

OGRDBstats Report

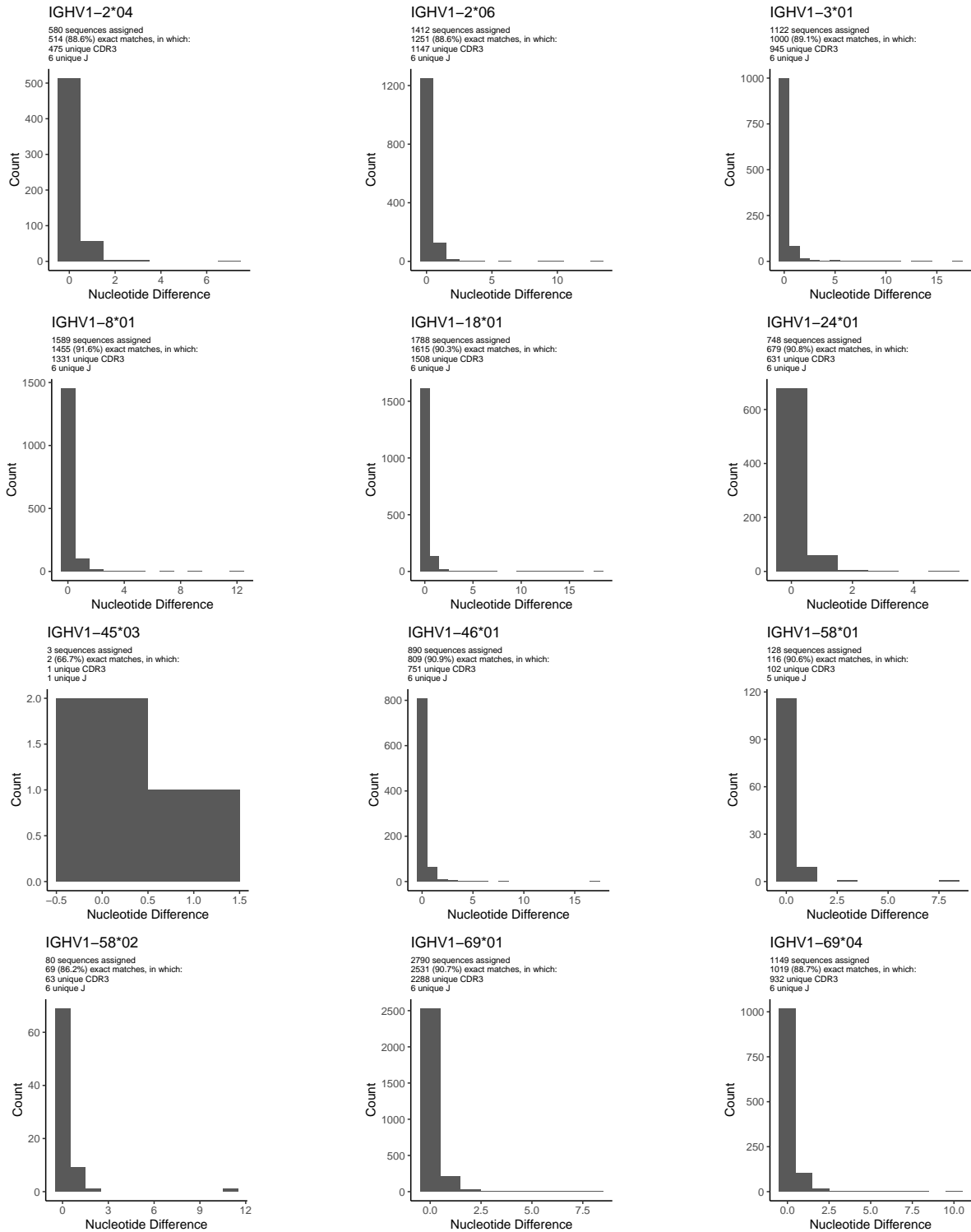
Contents

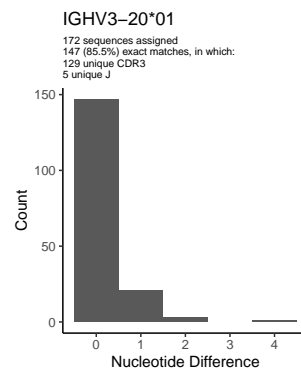
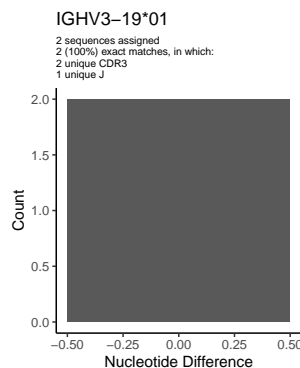
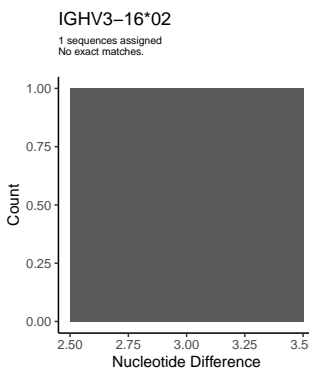
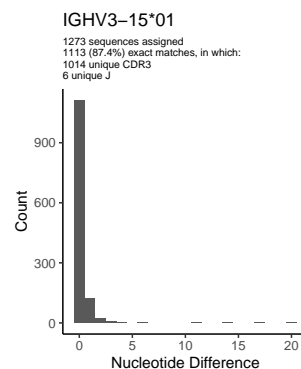
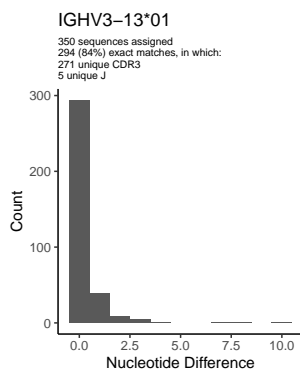
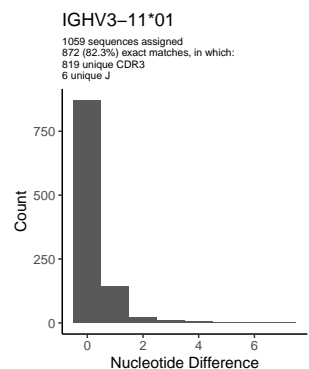
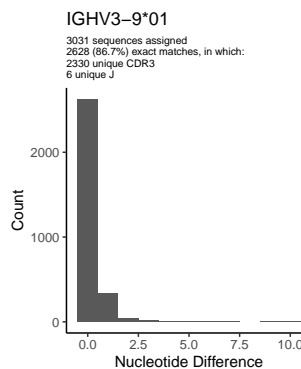
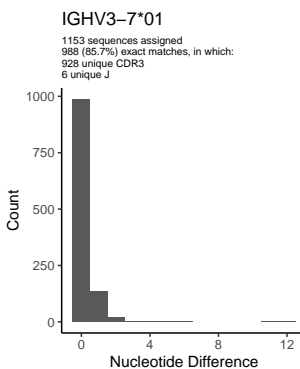
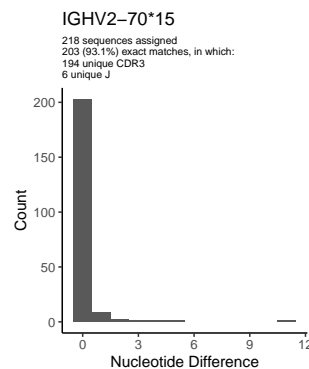
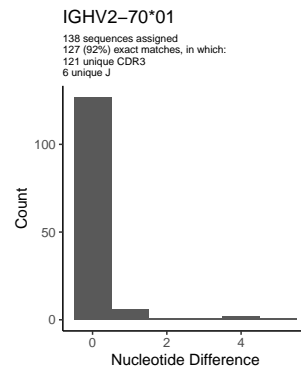
