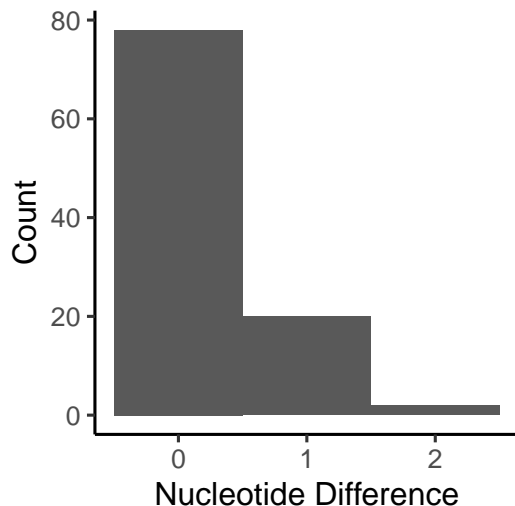


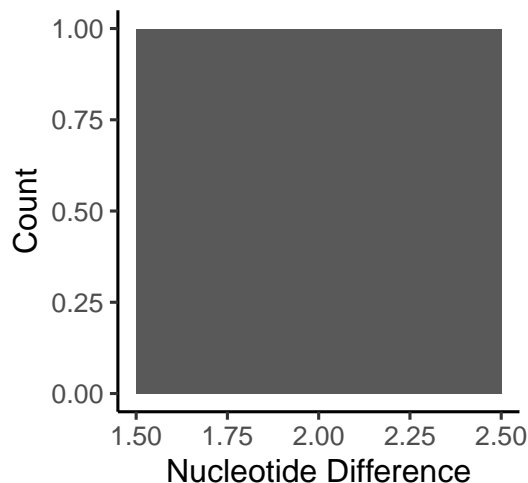
TRBV20–1*bp01

100 sequences assigned
78 (78%) exact matches, in which:
72 unique CDR3
12 unique J



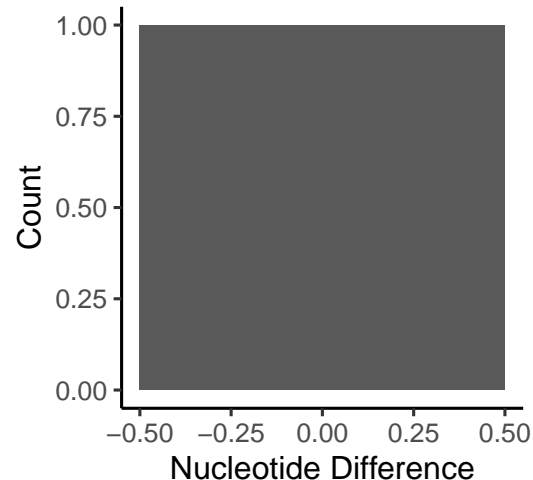
TRBV10–2*bp02

1 sequences assigned
No exact matches.



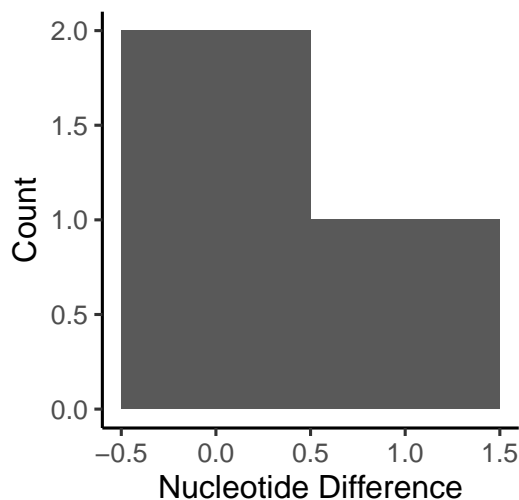
TRBV14*bp01

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



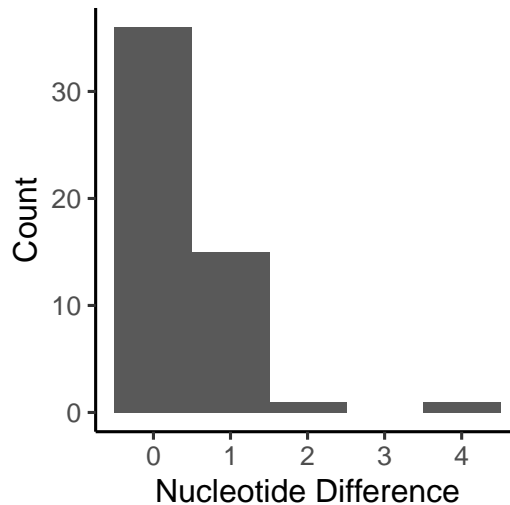
TRBV10–1*bp01

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



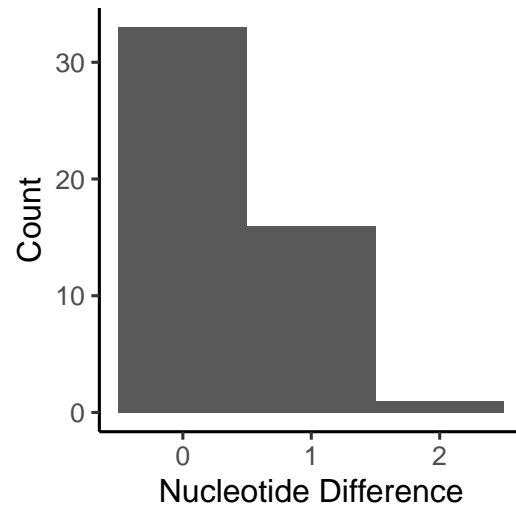
TRBV10–3*bp01

53 sequences assigned
36 (67.9%) exact matches, in which:
31 unique CDR3
10 unique J



