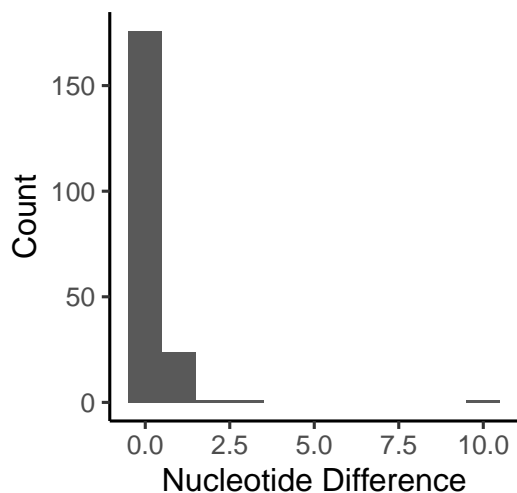


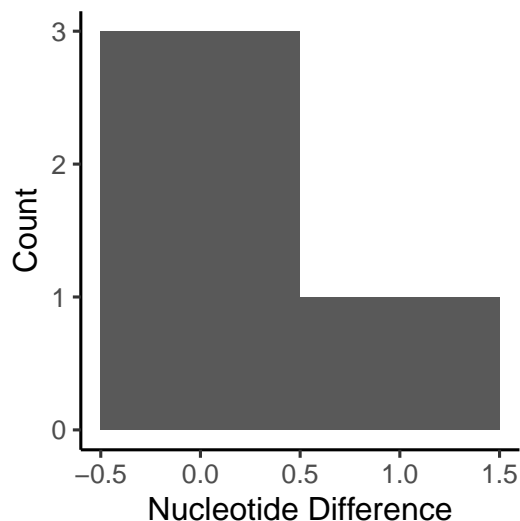
TRBV20–1*bp01

203 sequences assigned
176 (86.7%) exact matches, in which:
166 unique CDR3
13 unique J



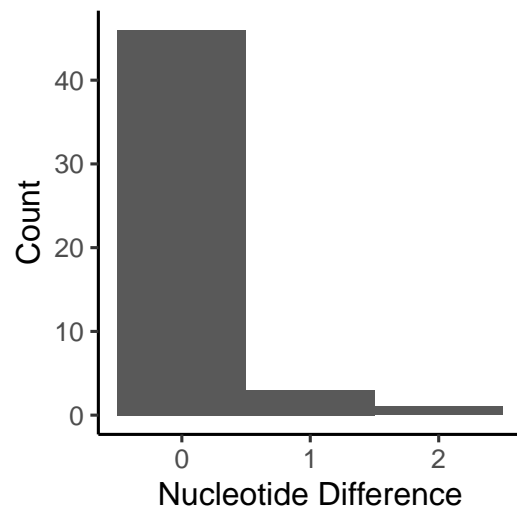
TRBV10–2*bp01

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



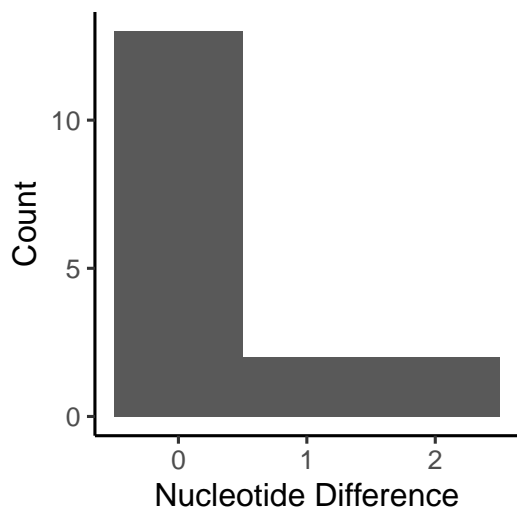
TRBV14*bp01

50 sequences assigned
46 (92%) exact matches, in which:
45 unique CDR3
10 unique J



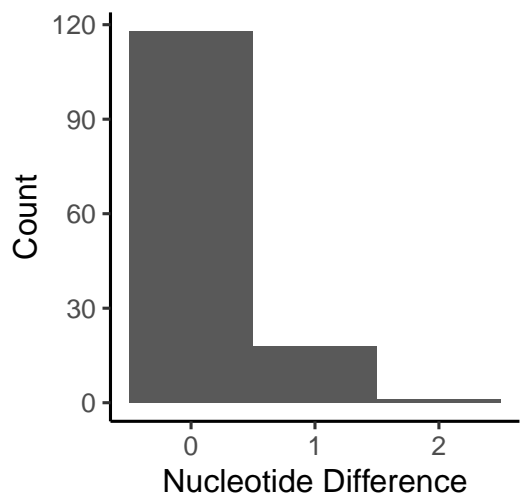
TRBV10–1*bp01

17 sequences assigned
13 (76.5%) exact matches, in which:
12 unique CDR3
5 unique J



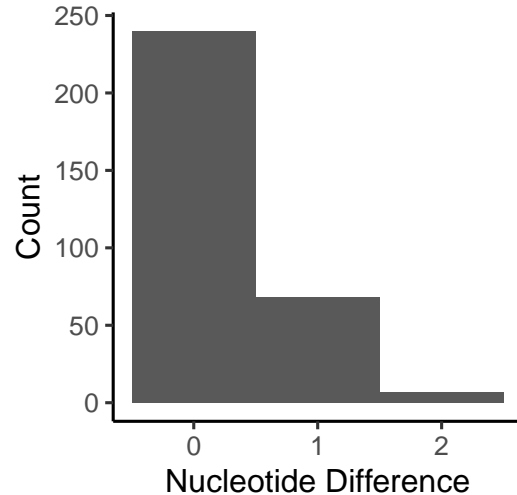
TRBV10–3*bp01

137 sequences assigned
118 (86.1%) exact matches, in which:
97 unique CDR3
13 unique J



