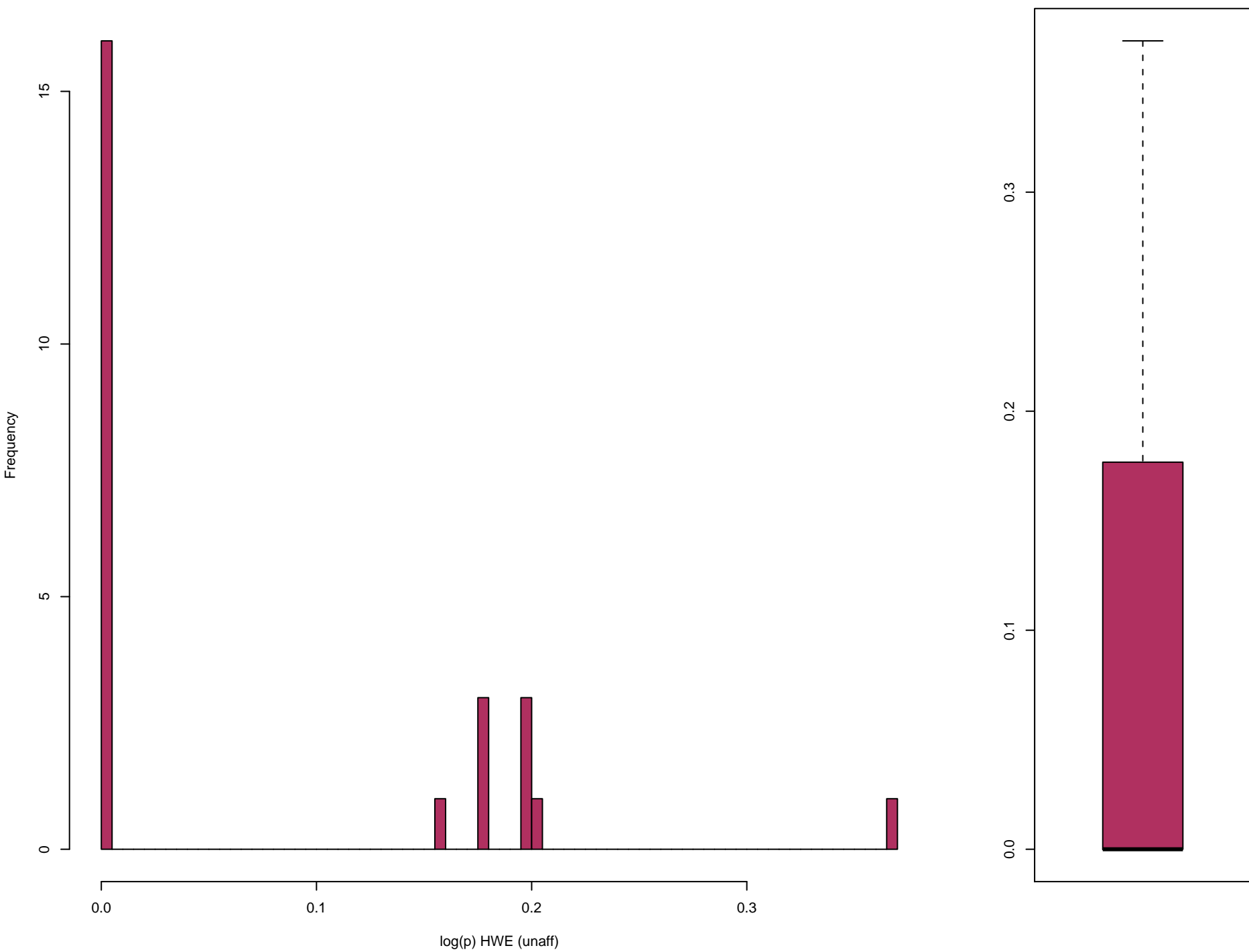
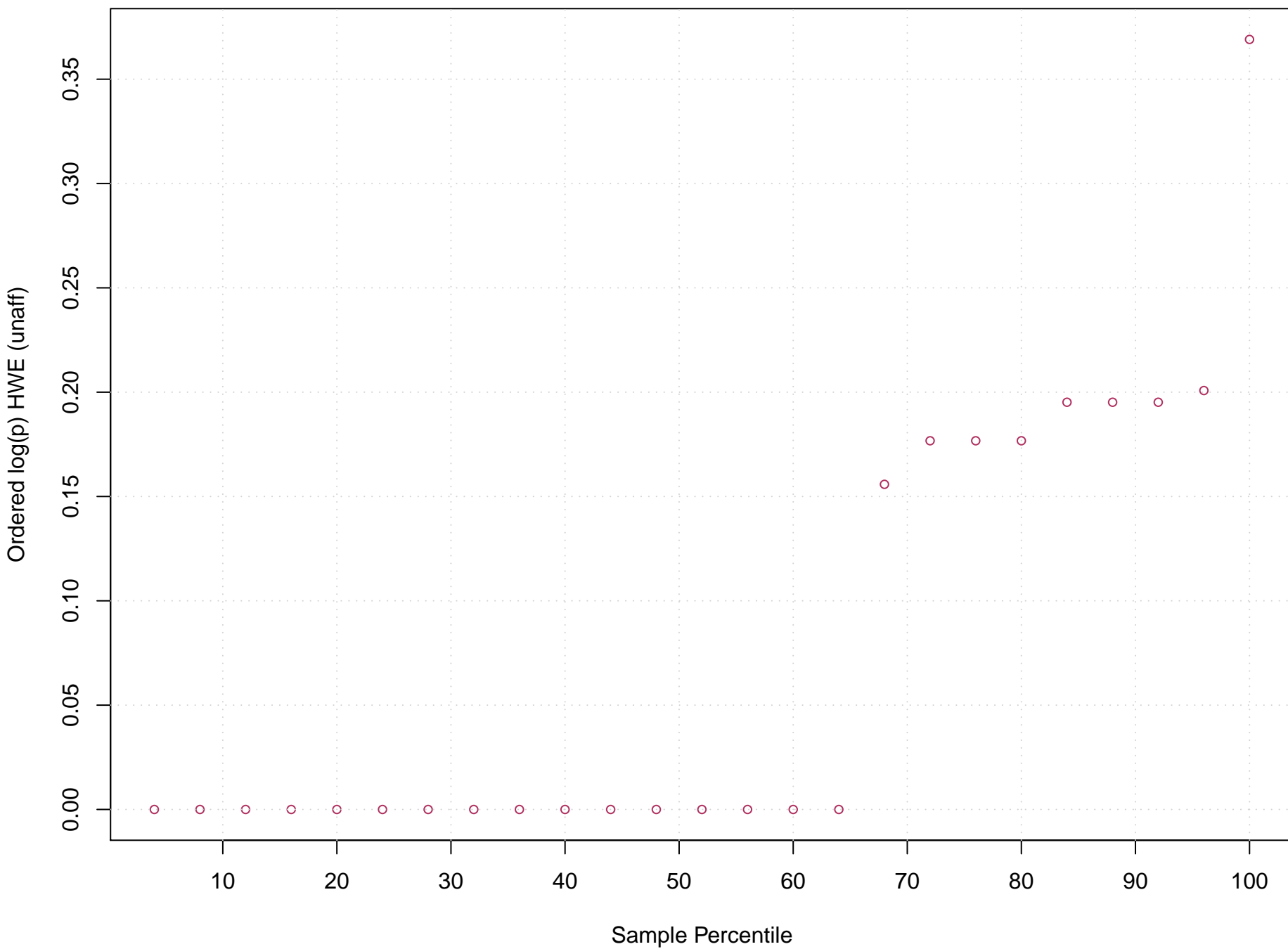


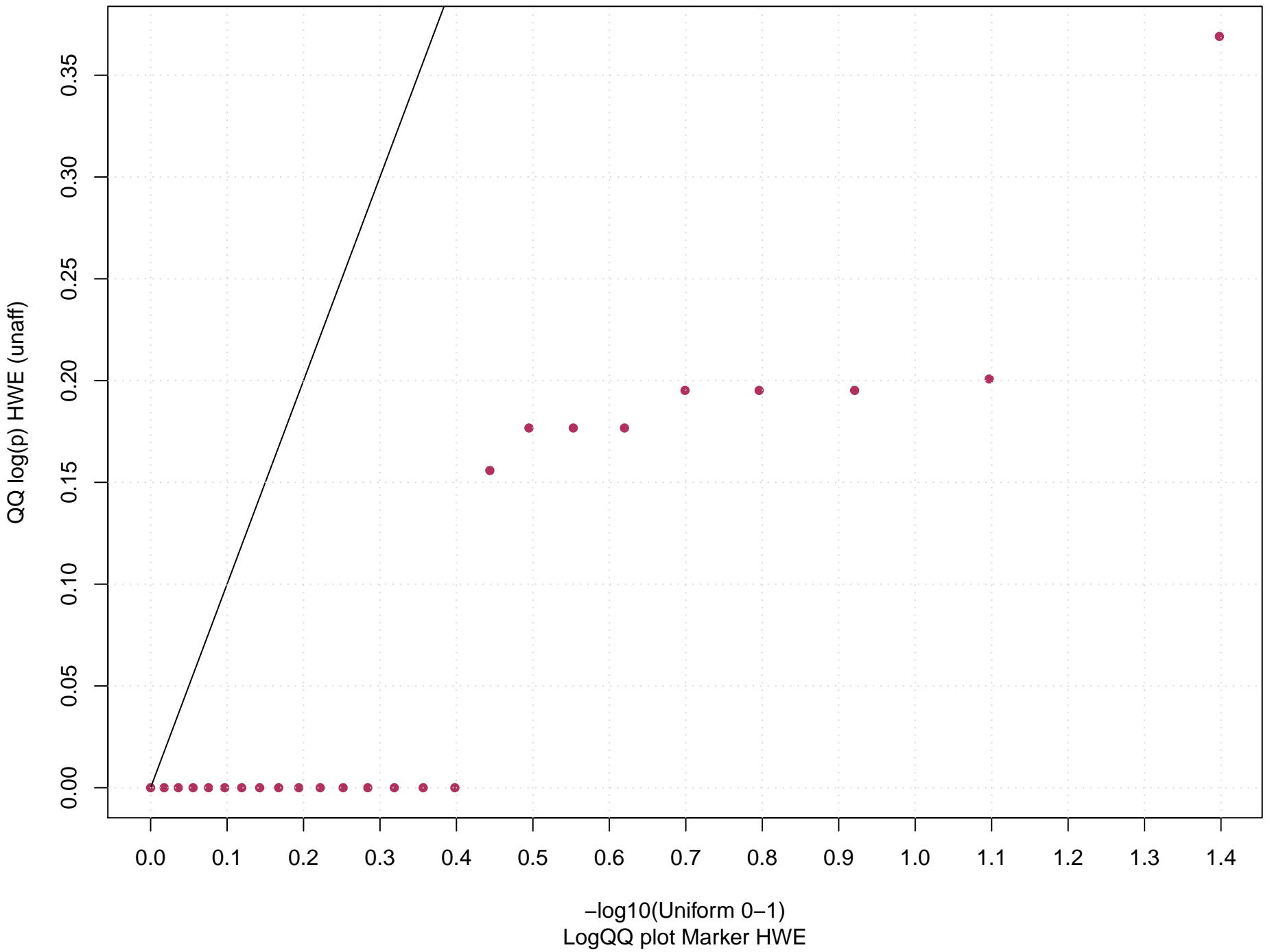
