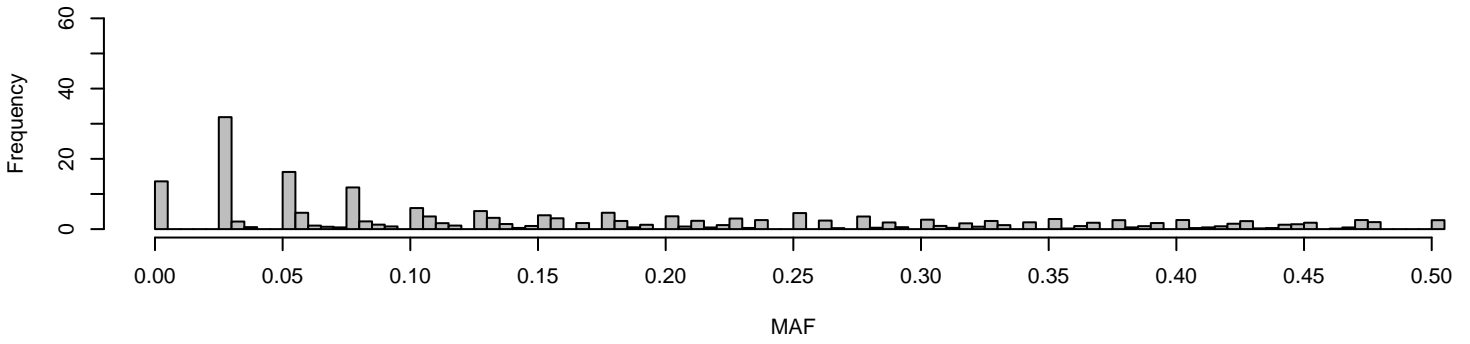
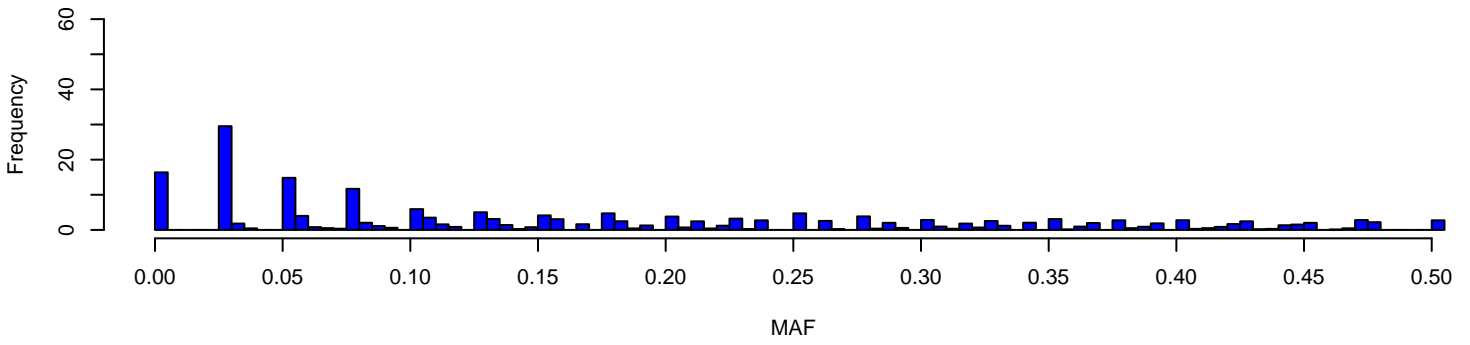


