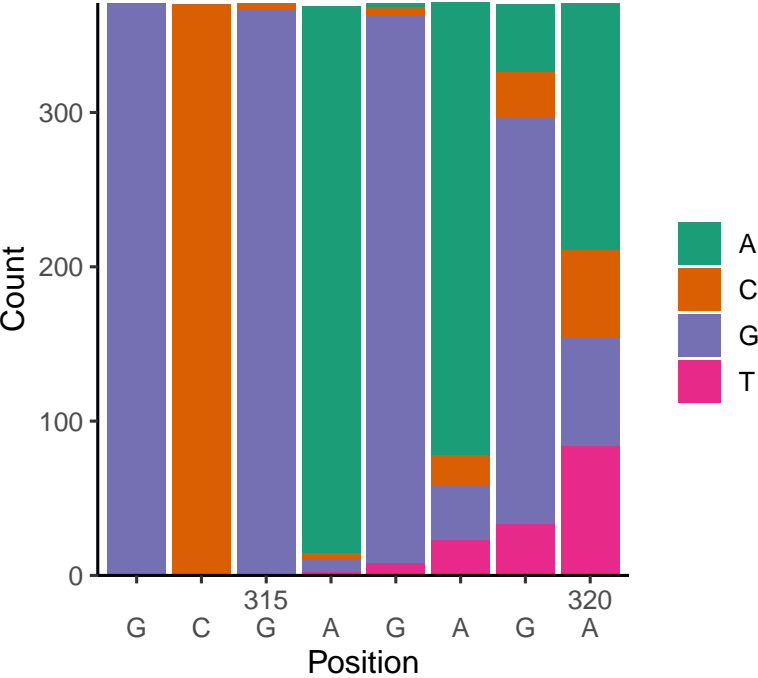
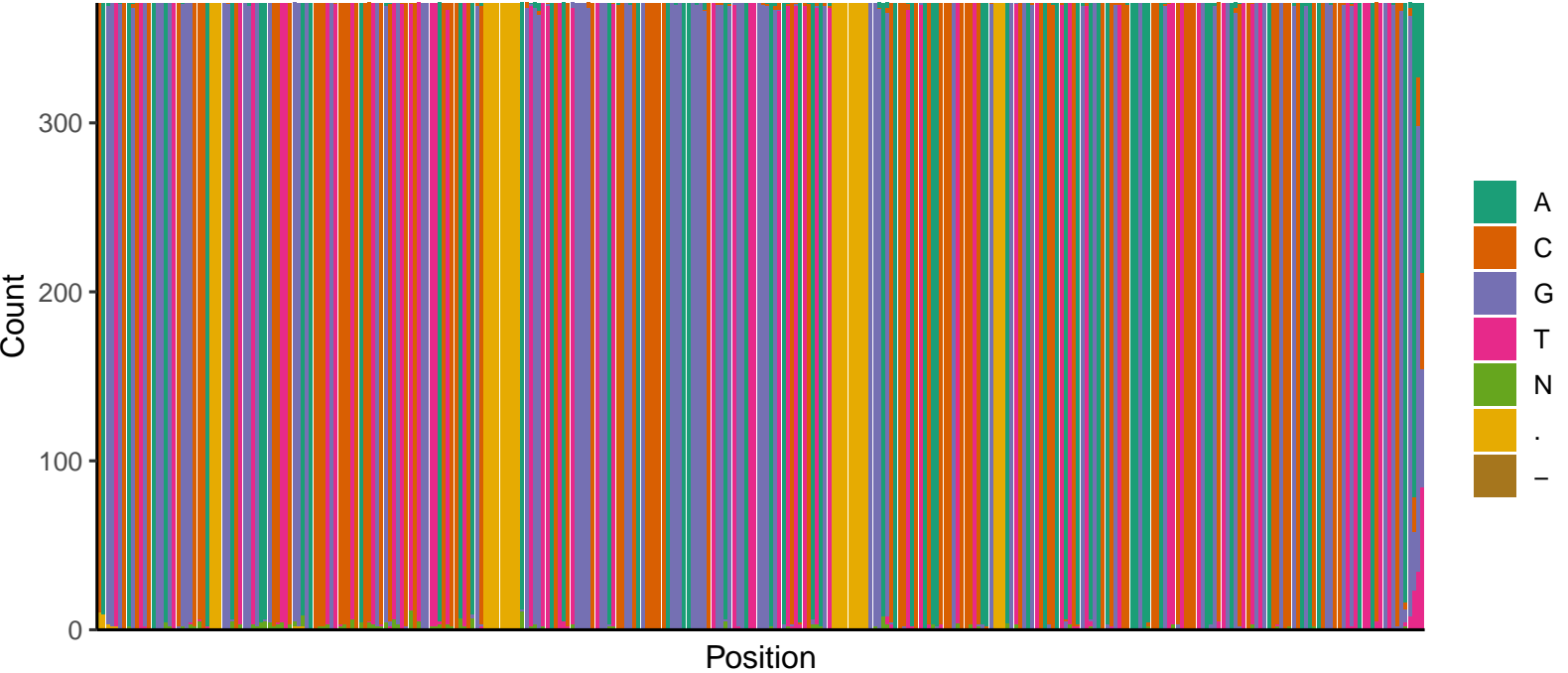


