# Summary Numbers

| Callset | SNPs | | | indels | | MNPs | others |
|---|---|---|---|---|---|---|---|
| | n | ts/tv | (1st ALT) | n | frm* | | |
| momia | 1,101 | 1.72 | 1.72 | 0 | – | 0 | 0 |
| | | | | | | * frameshift ratio: out/(out+in) |

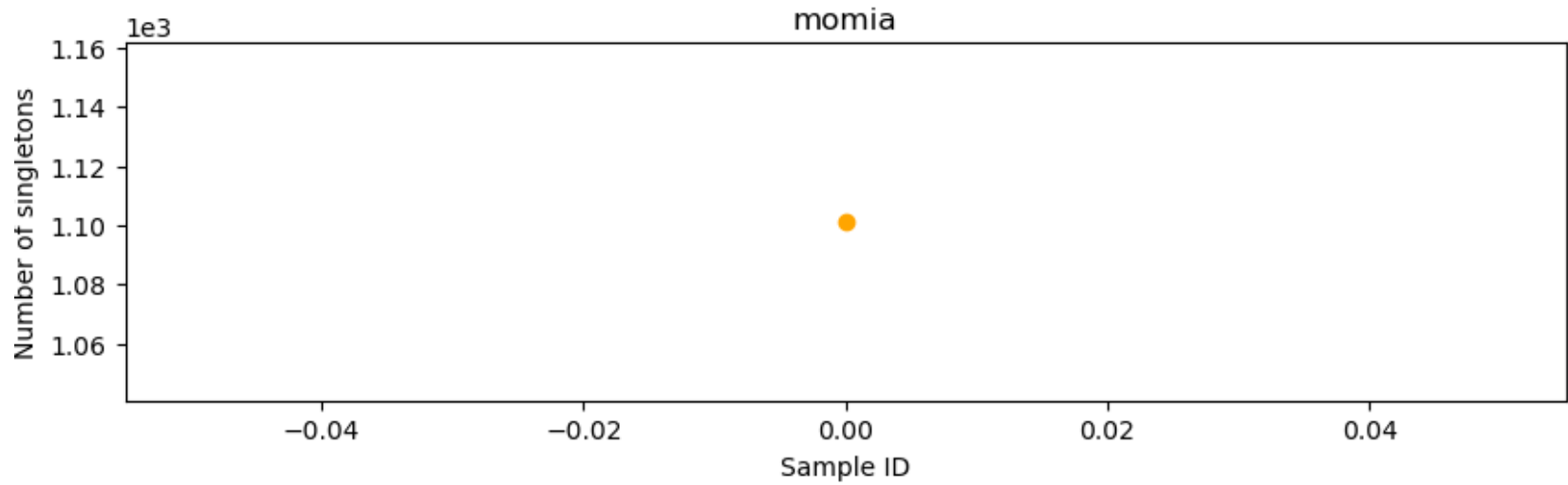| Callset | singletons (AC=1) | | | multiallelic | |
|---|---|---|---|---|---|
| | SNPs | ts/tv | indels | sites | SNPs |
| momia | 0.0% | 0.00 | 0.0% | 0 | 0 |

