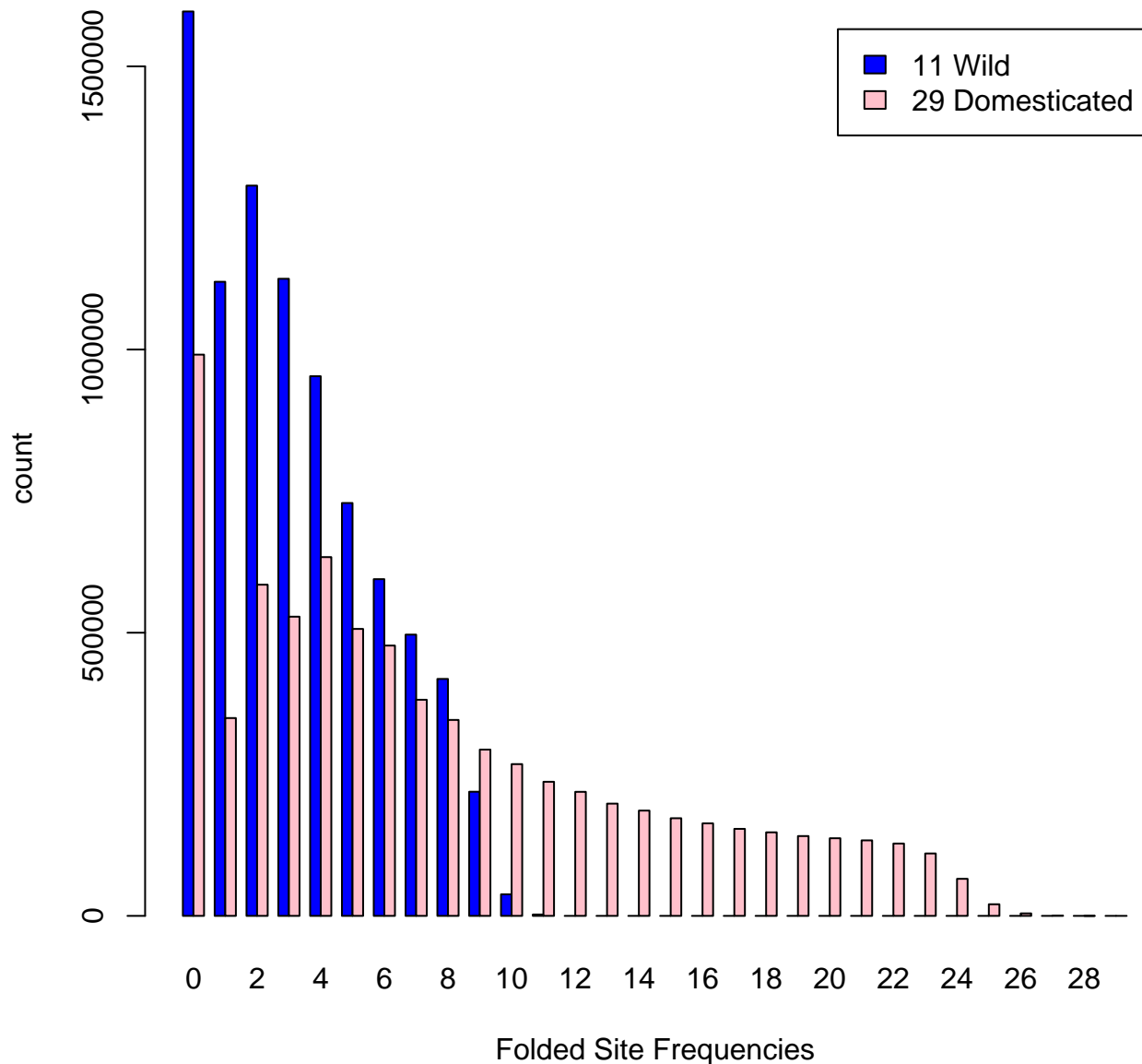
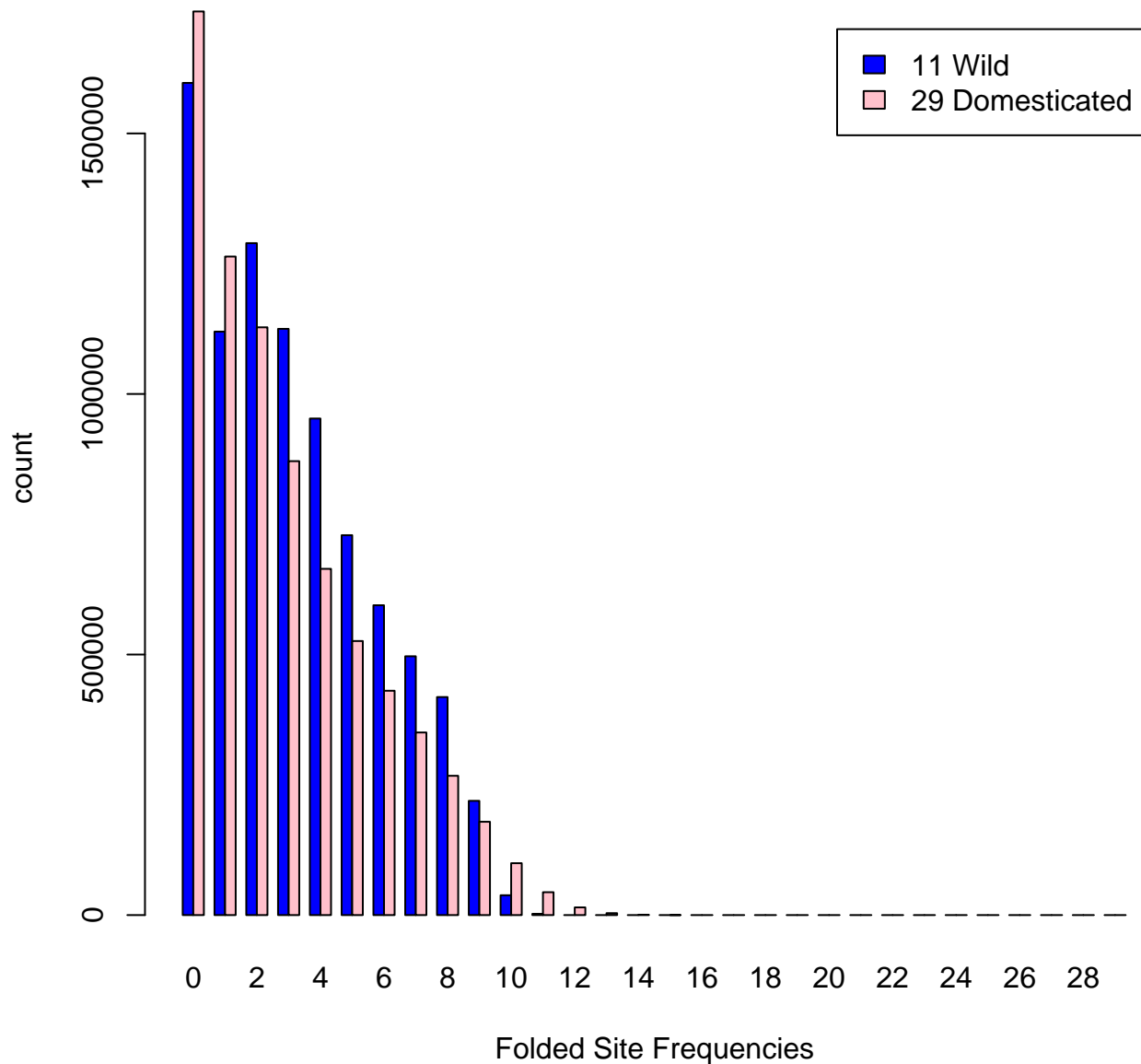


# Freq. on Autosomes, with <50% missing data



# Freq. on Autosomes (<50% MD), projected on 8 chr.



# Freq. on Autosomes, with <50% missing data

