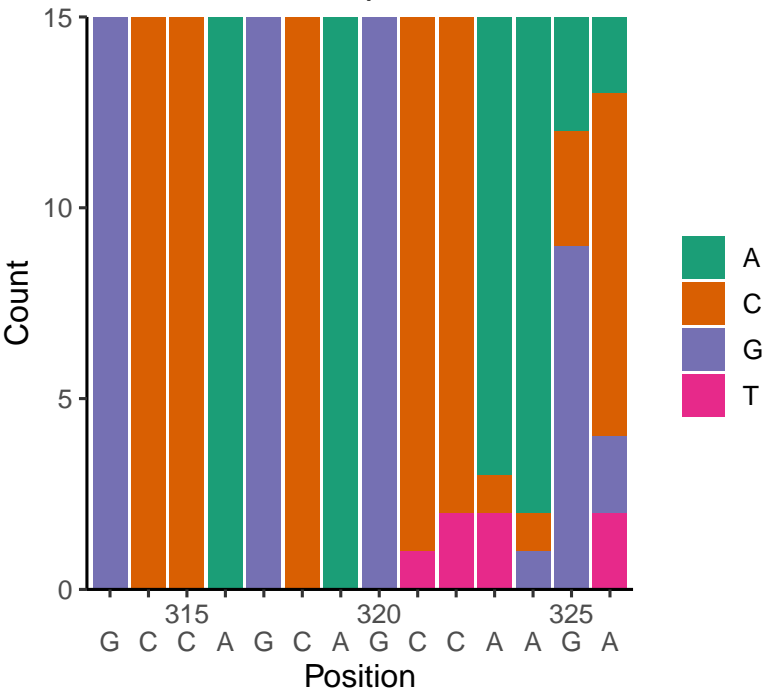
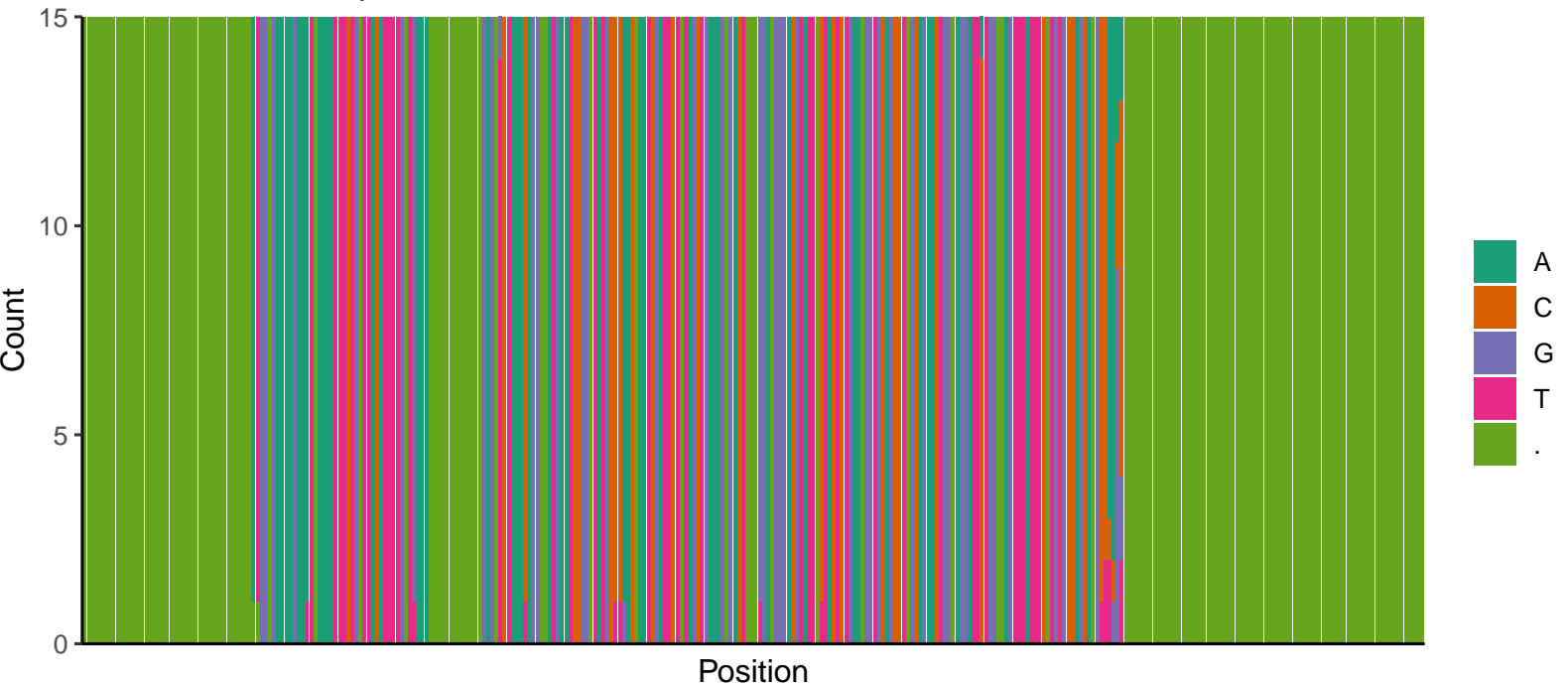