QC for tinywga – Marker HWE



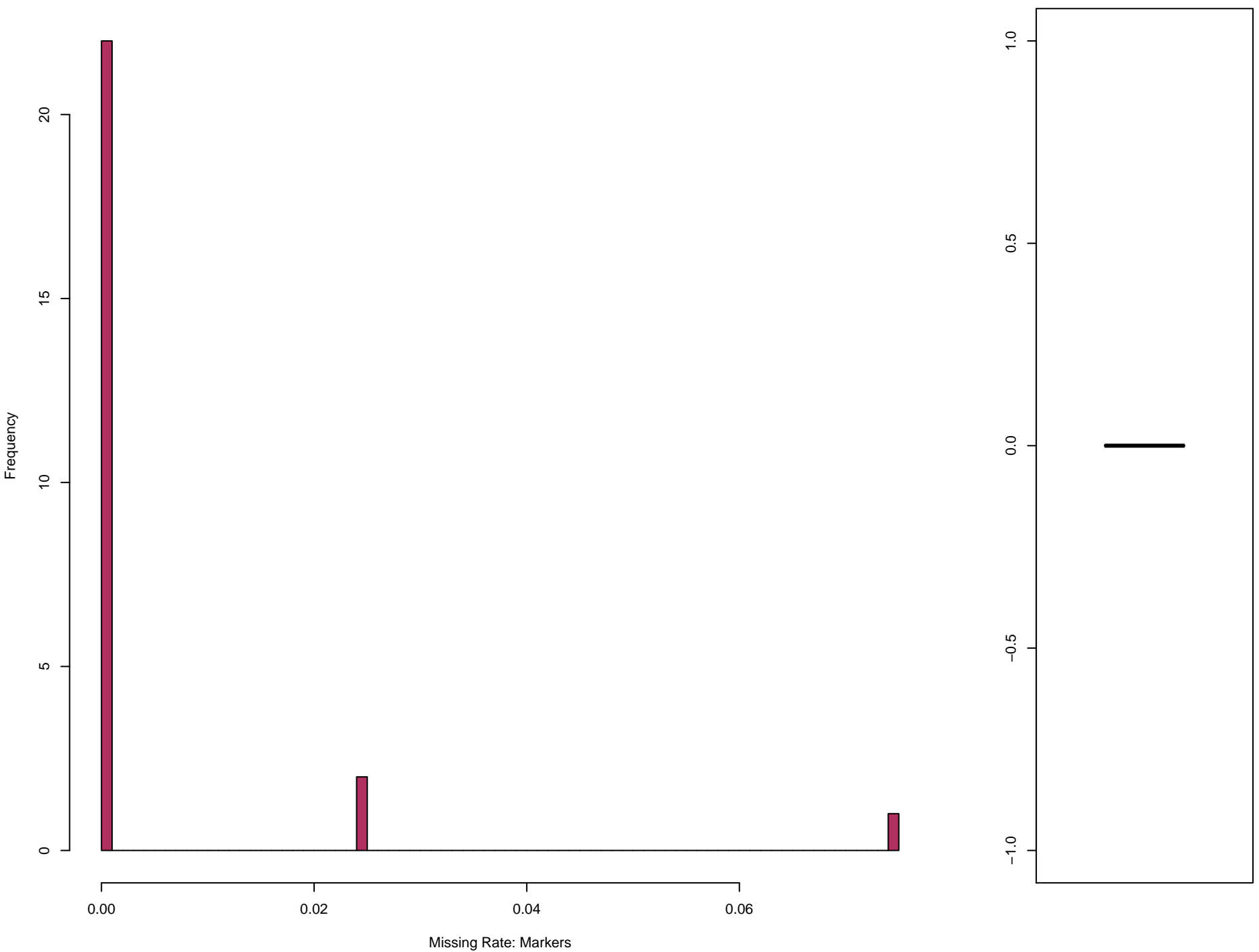
QC for tinywga – Ranked Marker HWE



