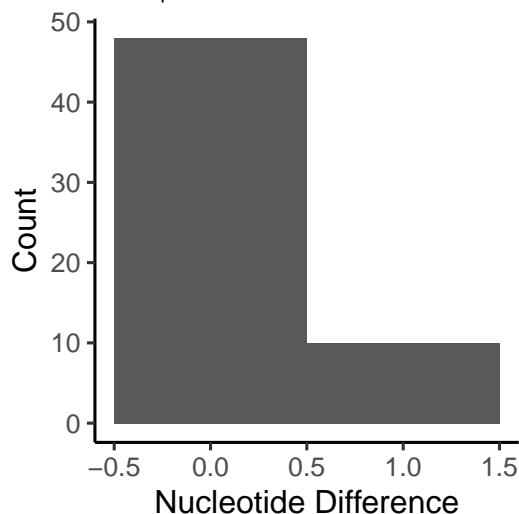


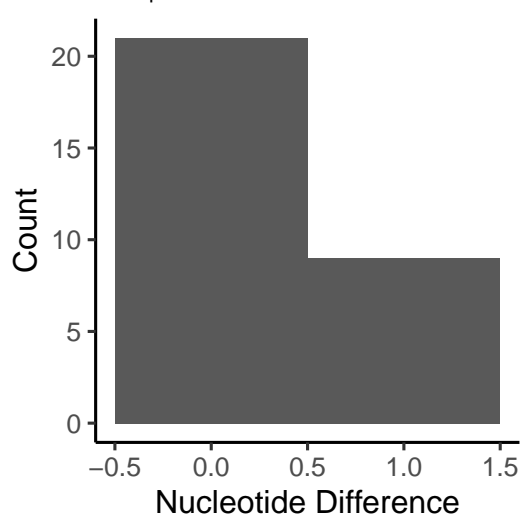
TRBV20–1*bp01

58 sequences assigned
48 (82.8%) exact matches, in which:
29 unique CDR3
10 unique J



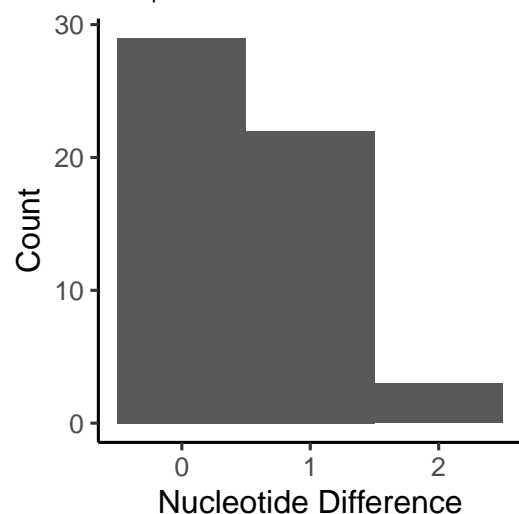
TRBV10–3*bp01

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



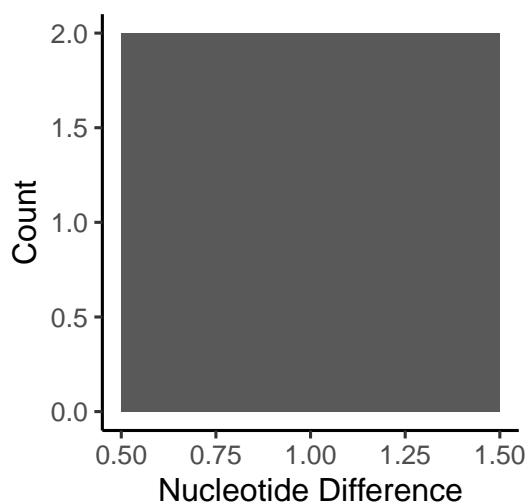
TRBV15*bp02

54 sequences assigned
29 (53.7%) exact matches, in which:
19 unique CDR3
8 unique J



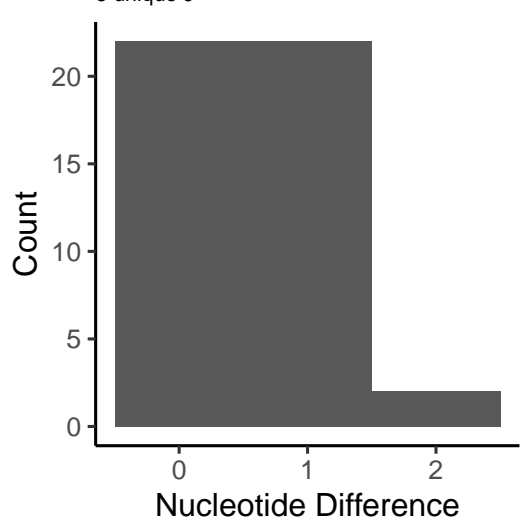
TRBV10–1*bp02

2 sequences assigned
No exact matches.



