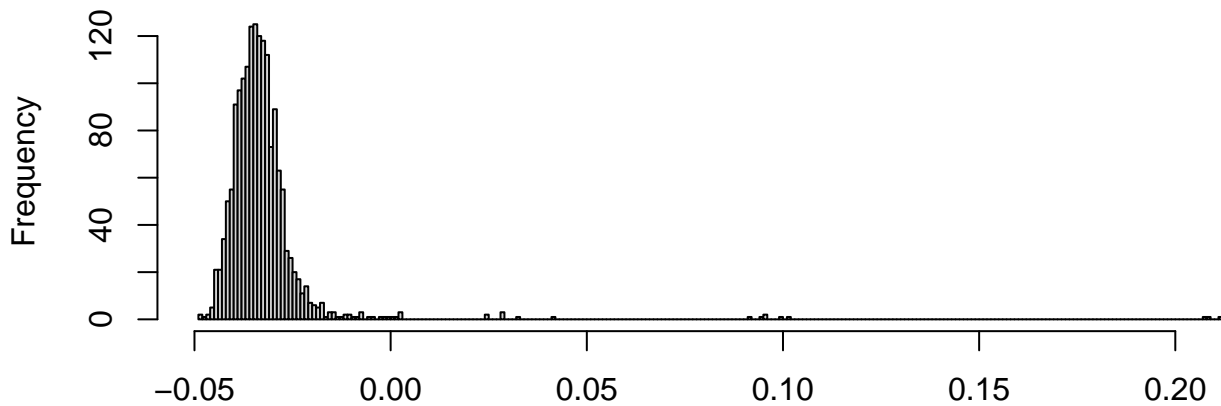


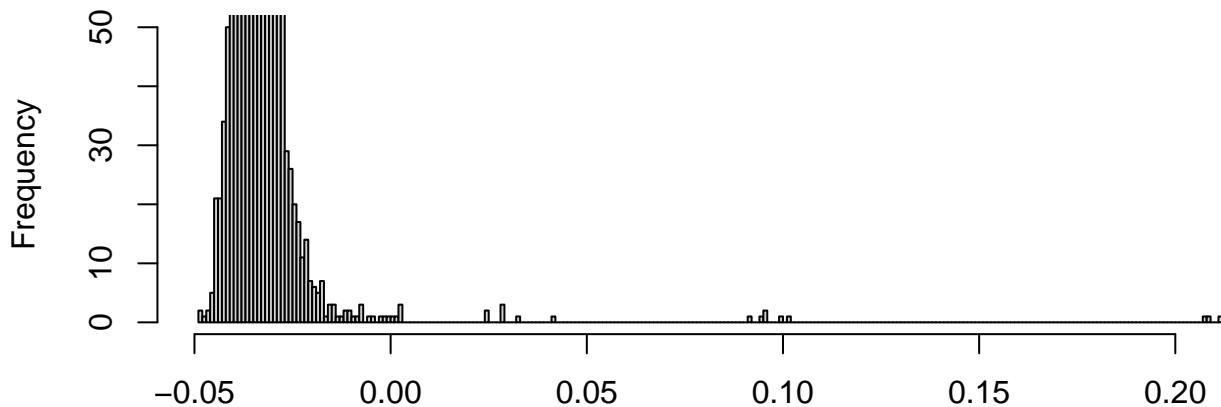
## HSP vs UP PLODs within ARG



PLOD

ale\_freqs\_below\_0.8 → 4594 loci

## Uncommon values suggest likely close-kin



PLOD