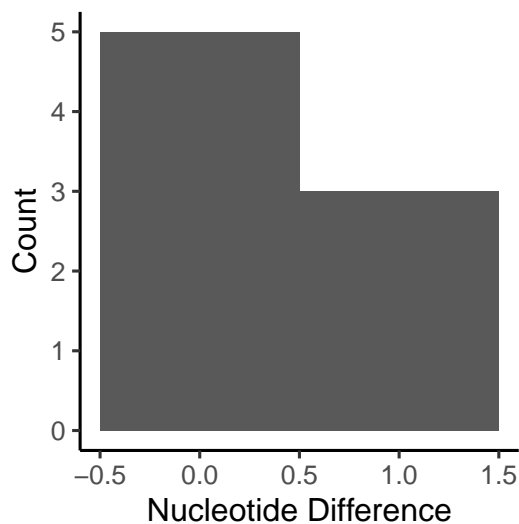
TRBV15*bp02

50 sequences assigned
33 (66%) exact matches, in which:
28 unique CDR3
10 unique J



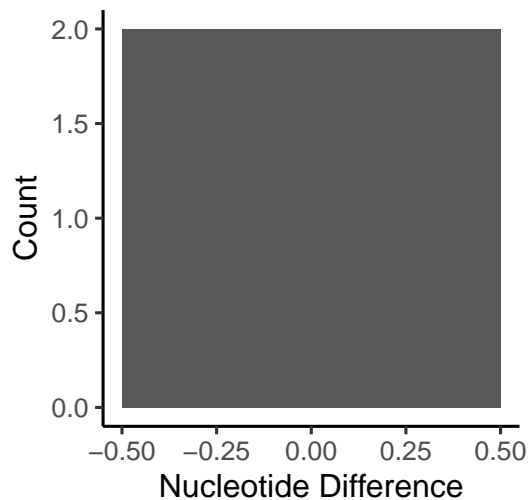
TRBV10–1*bp02

8 sequences assigned
5 (62.5%) exact matches, in which:
5 unique CDR3
4 unique J



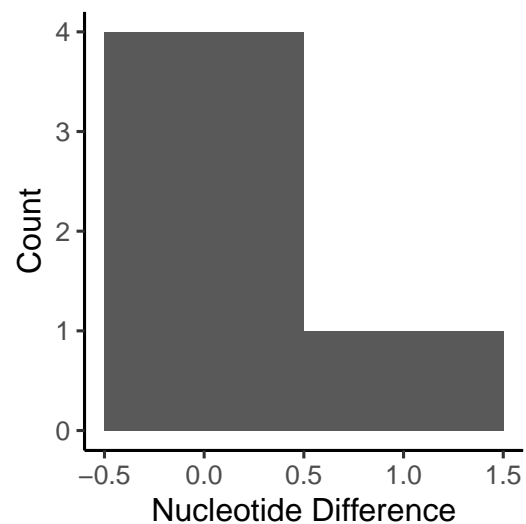
TRBV13*bp01

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



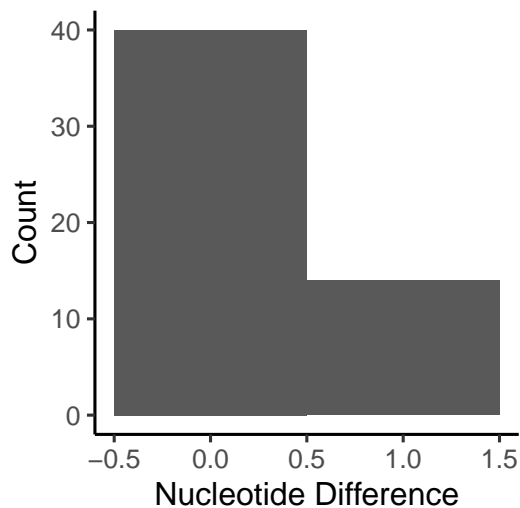
TRBV18*bp01

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



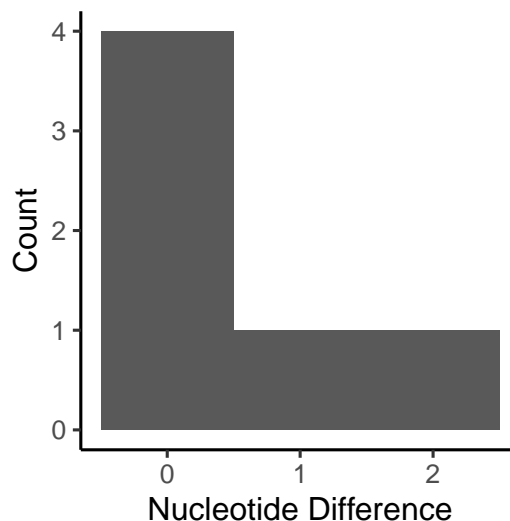
TRBV19*bp01

54 sequences assigned
40 (74.1%) exact matches, in which:
36 unique CDR3
11 unique J



