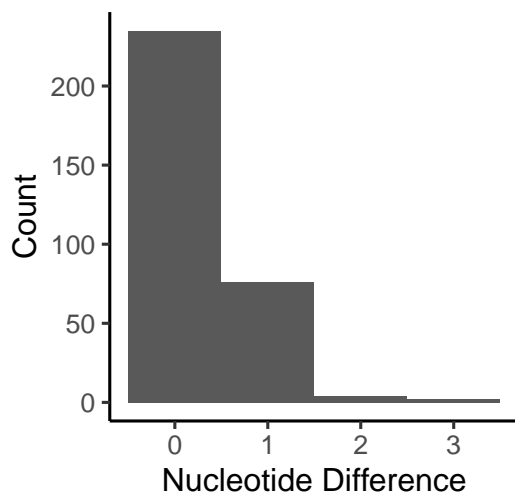


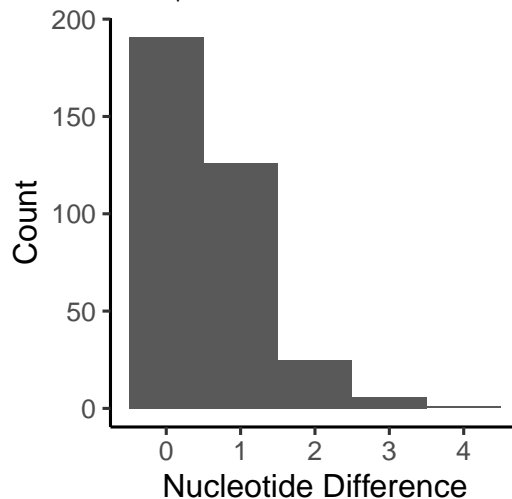
TRBV20–1*bp01

317 sequences assigned
235 (74.1%) exact matches, in which:
164 unique CDR3
12 unique J



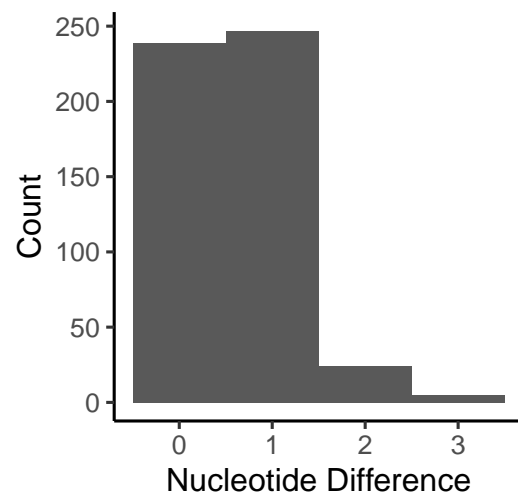
TRBV10–3*bp01

349 sequences assigned
191 (54.7%) exact matches, in which:
140 unique CDR3
13 unique J



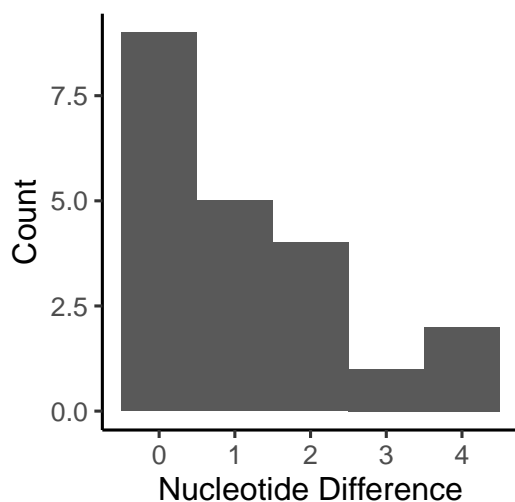
TRBV15*bp02

515 sequences assigned
239 (46.4%) exact matches, in which:
130 unique CDR3
13 unique J



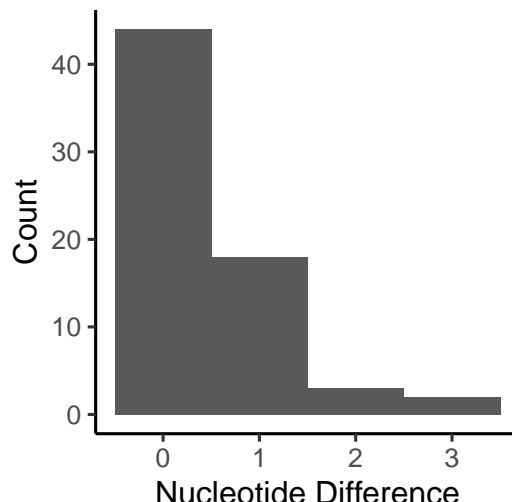
TRBV10–1*bp01

21 sequences assigned
9 (42.9%) exact matches, in which:
6 unique CDR3
4 unique J



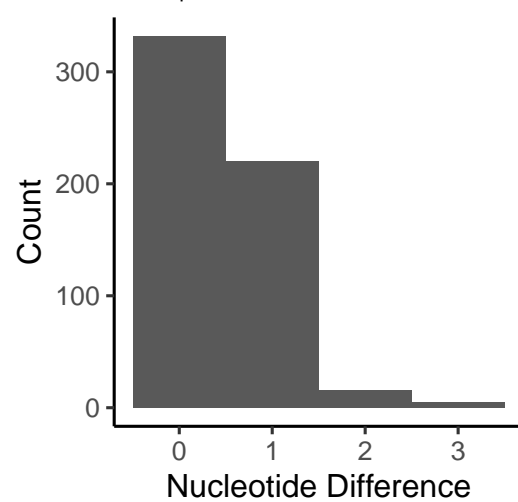
TRBV13*bp01

67 sequences assigned
44 (65.7%) exact matches, in which:
35 unique CDR3
9 unique J