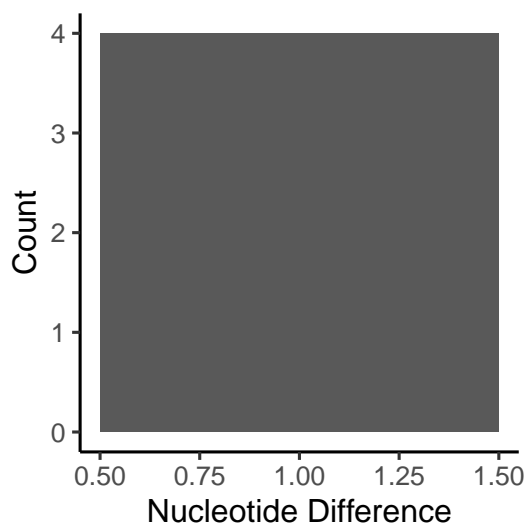
TRBV15*bp02

315 sequences assigned
240 (76.2%) exact matches, in which:
167 unique CDR3
13 unique J



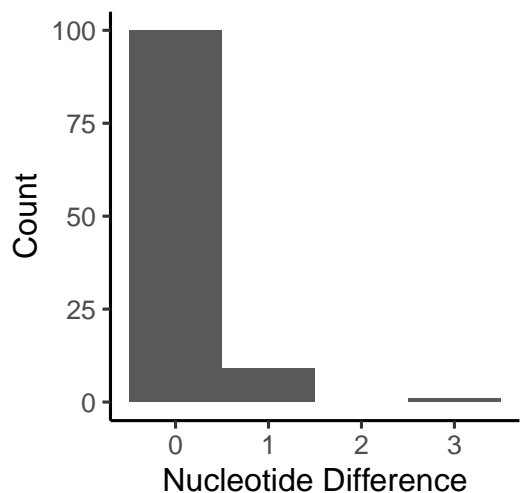
TRBV10–1*bp02

4 sequences assigned
No exact matches.



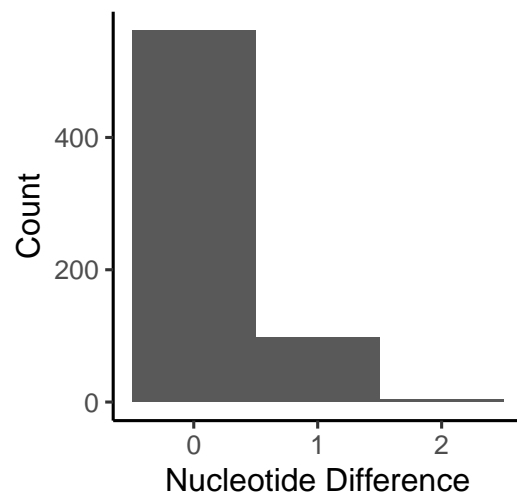
TRBV13*bp01

110 sequences assigned
100 (90.9%) exact matches, in which:
70 unique CDR3
11 unique J



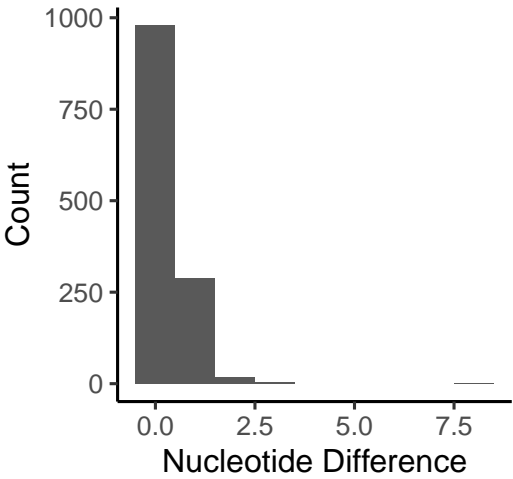
TRBV18*bp01

663 sequences assigned
562 (84.8%) exact matches, in which:
364 unique CDR3
13 unique J



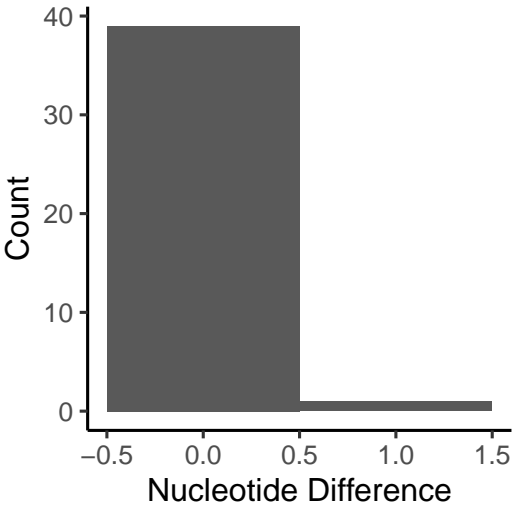
TRBV19*bp01

1290 sequences assigned
979 (75.9%) exact matches, in which:
649 unique CDR3
13 unique J



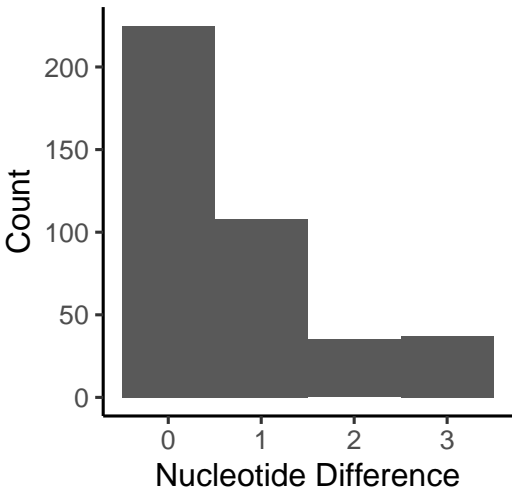
TRBV28*bp01

40 sequences assigned
39 (97.5%) exact matches, in which:
39 unique CDR3
11 unique J



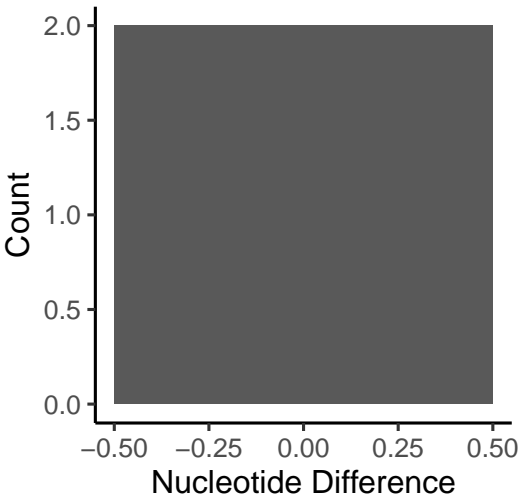
TRBV11-1*bp01

405 sequences assigned
225 (55.6%) exact matches, in which:
175 unique CDR3
13 unique J