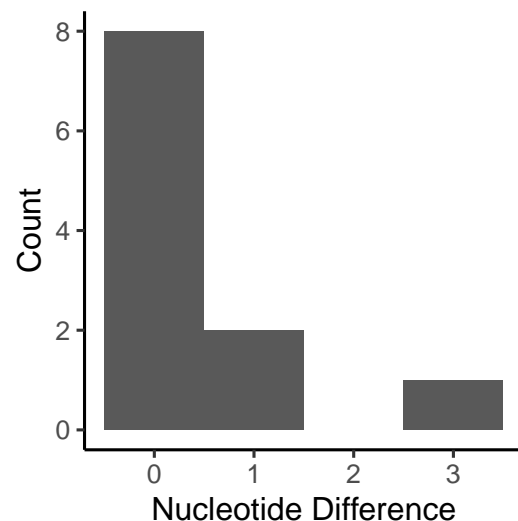
TRBV28*bp01

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



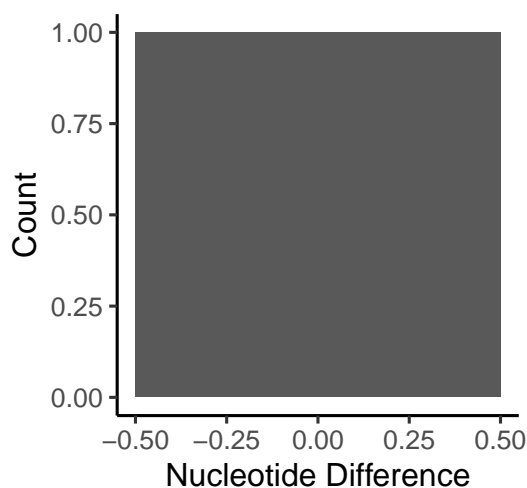
TRBV11-1*bp01

11 sequences assigned
8 (72.7%) exact matches, in which:
6 unique CDR3
3 unique J



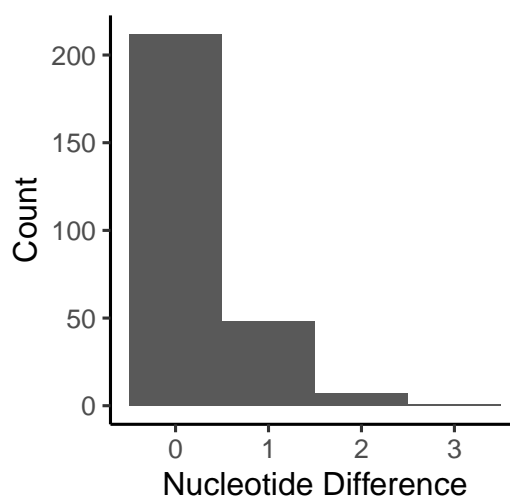
TRBV2*bp01

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



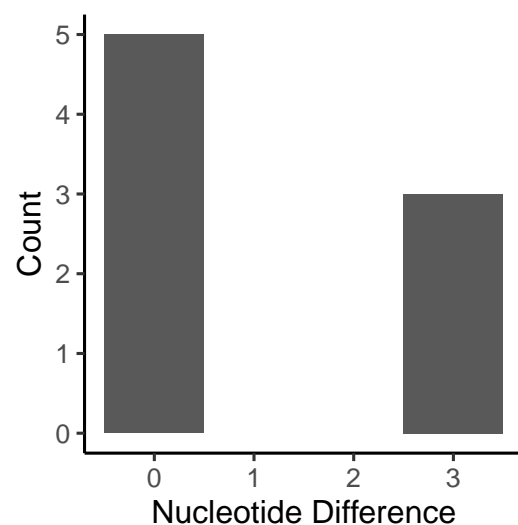
TRBV30*bp01

268 sequences assigned
212 (79.1%) exact matches, in which:
124 unique CDR3
13 unique J



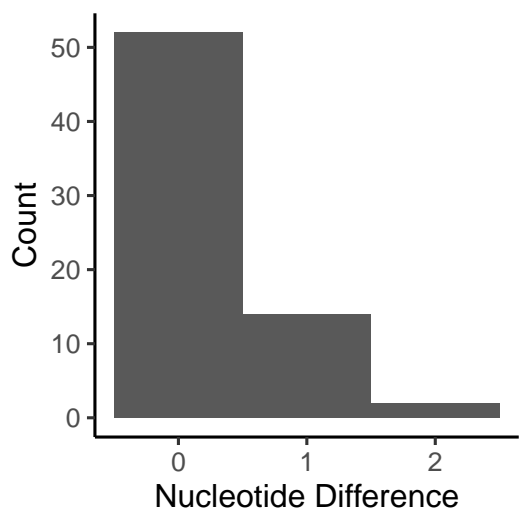
TRBV11-2*bp01

8 sequences assigned
5 (62.5%) exact matches, in which:
5 unique CDR3
5 unique J



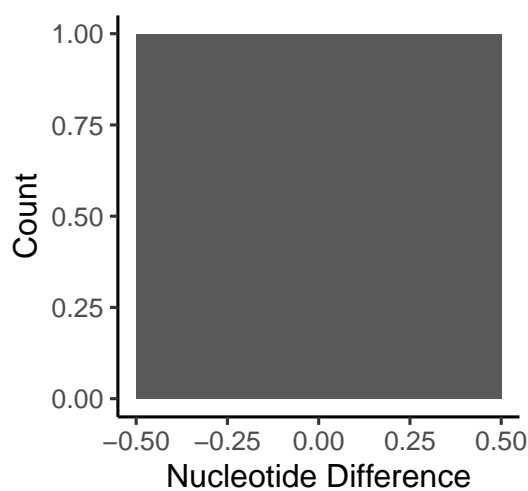
TRBV27*bp01

68 sequences assigned
52 (76.5%) exact matches, in which:
46 unique CDR3
9 unique J



