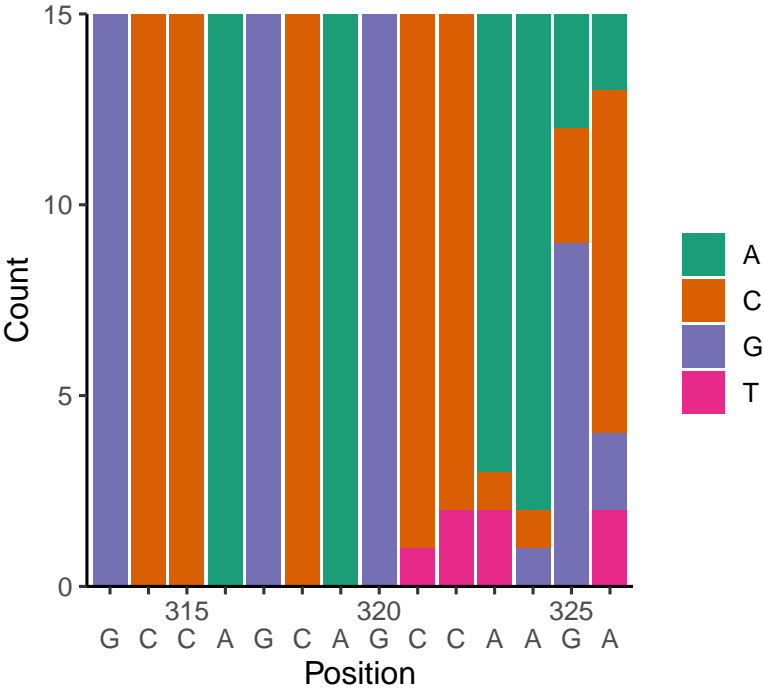
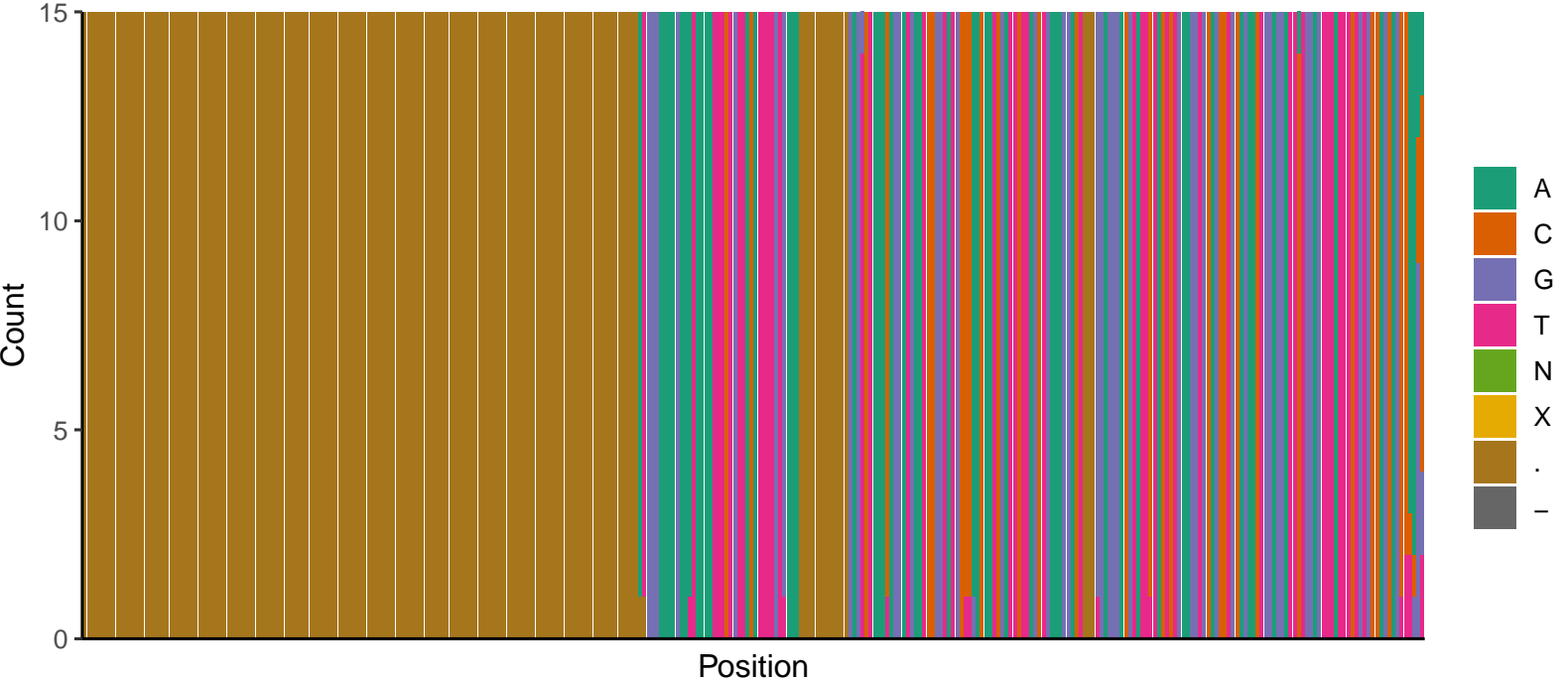


