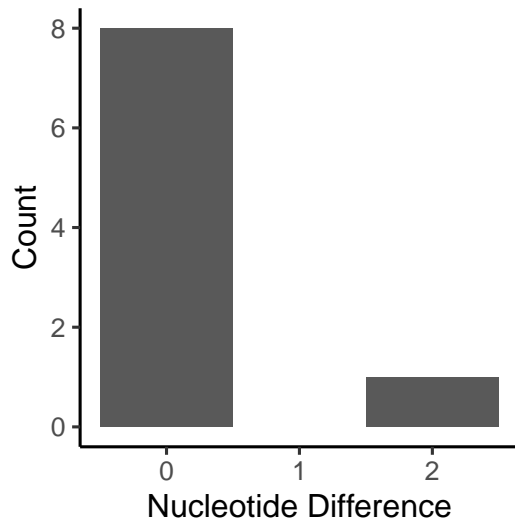


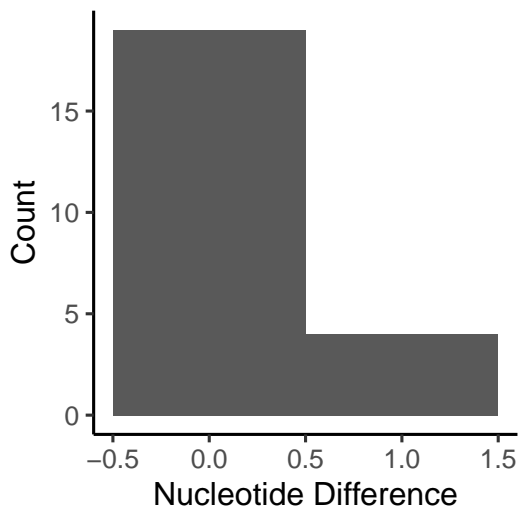
TRBV20–1*bp01

9 sequences assigned
8 (88.9%) exact matches, in which:
8 unique CDR3
5 unique J



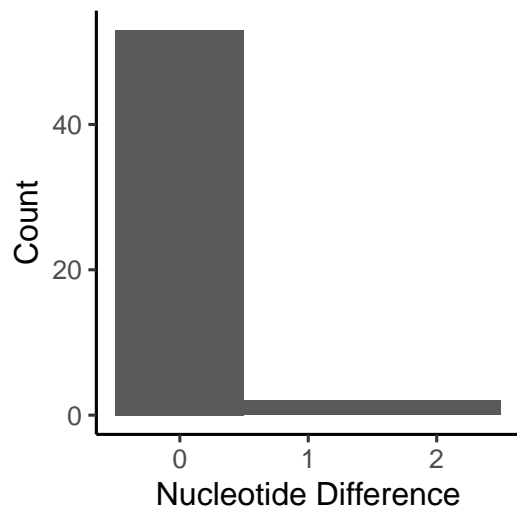
TRBV13*bp01

23 sequences assigned
19 (82.6%) exact matches, in which:
18 unique CDR3
9 unique J



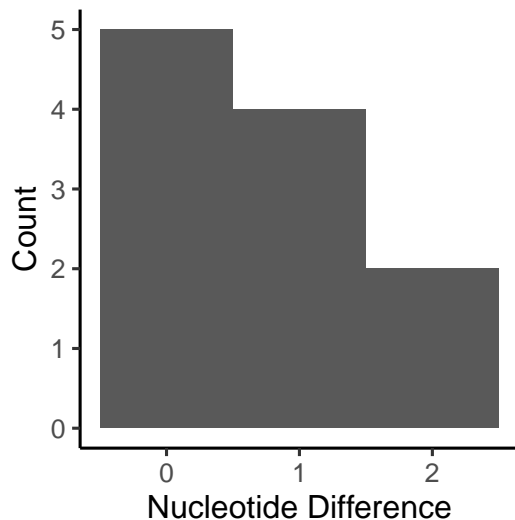
TRBV18*bp01

57 sequences assigned
53 (93%) exact matches, in which:
47 unique CDR3
11 unique J



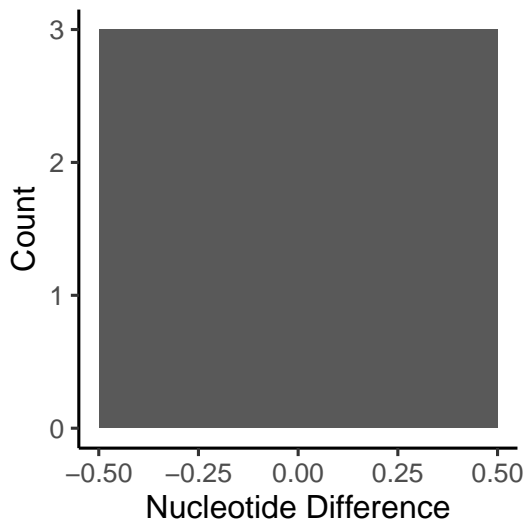
TRBV10–1*bp02

11 sequences assigned
5 (45.5%) exact matches, in which:
4 unique CDR3
4 unique J