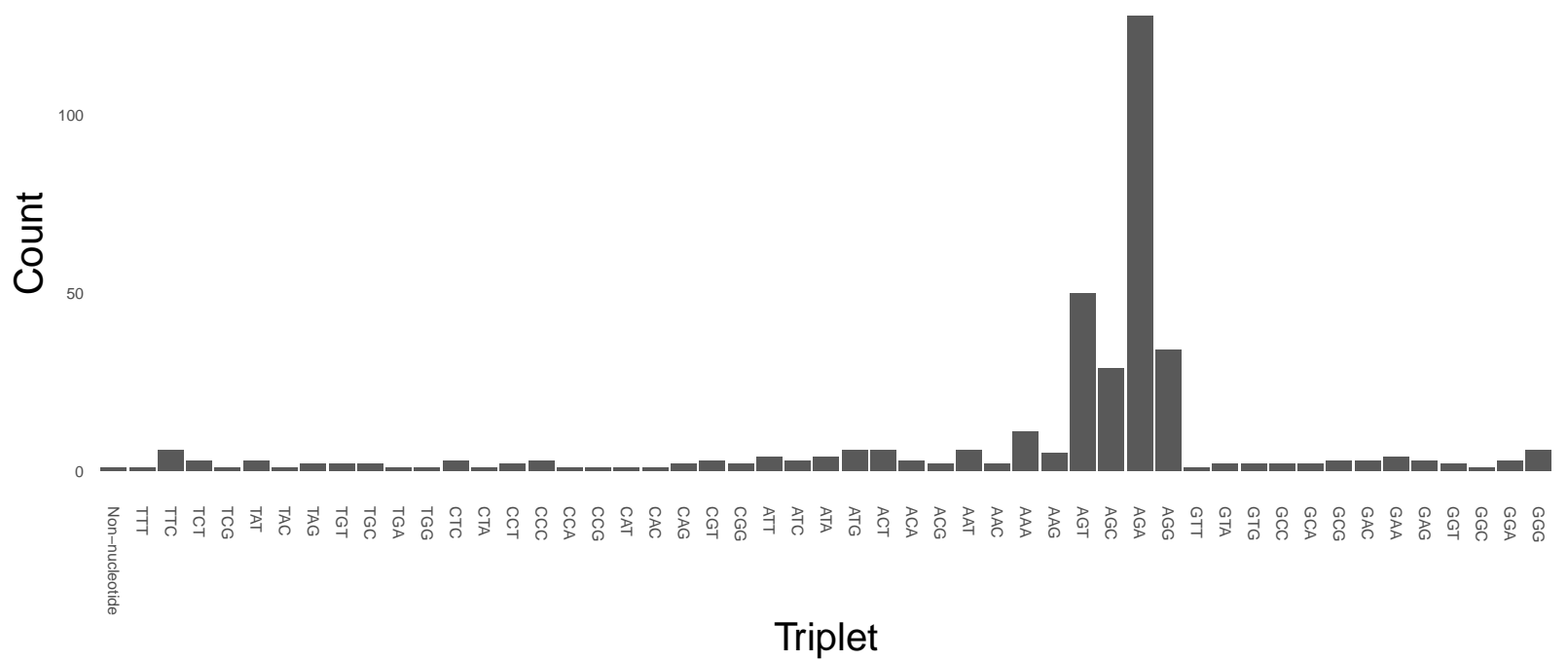
Gene IGHV4-38-2\*02\_A70G



Gene IGHV4-38-2\*02\_A70G

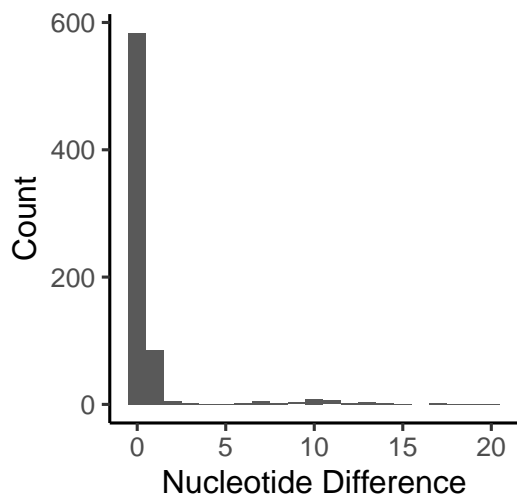


IGHV4-38-2\*02\_A70G- Final 3 nucleotides as a triplet



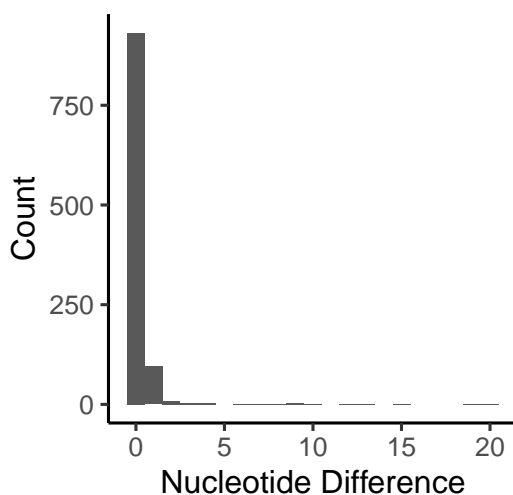
### IGHV1-2\*02

725 sequences assigned  
584 (80.6%) exact matches, in which:  
579 unique CDR3  
6 unique J



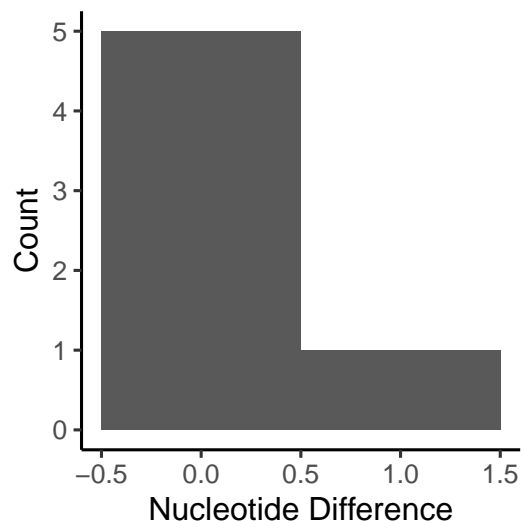
### IGHV1-8\*01

1059 sequences assigned  
932 (88%) exact matches, in which:  
924 unique CDR3  
7 unique J



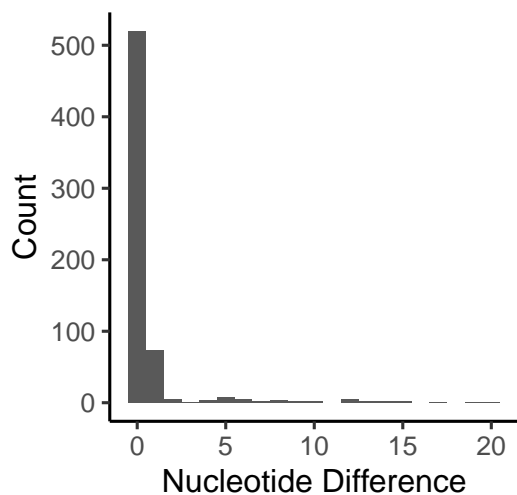
### IGHV1-45\*02

7 sequences assigned  
5 (71.4%) exact matches, in which:  
5 unique CDR3  
3 unique J



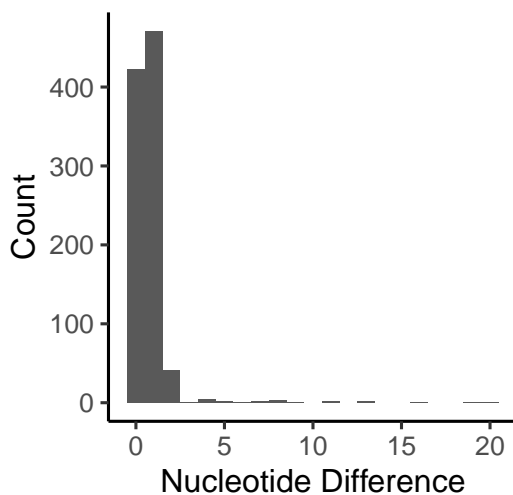
### IGHV1-2\*06

642 sequences assigned  
520 (81%) exact matches, in which:  
515 unique CDR3  
6 unique J



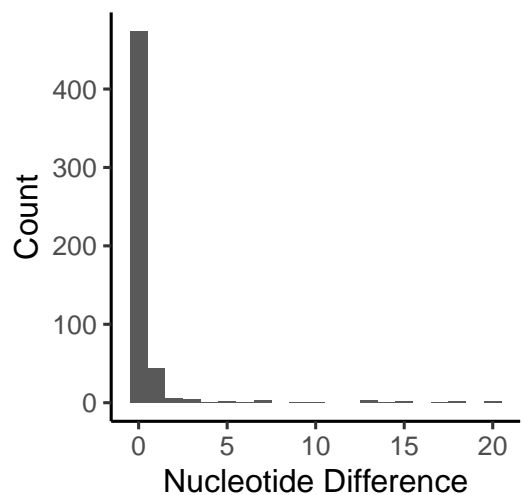
### IGHV1-18\*01

958 sequences assigned  
422 (44.1%) exact matches, in which:  
420 unique CDR3  
6 unique J



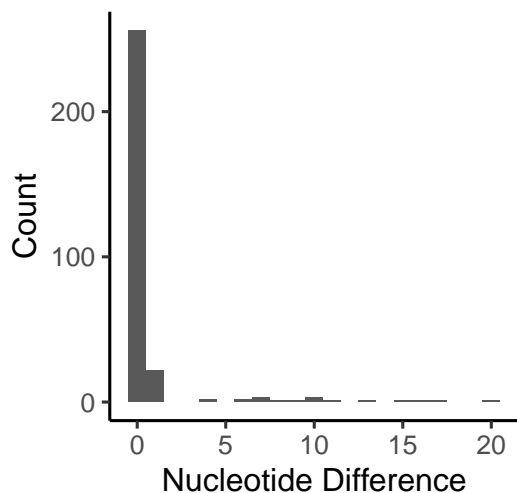
### IGHV1-46\*01

548 sequences assigned  
474 (86.5%) exact matches, in which:  
469 unique CDR3  
7 unique J



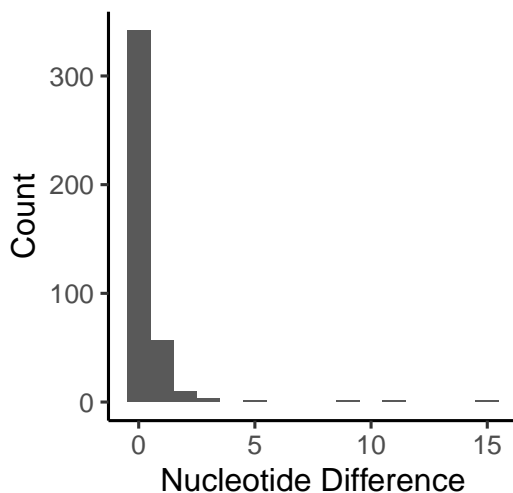
### IGHV1-3\*01

298 sequences assigned  
256 (85.9%) exact matches, in which:  
256 unique CDR3  
6 unique J



