

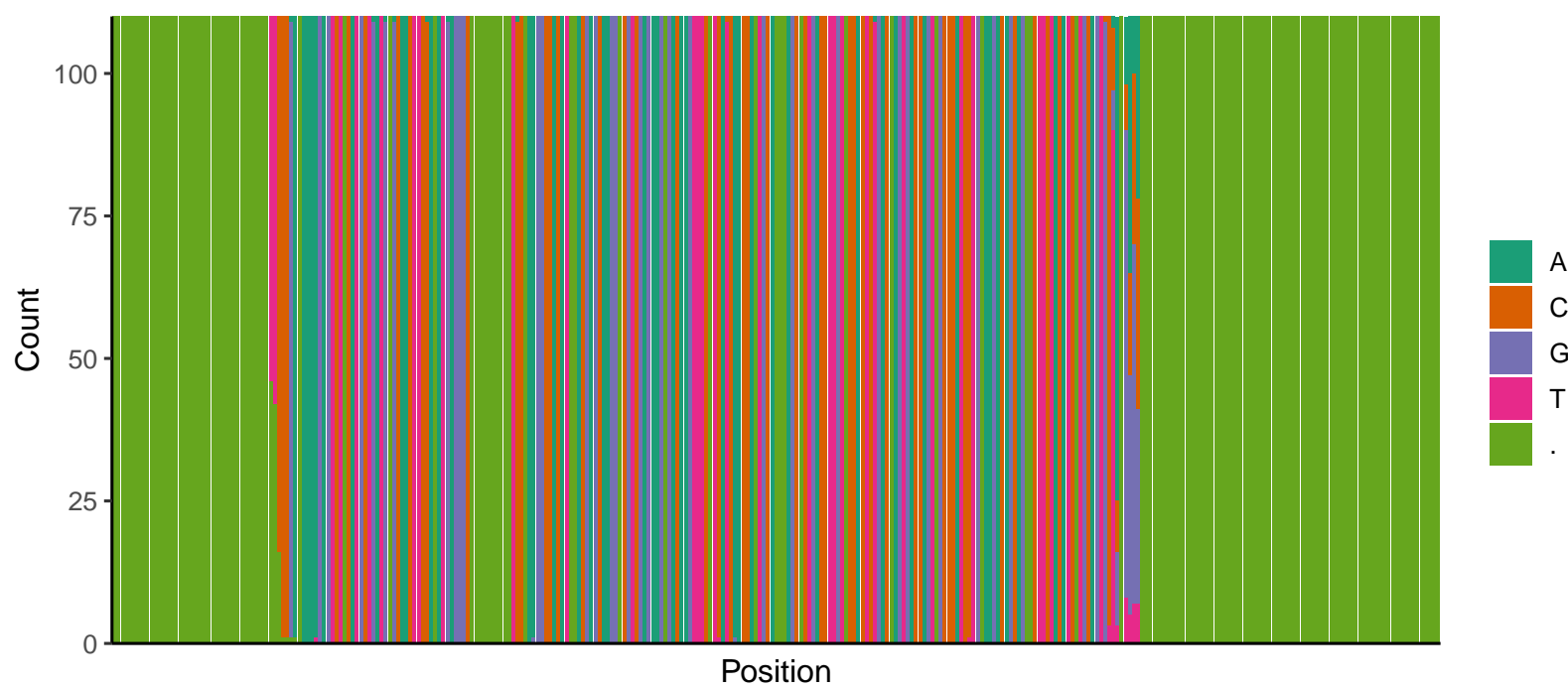
Gene TRBV10-3\*bp02\_313GCCATCAGTGAGTC326