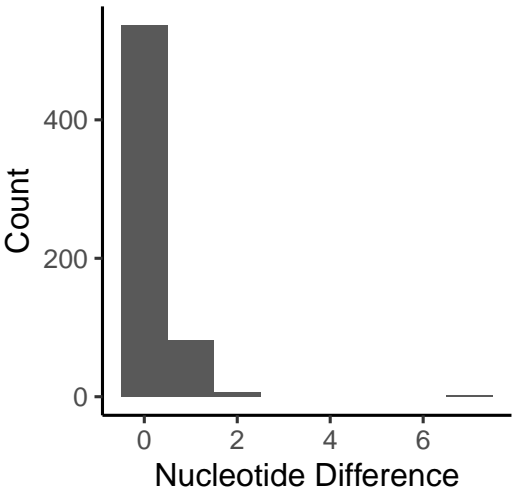
TRBV2*bp01

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



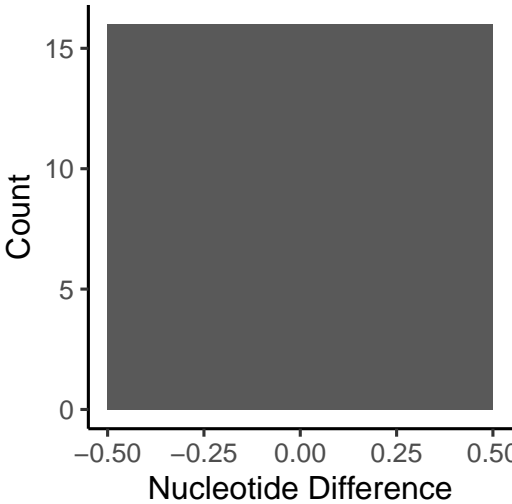
TRBV30*bp01

626 sequences assigned
537 (85.8%) exact matches, in which:
319 unique CDR3
13 unique J



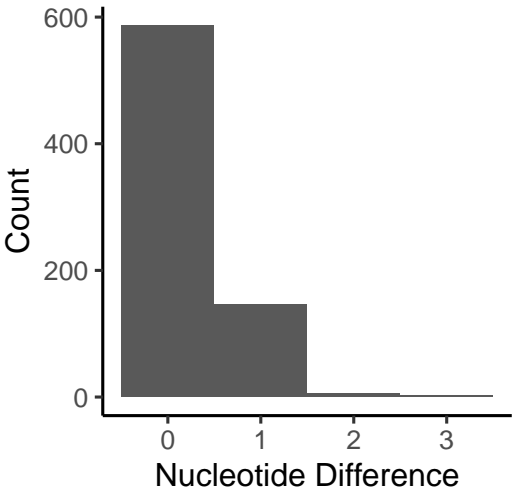
TRBV21-1*bp01

16 sequences assigned
16 (100%) exact matches, in which:
16 unique CDR3
9 unique J



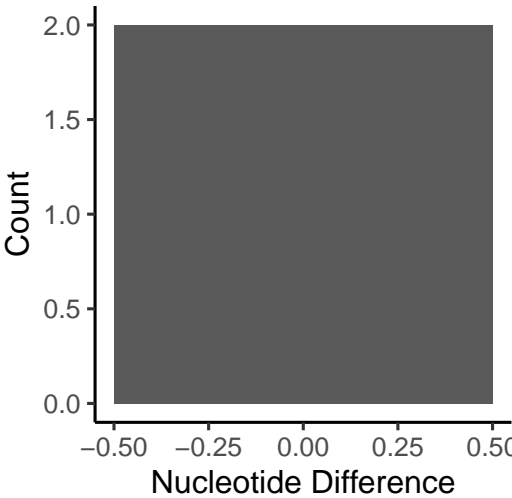
TRBV27*bp01

742 sequences assigned
587 (79.1%) exact matches, in which:
474 unique CDR3
13 unique J



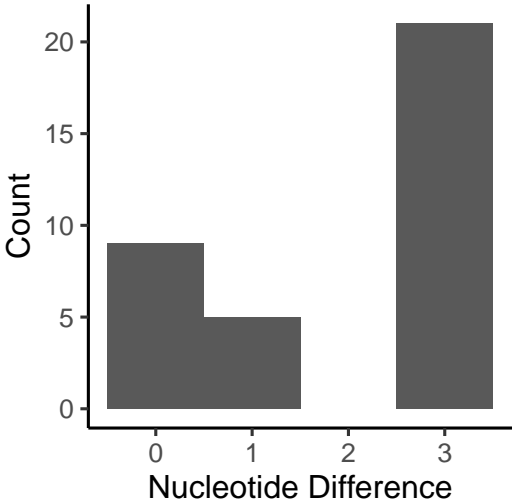
TRBV9*bp01

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



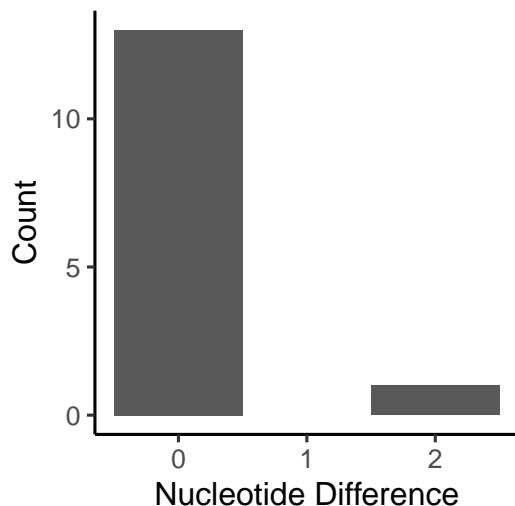
TRBV11-2*bp01

35 sequences assigned
9 (25.7%) exact matches, in which:
9 unique CDR3
7 unique J



