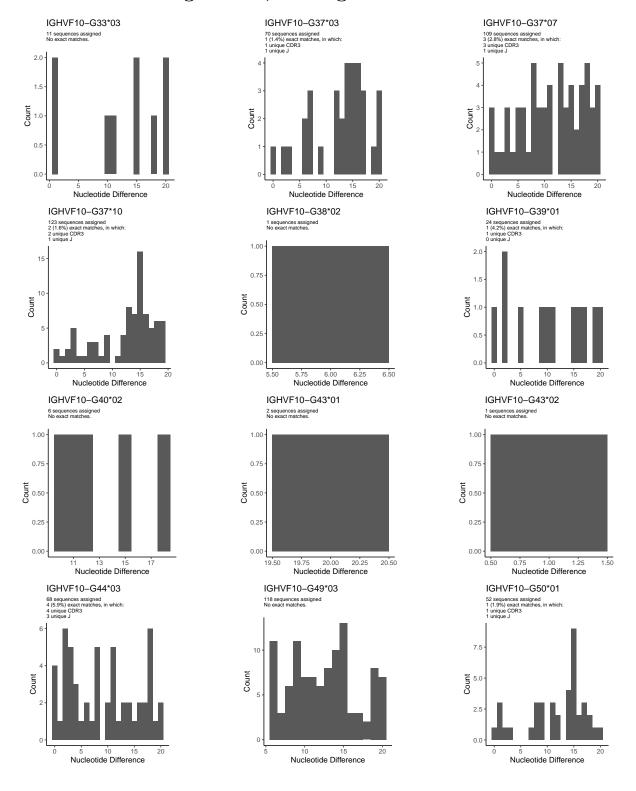
# OGRDBstats Report

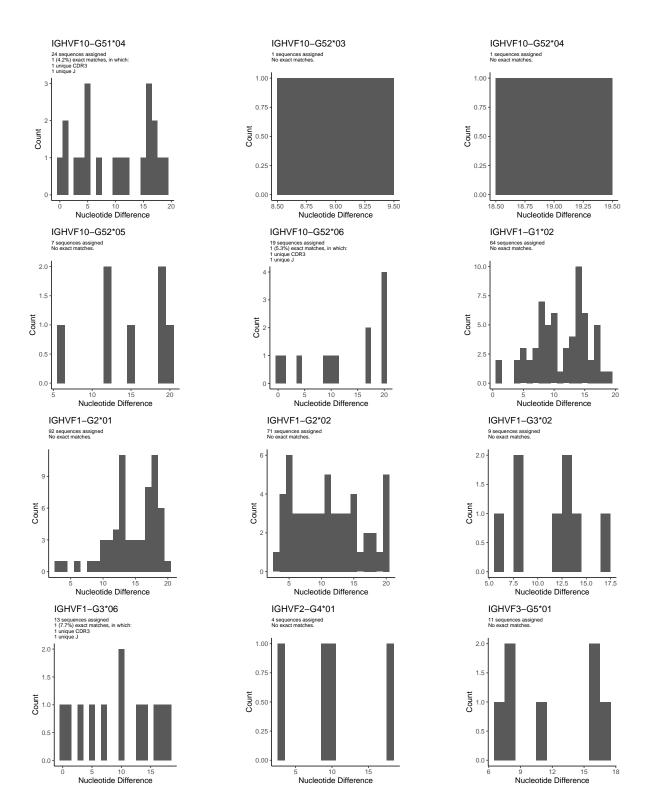
### Contents

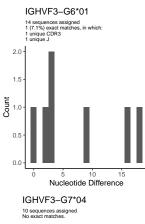
1	Novel sequence analysis 1.1 CDR3 length distribution, in assignments to novel alleles	<b>2</b> 2
2	Variation from germline, in assignments to each allele	3
3	Allele usage in potential haplotype anchor genes	8
4	Haplotype plots	9
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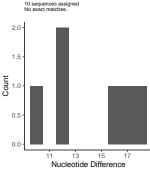
- 1 Novel sequence analysis
- 1.1 CDR3 length distribution, in assignments to novel alleles