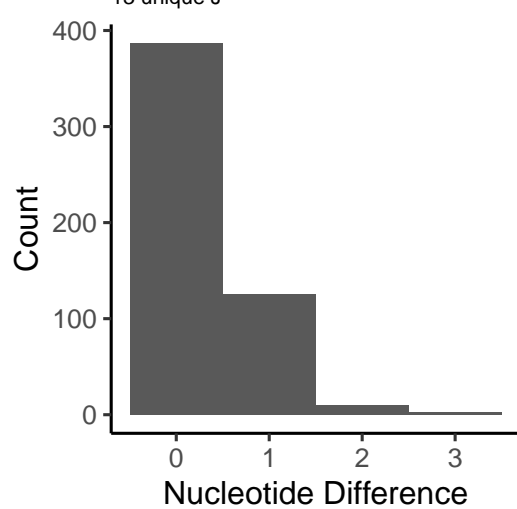
TRBV14*bp01

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



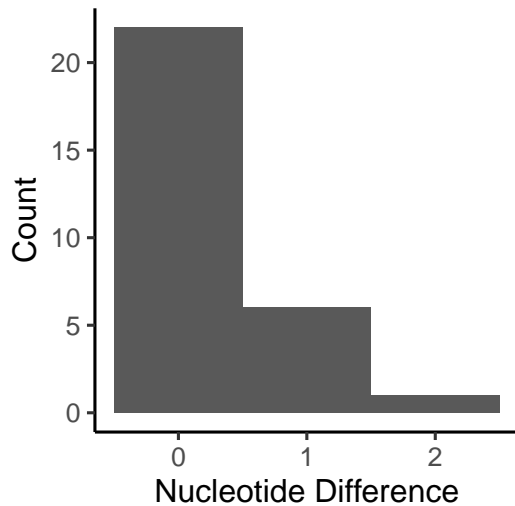
TRBV19*bp01

525 sequences assigned
387 (73.7%) exact matches, in which:
303 unique CDR3
13 unique J



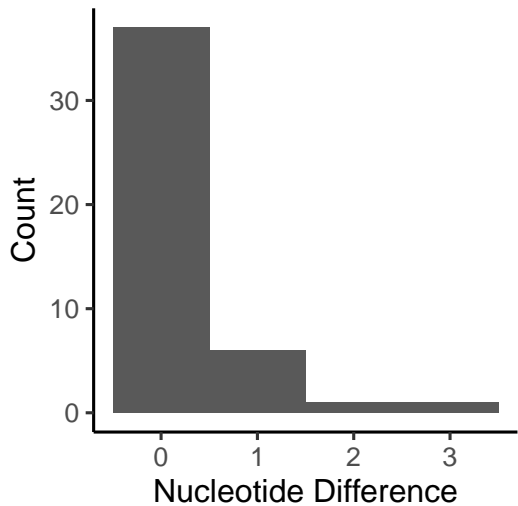
TRBV10–3*bp01

29 sequences assigned
22 (75.9%) exact matches, in which:
21 unique CDR3
11 unique J



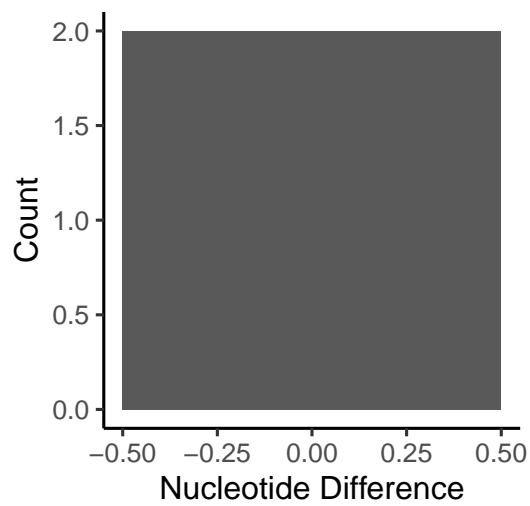
TRBV15*bp02

45 sequences assigned
37 (82.2%) exact matches, in which:
33 unique CDR3
10 unique J



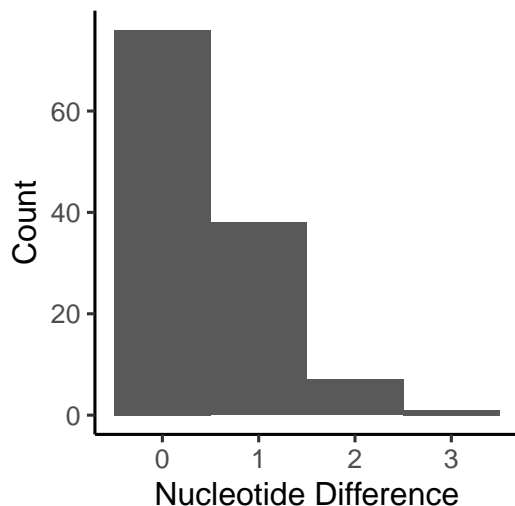
TRBV2*bp01

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



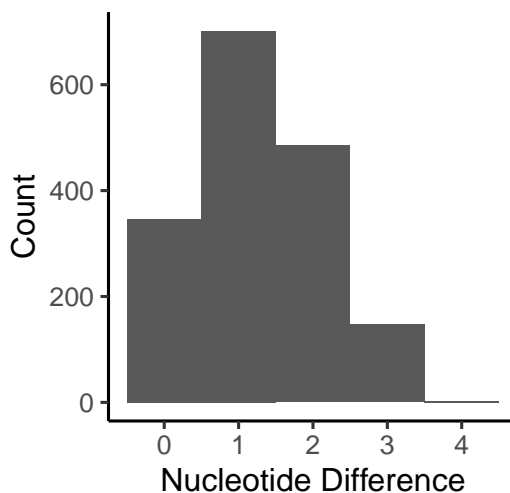
TRBV27*bp01

122 sequences assigned
76 (62.3%) exact matches, in which:
68 unique CDR3
12 unique J



