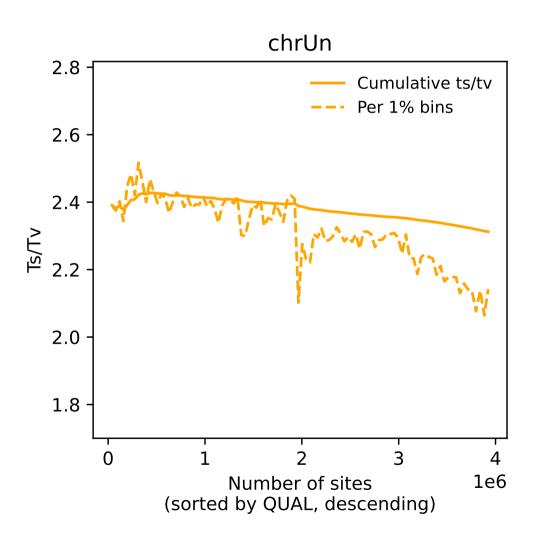
Summary Numbers

| | SNPs | | | indels | | MNPs | others | | | |
|----------------------------------|-----------|-------|-----------|---------|------|------|--------|--|--|--|
| Callset | n | ts/tv | (1st ALT) | n | frm* | | | | | |
| chrUn | 3,924,183 | 2.05 | 2.31 | 167,060 | - | 0 | 0 | | | |
| * frameshift ratio: out/(out+in) | | | | | | | | | | |

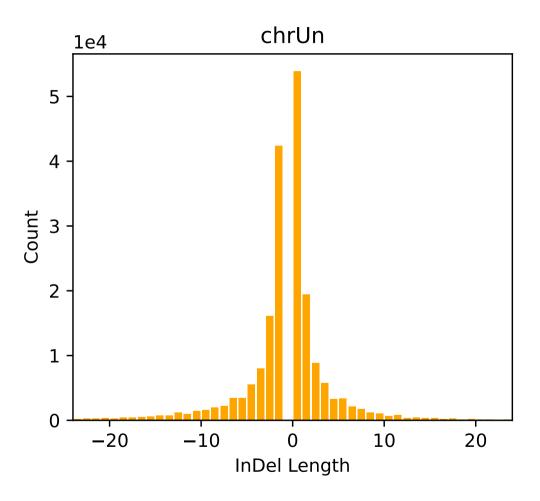
| | single | etons (A | C=1) | multiallelic | | |
|---------|--------|----------|--------|--------------|---------|--|
| Callset | SNPs | ts/tv | indels | sites | SNPs | |
| chrUn | 15.6% | 1.52 | 3.6% | 309,488 | 279,506 | |

⁻ chrUn .. /blue/soltis/kasey.pham/euc_hyb_reseq/call_snps/03.mpileup/chrUn.vcf.gz

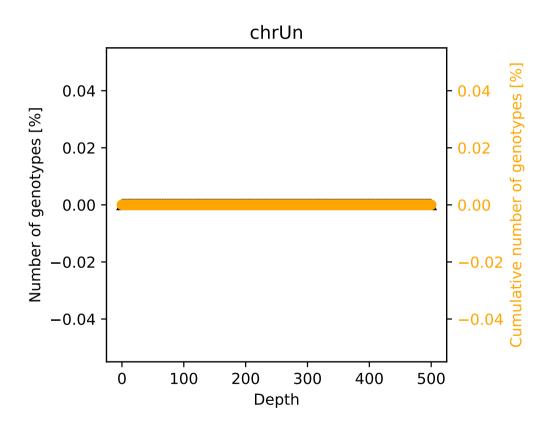
Ts/Tv stratified by QUAL



Indel distribution



Depth distribution



Substitution types