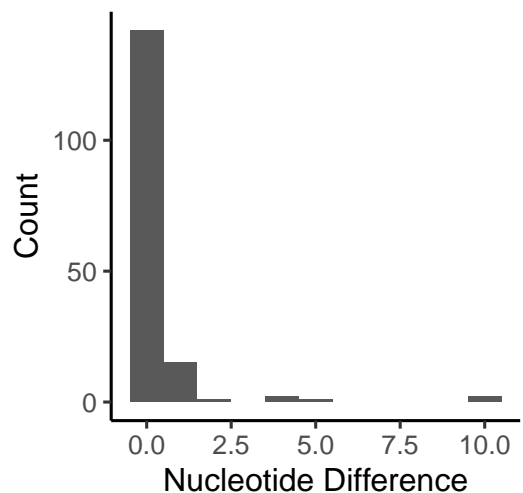
### IGHV1-24\*01

416 sequences assigned  
342 (82.2%) exact matches, in which:  
340 unique CDR3  
7 unique J



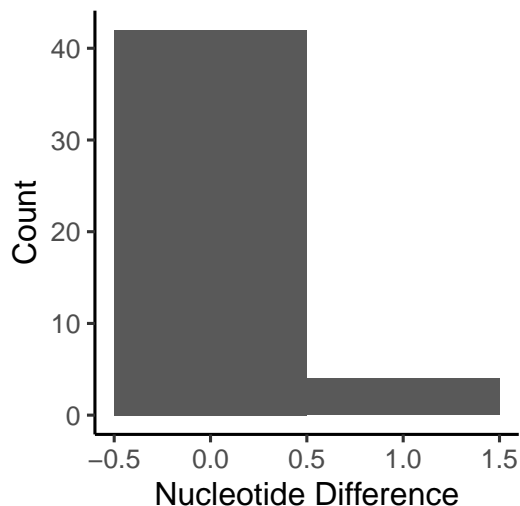
### IGHV1-58\*01

165 sequences assigned  
142 (86.1%) exact matches, in which:  
142 unique CDR3  
6 unique J



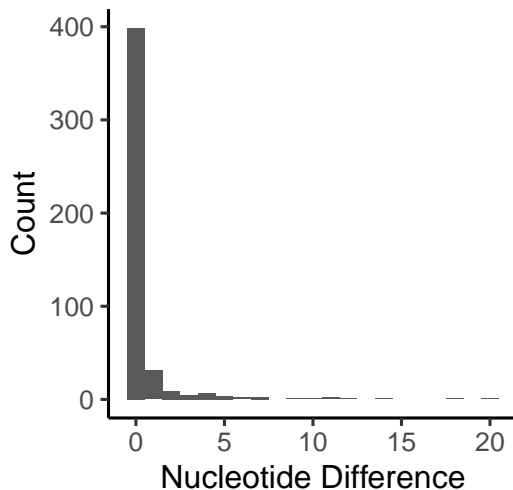
### IGHV1-69-2\*01

46 sequences assigned  
42 (91.3%) exact matches, in which:  
42 unique CDR3  
6 unique J



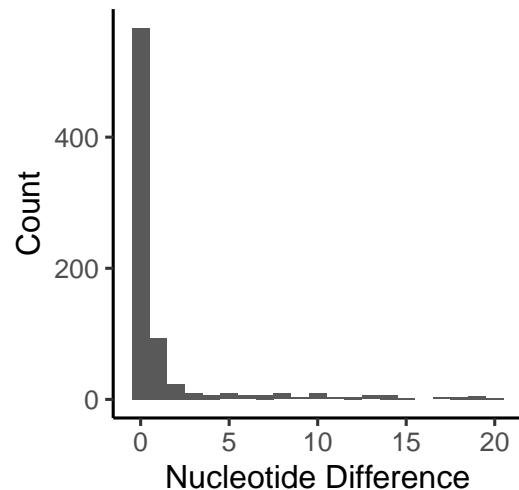
### IGHV2-5\*02

469 sequences assigned  
399 (85.1%) exact matches, in which:  
393 unique CDR3  
6 unique J



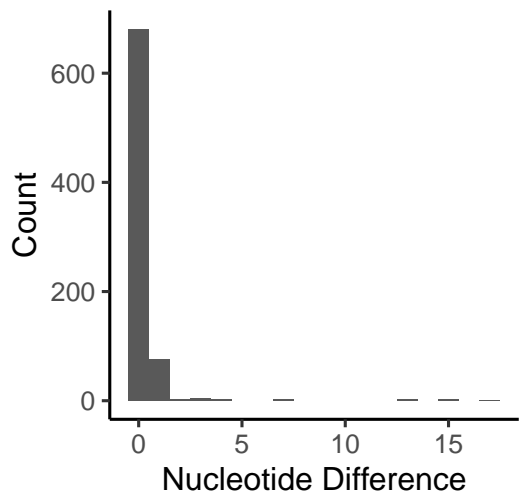
### IGHV3-7\*01

789 sequences assigned  
567 (71.9%) exact matches, in which:  
565 unique CDR3  
7 unique J



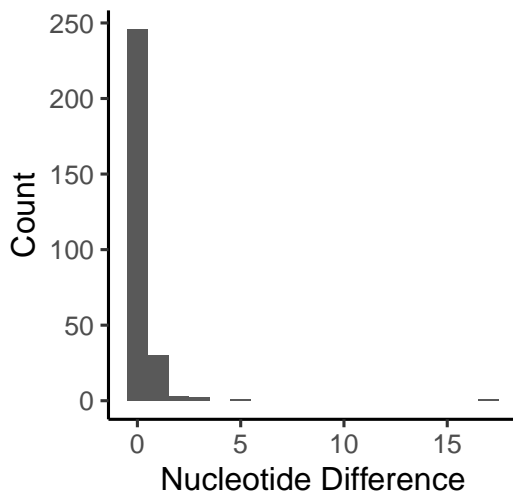
### IGHV1-69\*06

777 sequences assigned  
681 (87.6%) exact matches, in which:  
670 unique CDR3  
7 unique J



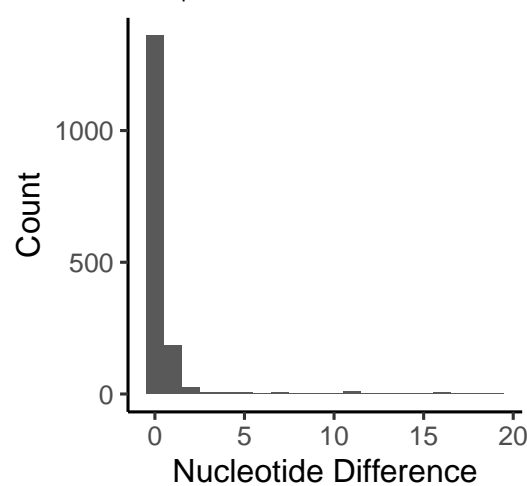
### IGHV2-26\*01

283 sequences assigned  
246 (86.9%) exact matches, in which:  
246 unique CDR3  
7 unique J



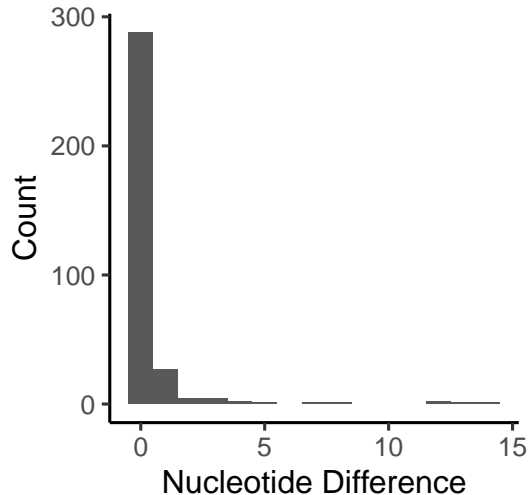
### IGHV3-9\*01

1629 sequences assigned  
1360 (83.5%) exact matches, in which:  
1349 unique CDR3  
7 unique J



