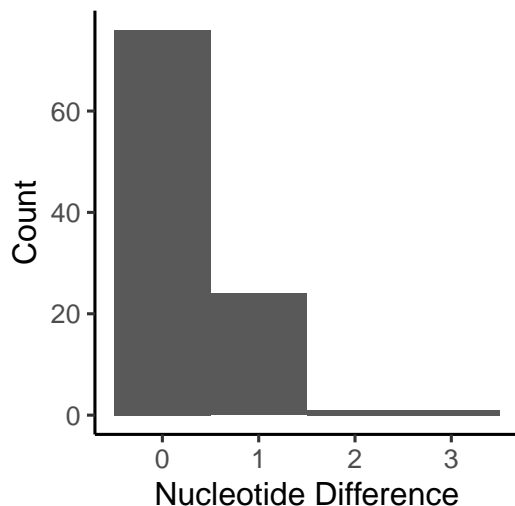


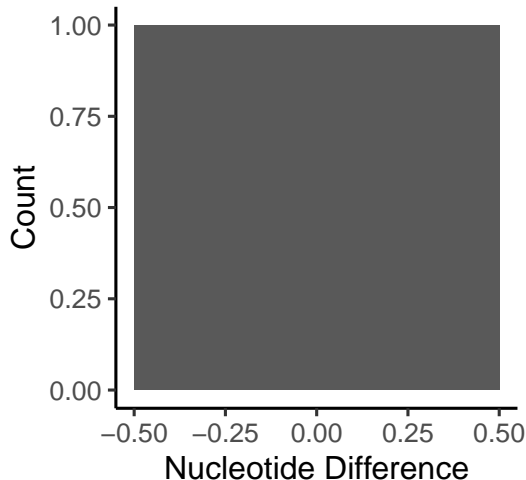
TRBV20-1*bp01

102 sequences assigned
76 (74.5%) exact matches, in which:
29 unique CDR3
9 unique J



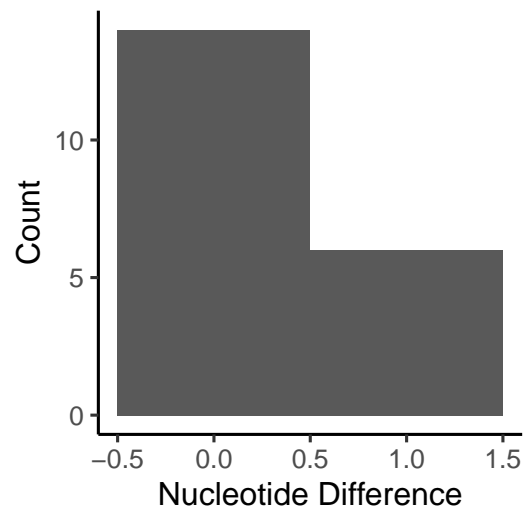
TRBV10-2*bp01

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



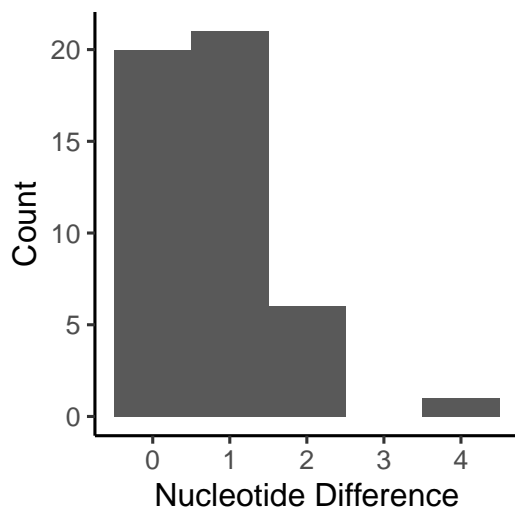
TRBV13*bp01

20 sequences assigned
14 (70%) exact matches, in which:
10 unique CDR3
4 unique J



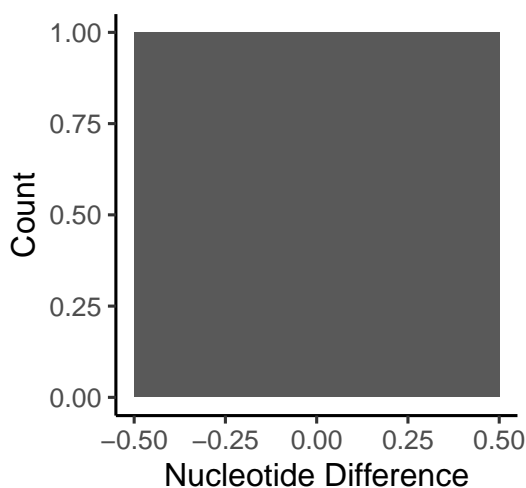
TRBV10-1*bp01

48 sequences assigned
20 (41.7%) exact matches, in which:
10 unique CDR3
1 unique J