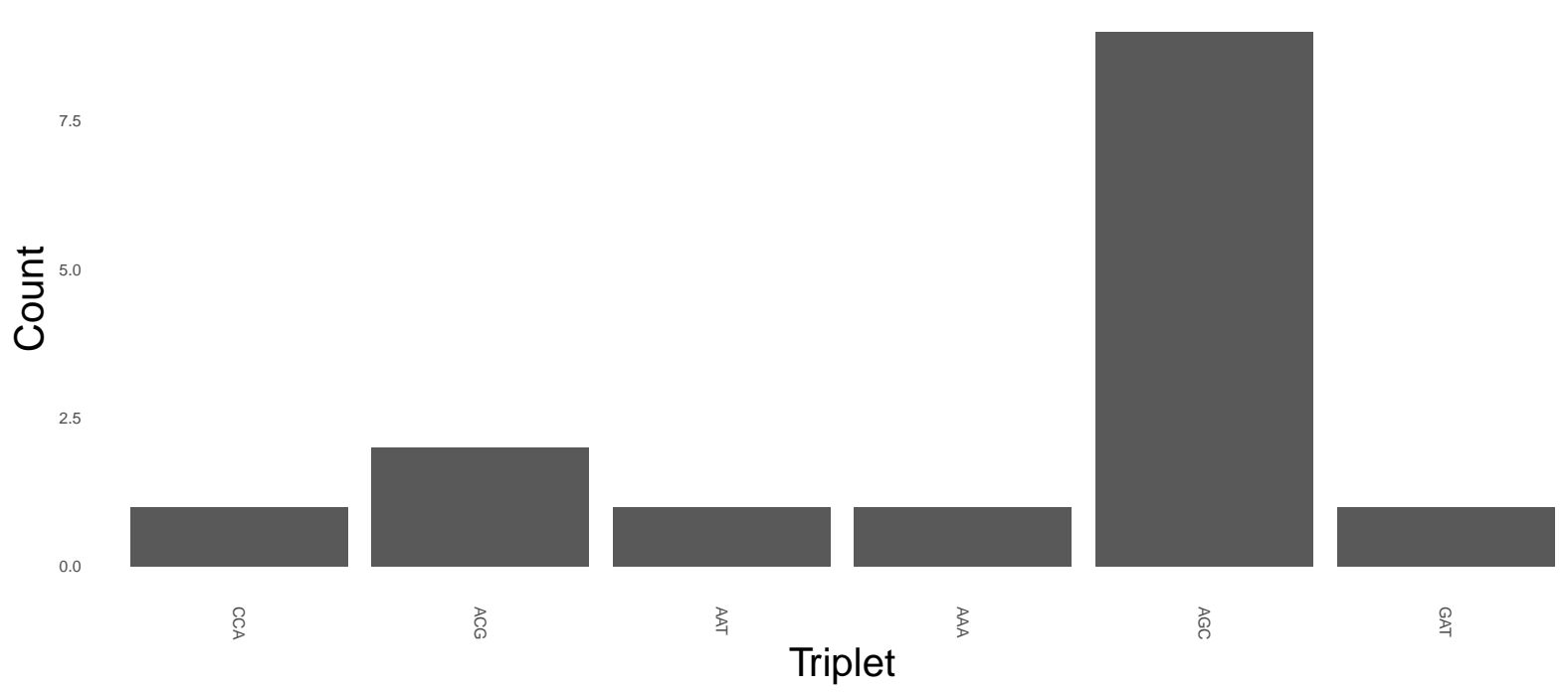
Gene TRBV14\*bp02\_322CAAGA326



Gene TRBV14\*bp02\_322CAAGA326

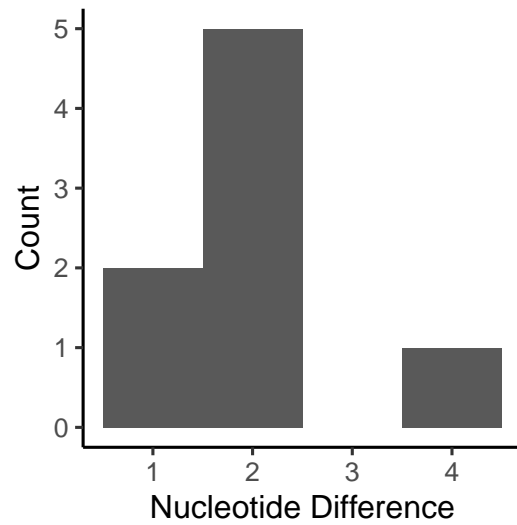


TRBV14\*bp02\_322CAAGA326– Final 3 nucleotides as a triplet



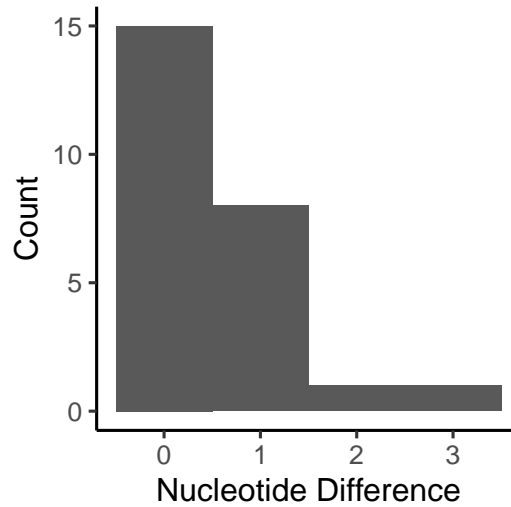
### TRBV10–1\*bp01

8 sequences assigned  
No exact matches.



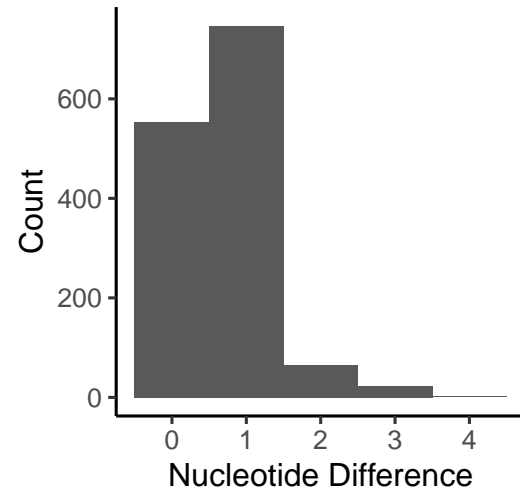
### TRBV14\*bp01

25 sequences assigned  
15 (60%) exact matches, in which:  
11 unique CDR3  
4 unique J



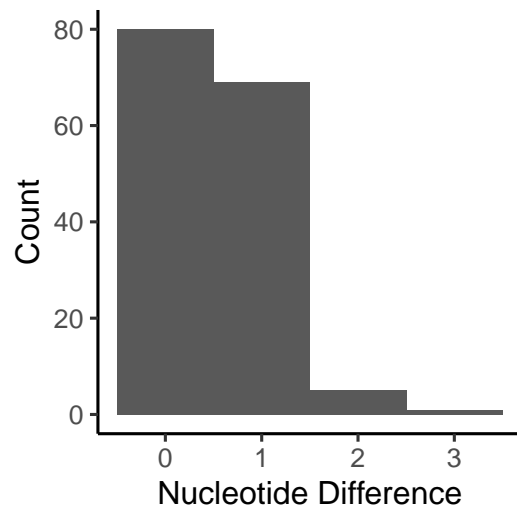
### TRBV19\*bp01

1391 sequences assigned  
554 (39.8%) exact matches, in which:  
266 unique CDR3  
11 unique J