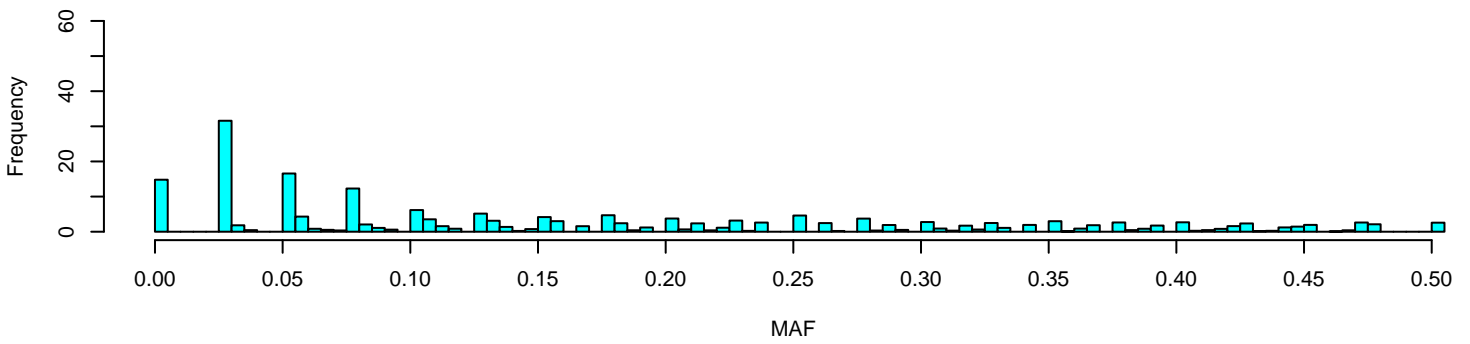
**Probability of MAFs for all SNPs on Chr 3
Filtered domesticus**