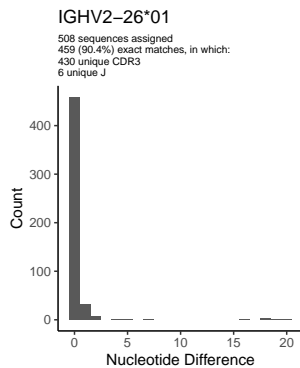
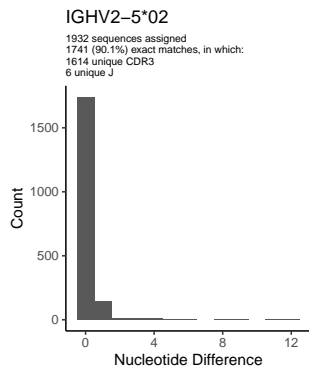
| | | |
|----------|---|-----------|
| 1 | Novel sequence analysis | 2 |
| 1.1 | CDR3 length distribution, in assignments to novel alleles | 2 |
| 2 | Variation from germline, in assignments to each allele | 3 |
| 3 | Allele usage in potential haplotype anchor genes | 8 |
| 4 | Haplotype plots | 9 |
| 5 | Configuration settings | 10 |

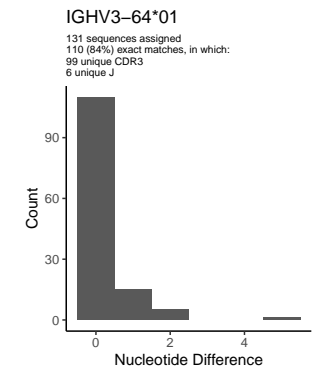
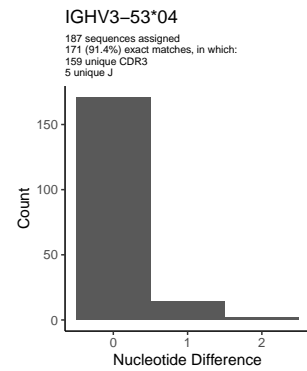