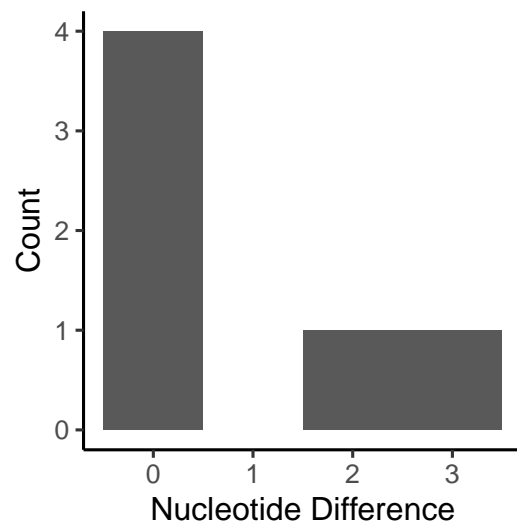
TRBV11–1*bp01

1684 sequences assigned
347 (20.6%) exact matches, in which:
184 unique CDR3
11 unique J



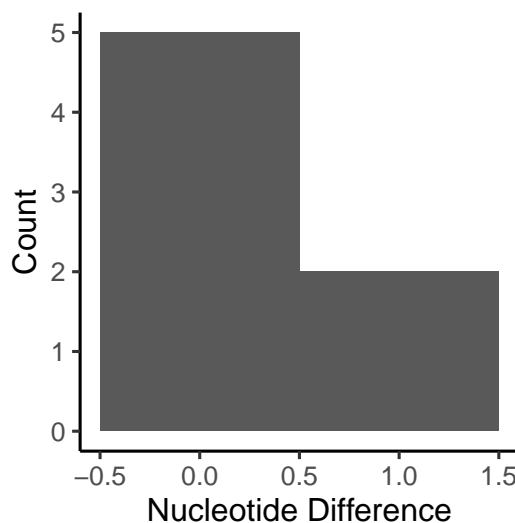
TRBV11–2*bp01

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



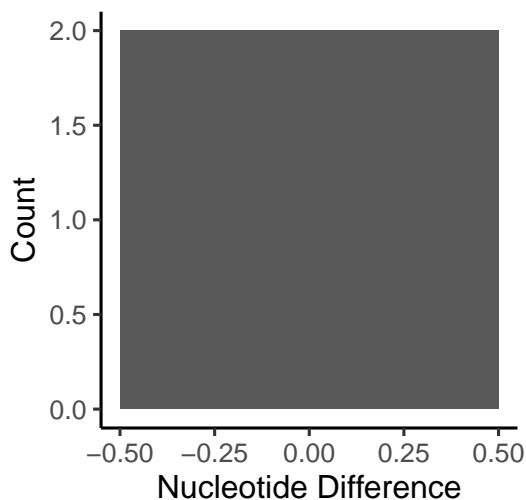
TRBV28*bp01

7 sequences assigned
5 (71.4%) exact matches, in which:
3 unique CDR3
2 unique J



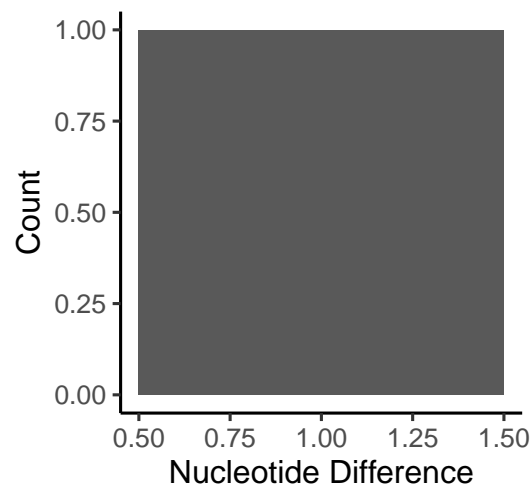
TRBV21–1*bp01

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



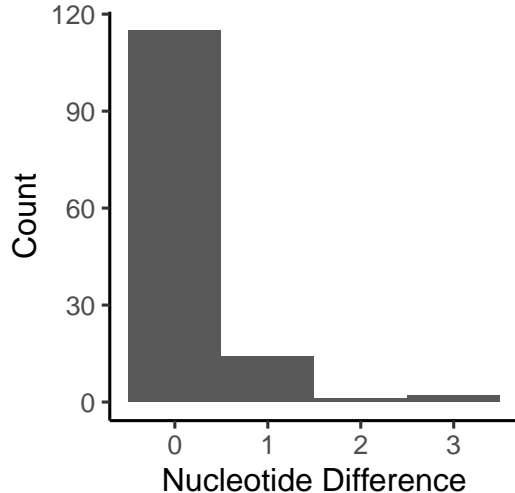
TRBV11–3*bp01

1 sequences assigned
No exact matches.