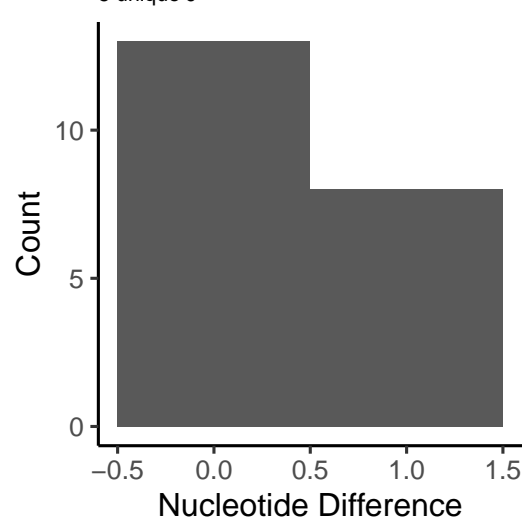
TRBV13*bp01

46 sequences assigned
22 (47.8%) exact matches, in which:
15 unique CDR3
6 unique J



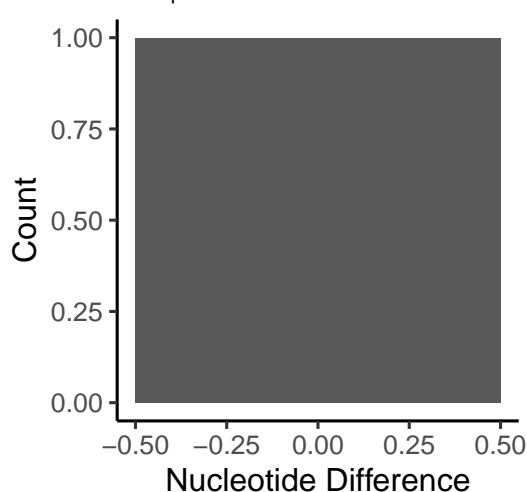
TRBV18*bp01

21 sequences assigned
13 (61.9%) exact matches, in which:
9 unique CDR3
5 unique J



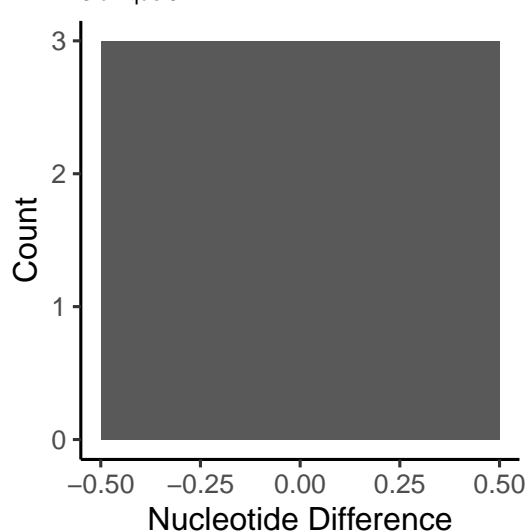
TRBV10–2*bp01

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



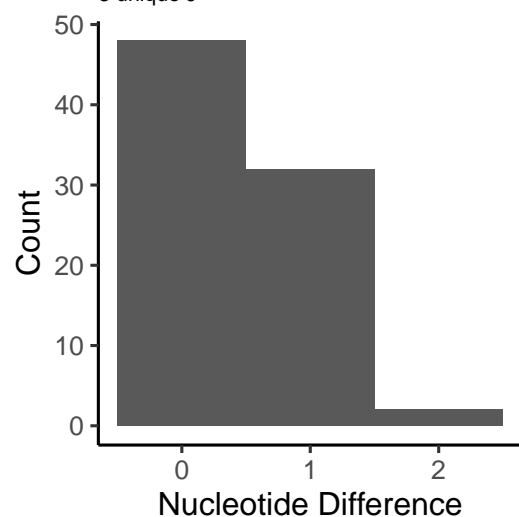
TRBV14*bp01

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



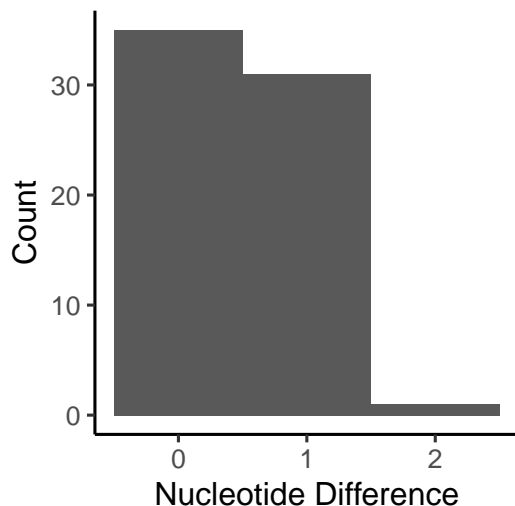
TRBV19*bp01

82 sequences assigned
48 (58.5%) exact matches, in which:
30 unique CDR3
8 unique J



