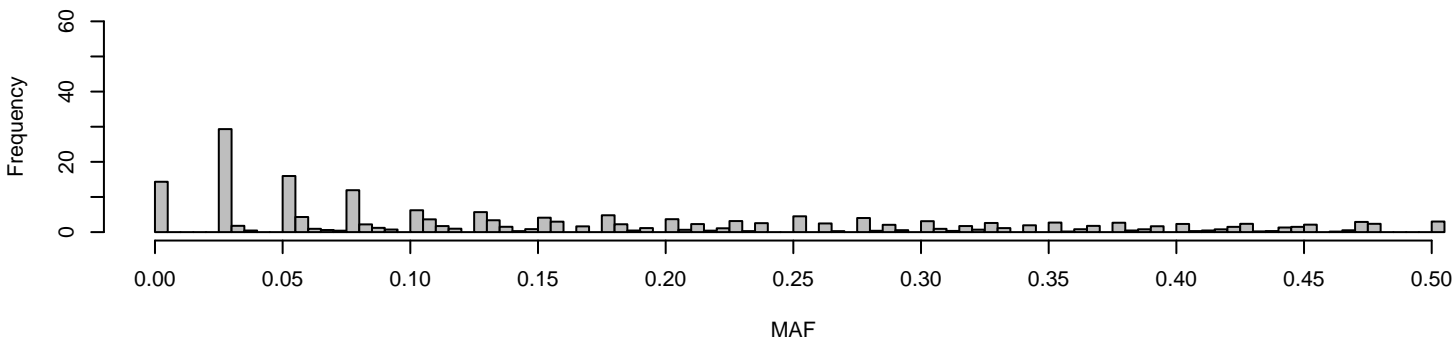
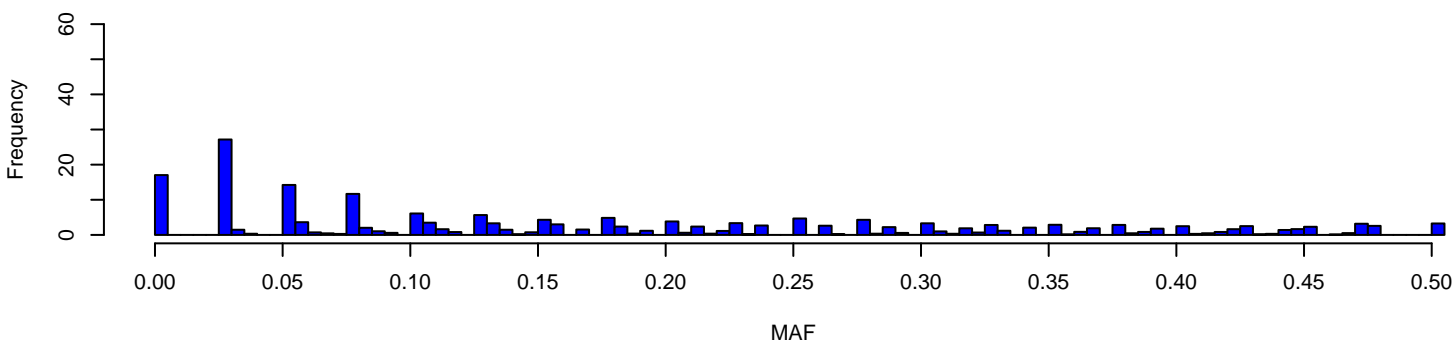