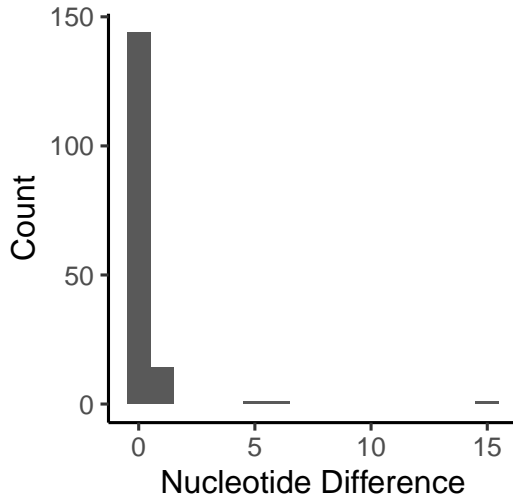
### IGHV2-5\*01

332 sequences assigned  
288 (86.7%) exact matches, in which:  
285 unique CDR3  
6 unique J



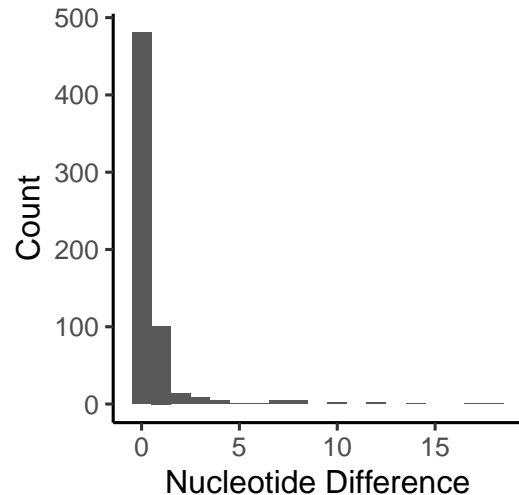
### IGHV2-70\*01

161 sequences assigned  
144 (89.4%) exact matches, in which:  
144 unique CDR3  
6 unique J



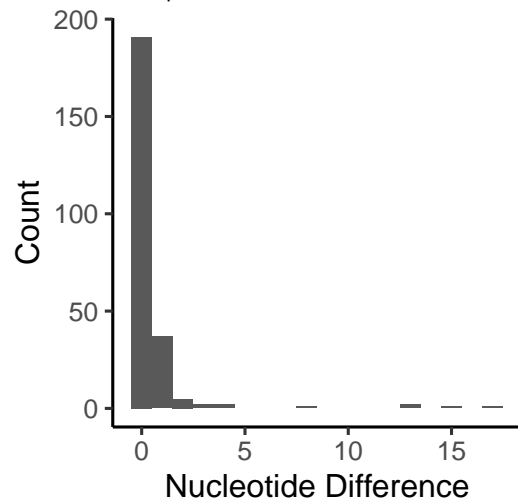
### IGHV3-11\*01

628 sequences assigned  
481 (76.6%) exact matches, in which:  
477 unique CDR3  
7 unique J



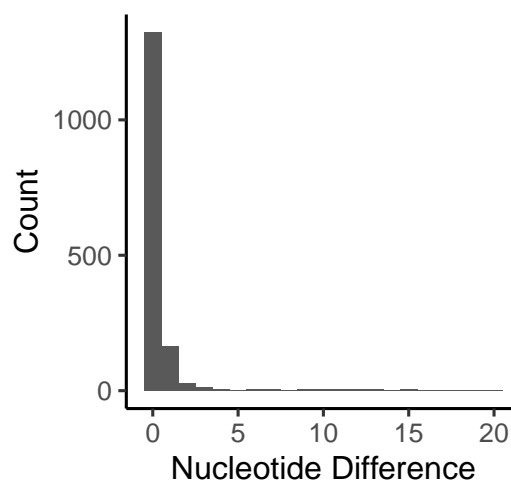
### IGHV3-13\*01

242 sequences assigned  
191 (78.9%) exact matches, in which:  
190 unique CDR3  
7 unique J



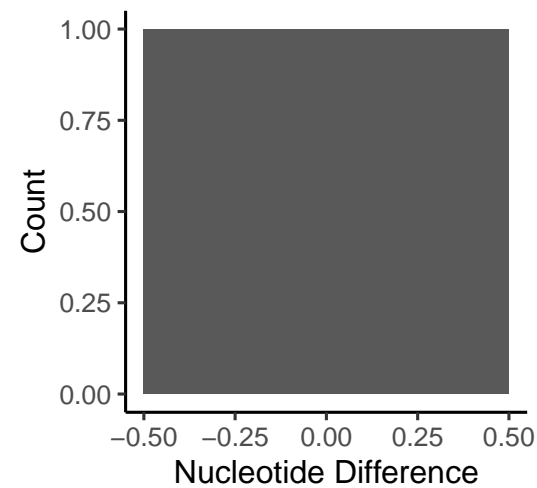
### IGHV3-21\*01

1588 sequences assigned  
1323 (83.3%) exact matches, in which:  
1316 unique CDR3  
7 unique J



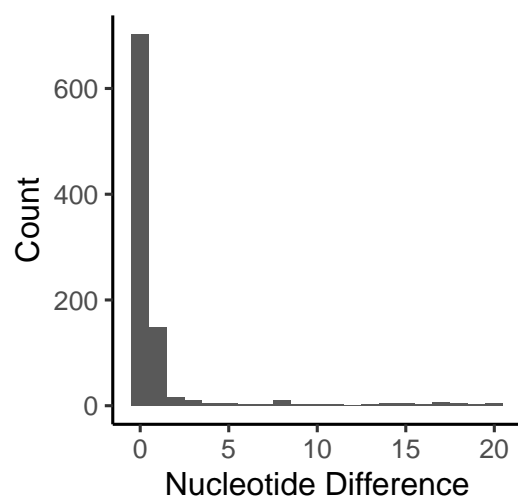
### IGHV3-35\*01

1 sequences assigned  
1 (100%) exact matches, in which:  
1 unique CDR3  
1 unique J



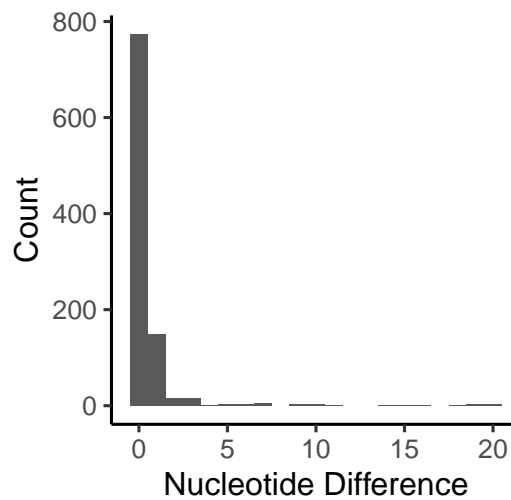
### IGHV3-15\*01

969 sequences assigned  
703 (72.5%) exact matches, in which:  
701 unique CDR3  
7 unique J



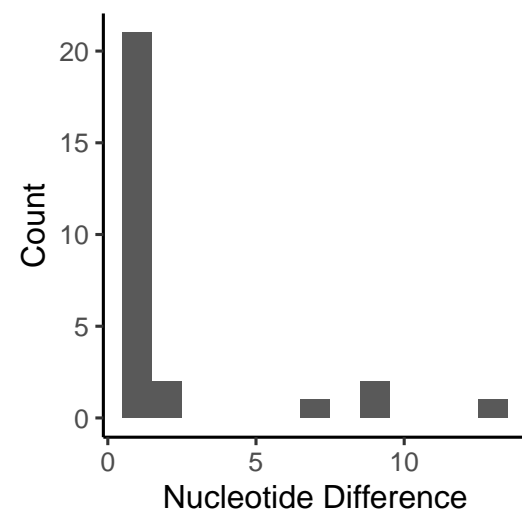
### IGHV3-30-3\*01

989 sequences assigned  
774 (78.3%) exact matches, in which:  
767 unique CDR3  
7 unique J



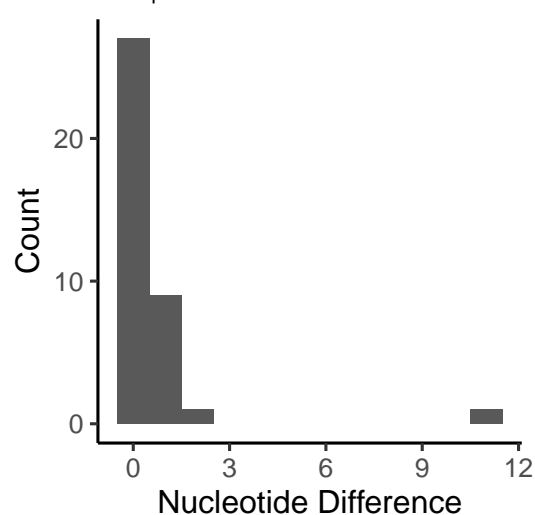
### IGHV3-43\*01

27 sequences assigned  
No exact matches.



