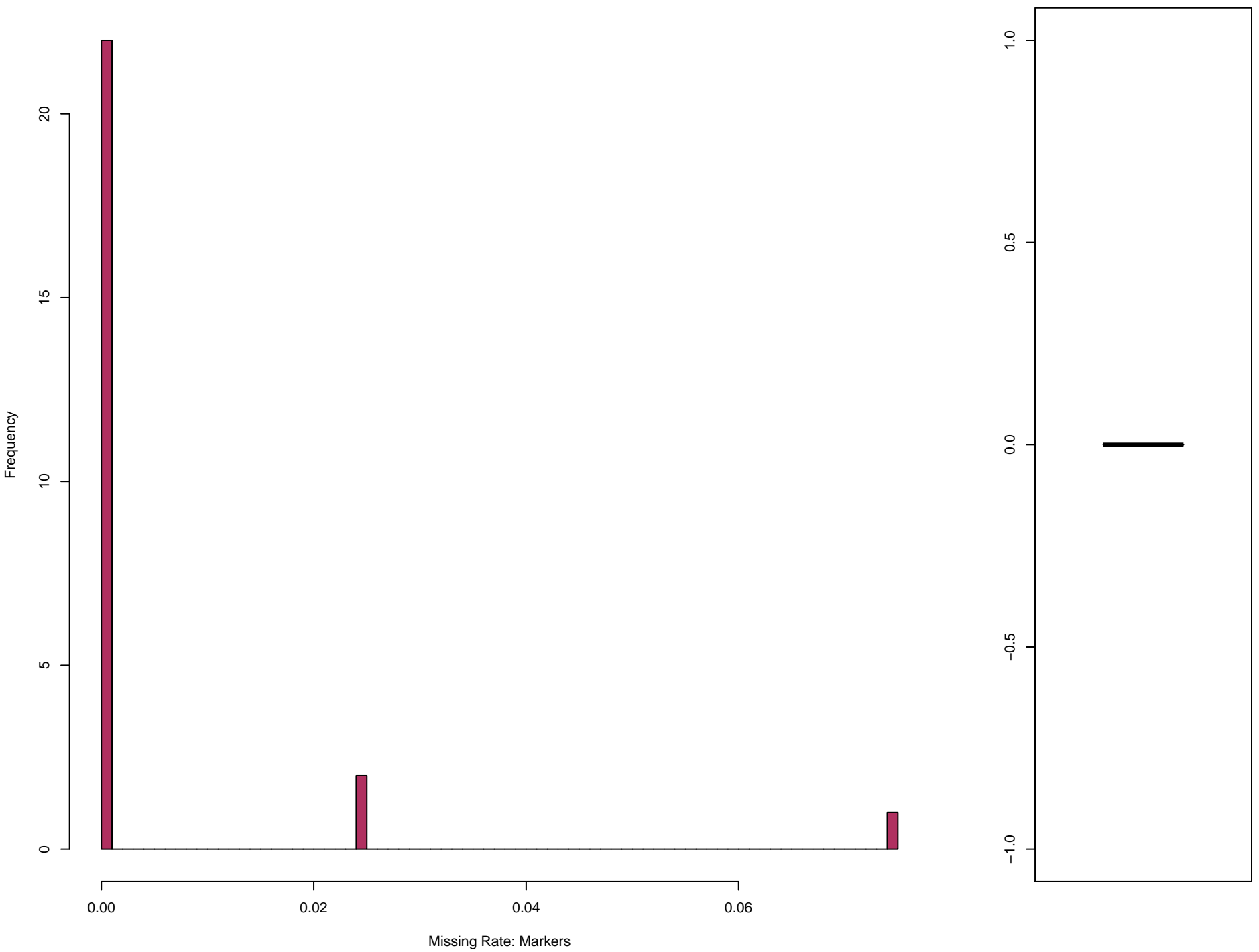
# QC for tinywga – Ranked Marker HWE



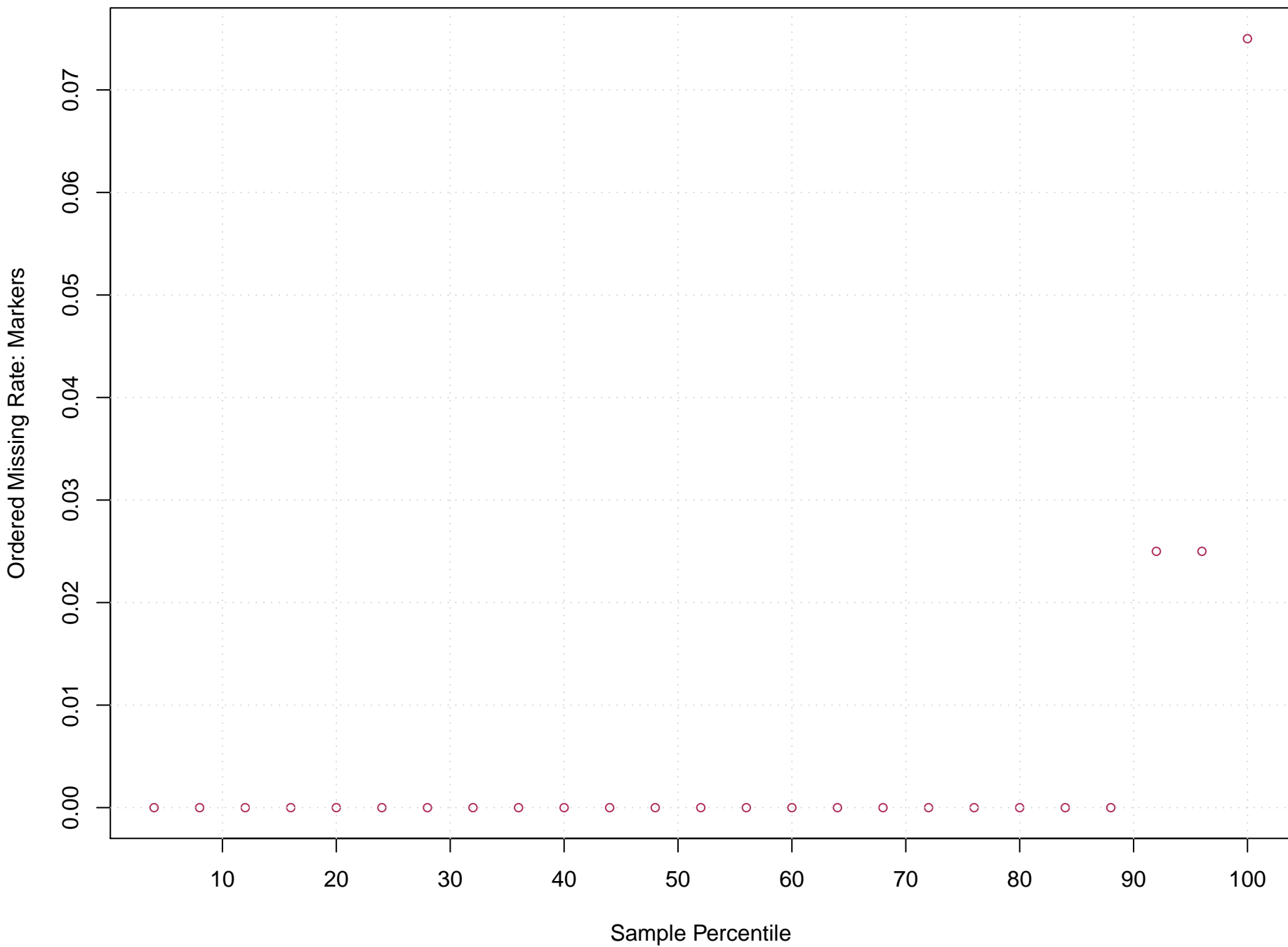