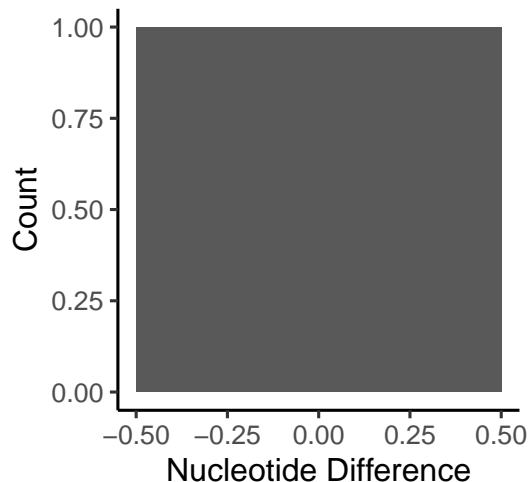
TRBV27*bp01

67 sequences assigned
35 (52.2%) exact matches, in which:
25 unique CDR3
9 unique J



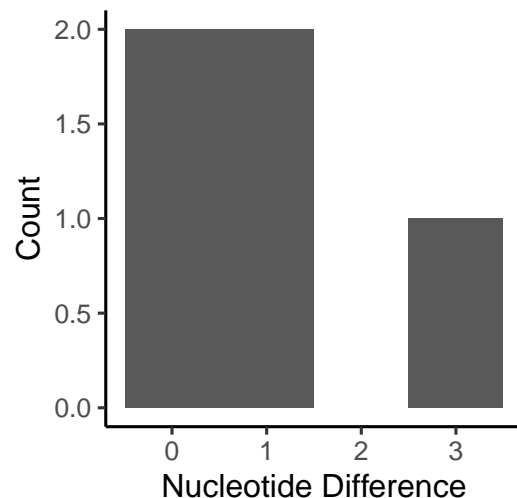
TRBV9*bp01

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



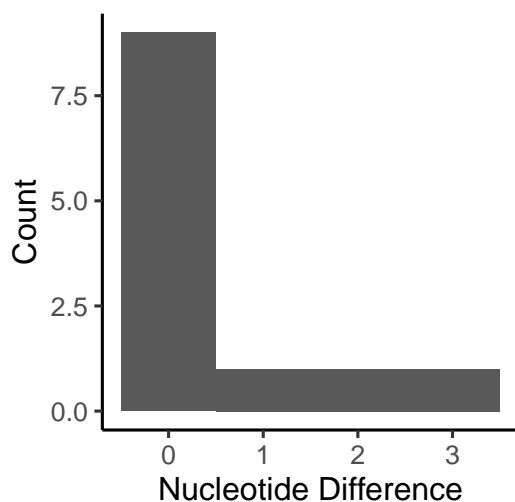
TRBV11-2*bp01

5 sequences assigned
2 (40%) exact matches, in which:
2 unique CDR3
1 unique J



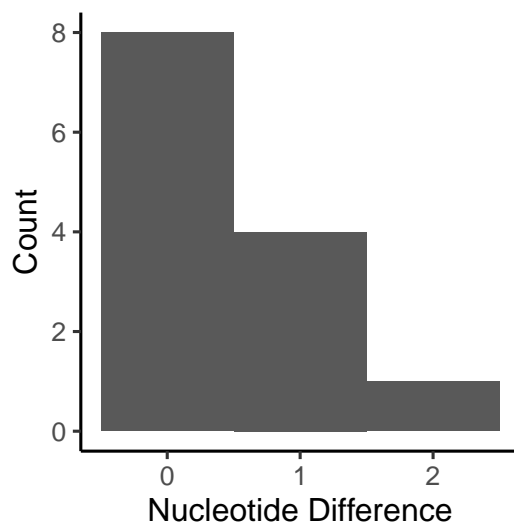
TRBV28*bp01

12 sequences assigned
9 (75%) exact matches, in which:
9 unique CDR3
5 unique J



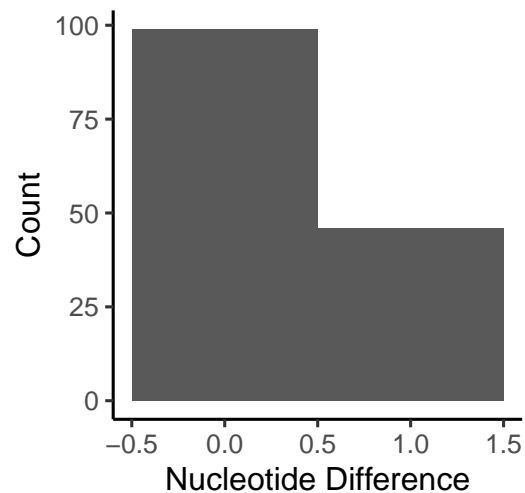
TRBV11-1*bp01

13 sequences assigned
8 (61.5%) exact matches, in which:
4 unique CDR3
3 unique J



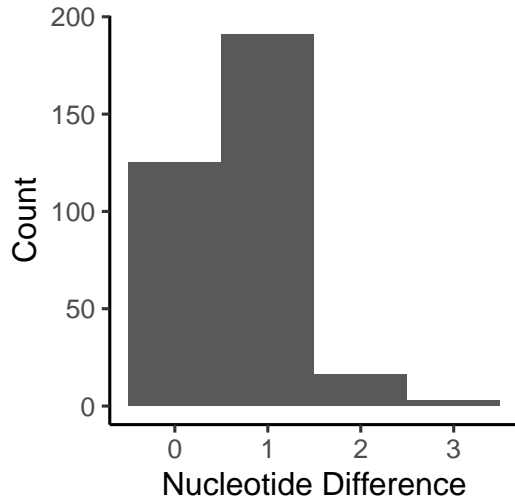
TRBV12-34*bp01

145 sequences assigned
99 (68.3%) exact matches, in which:
44 unique CDR3
9 unique J