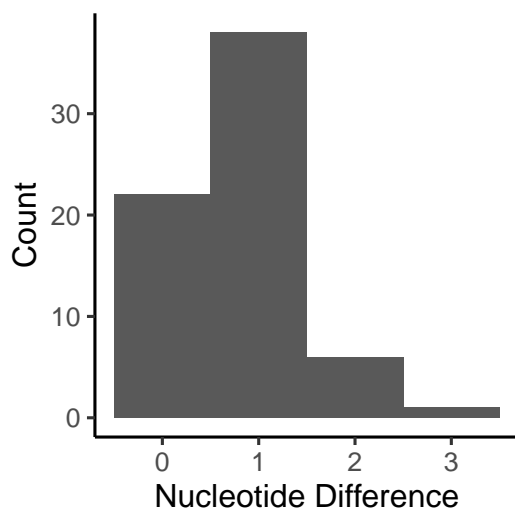
TRBV18*bp01

573 sequences assigned
332 (57.9%) exact matches, in which:
215 unique CDR3
12 unique J



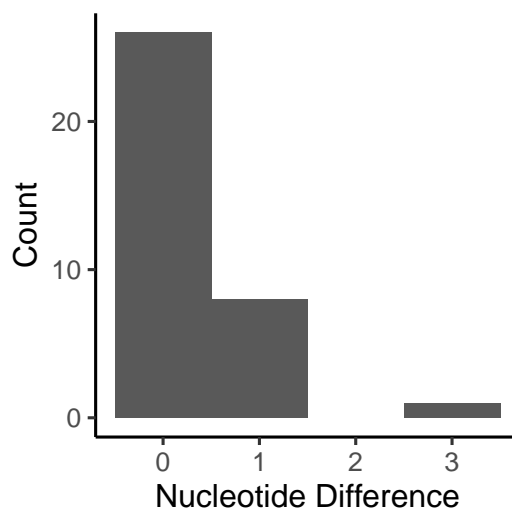
TRBV10–2*bp01

67 sequences assigned
22 (32.8%) exact matches, in which:
17 unique CDR3
7 unique J



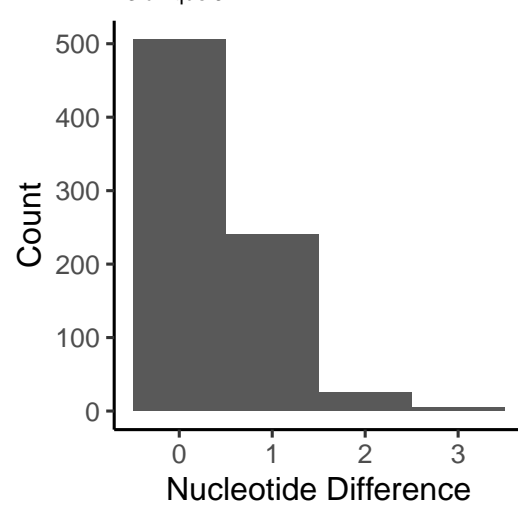
TRBV14*bp01

35 sequences assigned
26 (74.3%) exact matches, in which:
21 unique CDR3
9 unique J



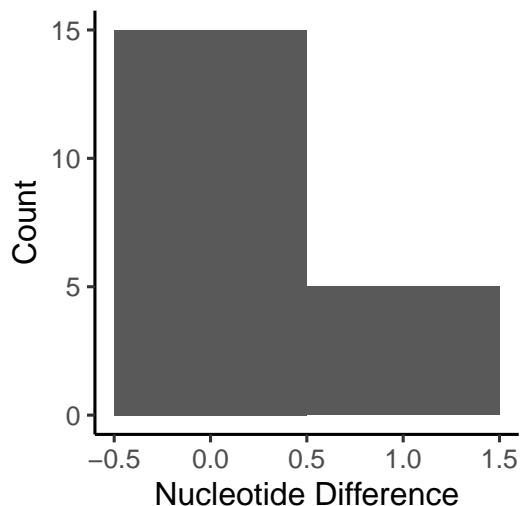
TRBV19*bp01

776 sequences assigned
506 (65.2%) exact matches, in which:
363 unique CDR3
13 unique J



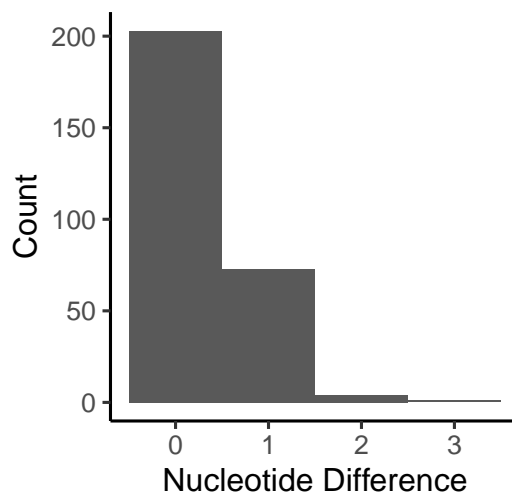
TRBV2*bp01

20 sequences assigned
15 (75%) exact matches, in which:
15 unique CDR3
8 unique J