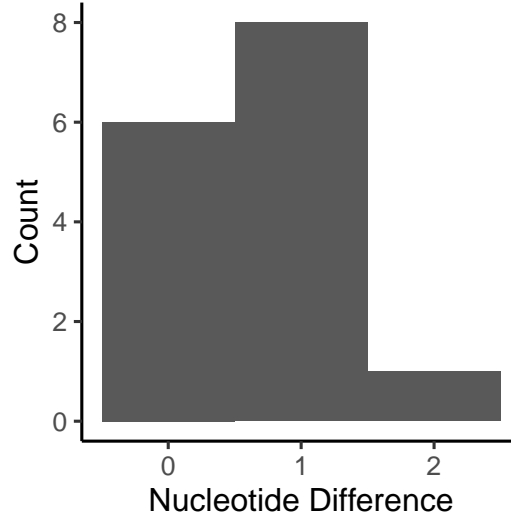
### TRBV20–1\*bp04

155 sequences assigned  
80 (51.6%) exact matches, in which:  
38 unique CDR3  
7 unique J



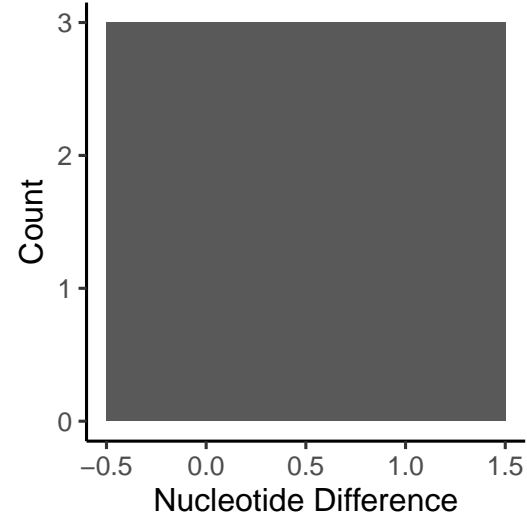
### TRBV14\*bp02\_322CAAGA:

15 sequences assigned  
6 (40%) exact matches, in which:  
5 unique CDR3  
3 unique J



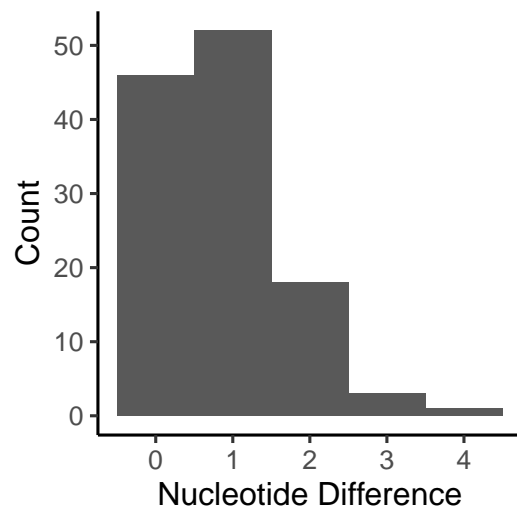
### TRBV2\*bp01

6 sequences assigned  
3 (50%) exact matches, in which:  
3 unique CDR3  
3 unique J



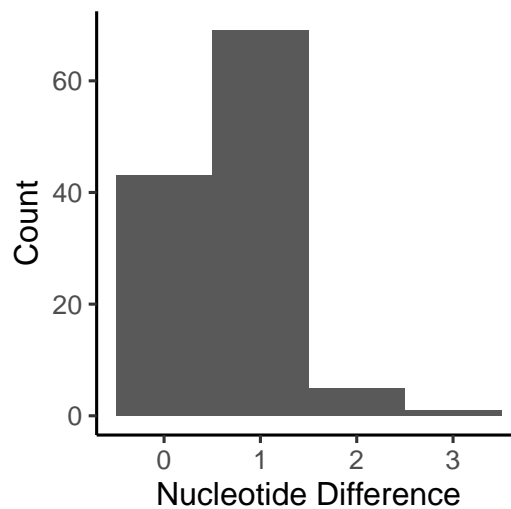
### TRBV10–3\*bp01

120 sequences assigned  
46 (38.3%) exact matches, in which:  
35 unique CDR3  
10 unique J



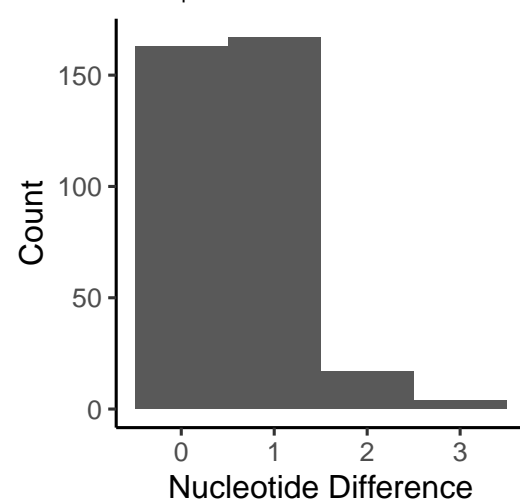
### TRBV15\*bp02

118 sequences assigned  
43 (36.4%) exact matches, in which:  
17 unique CDR3  
4 unique J



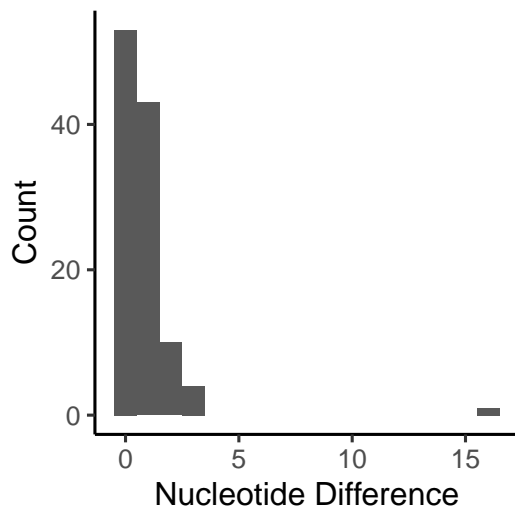
### TRBV27\*bp01

351 sequences assigned  
163 (46.4%) exact matches, in which:  
95 unique CDR3  
12 unique J