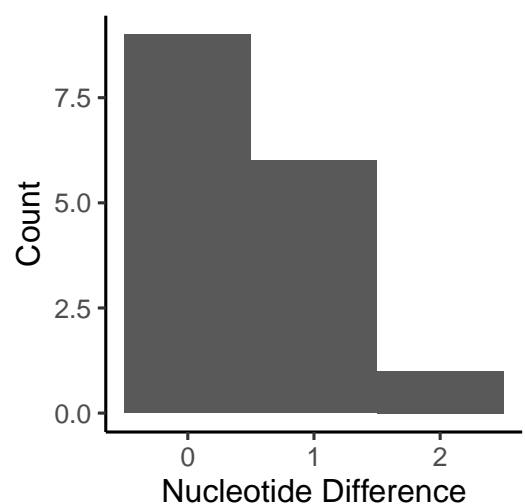
TRBV10-2*bp02

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



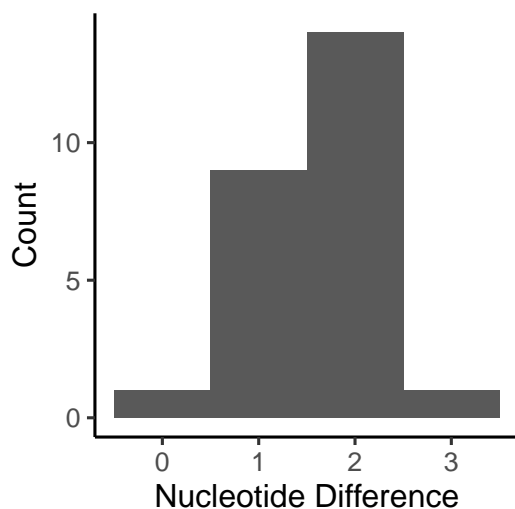
TRBV14*bp01

16 sequences assigned
9 (56.2%) exact matches, in which:
4 unique CDR3
2 unique J



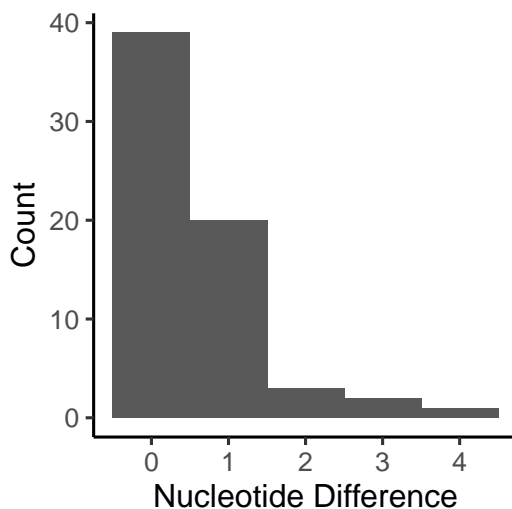
TRBV10-1*bp02

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



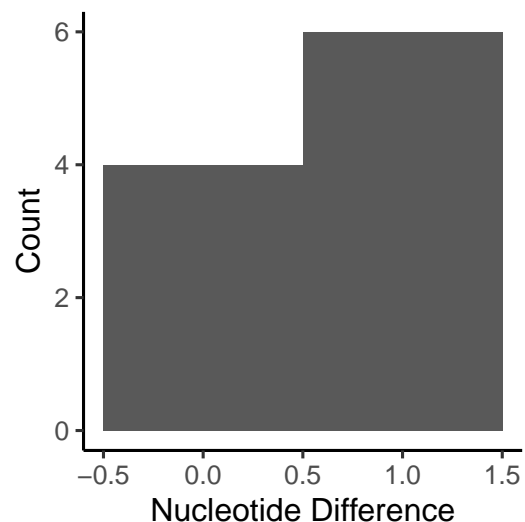
TRBV10-3*bp01

65 sequences assigned
39 (60%) exact matches, in which:
24 unique CDR3
10 unique J



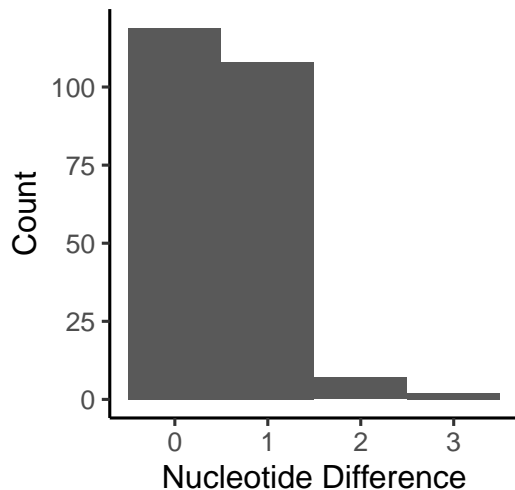
TRBV15*bp02

10 sequences assigned
4 (40%) exact matches, in which:
4 unique CDR3
4 unique J



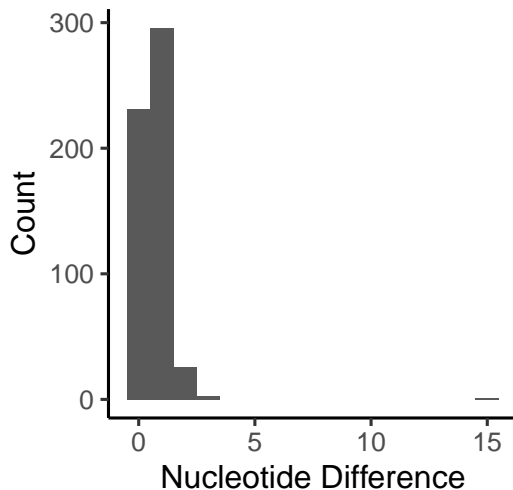
TRBV18*bp01

236 sequences assigned
119 (50.4%) exact matches, in which:
59 unique CDR3
9 unique J



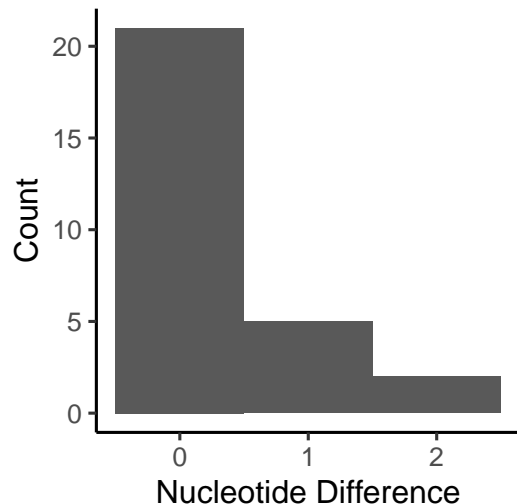
TRBV27*bp01

557 sequences assigned
231 (41.5%) exact matches, in which:
122 unique CDR3
13 unique J