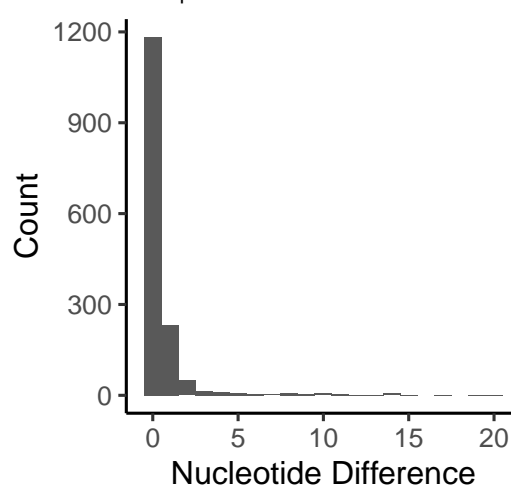
### IGHV3-20\*01

38 sequences assigned  
27 (71.1%) exact matches, in which:  
27 unique CDR3  
4 unique J



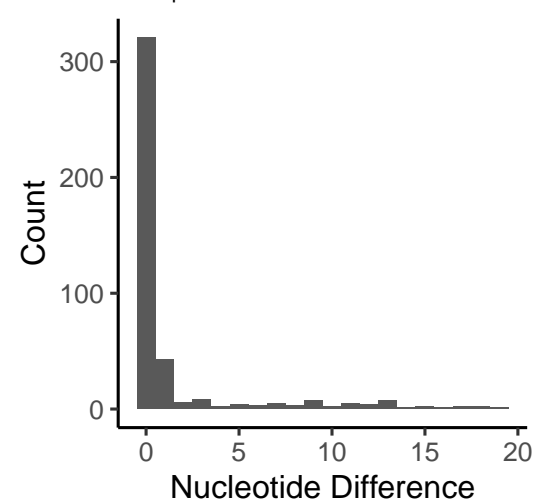
### IGHV3-33\*01

1548 sequences assigned  
1183 (76.4%) exact matches, in which:  
1172 unique CDR3  
7 unique J



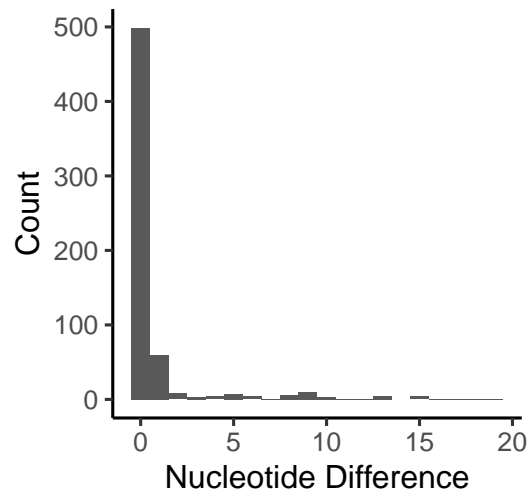
### IGHV3-48\*02

430 sequences assigned  
321 (74.7%) exact matches, in which:  
320 unique CDR3  
6 unique J



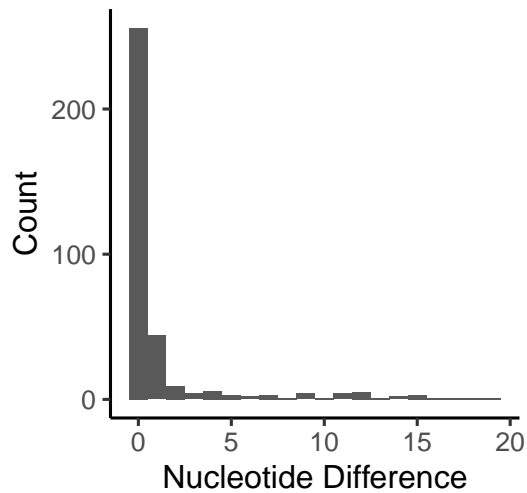
### IGHV3-48\*03

621 sequences assigned  
499 (80.4%) exact matches, in which:  
495 unique CDR3  
6 unique J



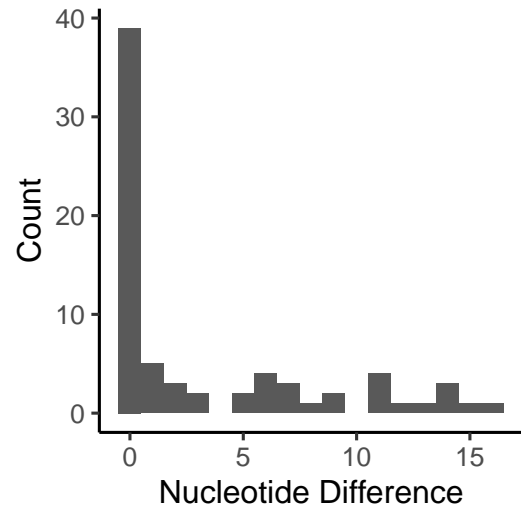
### IGHV3-53\*01

355 sequences assigned  
256 (72.1%) exact matches, in which:  
255 unique CDR3  
6 unique J



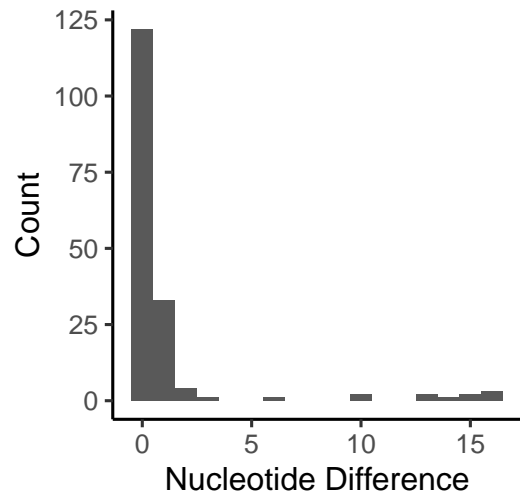
### IGHV3-72\*01

73 sequences assigned  
39 (53.4%) exact matches, in which:  
39 unique CDR3  
7 unique J



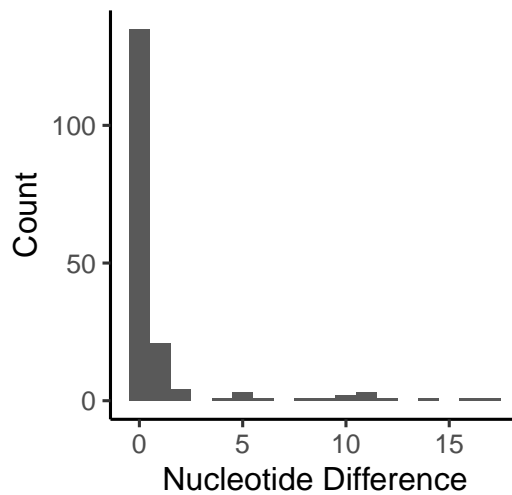
### IGHV3-49\*04

173 sequences assigned  
122 (70.5%) exact matches, in which:  
121 unique CDR3  
6 unique J



