

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

Population