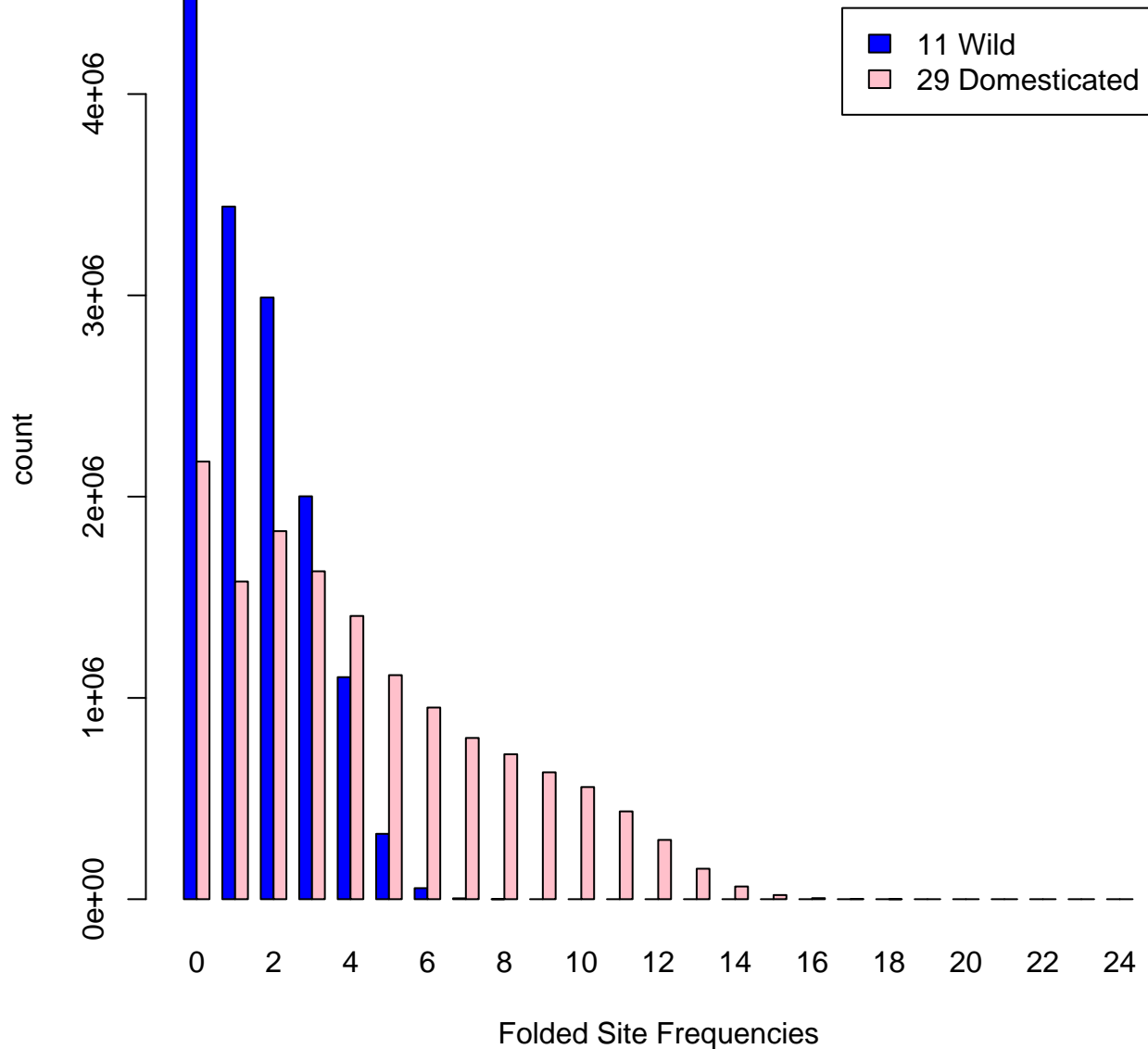
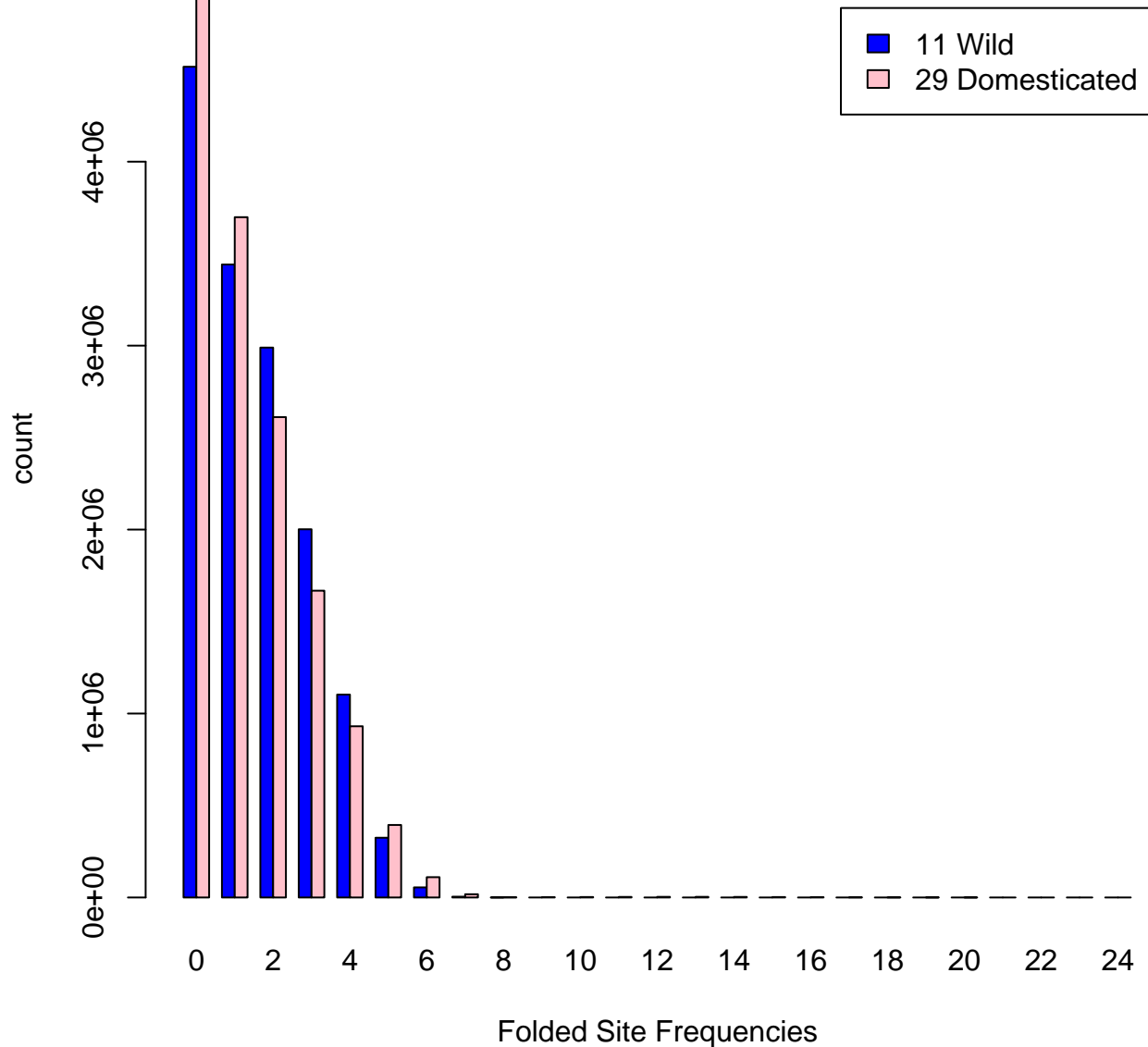


# Freq. on Autosomes + UN, with <95% missing data



# Freq. on Autosomes + UN (<95% MD), projected on 8 chr.



# Freq. on Autosomes + UN, with <95% missing data