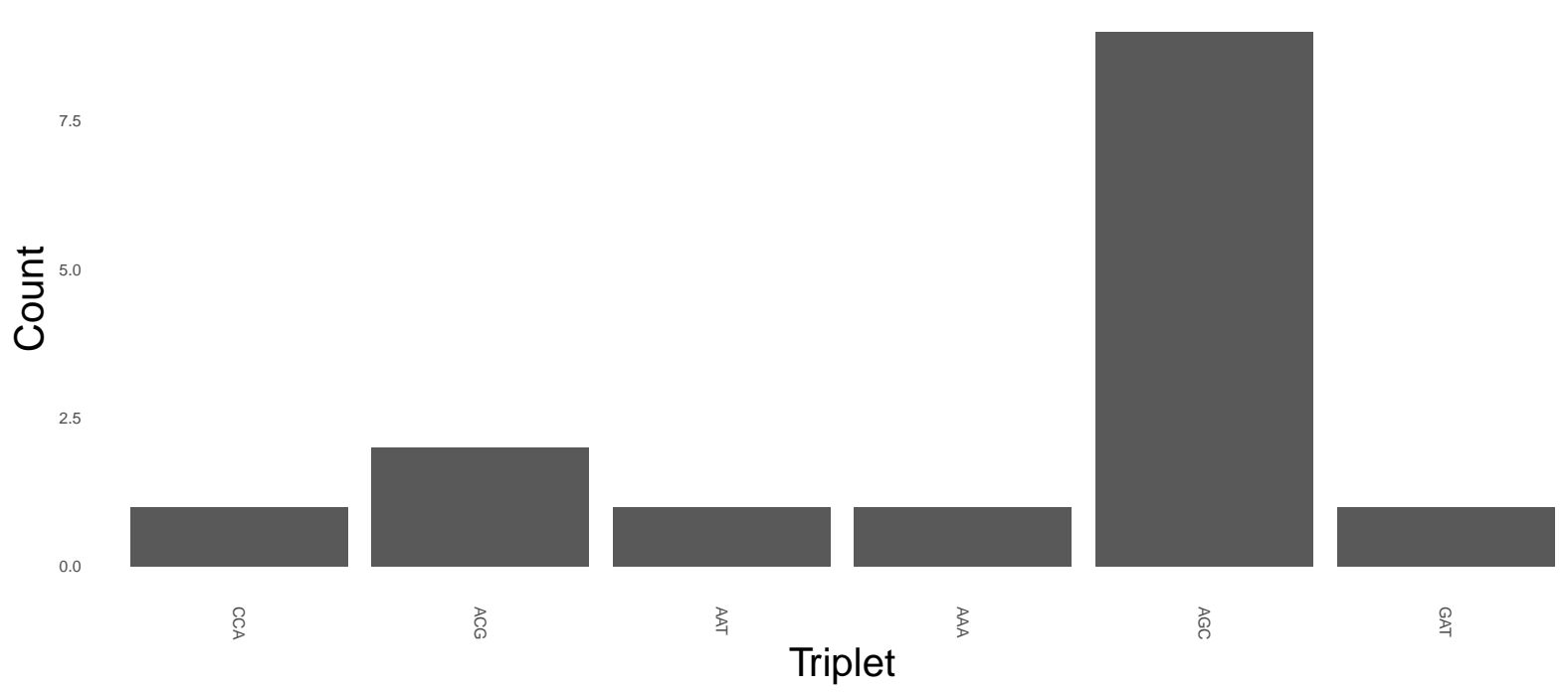
Gene TRBV14*bp02_322CAAGA326



Gene TRBV14*bp02_322CAAGA326

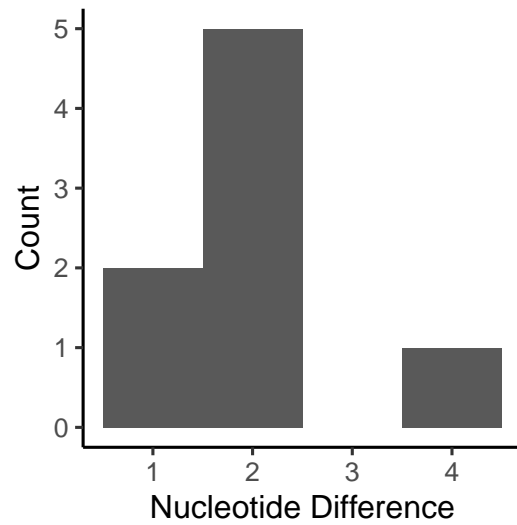


TRBV14*bp02_322CAAGA326– Final 3 nucleotides as a triplet



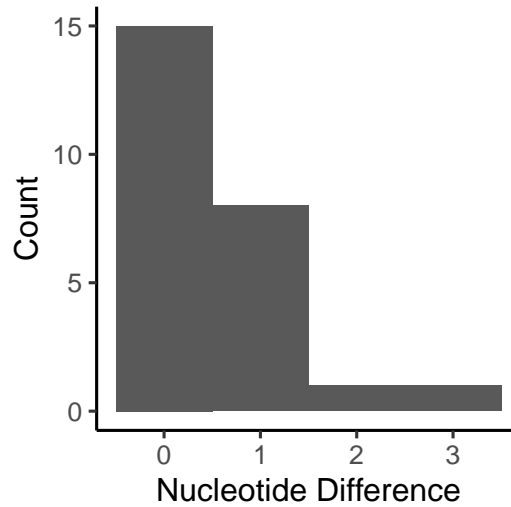
TRBV10–1*bp01

8 sequences assigned
No exact matches.