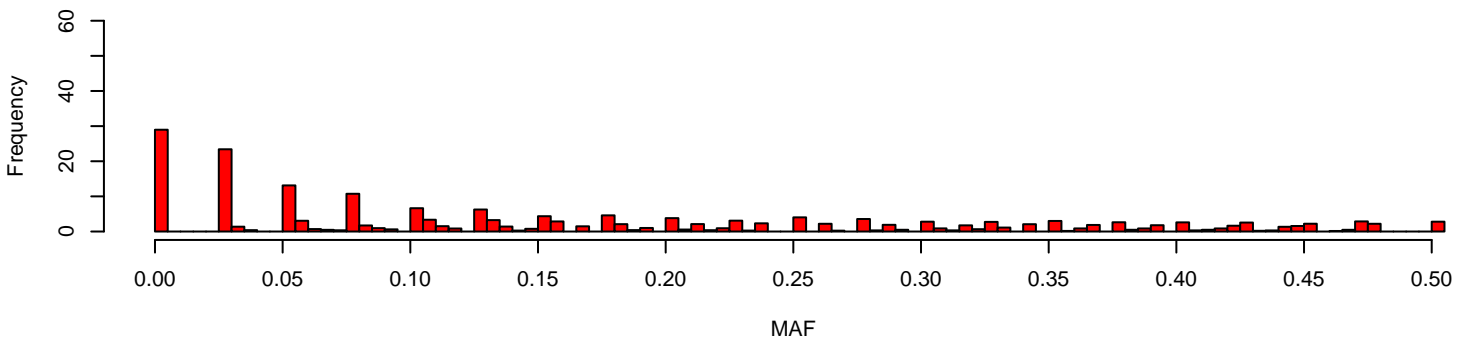
17 strains VQSR domesticus



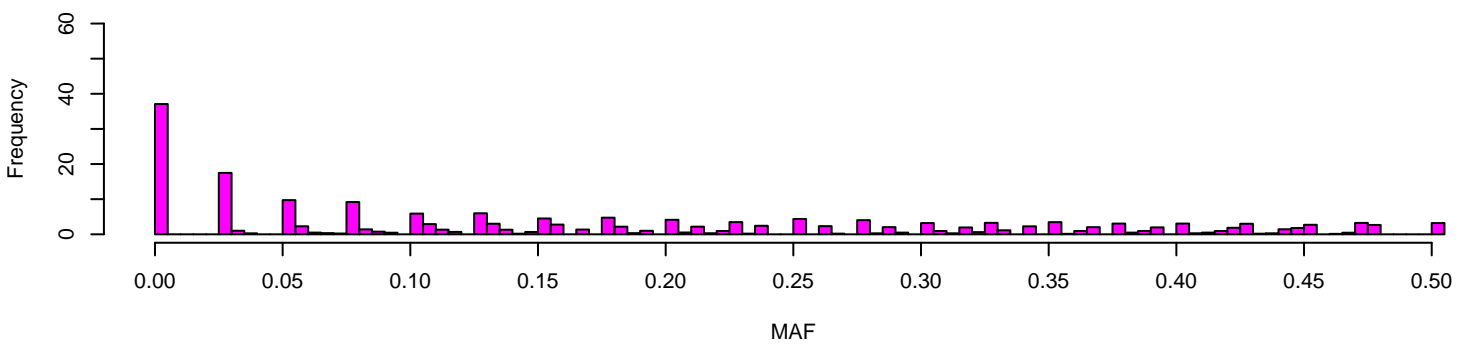
Array VQSR domesticus



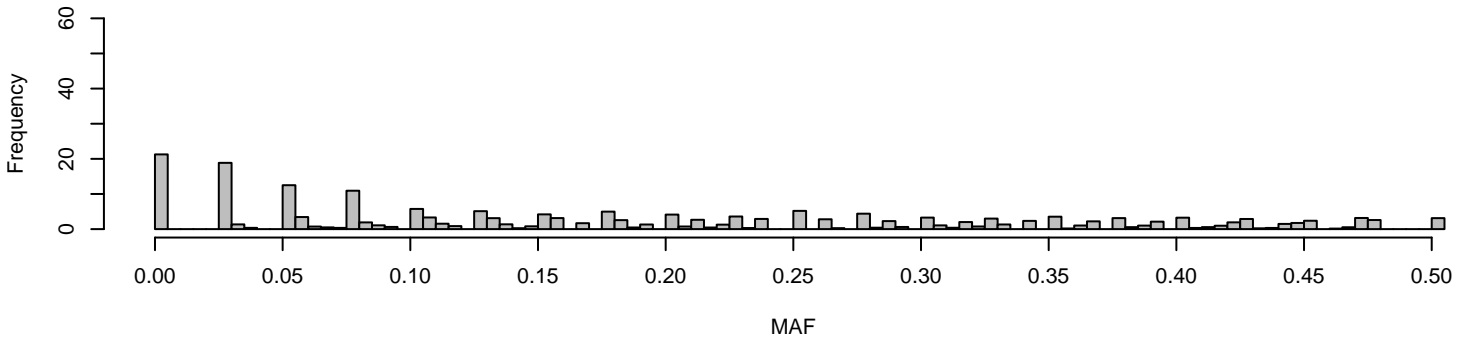
Filtered castaneus