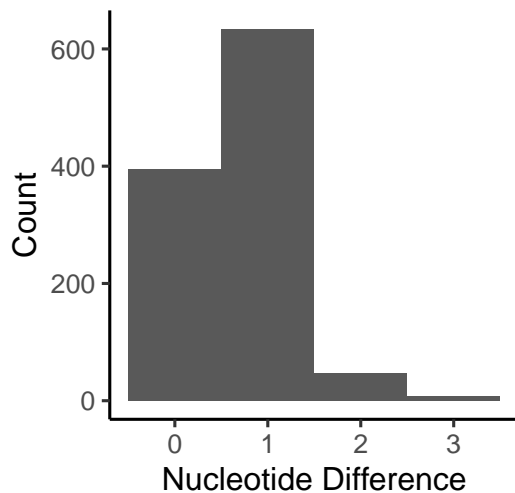
TRBV9*bp01

28 sequences assigned
21 (75%) exact matches, in which:
9 unique CDR3
6 unique J



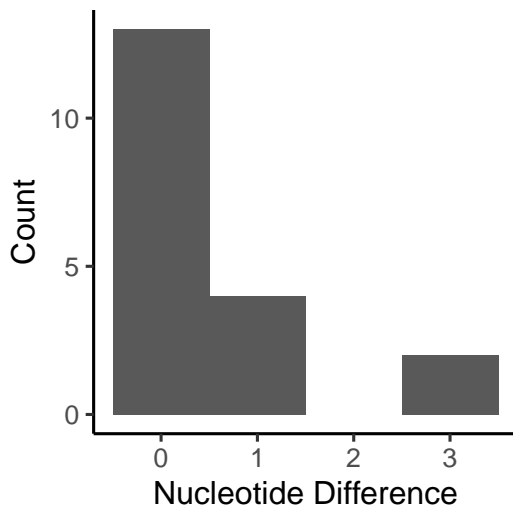
TRBV19*bp01

1083 sequences assigned
394 (36.4%) exact matches, in which:
206 unique CDR3
12 unique J



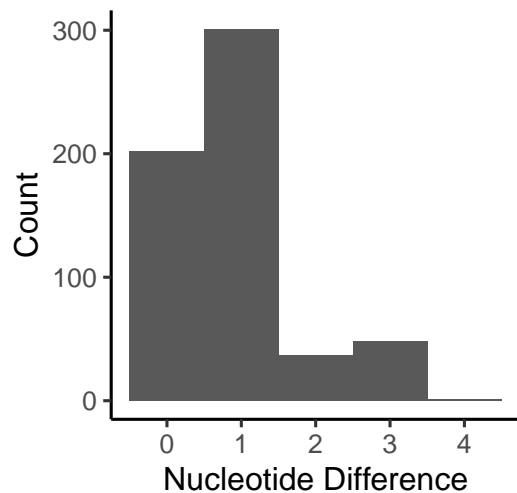
TRBV28*bp01

19 sequences assigned
13 (68.4%) exact matches, in which:
8 unique CDR3
3 unique J



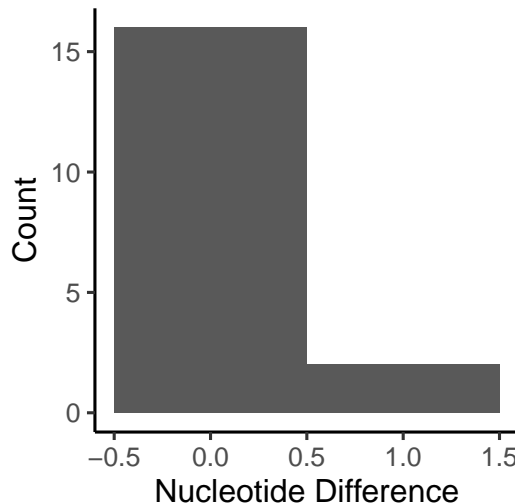
TRBV11-1*bp01

589 sequences assigned
202 (34.3%) exact matches, in which:
114 unique CDR3
10 unique J



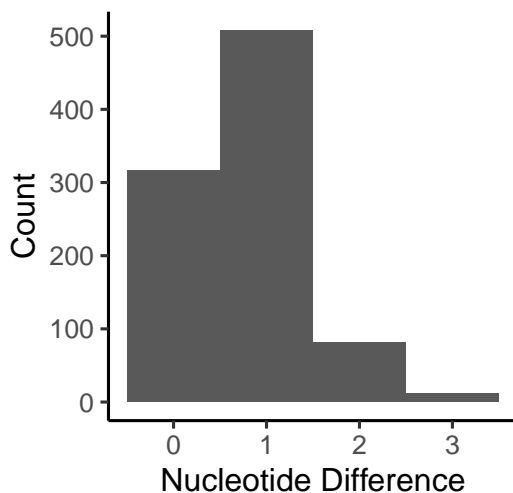
TRBV2*bp01

18 sequences assigned
16 (88.9%) exact matches, in which:
11 unique CDR3
6 unique J