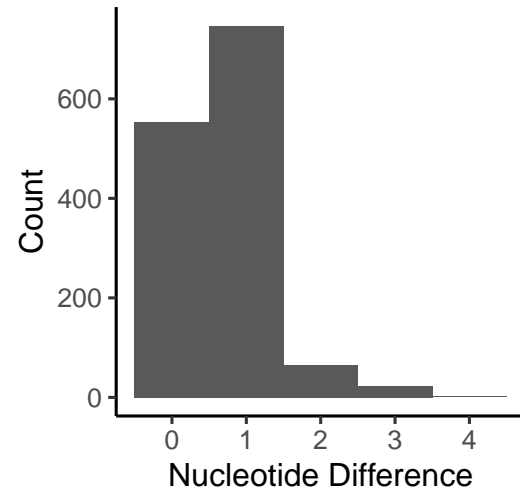
TRBV14*bp01

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



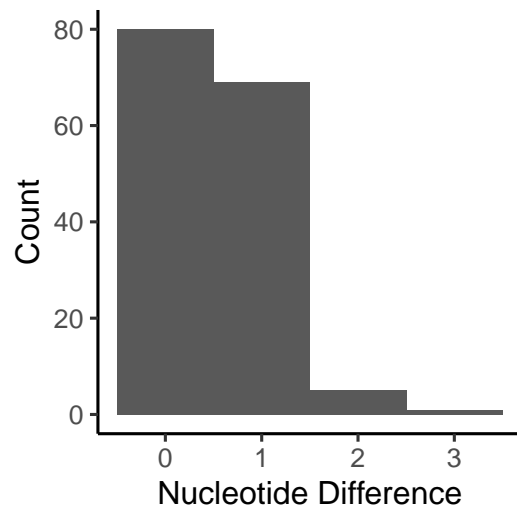
TRBV19*bp01

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



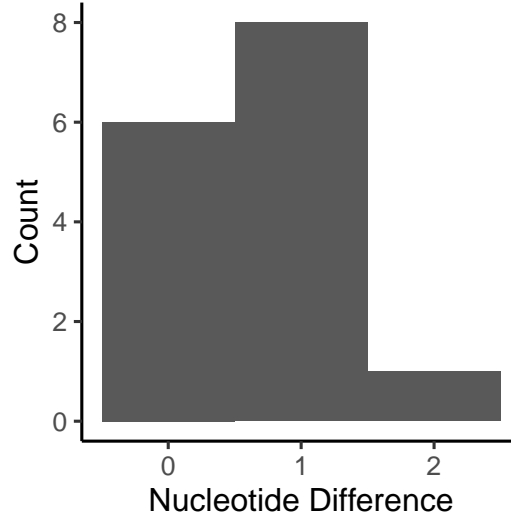
TRBV20–1*bp04

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



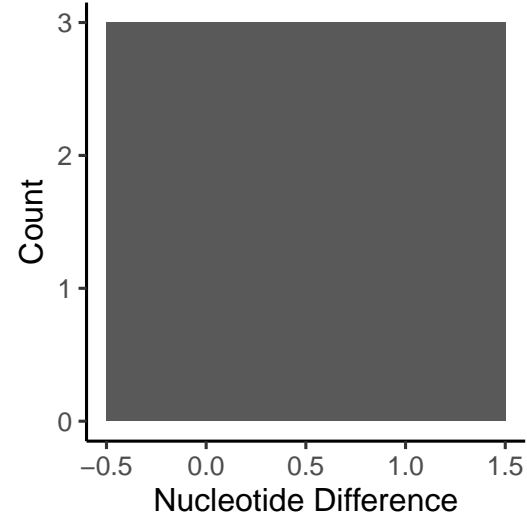
TRBV14*bp02_322CAAGA:

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



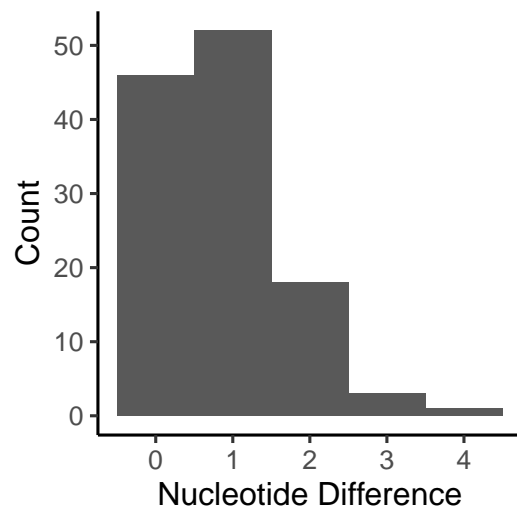
TRBV2*bp01

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



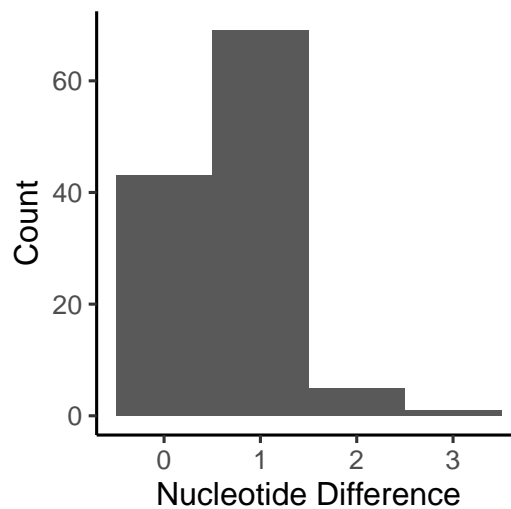
TRBV10–3*bp01

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



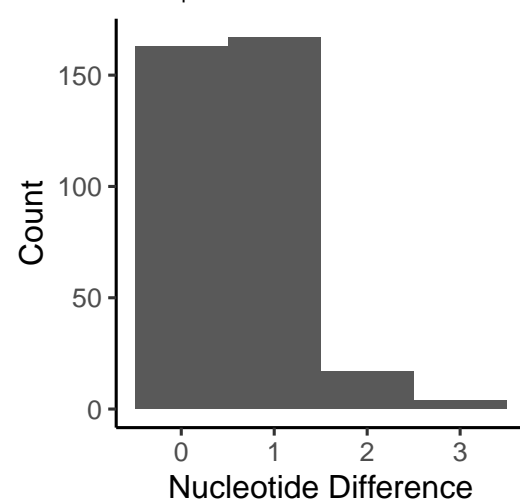
TRBV15*bp02

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



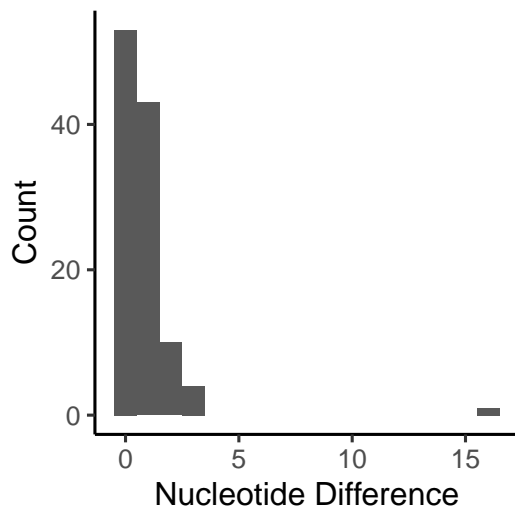
TRBV27*bp01

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



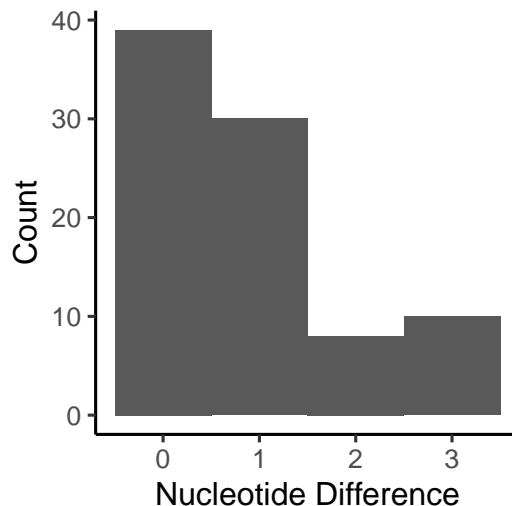
TRBV28*bp01

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



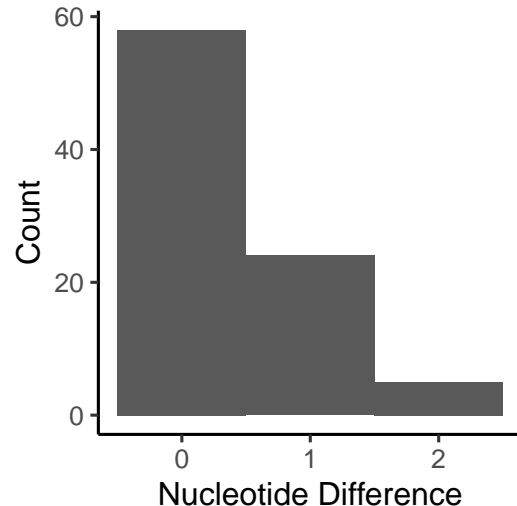
TRBV11-1*bp01

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



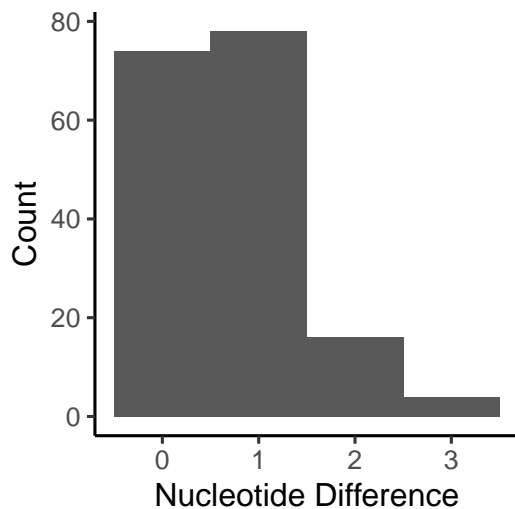