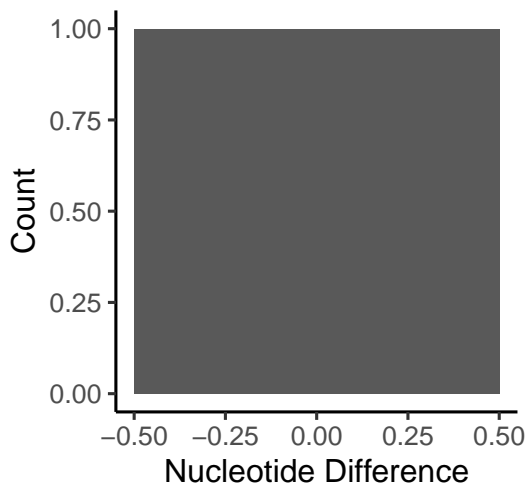
TRBV30*bp01

132 sequences assigned
115 (87.1%) exact matches, in which:
99 unique CDR3
13 unique J



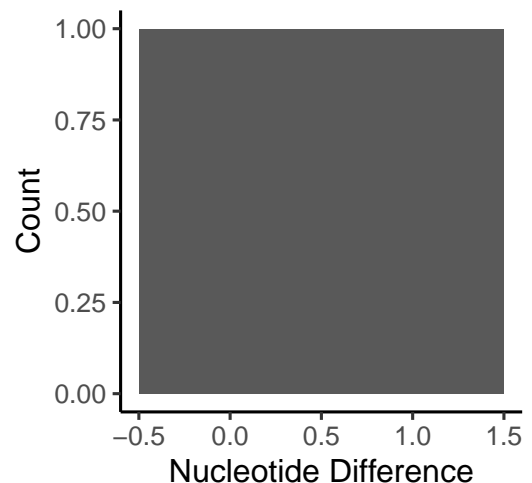
TRBV21–1*bp02

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



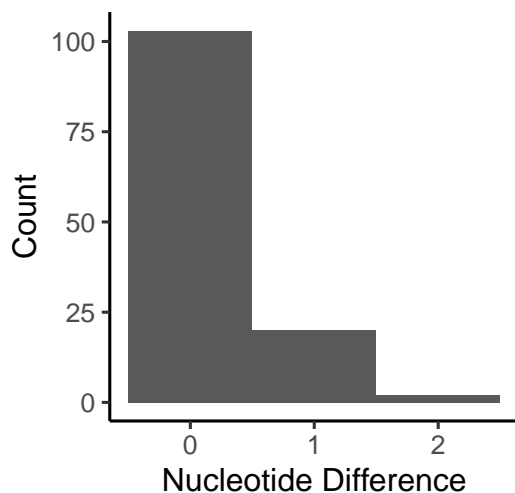
TRBV12–5*bp01

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



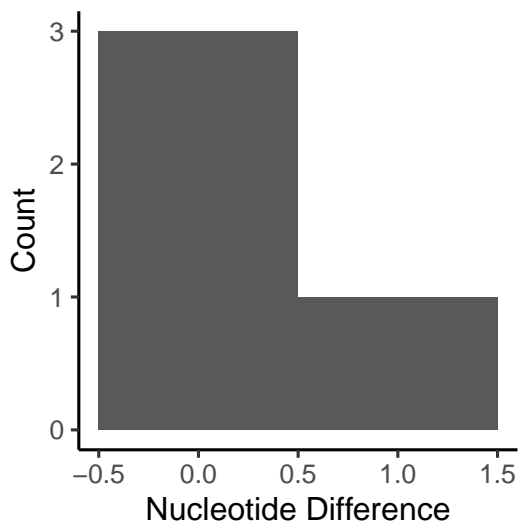
TRBV12–34*bp01

125 sequences assigned
103 (82.4%) exact matches, in which:
86 unique CDR3
13 unique J