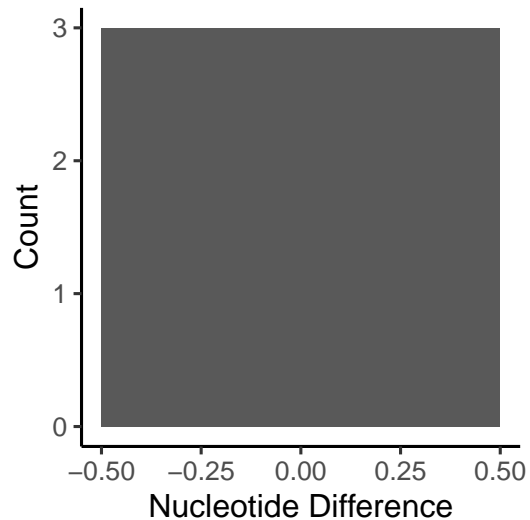
TRBV30*bp01

917 sequences assigned
316 (34.5%) exact matches, in which:
118 unique CDR3
9 unique J



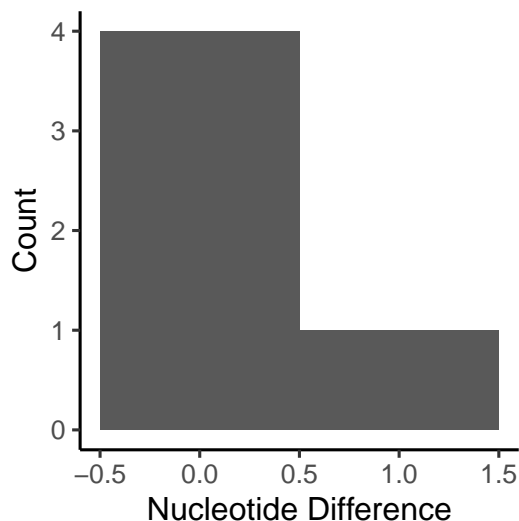
TRBV21-1*bp01

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



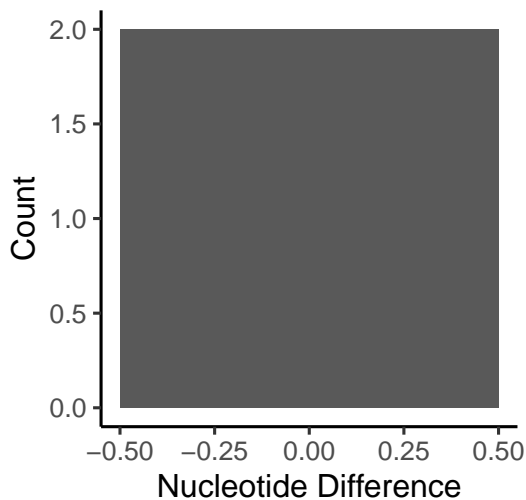
TRBV21–1*bp02

5 sequences assigned
4 (80%) exact matches, in which:
4 unique CDR3
4 unique J



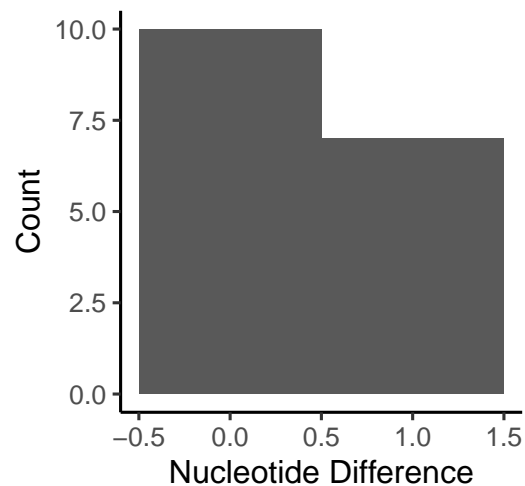
TRBV12–5*bp01

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



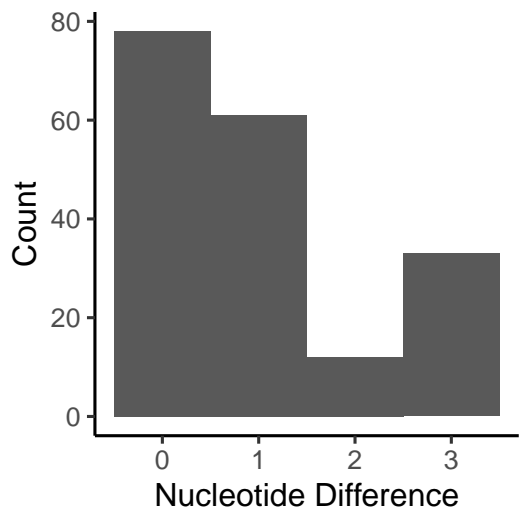
TRBV4–1*bp01

17 sequences assigned
10 (58.8%) exact matches, in which:
6 unique CDR3
4 unique J



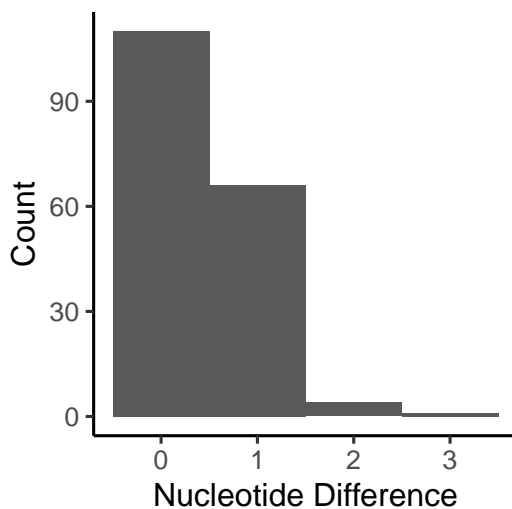
TRBV11–2*bp01

184 sequences assigned
78 (42.4%) exact matches, in which:
44 unique CDR3
11 unique J