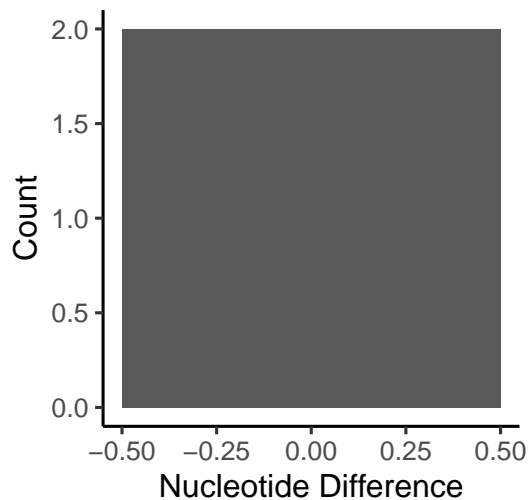
TRBV30*bp01

335 sequences assigned
125 (37.3%) exact matches, in which:
70 unique CDR3
7 unique J



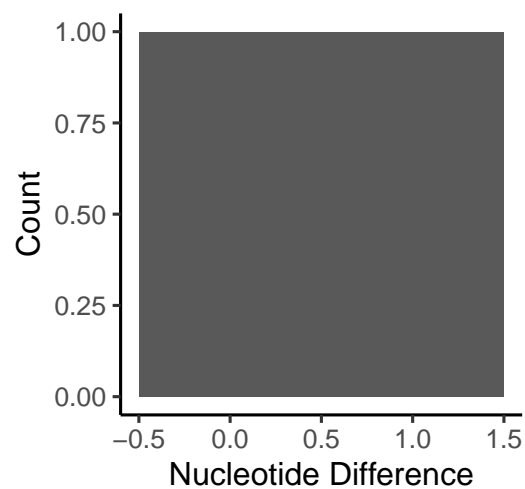
TRBV21-1*bp01

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



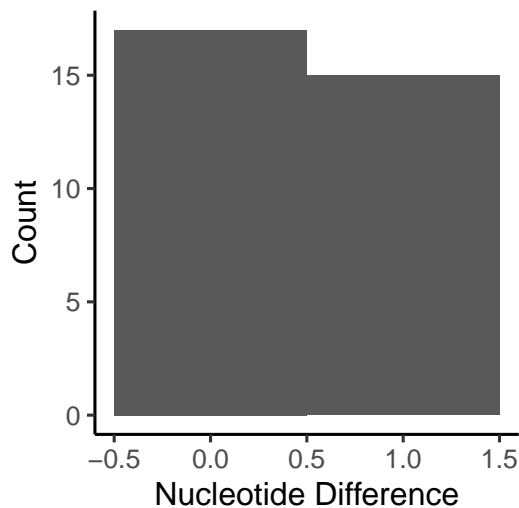
TRBV23-1*bp01

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



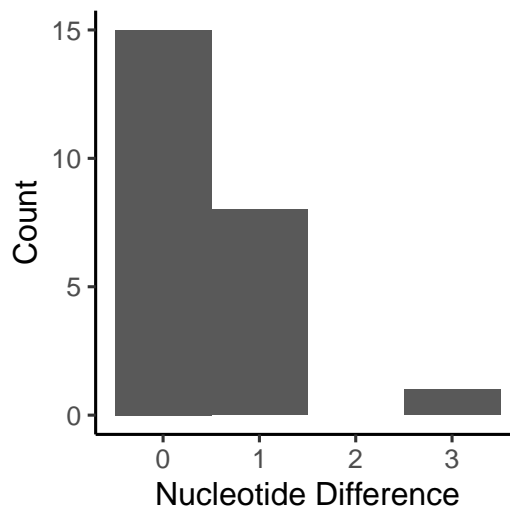
TRBV24-1*bp01

32 sequences assigned
17 (53.1%) exact matches, in which:
12 unique CDR3
5 unique J



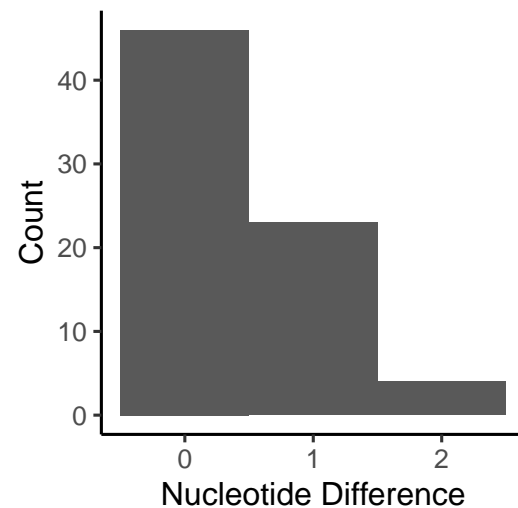
TRBV4-3*bp01

24 sequences assigned
15 (62.5%) exact matches, in which:
13 unique CDR3
5 unique J