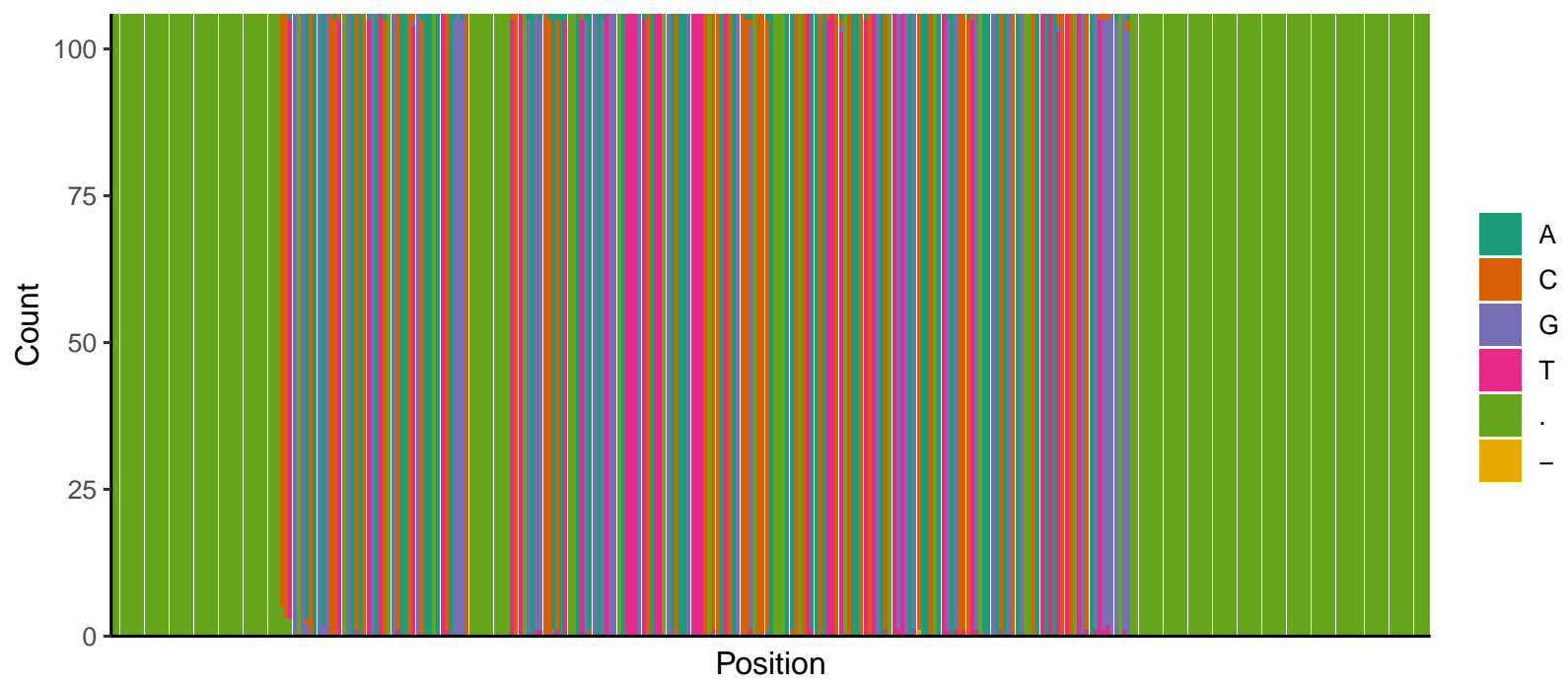
Gene TRBV14\*bp02\_322CAAGA326



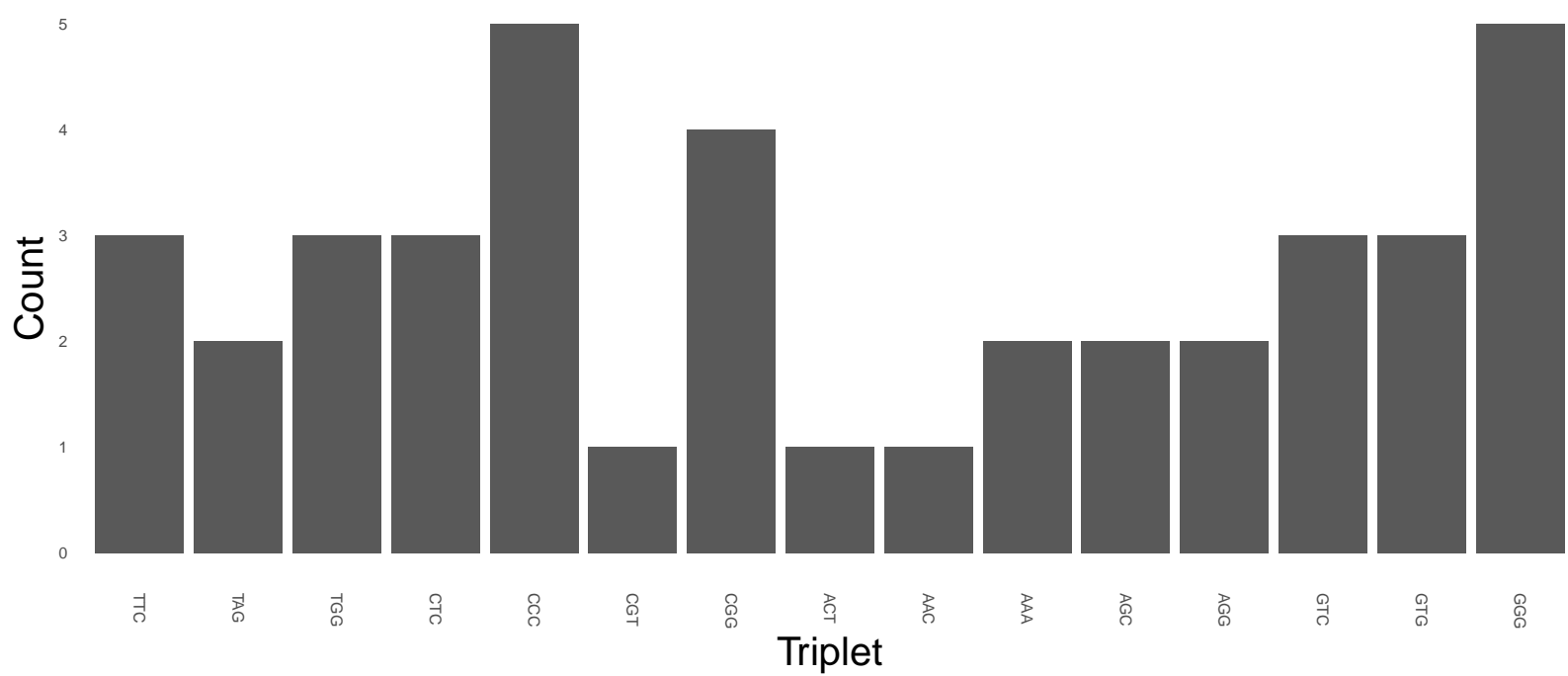
Gene TRBV20-1\*bp04\_322GA323



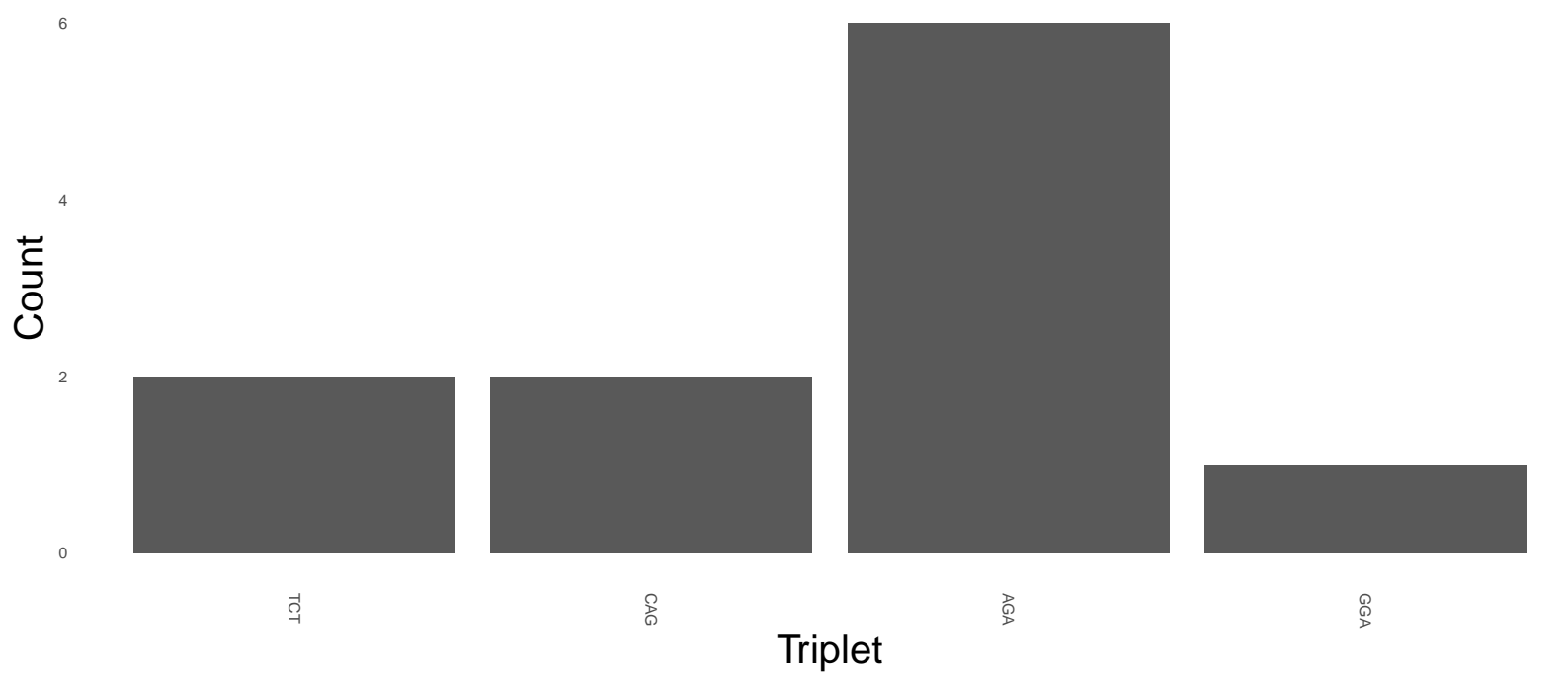
Gene TRBV29-1\*bp03\_C315T



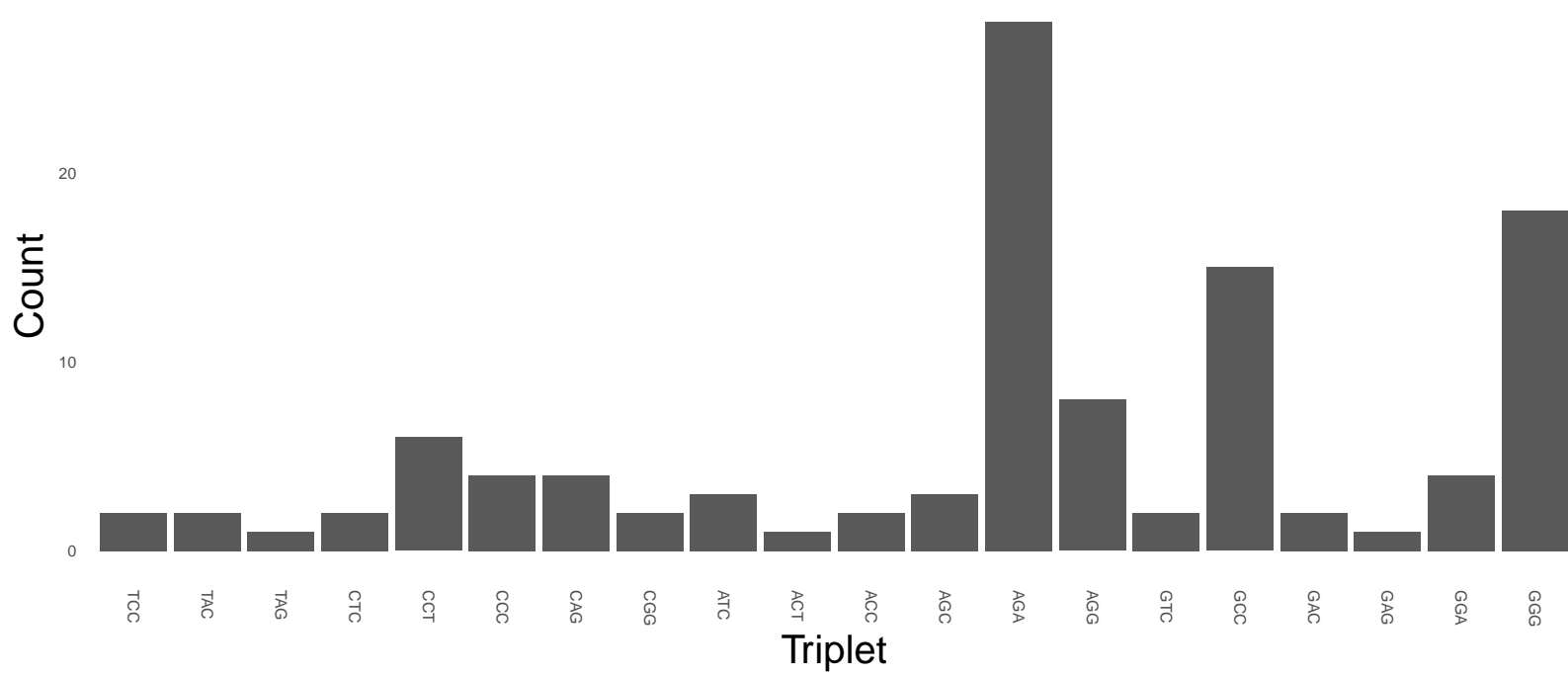
TRBV10–3\*bp02\_313GCCATCAGTGAGTC326– Final 3 nucleotides as a triplet