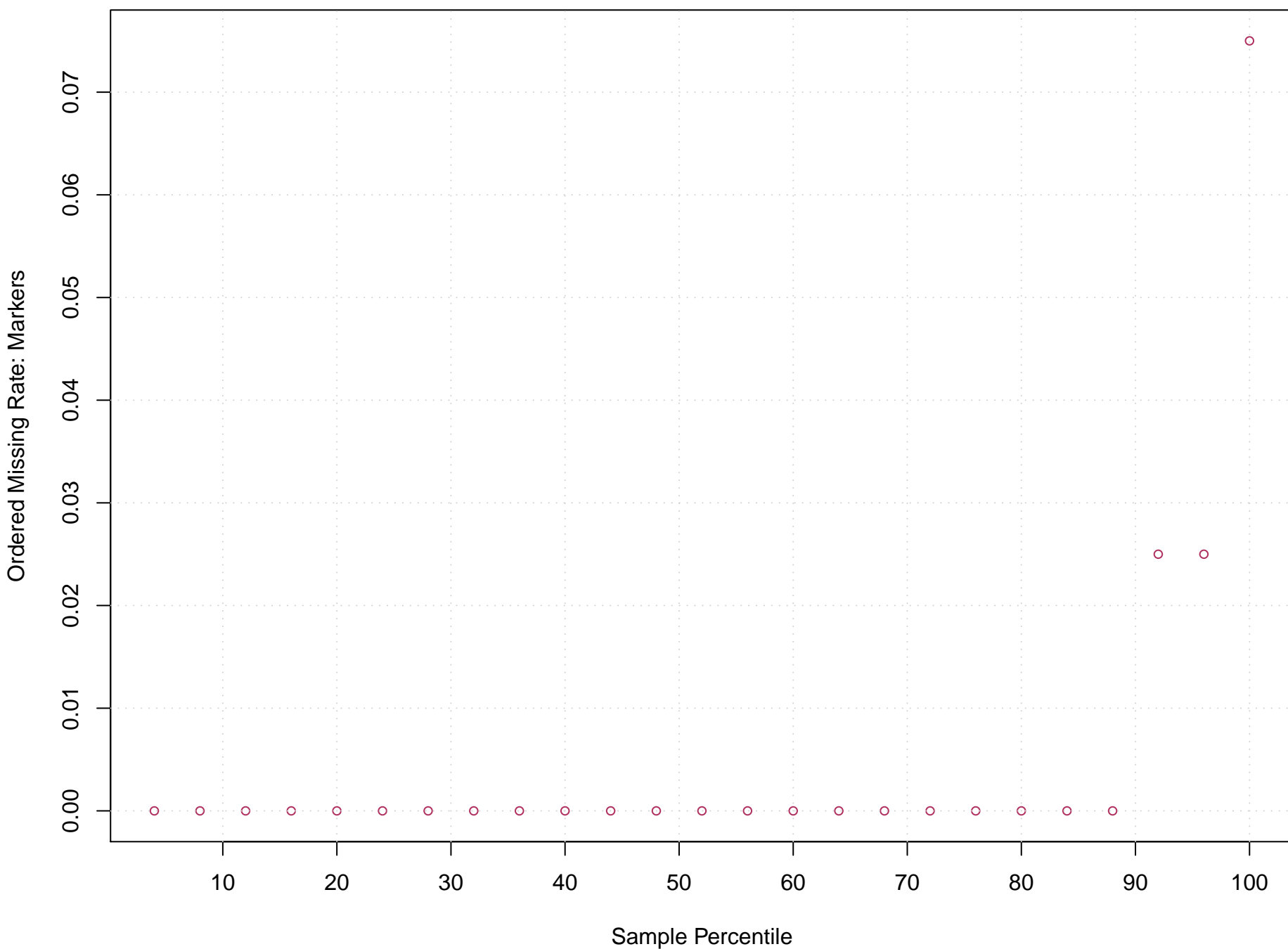
Log QQ Plot(n=25)



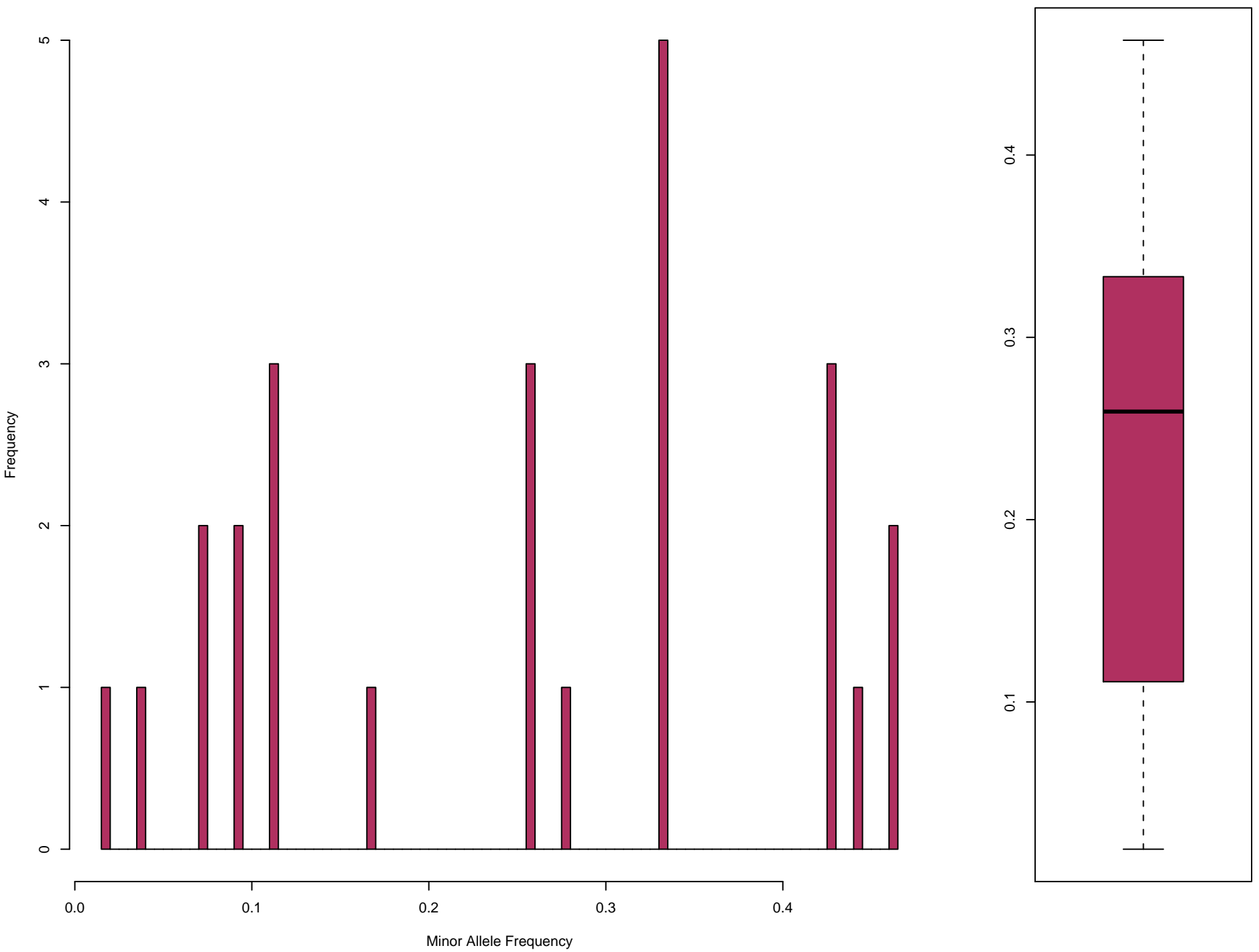
QC for tinywga – Marker Missing Genotype



