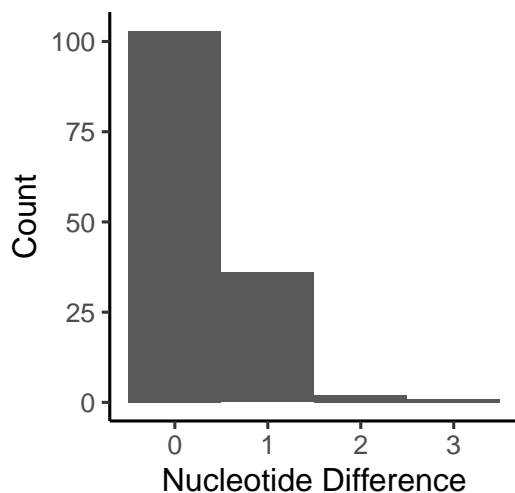


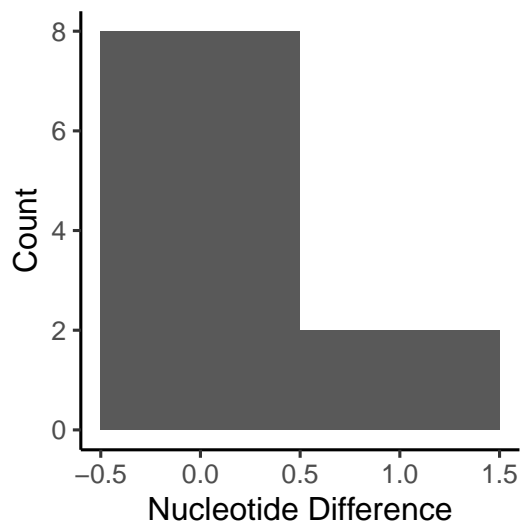
TRBV20–1*bp01

142 sequences assigned
103 (72.5%) exact matches, in which:
72 unique CDR3
12 unique J



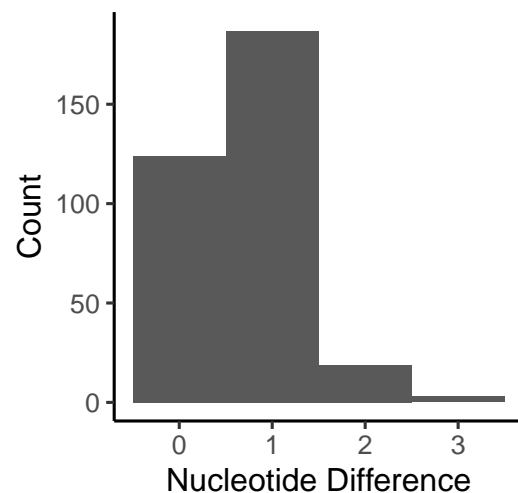
TRBV10–3*bp01

10 sequences assigned
8 (80%) exact matches, in which:
8 unique CDR3
6 unique J



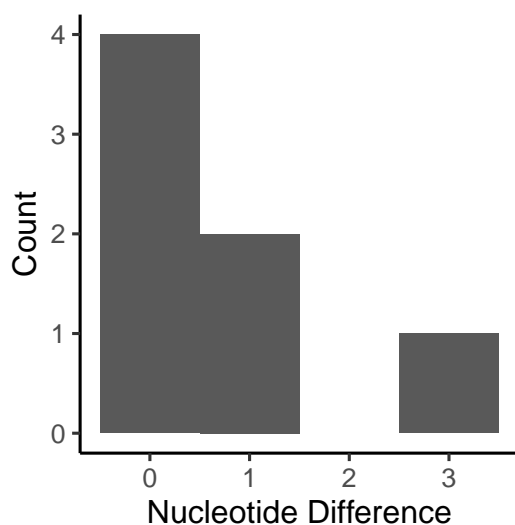
TRBV15*bp02

333 sequences assigned
124 (37.2%) exact matches, in which:
75 unique CDR3
11 unique J