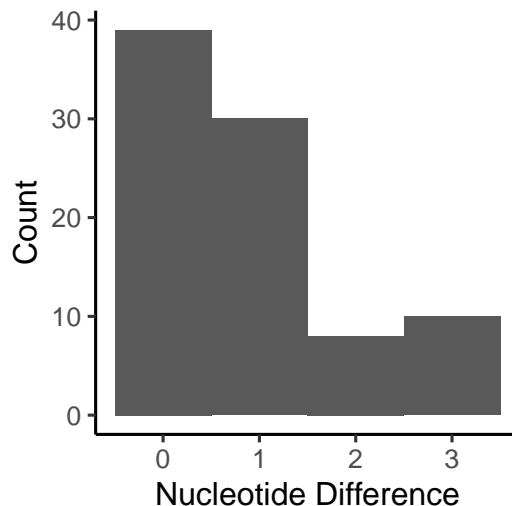
### TRBV28\*bp01

111 sequences assigned  
53 (47.7%) exact matches, in which:  
37 unique CDR3  
10 unique J



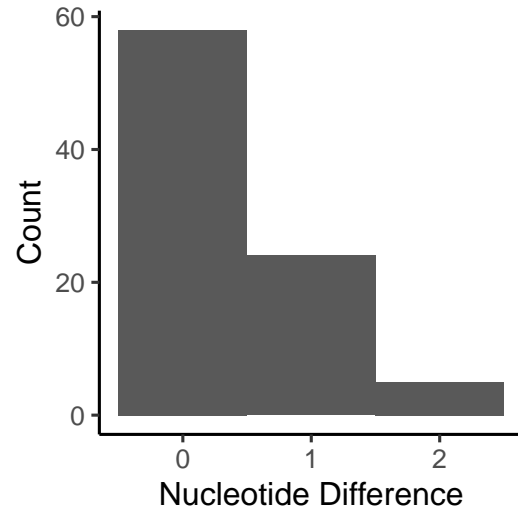
### TRBV11-1\*bp01

87 sequences assigned  
39 (44.8%) exact matches, in which:  
30 unique CDR3  
6 unique J



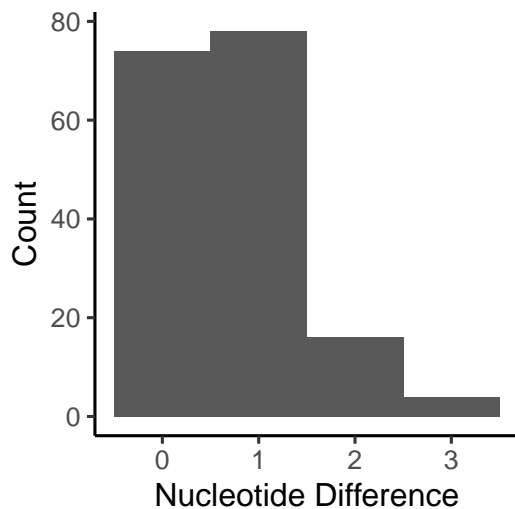
### TRBV12-34\*bp01

87 sequences assigned  
58 (66.7%) exact matches, in which:  
33 unique CDR3  
9 unique J



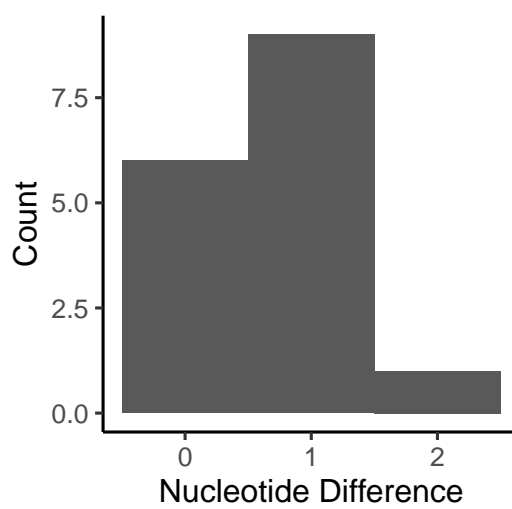
### TRBV30\*bp01

172 sequences assigned  
74 (43%) exact matches, in which:  
37 unique CDR3  
7 unique J



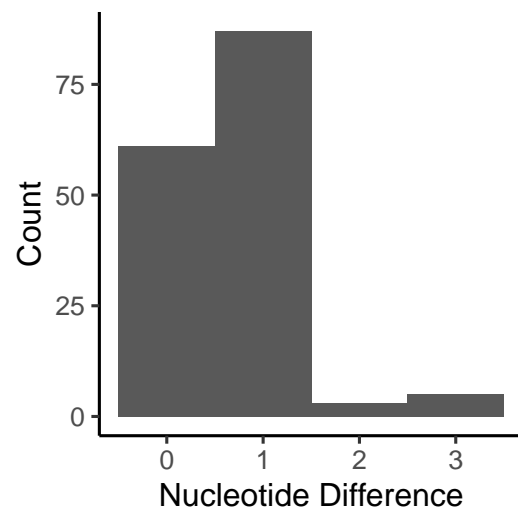
### TRBV21-1\*bp01

16 sequences assigned  
6 (37.5%) exact matches, in which:  
6 unique CDR3  
5 unique J



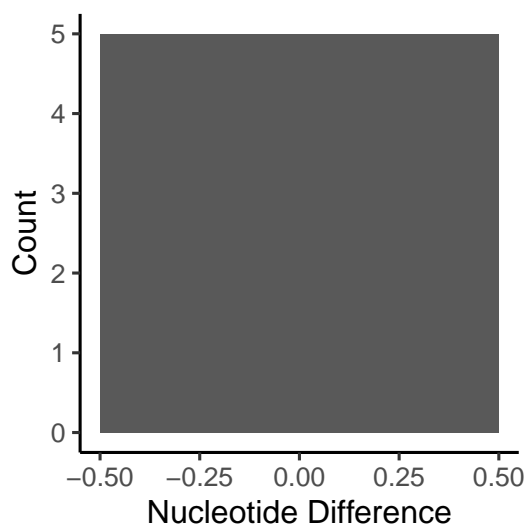
### TRBV24-1\*bp01

156 sequences assigned  
61 (39.1%) exact matches, in which:  
30 unique CDR3  
8 unique J



