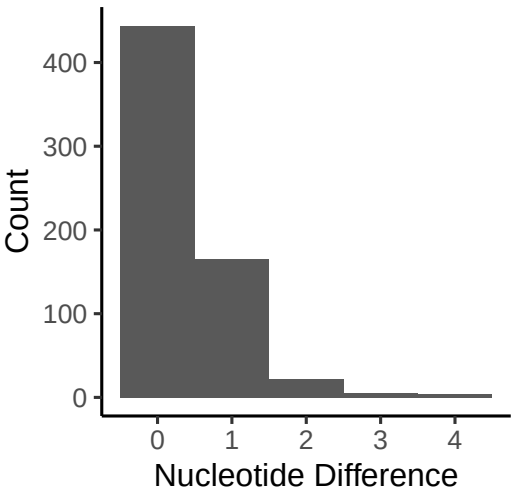
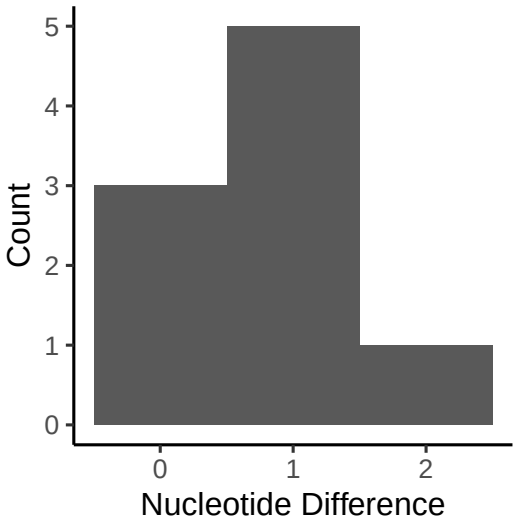
TRBV20−1*bp01

640 sequences assigned
444 (69.4%) exact matches, in which:
178 unique CDR3
12 unique J



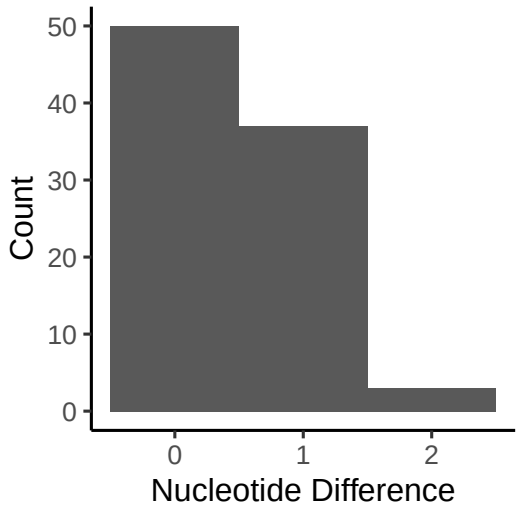
TRBV10−2*bp01

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



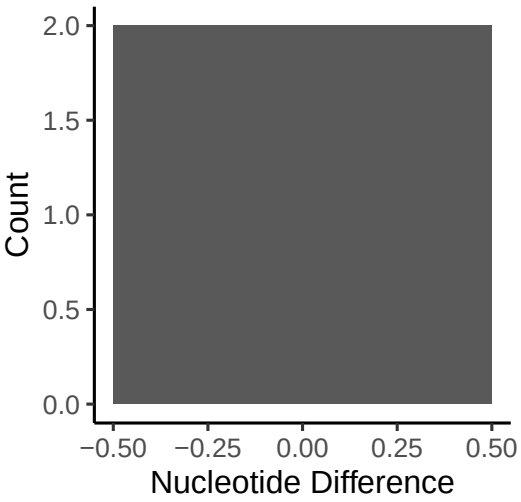
TRBV14*bp01

90 sequences assigned
50 (55.6%) exact matches, in which:
31 unique CDR3
6 unique J



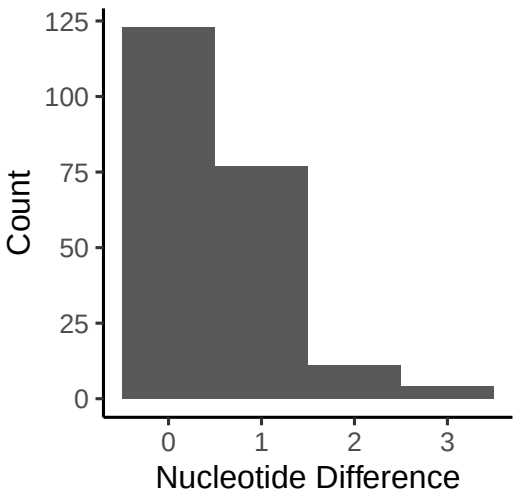
TRBV10−1*bp01

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



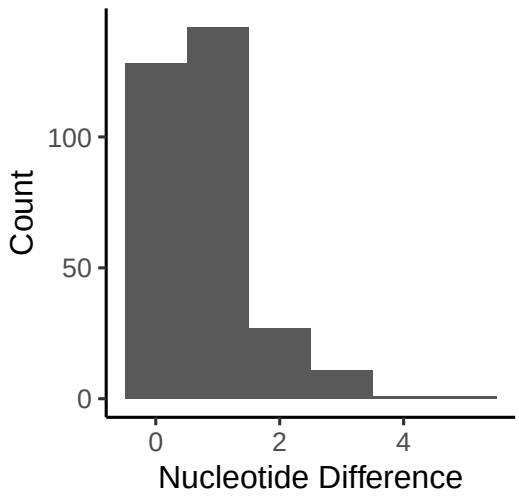
TRBV10−3*bp01

215 sequences assigned
123 (57.2%) exact matches, in which:
73 unique CDR3
10 unique J



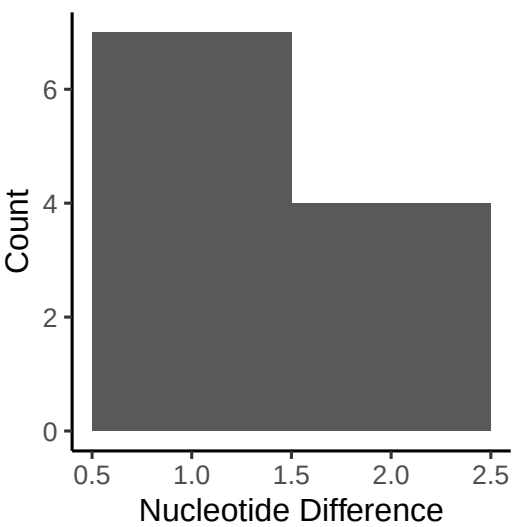
TRBV15*bp02

310 sequences assigned
128 (41.3%) exact matches, in which:
60 unique CDR3
7 unique J



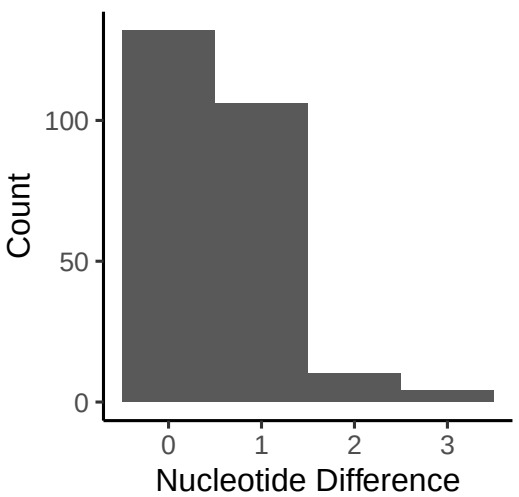
TRBV10−1*bp02

11 sequences assigned
No exact matches.