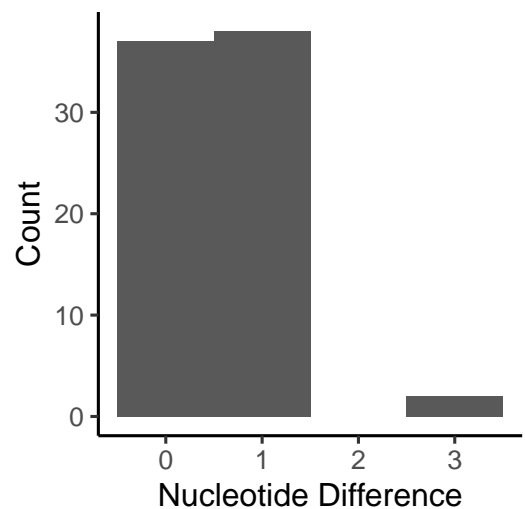
TRBV12–34*bp01

181 sequences assigned
110 (60.8%) exact matches, in which:
57 unique CDR3
12 unique J



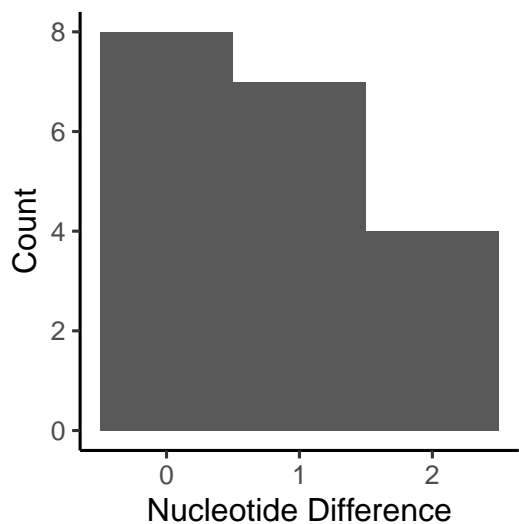
TRBV24–1*bp02

77 sequences assigned
37 (48.1%) exact matches, in which:
15 unique CDR3
7 unique J



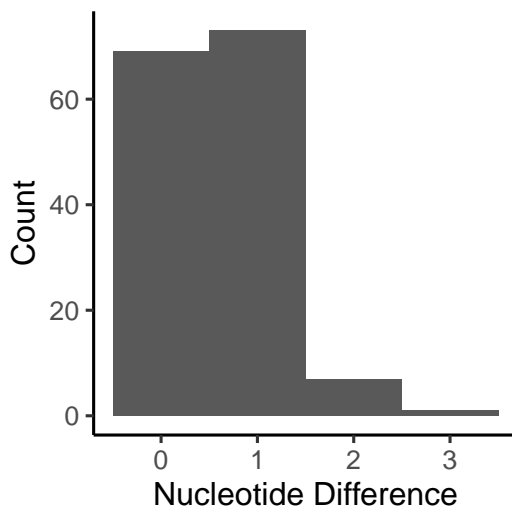
TRBV11–3*bp01

19 sequences assigned
8 (42.1%) exact matches, in which:
7 unique CDR3
4 unique J



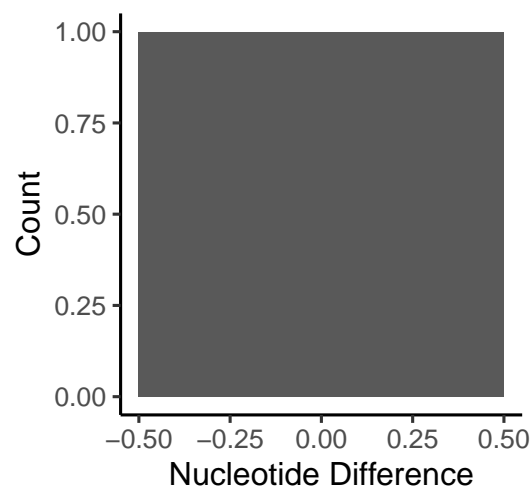
TRBV24–1*bp01

150 sequences assigned
69 (46%) exact matches, in which:
36 unique CDR3
7 unique J



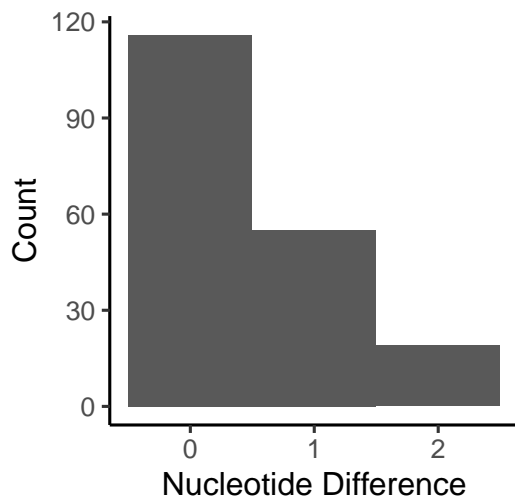
TRBV4–2*bp01

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



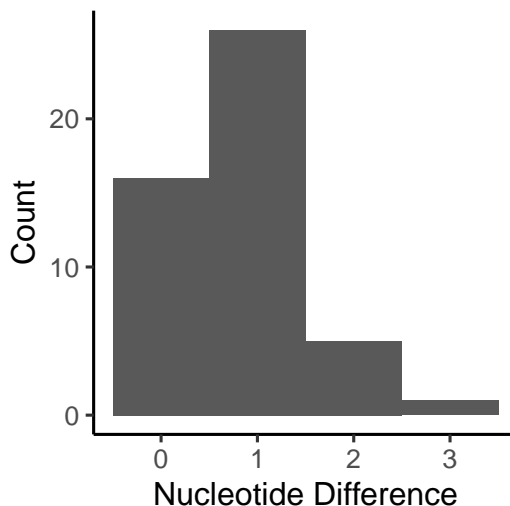
TRBV5-1*bp01

190 sequences assigned
116 (61.1%) exact matches, in which:
35 unique CDR3
7 unique J