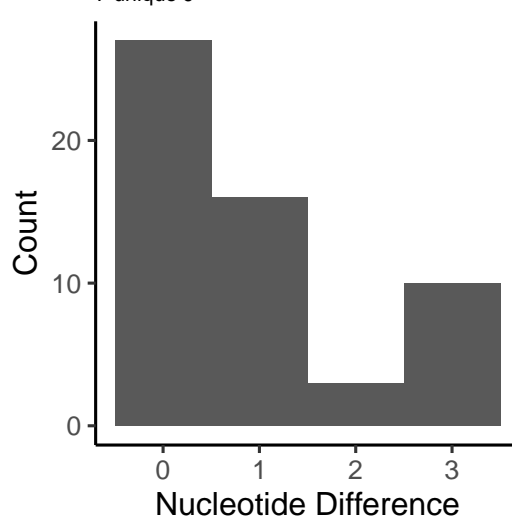
### TRBV9\*bp01

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



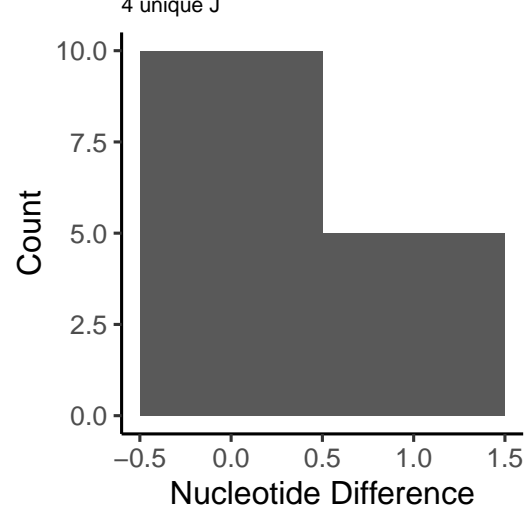
### TRBV11-2\*bp01

56 sequences assigned  
27 (48.2%) exact matches, in which:  
14 unique CDR3  
7 unique J



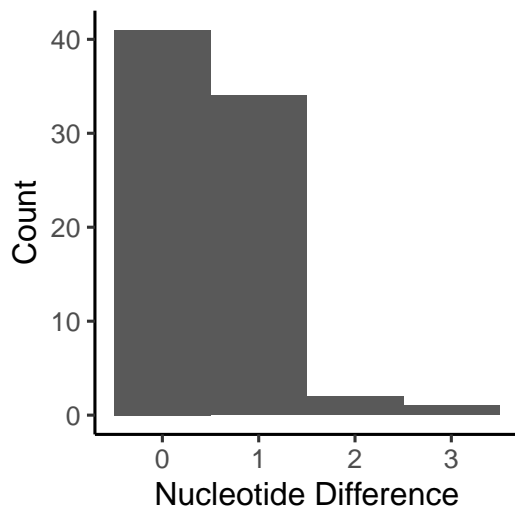
### TRBV4-1\*bp01

15 sequences assigned  
10 (66.7%) exact matches, in which:  
5 unique CDR3  
4 unique J



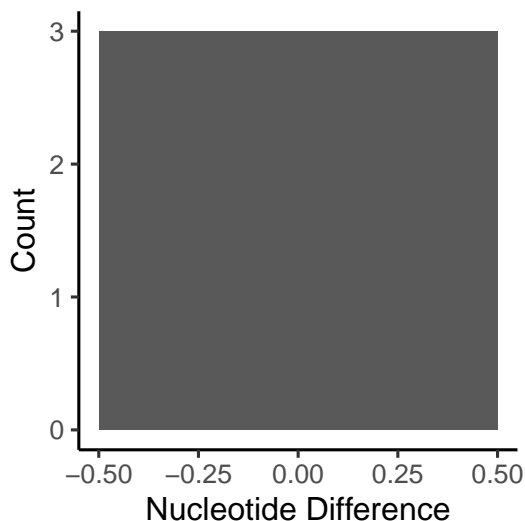
### TRBV4-2\*bp01

78 sequences assigned  
41 (52.6%) exact matches, in which:  
31 unique CDR3  
7 unique J



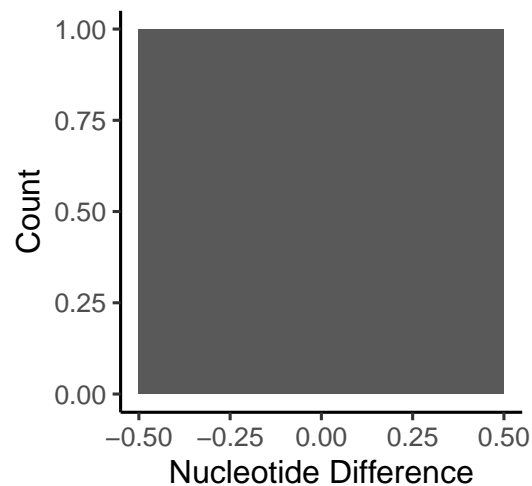
### TRBV25-1\*bp01

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



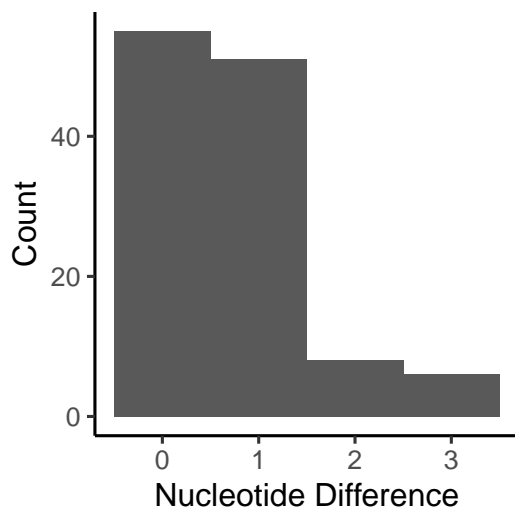
### TRBV5-5\*bp01

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



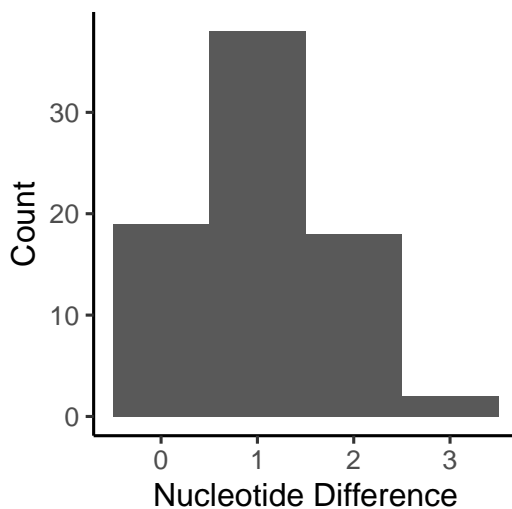
### TRBV4-3\*bp01

120 sequences assigned  
55 (45.8%) exact matches, in which:  
32 unique CDR3  
8 unique J