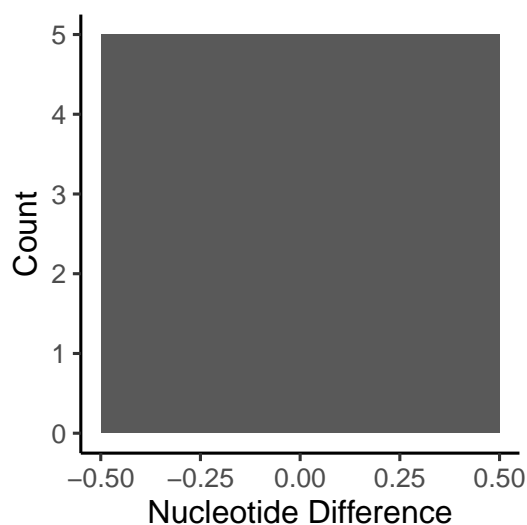
TRBV10–1*bp01

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



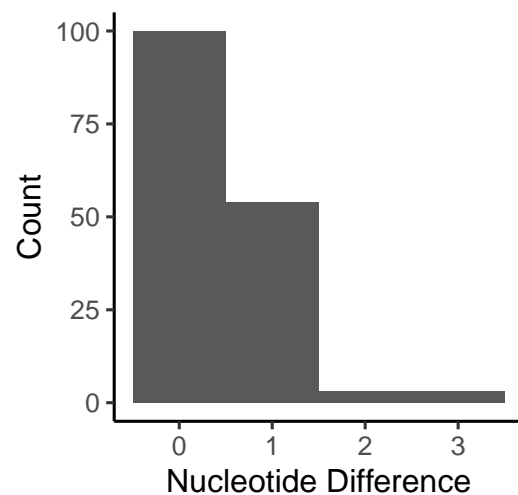
TRBV13*bp01

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



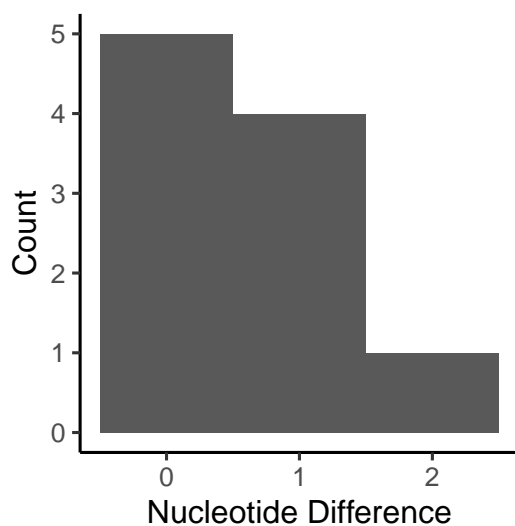
TRBV18*bp01

160 sequences assigned
100 (62.5%) exact matches, in which:
71 unique CDR3
12 unique J



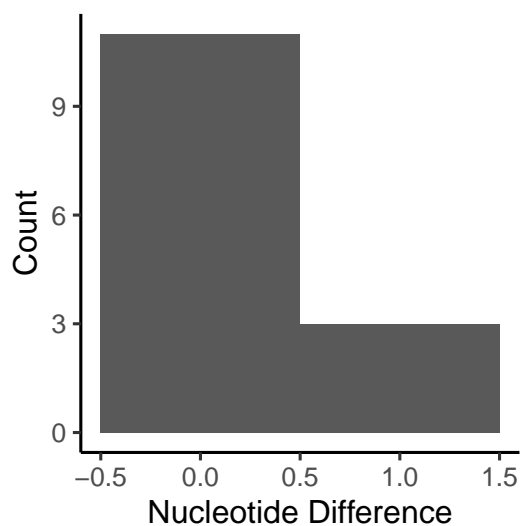
TRBV10–2*bp01

10 sequences assigned
5 (50%) exact matches, in which:
5 unique CDR3
3 unique J



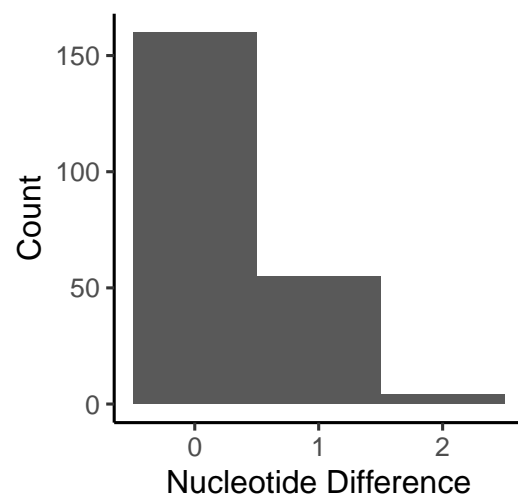
TRBV14*bp01

14 sequences assigned
11 (78.6%) exact matches, in which:
8 unique CDR3
5 unique J



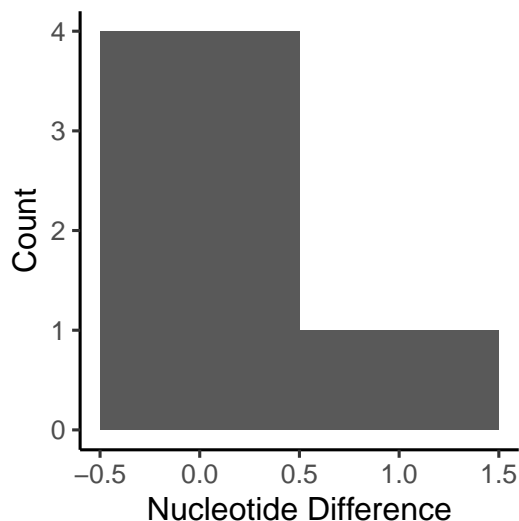
TRBV19*bp01

219 sequences assigned
160 (73.1%) exact matches, in which:
119 unique CDR3
13 unique J



