

# Skeletonema

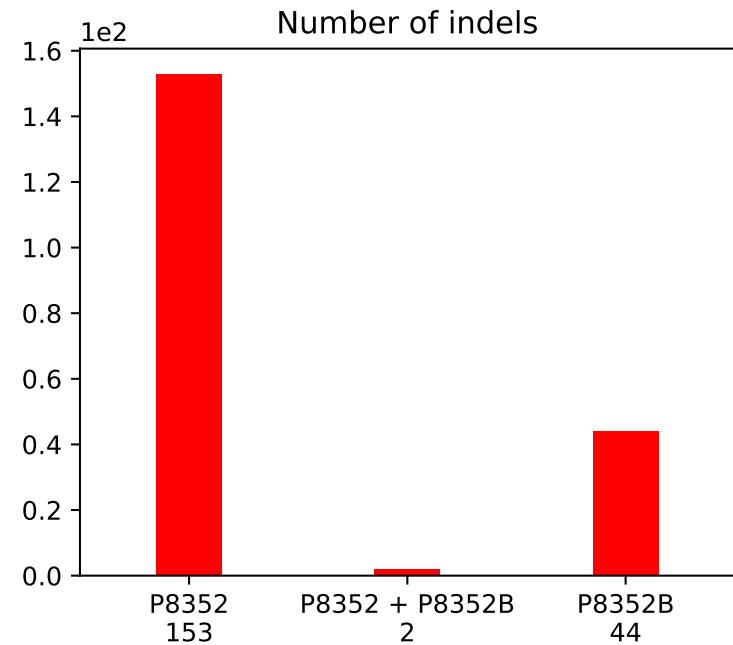
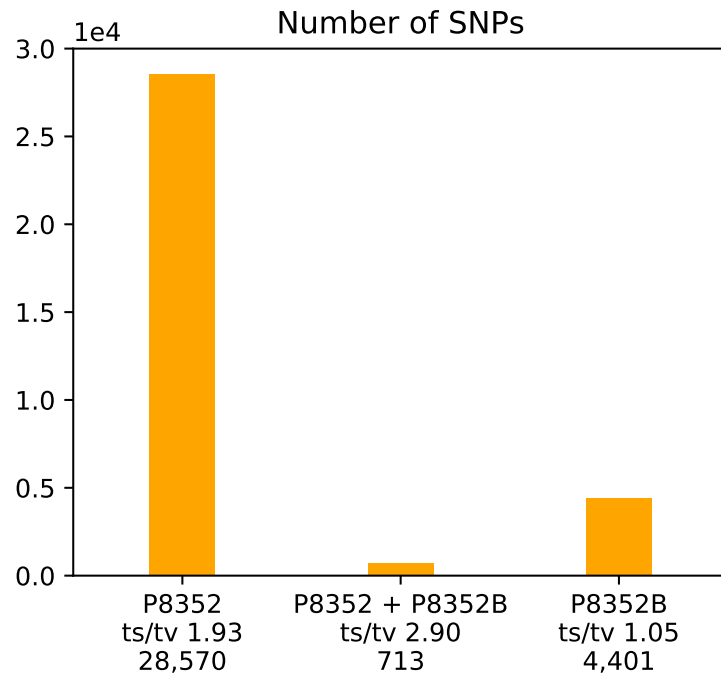
|                | SNPs   |       |           | indels |      | MNPs | others |
|----------------|--------|-------|-----------|--------|------|------|--------|
| Callset        | n      | ts/tv | (1st ALT) | n      | frm* |      |        |
| P8352          | 28,570 | 1.92  | 1.93      | 153    | –    | 0    | 0      |
| P8352B         | 4,401  | 1.05  | 1.05      | 44     | –    | 0    | 0      |
| P8352 + P8352B | 713    | 2.90  | 2.90      | 2      | –    | 0    | 0      |