TRBV12-34*bp01

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



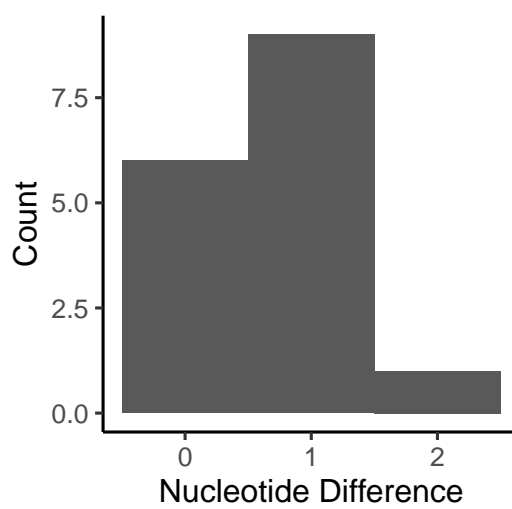
TRBV30*bp01

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



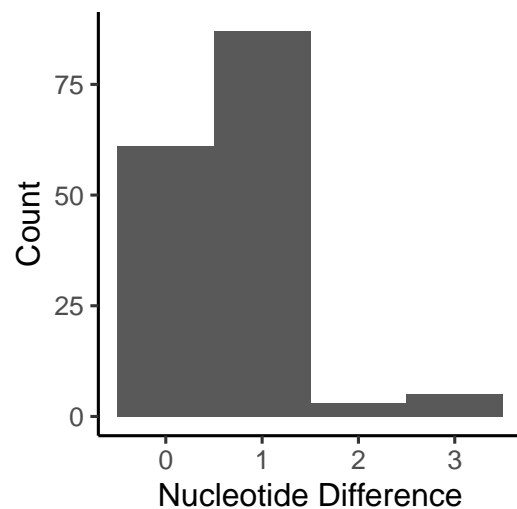
TRBV21-1*bp01

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



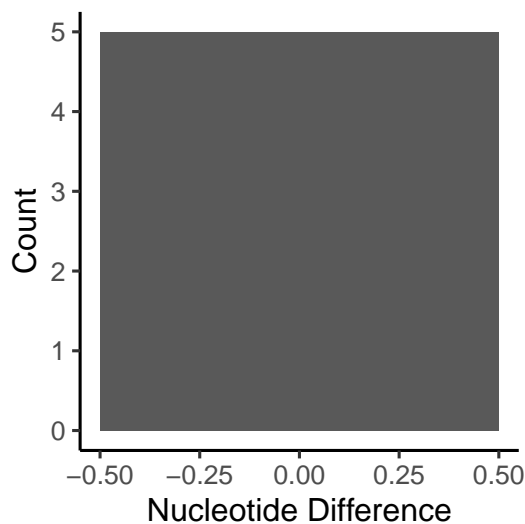
TRBV24-1*bp01

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



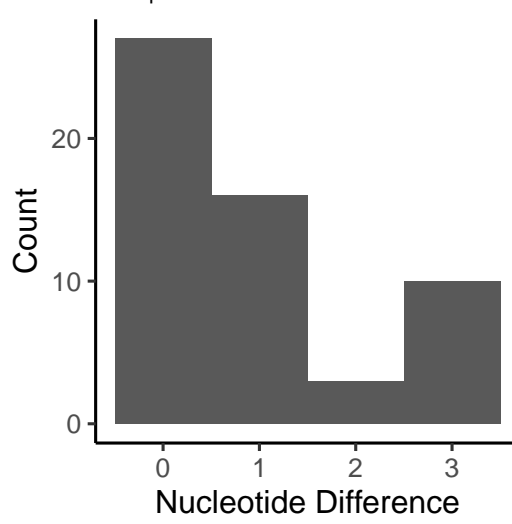
TRBV9*bp01

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



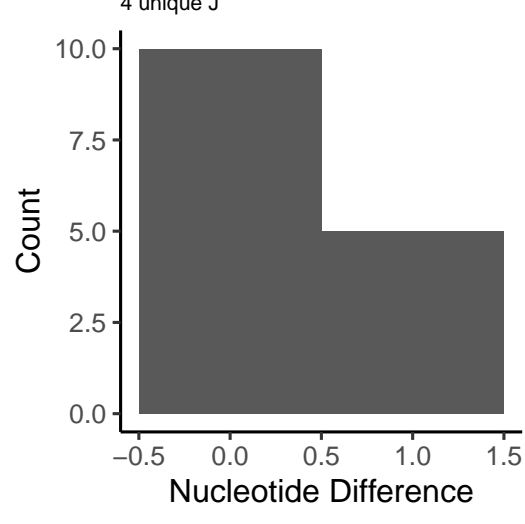
TRBV11-2*bp01

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



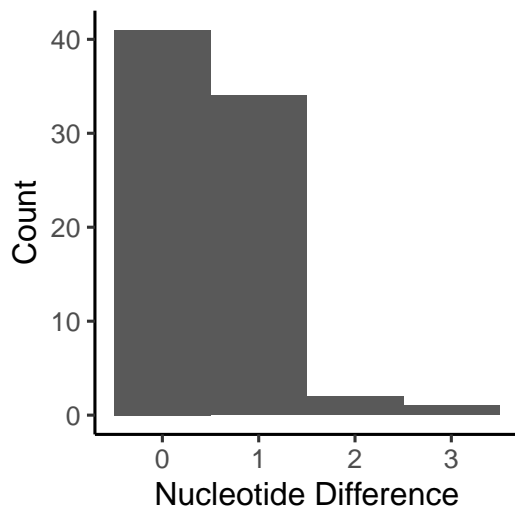
TRBV4-1*bp01

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