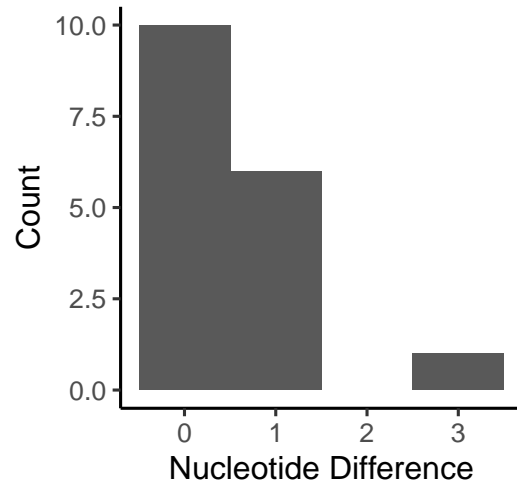
TRBV5-5*bp01

48 sequences assigned
16 (33.3%) exact matches, in which:
7 unique CDR3
3 unique J



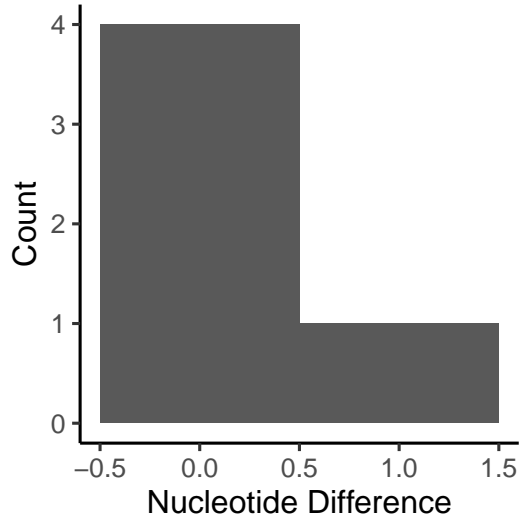
TRBV5-8*bp01

17 sequences assigned
10 (58.8%) exact matches, in which:
4 unique CDR3
4 unique J



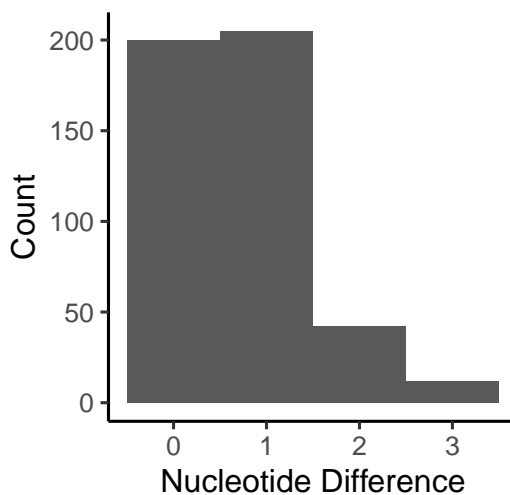
TRBV25-1*bp01

5 sequences assigned
4 (80%) exact matches, in which:
2 unique CDR3
2 unique J



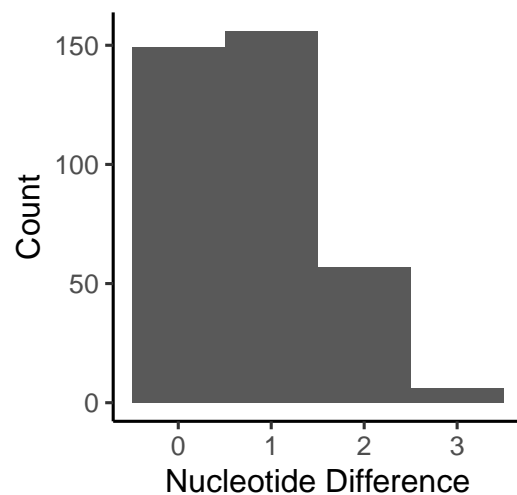
TRBV5-6*bp01

459 sequences assigned
200 (43.6%) exact matches, in which:
112 unique CDR3
11 unique J



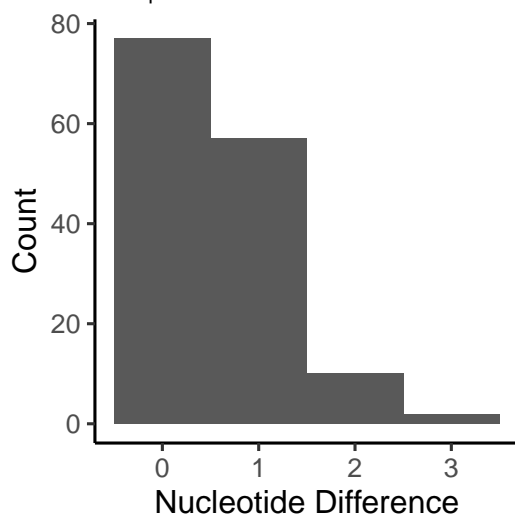
TRBV6-1*bp01

368 sequences assigned
149 (40.5%) exact matches, in which:
89 unique CDR3
11 unique J