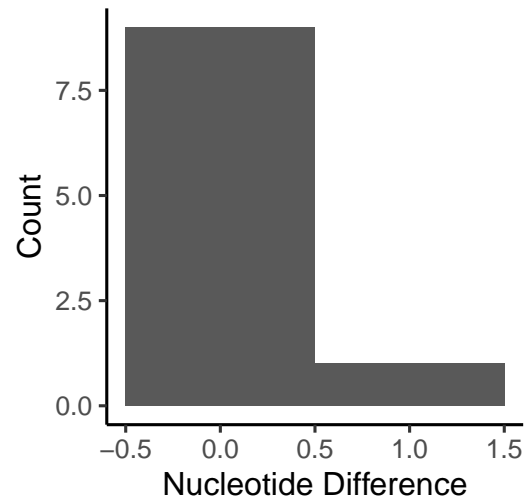
TRBV30*bp01

281 sequences assigned
203 (72.2%) exact matches, in which:
104 unique CDR3
12 unique J



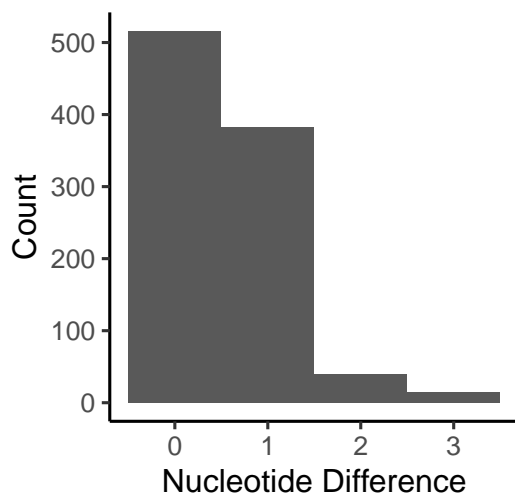
TRBV21-1*bp02

10 sequences assigned
9 (90%) exact matches, in which:
9 unique CDR3
5 unique J



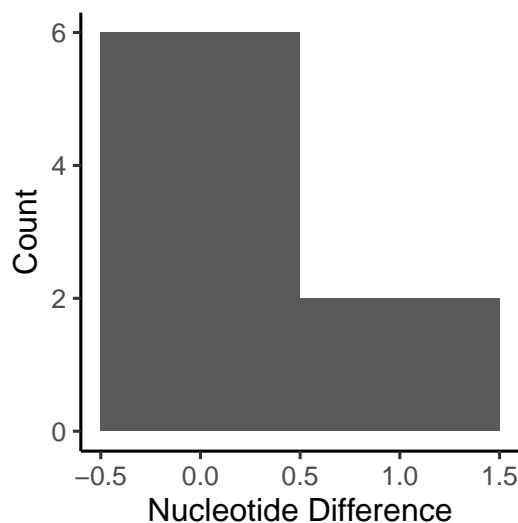
TRBV27*bp01

954 sequences assigned
516 (54.1%) exact matches, in which:
384 unique CDR3
13 unique J



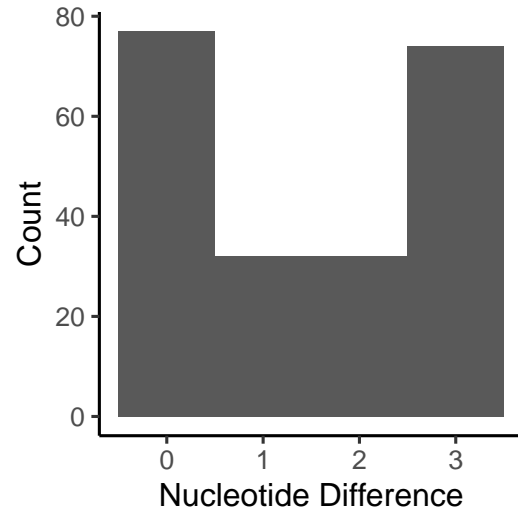
TRBV9*bp01

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



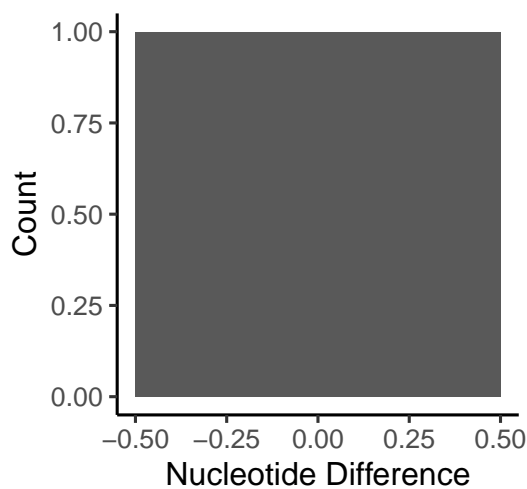
TRBV11-2*bp01

215 sequences assigned
77 (35.8%) exact matches, in which:
64 unique CDR3
12 unique J