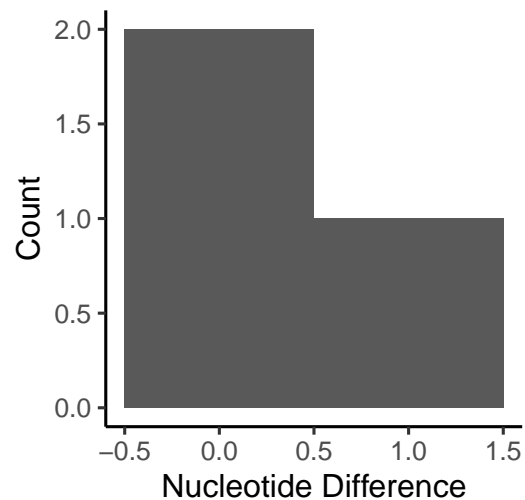
TRBV4–2*bp01

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



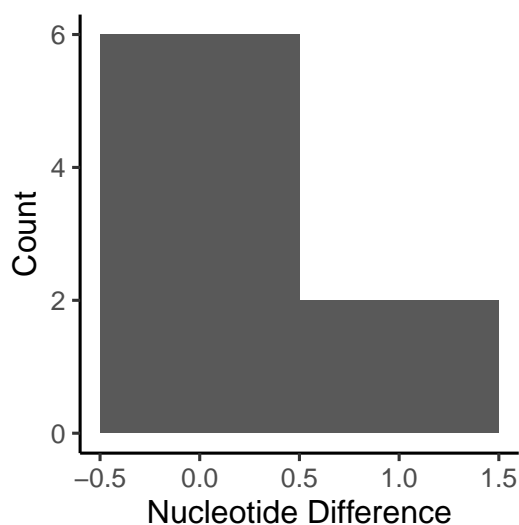
TRBV5–3*bp01

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



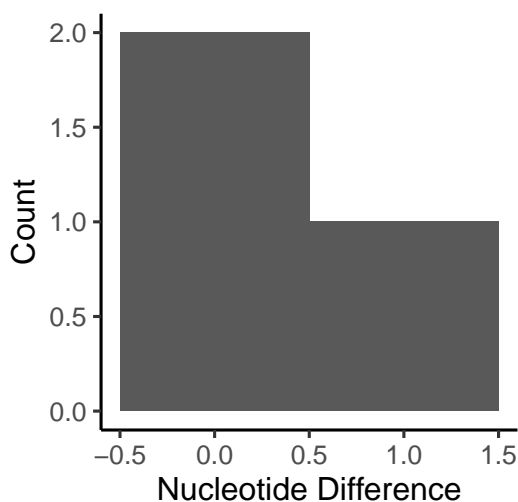
TRBV24–1*bp01

8 sequences assigned
6 (75%) exact matches, in which:
6 unique CDR3
4 unique J



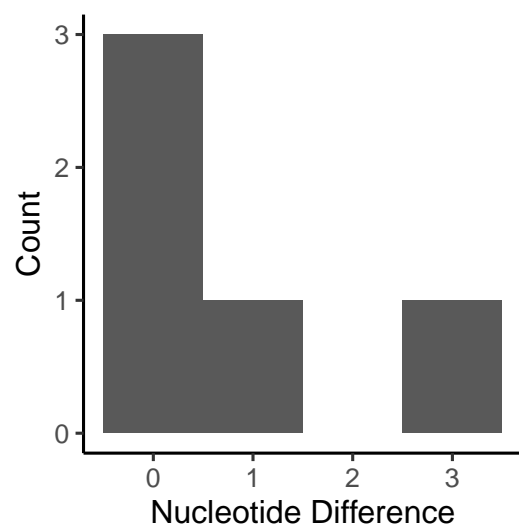
TRBV4–3*bp01

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



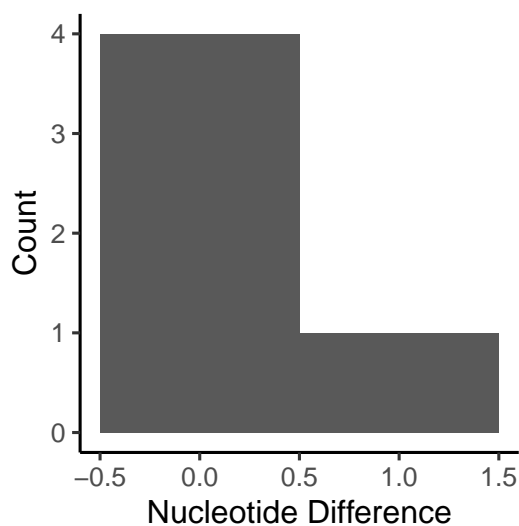
TRBV5–3*bp02

5 sequences assigned
3 (60%) exact matches, in which:
2 unique CDR3
2 unique J



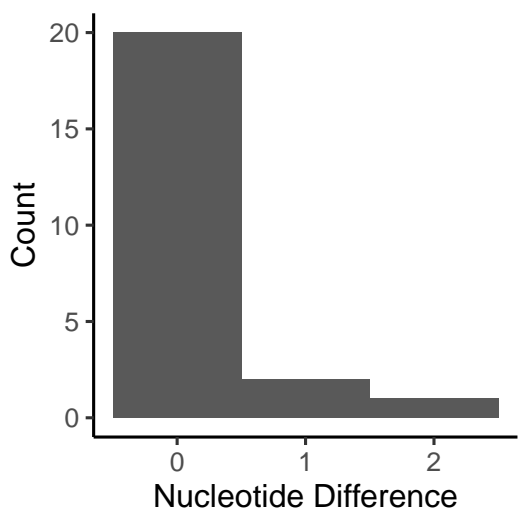
TRBV24–1*bp02

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



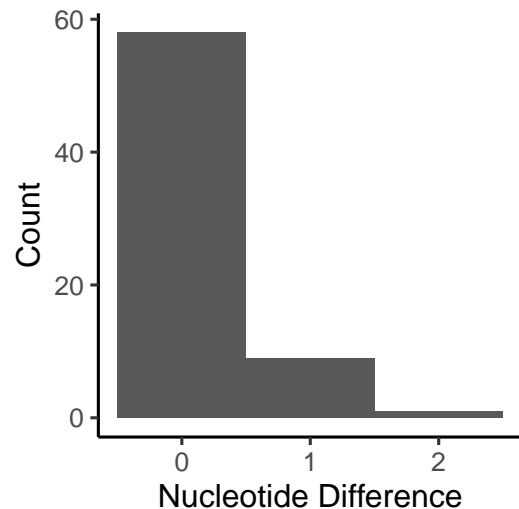
TRBV5–1*bp01

23 sequences assigned
20 (87%) exact matches, in which:
18 unique CDR3
7 unique J