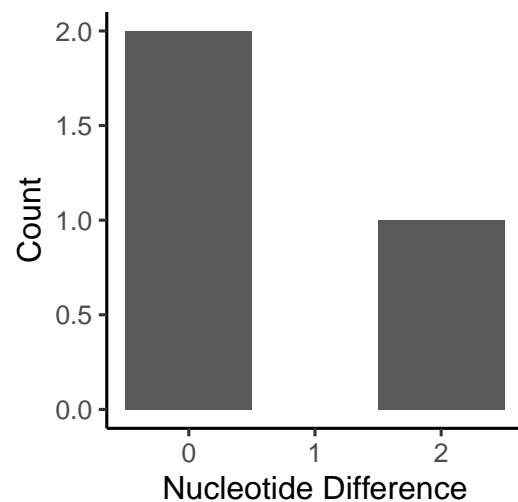
TRBV9*bp01

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



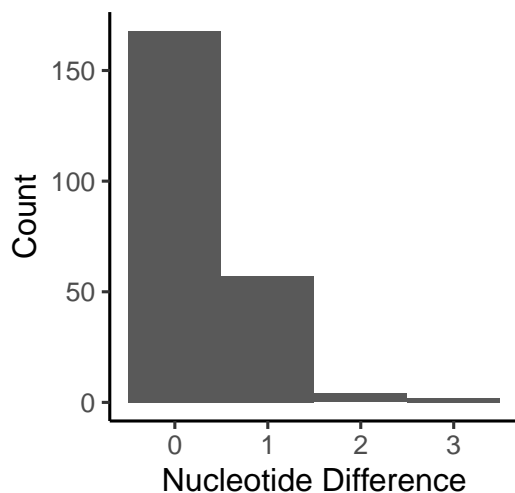
TRBV12-5*bp01

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



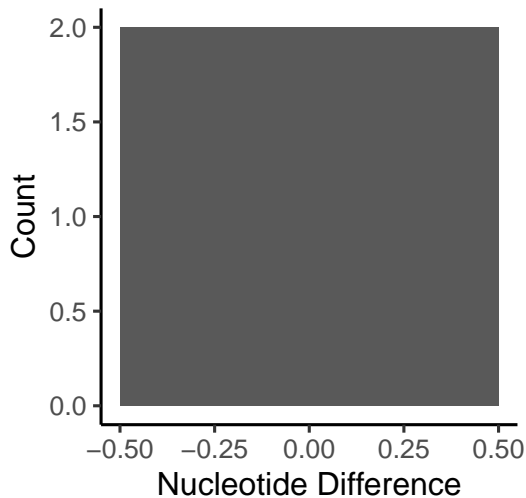
TRBV12–34*bp01

231 sequences assigned
168 (72.7%) exact matches, in which:
129 unique CDR3
14 unique J



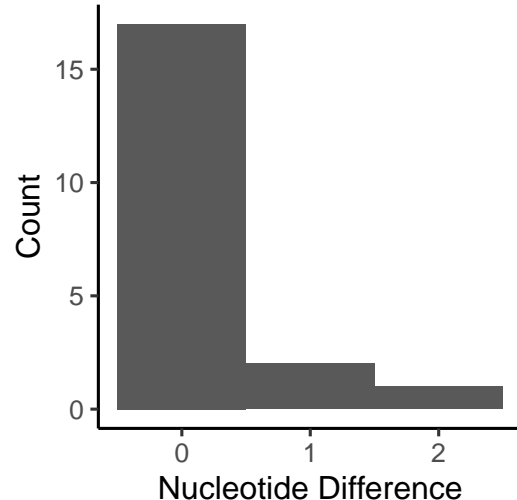
TRBV25–1*bp01

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



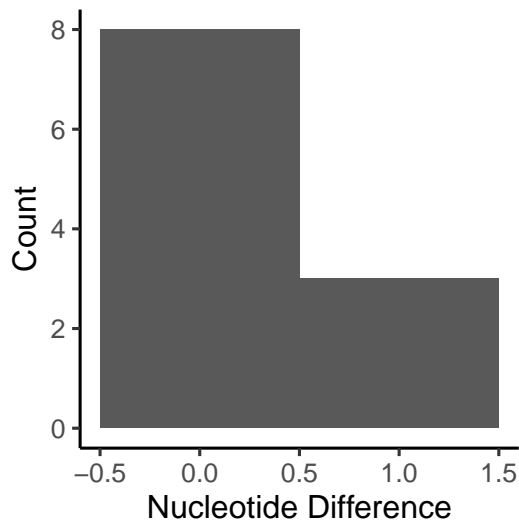
TRBV5–4*bp01

20 sequences assigned
17 (85%) exact matches, in which:
17 unique CDR3
8 unique J



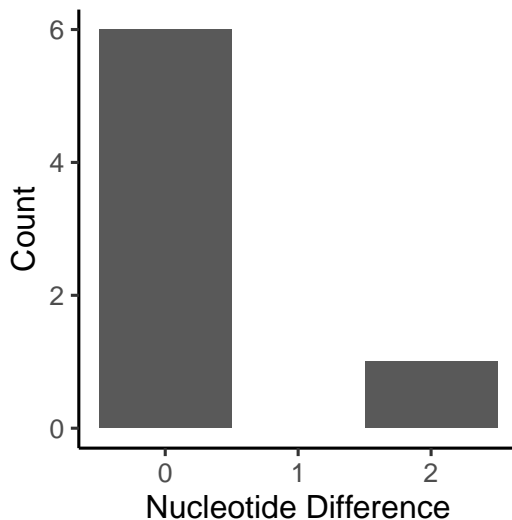
TRBV24–1*bp01

11 sequences assigned
8 (72.7%) exact matches, in which:
8 unique CDR3
5 unique J