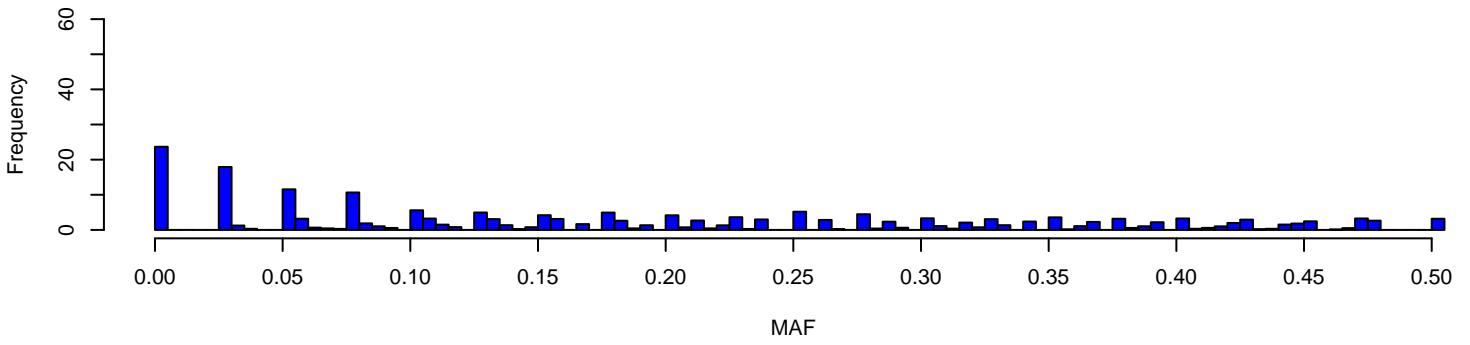
17 strains VQSR castaneus



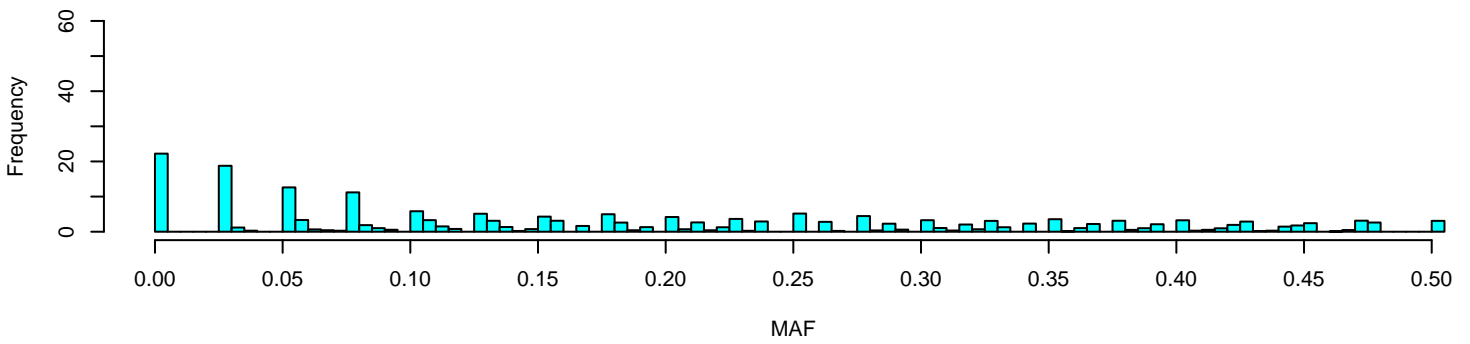
**Probability of MAFs for known SNPs on Chr 3
Filtered domesticus**