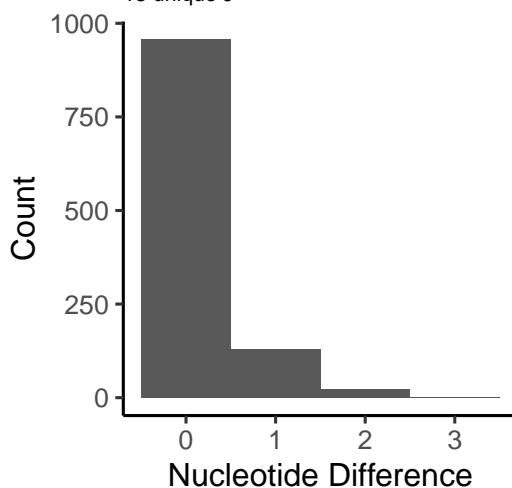
TRBV11-2*bp03

14 sequences assigned
13 (92.9%) exact matches, in which:
12 unique CDR3
8 unique J



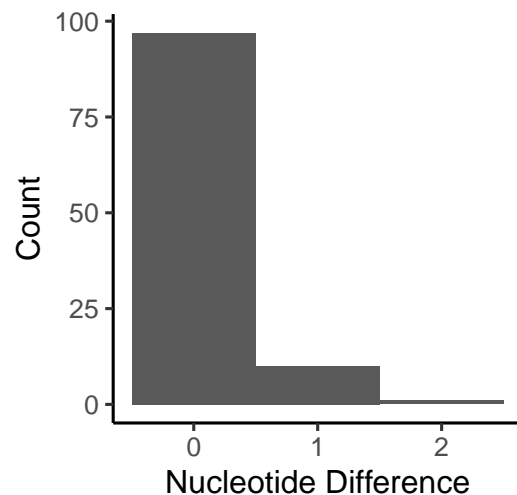
TRBV12-34*bp01

1110 sequences assigned
957 (86.2%) exact matches, in which:
640 unique CDR3
13 unique J



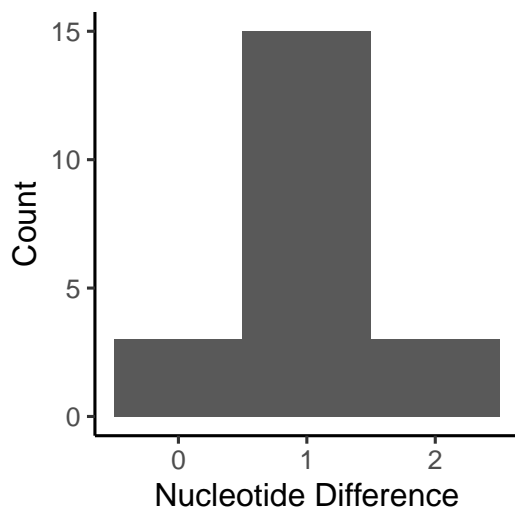
TRBV4-2*bp01

108 sequences assigned
97 (89.8%) exact matches, in which:
90 unique CDR3
10 unique J



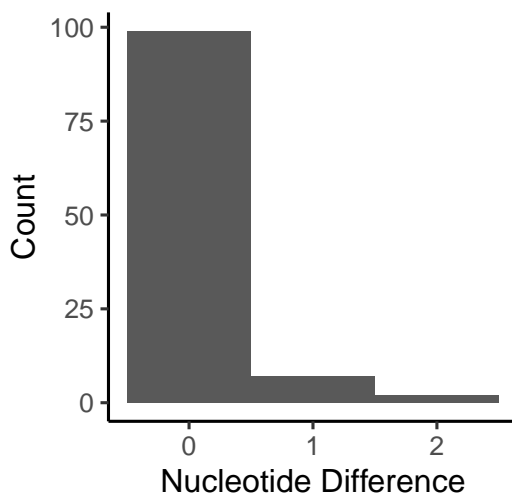
TRBV11-3*bp01

21 sequences assigned
3 (14.3%) exact matches, in which:
3 unique CDR3
2 unique J



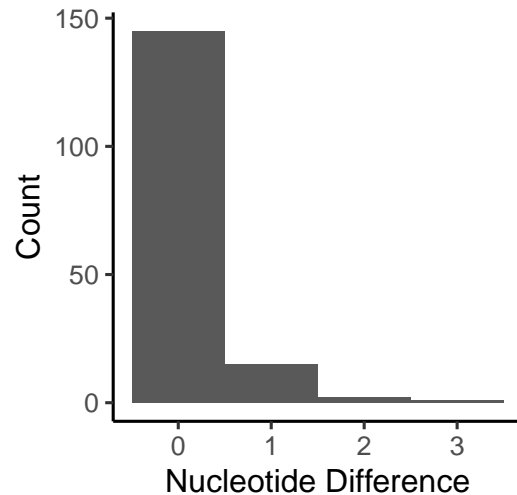
TRBV24-1*bp01

108 sequences assigned
99 (91.7%) exact matches, in which:
83 unique CDR3
11 unique J



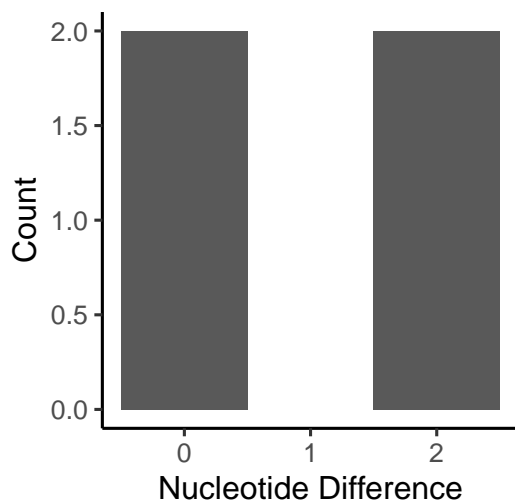
TRBV4-3*bp01

163 sequences assigned
145 (89%) exact matches, in which:
131 unique CDR3
13 unique J



