

ACMG-59 Pathogenic: Fraction in 1000 Genomes

Fraction with at least 1 non-reference site

0.6
0.4
0.2
0.0

Superpopulation



ACB ASW ESN GWD LWK MSL YRI CLM MXL PEL PUR CDX CHB CHS JPT KHV CEU FIN GBR IBS TSI BEB GIH ITU PJL STU

Population