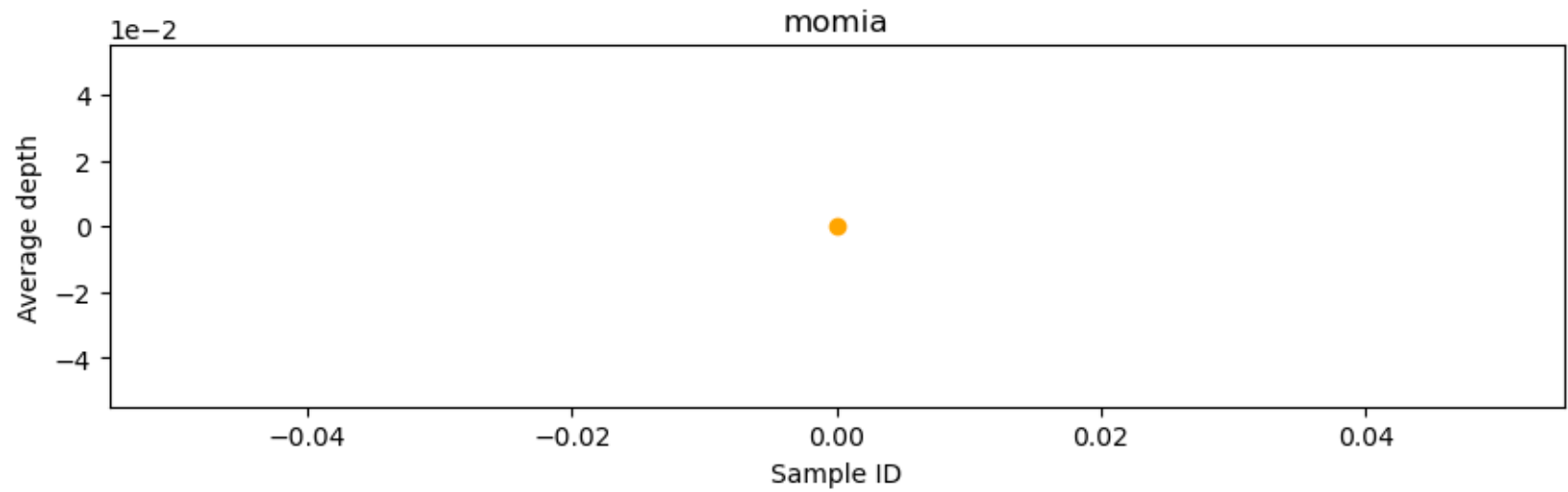- momia .. `momia6_hg38.vcf.gz`
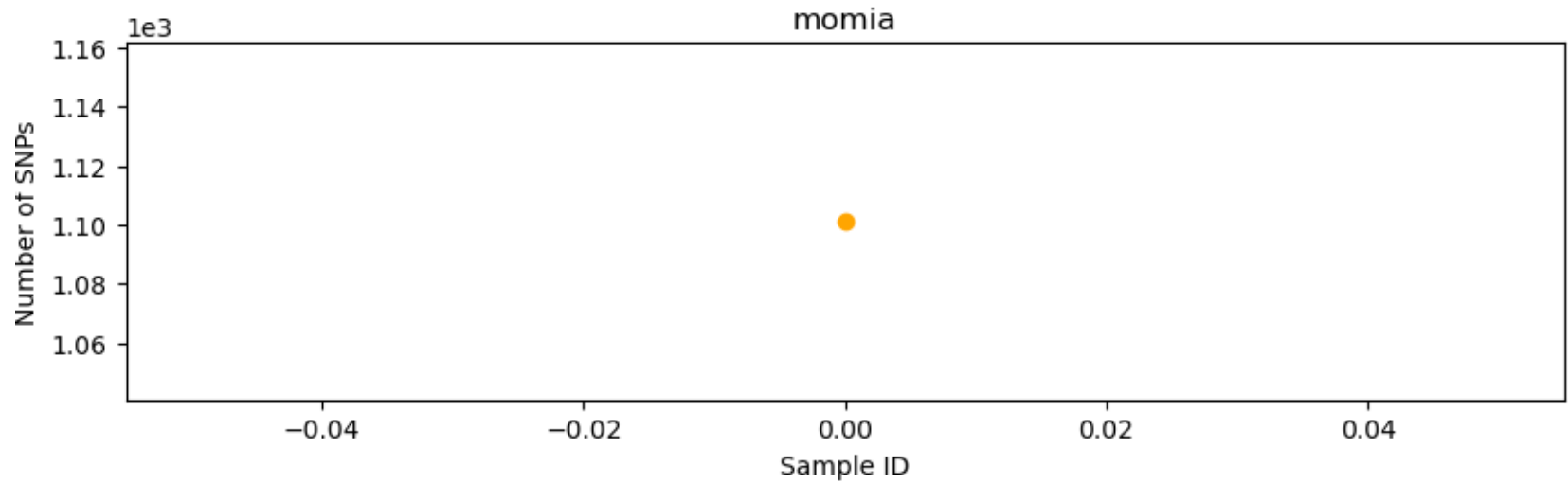
# Ts/Tv by sample