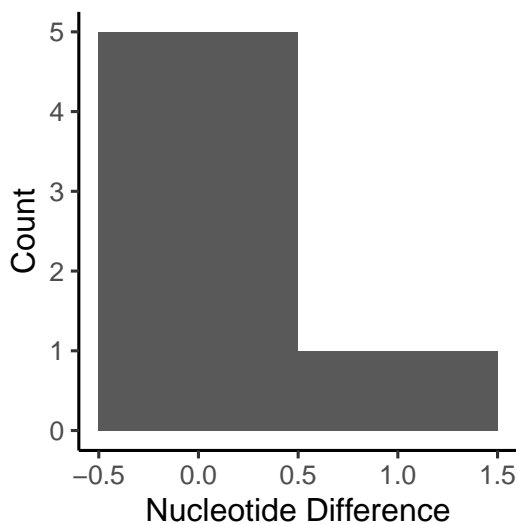
TRBV12-5*bp01

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



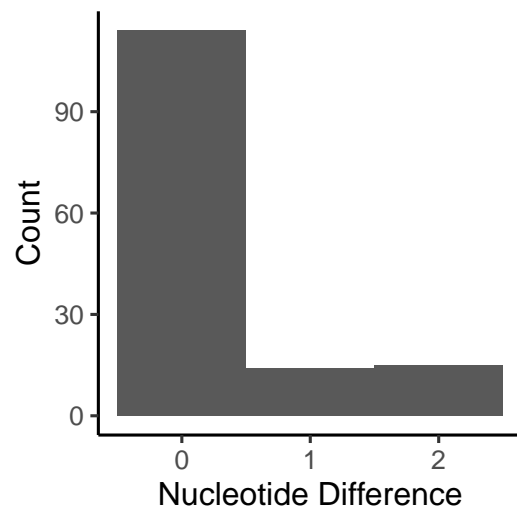
TRBV4-1*bp01

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



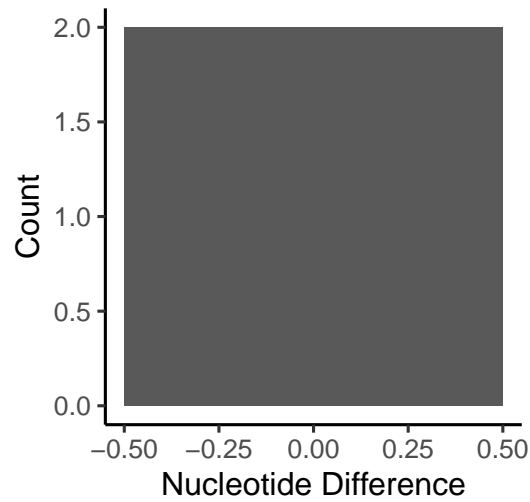
TRBV5-1*bp01

143 sequences assigned
114 (79.7%) exact matches, in which:
105 unique CDR3
12 unique J



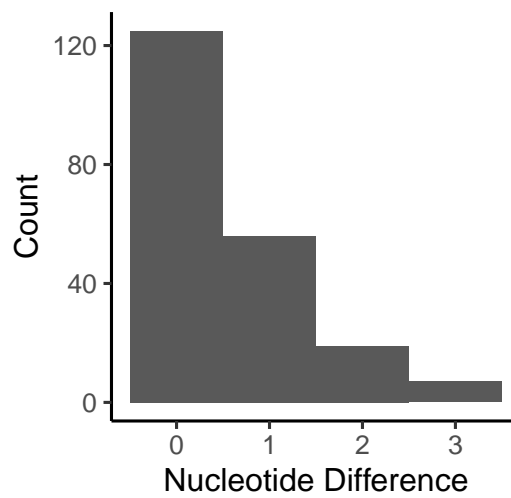
TRBV25-1*bp01

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



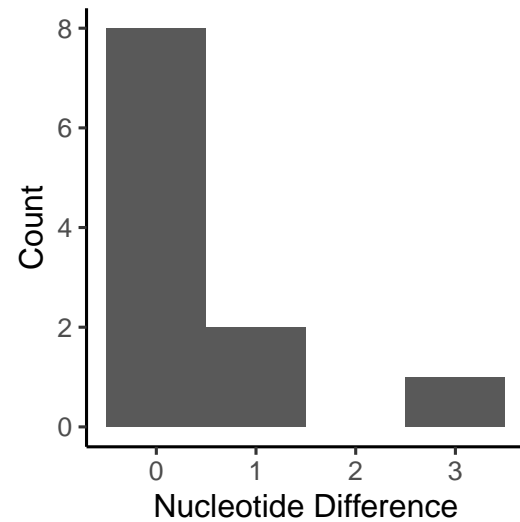
TRBV5-5*bp01

207 sequences assigned
125 (60.4%) exact matches, in which:
101 unique CDR3
12 unique J



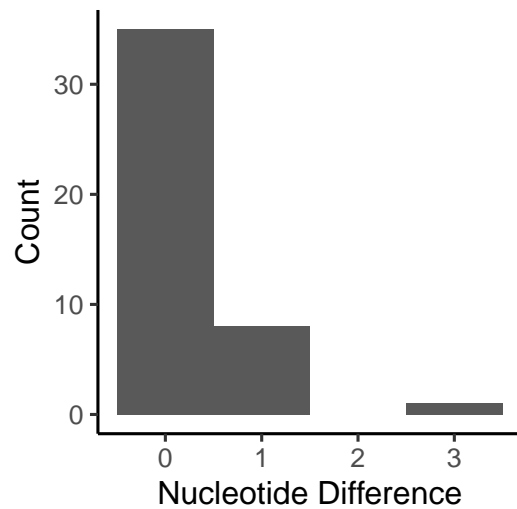
TRBV5-8*bp01

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



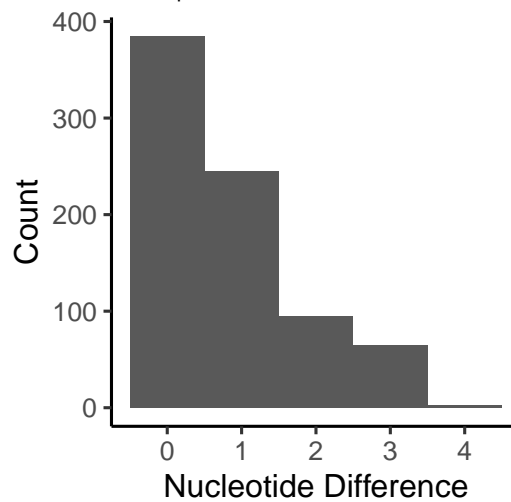
TRBV5-3*bp01

44 sequences assigned
35 (79.5%) exact matches, in which:
32 unique CDR3
11 unique J