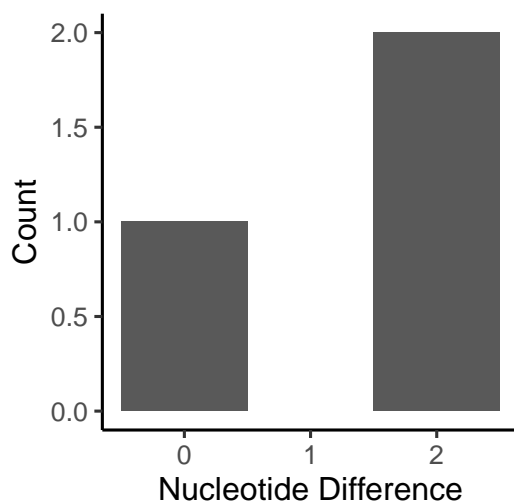
TRBV5-4*bp01

73 sequences assigned
46 (63%) exact matches, in which:
30 unique CDR3
10 unique J



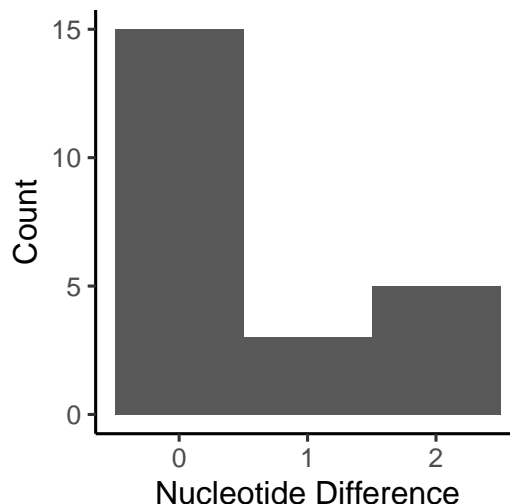
TRBV4-1*bp01

3 sequences assigned
1 (33.3%) exact matches, in which:
1 unique CDR3
1 unique J



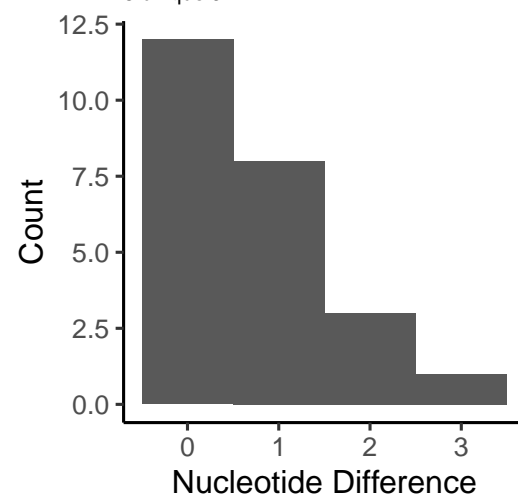
TRBV5-1*bp01

23 sequences assigned
15 (65.2%) exact matches, in which:
11 unique CDR3
6 unique J



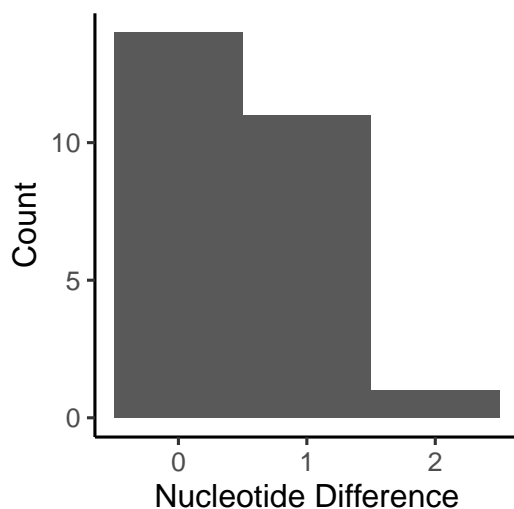
TRBV5-5*bp01

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



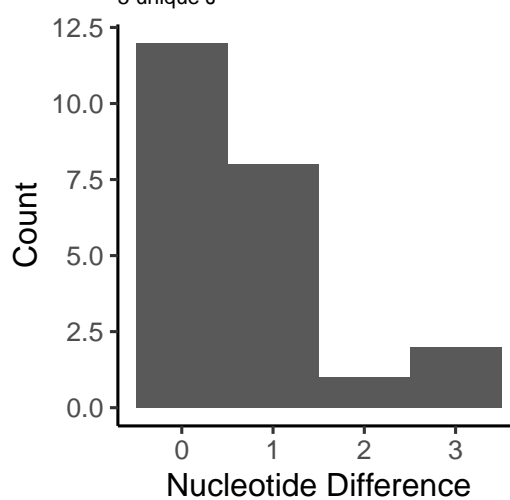
TRBV4-2*bp01

26 sequences assigned
14 (53.8%) exact matches, in which:
13 unique CDR3
6 unique J