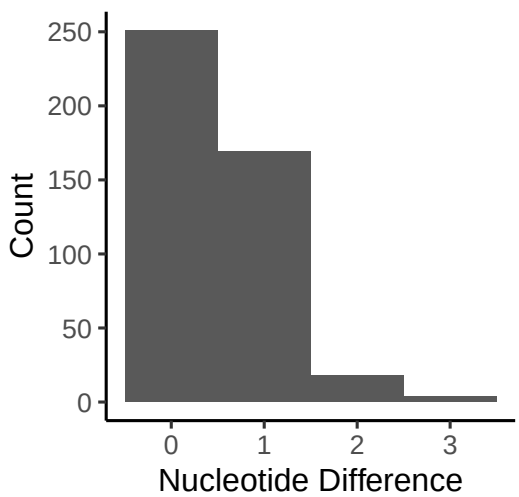
TRBV13*bp01

252 sequences assigned
132 (52.4%) exact matches, in which:
81 unique CDR3
10 unique J



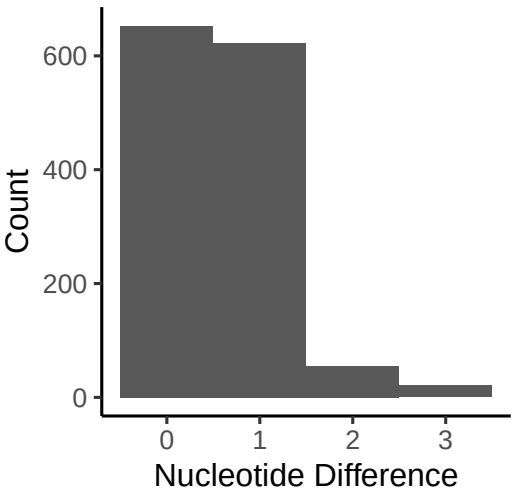
TRBV18*bp01

442 sequences assigned
251 (56.8%) exact matches, in which:
110 unique CDR3
13 unique J



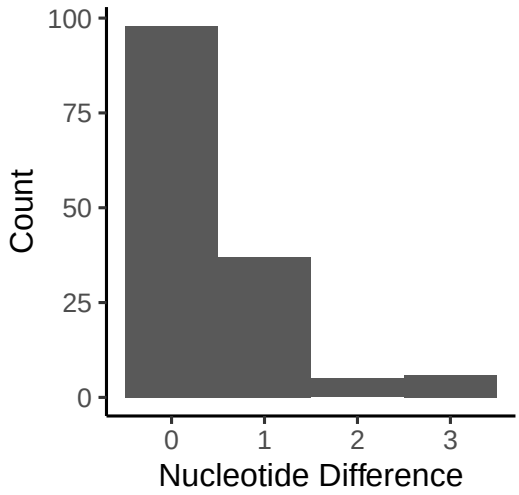
TRBV19*bp01

1351 sequences assigned
653 (48.3%) exact matches, in which:
316 unique CDR3
13 unique J



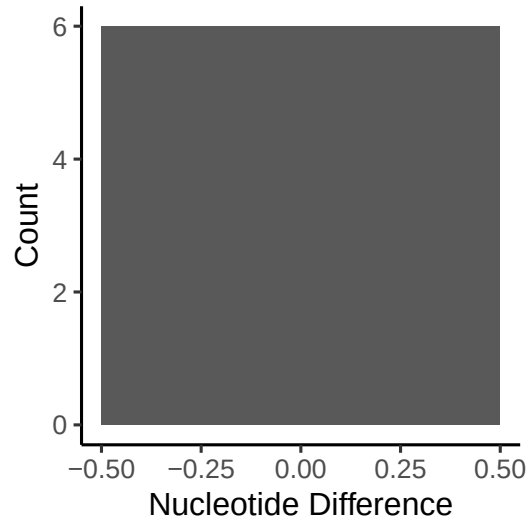
TRBV28*bp01

146 sequences assigned
98 (67.1%) exact matches, in which:
68 unique CDR3
13 unique J



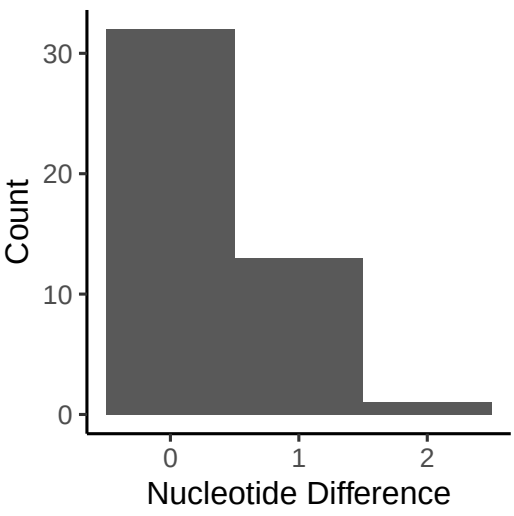
TRBV9*bp02

6 sequences assigned
6 (100%) exact matches, in which:
4 unique CDR3
3 unique J



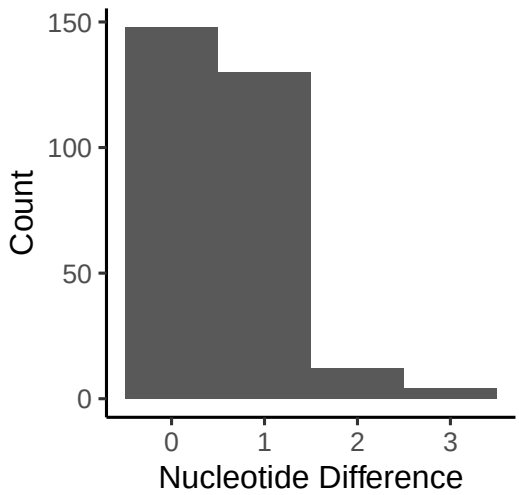
TRBV2*bp01

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



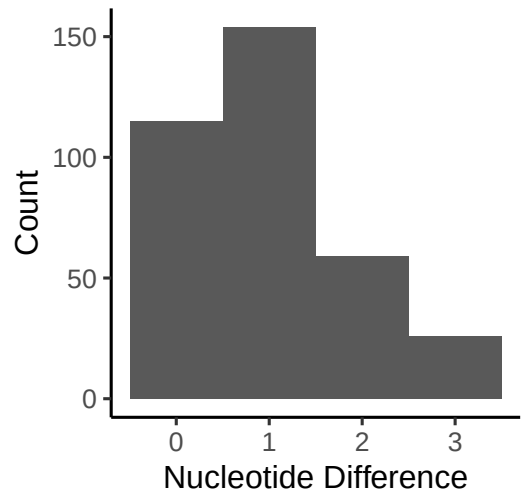
TRBV30*bp01

294 sequences assigned
148 (50.3%) exact matches, in which:
72 unique CDR3
11 unique J