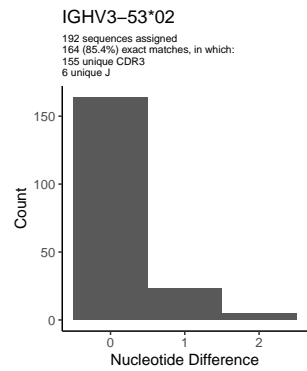
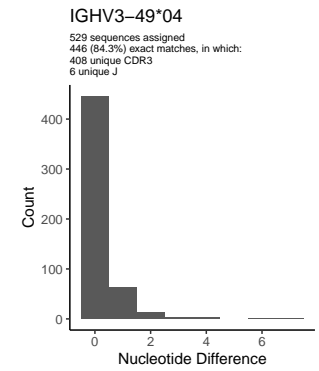
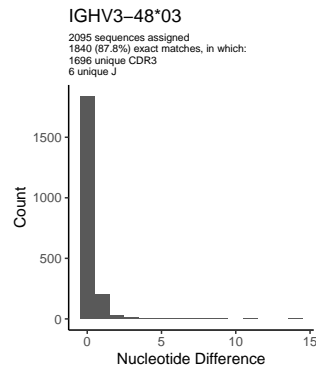
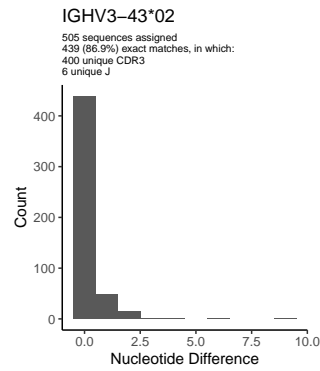
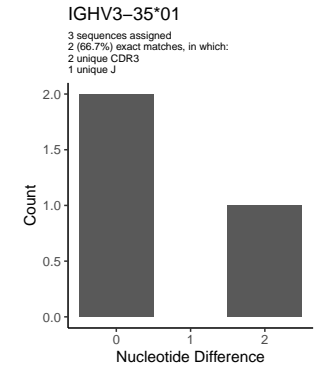
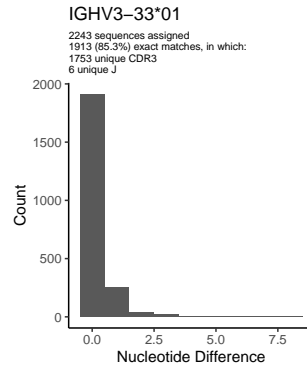
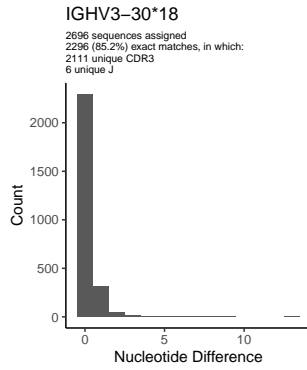
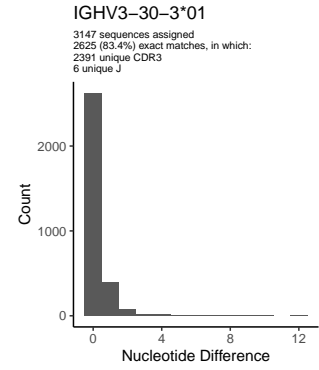
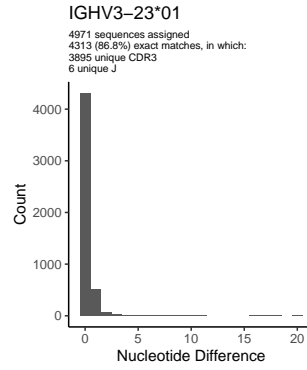
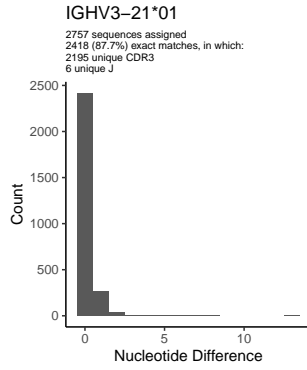
1 Novel sequence analysis

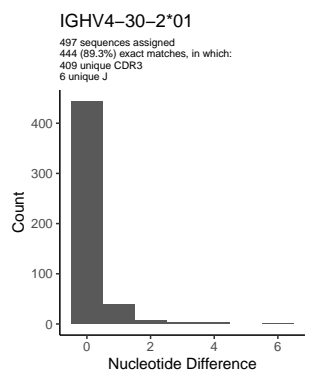