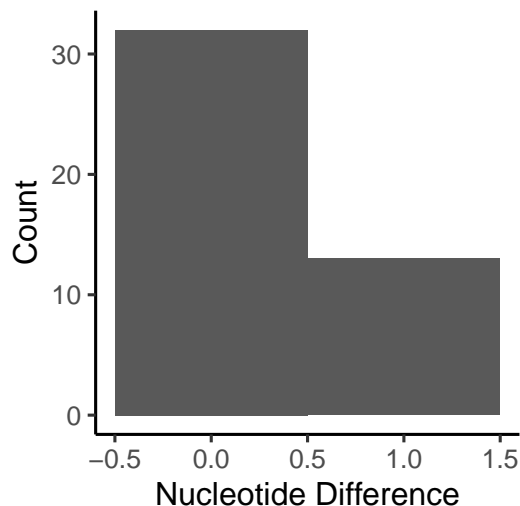
TRBV2*bp01

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



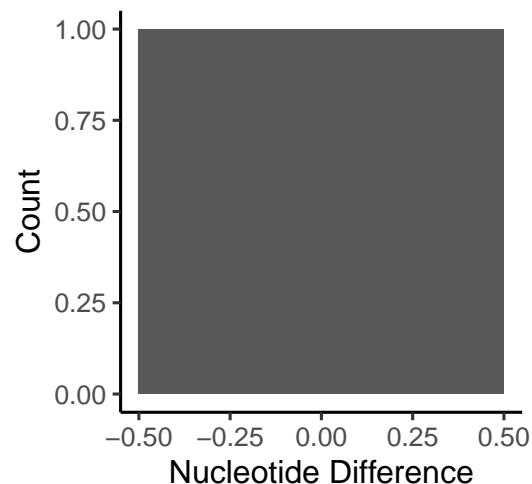
TRBV30*bp01

45 sequences assigned
32 (71.1%) exact matches, in which:
20 unique CDR3
11 unique J



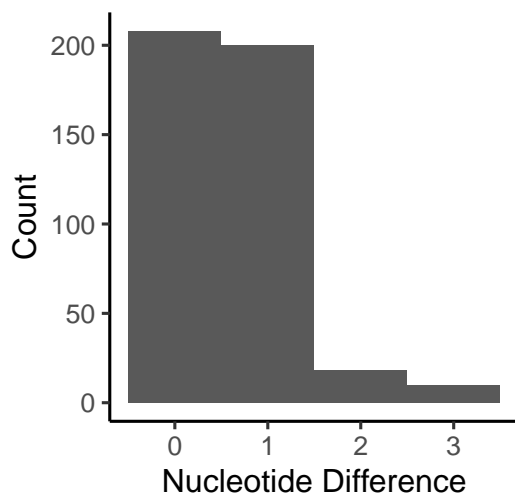
TRBV21-1*bp02

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



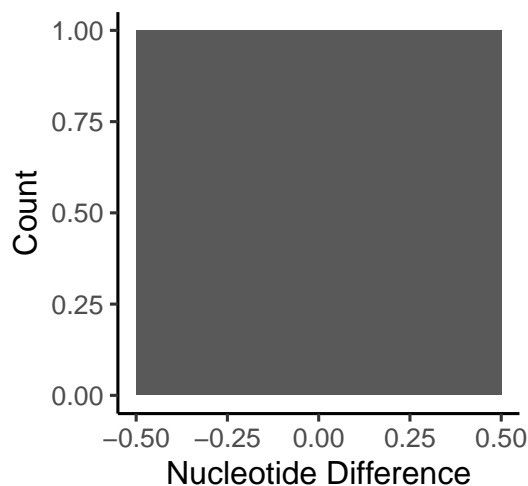
TRBV27*bp01

436 sequences assigned
208 (47.7%) exact matches, in which:
149 unique CDR3
12 unique J



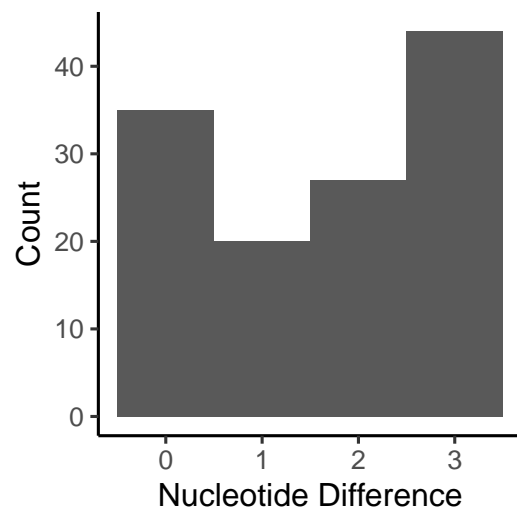
TRBV9*bp01

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



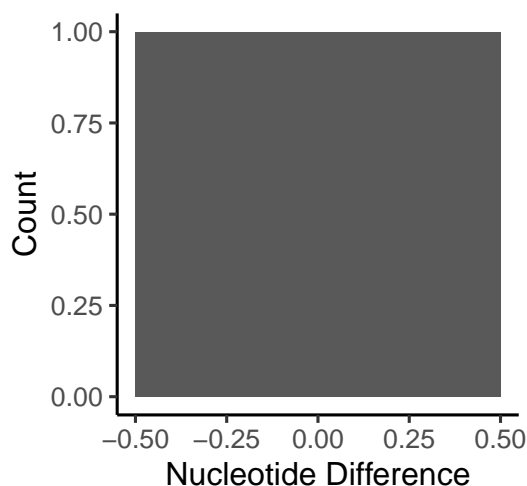
TRBV11-2*bp01

126 sequences assigned
35 (27.8%) exact matches, in which:
27 unique CDR3
8 unique J