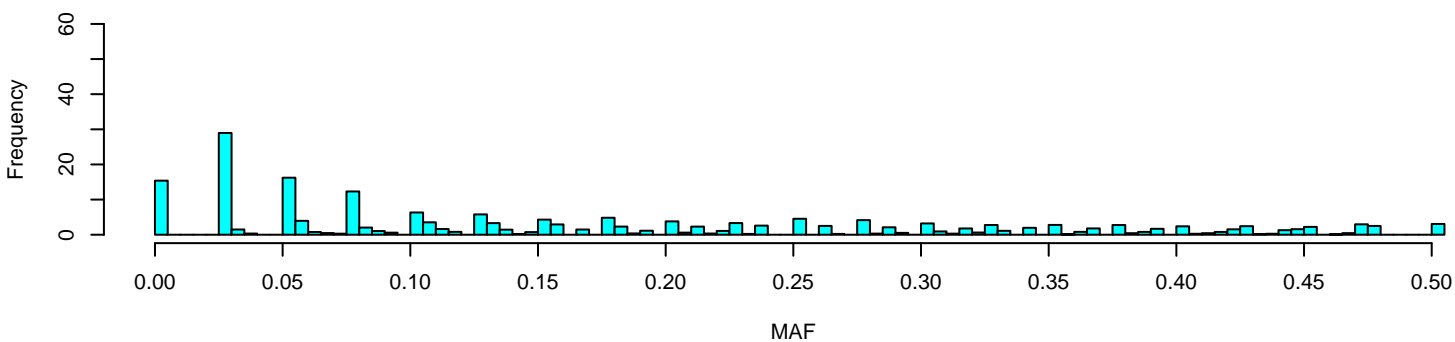
**Probability of MAFs for all SNPs on Chr 2  
Filtered domesticus**



