## Genome-wide expected heterozygosity (Hs) Regions 10000 -Hs\_CEU Counts of SNPs 7500 -Hs\_NE 5000 -Hs NEU Hs\_PNW 2500 -Hs\_SEU Hs WEU 0 -0.0 0.3 0.5 Gene diversity within Regions