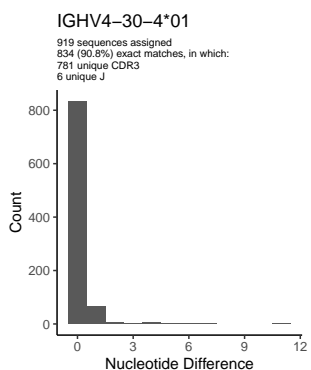
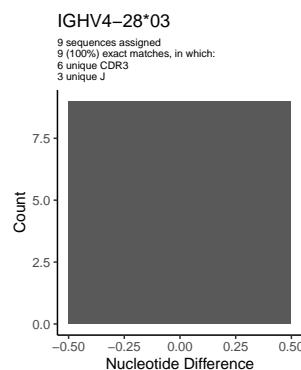
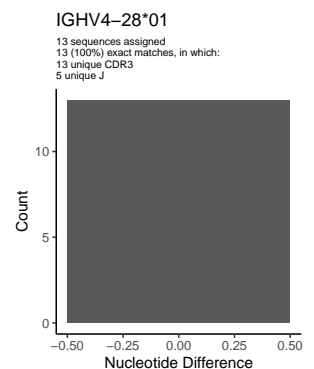
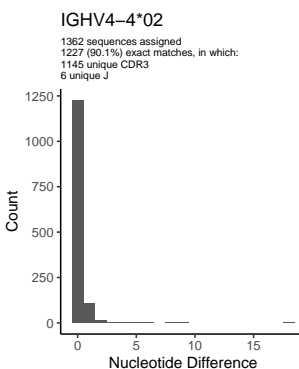
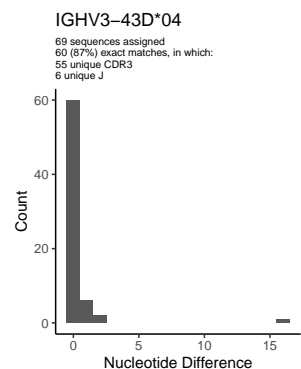
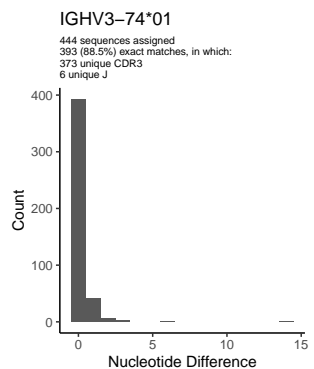
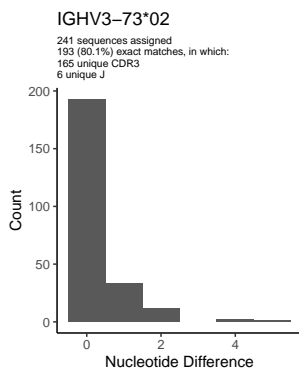
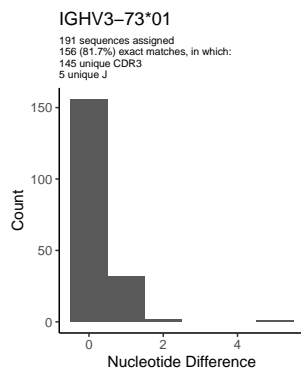
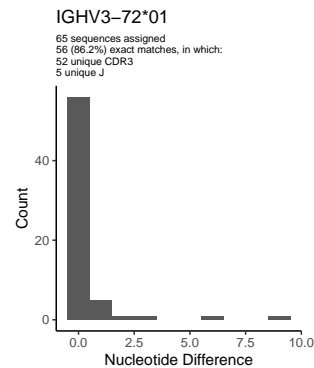
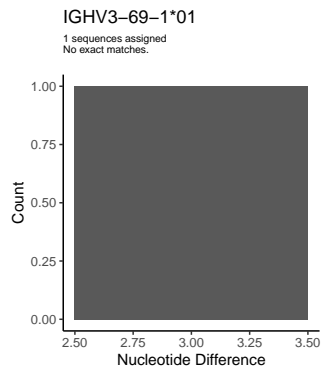
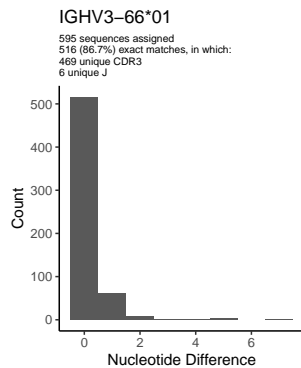