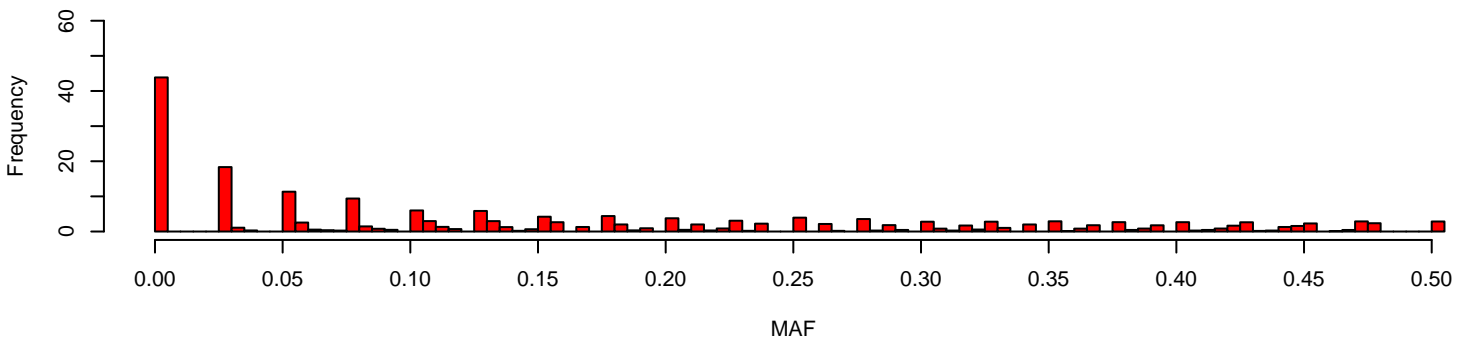
17 strains VQSR domesticus



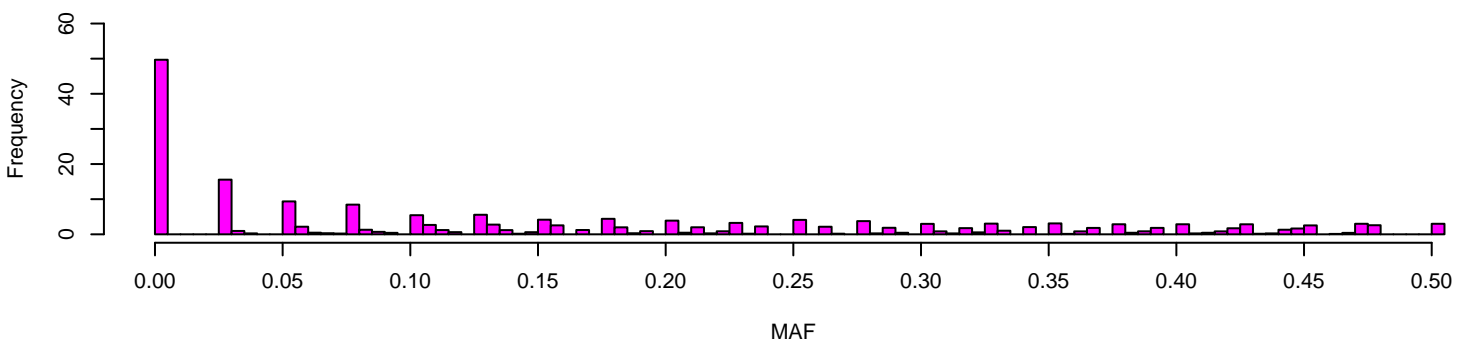
Array VQSR domesticus



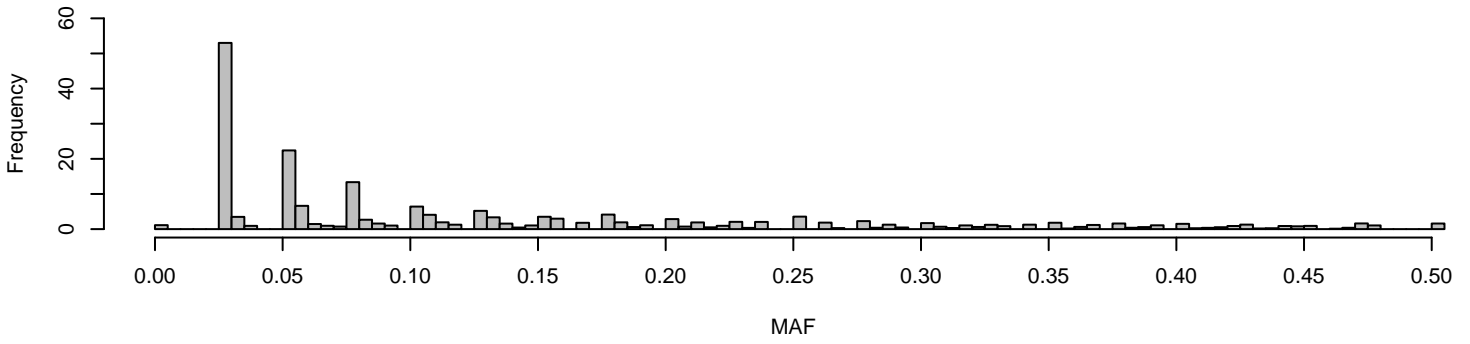
Filtered castaneus



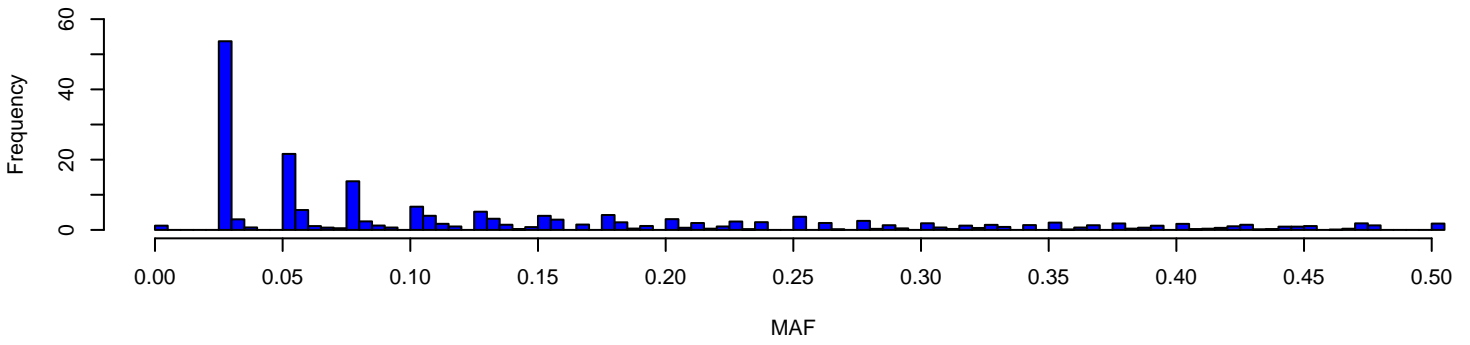
17 strains VQSR castaneus



