Regions 10000 -Counts of SNPs Hs_CEU Hs NE Hs NEU 5000 -Hs PNW Hs SEU Hs WEU 0.0 0.1 0.2 0.3 0.4 0.5 ene diversity (Hs) withing Regions (50% filter)

Genome-wide expected heterozy