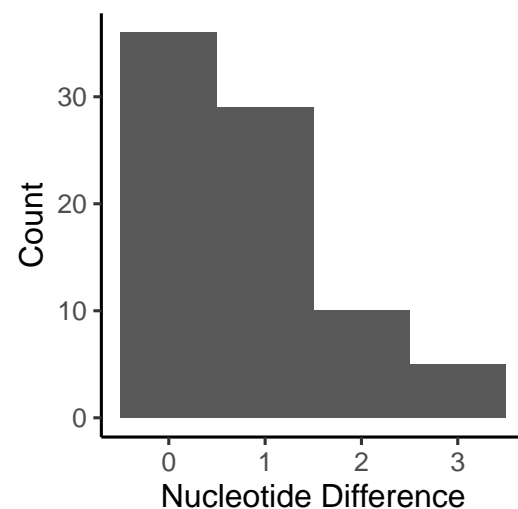
TRBV5-3*bp01

23 sequences assigned
12 (52.2%) exact matches, in which:
9 unique CDR3
5 unique J



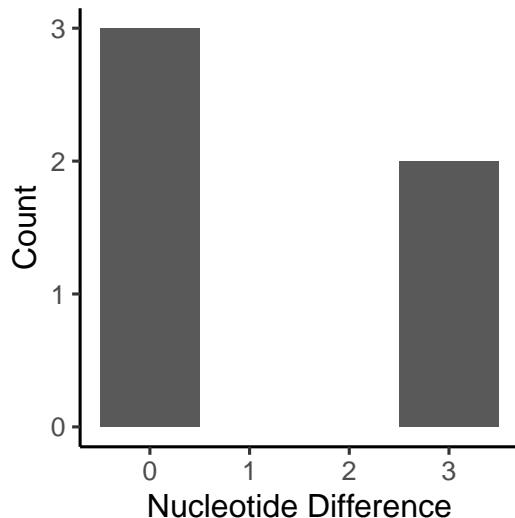
TRBV5-6*bp01

80 sequences assigned
36 (45%) exact matches, in which:
22 unique CDR3
7 unique J



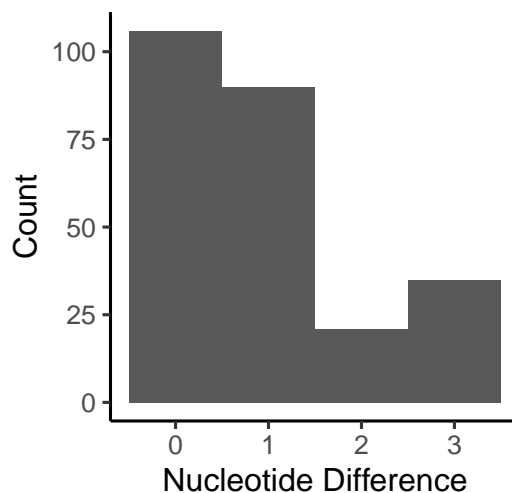
TRBV5–8*bp01

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



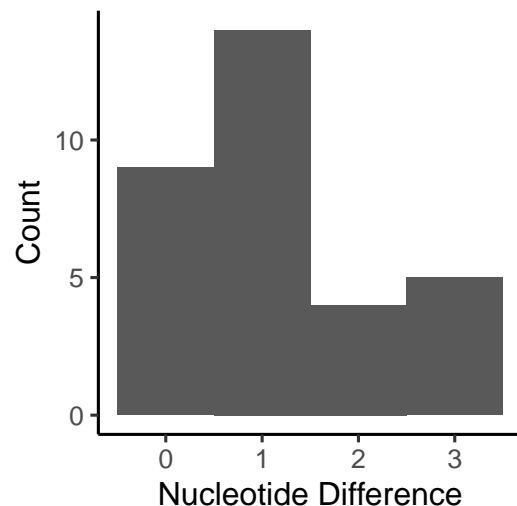
TRBV6–5*bp01

252 sequences assigned
106 (42.1%) exact matches, in which:
81 unique CDR3
13 unique J



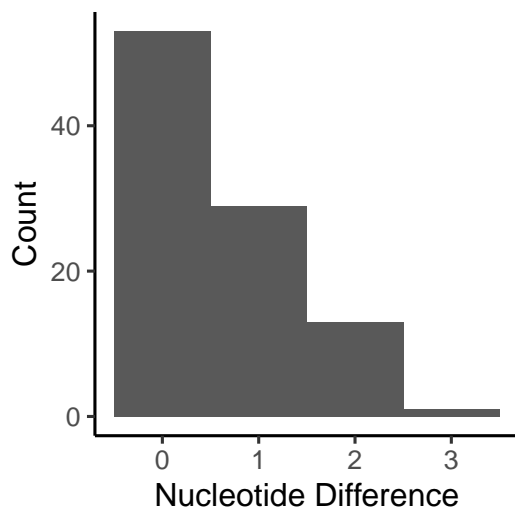
TRBV6–9*bp01

32 sequences assigned
9 (28.1%) exact matches, in which:
9 unique CDR3
6 unique J



