

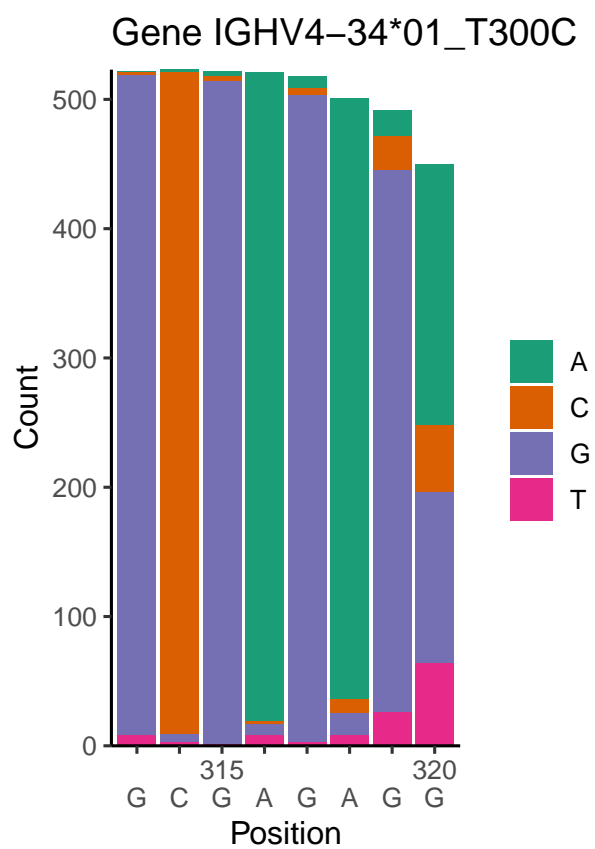
# OGRDBstats Report

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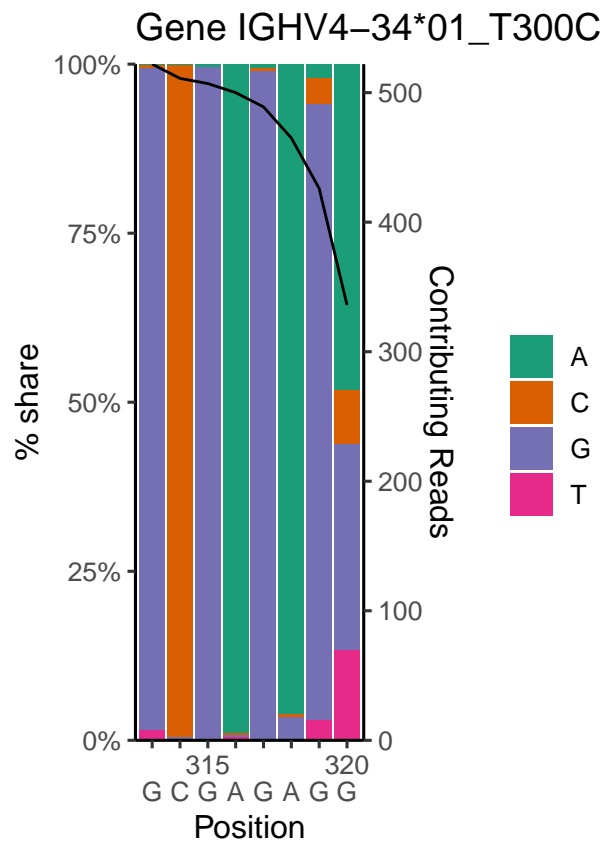
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# 1 Novel sequence analysis

