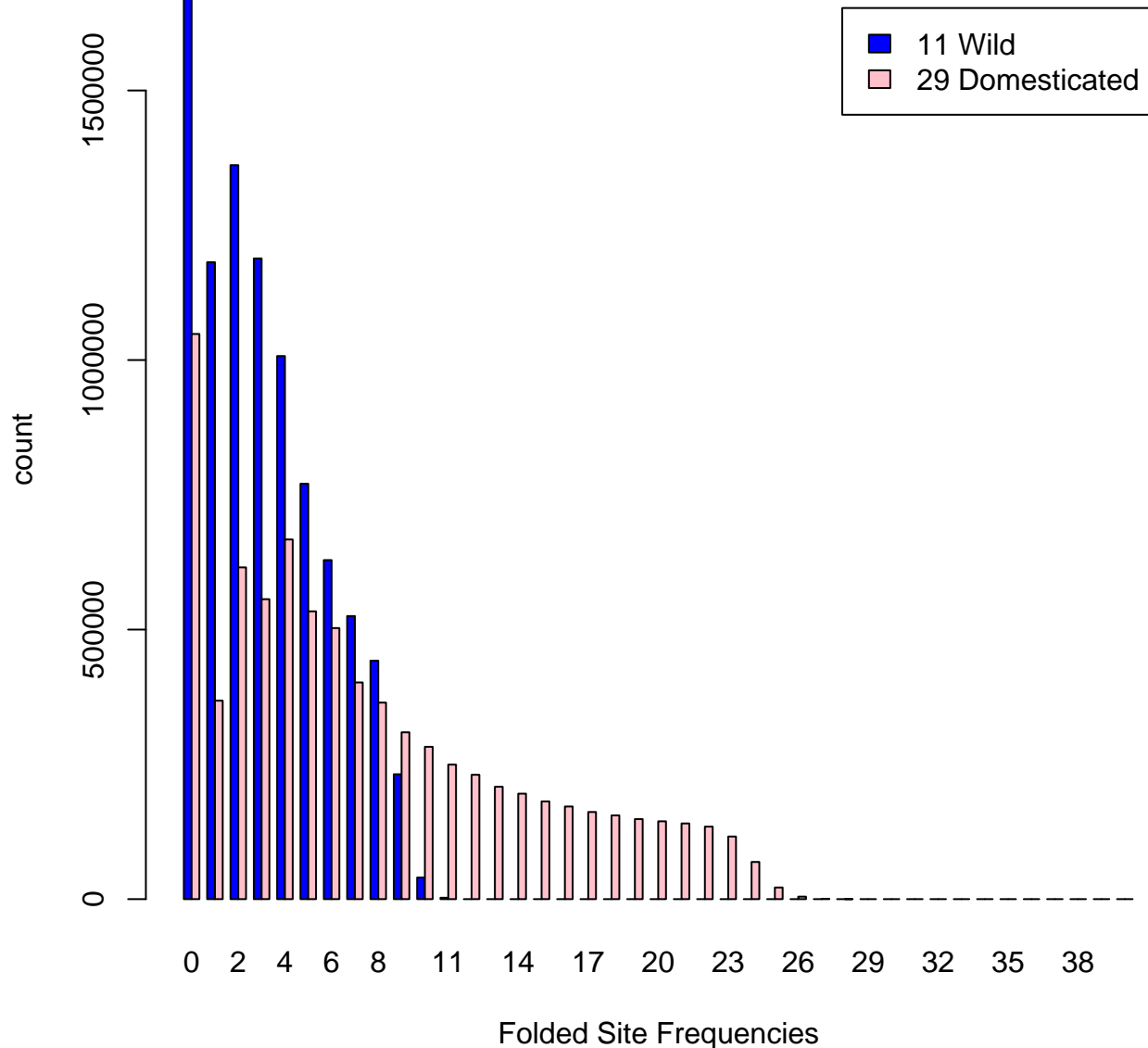
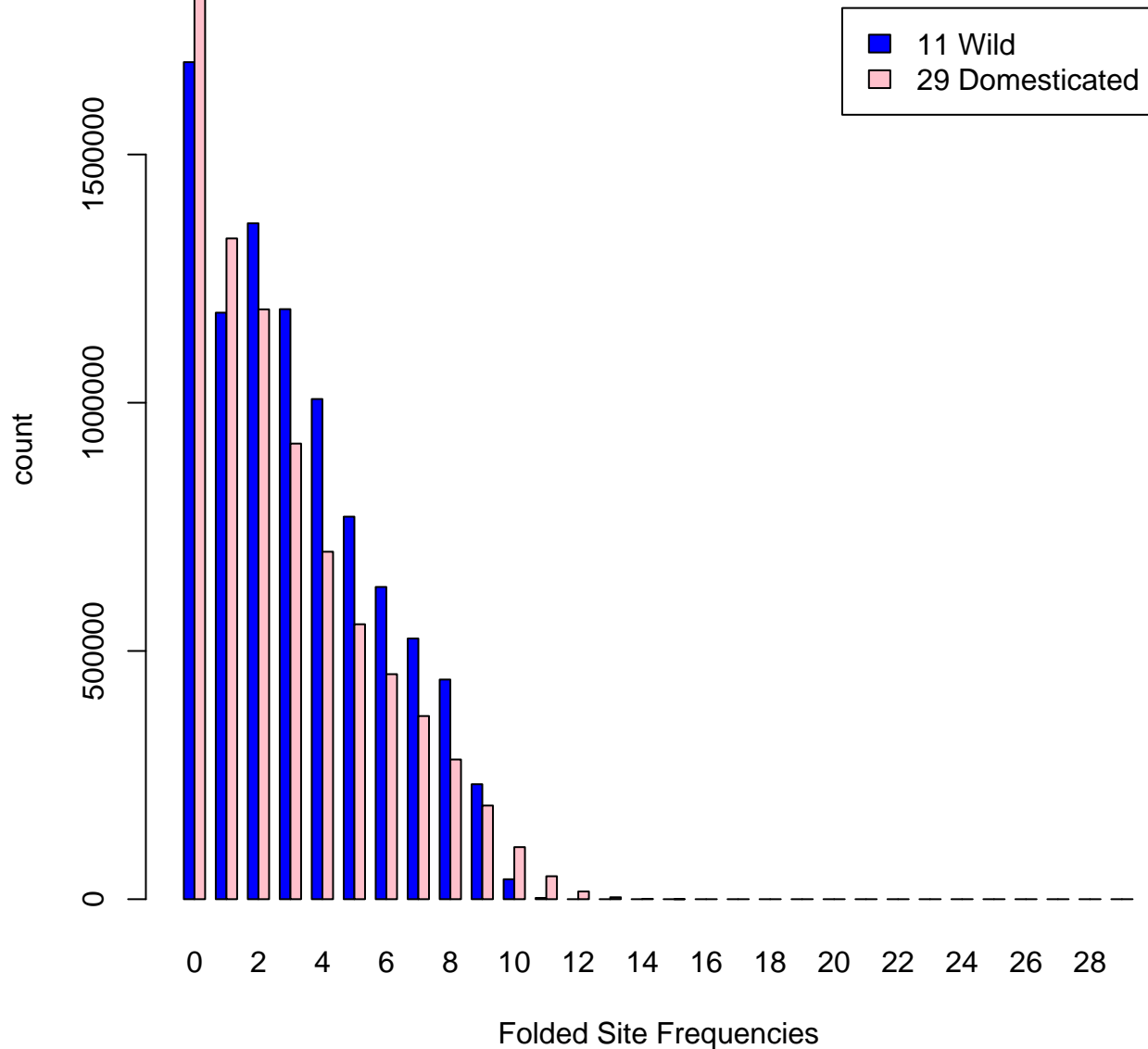


Freq. on Autosomes + UN, with <50% missing data



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



Freq. on Autosomes + UN, with <50% missing data