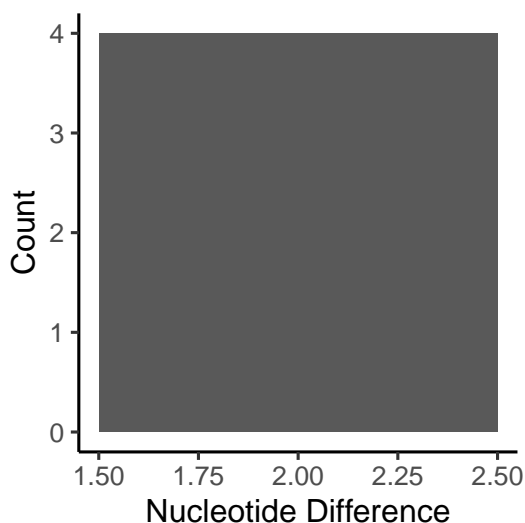
TRBV5-4*bp01

146 sequences assigned
77 (52.7%) exact matches, in which:
37 unique CDR3
7 unique J



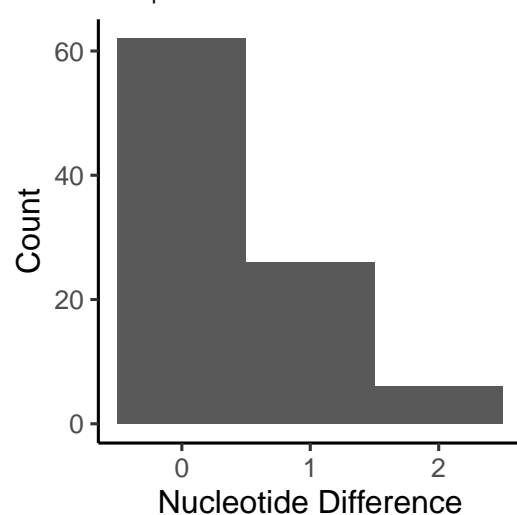
TRBV5-7*bp01

4 sequences assigned
No exact matches.



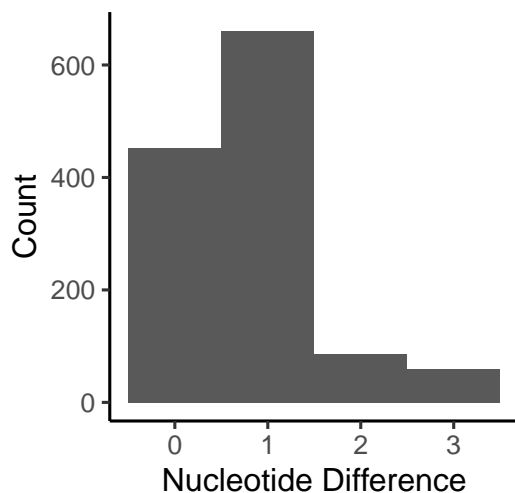
TRBV6-4*bp01

94 sequences assigned
62 (66%) exact matches, in which:
31 unique CDR3
9 unique J



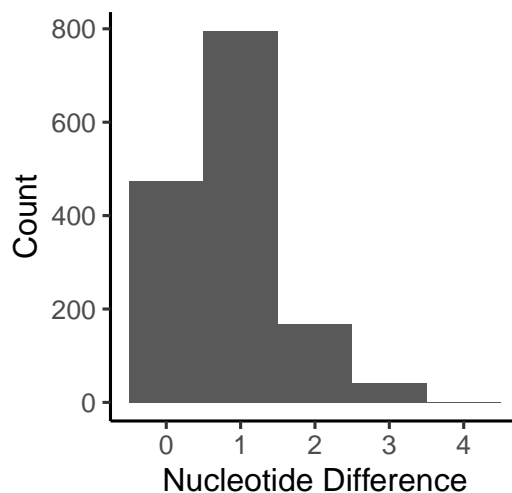
TRBV6-5*bp01

1259 sequences assigned
453 (36%) exact matches, in which:
245 unique CDR3
12 unique J



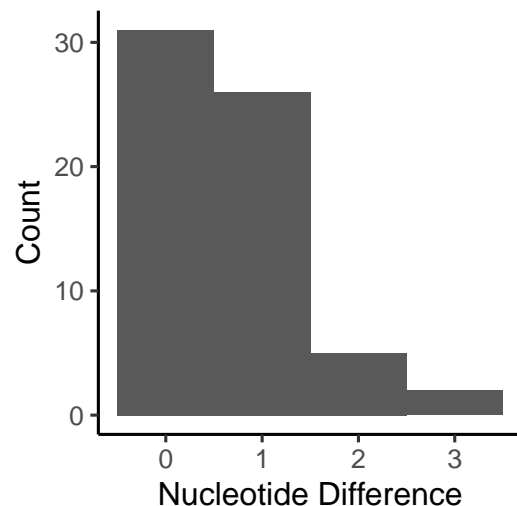
TRBV6-23*bp01

1481 sequences assigned
474 (32%) exact matches, in which:
228 unique CDR3
12 unique J



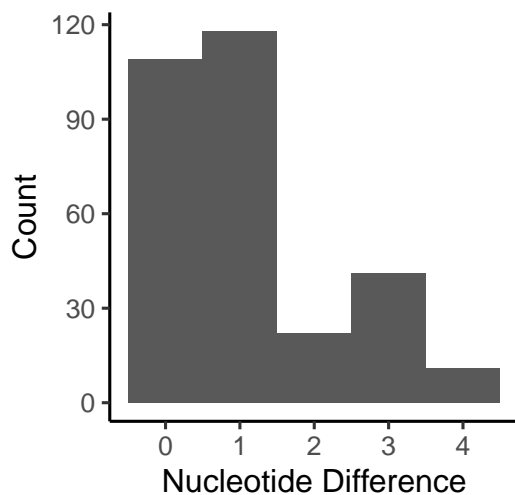
TRBV7-6*bp01

64 sequences assigned
31 (48.4%) exact matches, in which:
18 unique CDR3
5 unique J