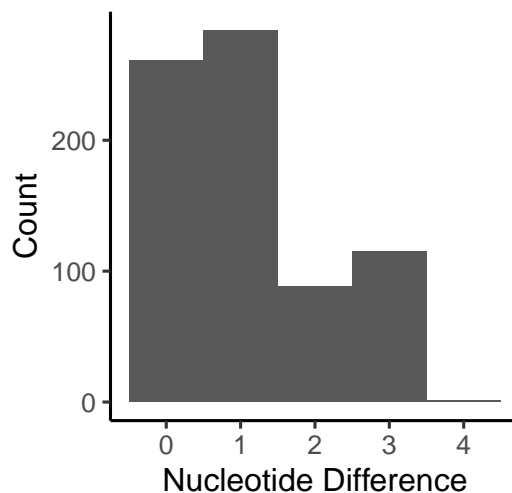
TRBV28*bp01

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



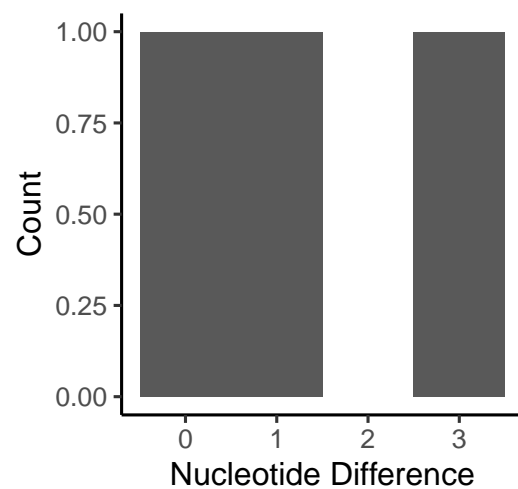
TRBV11-1*bp01

749 sequences assigned
261 (34.8%) exact matches, in which:
168 unique CDR3
13 unique J



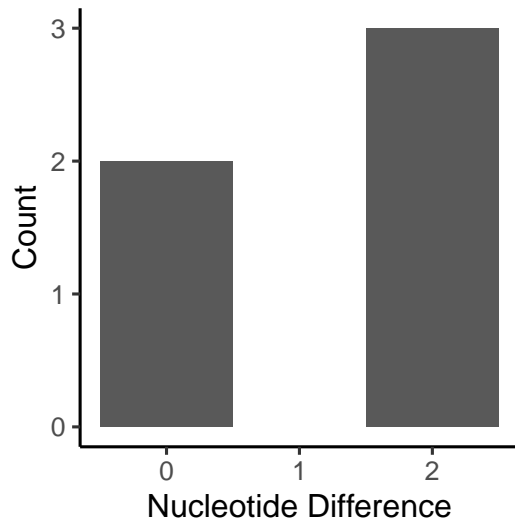
TRBV11-3*bp01

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



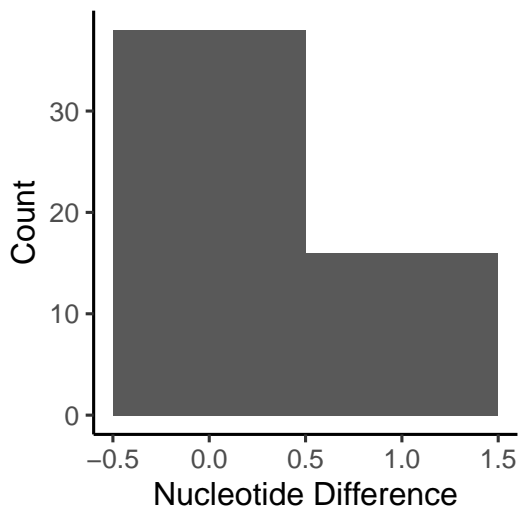
TRBV12–5*bp01

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



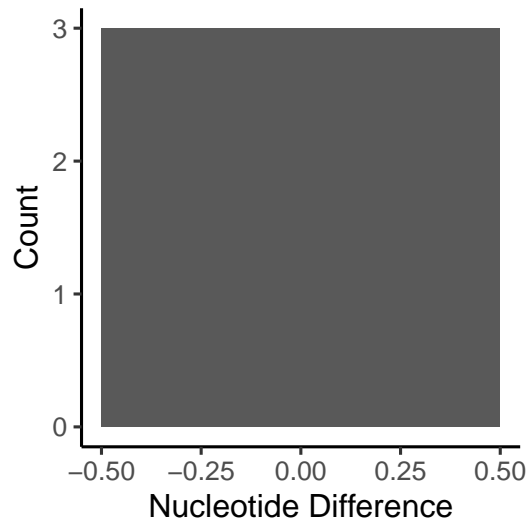
TRBV24–1*bp02

54 sequences assigned
38 (70.4%) exact matches, in which:
34 unique CDR3
8 unique J



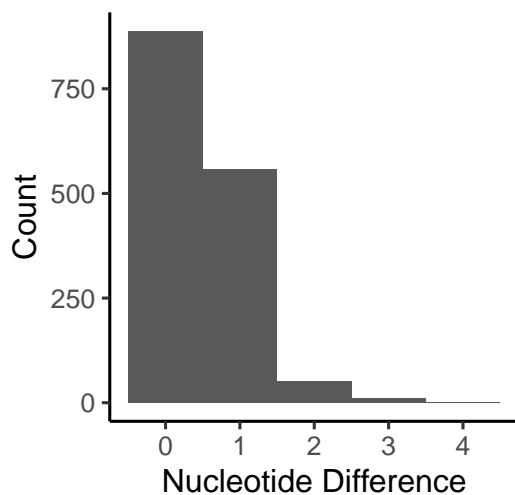
TRBV25–1*bp01

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



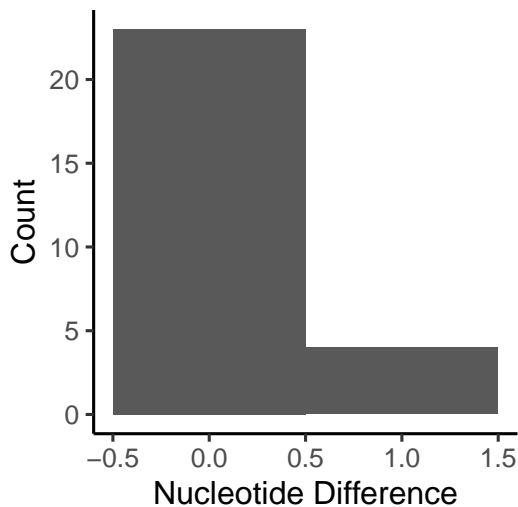
TRBV12–34*bp01

1511 sequences assigned
888 (58.8%) exact matches, in which:
501 unique CDR3
13 unique J