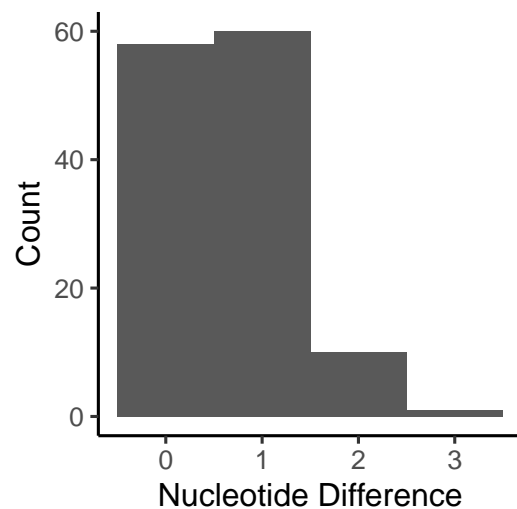
### TRBV5-3\*bp01

77 sequences assigned  
19 (24.7%) exact matches, in which:  
14 unique CDR3  
4 unique J



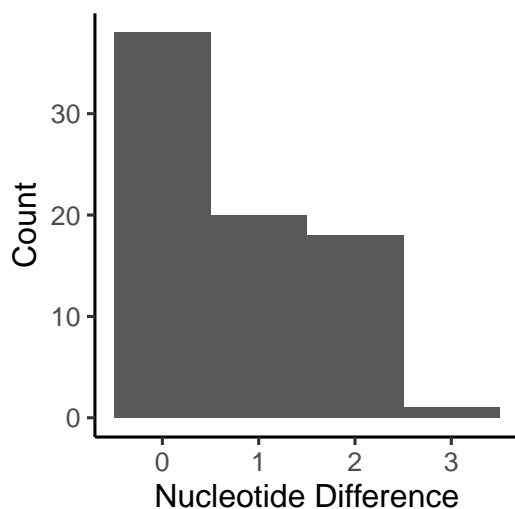
### TRBV5-6\*bp01

129 sequences assigned  
58 (45%) exact matches, in which:  
40 unique CDR3  
6 unique J



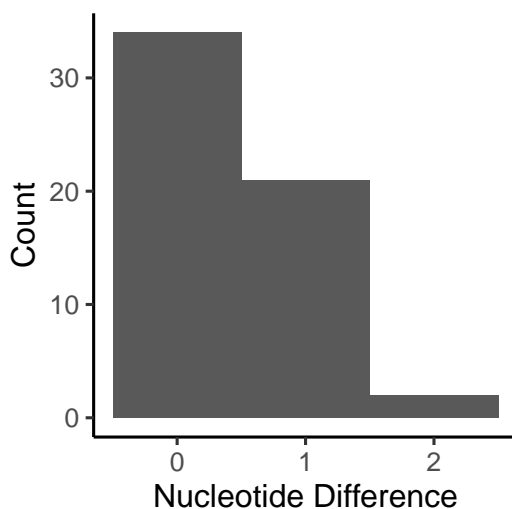
### TRBV5-1\*bp01

77 sequences assigned  
38 (49.4%) exact matches, in which:  
19 unique CDR3  
6 unique J



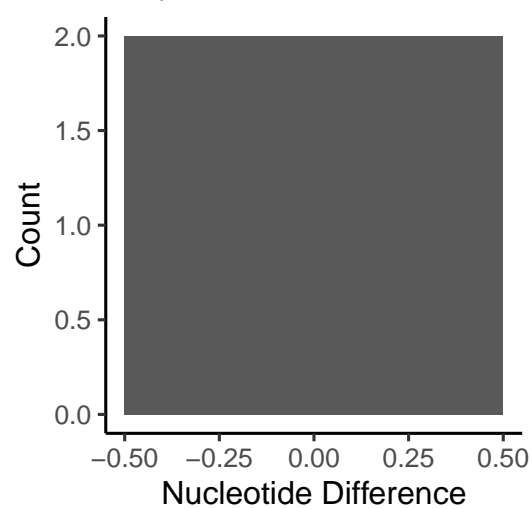
### TRBV5-4\*bp01

57 sequences assigned  
34 (59.6%) exact matches, in which:  
19 unique CDR3  
6 unique J



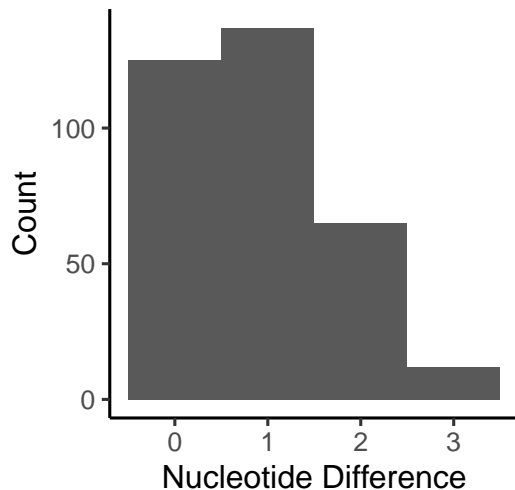
### TRBV5-8\*bp01

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



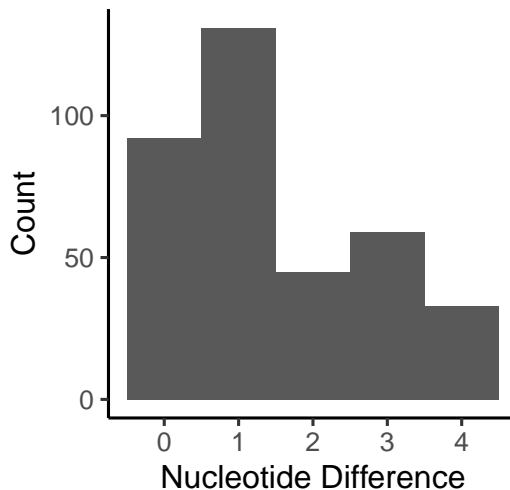
### TRBV6-1\*bp01

339 sequences assigned  
125 (36.9%) exact matches, in which:  
85 unique CDR3  
11 unique J



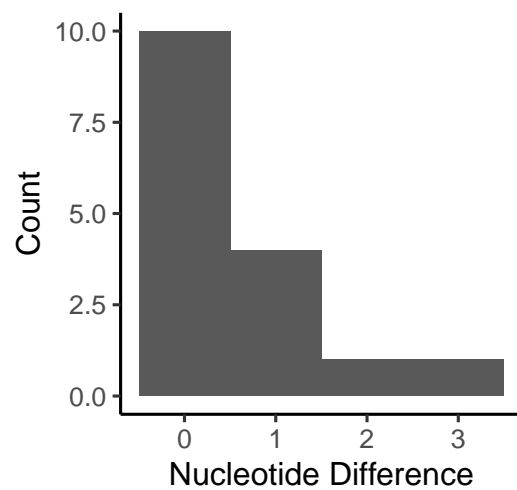
### TRBV6-6\*bp01

360 sequences assigned  
92 (25.6%) exact matches, in which:  
61 unique CDR3  
12 unique J