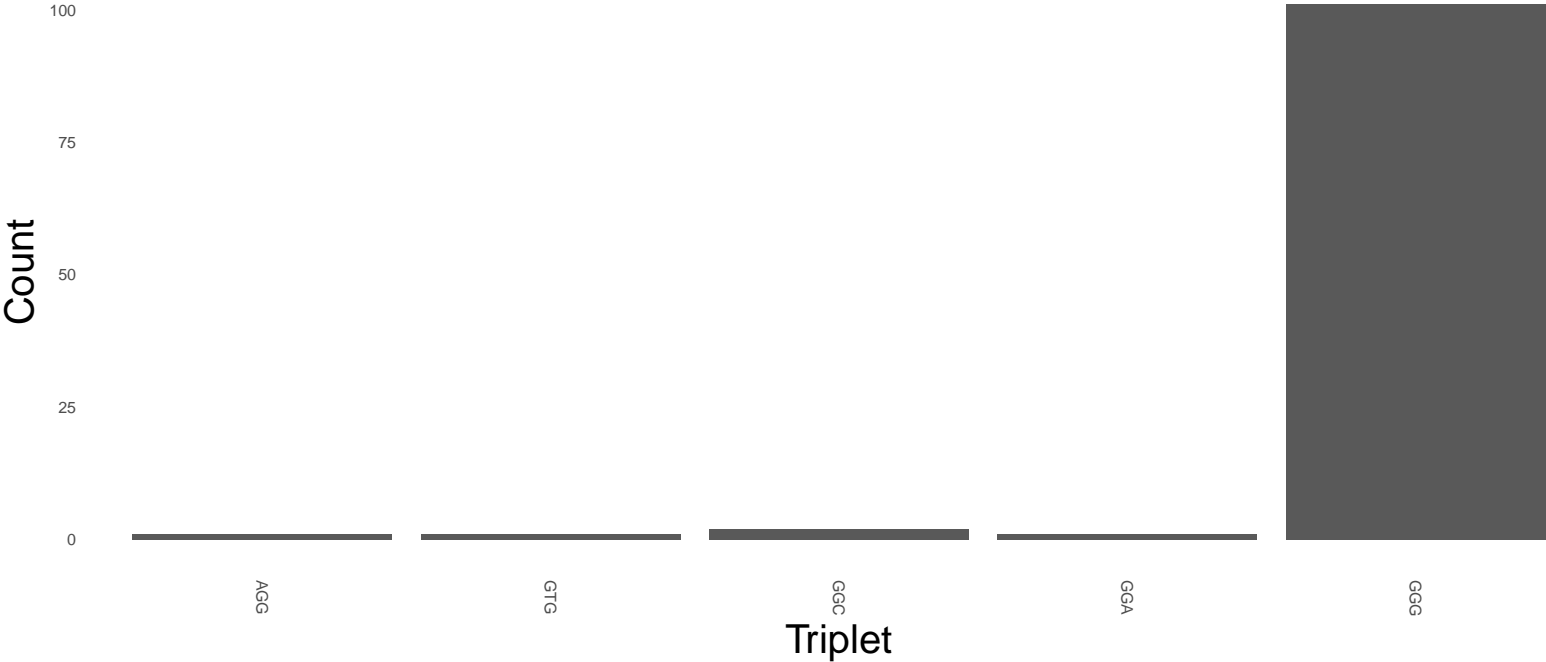
TRBV14\*bp02\_322CAAGA326– Final 3 nucleotides as a triplet



TRBV20–1\*bp04\_322GA323– Final 3 nucleotides as a triplet

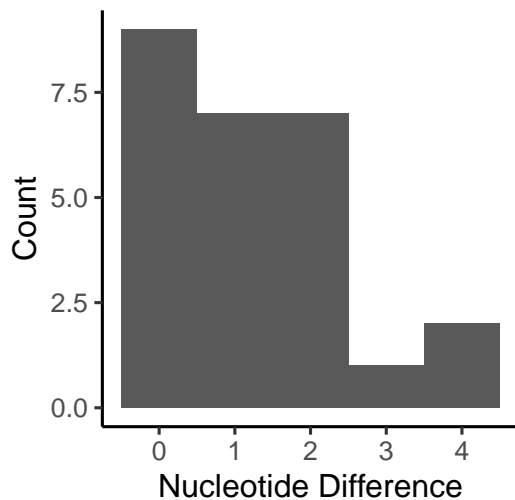


TRBV29-1\*bp03\_C315T- Final 3 nucleotides as a triplet



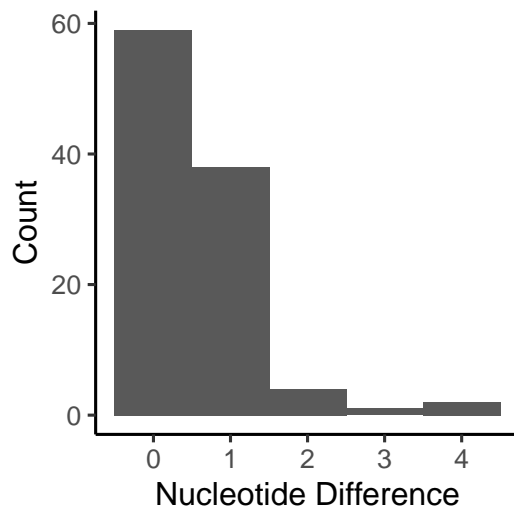
### TRBV10–1\*bp01

26 sequences assigned  
9 (34.6%) exact matches, in which:  
5 unique CDR3  
3 unique J



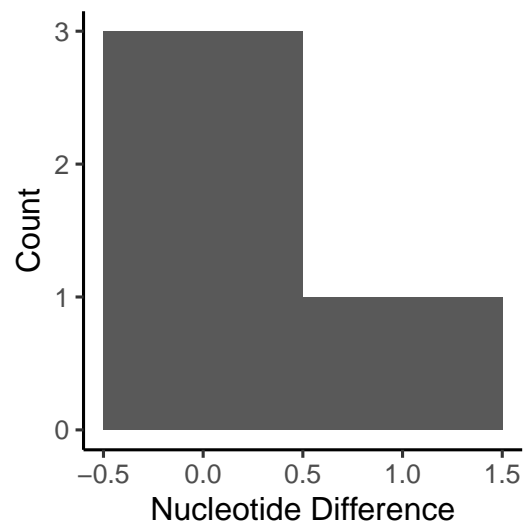
### TRBV10–3\*bp01

104 sequences assigned  
59 (56.7%) exact matches, in which:  
49 unique CDR3  
10 unique J



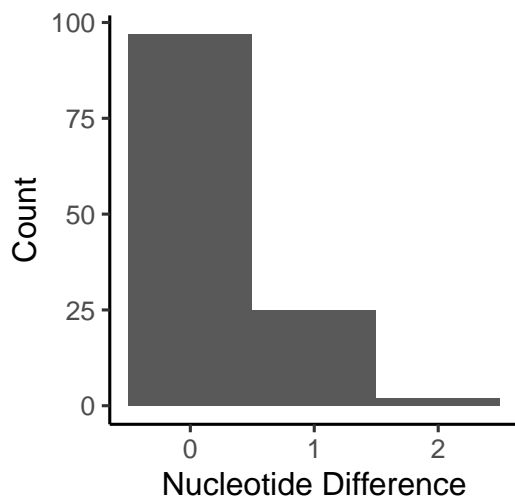
### TRBV14\*bp01

4 sequences assigned  
3 (75%) exact matches, in which:  
2 unique CDR3  
2 unique J



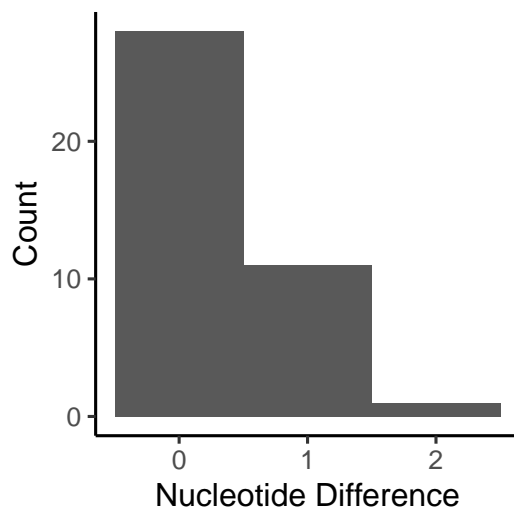
### TRBV20–1\*bp01

124 sequences assigned  
97 (78.2%) exact matches, in which:  
45 unique CDR3  
9 unique J



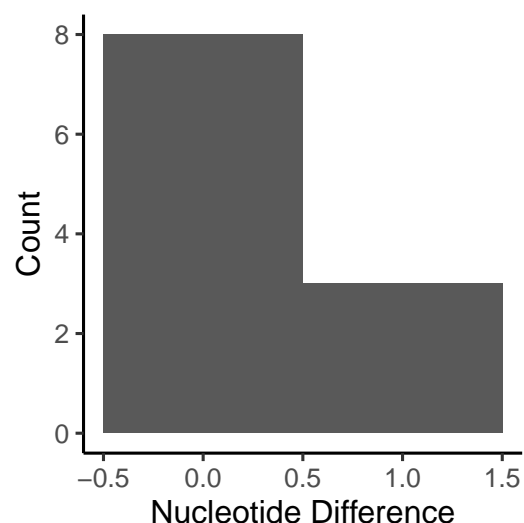
### TRBV10–3\*bp02\_313GCC

40 sequences assigned  
28 (70%) exact matches, in which:  
21 unique CDR3  
7 unique J



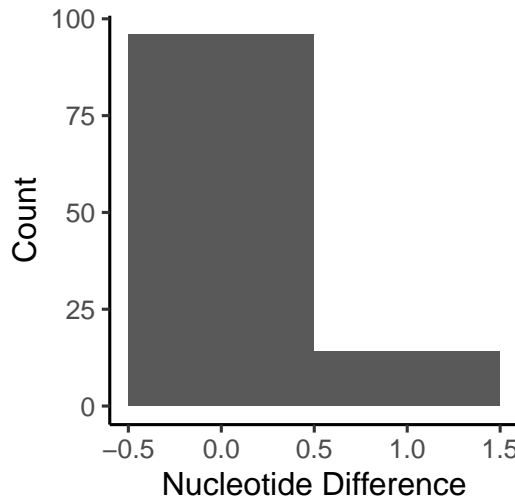
### TRBV14\*bp02\_322CAAGA

11 sequences assigned  
8 (72.7%) exact matches, in which:  
5 unique CDR3  
3 unique J



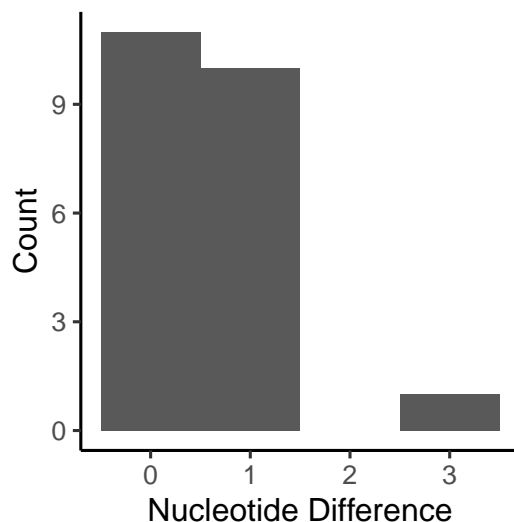
### TRBV20–1\*bp04\_322GA:

110 sequences assigned  
96 (87.3%) exact matches, in which:  
46 unique CDR3  
11 unique J



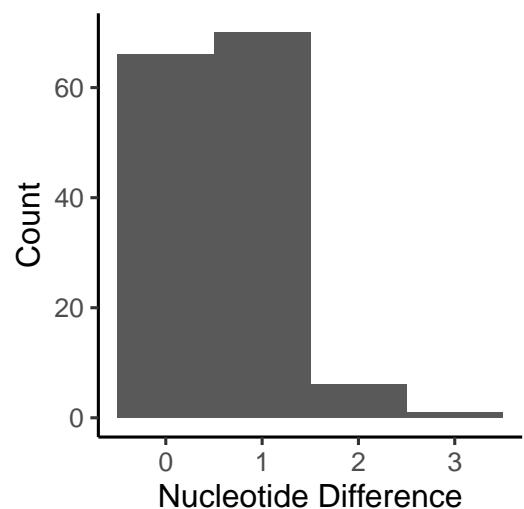
### TRBV13\*bp01

22 sequences assigned  
11 (50%) exact matches, in which:  
10 unique CDR3  
8 unique J



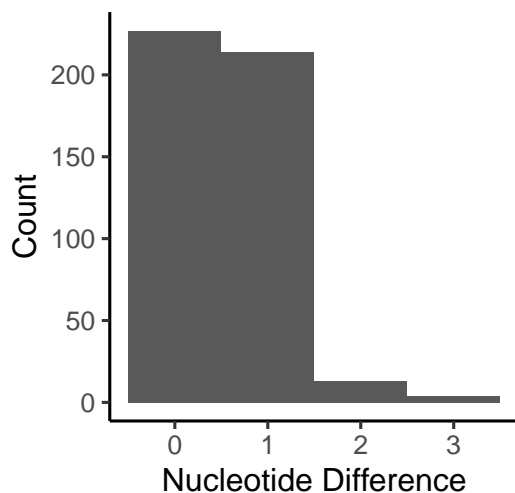
### TRBV15\*bp02

143 sequences assigned  
66 (46.2%) exact matches, in which:  
29 unique CDR3  
9 unique J



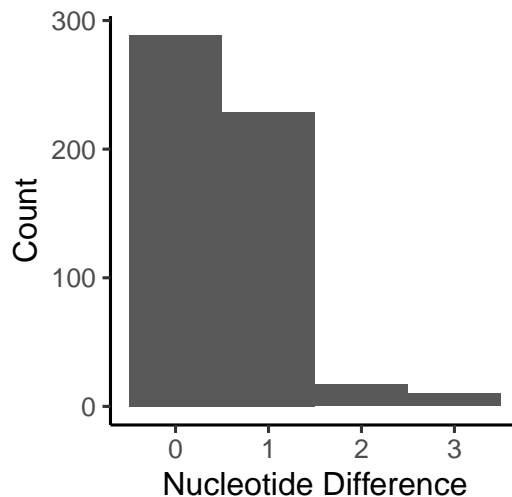
### TRBV18\*bp01

458 sequences assigned  
227 (49.6%) exact matches, in which:  
145 unique CDR3  
12 unique J



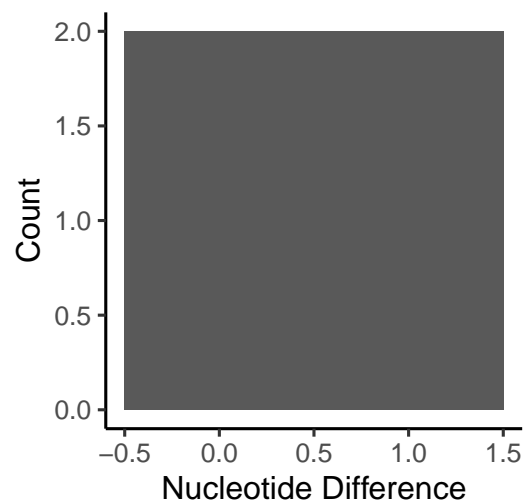
### TRBV27\*bp01

545 sequences assigned  
289 (53%) exact matches, in which:  
182 unique CDR3  
13 unique J



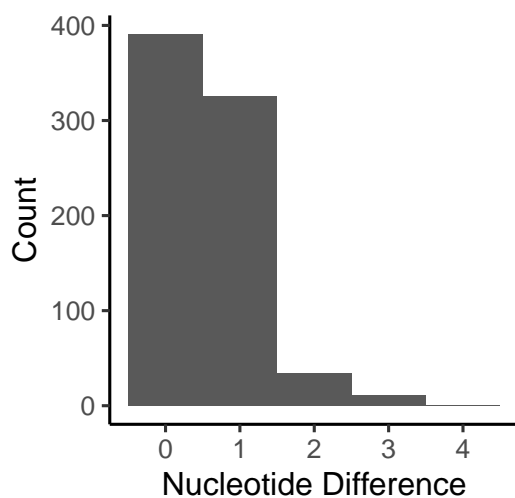
### TRBV9\*bp01

4 sequences assigned  
2 (50%) exact matches, in which:  
2 unique CDR3  
1 unique J



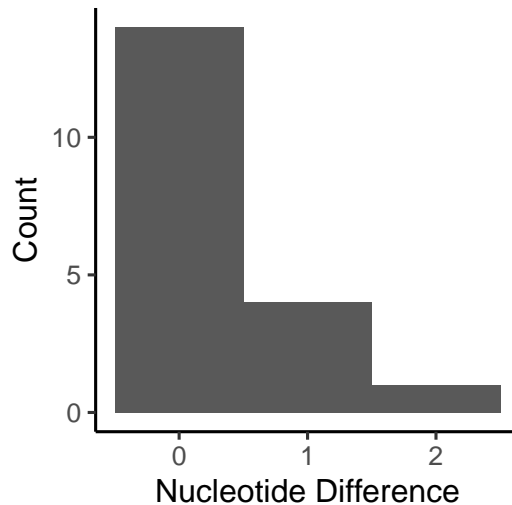
### TRBV19\*bp01

763 sequences assigned  
391 (51.2%) exact matches, in which:  
203 unique CDR3  
12 unique J



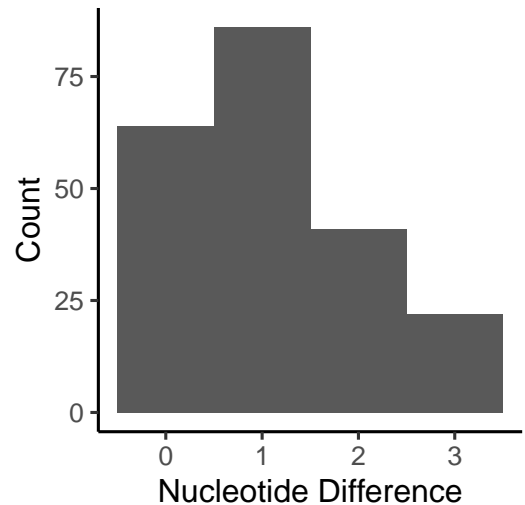
### TRBV28\*bp01

19 sequences assigned  
14 (73.7%) exact matches, in which:  
11 unique CDR3  
5 unique J



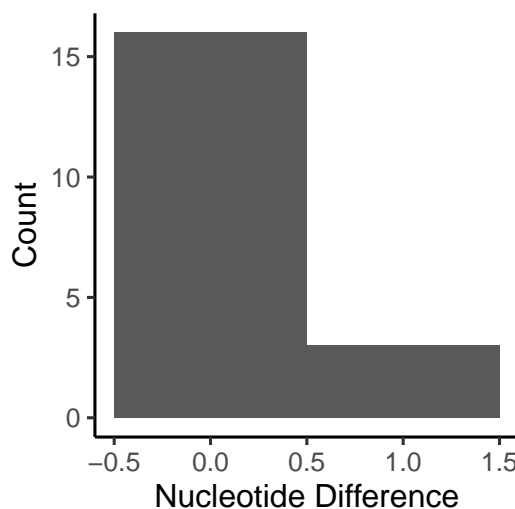
### TRBV11-1\*bp01

213 sequences assigned  
64 (30%) exact matches, in which:  
45 unique CDR3  
10 unique J



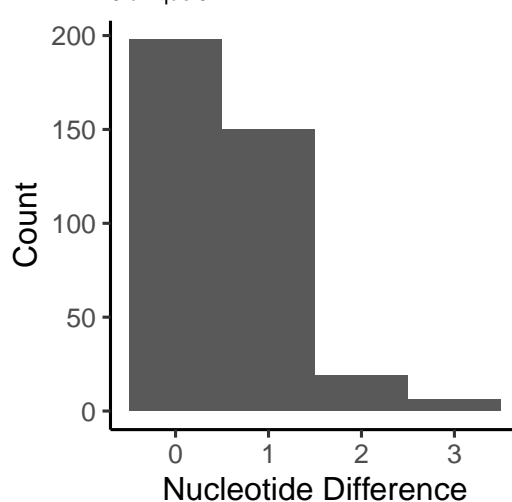
### TRBV2\*bp01

19 sequences assigned  
16 (84.2%) exact matches, in which:  
13 unique CDR3  
6 unique J