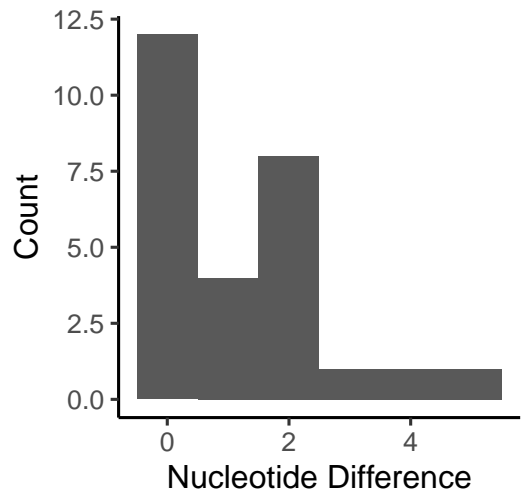
TRBV4–2*bp01

27 sequences assigned
23 (85.2%) exact matches, in which:
20 unique CDR3
7 unique J



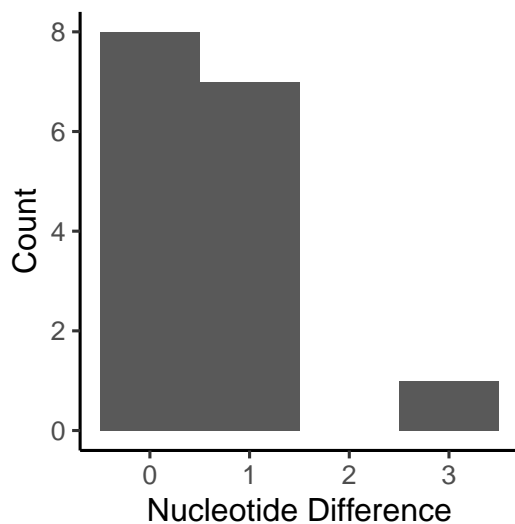
TRBV5–3*bp01

27 sequences assigned
12 (44.4%) exact matches, in which:
11 unique CDR3
8 unique J



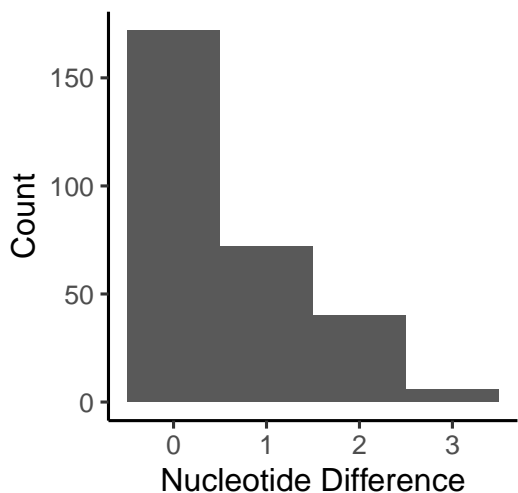
TRBV4–1*bp01

16 sequences assigned
8 (50%) exact matches, in which:
8 unique CDR3
5 unique J



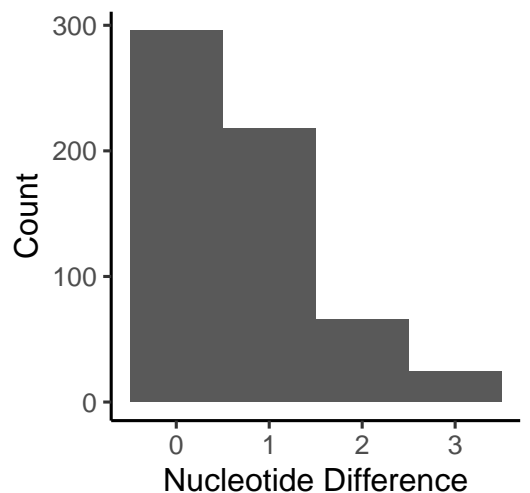
TRBV5–1*bp01

290 sequences assigned
172 (59.3%) exact matches, in which:
137 unique CDR3
12 unique J



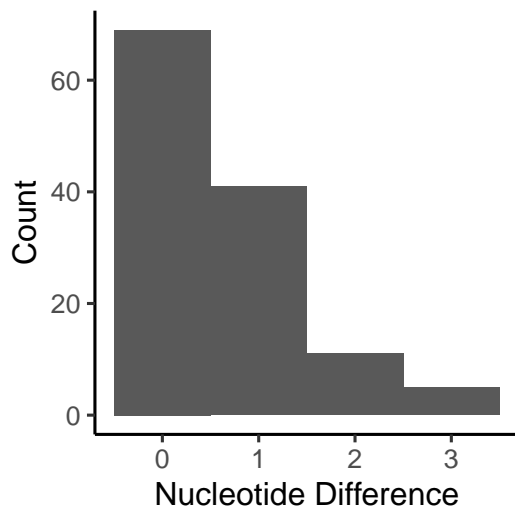
TRBV5–4*bp01

604 sequences assigned
296 (49%) exact matches, in which:
226 unique CDR3
13 unique J