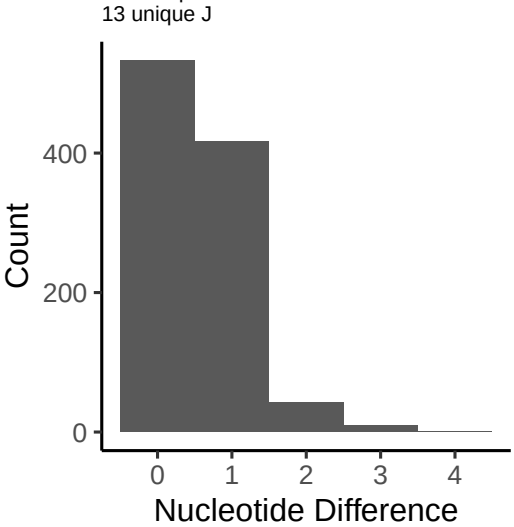
TRBV11−1*bp01

354 sequences assigned
115 (32.5%) exact matches, in which:
80 unique CDR3
10 unique J



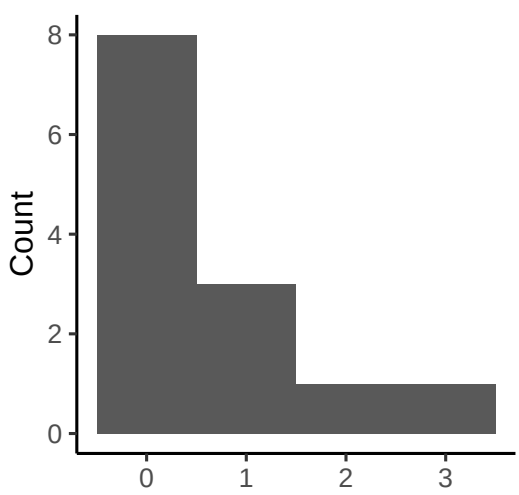
TRBV27*bp01

1003 sequences assigned
533 (53.1%) exact matches, in which:
346 unique CDR3
13 unique J



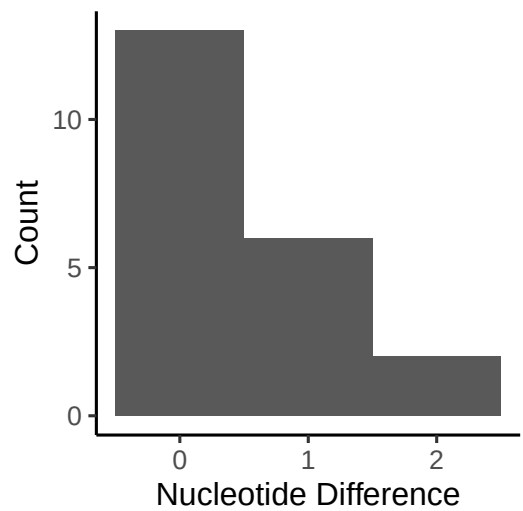
TRBV9*bp01

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



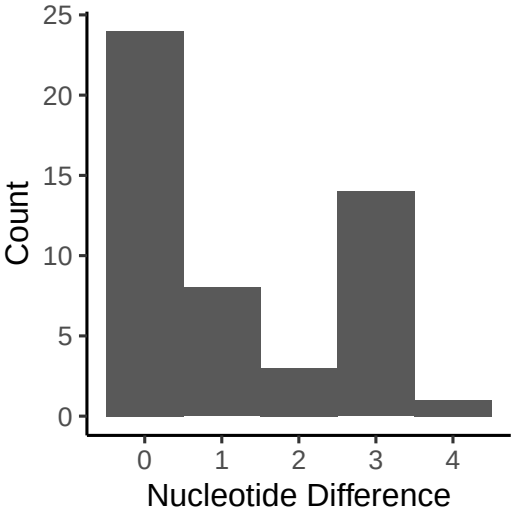
TRBV21−1*bp01

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



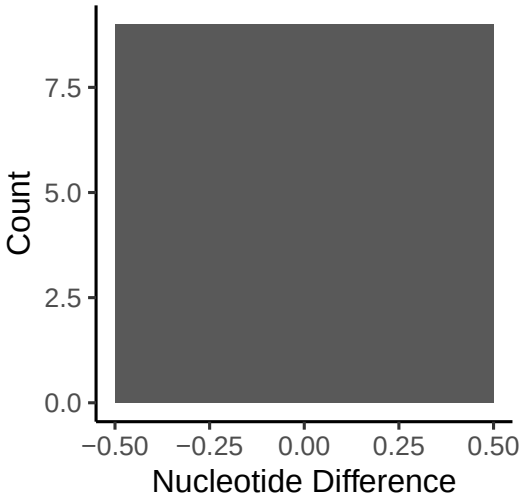
TRBV11−2*bp01

50 sequences assigned
24 (48%) exact matches, in which:
14 unique CDR3
7 unique J



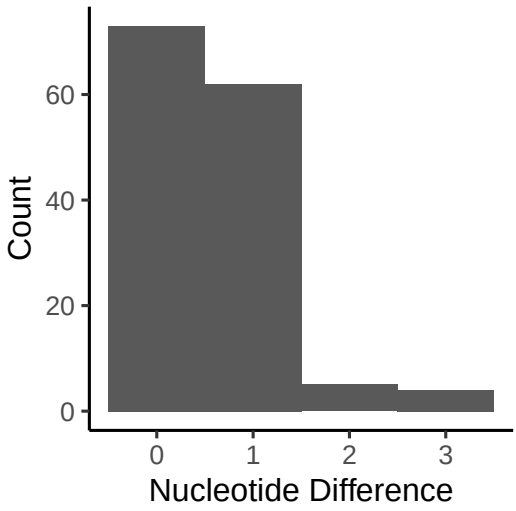
TRBV12−5*bp01

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



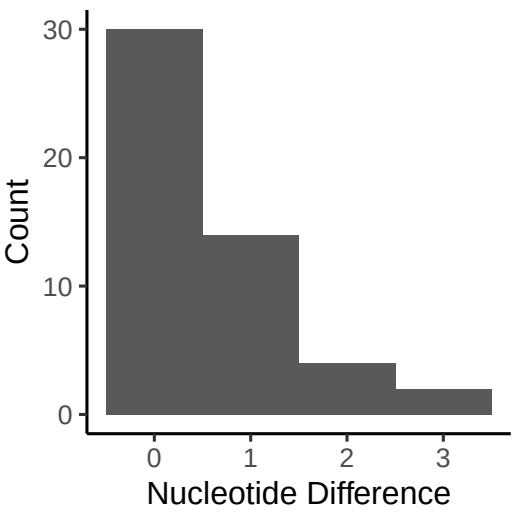
TRBV24−1*bp01

144 sequences assigned
73 (50.7%) exact matches, in which:
46 unique CDR3
6 unique J