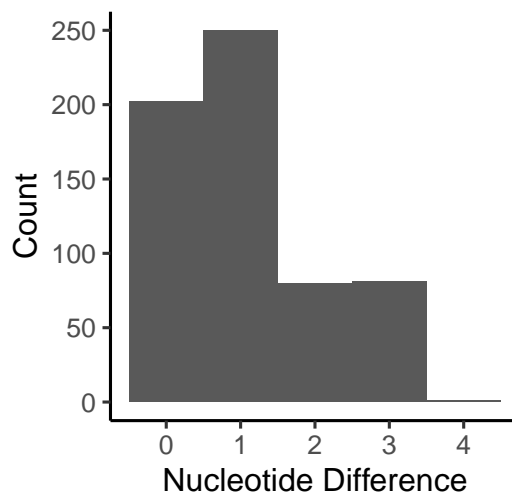
TRBV28*bp01

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



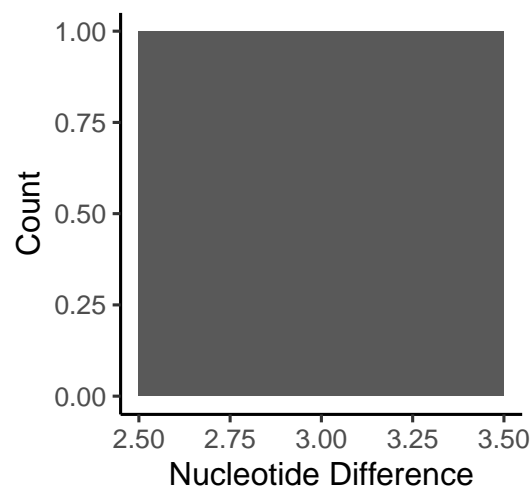
TRBV11-1*bp01

614 sequences assigned
202 (32.9%) exact matches, in which:
122 unique CDR3
12 unique J



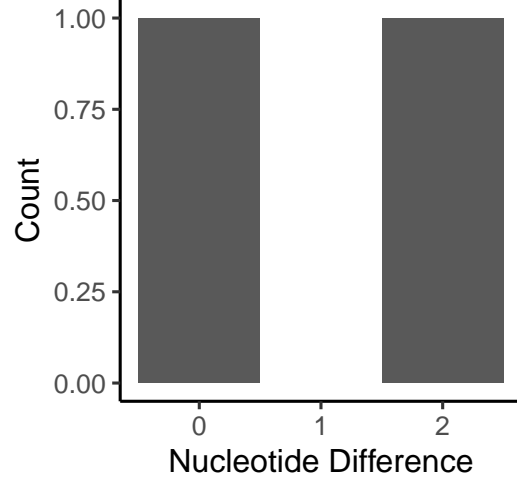
TRBV11-3*bp01

1 sequences assigned
No exact matches.



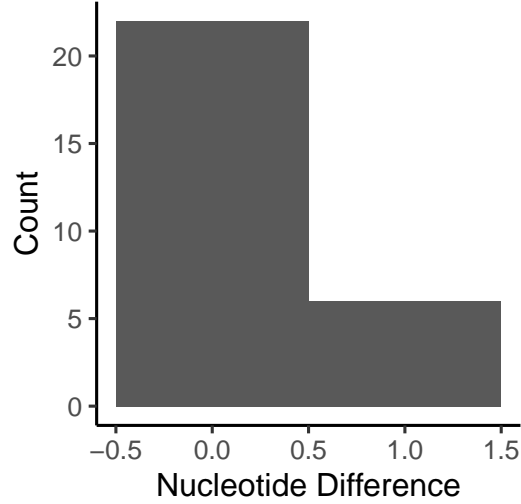
TRBV12–5*bp01

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



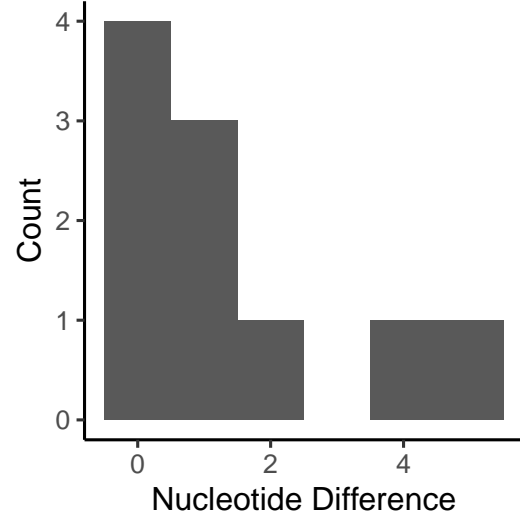
TRBV24–1*bp02

28 sequences assigned
22 (78.6%) exact matches, in which:
18 unique CDR3
6 unique J