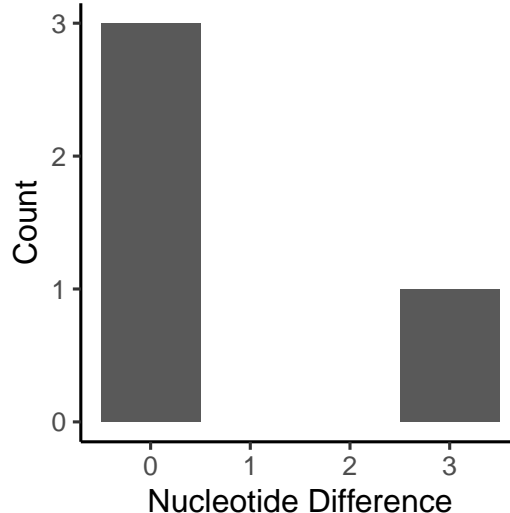
TRBV5-5*bp01

126 sequences assigned
69 (54.8%) exact matches, in which:
52 unique CDR3
12 unique J



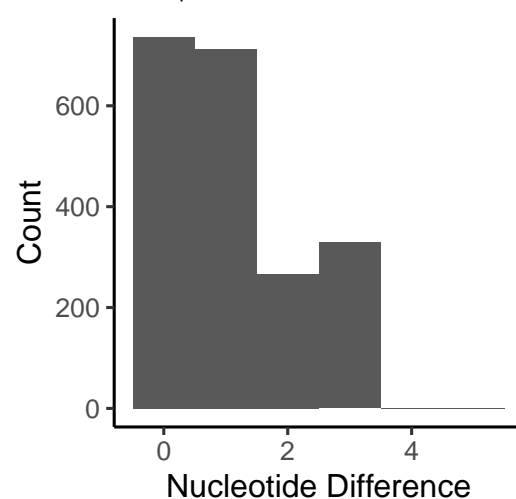
TRBV5-8*bp01

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



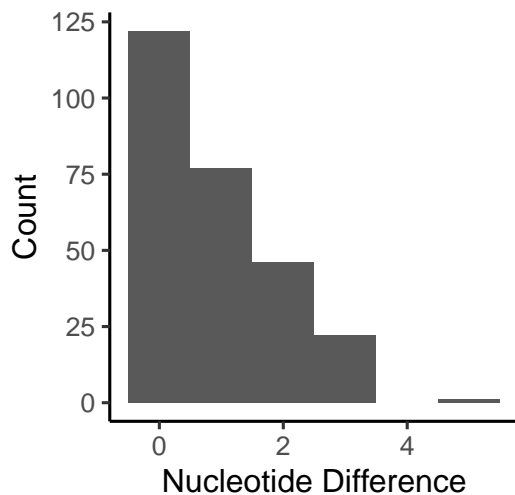
TRBV6-5*bp01

2048 sequences assigned
737 (36%) exact matches, in which:
560 unique CDR3
13 unique J



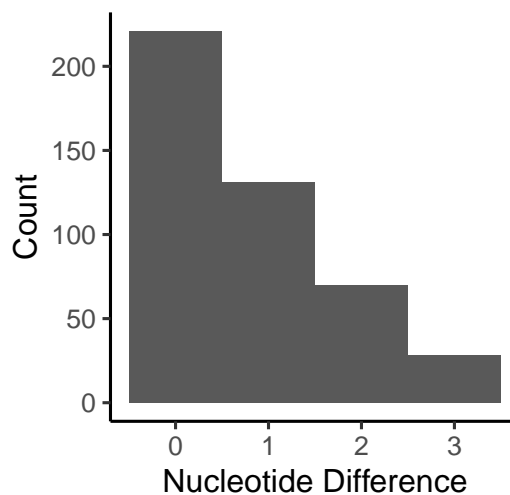
TRBV5-6*bp01

268 sequences assigned
122 (45.5%) exact matches, in which:
94 unique CDR3
12 unique J



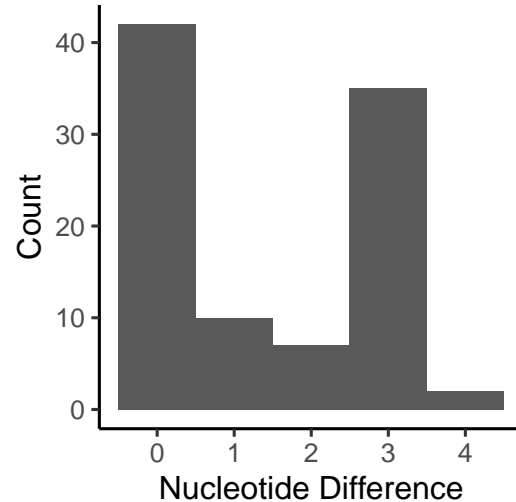
TRBV6-1*bp01

450 sequences assigned
221 (49.1%) exact matches, in which:
180 unique CDR3
13 unique J



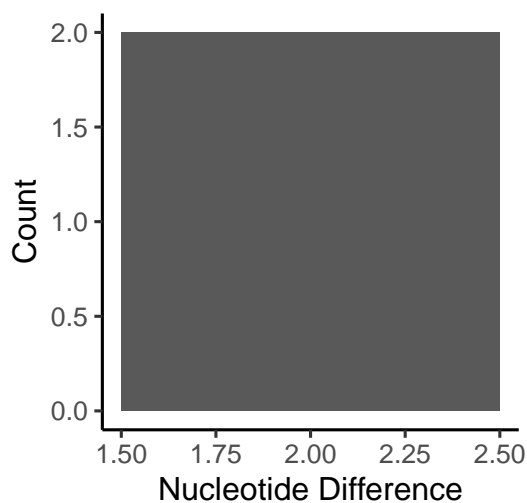
TRBV6-6*bp01

96 sequences assigned
42 (43.8%) exact matches, in which:
39 unique CDR3
12 unique J