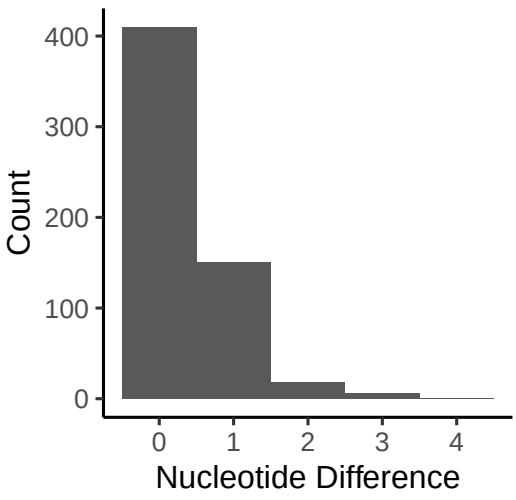
TRBV11−2*bp03

50 sequences assigned
30 (60%) exact matches, in which:
18 unique CDR3
7 unique J



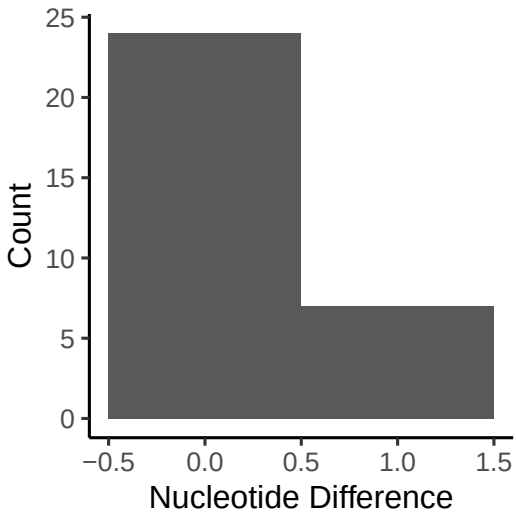
TRBV12−34*bp01

586 sequences assigned
410 (70%) exact matches, in which:
189 unique CDR3
12 unique J



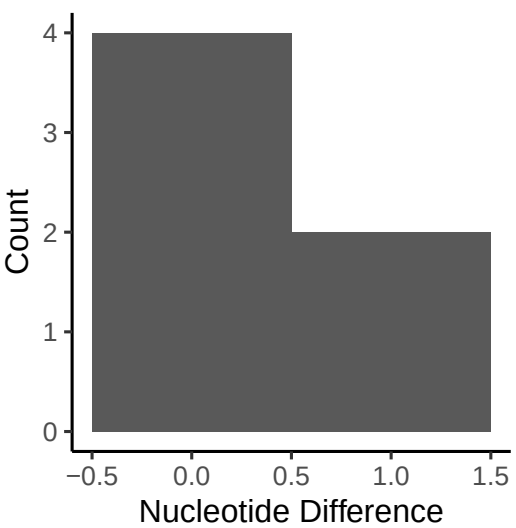
TRBV4−1*bp01

31 sequences assigned
24 (77.4%) exact matches, in which:
15 unique CDR3
7 unique J



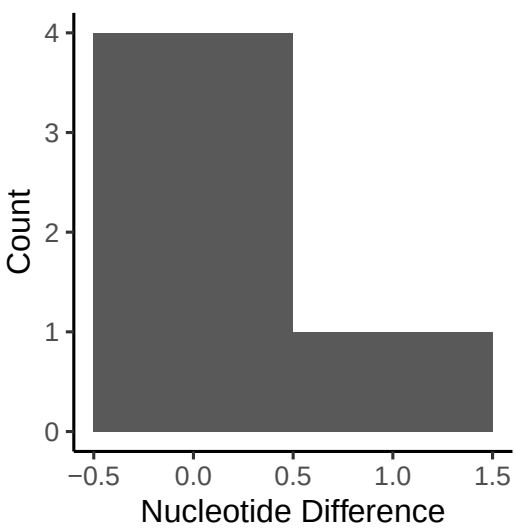
TRBV11−3*bp01

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



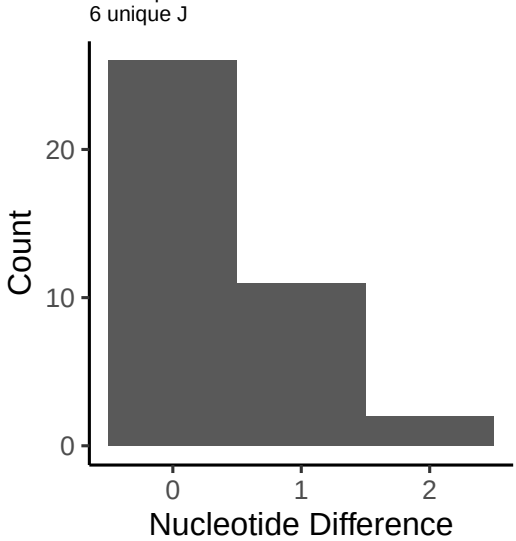
TRBV23−1*bp01

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



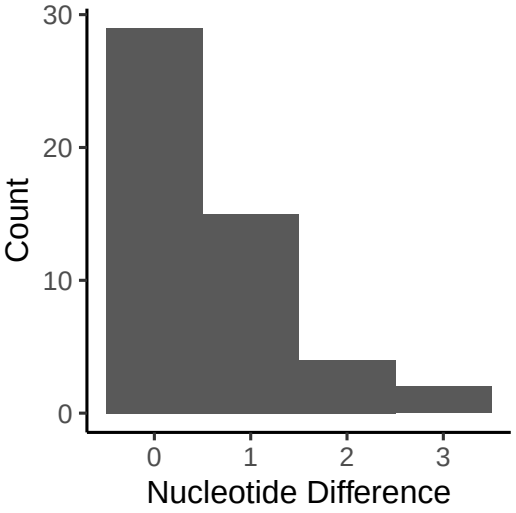
TRBV4−2*bp01

39 sequences assigned
26 (66.7%) exact matches, in which:
21 unique CDR3
6 unique J