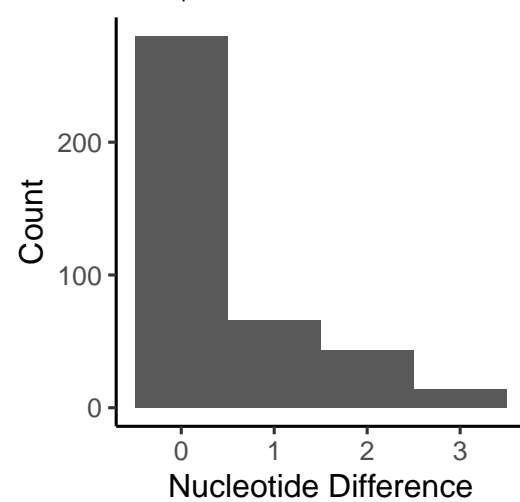
TRBV5-6*bp01

793 sequences assigned
385 (48.5%) exact matches, in which:
294 unique CDR3
13 unique J



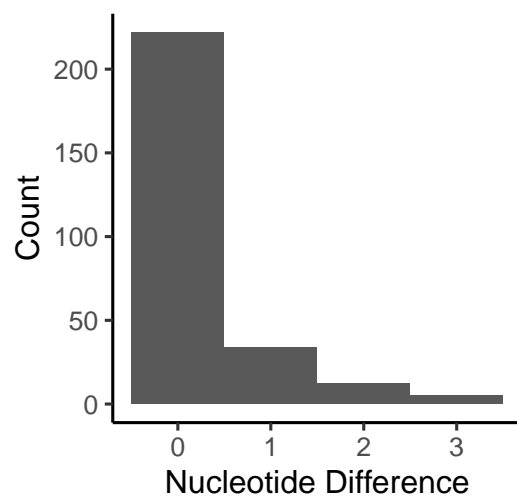
TRBV6-1*bp01

403 sequences assigned
280 (69.5%) exact matches, in which:
243 unique CDR3
13 unique J



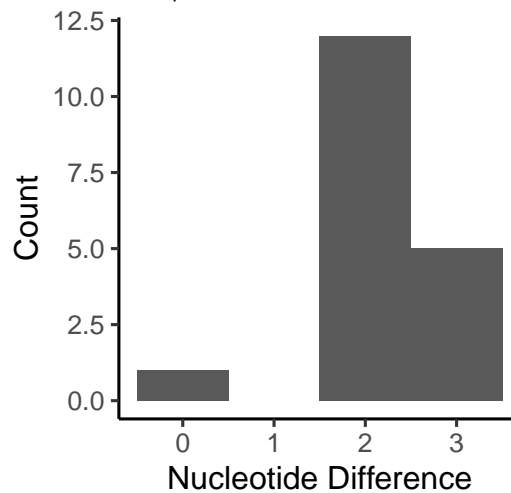
TRBV5-4*bp01

273 sequences assigned
222 (81.3%) exact matches, in which:
171 unique CDR3
13 unique J



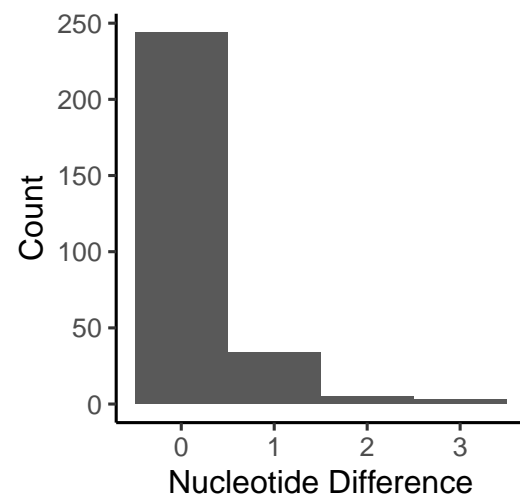
TRBV5-7*bp01

18 sequences assigned
1 (5.6%) exact matches, in which:
1 unique CDR3
1 unique J



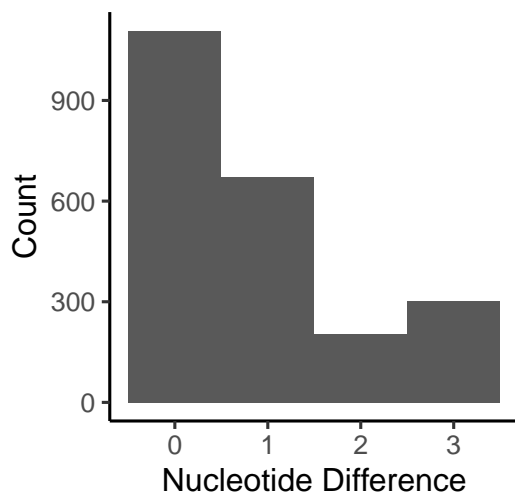
TRBV6-4*bp01

286 sequences assigned
244 (85.3%) exact matches, in which:
196 unique CDR3
13 unique J



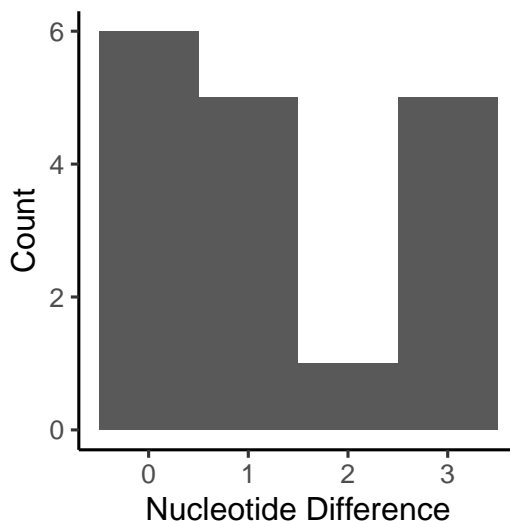
TRBV6-5*bp01

2286 sequences assigned
1108 (48.5%) exact matches, in which:
860 unique CDR3
13 unique J