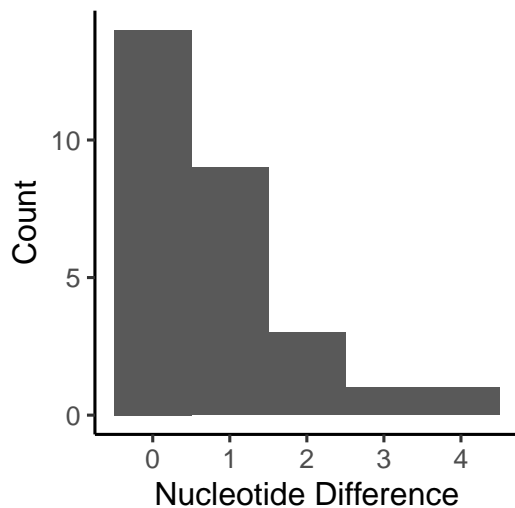
TRBV5–4*bp01

68 sequences assigned
58 (85.3%) exact matches, in which:
53 unique CDR3
11 unique J



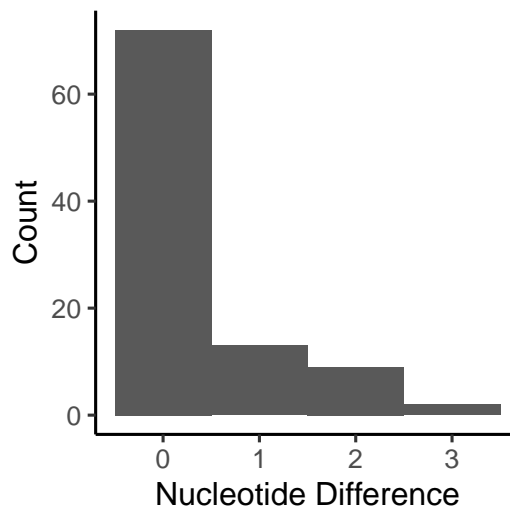
TRBV5-5*bp01

28 sequences assigned
14 (50%) exact matches, in which:
12 unique CDR3
9 unique J



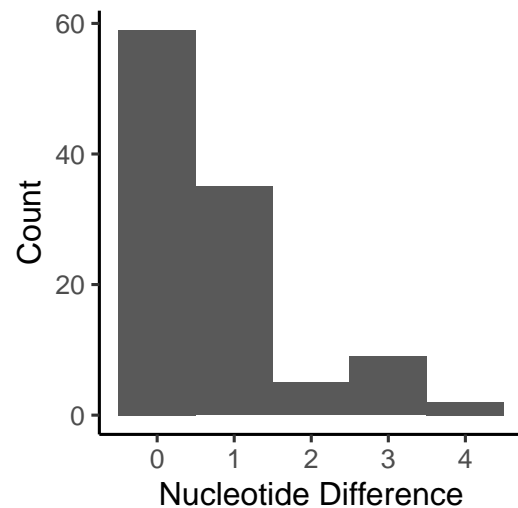
TRBV6-1*bp01

96 sequences assigned
72 (75%) exact matches, in which:
67 unique CDR3
12 unique J



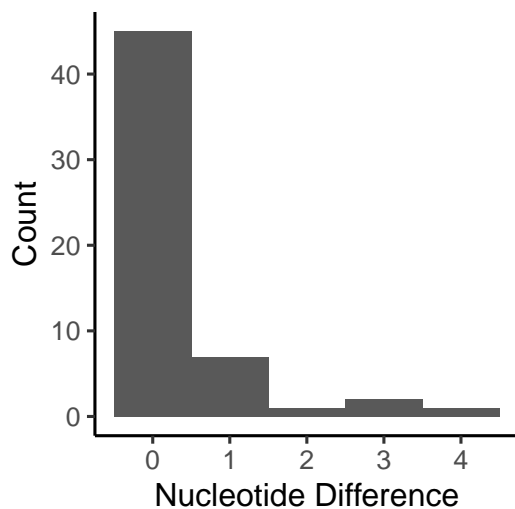
TRBV6-6*bp01

110 sequences assigned
59 (53.6%) exact matches, in which:
54 unique CDR3
12 unique J



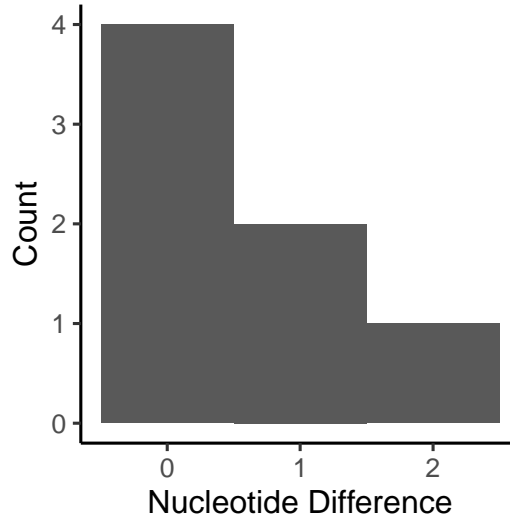
TRBV5-6*bp01

56 sequences assigned
45 (80.4%) exact matches, in which:
39 unique CDR3
12 unique J