## 1.1 End-nucleotide composition

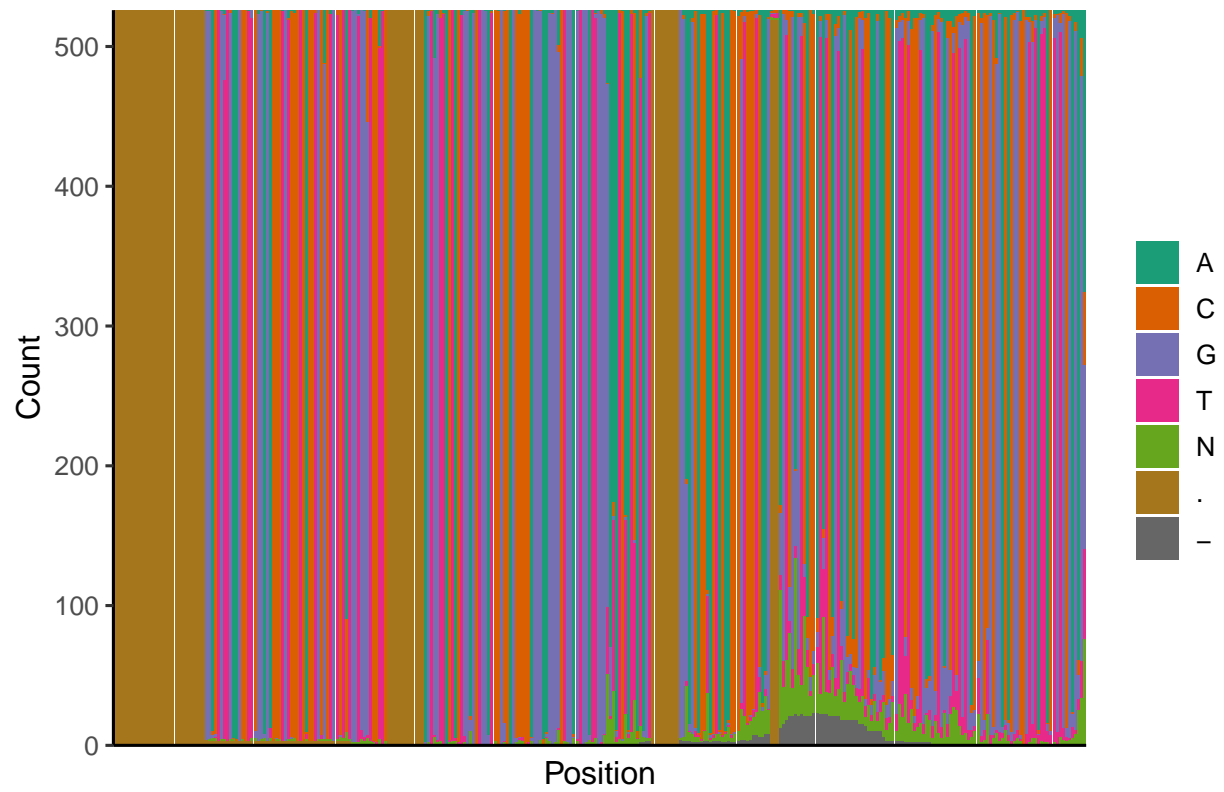


## 1.2 Per-nucleotide consensus where previous nucleotides match the consensus

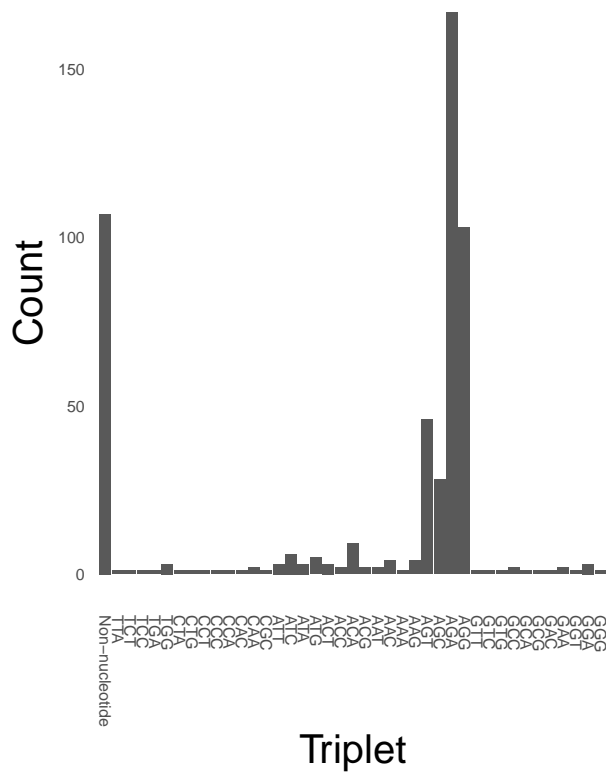


### 1.3 Whole-sequence composition of each assigned read

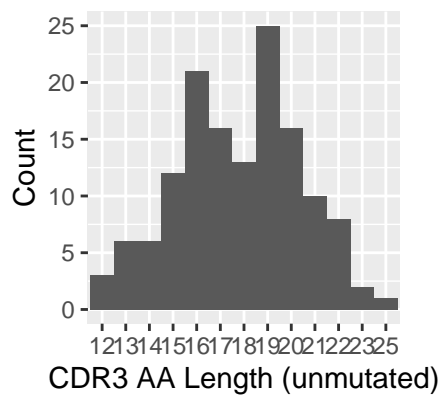
Gene IGHV4-34\*01\_T300C



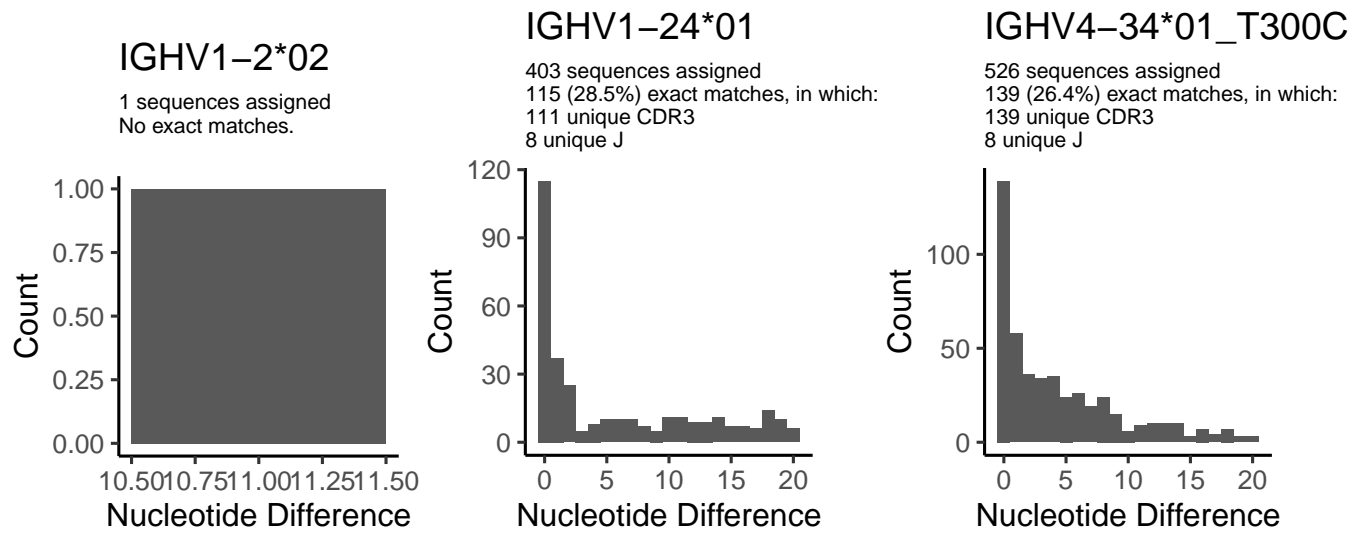
#### 1.4 Final three nucleotides: frequency of each observed triplet IGHV4-34\*01\_T300C- Final 3 nucleotides as a triplet



#### 1.5 CDR3 length distribution, in assignments to novel alleles IGHV4-34\*01\_T300C



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



### 3 Allele usage in potential haplotype anchor genes

## 4 Configuration settings