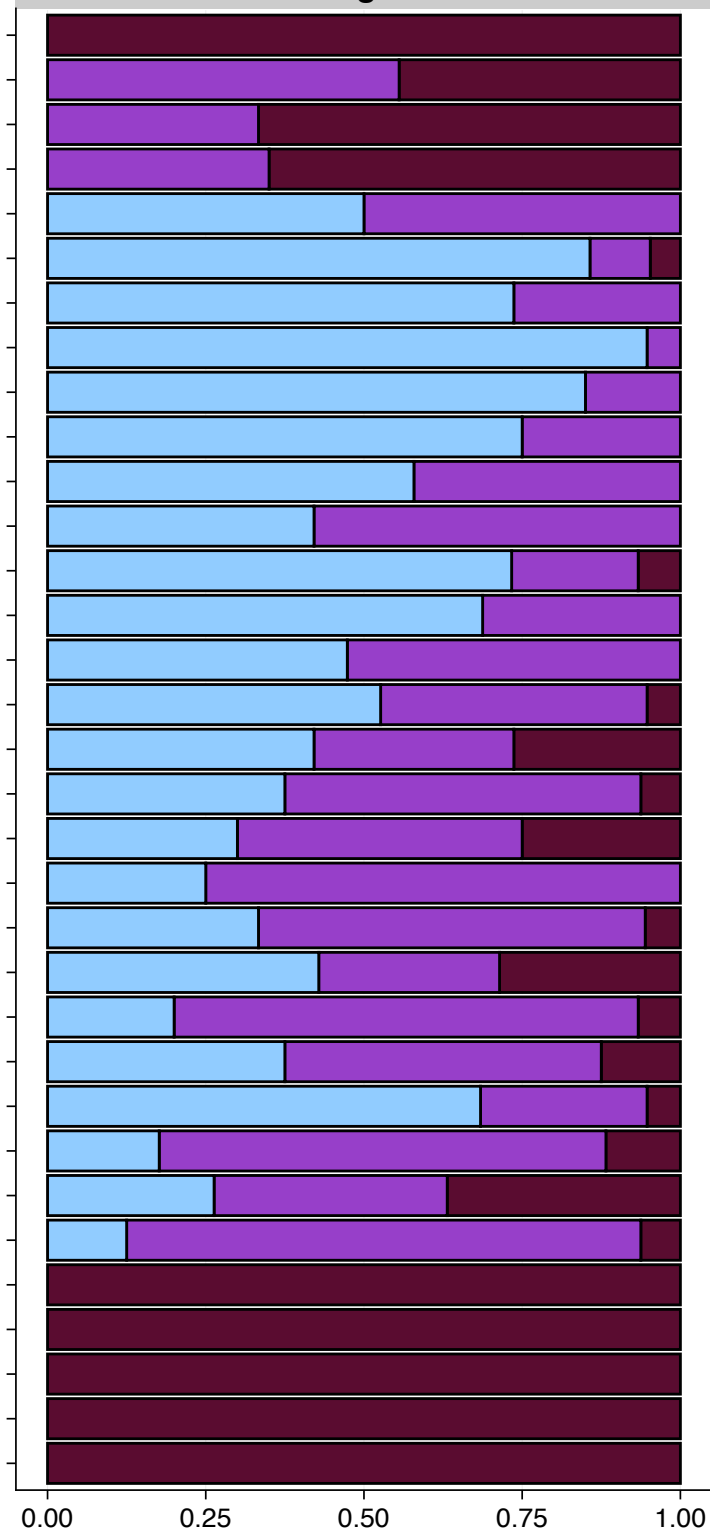


Haplotypes for putative inversion regions

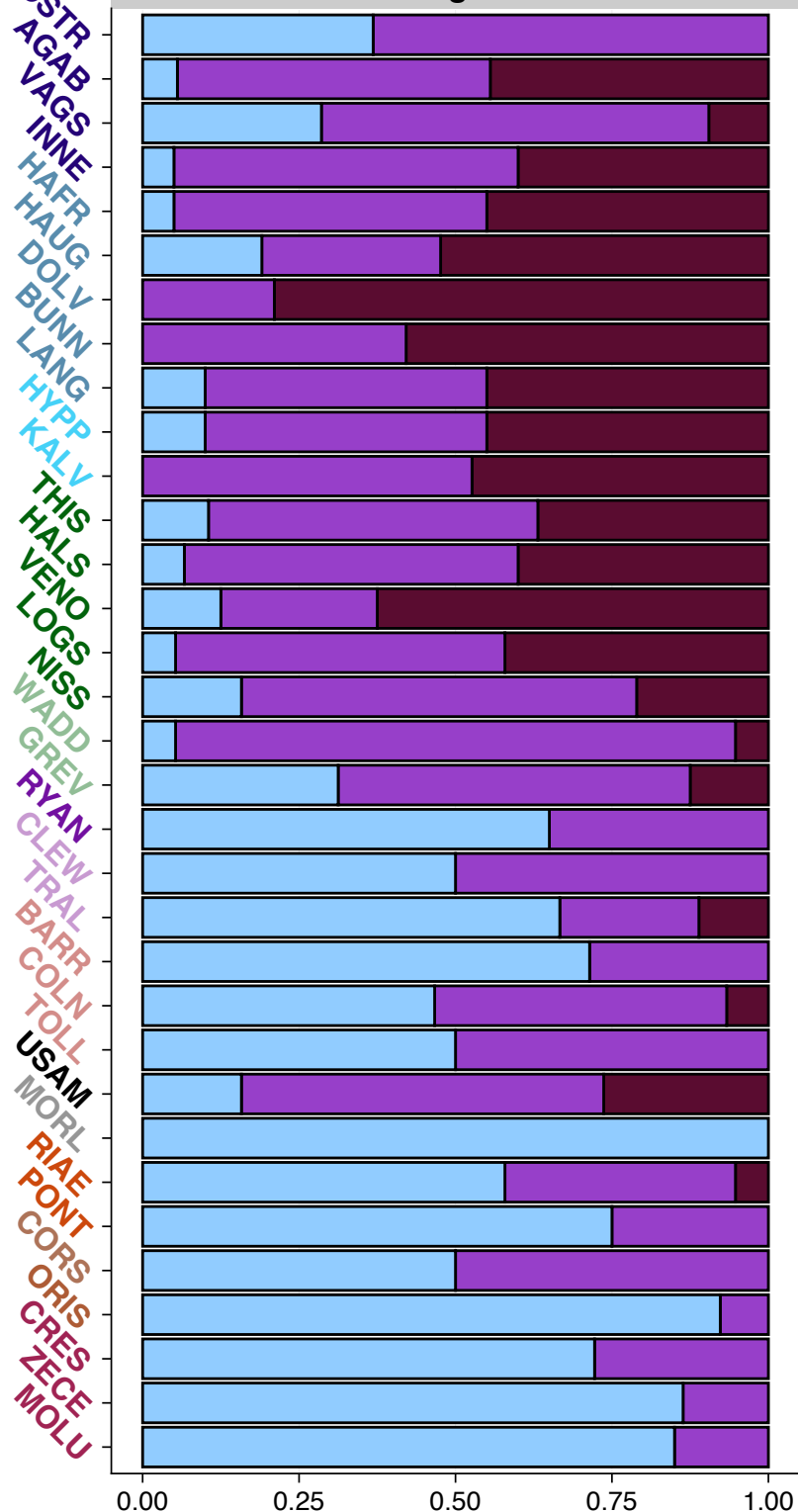
homo_1 het homo_2

population

Reg04



Reg05



Reg08