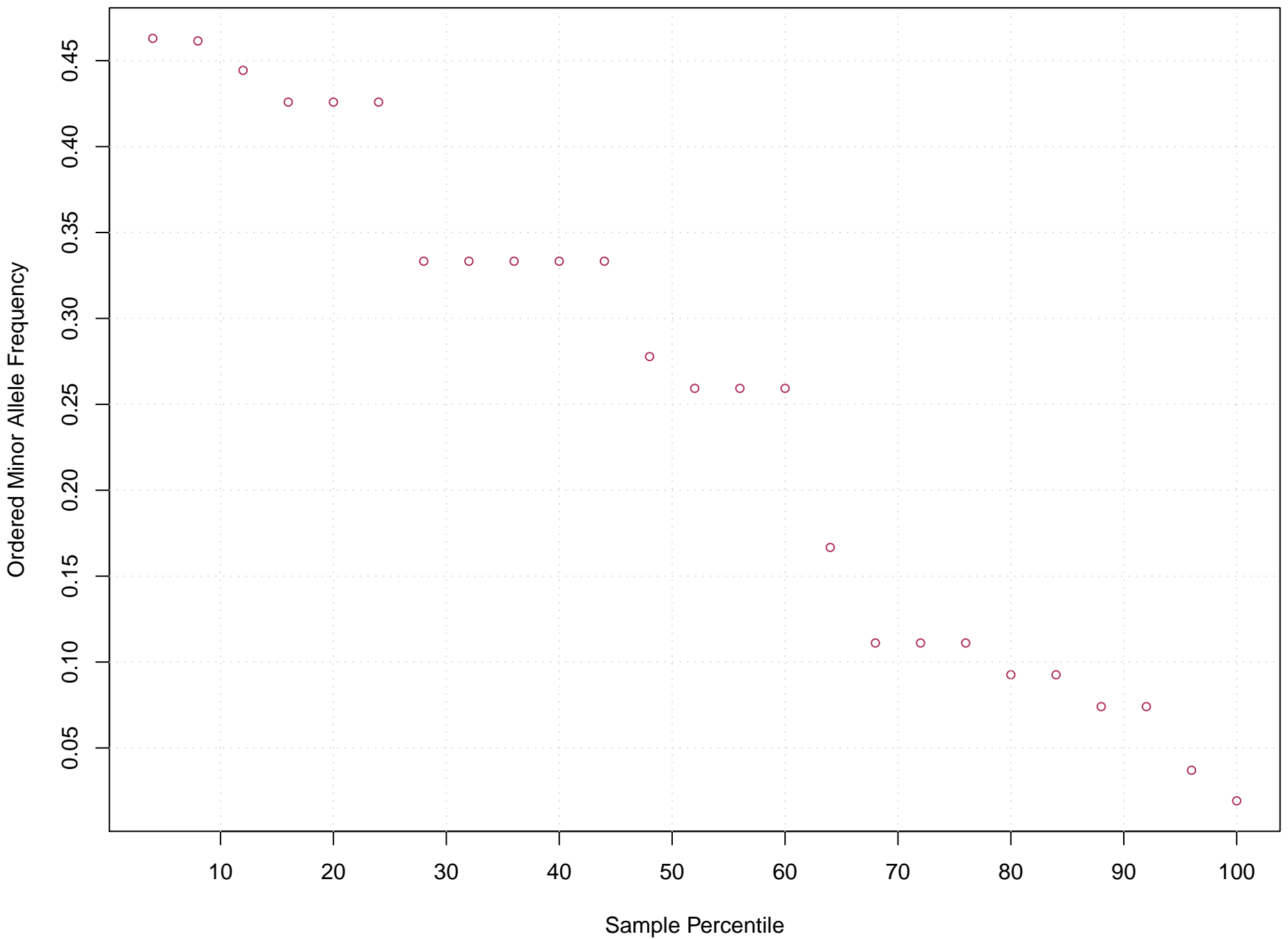
QC for tinywga – Ranked Marker Missing Genotype



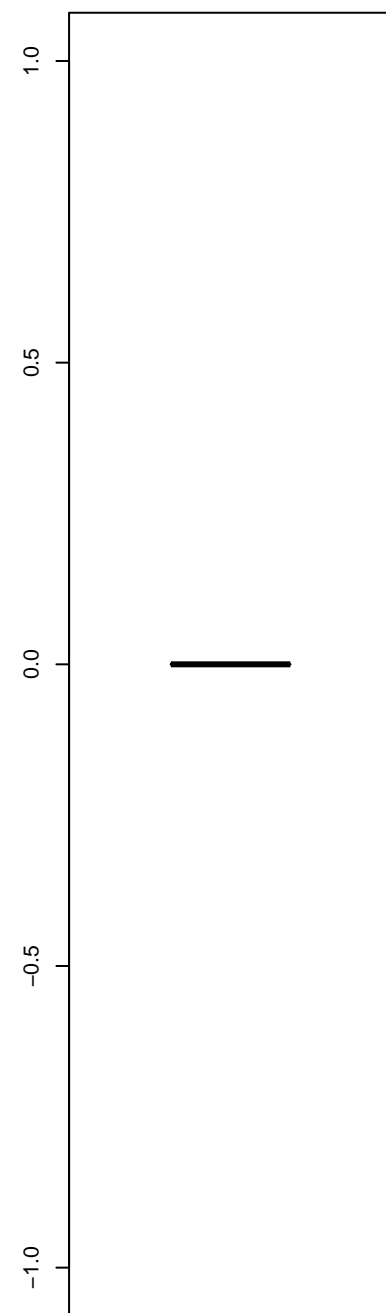