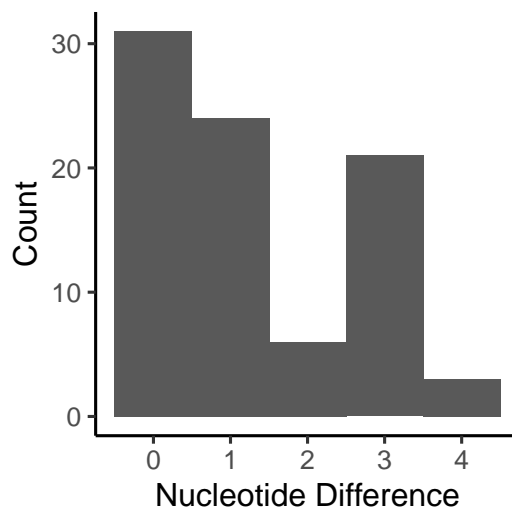
TRBV6–1*bp01

96 sequences assigned
53 (55.2%) exact matches, in which:
41 unique CDR3
13 unique J



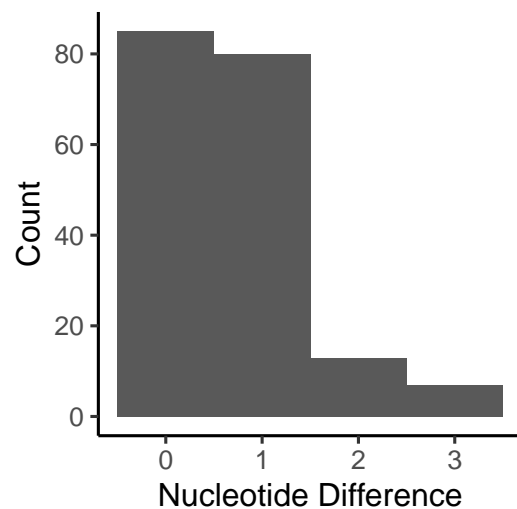
TRBV6–6*bp01

85 sequences assigned
31 (36.5%) exact matches, in which:
25 unique CDR3
8 unique J



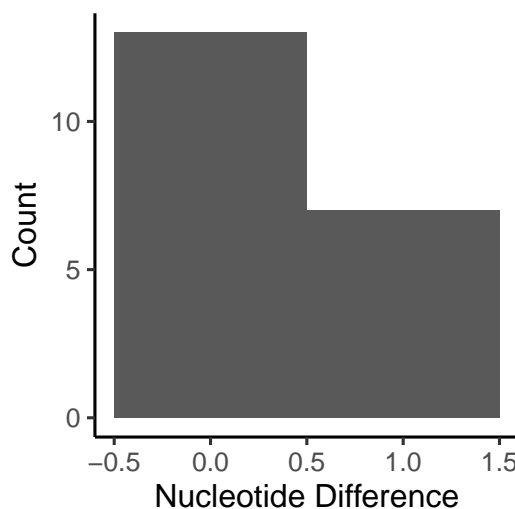
TRBV6–23*bp01

185 sequences assigned
85 (45.9%) exact matches, in which:
60 unique CDR3
12 unique J



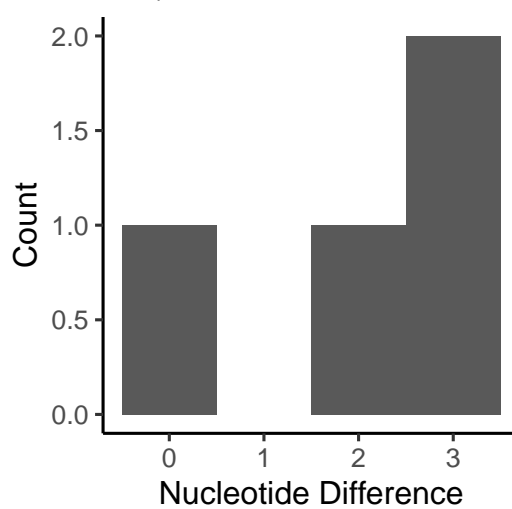
TRBV6–4*bp01

20 sequences assigned
13 (65%) exact matches, in which:
11 unique CDR3
5 unique J