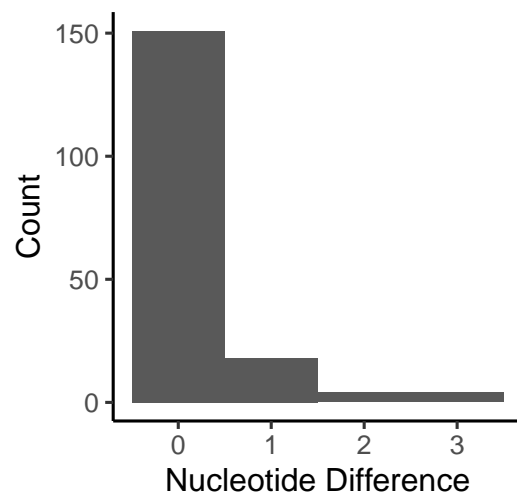
TRBV6-8*bp01

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



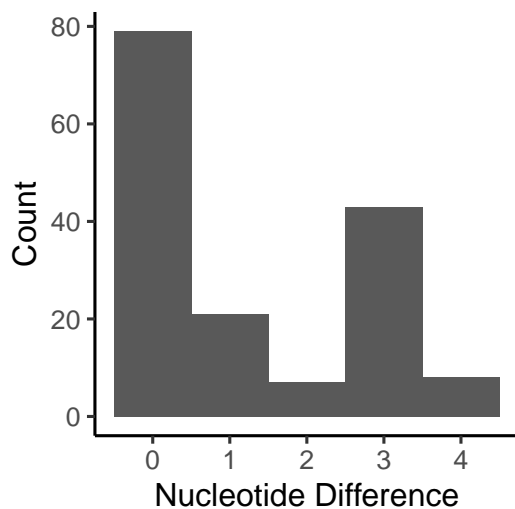
TRBV7-2*bp01

177 sequences assigned
151 (85.3%) exact matches, in which:
136 unique CDR3
13 unique J



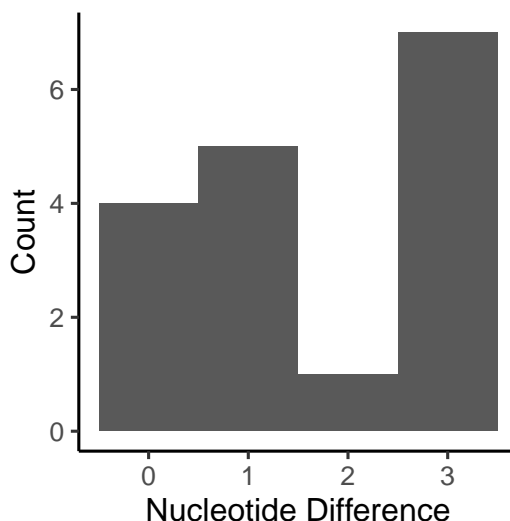
TRBV6-6*bp01

158 sequences assigned
79 (50%) exact matches, in which:
73 unique CDR3
12 unique J



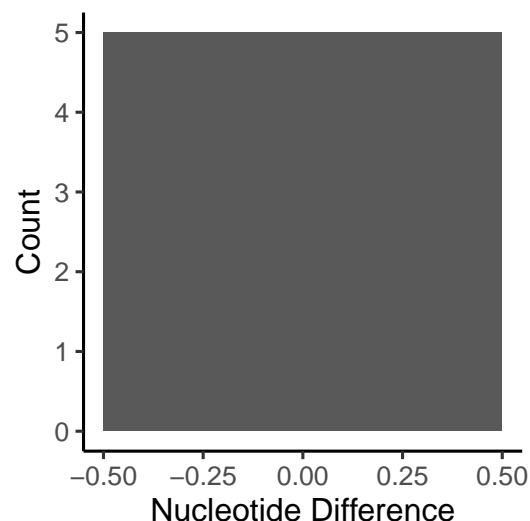
TRBV6-9*bp01

17 sequences assigned
4 (23.5%) exact matches, in which:
4 unique CDR3
4 unique J



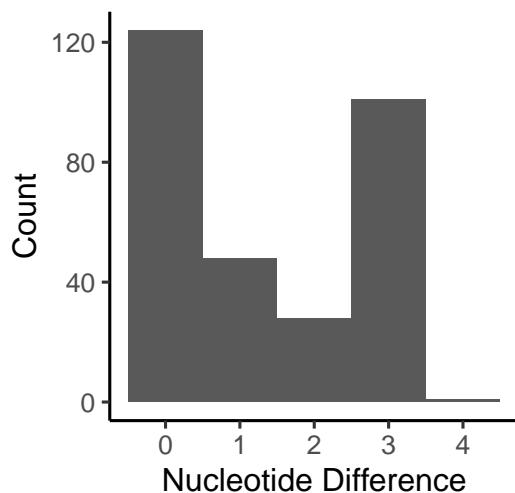
TRBV7-3*bp01

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



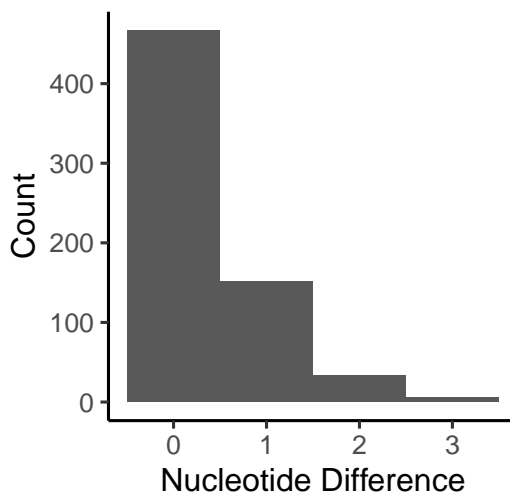
TRBV6-6*bp02

302 sequences assigned
124 (41.1%) exact matches, in which:
112 unique CDR3
12 unique J