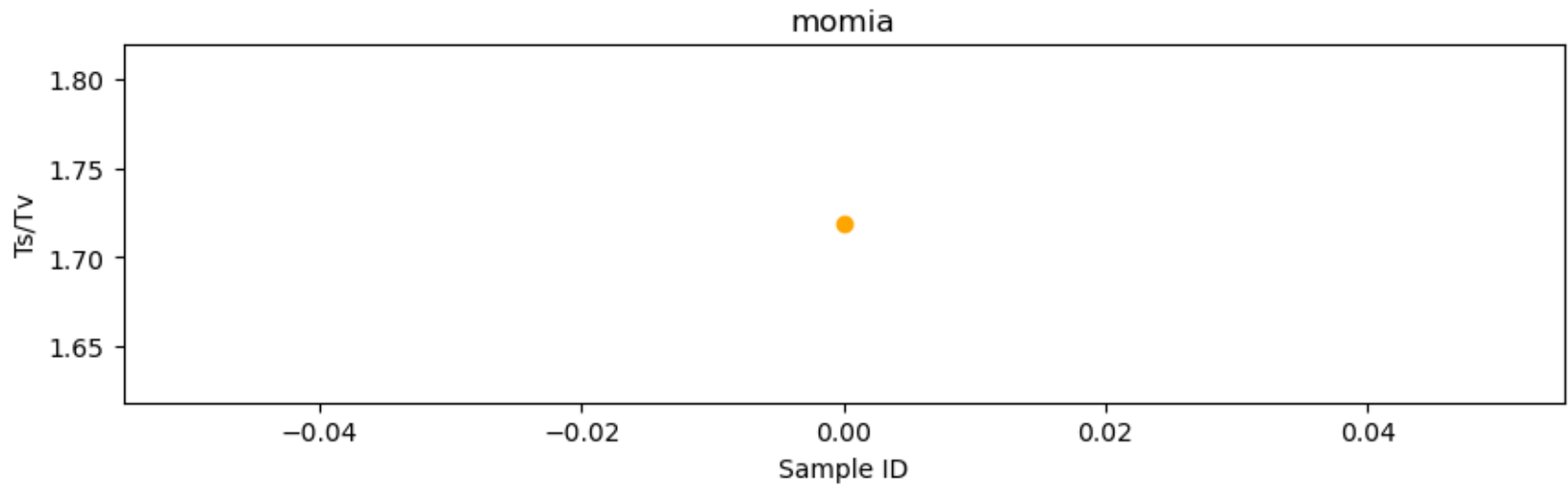
# Hets vs non-ref Homs by sample
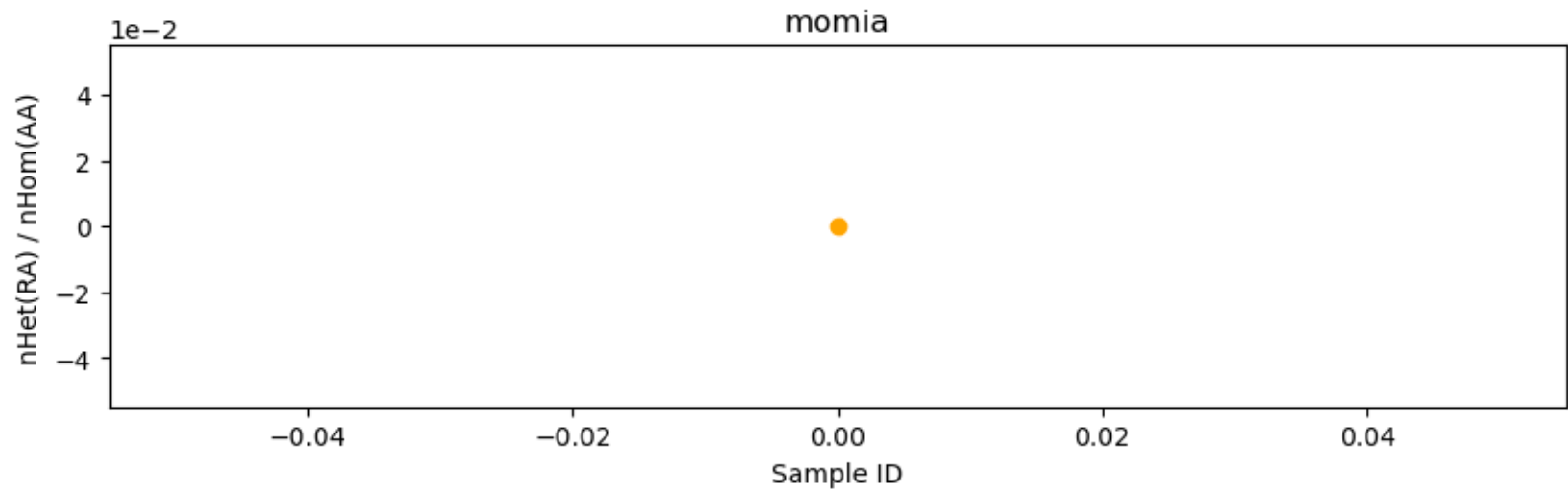
# Singletons by sample (hets and homs)