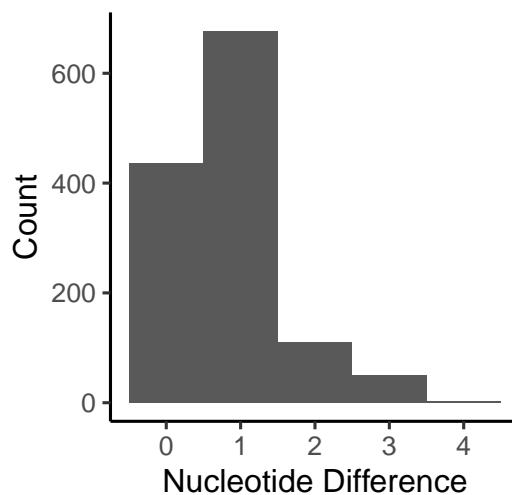
TRBV6-6*bp01

301 sequences assigned
109 (36.2%) exact matches, in which:
68 unique CDR3
12 unique J



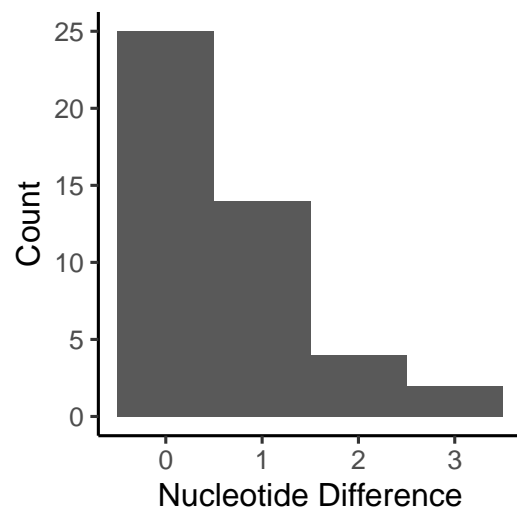
TRBV7-2*bp01

1276 sequences assigned
436 (34.2%) exact matches, in which:
151 unique CDR3
13 unique J



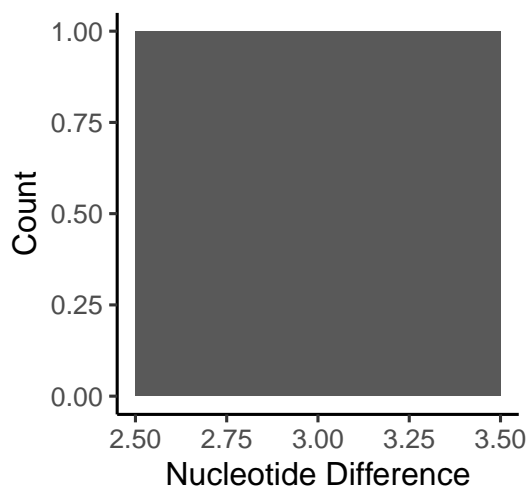
TRBV7-7*bp01

45 sequences assigned
25 (55.6%) exact matches, in which:
4 unique CDR3
2 unique J



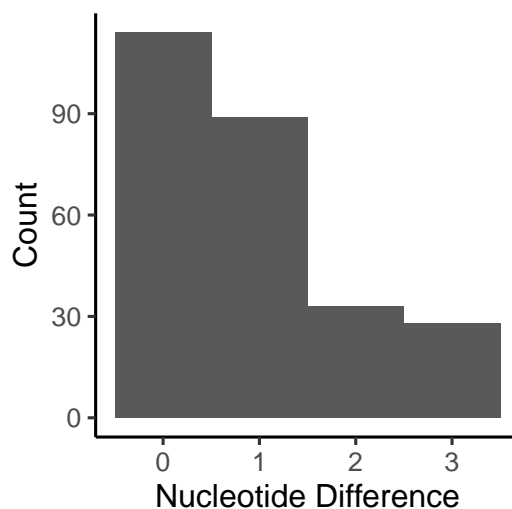
TRBV6-9*bp01

1 sequences assigned
No exact matches.



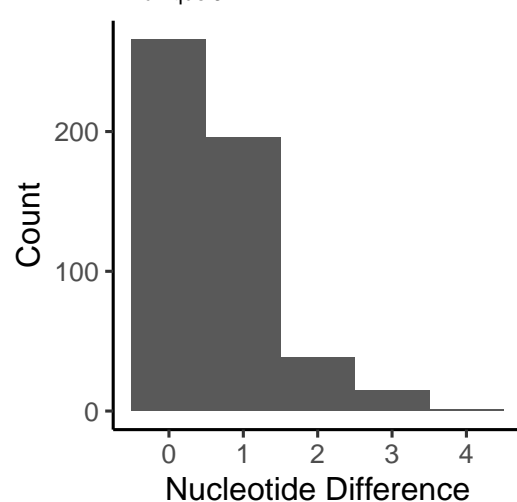
TRBV7-3*bp01

264 sequences assigned
114 (43.2%) exact matches, in which:
50 unique CDR3
11 unique J



TRBV7-9*bp01

516 sequences assigned
266 (51.6%) exact matches, in which:
102 unique CDR3
11 unique J



TRBV29-1*bp01

295 sequences assigned
96 (32.5%) exact matches, in which:
67 unique CDR3
10 unique J