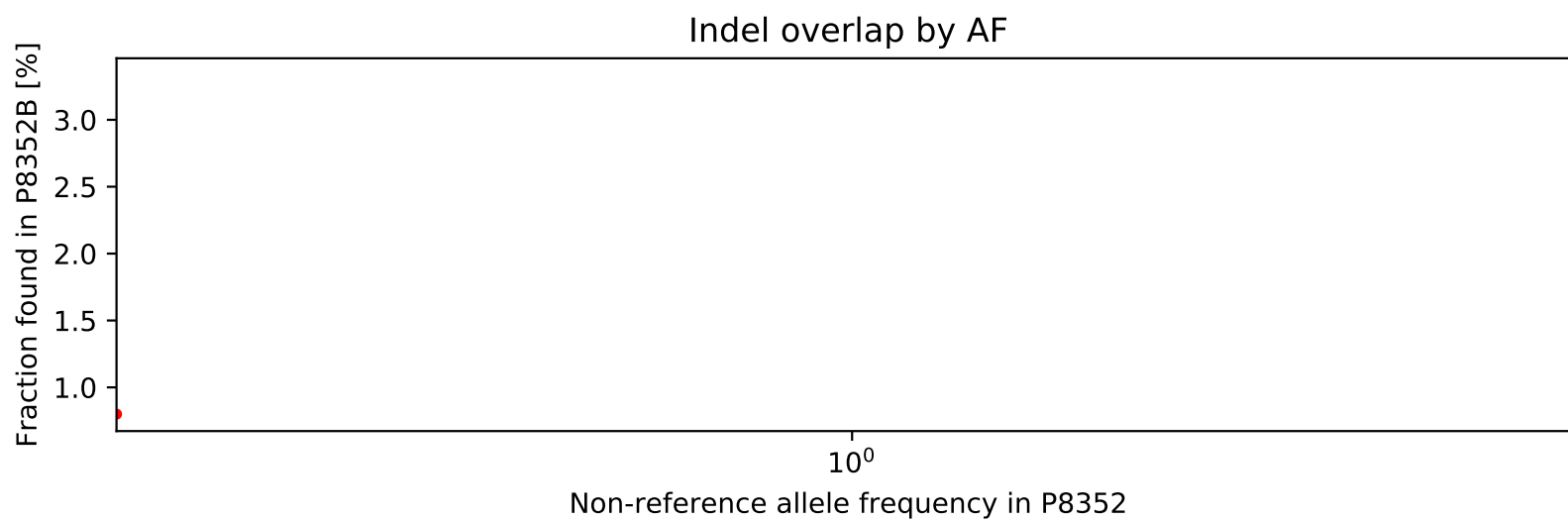
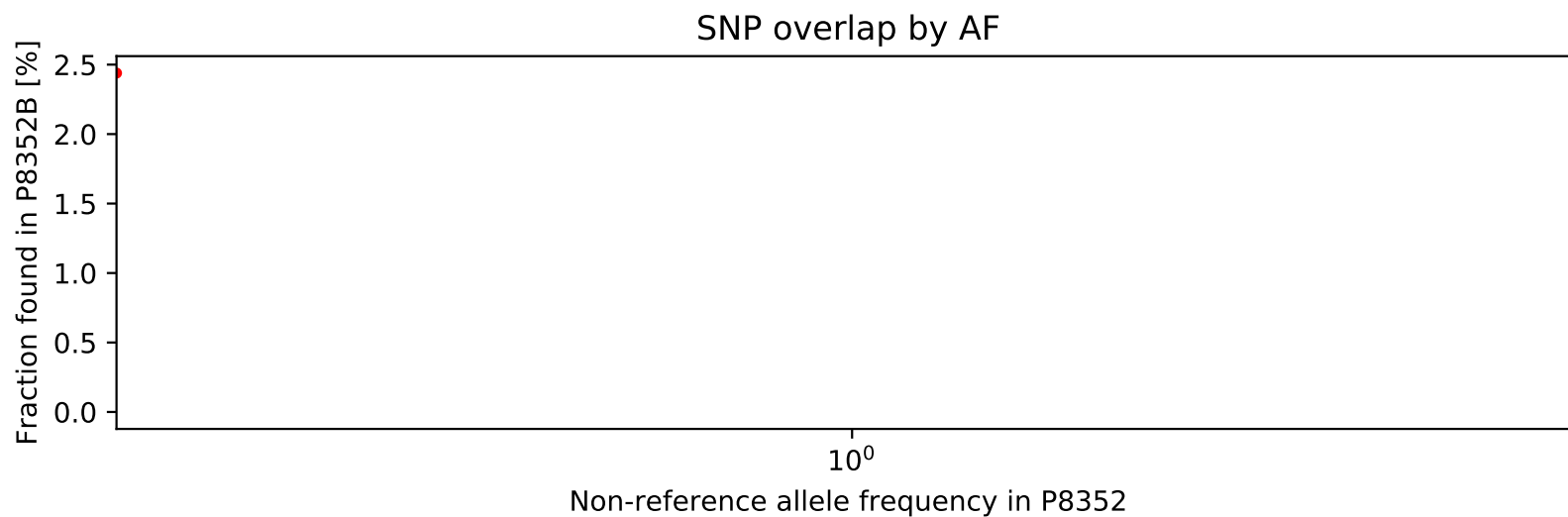
\* frameshift ratio: out/(out+in)