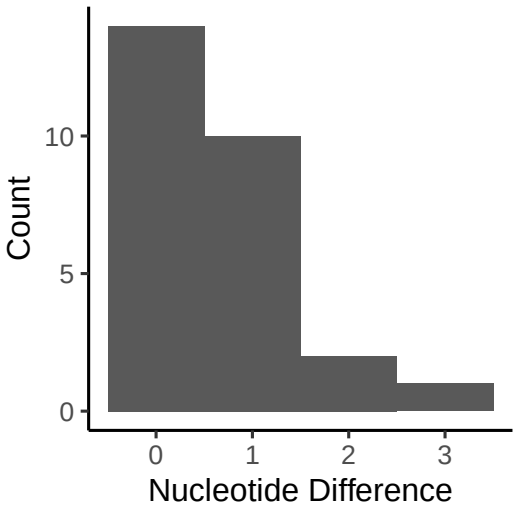
TRBV4−3*bp01

50 sequences assigned
29 (58%) exact matches, in which:
19 unique CDR3
9 unique J



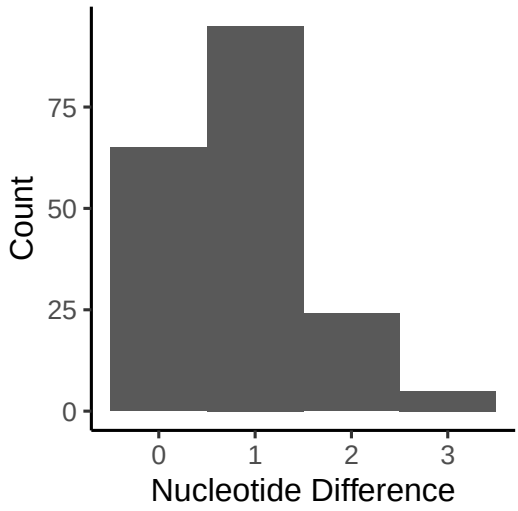
TRBV5−3*bp01

27 sequences assigned
14 (51.9%) exact matches, in which:
6 unique CDR3
5 unique J



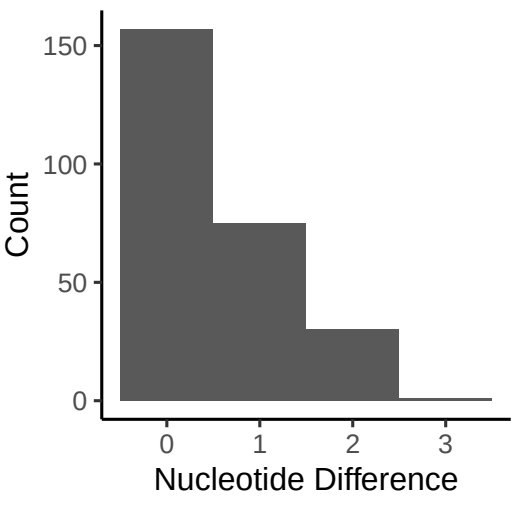
TRBV5−5*bp01

189 sequences assigned
65 (34.4%) exact matches, in which:
41 unique CDR3
11 unique J



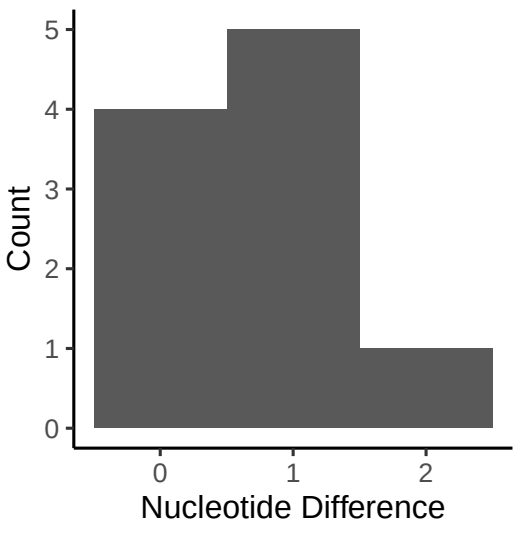
TRBV5−1*bp01

263 sequences assigned
157 (59.7%) exact matches, in which:
84 unique CDR3
12 unique J



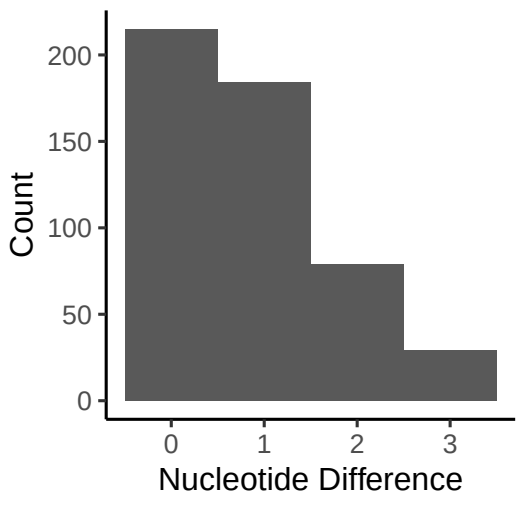
TRBV5−3*bp02

10 sequences assigned
4 (40%) exact matches, in which:
2 unique CDR3
2 unique J



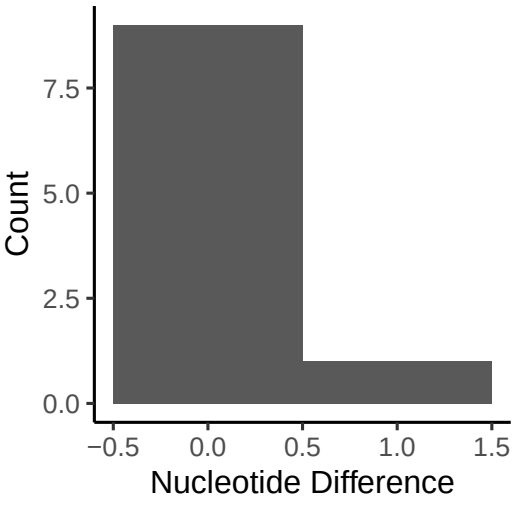
TRBV5−6*bp01

507 sequences assigned
215 (42.4%) exact matches, in which:
116 unique CDR3
12 unique J



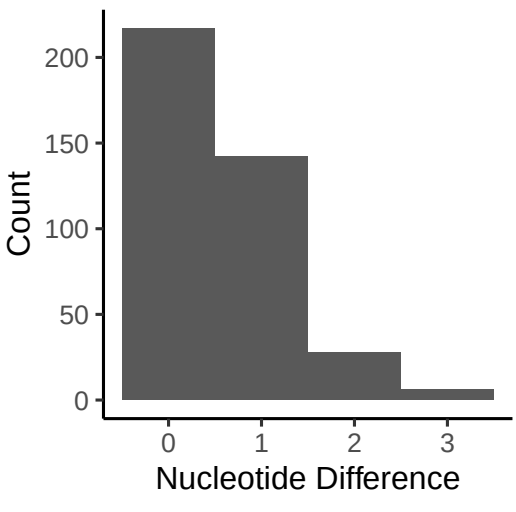
TRBV25−1*bp01

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



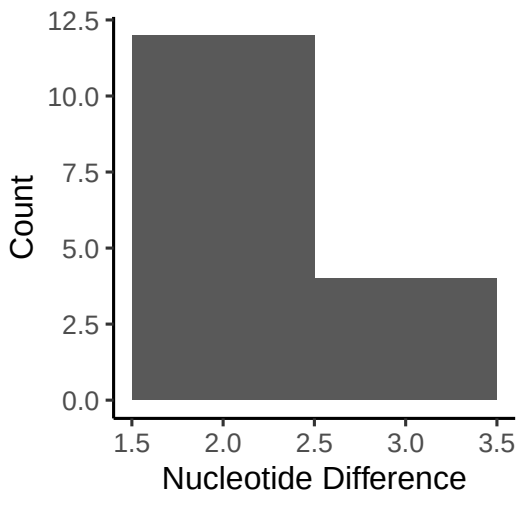
TRBV5−4*bp01

393 sequences assigned
217 (55.2%) exact matches, in which:
128 unique CDR3
13 unique J