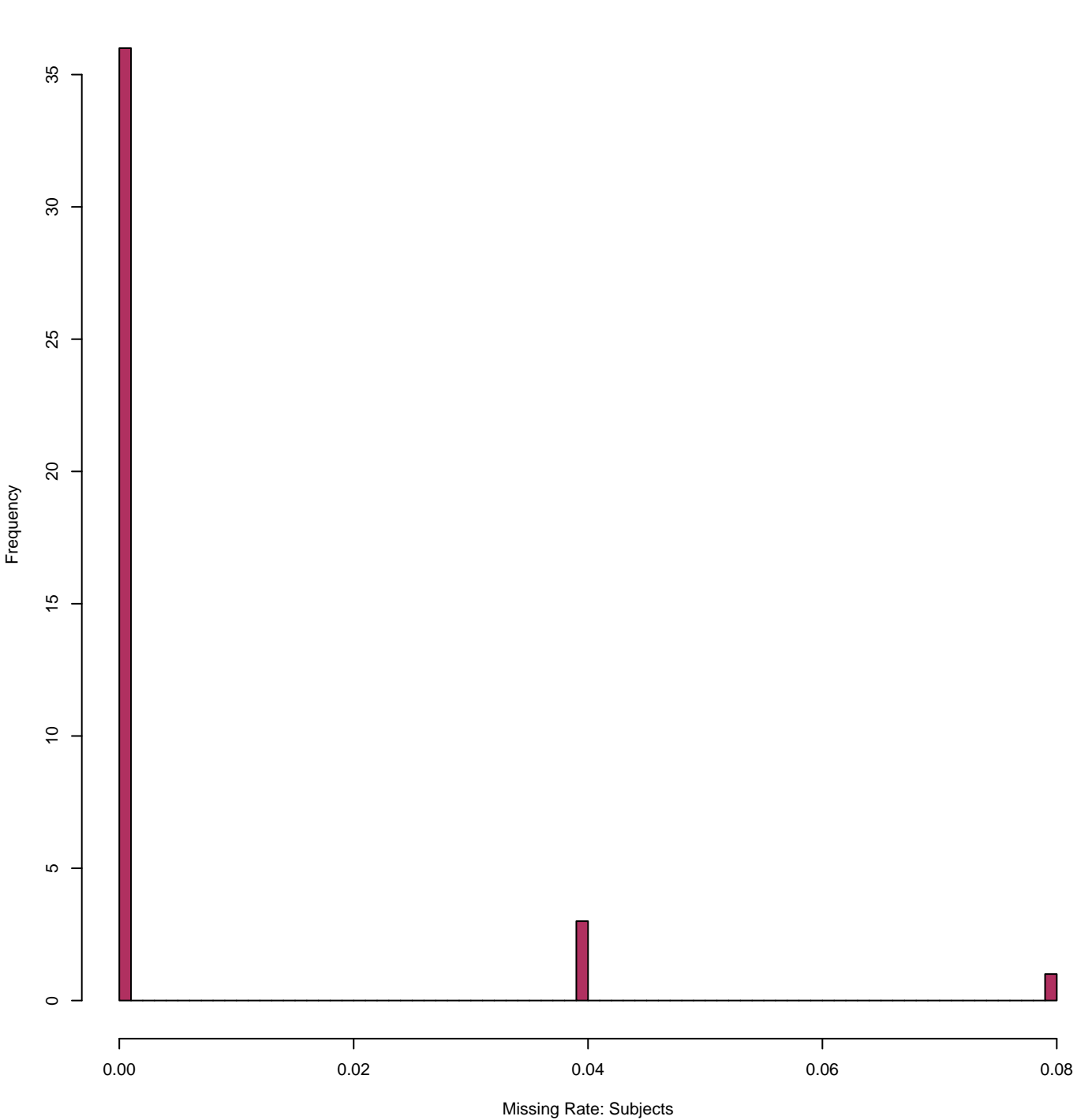
QC for tinywga – Marker MAF



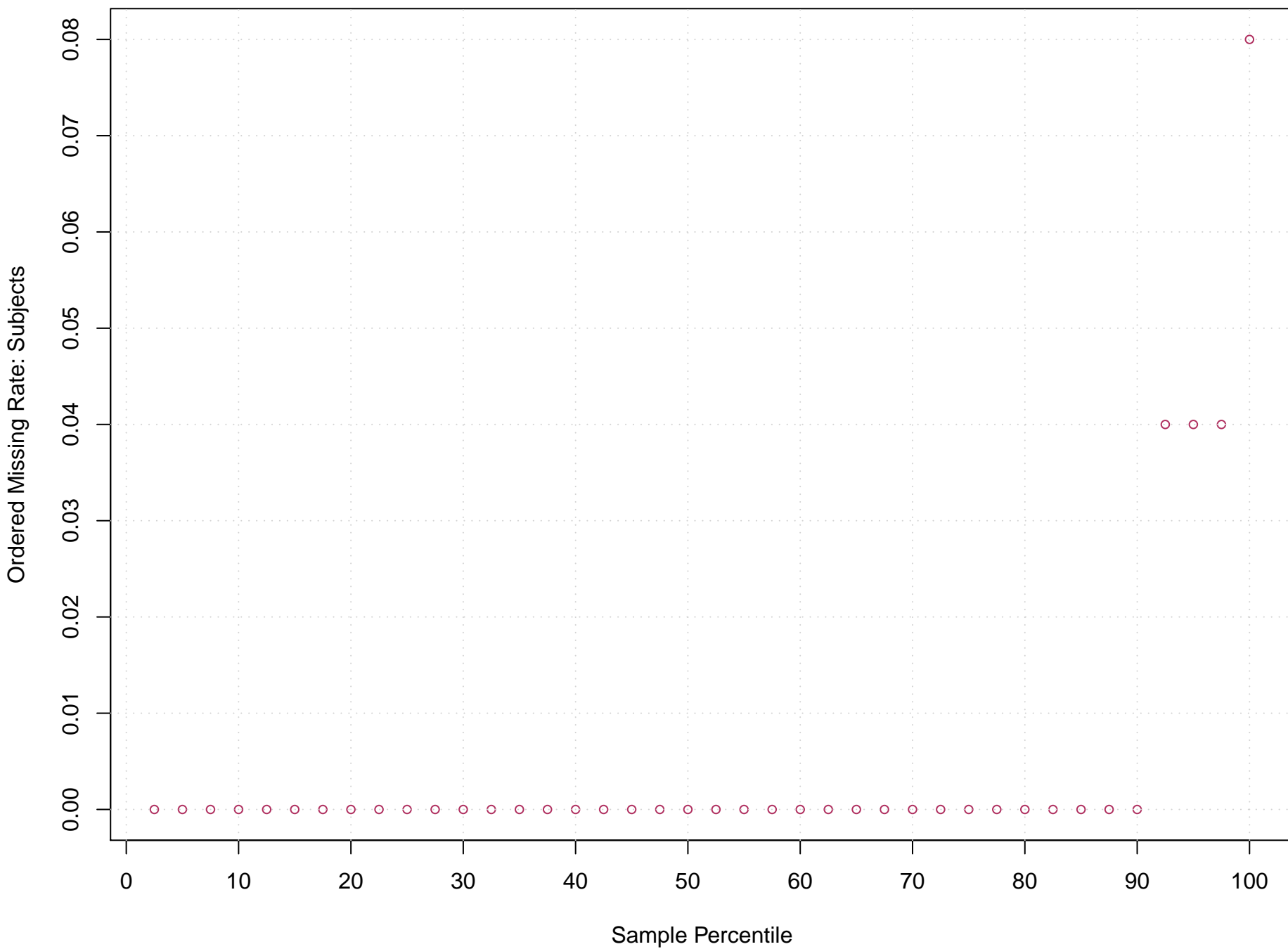