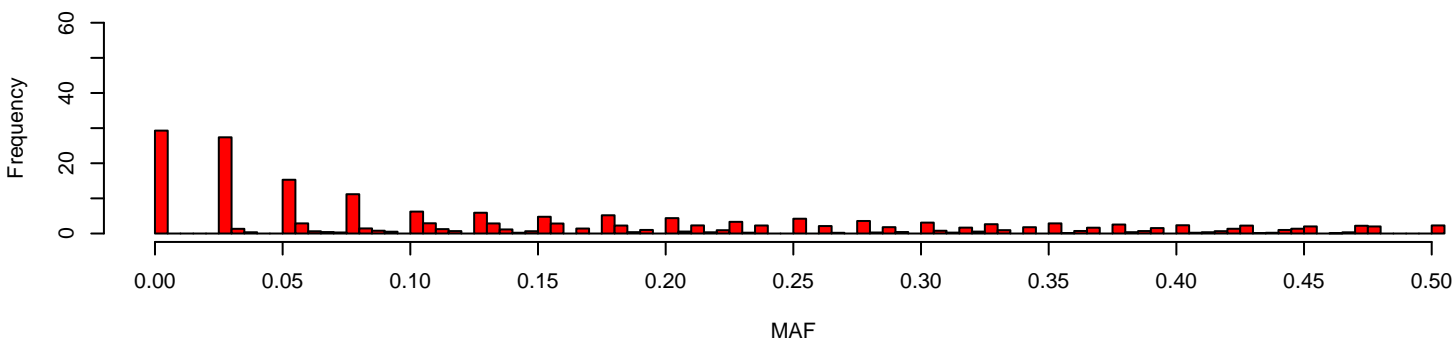
**17 strains VQSR domesticus**



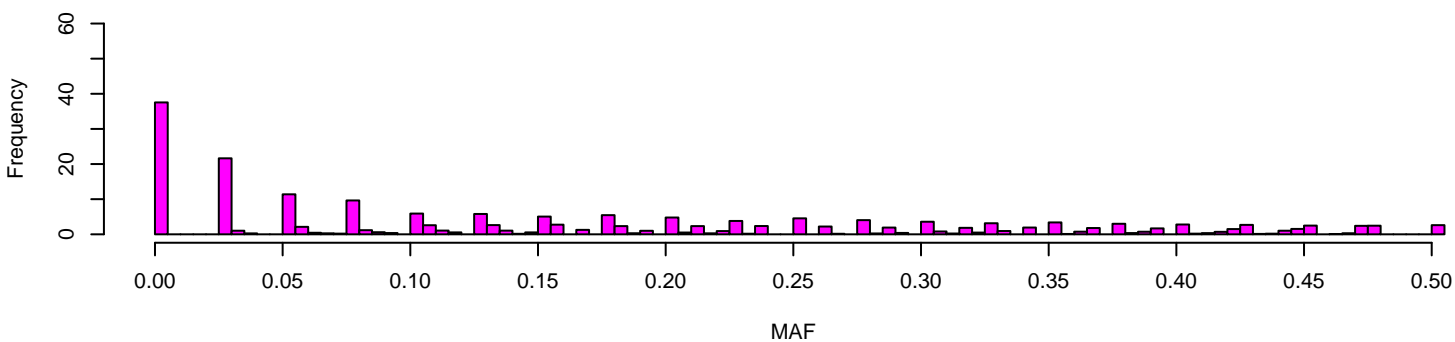
**Array VQSR domesticus**



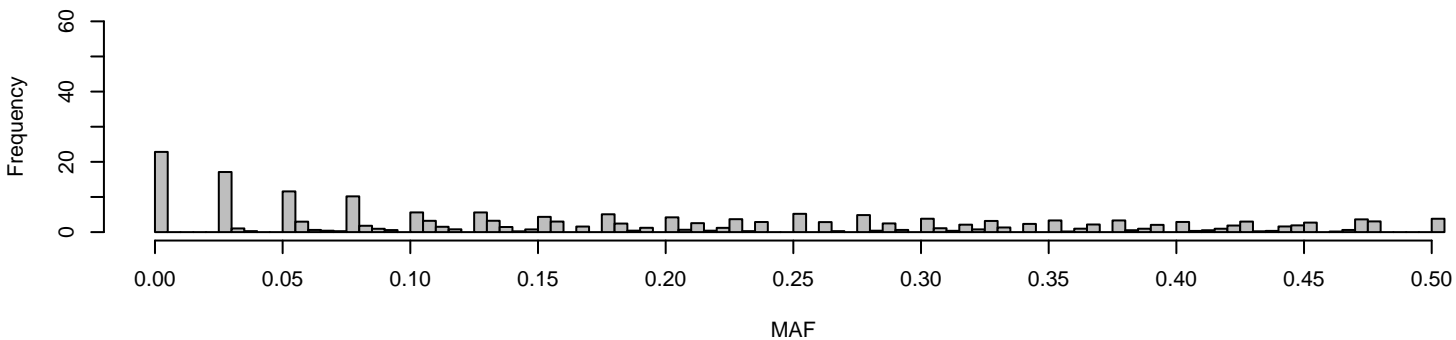