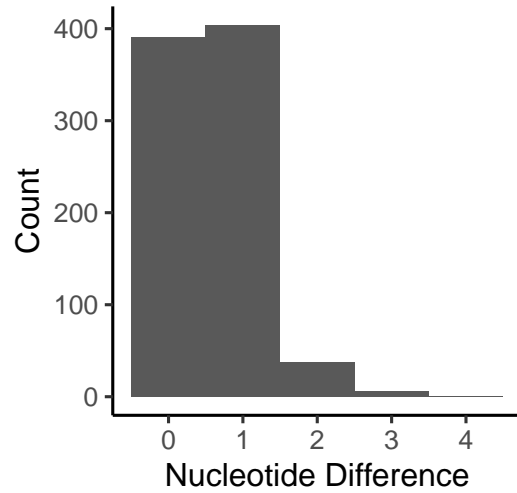
TRBV5–3*bp01

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



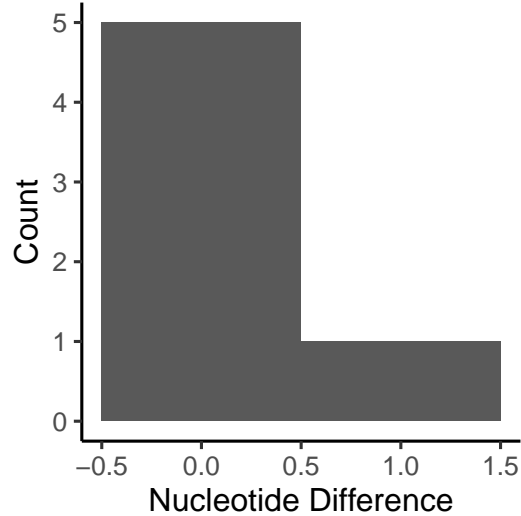
TRBV12–34*bp01

839 sequences assigned
391 (46.6%) exact matches, in which:
209 unique CDR3
12 unique J



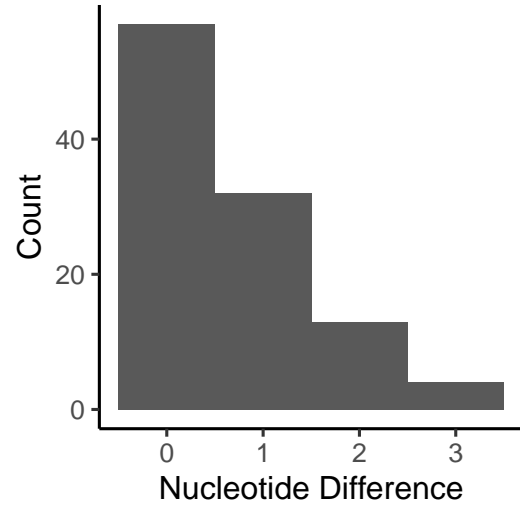
TRBV4–2*bp01

6 sequences assigned
5 (83.3%) exact matches, in which:
4 unique CDR3
3 unique J



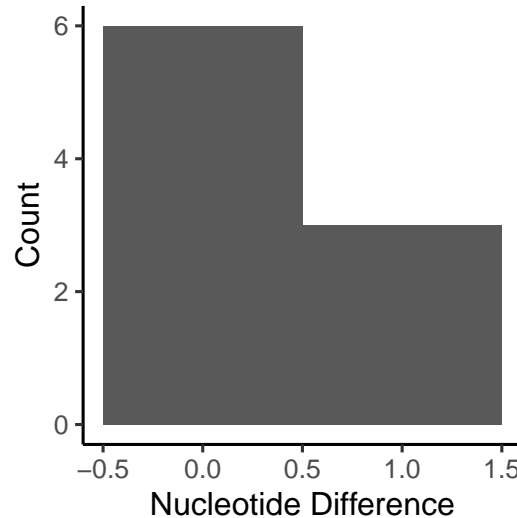
TRBV5–4*bp01

106 sequences assigned
57 (53.8%) exact matches, in which:
49 unique CDR3
10 unique J