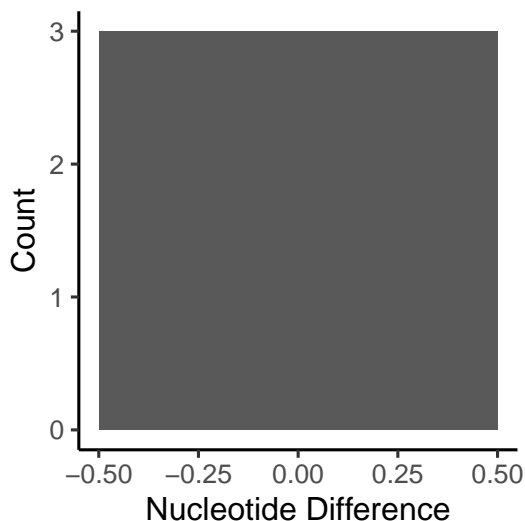
TRBV4-2*bp01

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



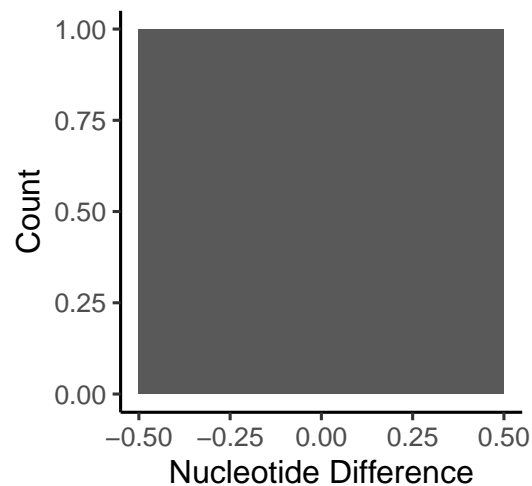
TRBV25-1*bp01

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



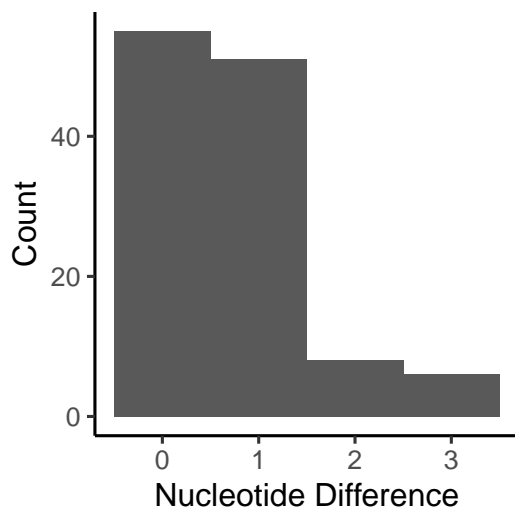
TRBV5-5*bp01

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



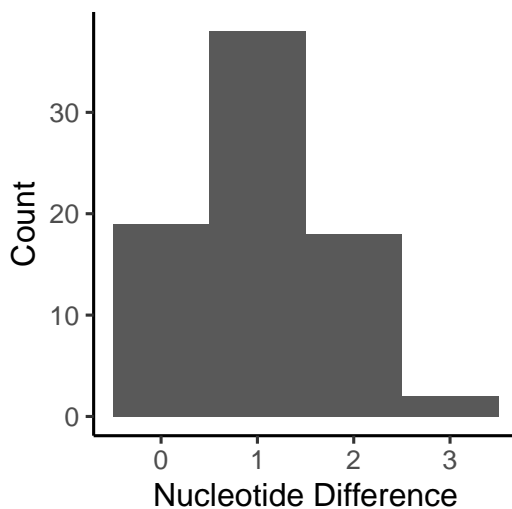
TRBV4-3*bp01

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



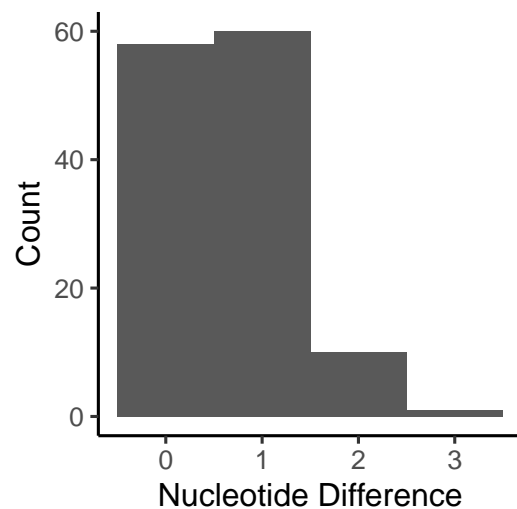
TRBV5-3*bp01

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



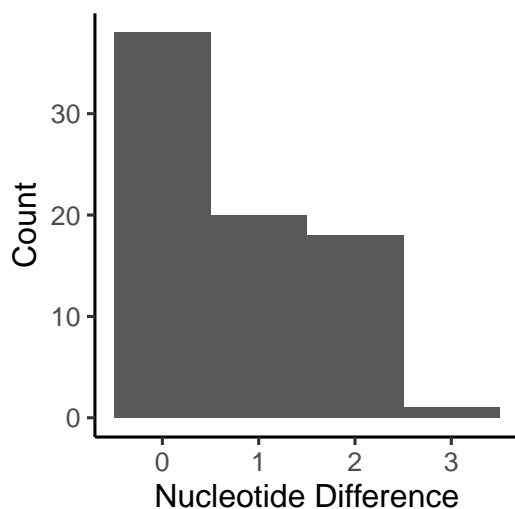
TRBV5-6*bp01

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



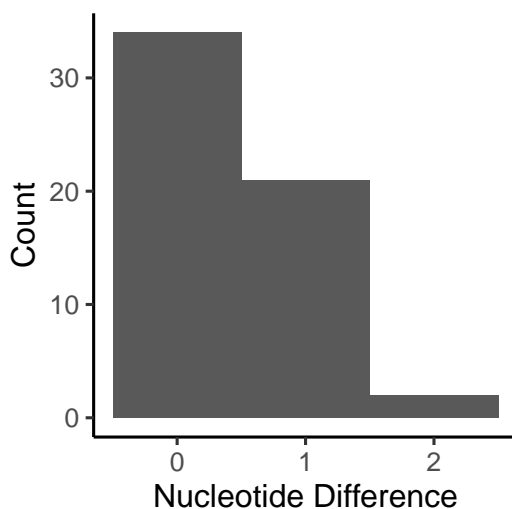
TRBV5-1*bp01