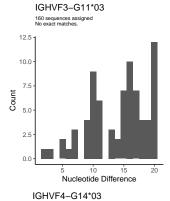
### 2 Variation from germline, in assignments to each allele

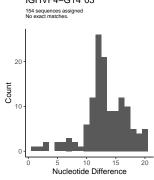


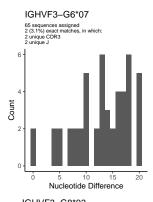


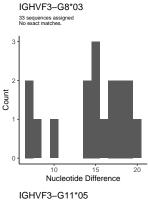


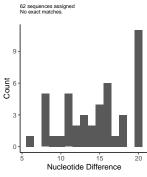


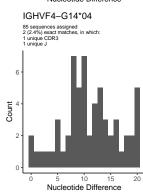


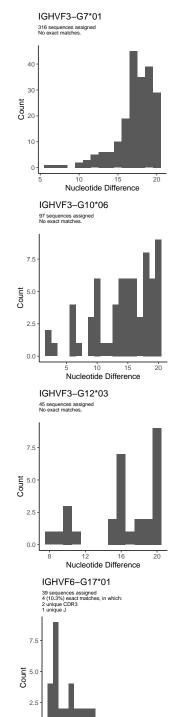


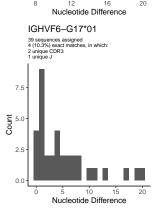


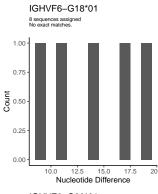




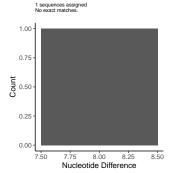




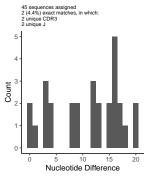




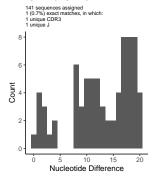




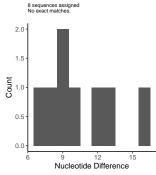
IGHVF6-G22\*02



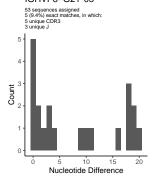
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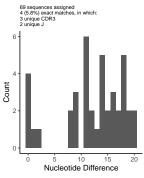
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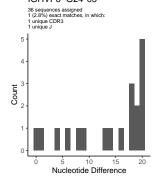
#### IGHVF6-G21\*03



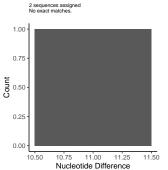
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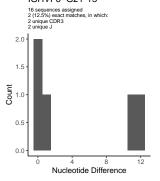
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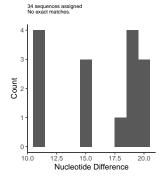
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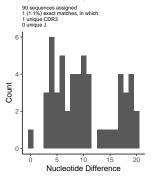
#### IGHVF6-G21\*13

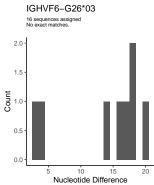


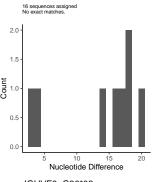
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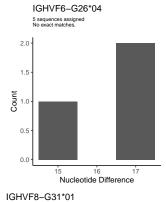


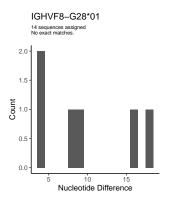
IGHVF6-G25\*02





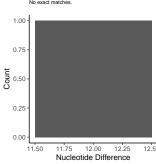


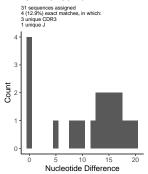






1 sequences assigned No exact matches.





## 3 Allele usage in potential haplotype anchor genes



4 Haplotype plots

### 5 Configuration settings

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## Repertoire file: /misc/work/jenkins/PRJNA248475/M3/M3_cln/M3_cln/M3_cln/24/a209f81cb8d29fbb334cd10c1
##
## Germline reference file: /misc/work/jenkins/PRJNA248475/M3/M3_cln/M3_cln/M3_cln/54/1efbe6c92217f71bd
##
## Novel allele file:
##
## Species: Homosapiens
##
## Chain: IGHV
##
## Segment: V
##
## Warning no inferred sequences found.
```