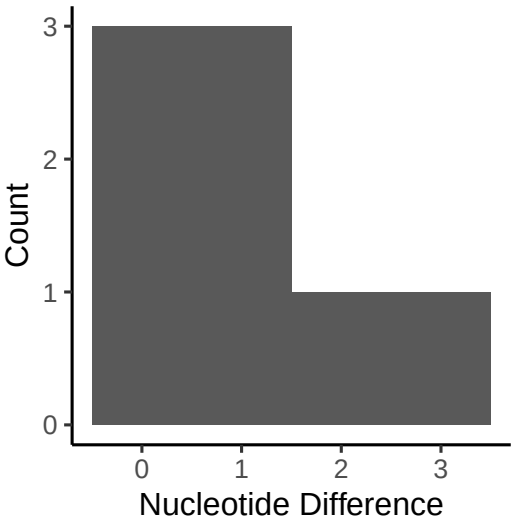
TRBV5−7*bp01

16 sequences assigned
No exact matches.



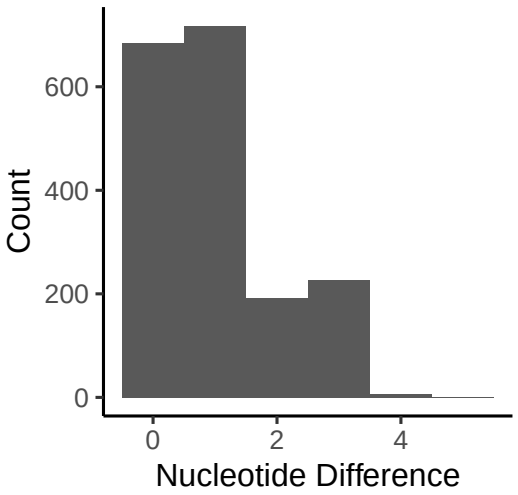
TRBV5−8*bp01

8 sequences assigned
3 (37.5%) exact matches, in which:
1 unique CDR3
1 unique J



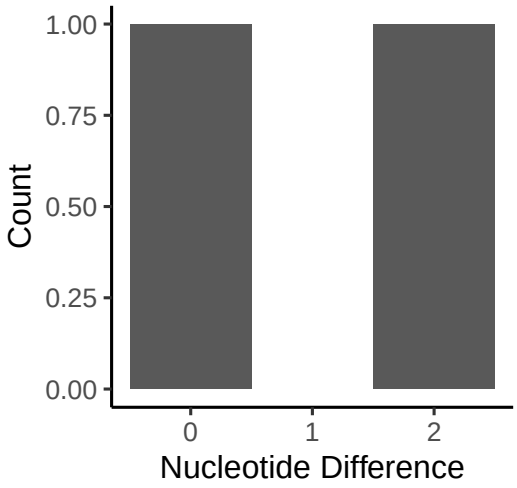
TRBV6−5*bp01

1828 sequences assigned
684 (37.4%) exact matches, in which:
407 unique CDR3
13 unique J



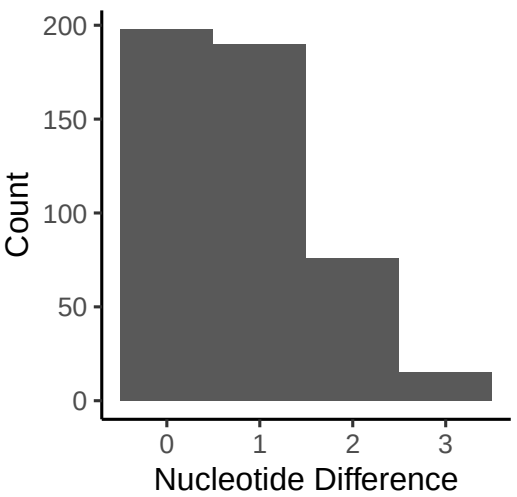
TRBV6−7*bp01

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



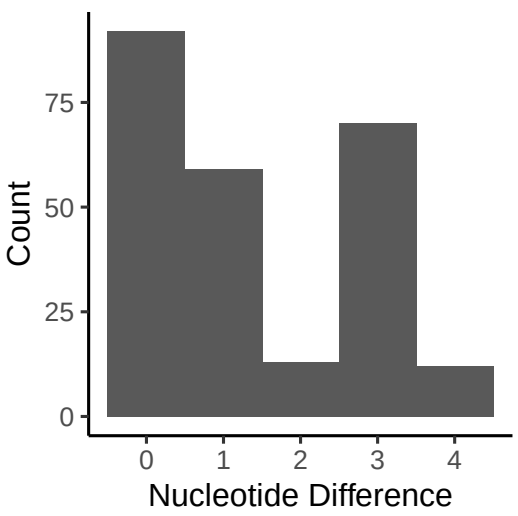
TRBV6−1*bp01

479 sequences assigned
198 (41.3%) exact matches, in which:
146 unique CDR3
13 unique J



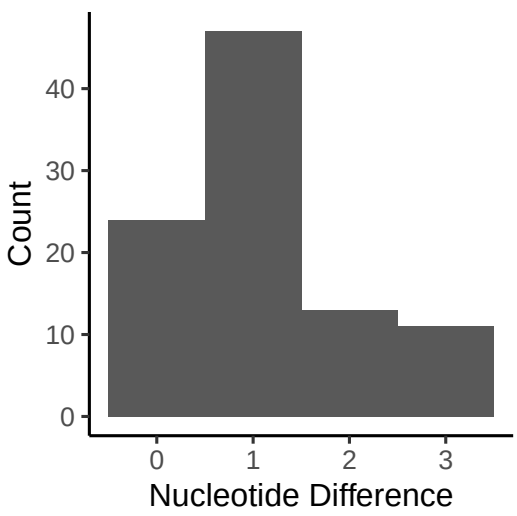
TRBV6−6*bp01

246 sequences assigned
92 (37.4%) exact matches, in which:
72 unique CDR3
12 unique J