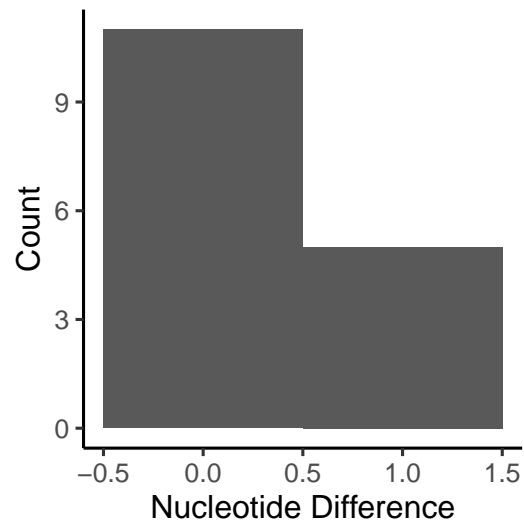
TRBV5–1*bp01

7 sequences assigned
6 (85.7%) exact matches, in which:
5 unique CDR3
4 unique J



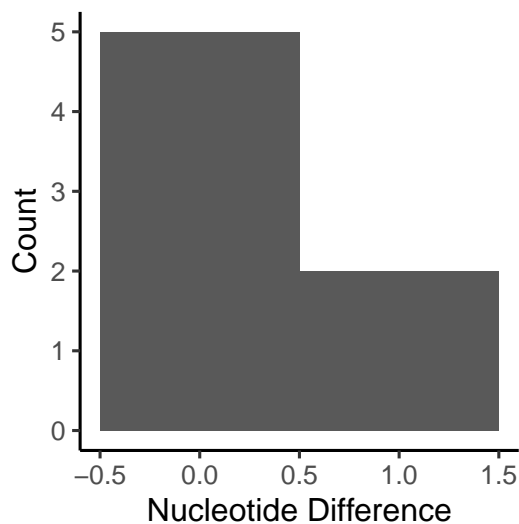
TRBV5–5*bp01

16 sequences assigned
11 (68.8%) exact matches, in which:
11 unique CDR3
7 unique J



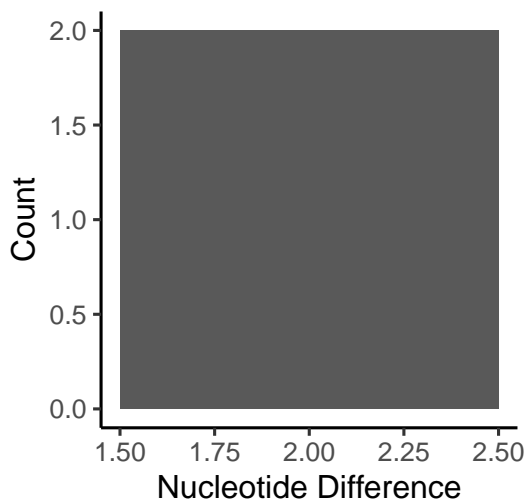
TRBV4–3*bp01

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



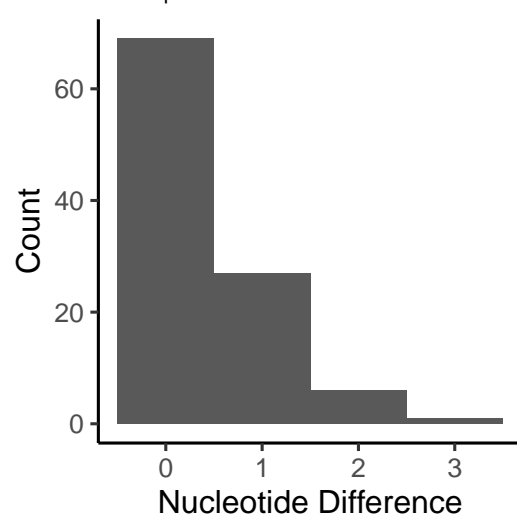
TRBV5–3*bp01

2 sequences assigned
No exact matches.



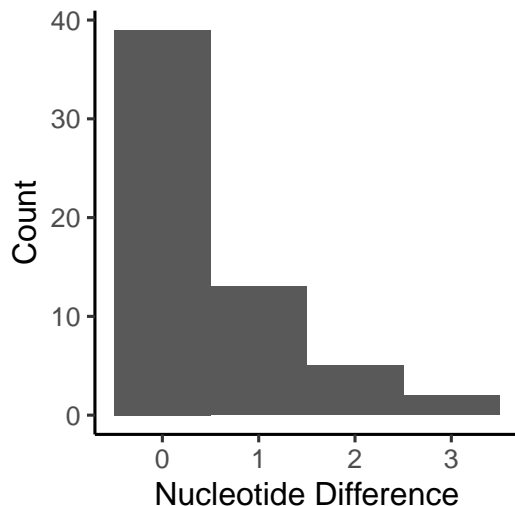
TRBV5–6*bp01

103 sequences assigned
69 (67%) exact matches, in which:
57 unique CDR3
10 unique J



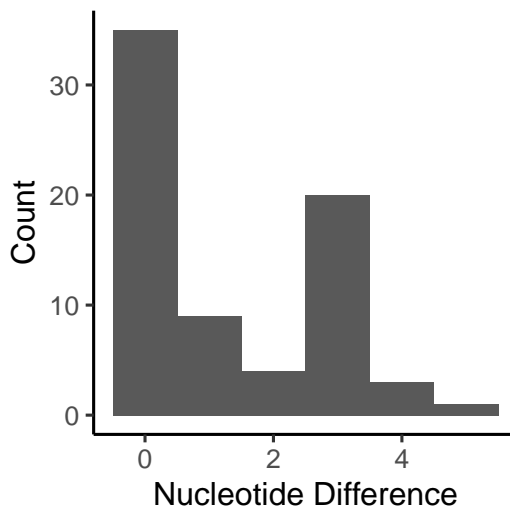
TRBV6-1*bp01

59 sequences assigned
39 (66.1%) exact matches, in which:
35 unique CDR3
10 unique J



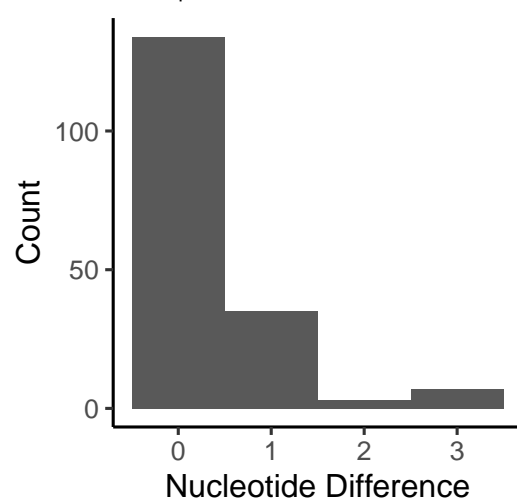
TRBV6-6*bp01

72 sequences assigned
35 (48.6%) exact matches, in which:
33 unique CDR3
9 unique J