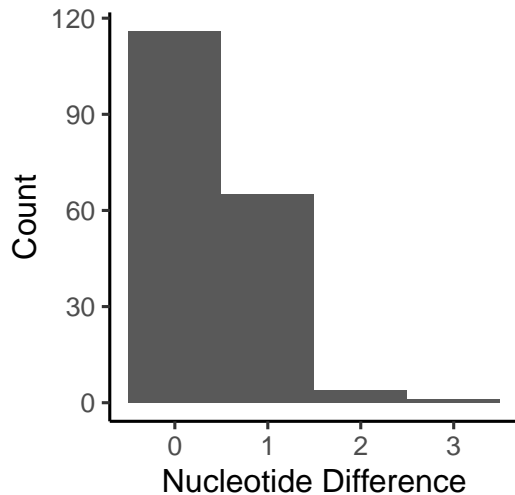
### TRBV7-2\*bp02

16 sequences assigned  
10 (62.5%) exact matches, in which:  
4 unique CDR3  
4 unique J



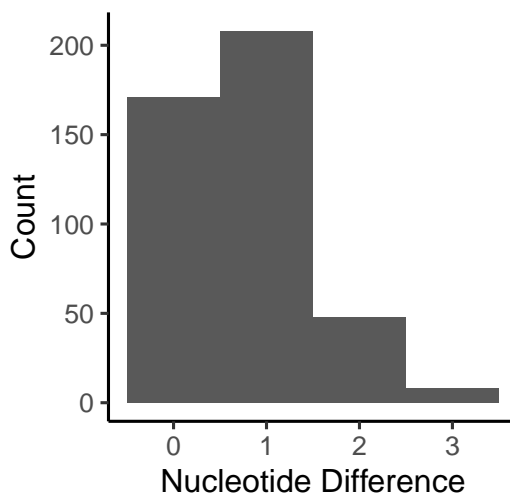
### TRBV6-4\*bp01

186 sequences assigned  
116 (62.4%) exact matches, in which:  
70 unique CDR3  
11 unique J



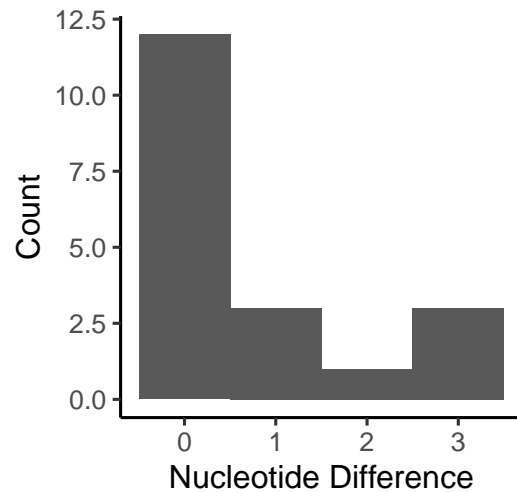
### TRBV6-23\*bp01

435 sequences assigned  
171 (39.3%) exact matches, in which:  
100 unique CDR3  
12 unique J



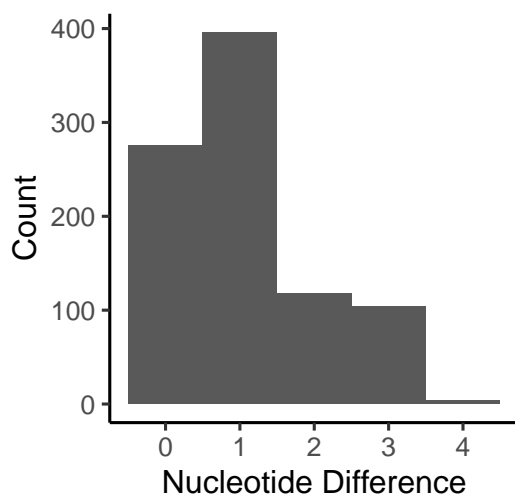
### TRBV7-3\*bp01

19 sequences assigned  
12 (63.2%) exact matches, in which:  
9 unique CDR3  
4 unique J



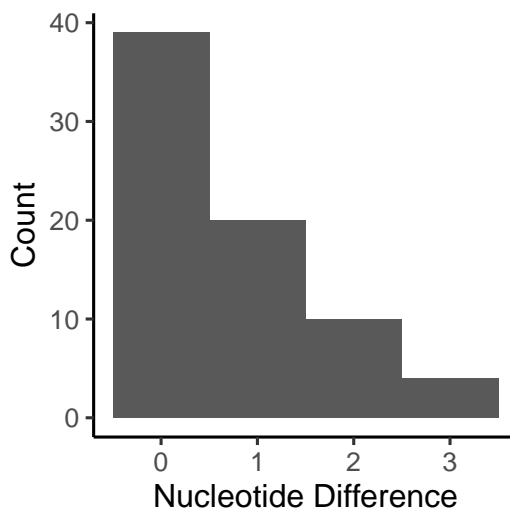
### TRBV6-5\*bp01

897 sequences assigned  
275 (30.7%) exact matches, in which:  
181 unique CDR3  
12 unique J



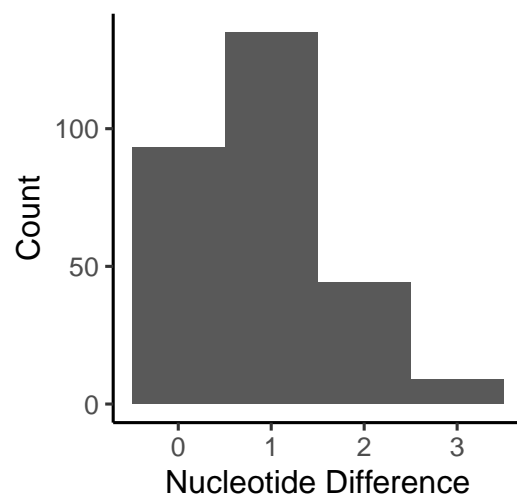
### TRBV7-2\*bp01

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



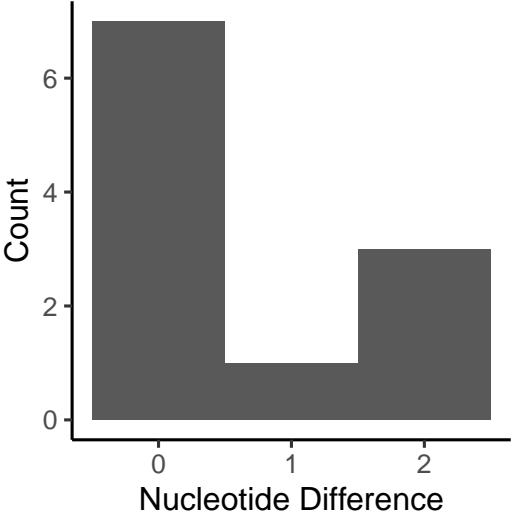
### TRBV7-6\*bp01

281 sequences assigned  
93 (33.1%) exact matches, in which:  
53 unique CDR3  
8 unique J