**Filtered castaneus**



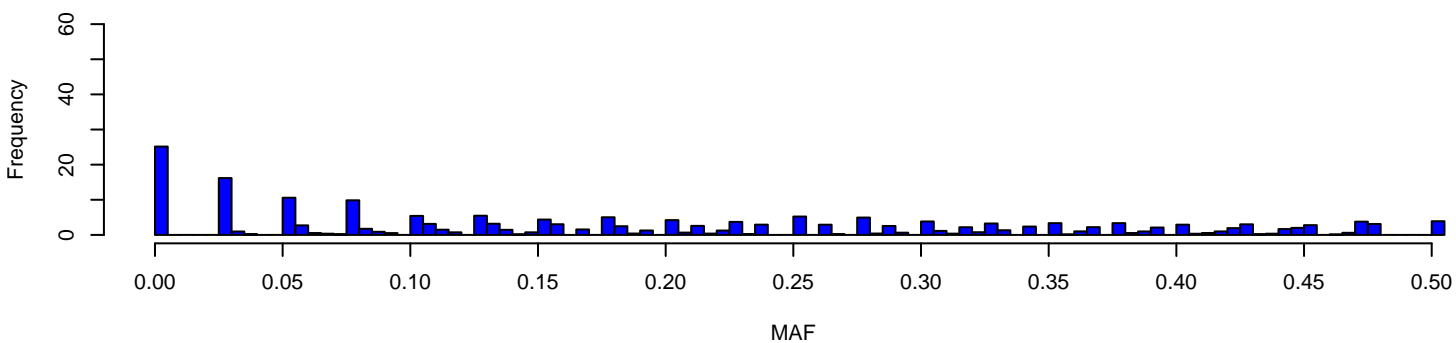
**17 strains VQSR castaneus**



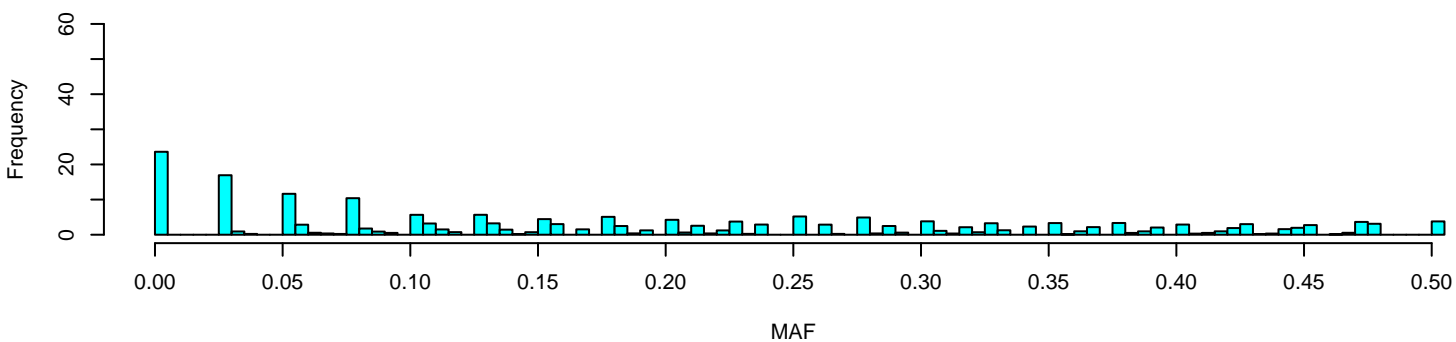
**Probability of MAFs for known SNPs on Chr 2**  
**Filtered domesticus**