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



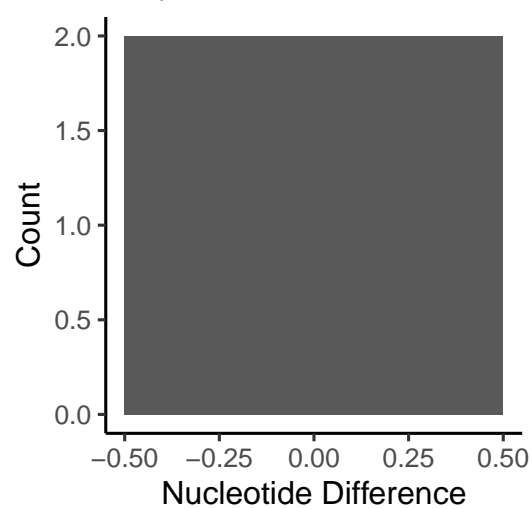
TRBV5-4*bp01

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



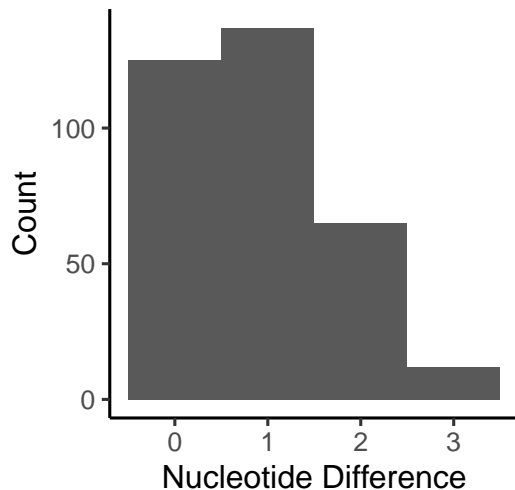
TRBV5-8*bp01

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



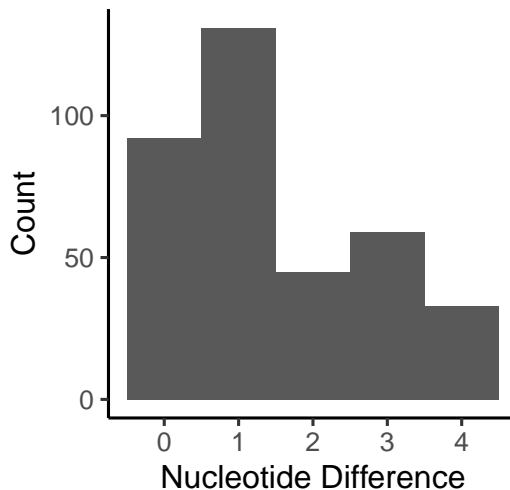
TRBV6-1*bp01

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



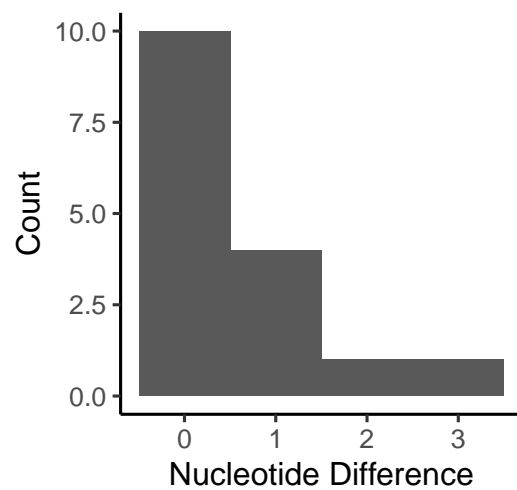
TRBV6-6*bp01

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



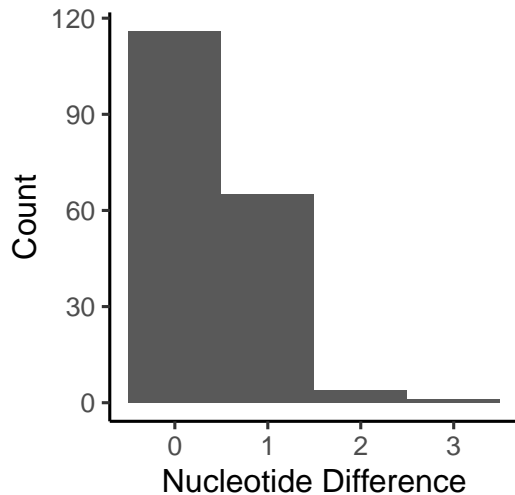
TRBV7-2*bp02

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



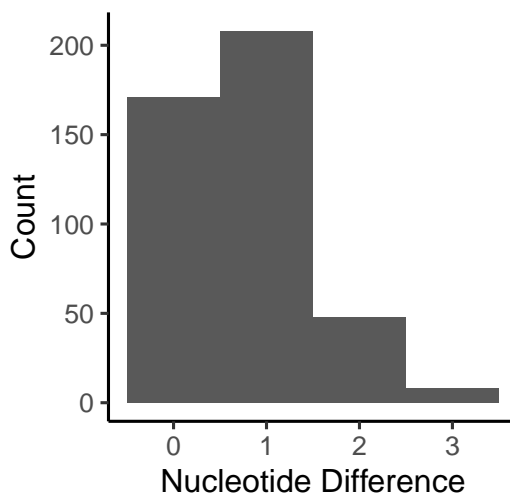
TRBV6-4*bp01

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



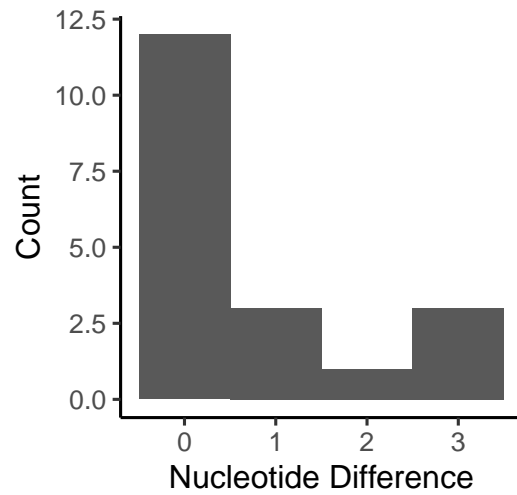