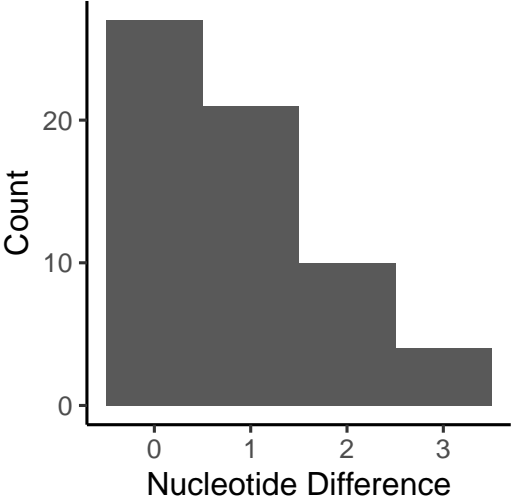
TRBV7-7\*bp01

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



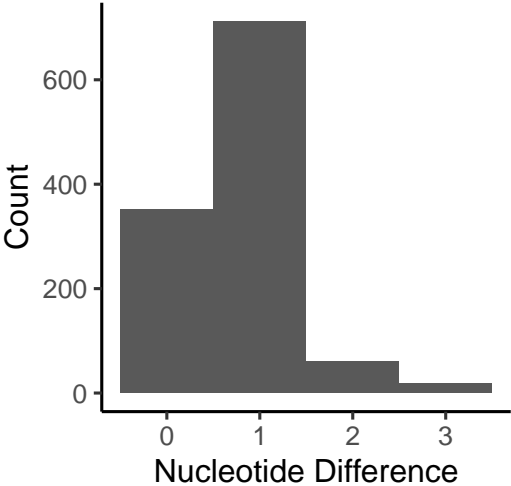
TRBV7-9\*bp01

62 sequences assigned  
27 (43.5%) exact matches, in which:  
12 unique CDR3  
6 unique J



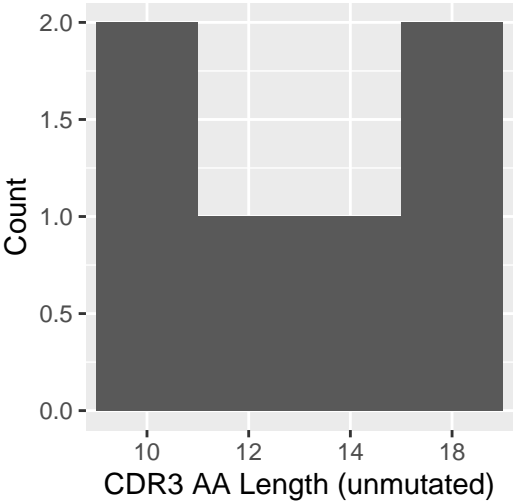
TRBV29-1\*bp01

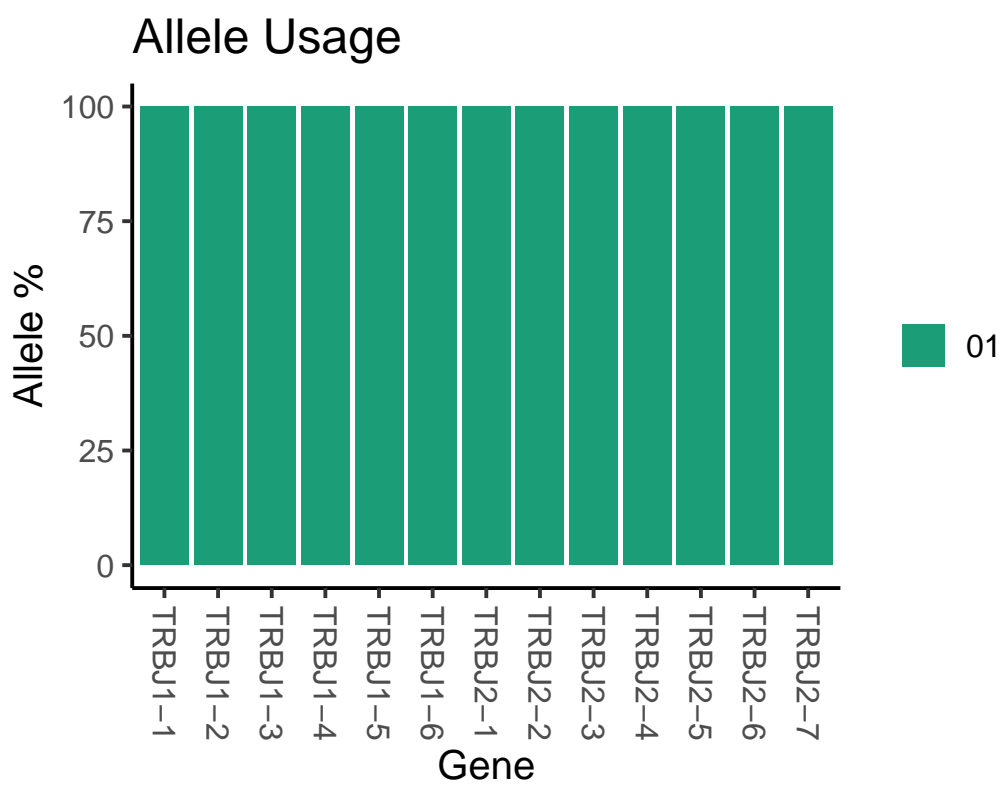
1142 sequences assigned  
351 (30.7%) exact matches, in which:  
186 unique CDR3  
6 unique J





TRBV14\*bp02\_322CAAGA326





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