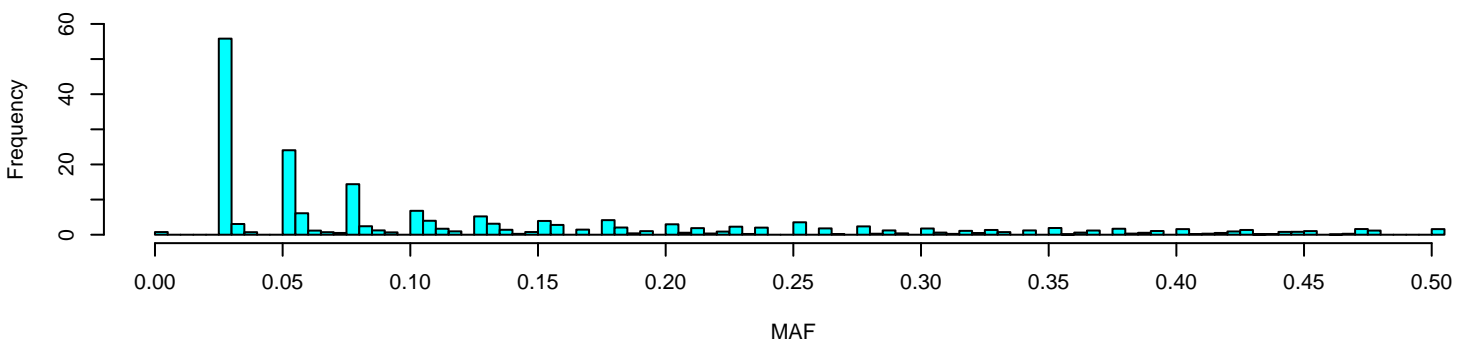
**Probability of MAFs for new SNPs on Chr 3
Filtered domesticus**



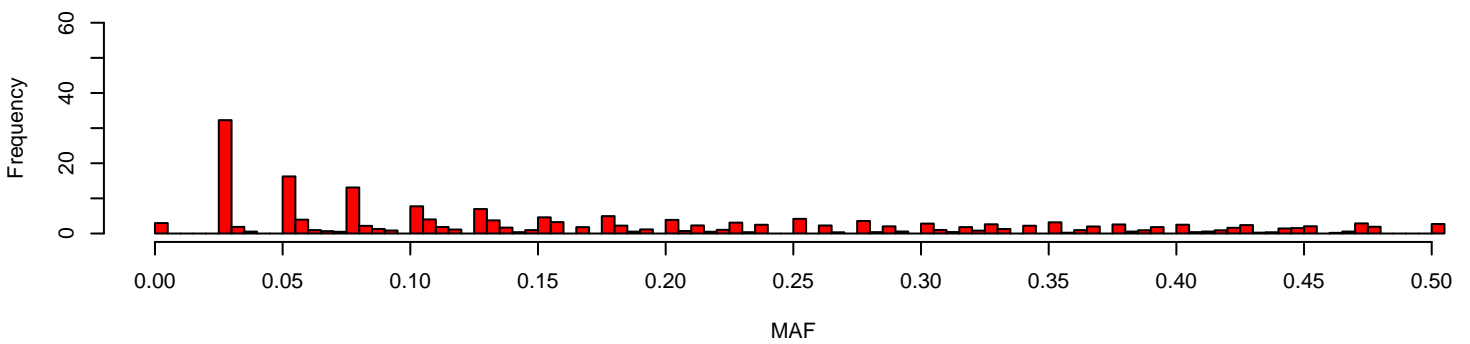
17 strains VQSR domesticus



Array VQSR domesticus



Filtered castaneus



17 strains VQSR castaneus