|                | singletons (AC=1) |       |        | multiallelic |      |
|----------------|-------------------|-------|--------|--------------|------|
| Callset        | SNPs              | ts/tv | indels | sites        | SNPs |
| P8352          | 0.2%              | 0.96  | 19.0%  | 2            | 2    |
| P8352B         | 0.0%              | 0.00  | 15.6%  | 1            | 0    |
| P8352 + P8352B | 0.0%              | 0.00  | 50.0%  | 0            | 0    |

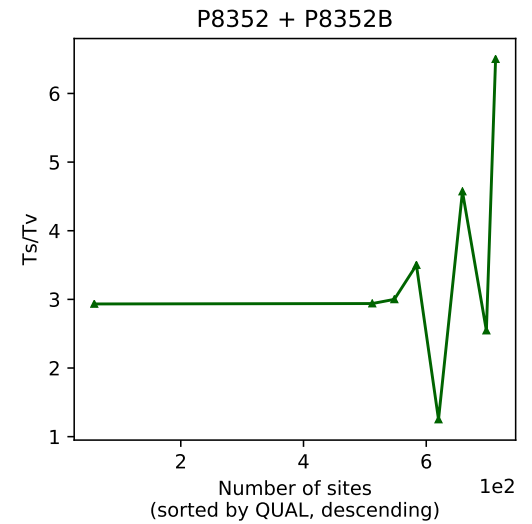
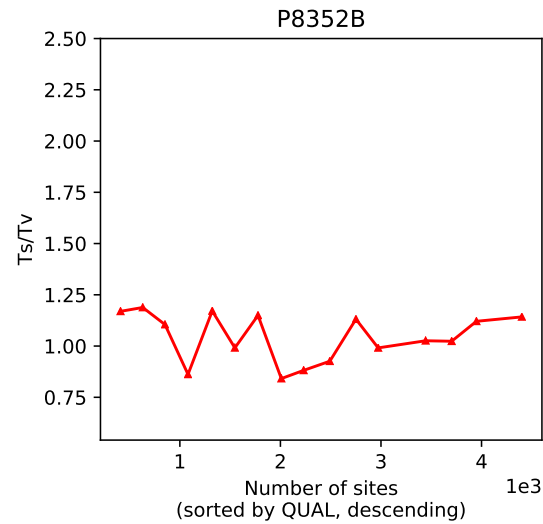
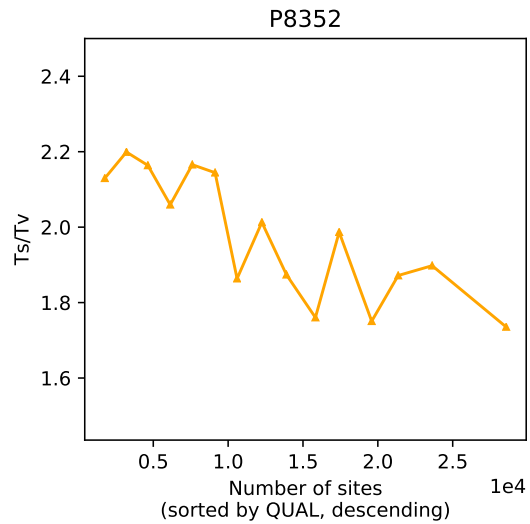
- P8352 .. P8352\_102/Bcftools/P8352\_102.bcftools\_filtered\_new.vcf.gz
- P8352B .. P8352\_104/Bcftools/P8352\_104.bcftools\_filtered\_new.vcf.gz