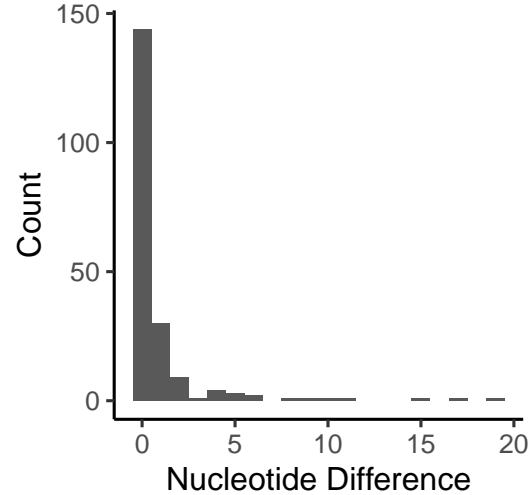
### IGHV3-53\*02

176 sequences assigned  
135 (76.7%) exact matches, in which:  
135 unique CDR3  
6 unique J



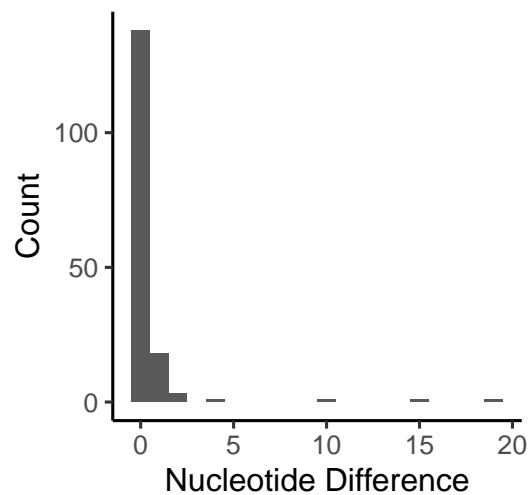
### IGHV3-73\*02

200 sequences assigned  
144 (72%) exact matches, in which:  
143 unique CDR3  
7 unique J



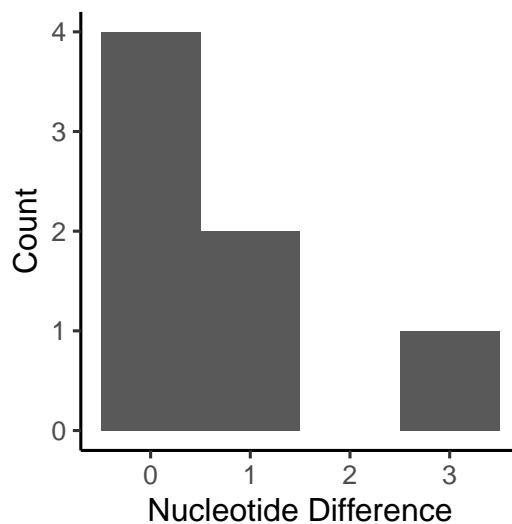
### IGHV3-49\*05

166 sequences assigned  
138 (83.1%) exact matches, in which:  
136 unique CDR3  
6 unique J



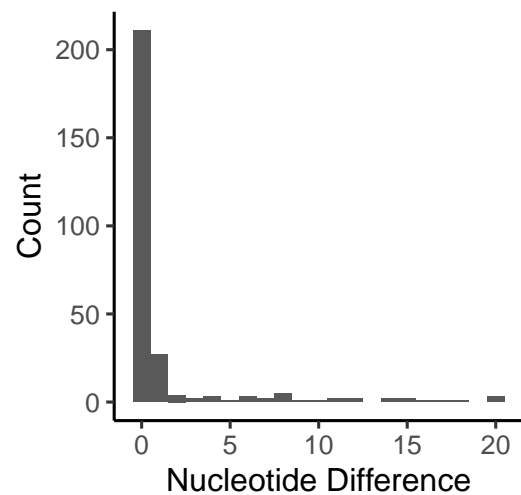
### IGHV3-64\*02

7 sequences assigned  
4 (57.1%) exact matches, in which:  
4 unique CDR3  
2 unique J



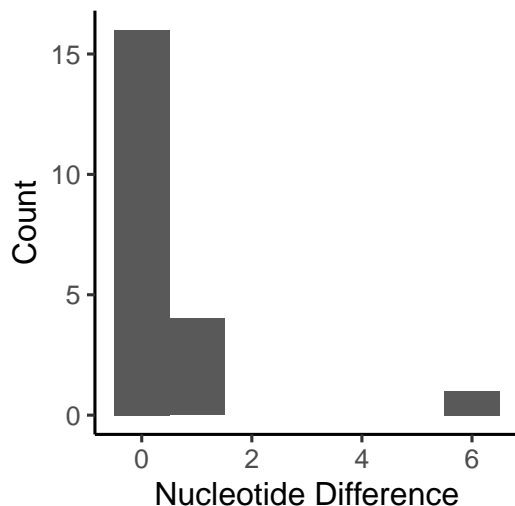
### IGHV3-74\*01

280 sequences assigned  
211 (75.4%) exact matches, in which:  
211 unique CDR3  
7 unique J



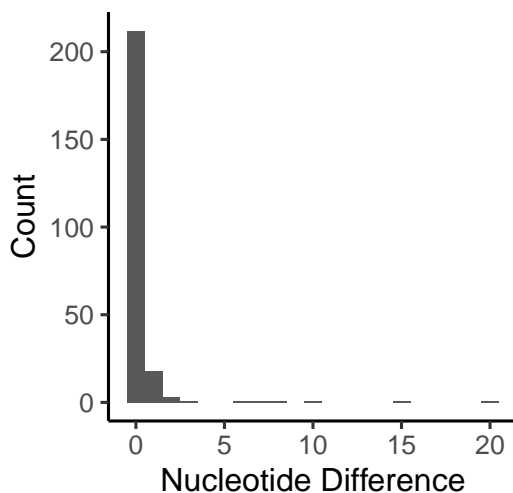
### IGHV3-43D\*04

21 sequences assigned  
16 (76.2%) exact matches, in which:  
16 unique CDR3  
6 unique J



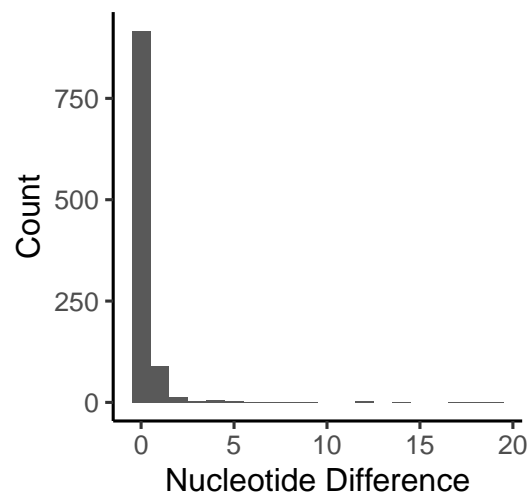
### IGHV4-30-4\*01

240 sequences assigned  
212 (88.3%) exact matches, in which:  
210 unique CDR3  
7 unique J



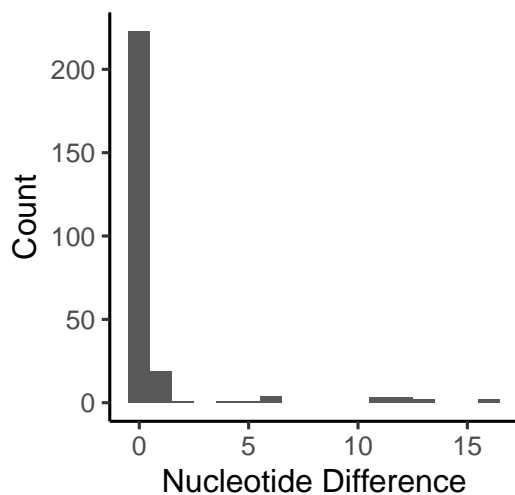
### IGHV4-34\*01

1052 sequences assigned  
917 (87.2%) exact matches, in which:  
911 unique CDR3  
7 unique J