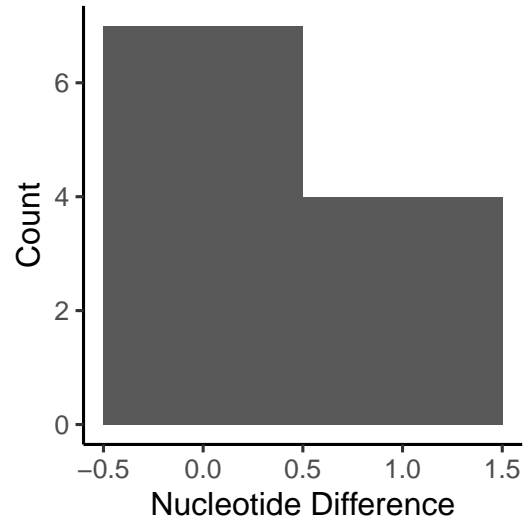
### TRBV30\*bp01

373 sequences assigned  
198 (53.1%) exact matches, in which:  
85 unique CDR3  
9 unique J



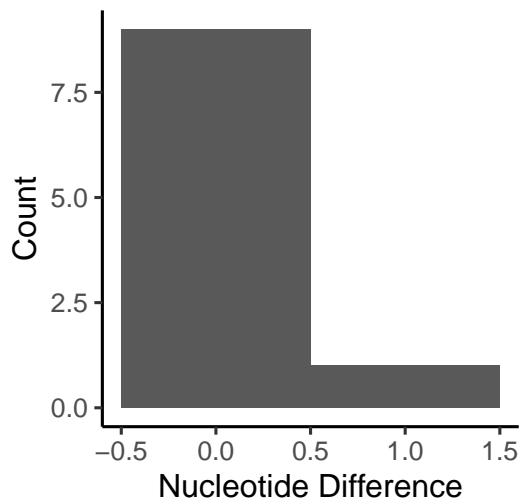
### TRBV21-1\*bp01

11 sequences assigned  
7 (63.6%) exact matches, in which:  
6 unique CDR3  
5 unique J



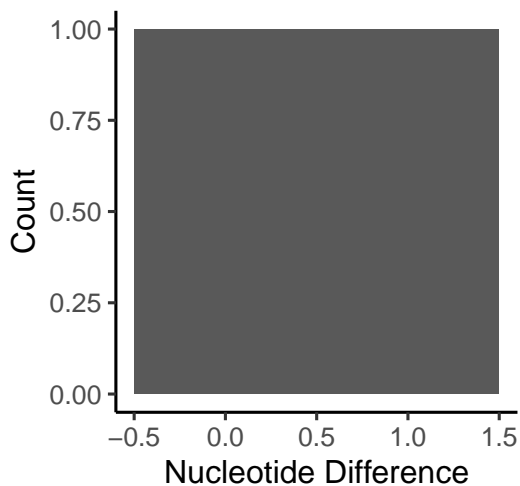
### TRBV21-1\*bp02

10 sequences assigned  
9 (90%) exact matches, in which:  
6 unique CDR3  
4 unique J



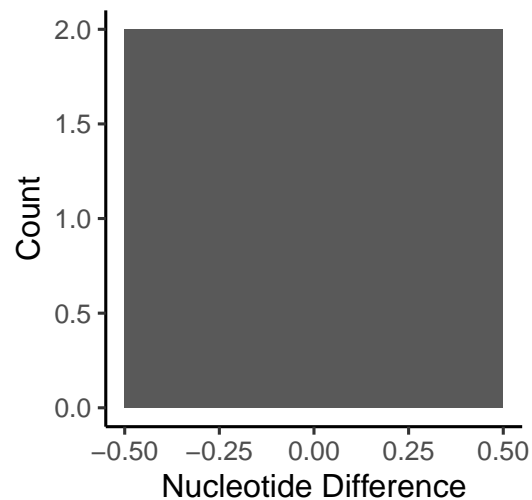
### TRBV12-5\*bp01

2 sequences assigned  
1 (50%) exact matches, in which:  
1 unique CDR3  
1 unique J



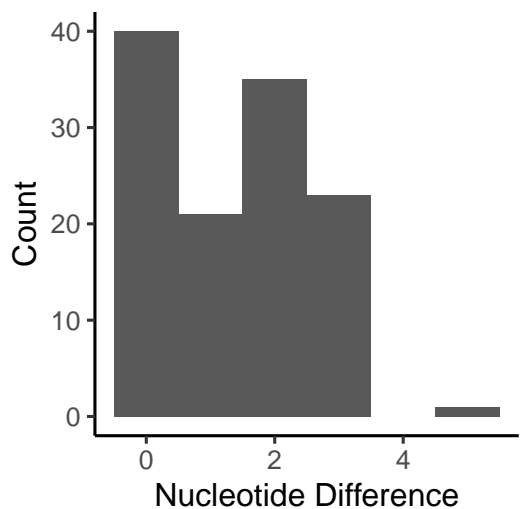
### TRBV4-1\*bp01

2 sequences assigned  
2 (100%) exact matches, in which:  
2 unique CDR3  
2 unique J



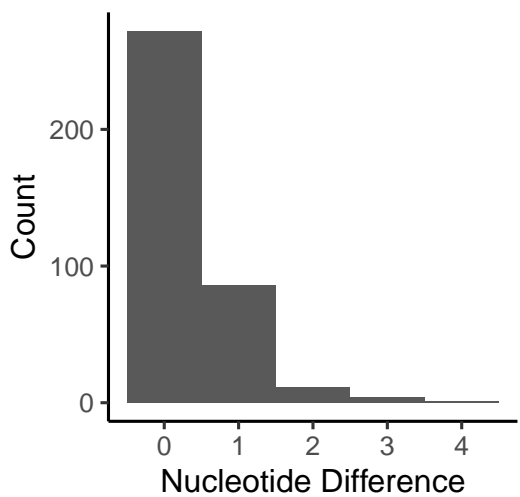
### TRBV11-2\*bp01

120 sequences assigned  
40 (33.3%) exact matches, in which:  
21 unique CDR3  
9 unique J



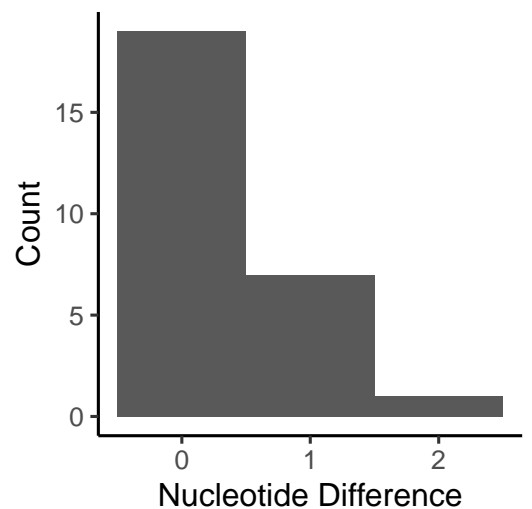
### TRBV12-34\*bp01

374 sequences assigned  
272 (72.7%) exact matches, in which:  
118 unique CDR3  
13 unique J



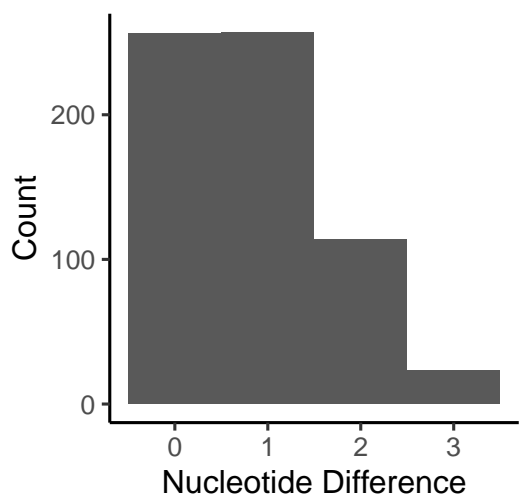
### TRBV4-2\*bp01

27 sequences assigned  
19 (70.4%) exact matches, in which:  
14 unique CDR3  
7 unique J



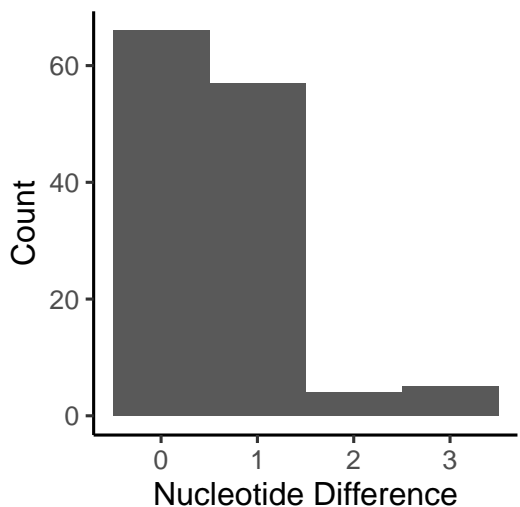
### TRBV11-3\*bp01

650 sequences assigned  
256 (39.4%) exact matches, in which:  
153 unique CDR3  
13 unique J



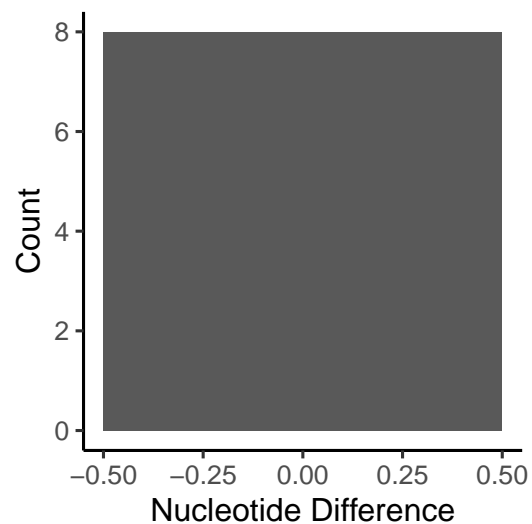
### TRBV24-1\*bp01

132 sequences assigned  
66 (50%) exact matches, in which:  
44 unique CDR3  
11 unique J