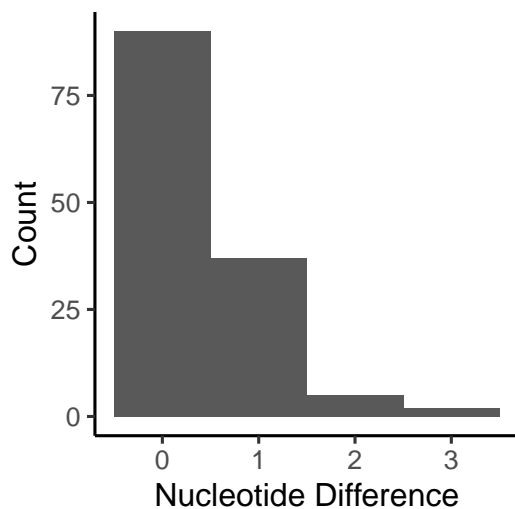
TRBV7-2*bp01

179 sequences assigned
134 (74.9%) exact matches, in which:
107 unique CDR3
14 unique J



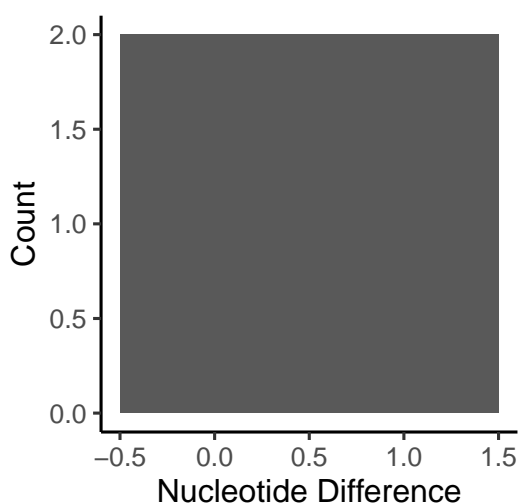
TRBV6-4*bp01

134 sequences assigned
90 (67.2%) exact matches, in which:
64 unique CDR3
11 unique J



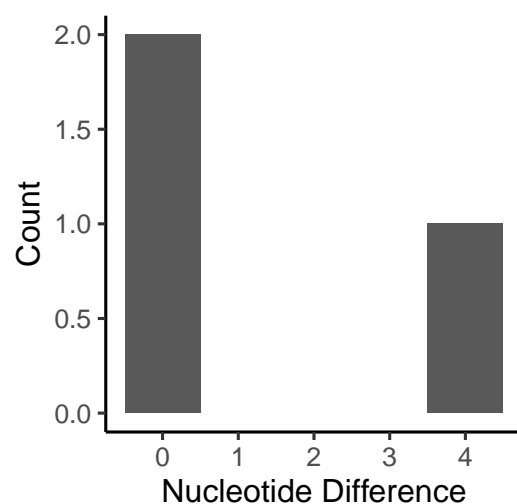
TRBV6-9*bp01

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



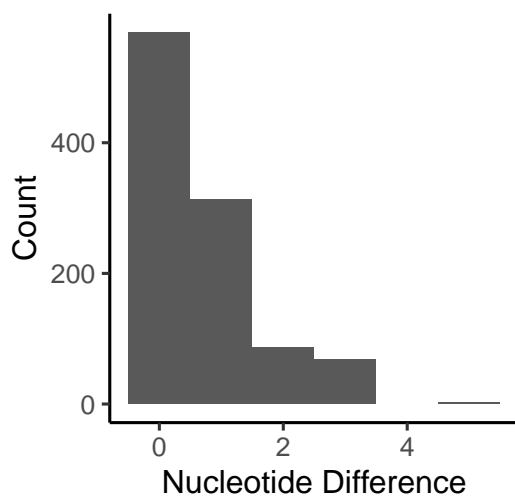
TRBV7-3*bp02

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



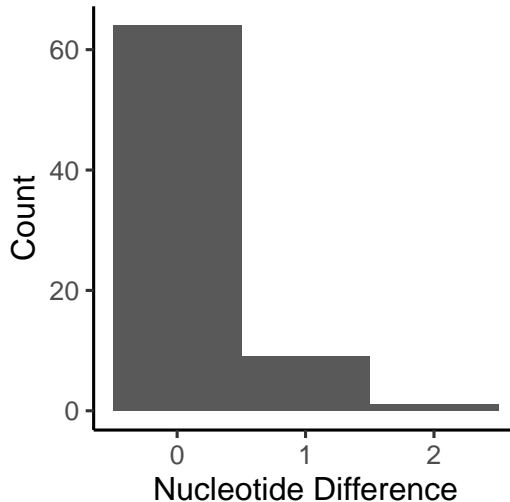
TRBV6-5*bp01

1039 sequences assigned
569 (54.8%) exact matches, in which:
429 unique CDR3
14 unique J



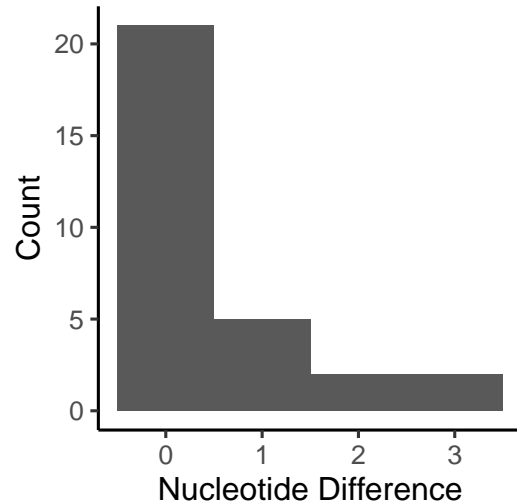
TRBV6-23*bp01

74 sequences assigned
64 (86.5%) exact matches, in which:
62 unique CDR3
10 unique J