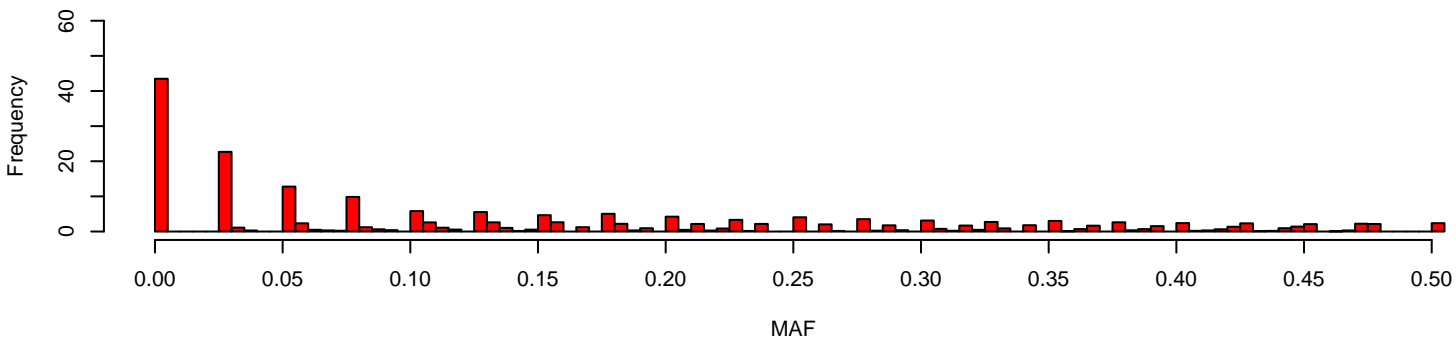
**17 strains VQSR domesticus**



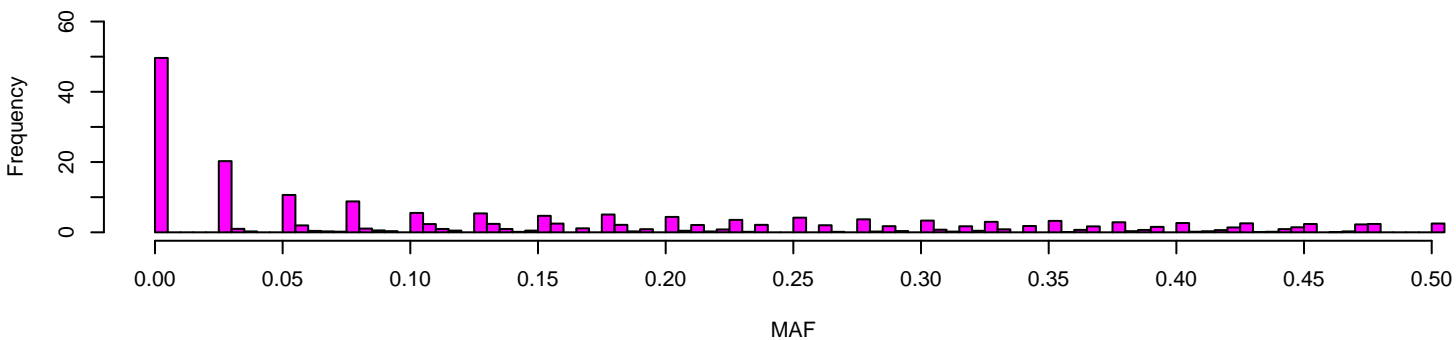
**Array VQSR domesticus**



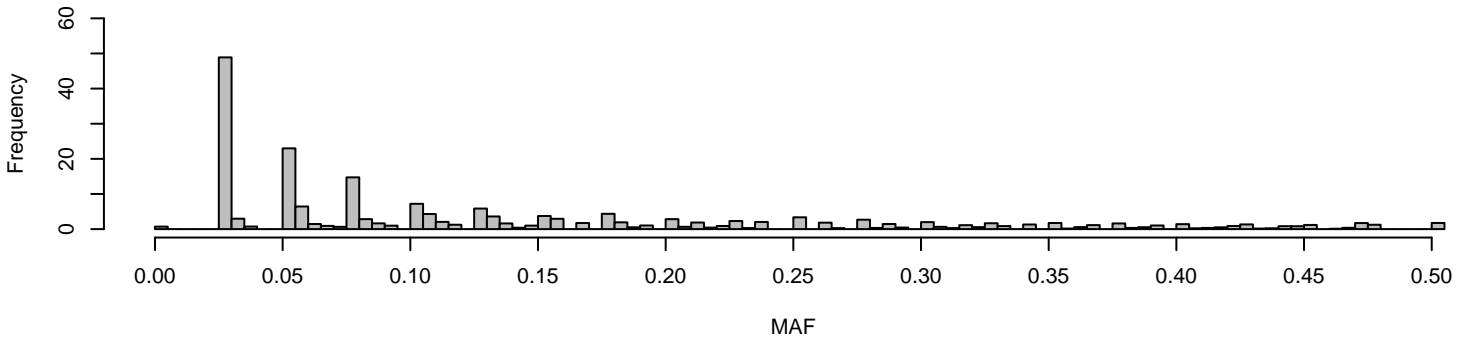
**Filtered castaneus**



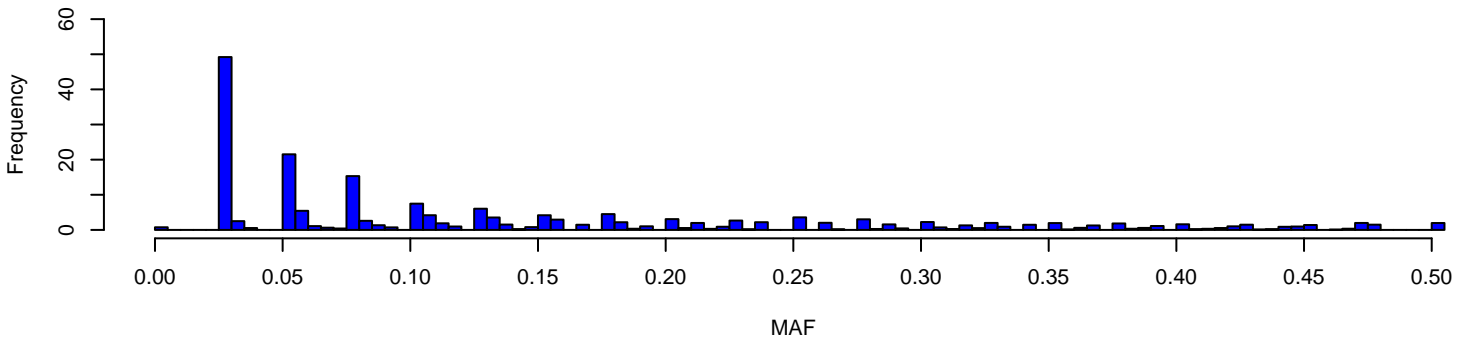