### TRBV4-3\*bp01

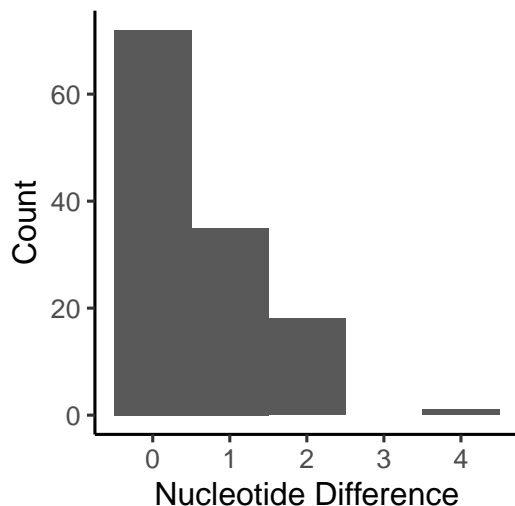
8 sequences assigned  
8 (100%) exact matches, in which:  
7 unique CDR3  
6 unique J





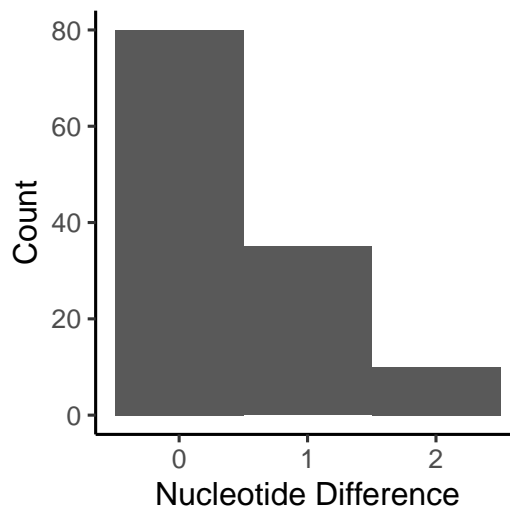
### TRBV5-1\*bp01

126 sequences assigned  
72 (57.1%) exact matches, in which:  
41 unique CDR3  
10 unique J



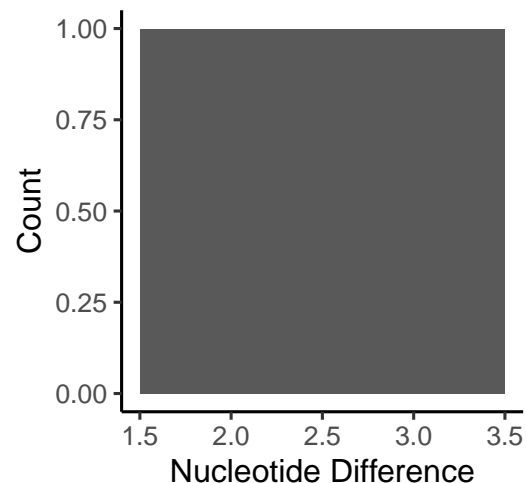
### TRBV5-4\*bp01

125 sequences assigned  
80 (64%) exact matches, in which:  
47 unique CDR3  
10 unique J



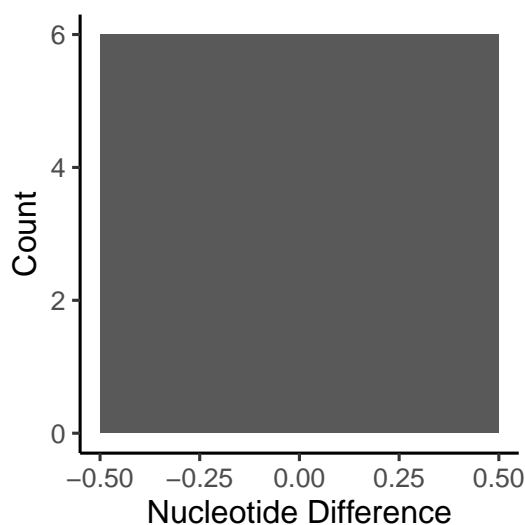
### TRBV5-7\*bp01

2 sequences assigned  
No exact matches.



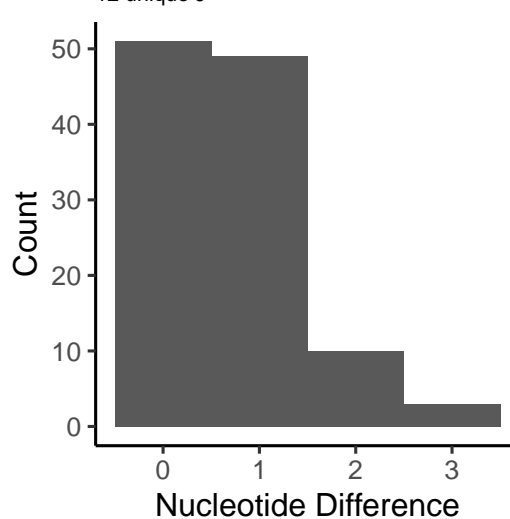
### TRBV25-1\*bp01

6 sequences assigned  
6 (100%) exact matches, in which:  
5 unique CDR3  
3 unique J



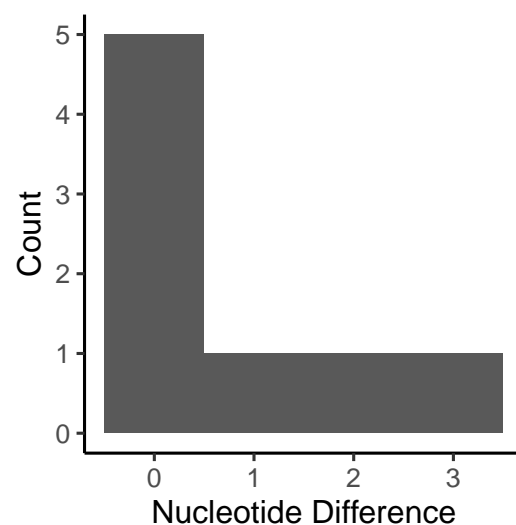
### TRBV5-5\*bp01

113 sequences assigned  
51 (45.1%) exact matches, in which:  
31 unique CDR3  
12 unique J



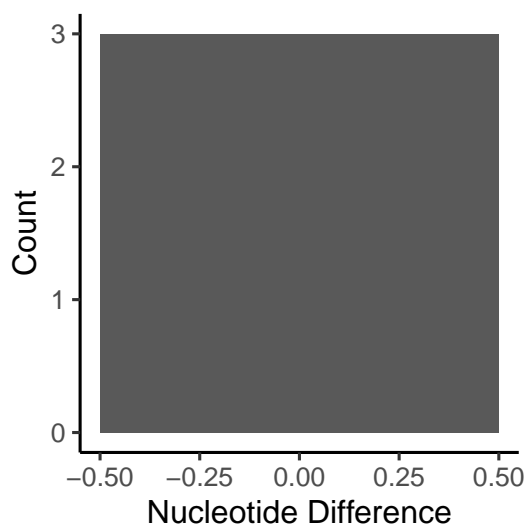
### TRBV5-8\*bp01

8 sequences assigned  
5 (62.5%) exact matches, in which:  
4 unique CDR3  
4 unique J



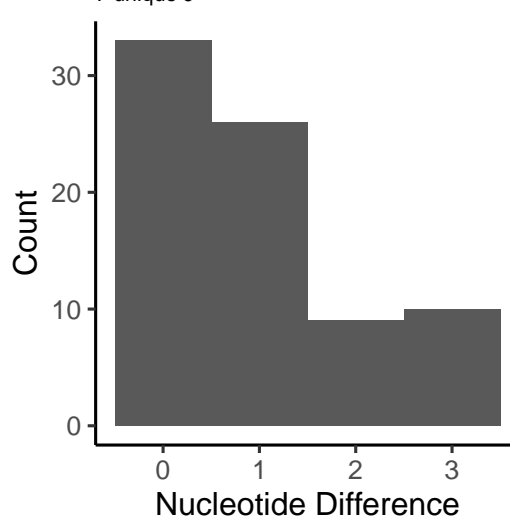
### TRBV5-3\*bp01

3 sequences assigned  
3 (100%) exact matches, in which:  
2 unique CDR3  
1 unique J



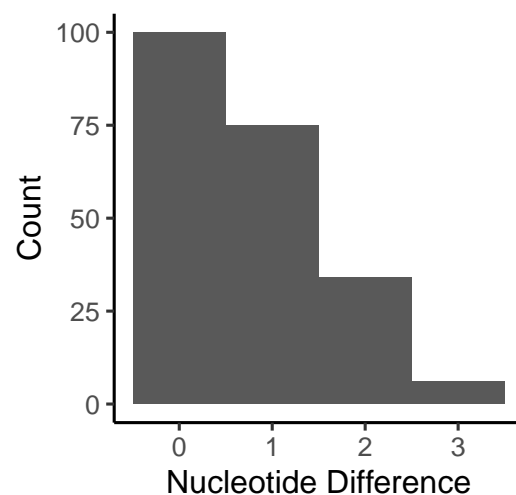
### TRBV5-6\*bp01

78 sequences assigned  
33 (42.3%) exact matches, in which:  
23 unique CDR3  
7 unique J