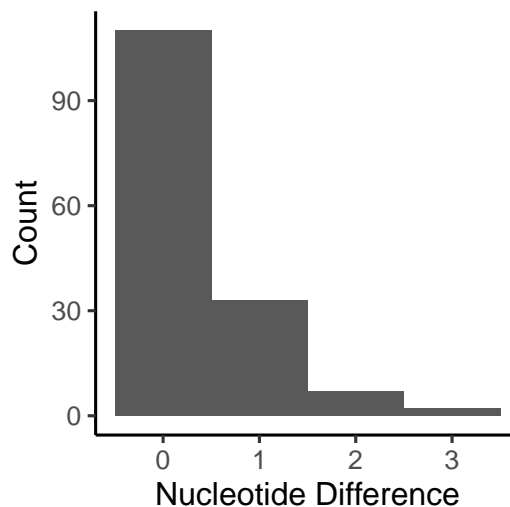
TRBV5-7*bp01

2 sequences assigned
No exact matches.



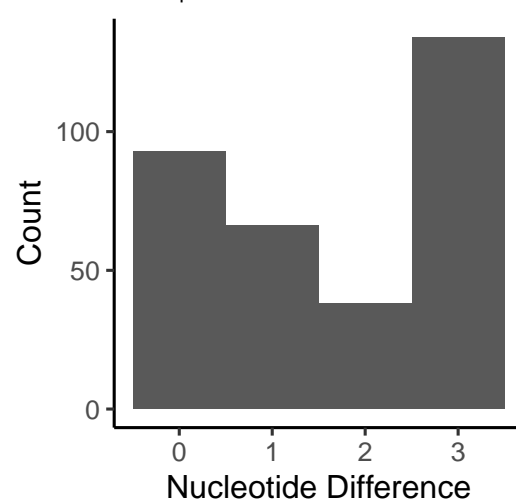
TRBV6-4*bp01

152 sequences assigned
110 (72.4%) exact matches, in which:
82 unique CDR3
9 unique J



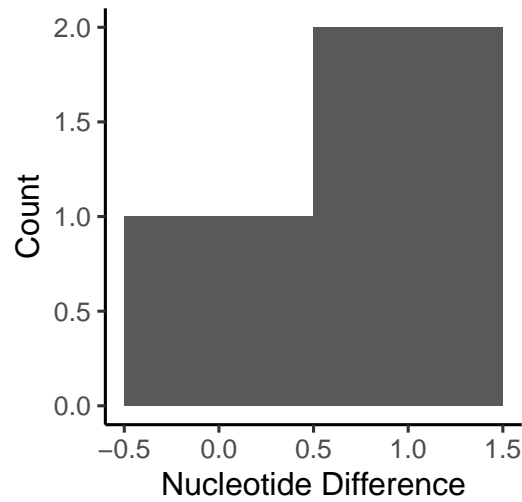
TRBV6-6*bp02

331 sequences assigned
93 (28.1%) exact matches, in which:
83 unique CDR3
12 unique J



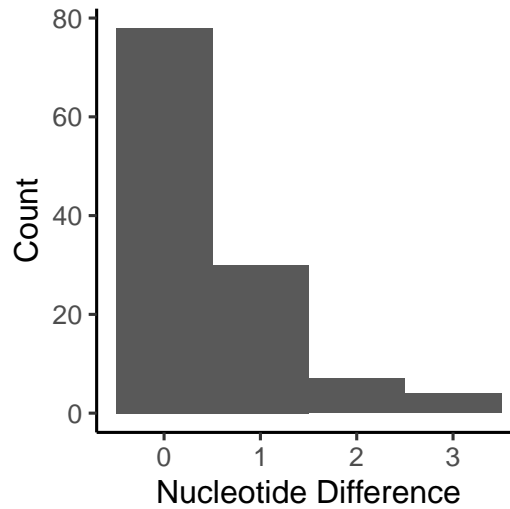
TRBV6–8*bp01

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



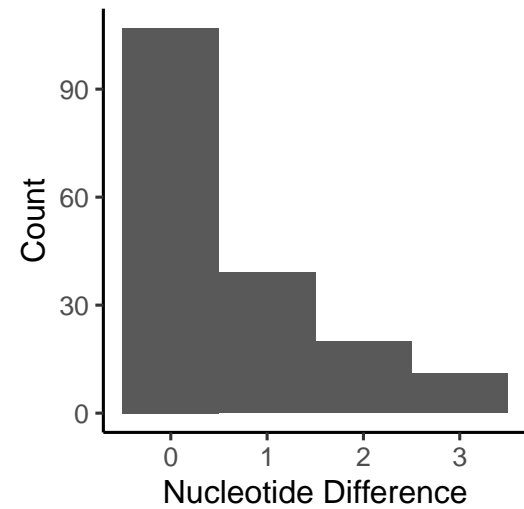
TRBV7–2*bp02

119 sequences assigned
78 (65.5%) exact matches, in which:
60 unique CDR3
13 unique J



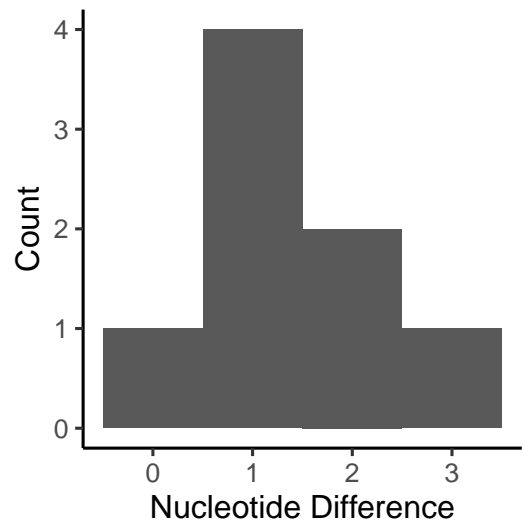
TRBV7–6*bp01

177 sequences assigned
107 (60.5%) exact matches, in which:
71 unique CDR3
12 unique J



