ERR2188722

Variant types

Depth

Quality score

Genotype quality

Main genotypes

Ref (0/0)
Het (0/x)
Hom (x/x)

Other genotypes

Het - two variants (x/y)
Uncalled (./.)

Biallelic base changes from reference

Biallelic Ti/Tv ratio: 5.48

Biallelic indel size distribution