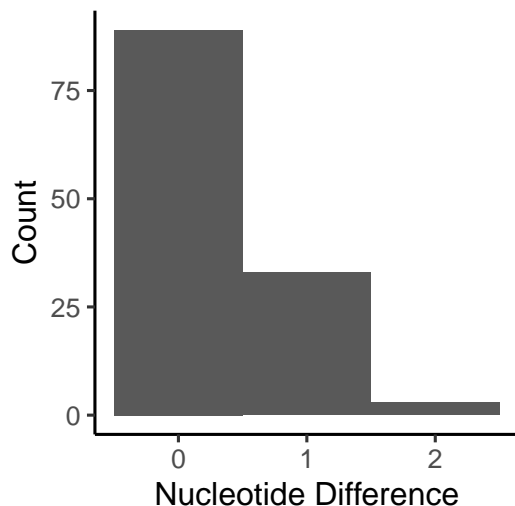
### TRBV6-1\*bp01

215 sequences assigned  
100 (46.5%) exact matches, in which:  
71 unique CDR3  
13 unique J



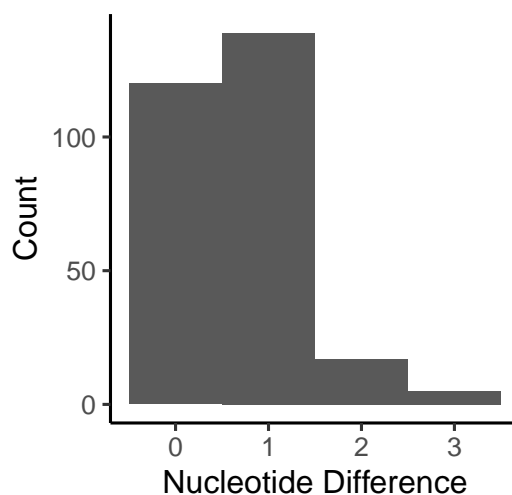
### TRBV6-4\*bp01

125 sequences assigned  
89 (71.2%) exact matches, in which:  
58 unique CDR3  
10 unique J



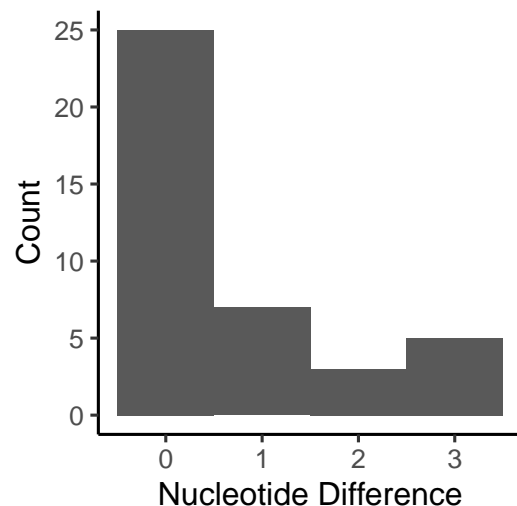
### TRBV6-23\*bp01

281 sequences assigned  
120 (42.7%) exact matches, in which:  
84 unique CDR3  
9 unique J



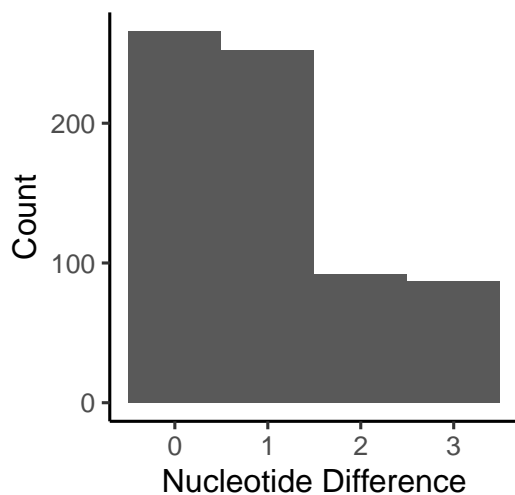
### TRBV7-3\*bp01

40 sequences assigned  
25 (62.5%) exact matches, in which:  
21 unique CDR3  
9 unique J



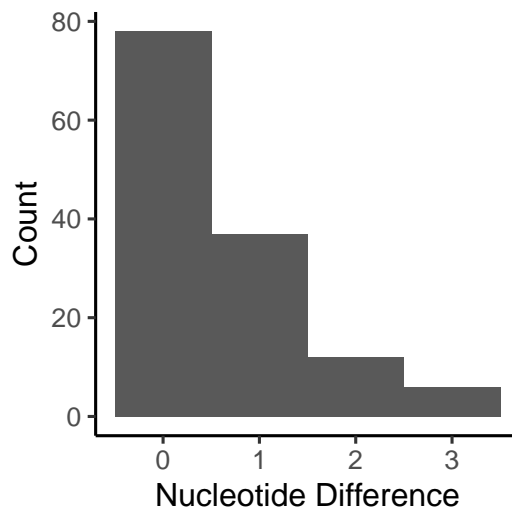
### TRBV6-5\*bp01

698 sequences assigned  
266 (38.1%) exact matches, in which:  
176 unique CDR3  
12 unique J



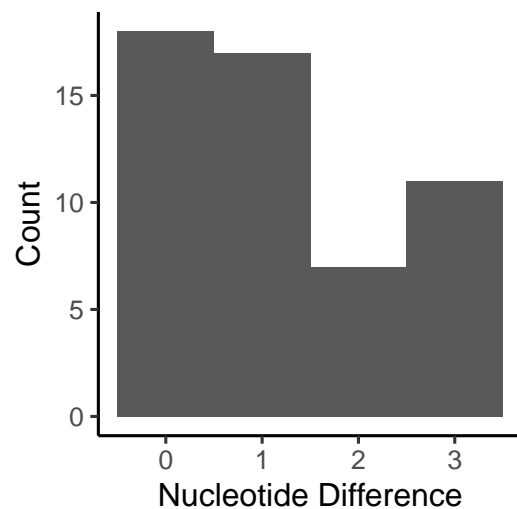
### TRBV7-2\*bp01

133 sequences assigned  
78 (58.6%) exact matches, in which:  
41 unique CDR3  
12 unique J



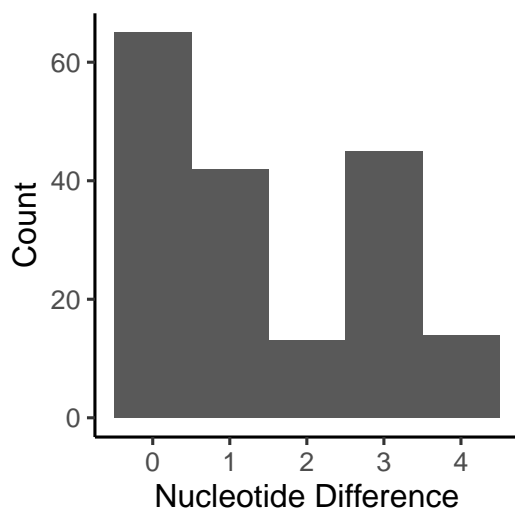
### TRBV7-4\*bp01

53 sequences assigned  
18 (34%) exact matches, in which:  
14 unique CDR3  
7 unique J



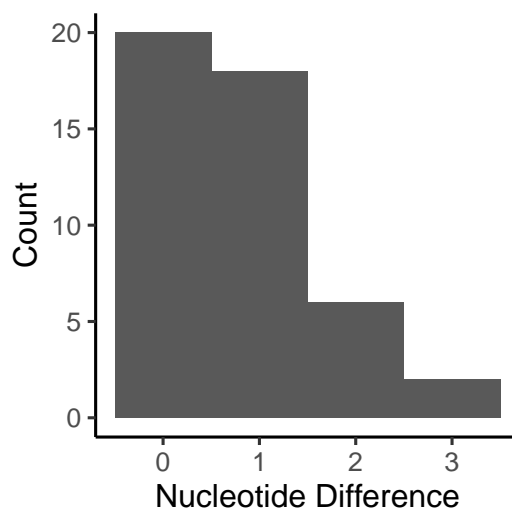
### TRBV6-6\*bp01

179 sequences assigned  
65 (36.3%) exact matches, in which:  
49 unique CDR3  
12 unique J



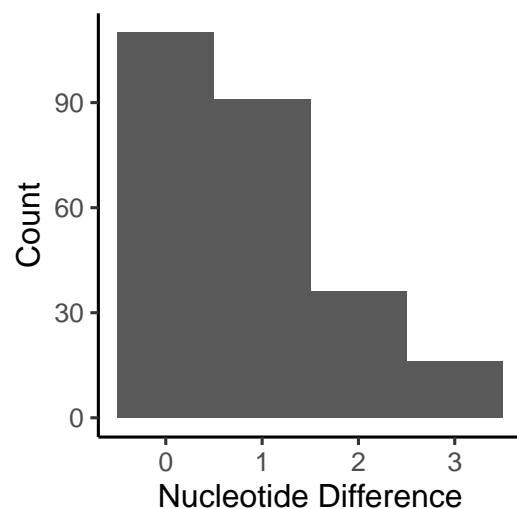
### TRBV7-2\*bp02

46 sequences assigned  
20 (43.5%) exact matches, in which:  
15 unique CDR3  
8 unique J



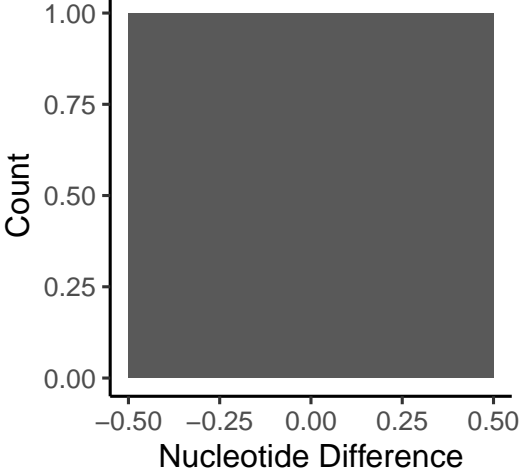
### TRBV7-6\*bp01

253 sequences assigned  
110 (43.5%) exact matches, in which:  
63 unique CDR3  
11 unique J