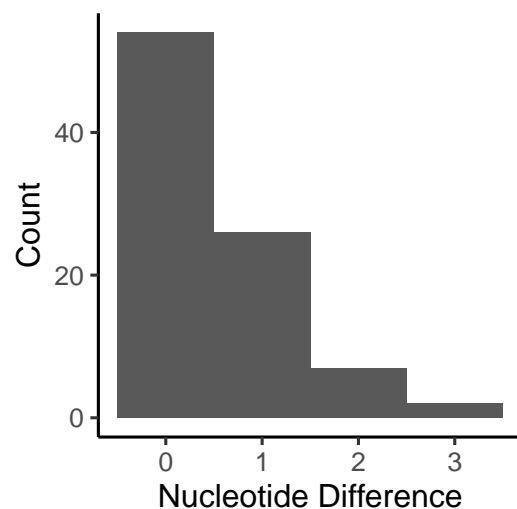
TRBV6–8*bp01

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



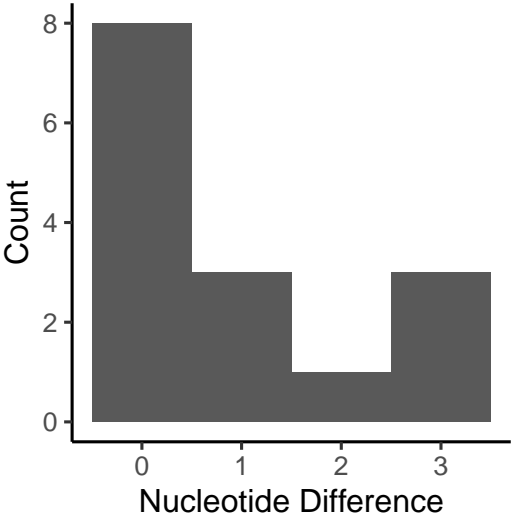
TRBV7–2*bp01

89 sequences assigned
54 (60.7%) exact matches, in which:
32 unique CDR3
11 unique J



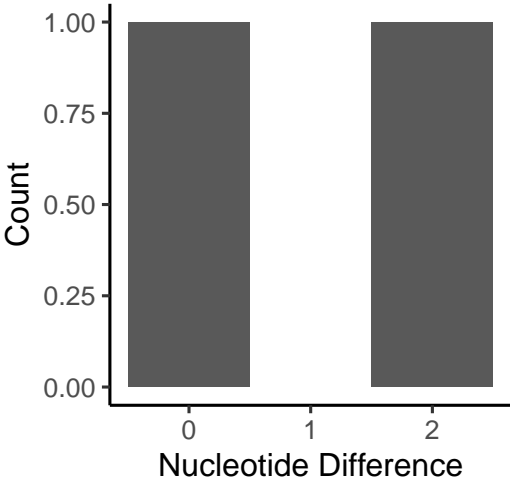
TRBV7-3*bp01

15 sequences assigned
8 (53.3%) exact matches, in which:
6 unique CDR3
4 unique J



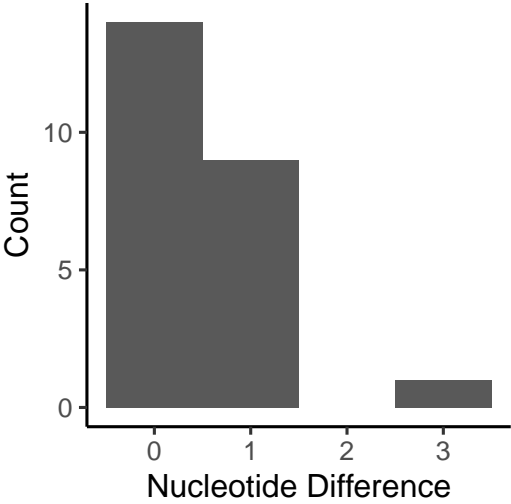
TRBV7-7*bp01

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



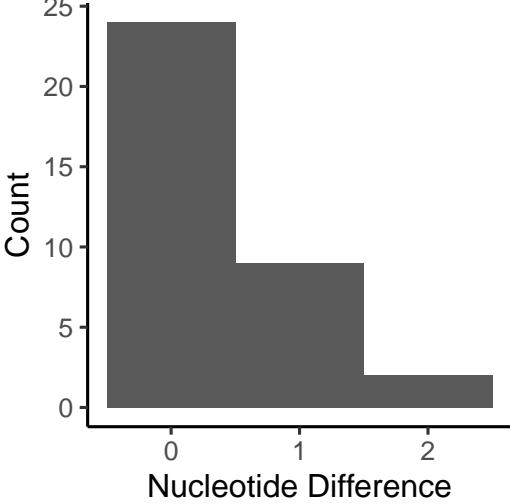
TRBV7-4*bp01

24 sequences assigned
14 (58.3%) exact matches, in which:
7 unique CDR3
3 unique J



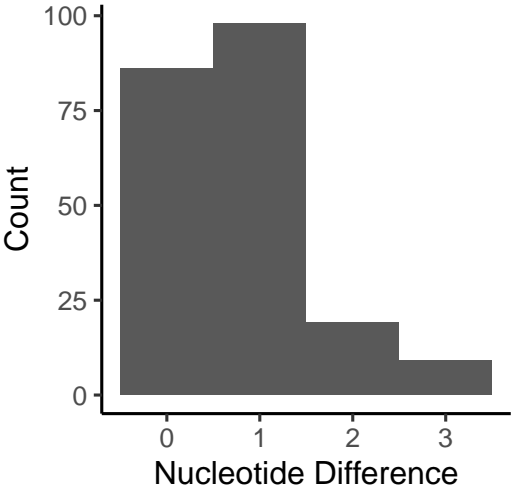
TRBV7-9*bp01

35 sequences assigned
24 (68.6%) exact matches, in which:
14 unique CDR3
8 unique J



TRBV7-6*bp01

212 sequences assigned
86 (40.6%) exact matches, in which:
46 unique CDR3
7 unique J



TRBV29-1*bp01

336 sequences assigned
108 (32.1%) exact matches, in which:
66 unique CDR3
8 unique J