1.1 CDR3 length distribution, in assignments to novel alleles

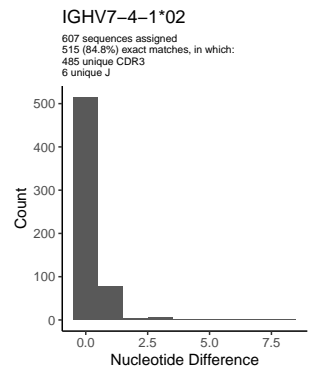
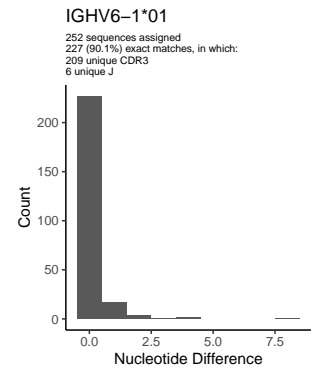
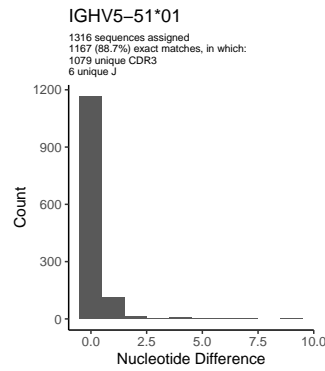
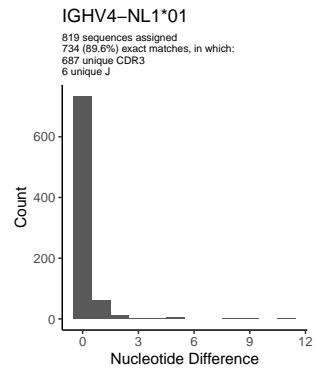
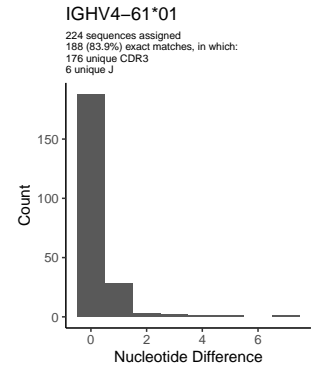
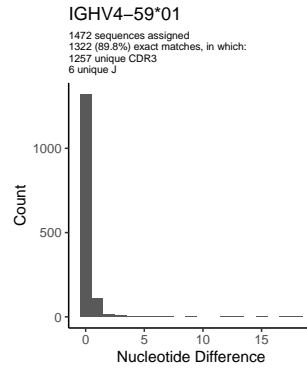
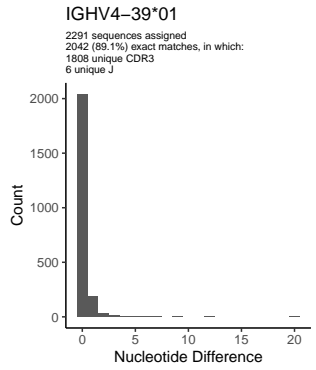
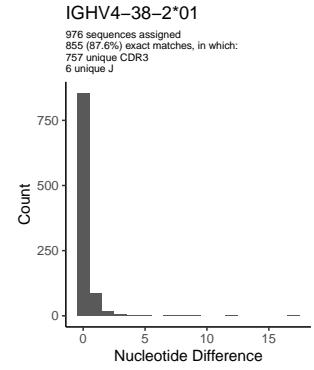
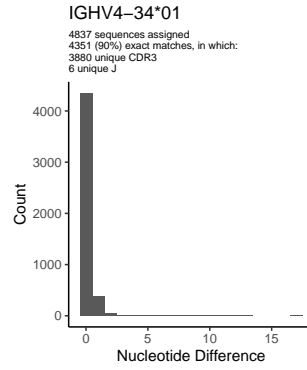
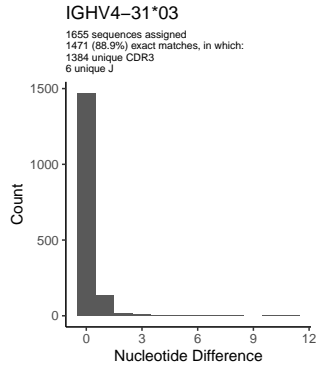
2 Variation from germline, in assignments to each allele











3 Allele usage in potential haplotype anchor genes



4 Haplotype plots

5 Configuration settings

```
## Repertoire file: /misc/work/jenkins/PRJEB26509/S67/S67/71/71/54/e7c68ffaae5ed08fc0341f23827552/71_Fi
##
## Germline reference file: /misc/work/jenkins/PRJEB26509/S67/S67/71/71/aa/b02cdc29a7724ebbf57ab68827ad
##
## Novel allele file:
##
## Species: Homosapiens
##
## Chain: IGHV
##
## Segment: V
##
## Warning    no inferred sequences found.
```