QC for tinywga – Ranked Marker MAF



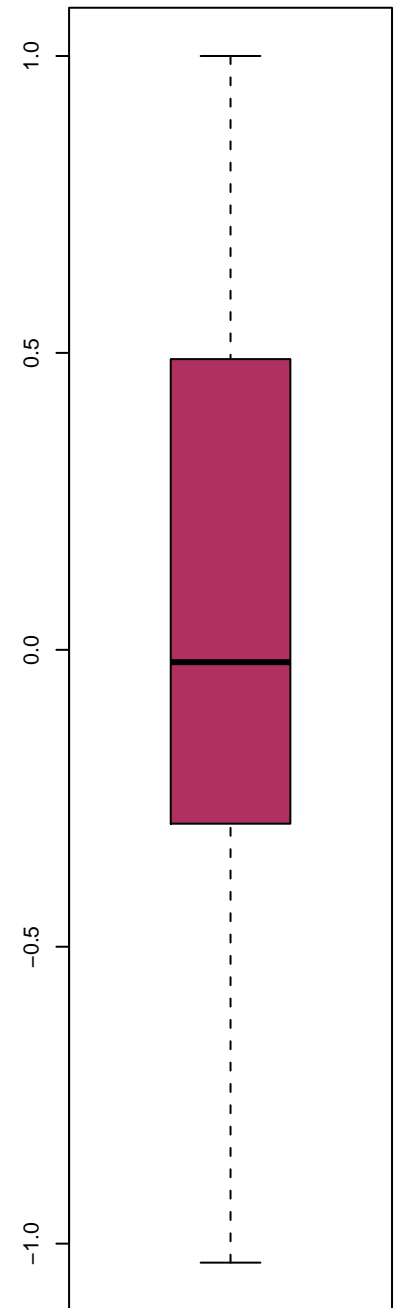
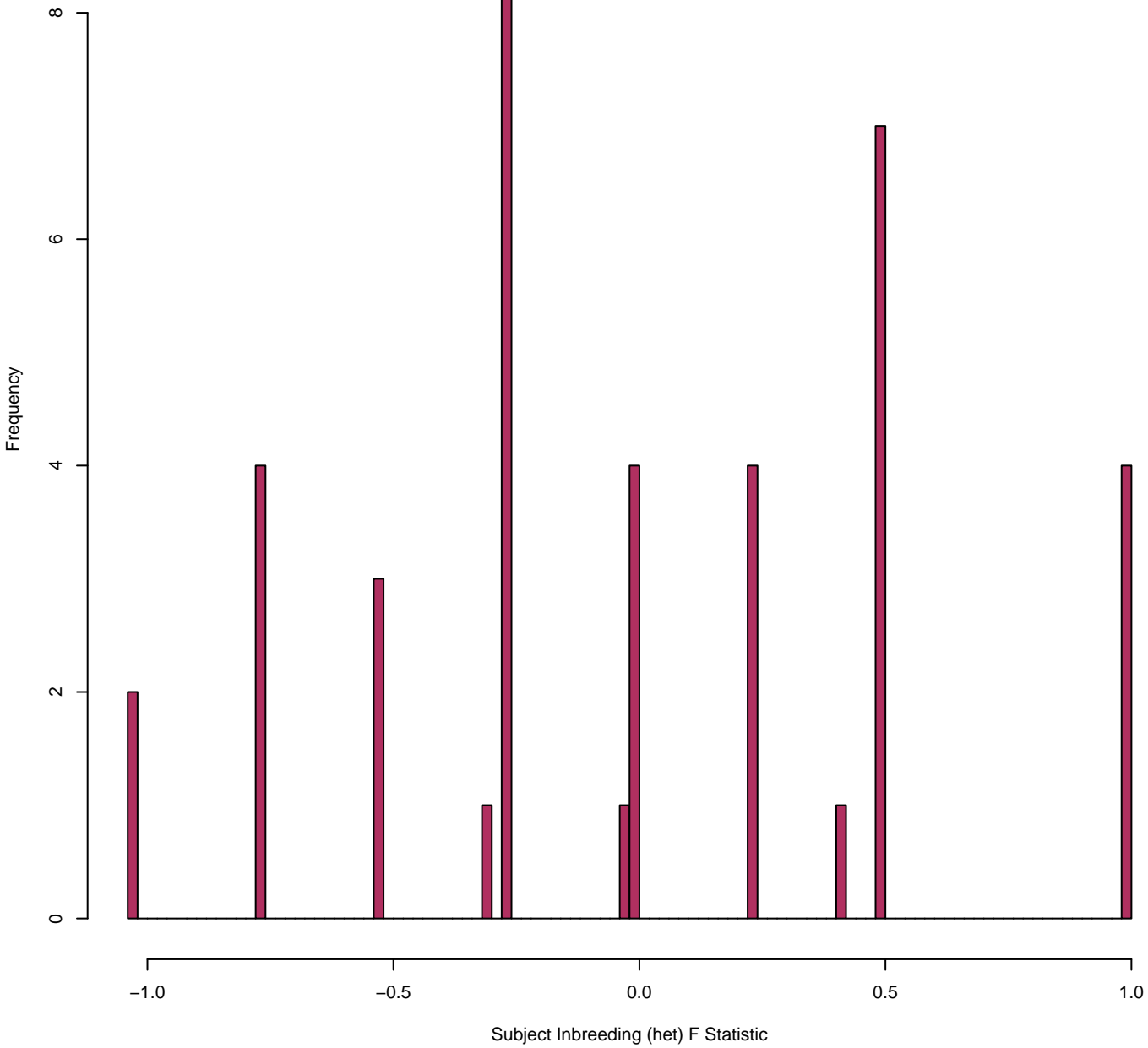