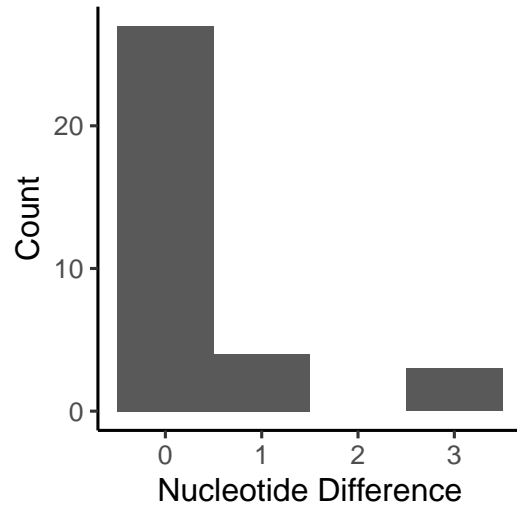
TRBV7-4*bp01

30 sequences assigned
21 (70%) exact matches, in which:
17 unique CDR3
9 unique J



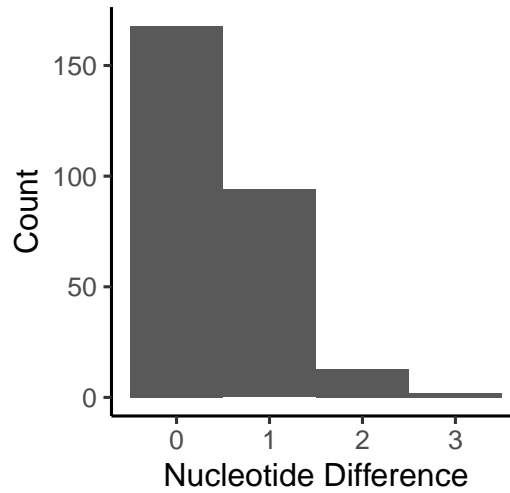
TRBV7-6*bp01

34 sequences assigned
27 (79.4%) exact matches, in which:
24 unique CDR3
7 unique J



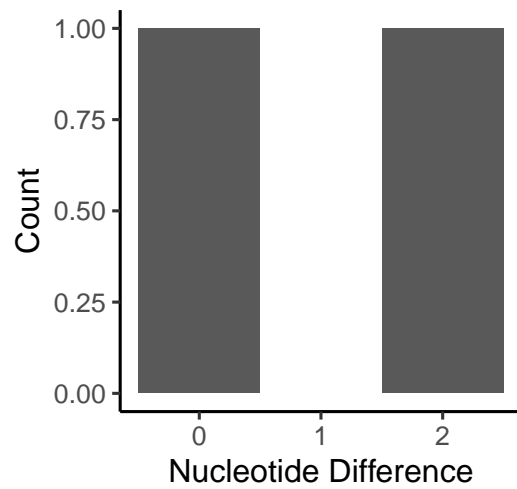
TRBV29-1*bp01

277 sequences assigned
168 (60.6%) exact matches, in which:
115 unique CDR3
13 unique J