Log QQ Plot(n=25)



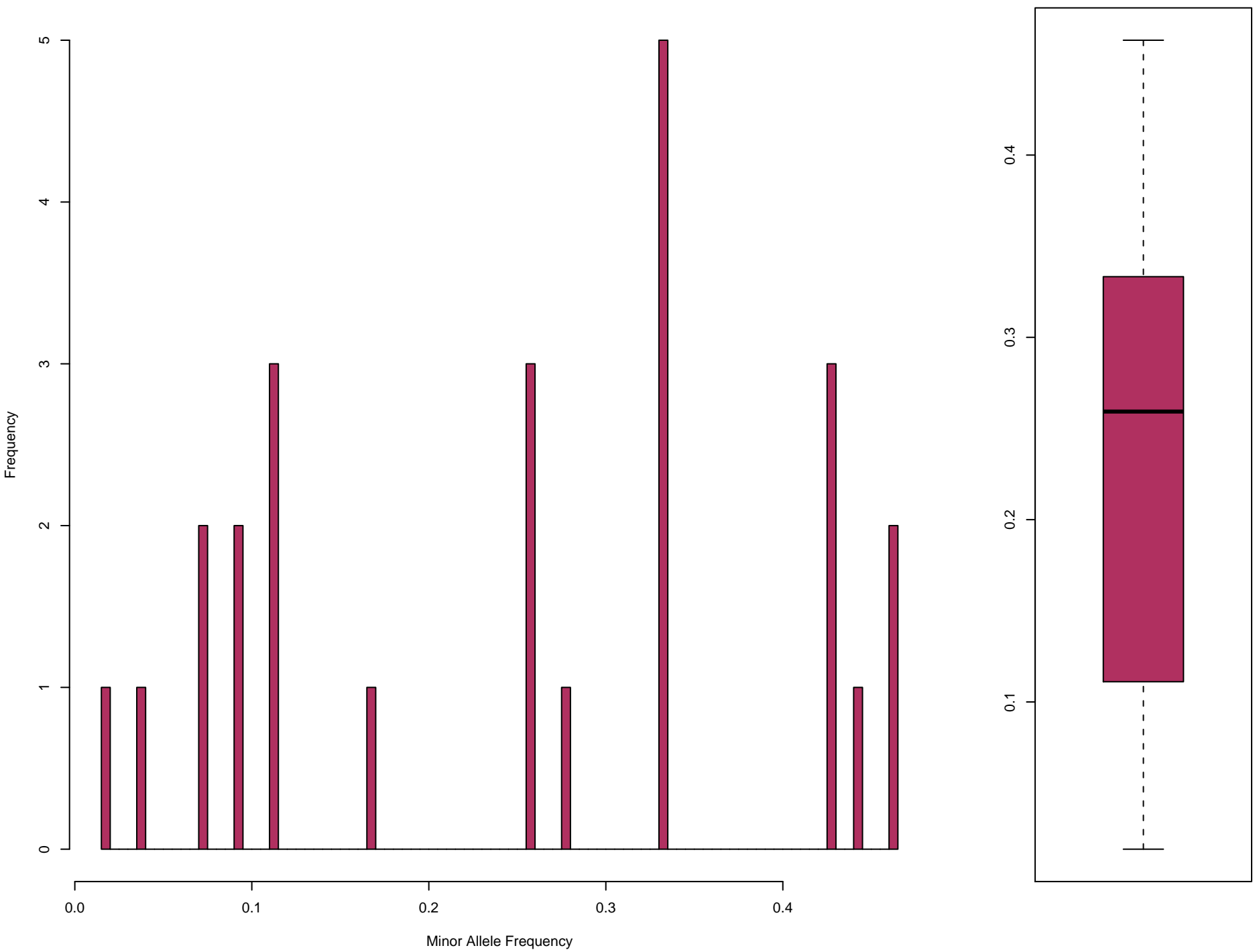
# QC for tinywga – Marker