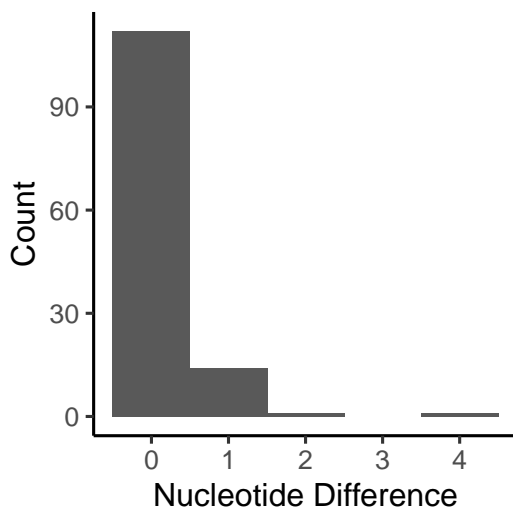
### IGHV4-4\*02

260 sequences assigned  
223 (85.8%) exact matches, in which:  
222 unique CDR3  
6 unique J



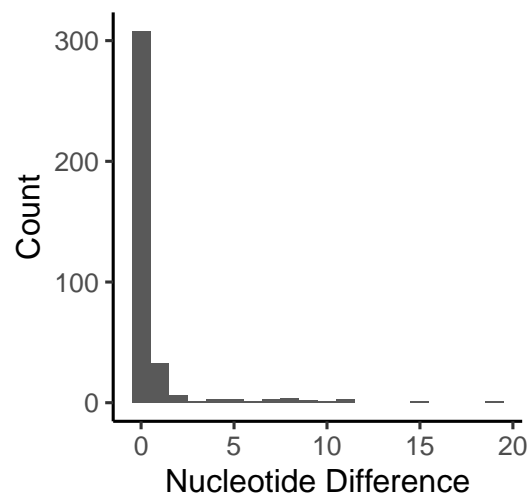
### IGHV4-30-2\*01

128 sequences assigned  
112 (87.5%) exact matches, in which:  
110 unique CDR3  
5 unique J



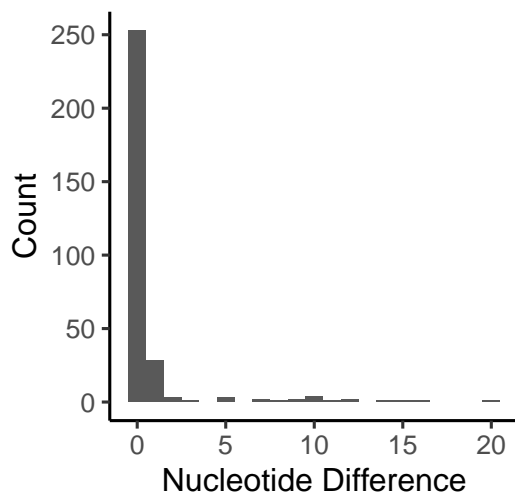
### IGHV4-38-2\*02\_A70G

371 sequences assigned  
308 (83%) exact matches, in which:  
302 unique CDR3  
6 unique J



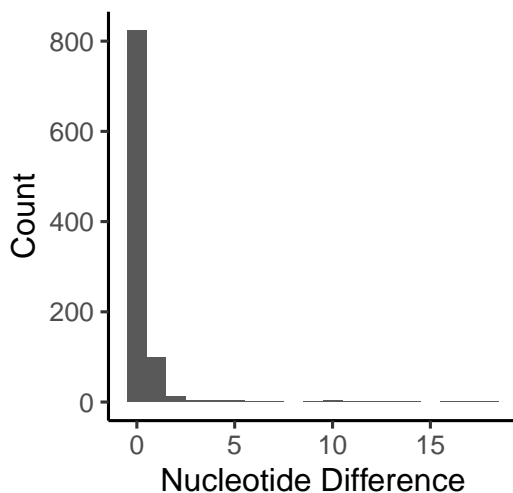
### IGHV4-4\*07

307 sequences assigned  
253 (82.4%) exact matches, in which:  
252 unique CDR3  
6 unique J



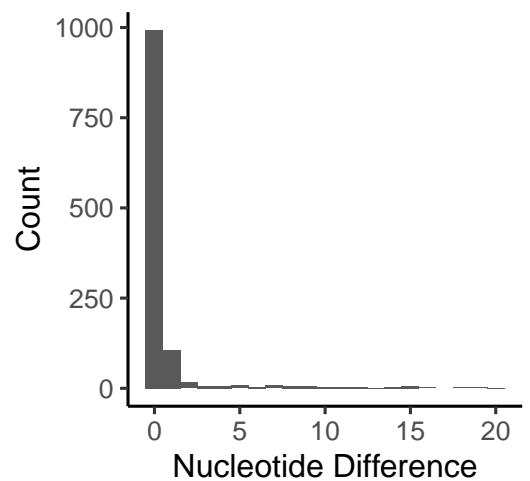
### IGHV4-31\*03

965 sequences assigned  
824 (85.4%) exact matches, in which:  
817 unique CDR3  
7 unique J



### IGHV4-39\*01

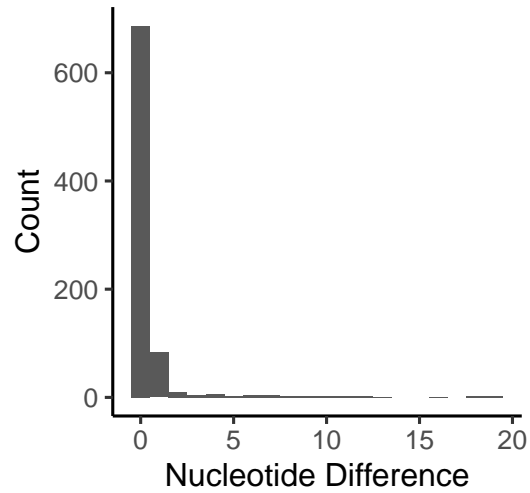
1197 sequences assigned  
993 (83%) exact matches, in which:  
988 unique CDR3  
7 unique J





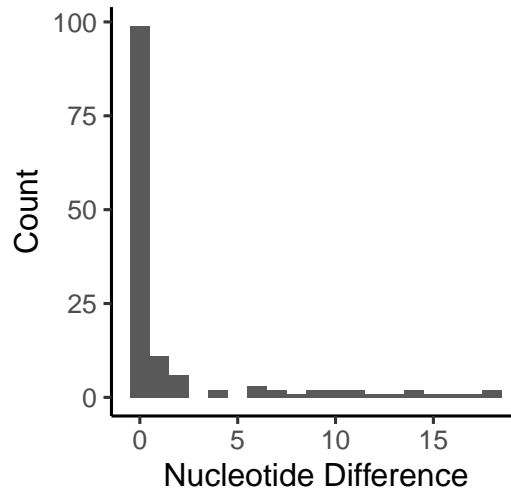
### IGHV4–59\*01

831 sequences assigned  
687 (82.7%) exact matches, in which:  
685 unique CDR3  
7 unique J



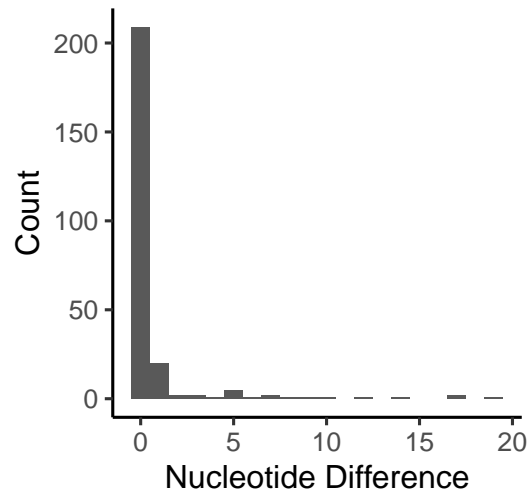
### IGHV6–1\*01

139 sequences assigned  
99 (71.2%) exact matches, in which:  
97 unique CDR3  
6 unique J



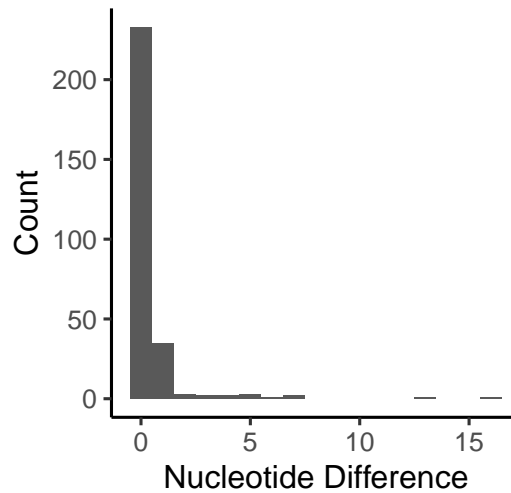
### IGHV4–61\*01

251 sequences assigned  
209 (83.3%) exact matches, in which:  
209 unique CDR3  
7 unique J