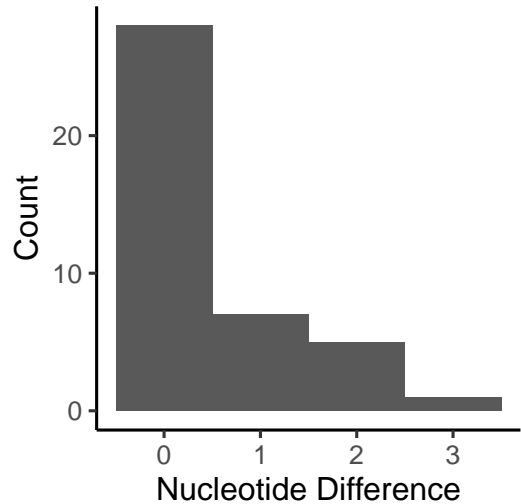
TRBV4–1*bp01

9 sequences assigned
6 (66.7%) exact matches, in which:
6 unique CDR3
4 unique J



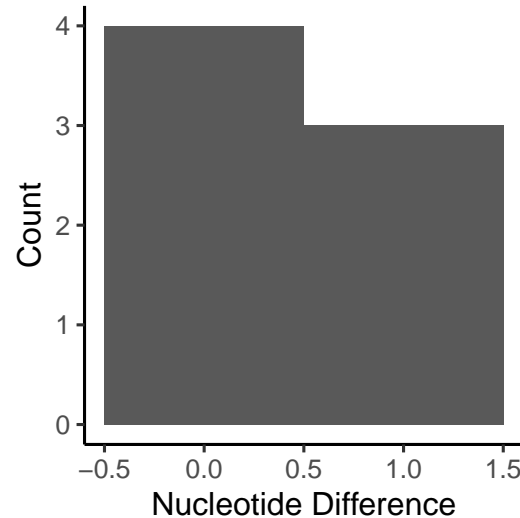
TRBV5–1*bp01

41 sequences assigned
28 (68.3%) exact matches, in which:
24 unique CDR3
10 unique J



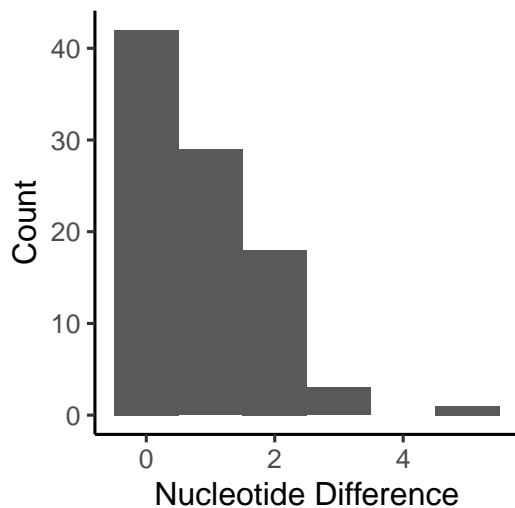
TRBV5–5*bp01

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



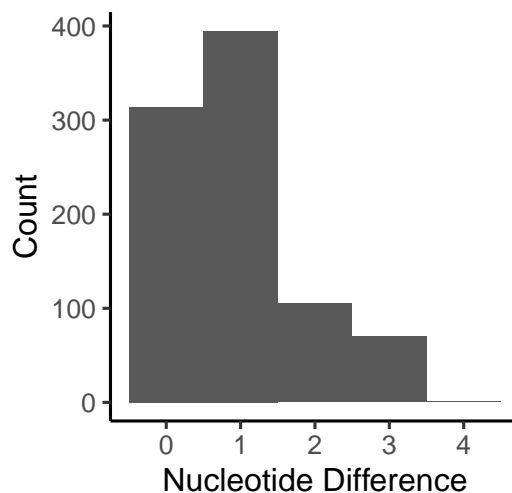
TRBV5-6*bp01

93 sequences assigned
42 (45.2%) exact matches, in which:
35 unique CDR3
10 unique J



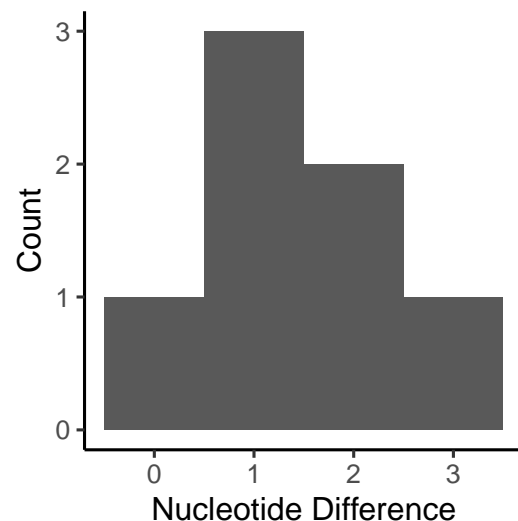
TRBV6-5*bp01

885 sequences assigned
314 (35.5%) exact matches, in which:
238 unique CDR3
12 unique J