QC for tinywga – Subject Missing Genotype



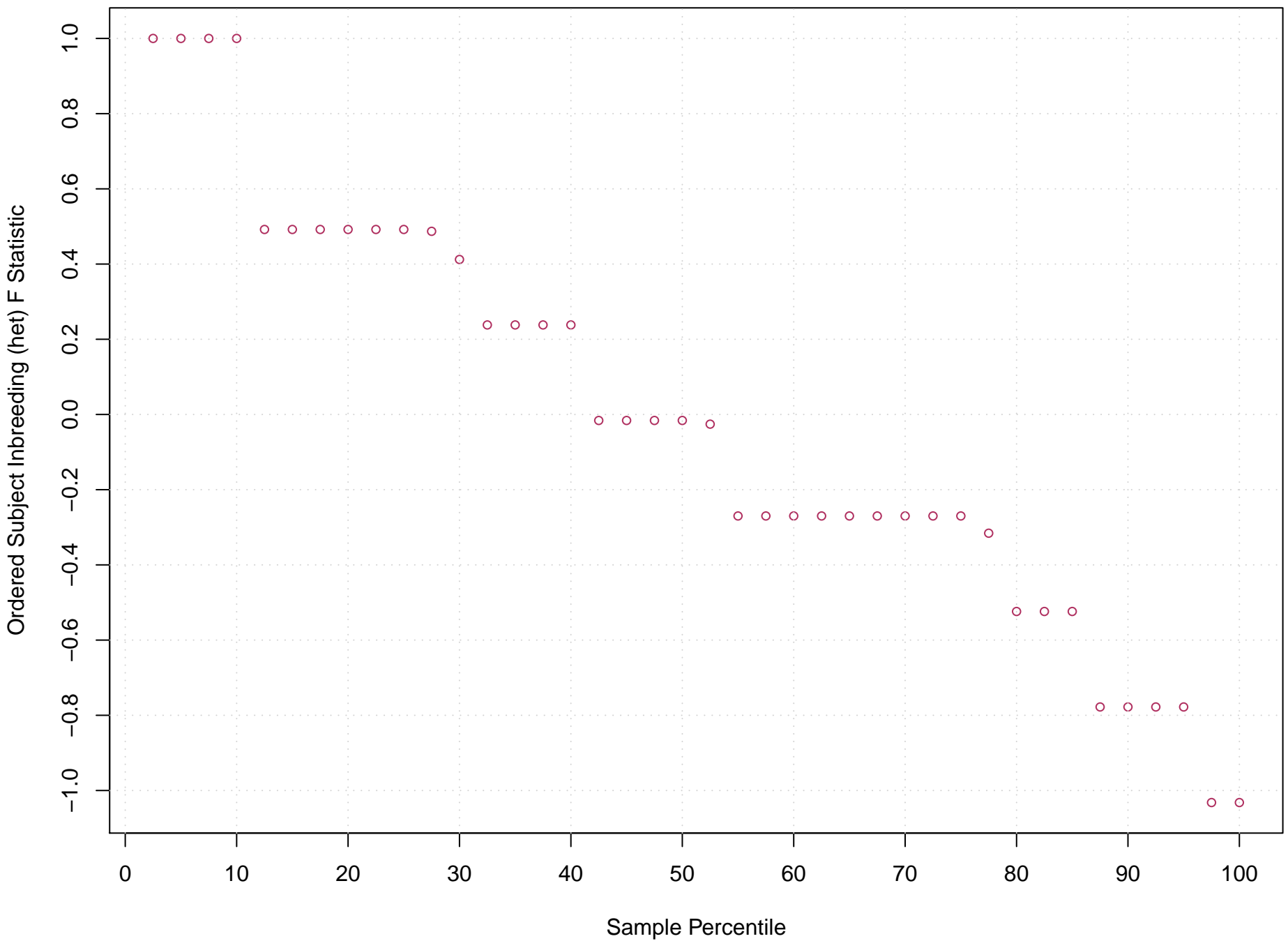
QC for tinywga – Ranked Subject Missing Genotype



QC for tinywga – Subject F Statistic



QC for tinywga – Ranked Subject F Statistic



Log QQ Plot(n=40)