# Total counts

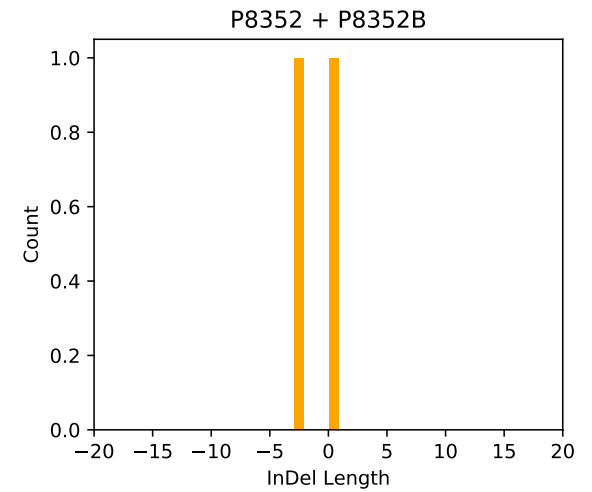
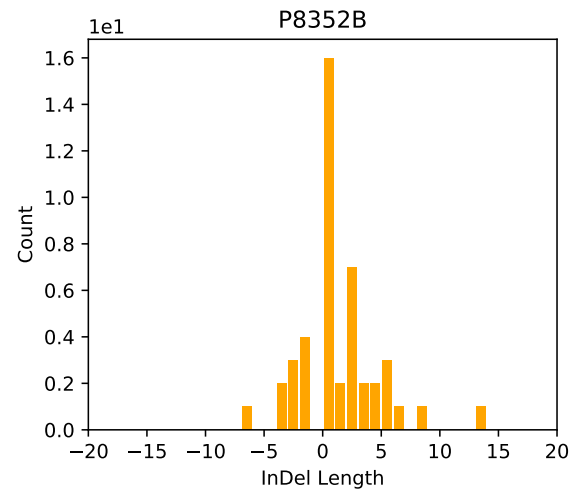
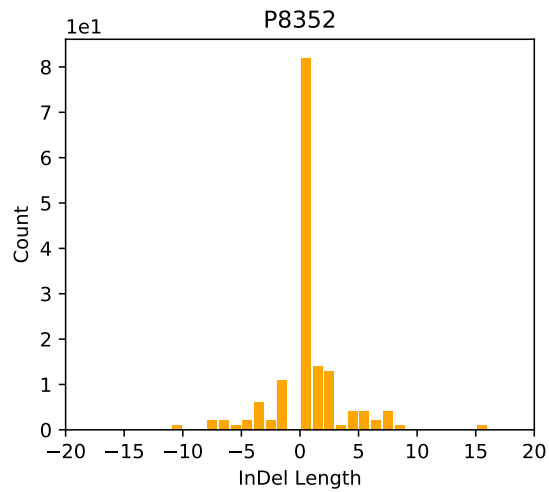




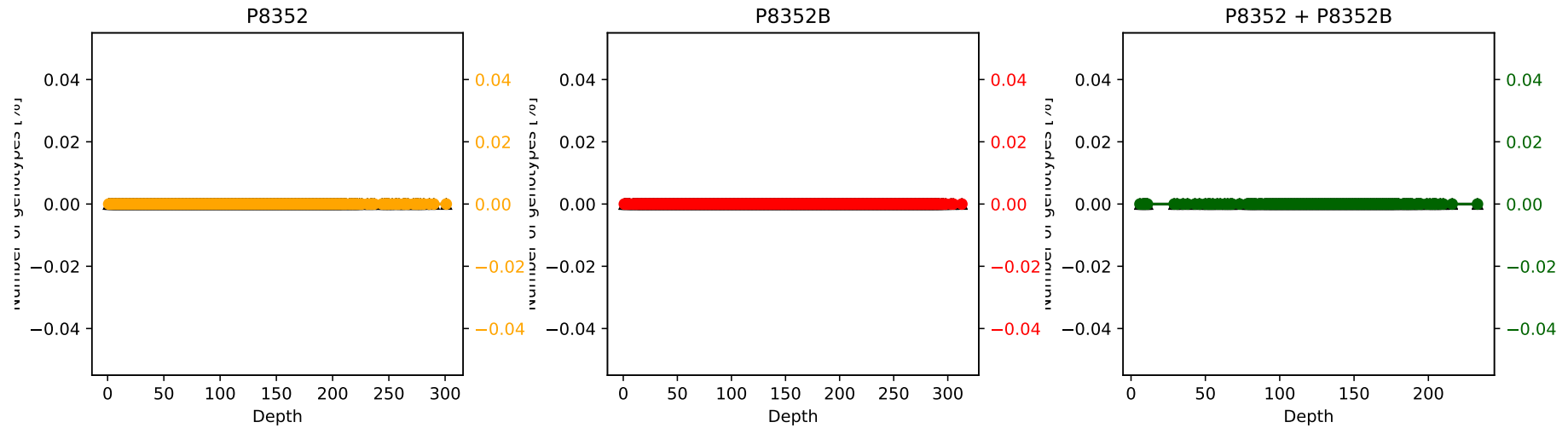
# Ts/Tv stratified by QUAL



# Indel distribution



# Depth distribution



# Substitution types