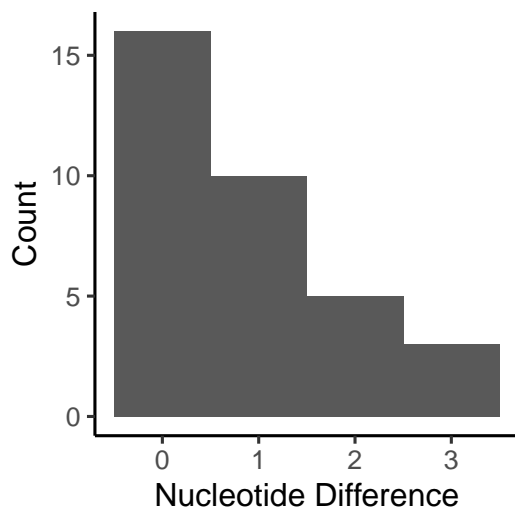
TRBV6-9*bp01

7 sequences assigned
1 (14.3%) exact matches, in which:
1 unique CDR3
1 unique J



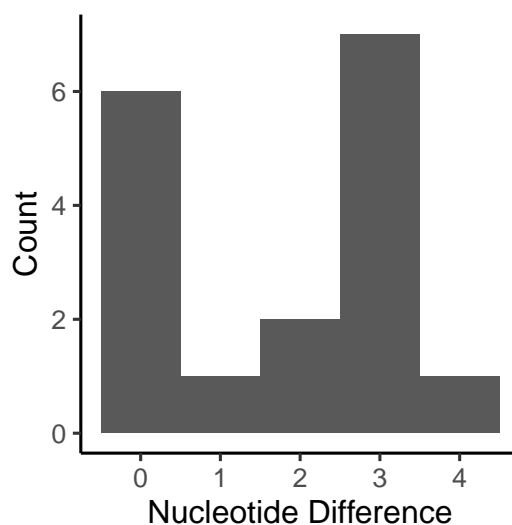
TRBV6-1*bp01

34 sequences assigned
16 (47.1%) exact matches, in which:
15 unique CDR3
6 unique J



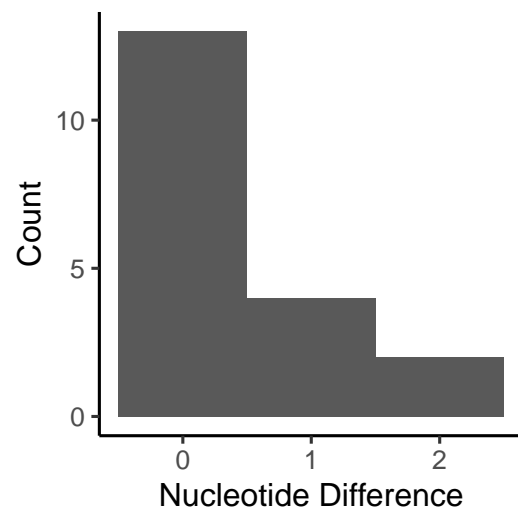
TRBV6-6*bp01

17 sequences assigned
6 (35.3%) exact matches, in which:
6 unique CDR3
4 unique J



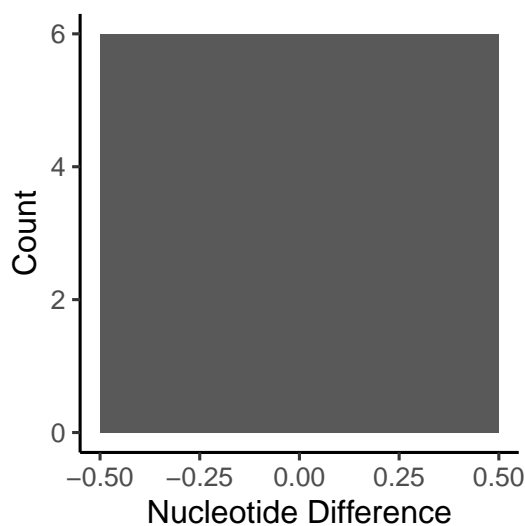
TRBV6-23*bp01

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



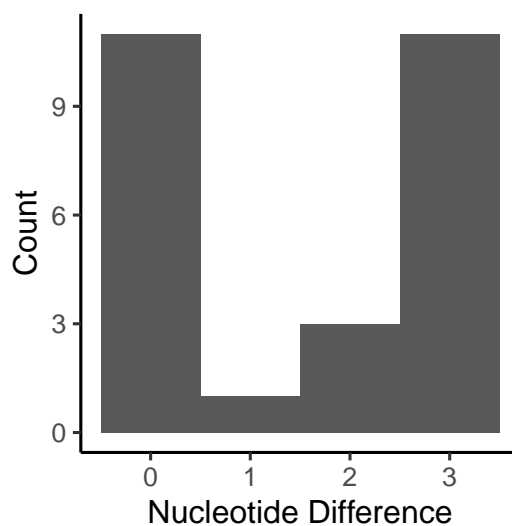
TRBV6-4*bp01

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



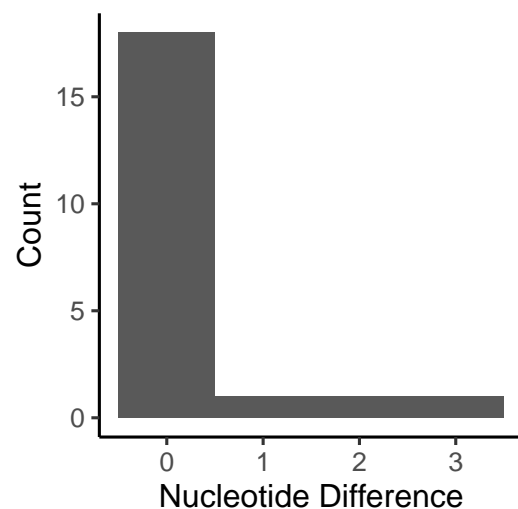
TRBV6-6*bp02

26 sequences assigned
11 (42.3%) exact matches, in which:
10 unique CDR3
6 unique J