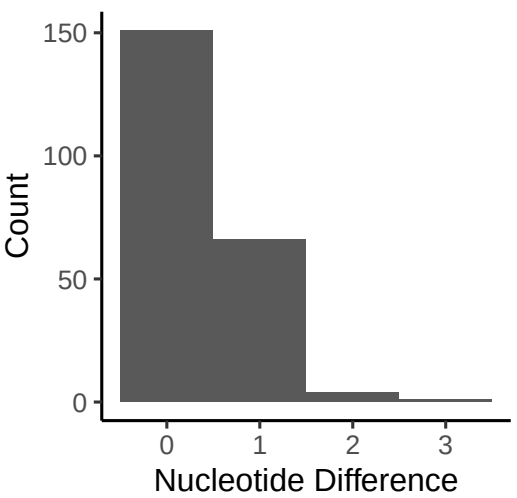
TRBV6−9*bp01

95 sequences assigned
24 (25.3%) exact matches, in which:
15 unique CDR3
6 unique J



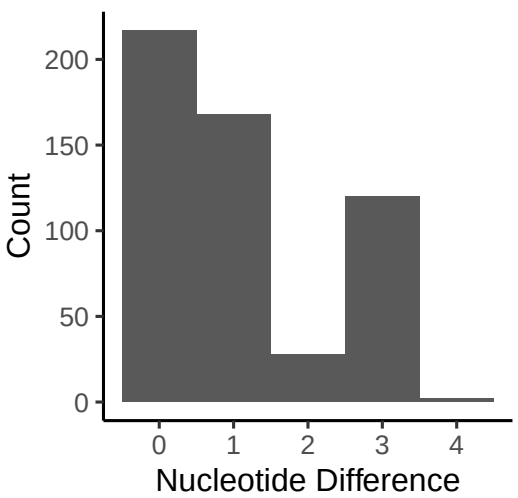
TRBV6−4*bp01

222 sequences assigned
151 (68%) exact matches, in which:
90 unique CDR3
12 unique J



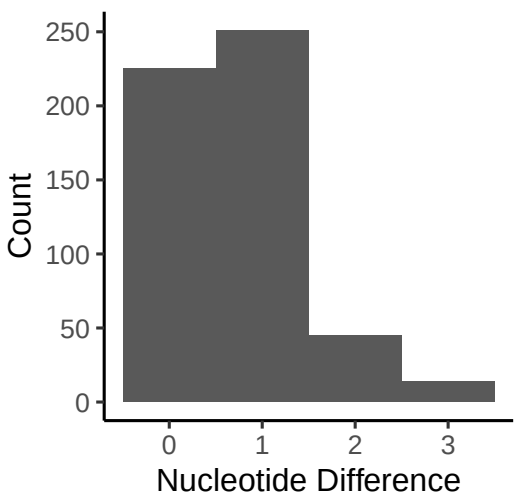
TRBV6−6*bp02

535 sequences assigned
217 (40.6%) exact matches, in which:
152 unique CDR3
13 unique J



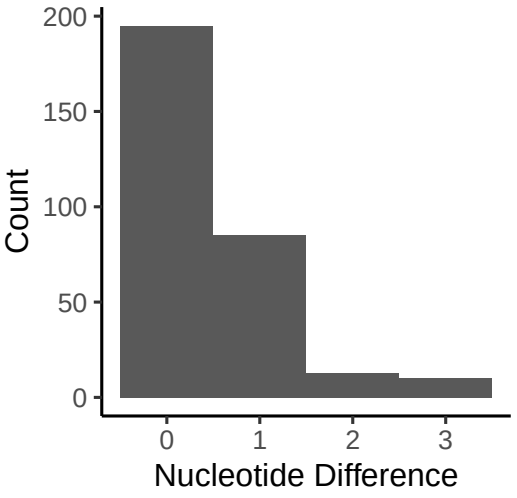
TRBV6−23*bp01

535 sequences assigned
225 (42.1%) exact matches, in which:
149 unique CDR3
13 unique J



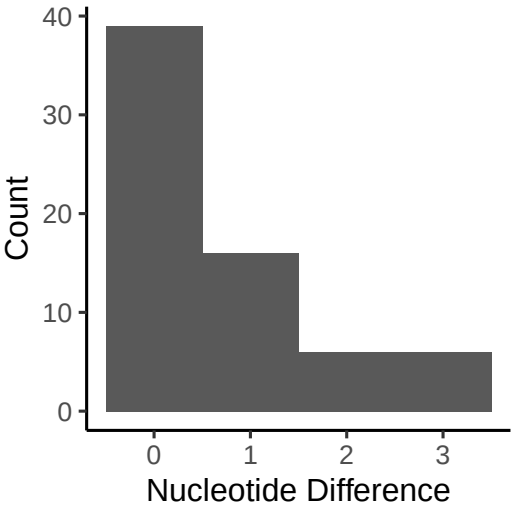
TRBV7−2*bp01

303 sequences assigned
195 (64.4%) exact matches, in which:
90 unique CDR3
12 unique J



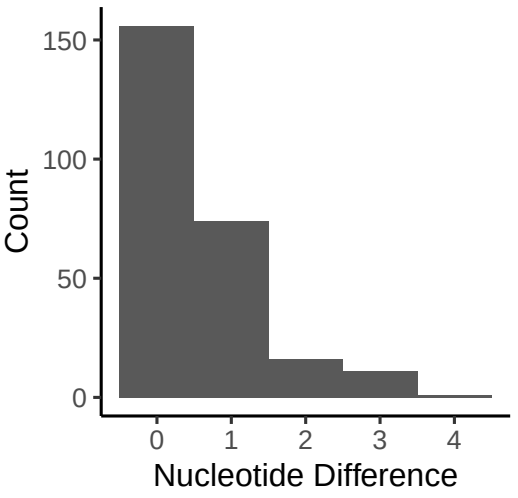
TRBV7−4*bp01

67 sequences assigned
39 (58.2%) exact matches, in which:
22 unique CDR3
5 unique J



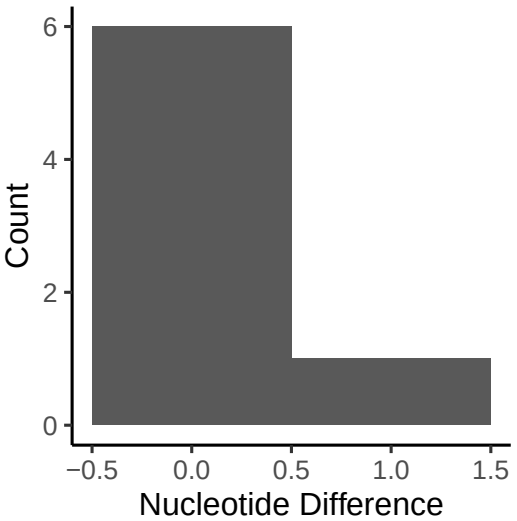
TRBV7−9*bp01

258 sequences assigned
156 (60.5%) exact matches, in which:
72 unique CDR3
9 unique J