Missing Genotype



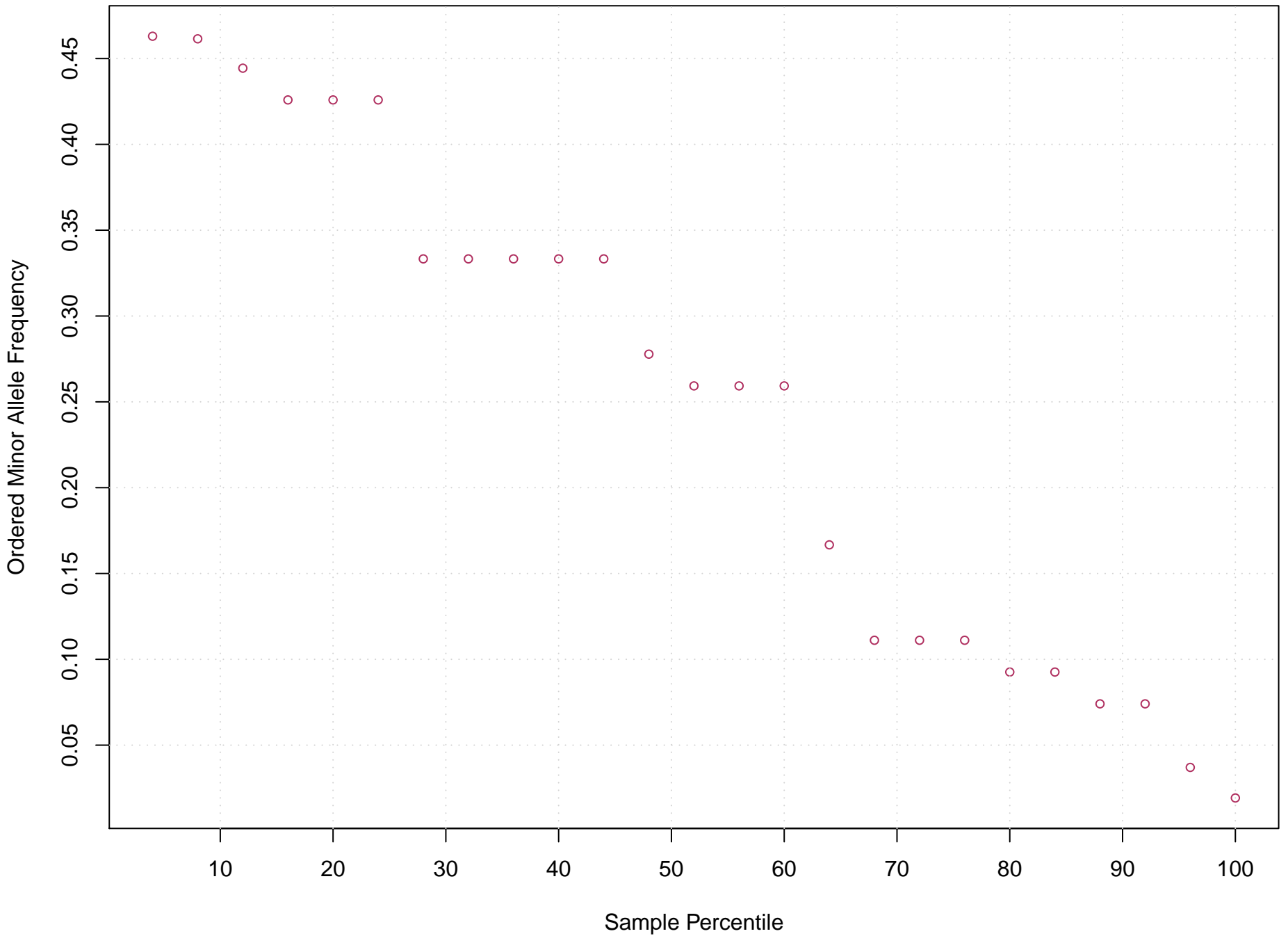
# QC for tinywga – Ranked Marker Missing Genotype