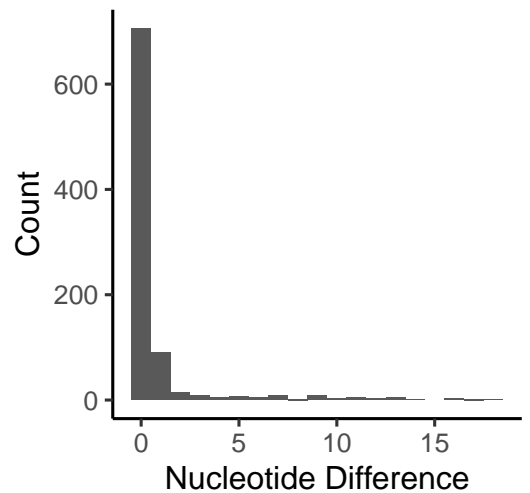
### IGHV7–4–1\*02

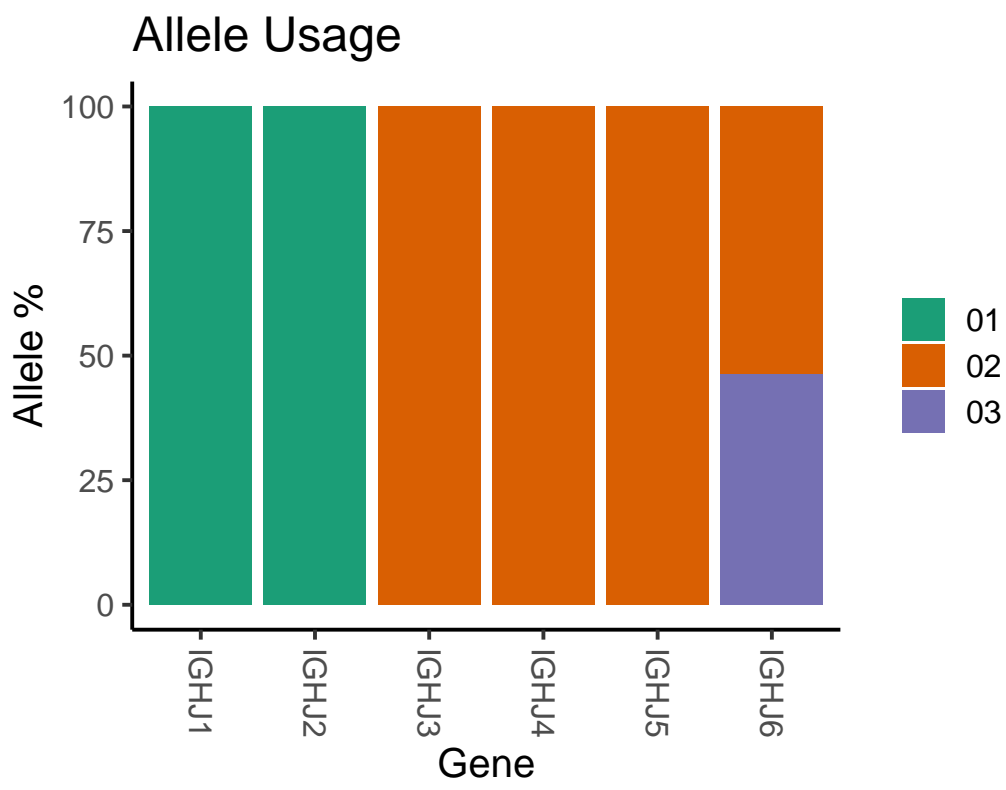
283 sequences assigned  
233 (82.3%) exact matches, in which:  
232 unique CDR3  
6 unique J



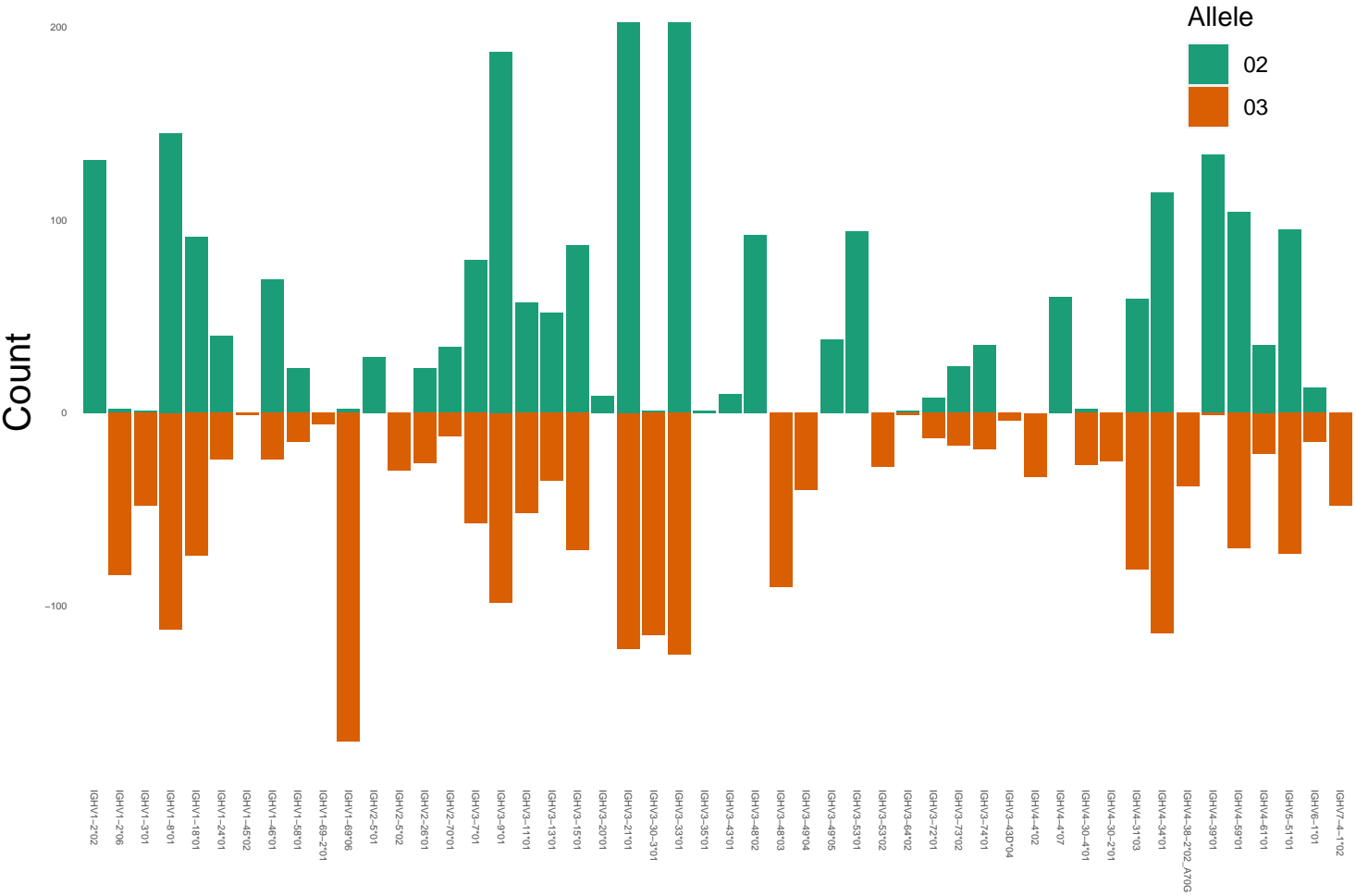
### IGHV5–51\*01

882 sequences assigned  
706 (80%) exact matches, in which:  
700 unique CDR3  
7 unique J





Sequence Count by IGHJ6 allele usage



Warning – no inferred sequences found.

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