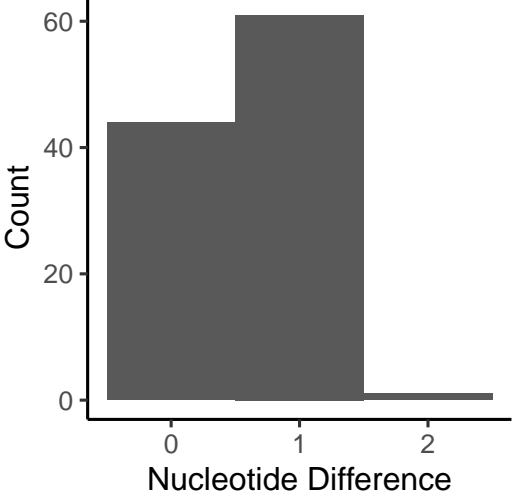
TRBV7–8\*bp01

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



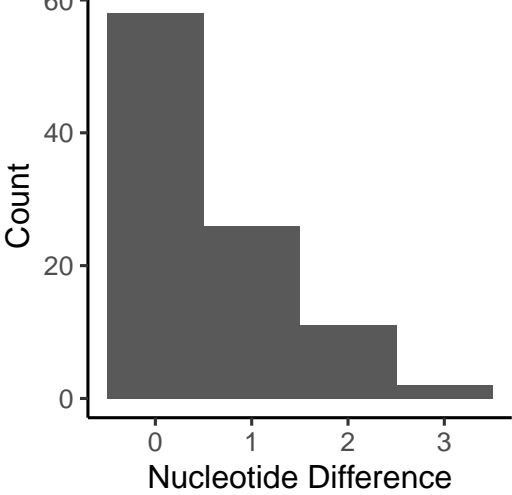
TRBV29–1\*bp03\_C315T

106 sequences assigned  
44 (41.5%) exact matches, in which:  
22 unique CDR3  
2 unique J



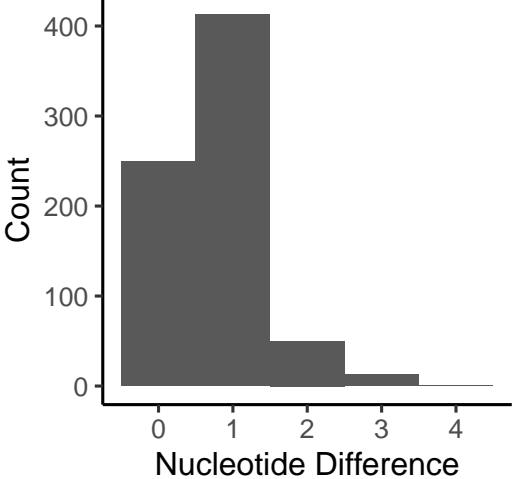
TRBV7–9\*bp01

97 sequences assigned  
58 (59.8%) exact matches, in which:  
38 unique CDR3  
11 unique J

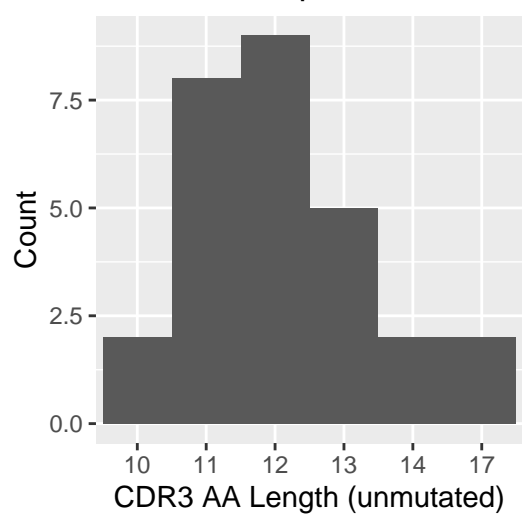


TRBV29–1\*bp01

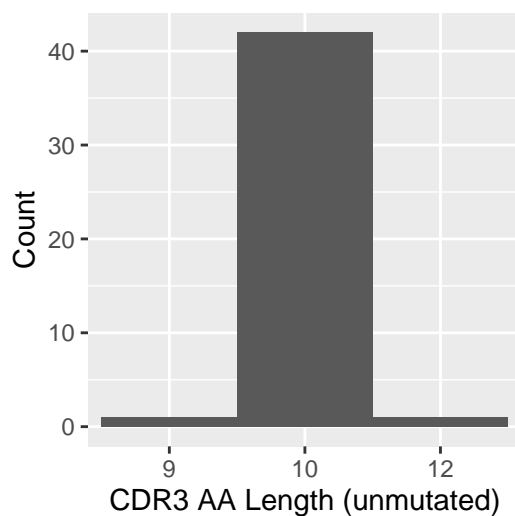
726 sequences assigned  
249 (34.3%) exact matches, in which:  
143 unique CDR3  
9 unique J



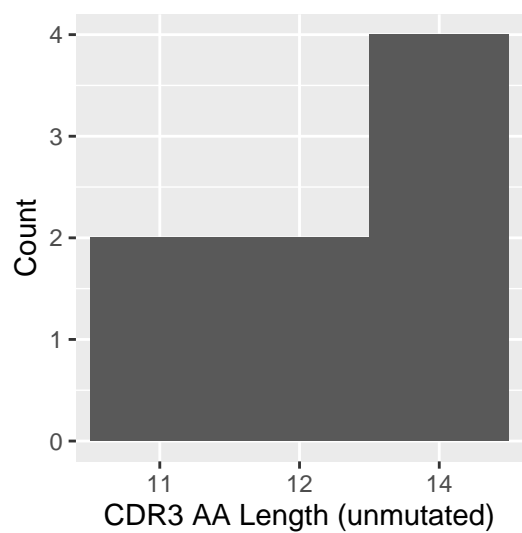
TRBV10-3\*bp02\_313GCCA<sup>-</sup>



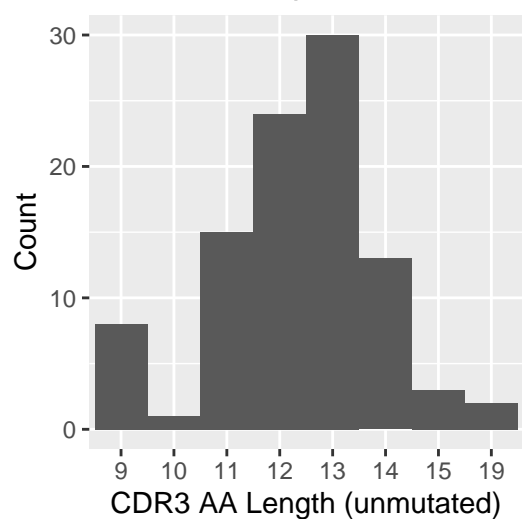
TRBV29-1\*bp03\_C315T

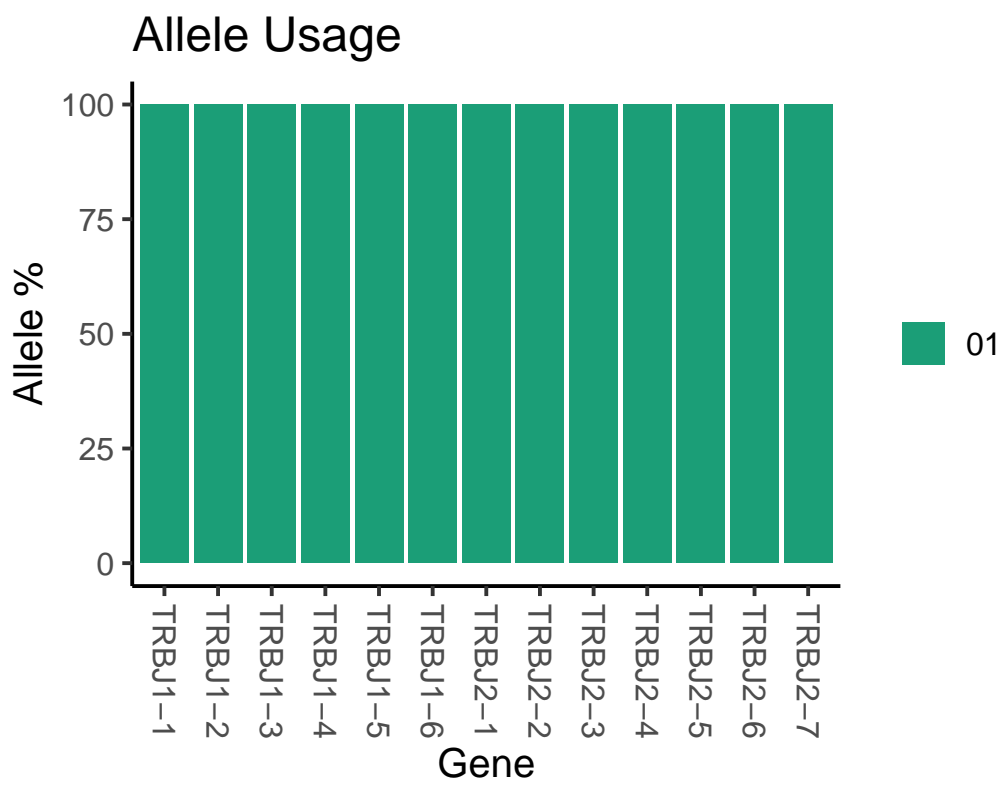


TRBV14\*bp02\_322CAAGA326



TRBV20-1\*bp04\_322GA323





**Warning** – No inferred sequence is found.