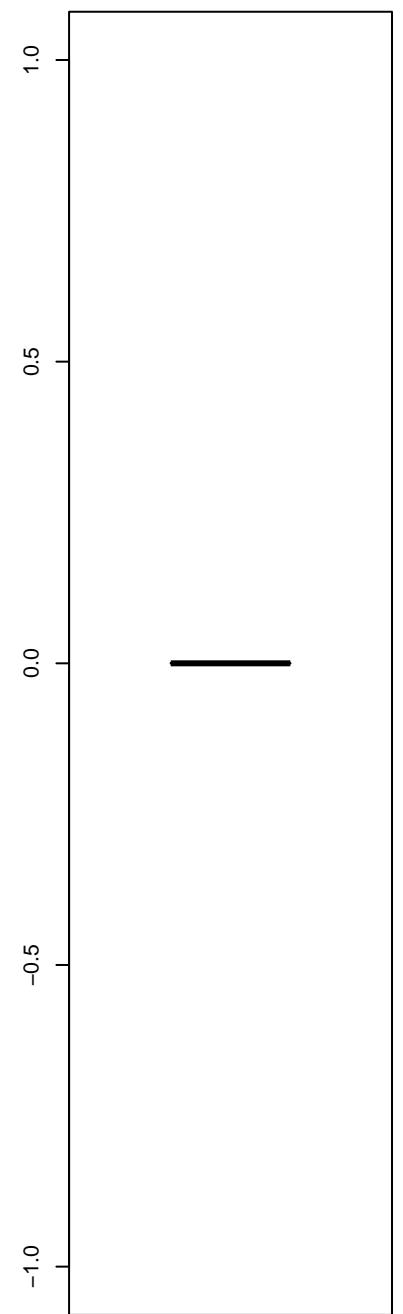
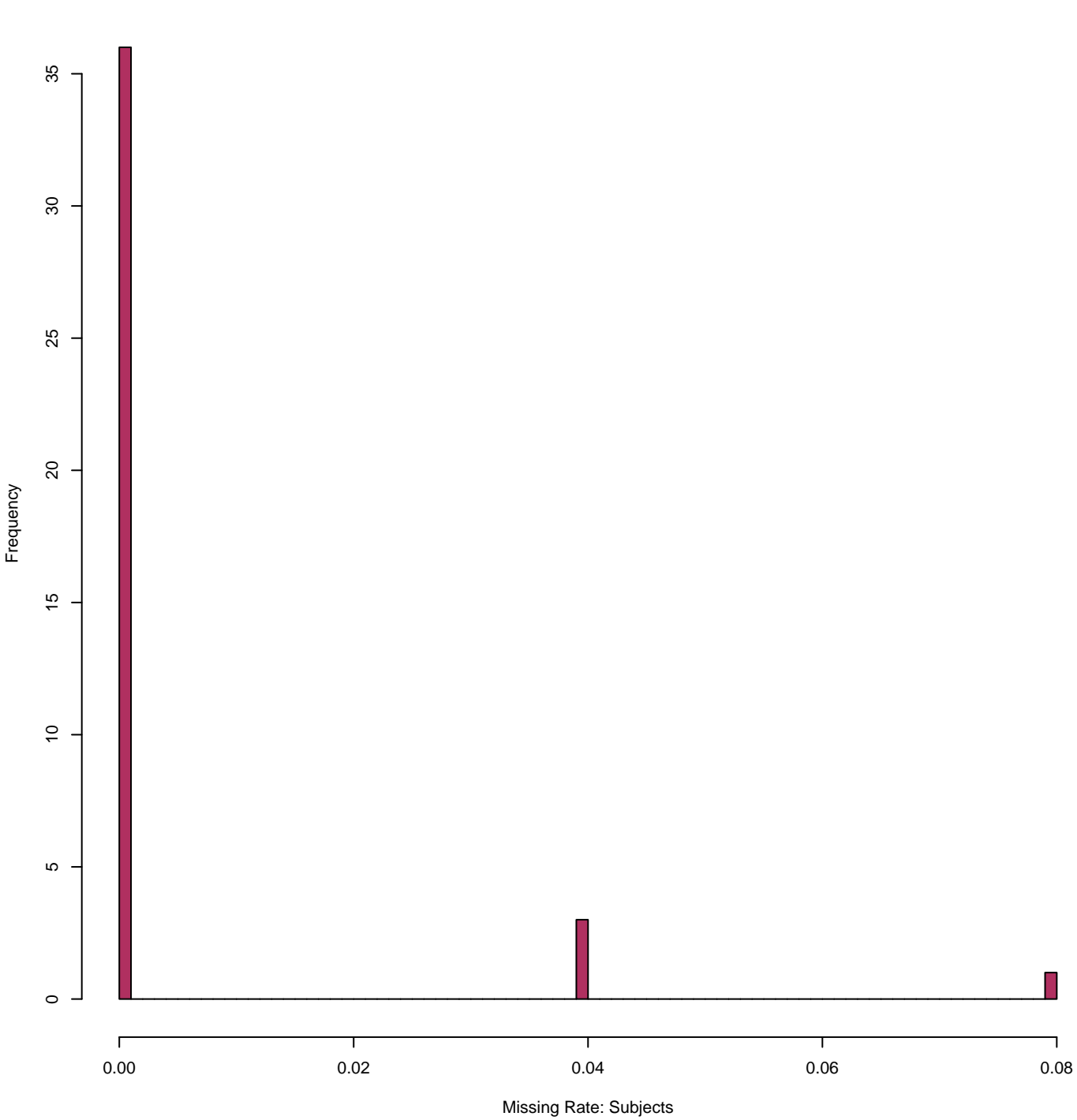
### QC for tinywga – Marker MAF



# QC for tinywga – Ranked Marker MAF

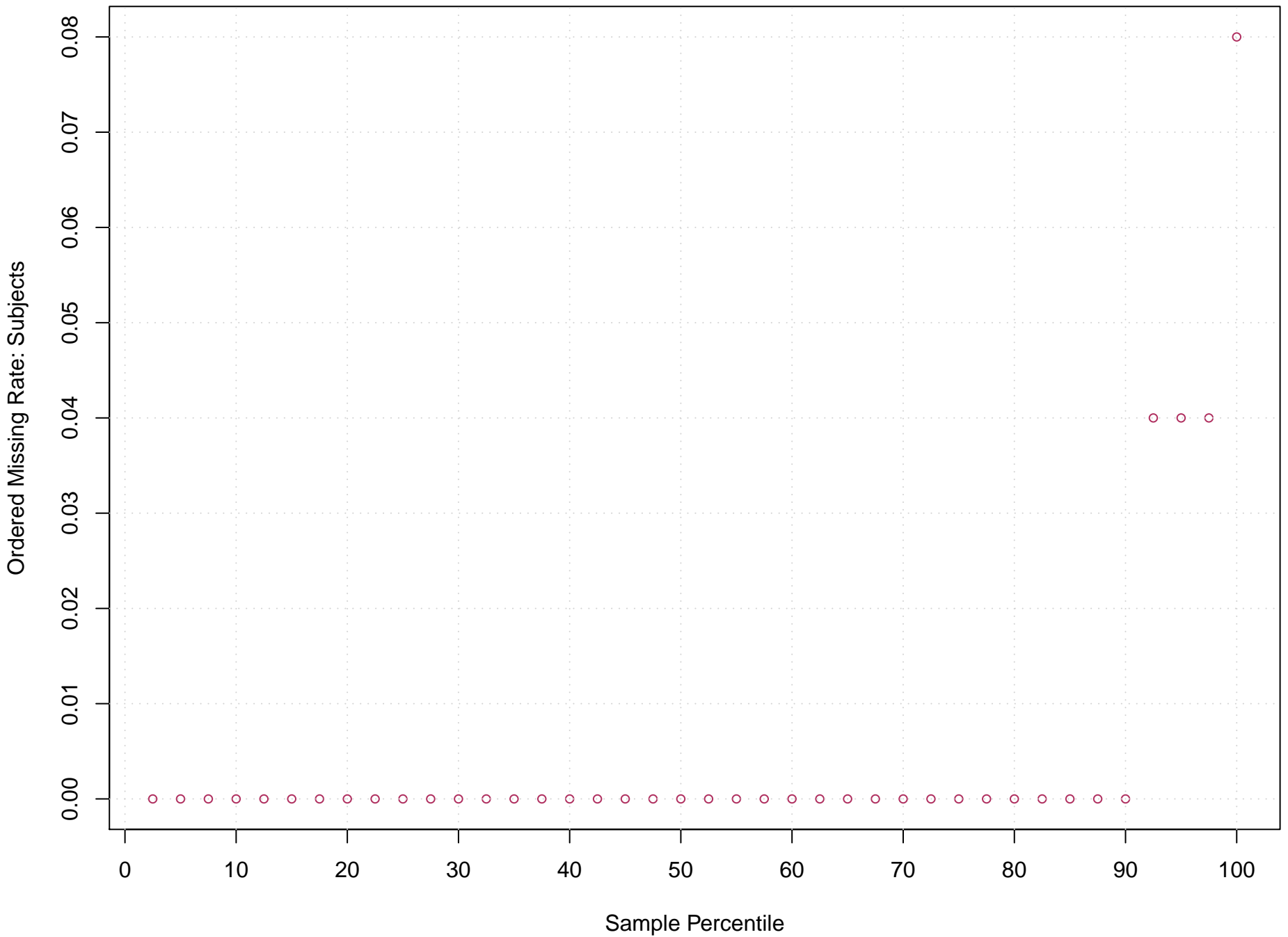