TRBV6-23*bp01

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



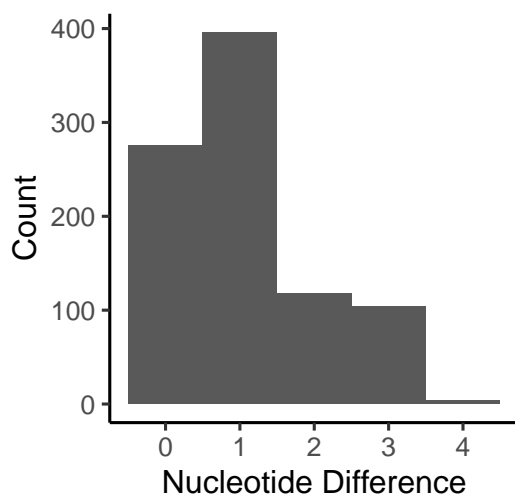
TRBV7-3*bp01

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



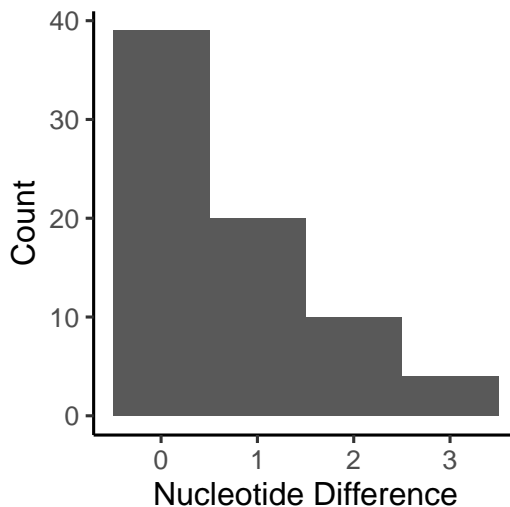
TRBV6-5*bp01

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



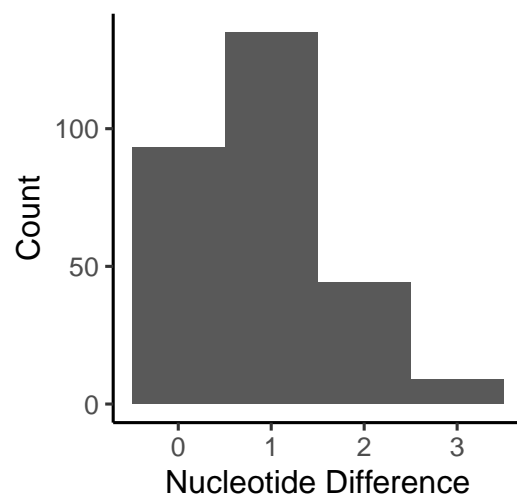
TRBV7-2*bp01

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



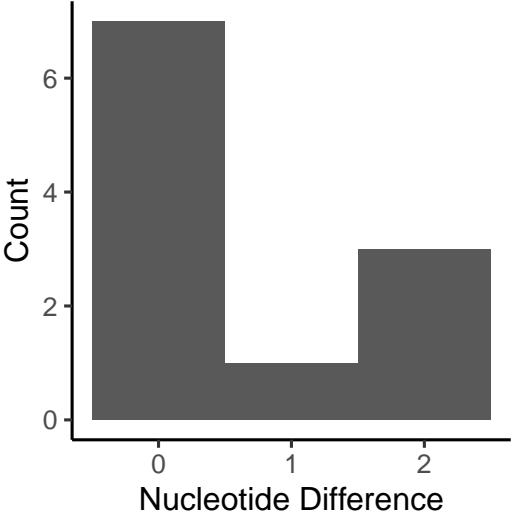
TRBV7-6*bp01

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



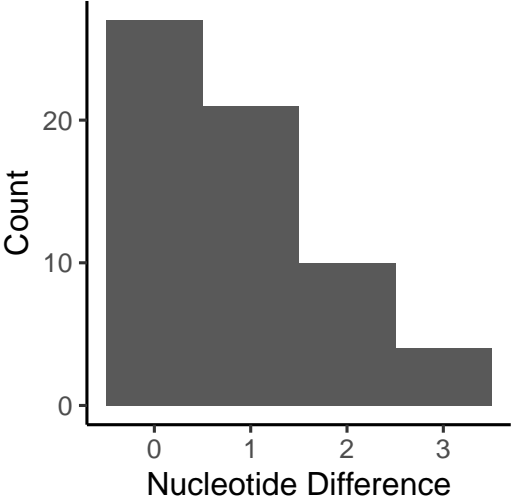
TRBV7-7*bp01

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



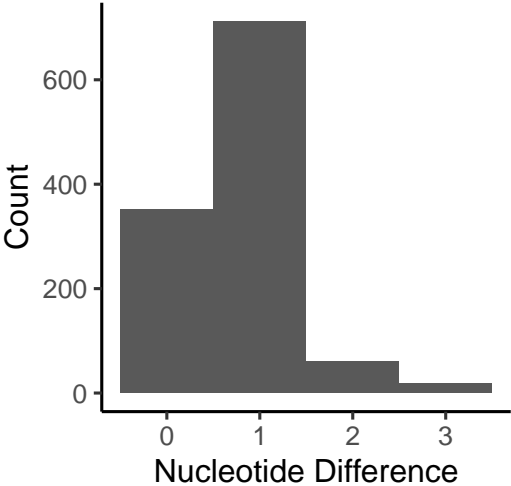
TRBV7-9*bp01

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