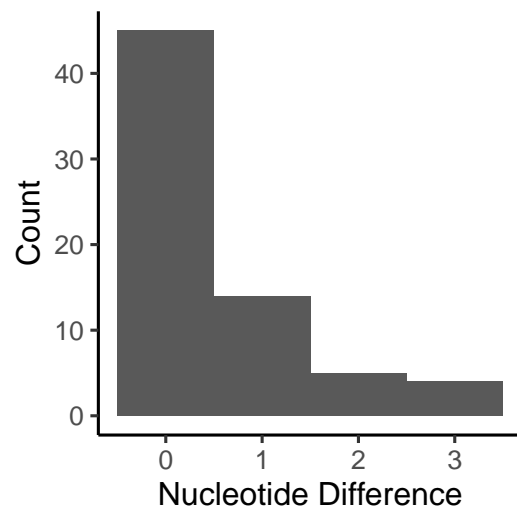
TRBV6-23*bp01

657 sequences assigned
467 (71.1%) exact matches, in which:
405 unique CDR3
13 unique J



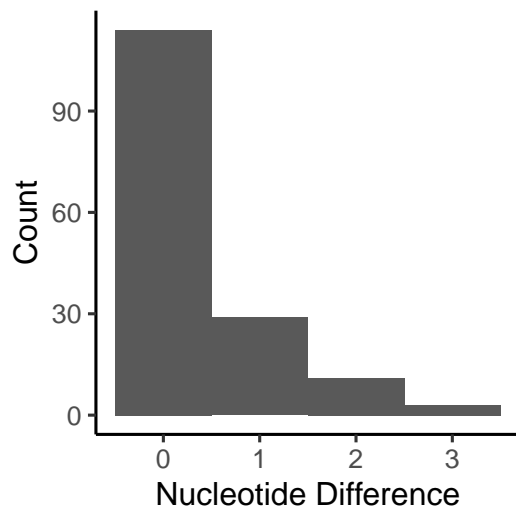
TRBV7-4*bp01

68 sequences assigned
45 (66.2%) exact matches, in which:
36 unique CDR3
13 unique J



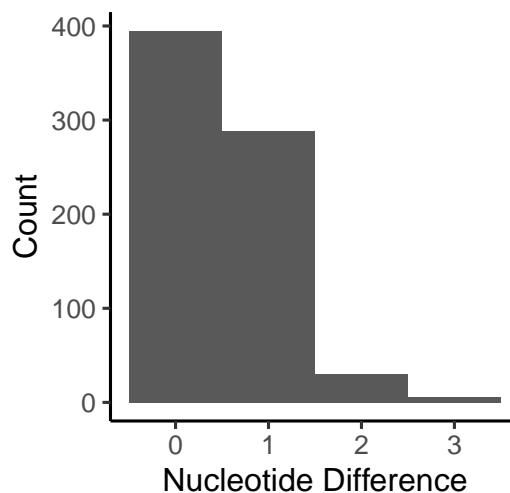
TRBV7-6*bp01

157 sequences assigned
114 (72.6%) exact matches, in which:
91 unique CDR3
12 unique J



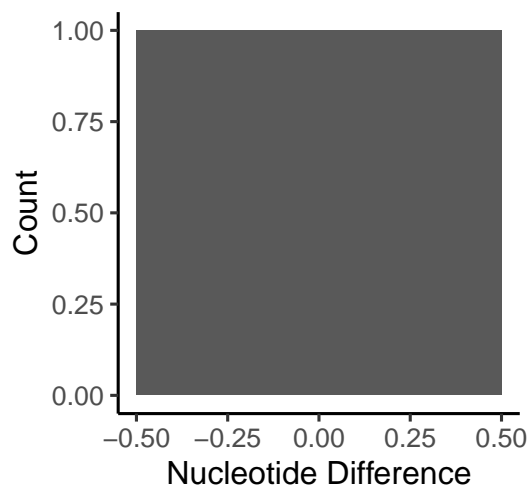
TRBV29-1*bp01

719 sequences assigned
395 (54.9%) exact matches, in which:
286 unique CDR3
13 unique J



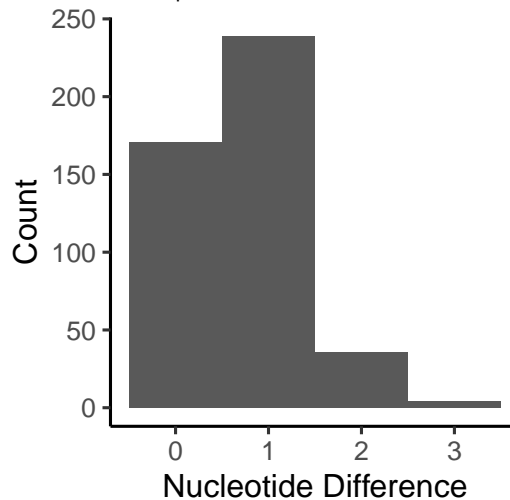
TRBV7-7*bp01

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



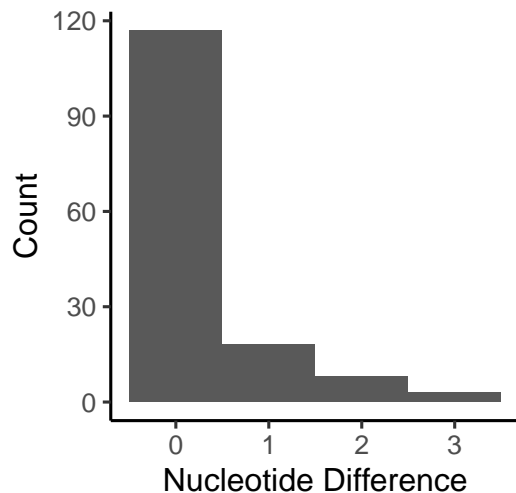
TRBV29-1*bp03

450 sequences assigned
171 (38%) exact matches, in which:
102 unique CDR3
12 unique J



TRBV7-9*bp01

146 sequences assigned
117 (80.1%) exact matches, in which:
108 unique CDR3
12 unique J





Warning – no inferred sequences found.