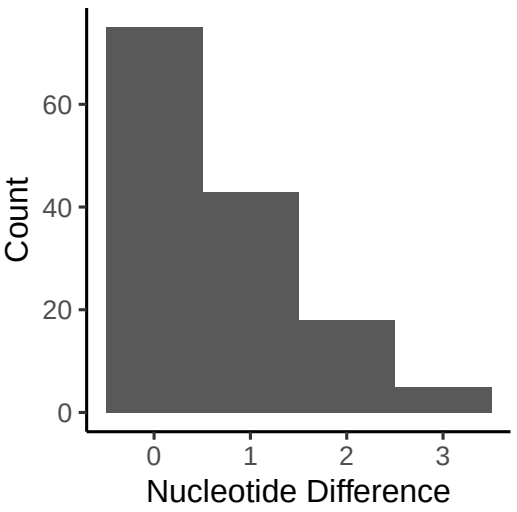
TRBV7−3*bp01

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



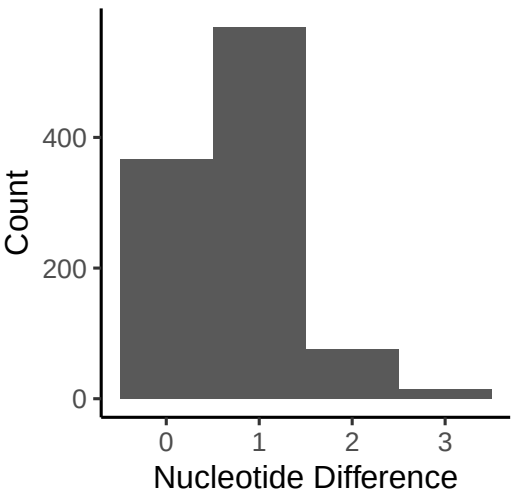
TRBV7−6*bp01

141 sequences assigned
75 (53.2%) exact matches, in which:
33 unique CDR3
6 unique J



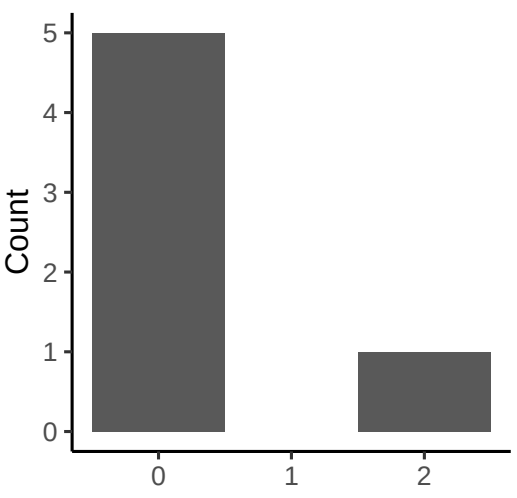
TRBV29−1*bp01

1026 sequences assigned
367 (35.8%) exact matches, in which:
210 unique CDR3
11 unique J



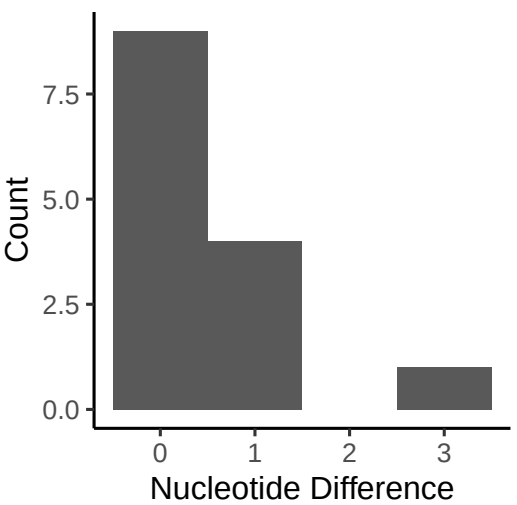
TRBV7−3*bp03

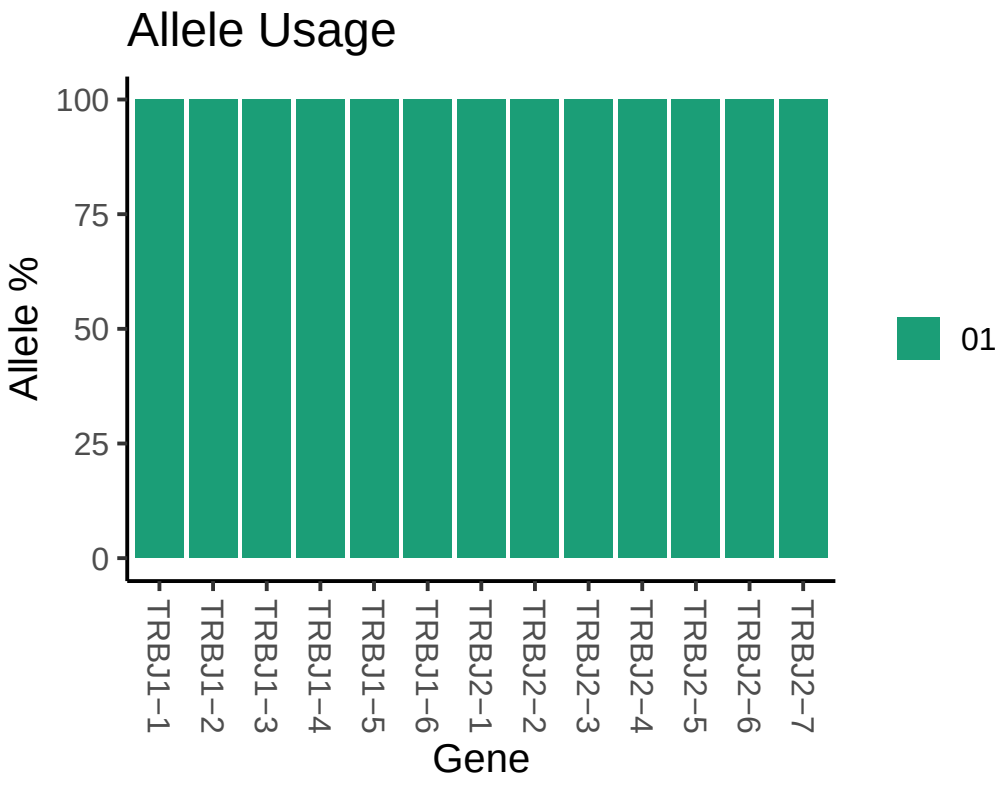
6 sequences assigned
5 (83.3%) exact matches, in which:
4 unique CDR3
4 unique J



TRBV7−7*bp01

14 sequences assigned
9 (64.3%) exact matches, in which:
3 unique CDR3
1 unique J




Allele Usage
Allele %
100
75
50
25
0
TRBJ1−1
TRBJ1−2
TRBJ1−3
TRBJ1−4
TRBJ1−5
TRBJ1−6
TRBJ2−1
TRBJ2−2
TRBJ2−3
TRBJ2−4
TRBJ2−5
TRBJ2−6
TRBJ2−7
Gene
01

Warning − no inferred sequences found.

Warning − no inferred sequences found.

Warning − no inferred sequences found.