**17 strains VQSR castaneus**



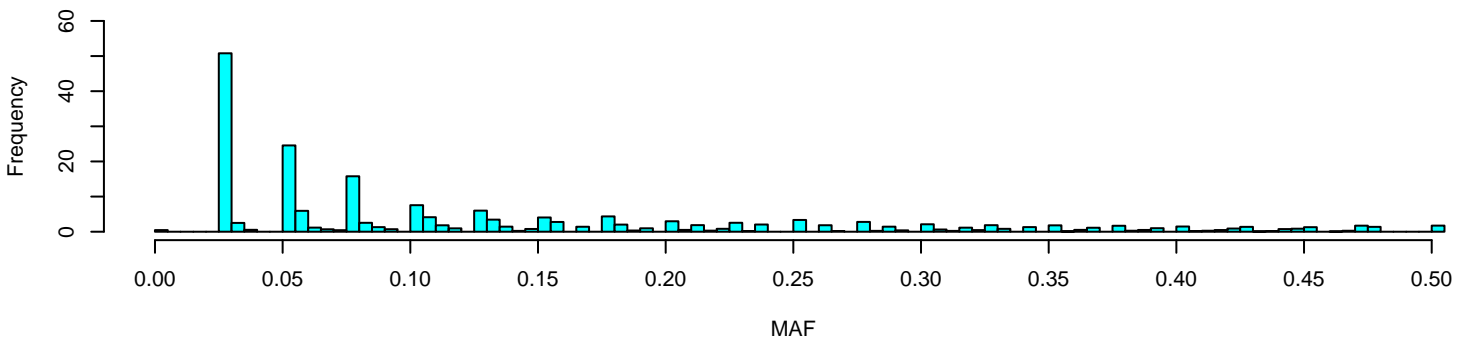
**Probability of MAFs for new SNPs on Chr 2  
Filtered domesticus**



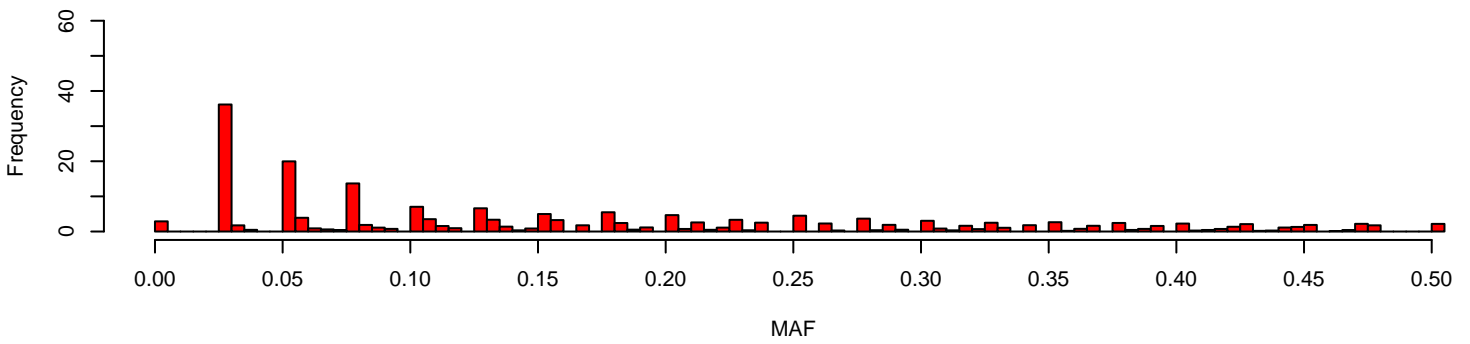
**17 strains VQSR domesticus**



**Array VQSR domesticus**



**Filtered castaneus**



**17 strains VQSR castaneus**