# QC for tinywga – Subject Missing Genotype

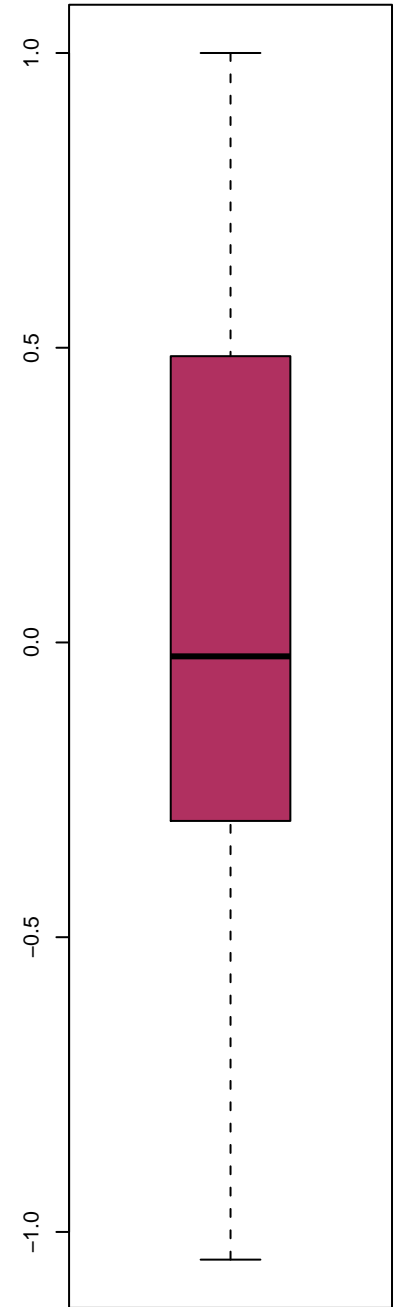
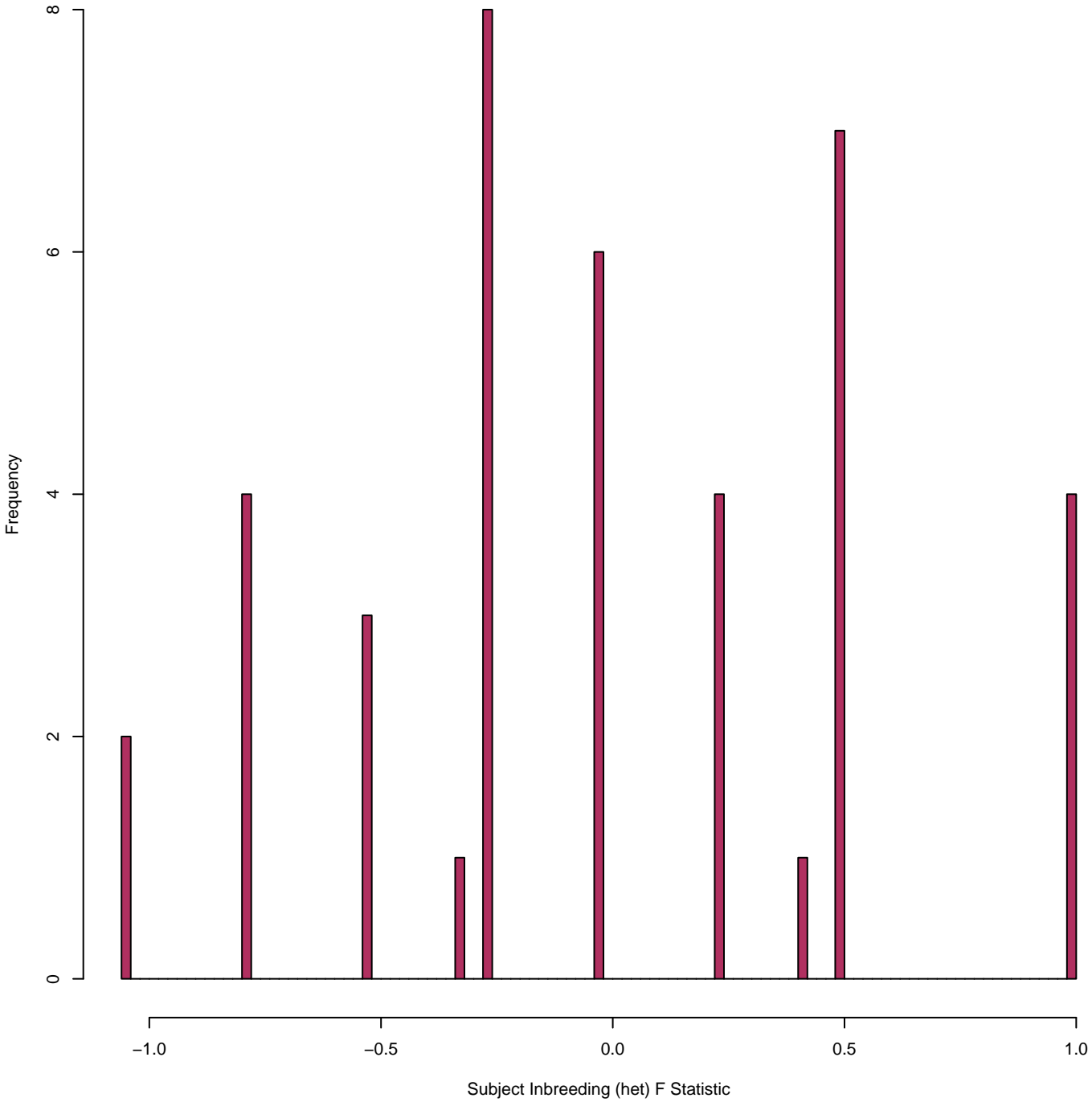