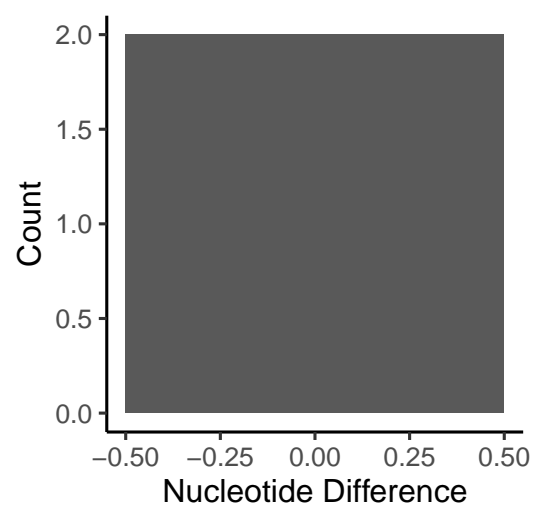
TRBV6-4*bp01

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



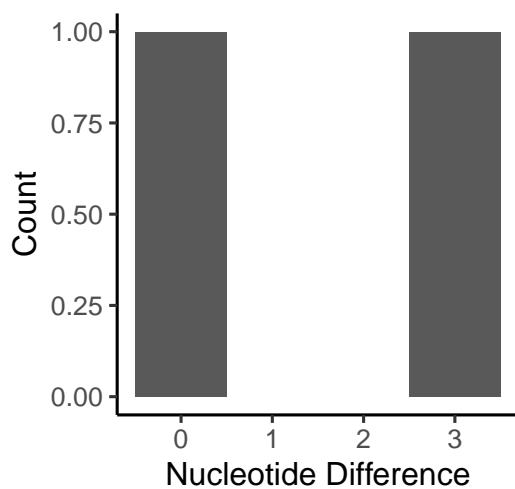
TRBV6-8*bp01

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



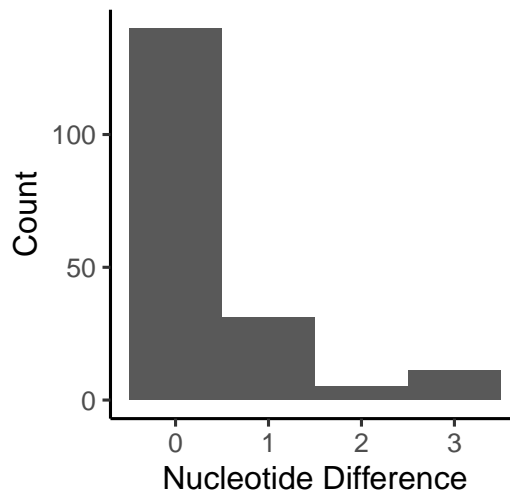
TRBV5-8*bp01

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



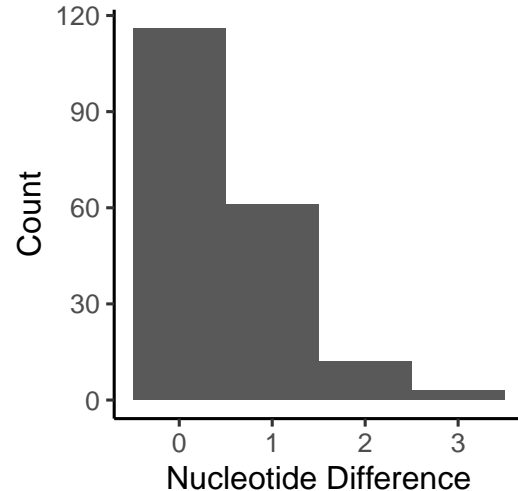
TRBV6-5*bp01

187 sequences assigned
140 (74.9%) exact matches, in which:
125 unique CDR3
13 unique J



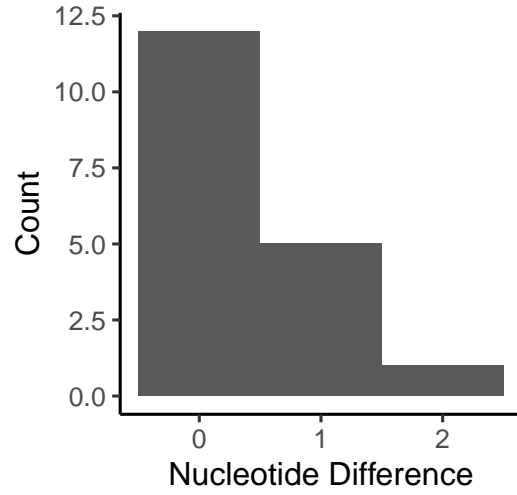
TRBV6-23*bp01

193 sequences assigned
116 (60.1%) exact matches, in which:
94 unique CDR3
12 unique J



