Sample

Variant types

Depth

Quality score

Genotype quality

Main genotypes

Ref (0/0)

Het (0/x)
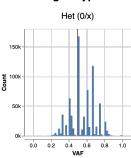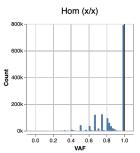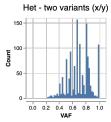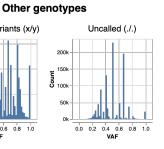

Hom (x/x)

Other genotypes

Het - two variants (x/y)

Uncalled (./.)

Biallelic base changes from reference

Biallelic Ti/Tv ratio: 2.07

Biallelic indel size distribution