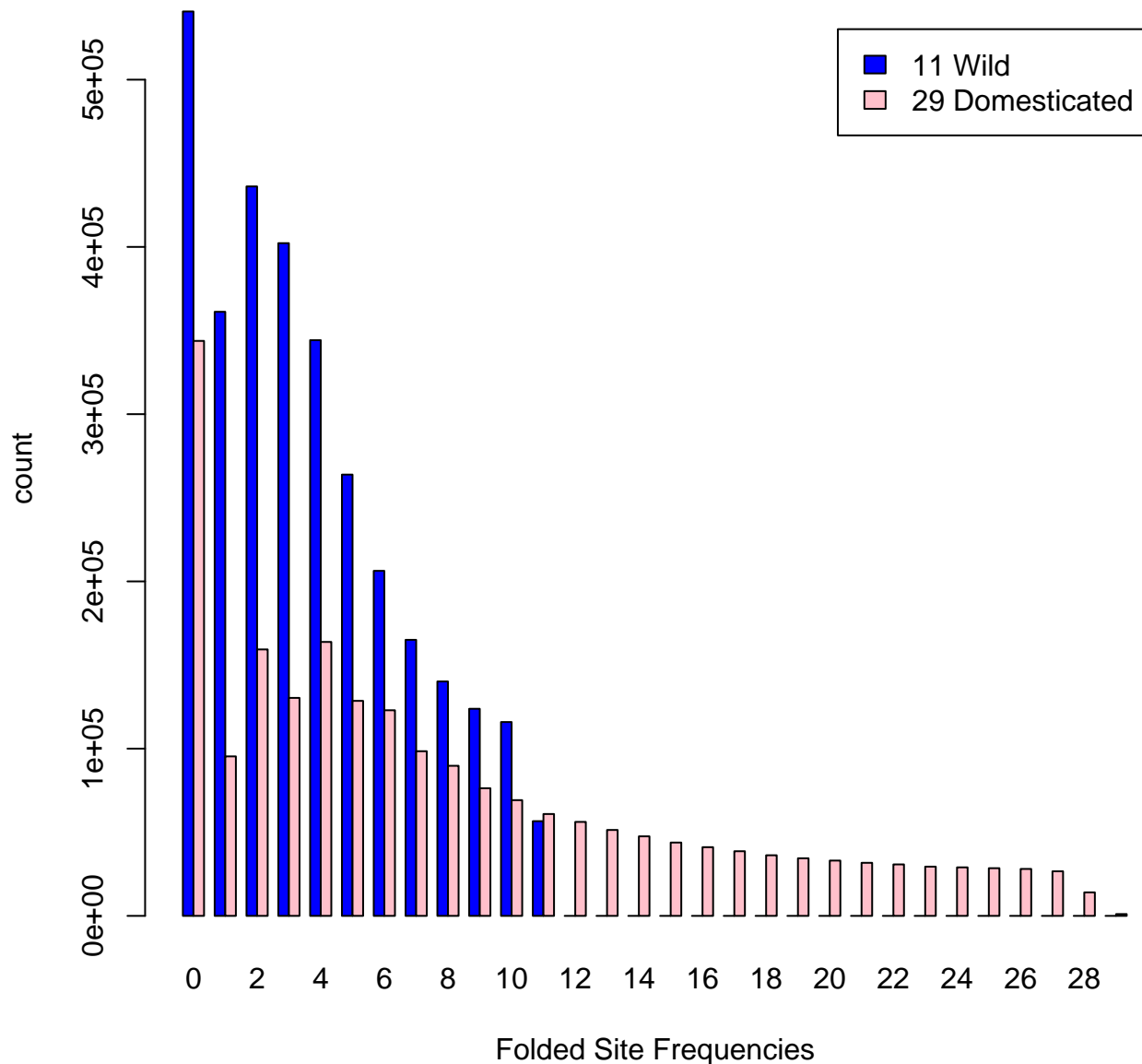
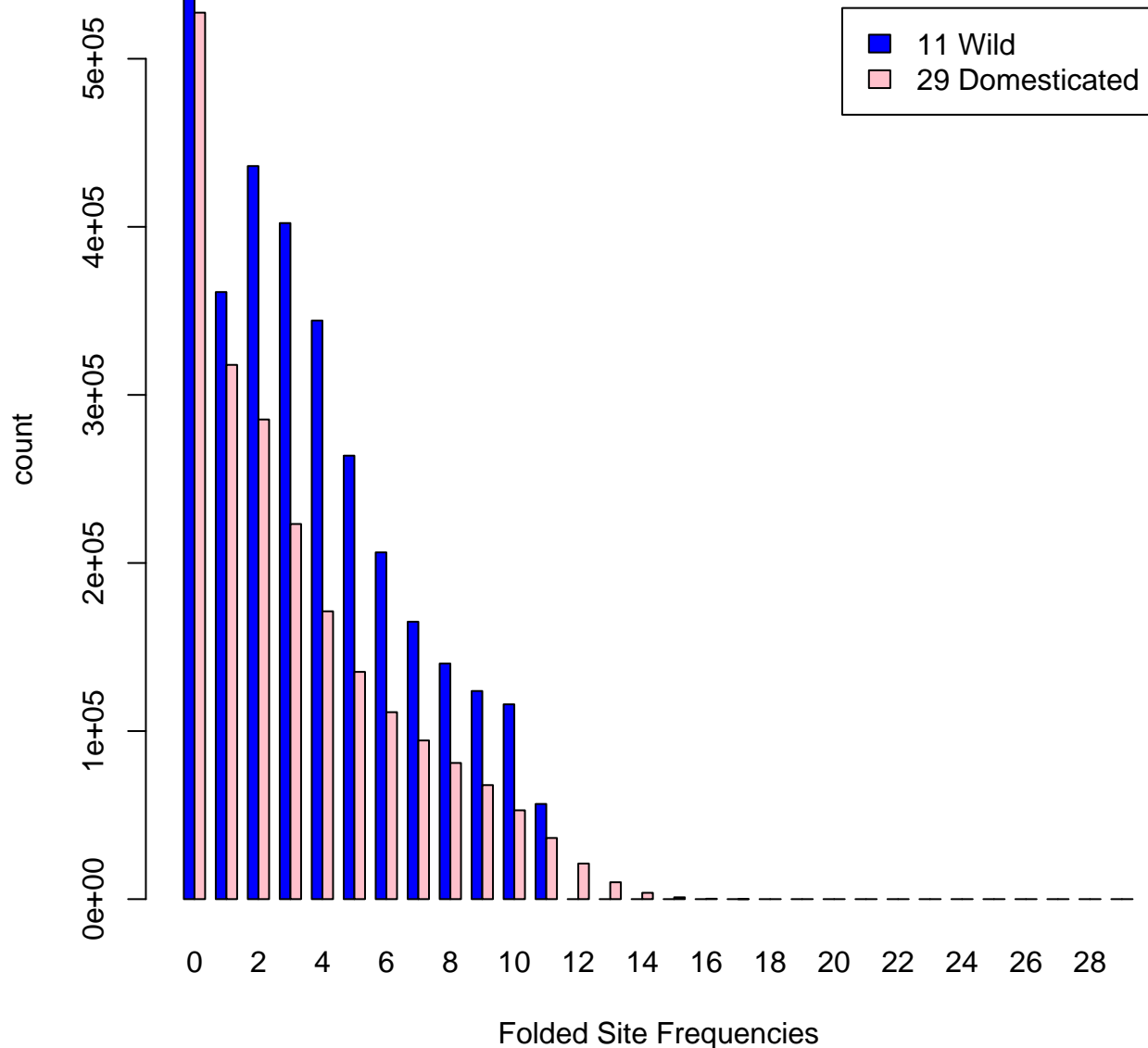


Freq. on Autosomes + UN, with <5% missing data



Freq. on Autosomes + UN (<5% MD), projected on 8 chr.



Freq. on Autosomes + UN, with <5% missing data