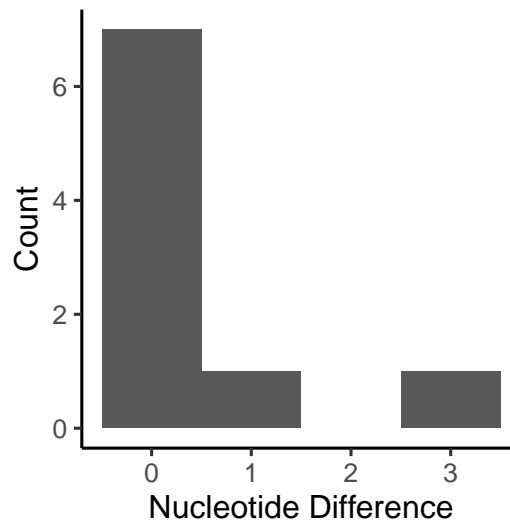
TRBV6–9*bp01

8 sequences assigned
1 (12.5%) exact matches, in which:
1 unique CDR3
1 unique J



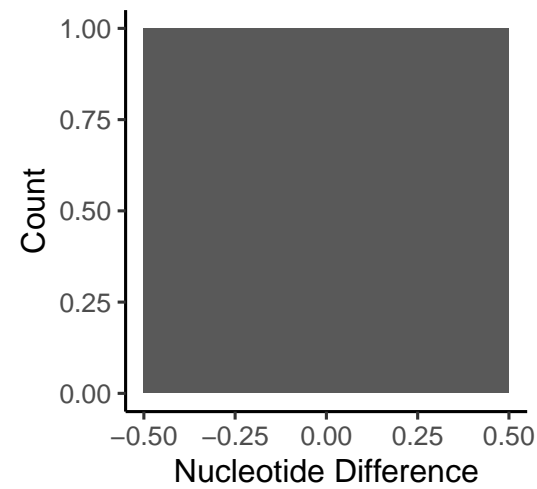
TRBV7–3*bp01

9 sequences assigned
7 (77.8%) exact matches, in which:
7 unique CDR3
5 unique J



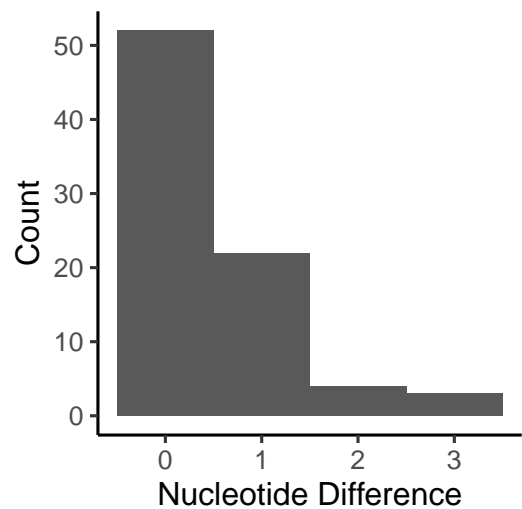
TRBV7–7*bp01

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



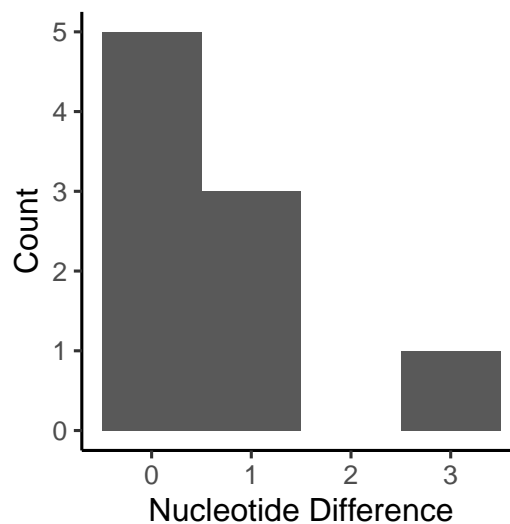
TRBV6–23*bp01

81 sequences assigned
52 (64.2%) exact matches, in which:
47 unique CDR3
12 unique J



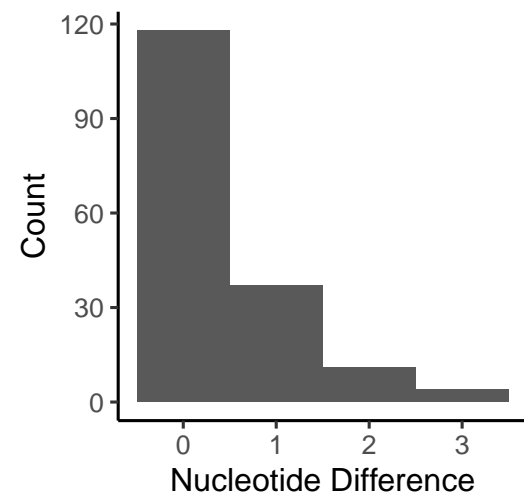
TRBV7–4*bp01

9 sequences assigned
5 (55.6%) exact matches, in which:
4 unique CDR3
2 unique J



TRBV7–9*bp01

170 sequences assigned
118 (69.4%) exact matches, in which:
97 unique CDR3
13 unique J



TRBV29-1*bp01

985 sequences assigned
437 (44.4%) exact matches, in which:
275 unique CDR3
12 unique J