# Average depth by sample

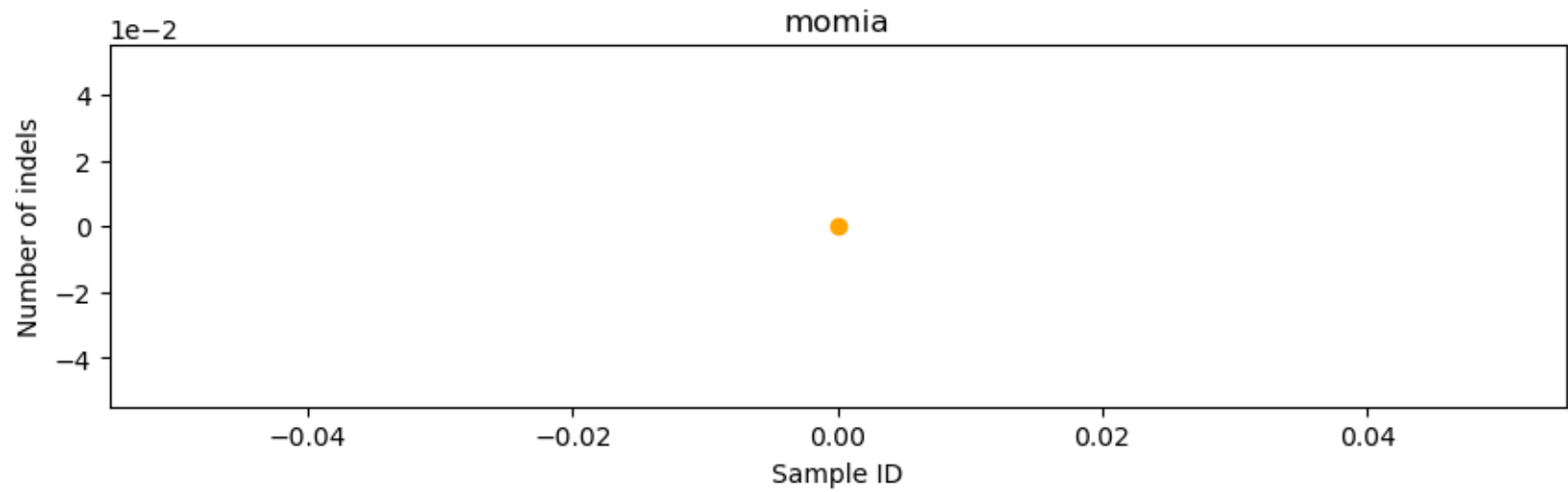# Number of SNPs by sample

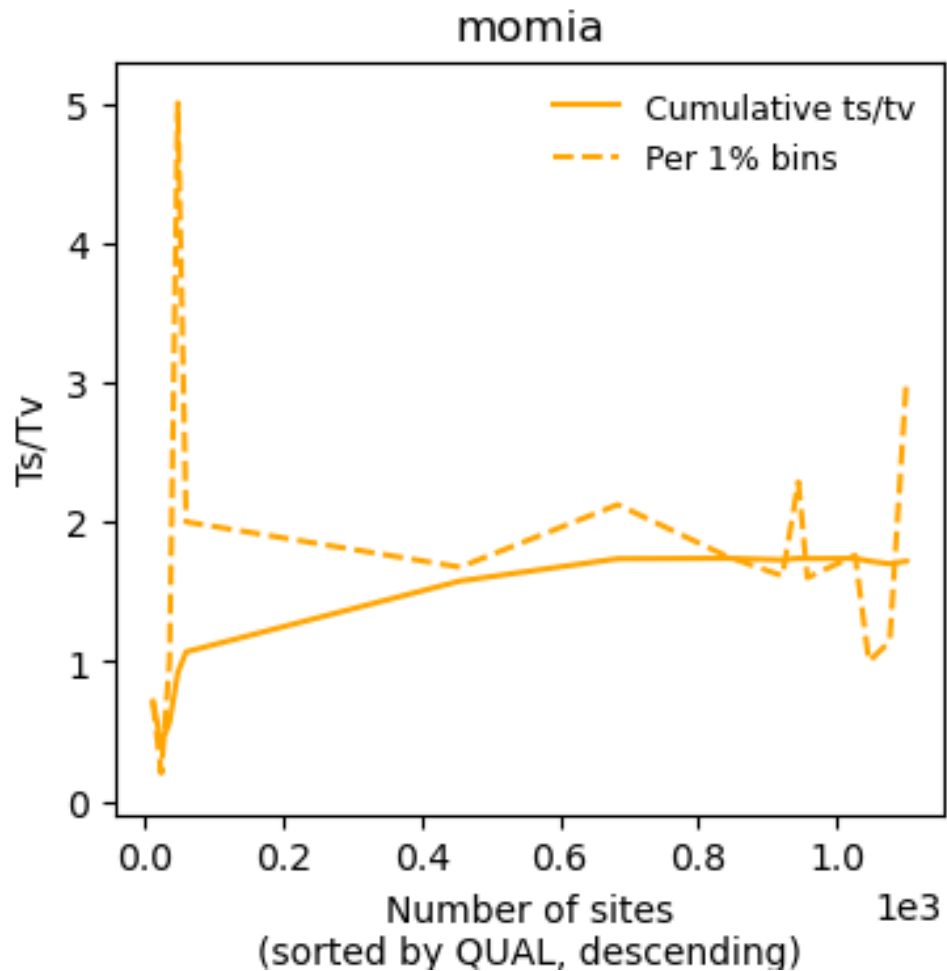