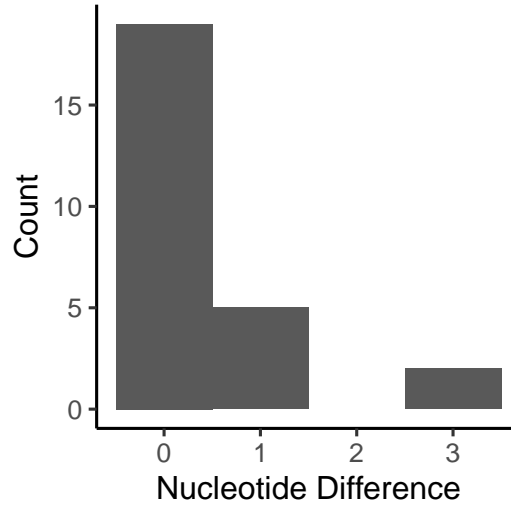
TRBV7-2*bp01

18 sequences assigned
12 (66.7%) exact matches, in which:
11 unique CDR3
9 unique J



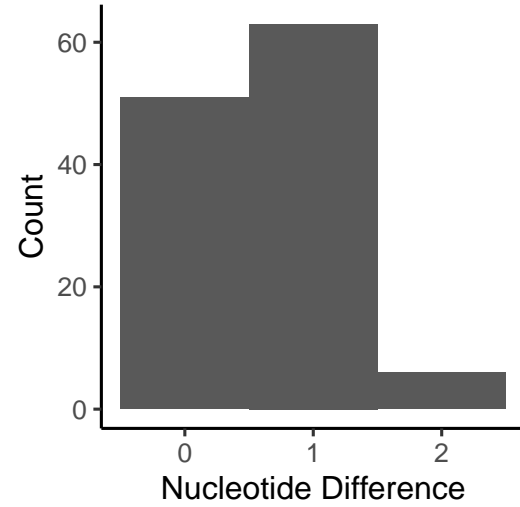
TRBV7-6*bp01

26 sequences assigned
19 (73.1%) exact matches, in which:
16 unique CDR3
9 unique J



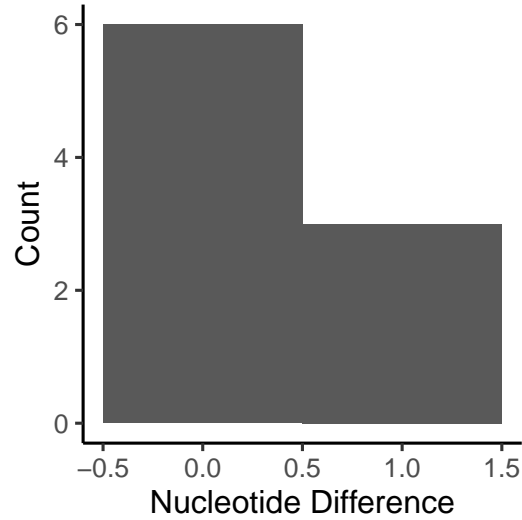
TRBV29-1*bp03

120 sequences assigned
51 (42.5%) exact matches, in which:
31 unique CDR3
8 unique J



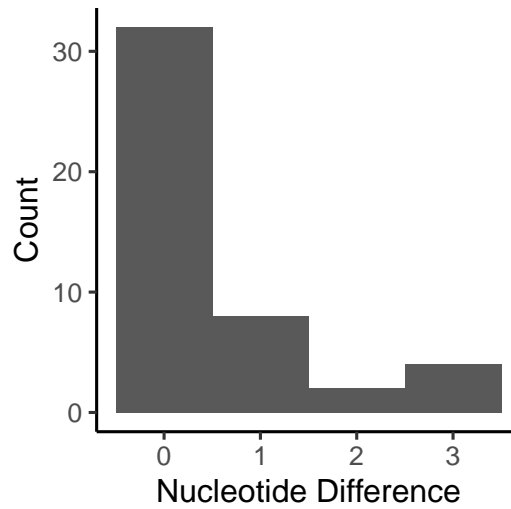
TRBV7-2*bp02

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



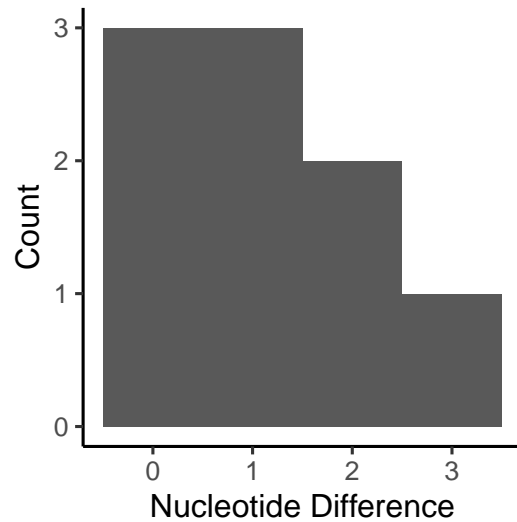
TRBV7-9*bp01

46 sequences assigned
32 (69.6%) exact matches, in which:
28 unique CDR3
9 unique J



TRBV7-4*bp01

9 sequences assigned
3 (33.3%) exact matches, in which:
2 unique CDR3
2 unique J



TRBV29-1*bp01

207 sequences assigned
140 (67.6%) exact matches, in which:
106 unique CDR3
13 unique J