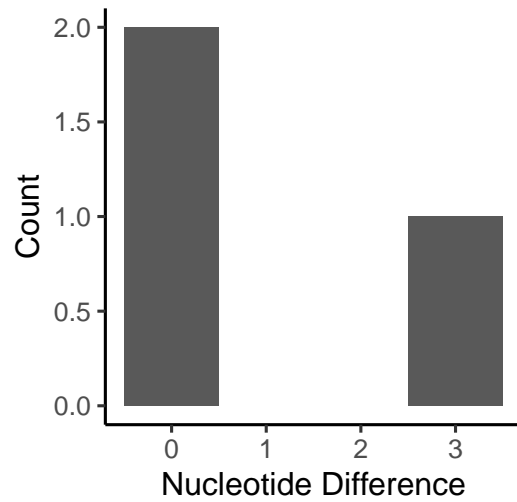
TRBV7-2*bp02

21 sequences assigned
18 (85.7%) exact matches, in which:
15 unique CDR3
9 unique J



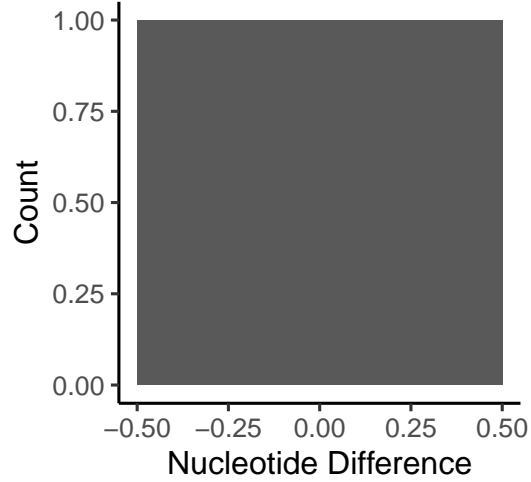
TRBV7-3*bp01

3 sequences assigned
2 (66.7%) exact matches, in which:
2 unique CDR3
2 unique J



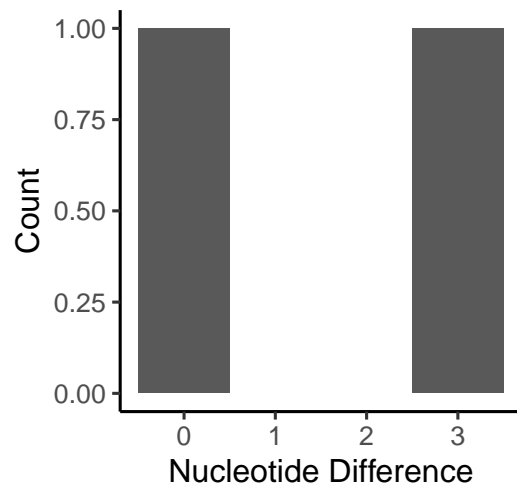
TRBV7-7*bp01

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



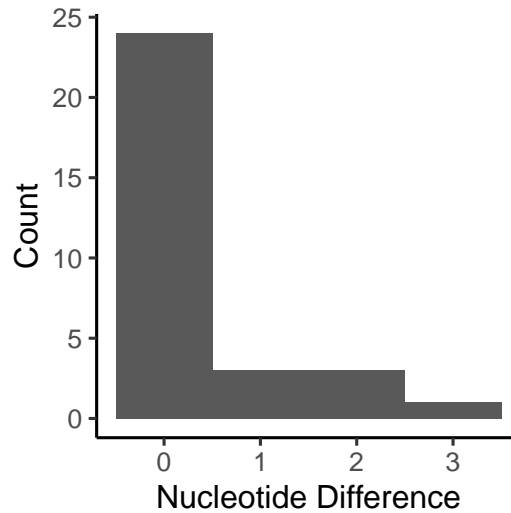
TRBV7-4*bp01

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



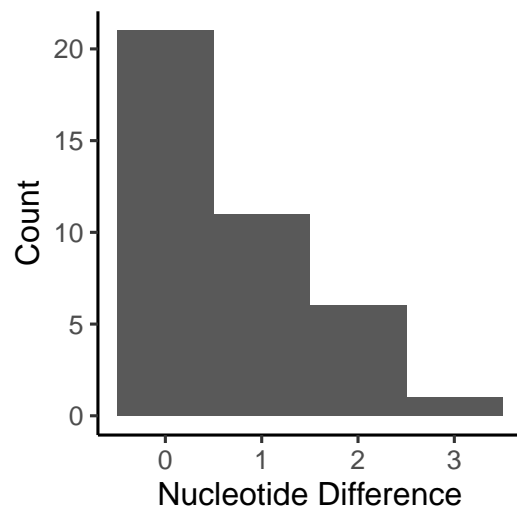
TRBV7-9*bp01

31 sequences assigned
24 (77.4%) exact matches, in which:
21 unique CDR3
9 unique J



TRBV7-6*bp01

39 sequences assigned
21 (53.8%) exact matches, in which:
13 unique CDR3
7 unique J



TRBV29-1*bp01

312 sequences assigned
160 (51.3%) exact matches, in which:
106 unique CDR3
11 unique J