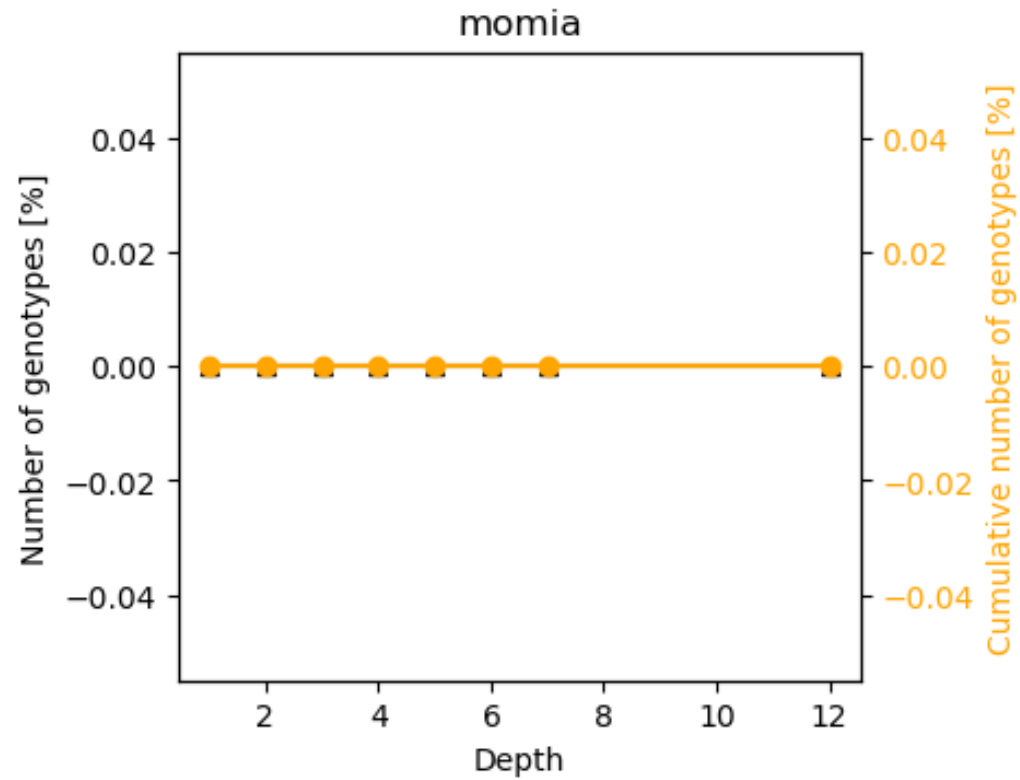# Number of indels by sample

# Ts/Tv stratified by QUAL

# Depth distribution

# Substitution types