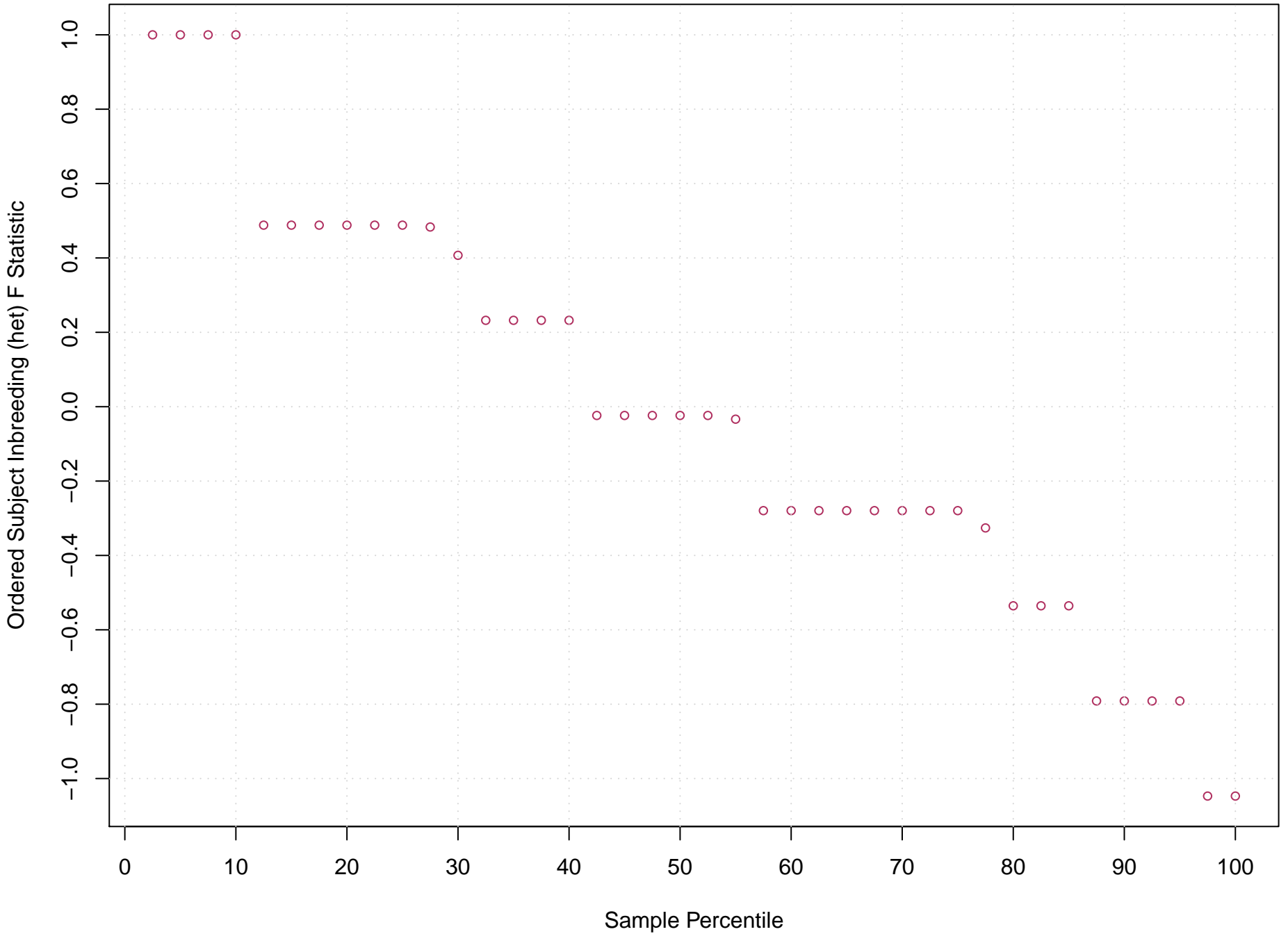
# QC for tinywga – Ranked Subject Missing Genotype



QC for tinywga – Subject F Statistic



# QC for tinywga – Ranked Subject F Statistic



Log QQ Plot(n=40)