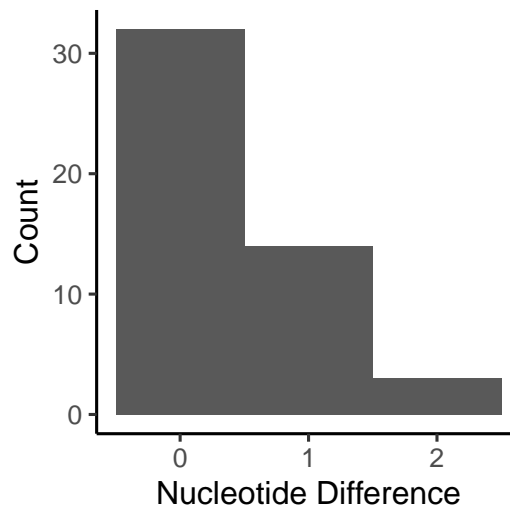
TRBV7-7*bp01

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



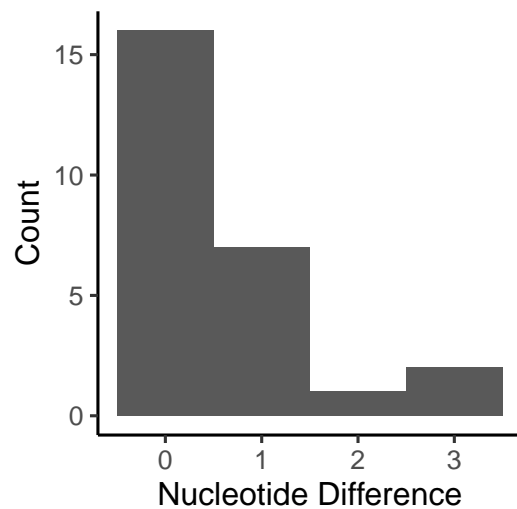
TRBV29-1*bp03

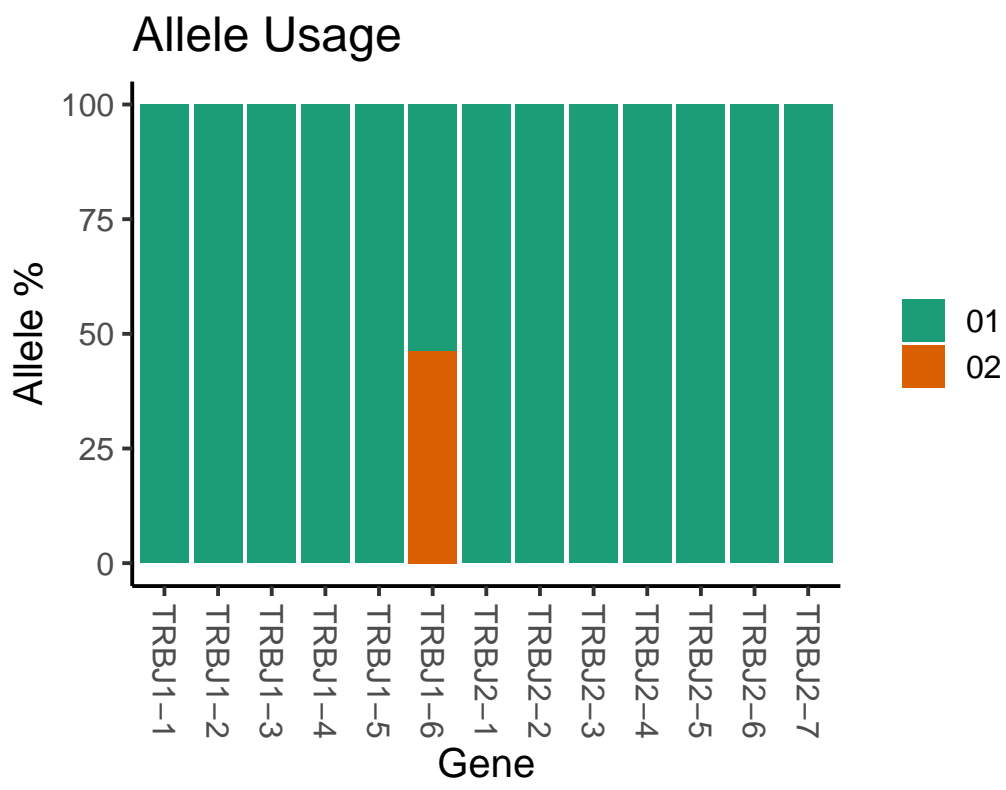
49 sequences assigned
32 (65.3%) exact matches, in which:
19 unique CDR3
11 unique J



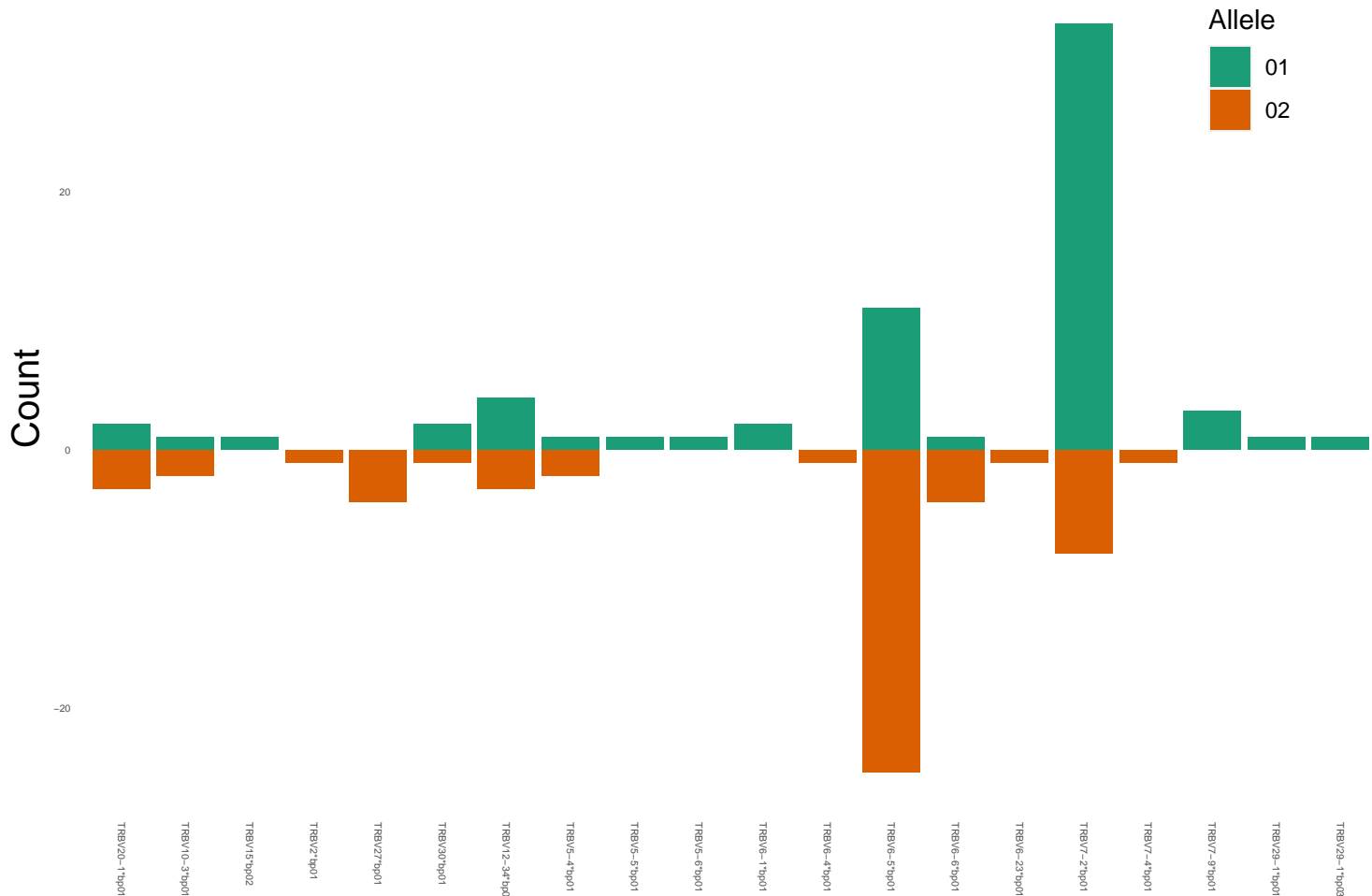
TRBV7-9*bp01

26 sequences assigned
16 (61.5%) exact matches, in which:
16 unique CDR3
7 unique J





Sequence Count by TRBJ1–6 allele usage



[illegible]