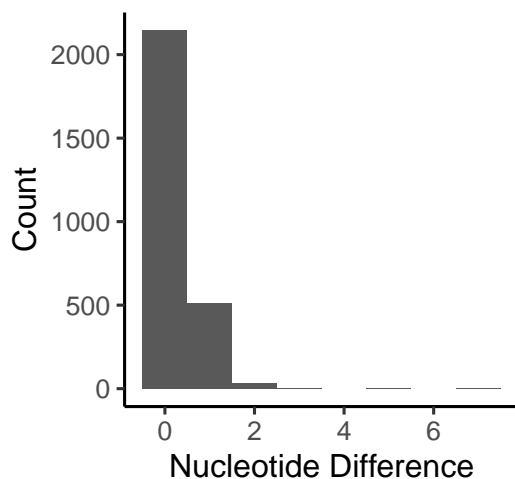


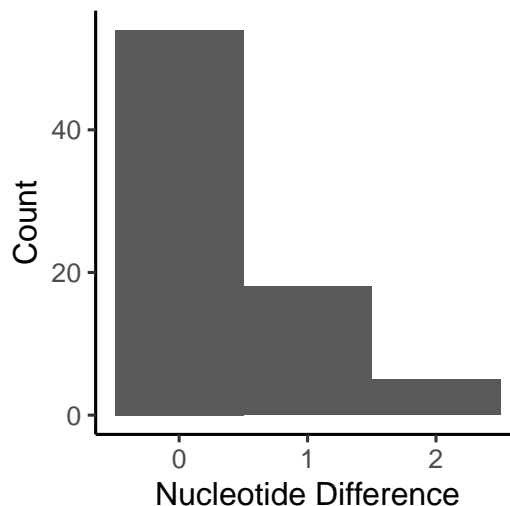
### TRBV20-1\*bp01

2700 sequences assigned  
2145 (79.4%) exact matches, in which:  
1215 unique CDR3  
14 unique J



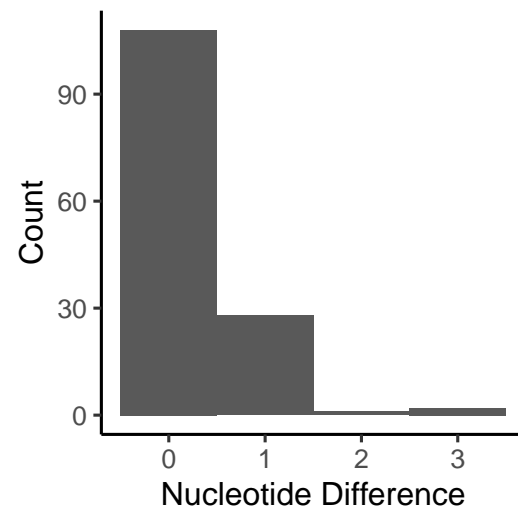
### TRBV10-2\*bp01

77 sequences assigned  
54 (70.1%) exact matches, in which:  
46 unique CDR3  
11 unique J



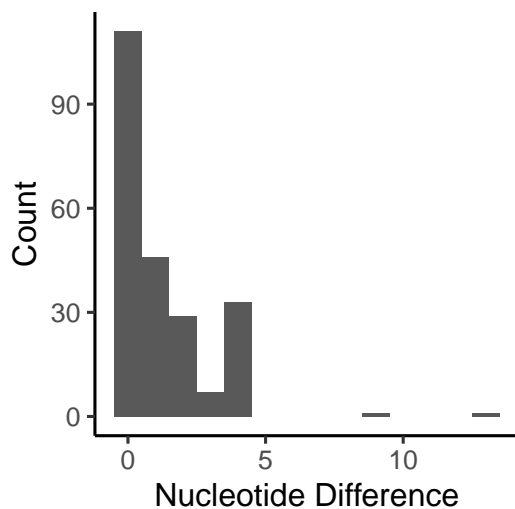
### TRBV13\*bp01

139 sequences assigned  
108 (77.7%) exact matches, in which:  
98 unique CDR3  
14 unique J



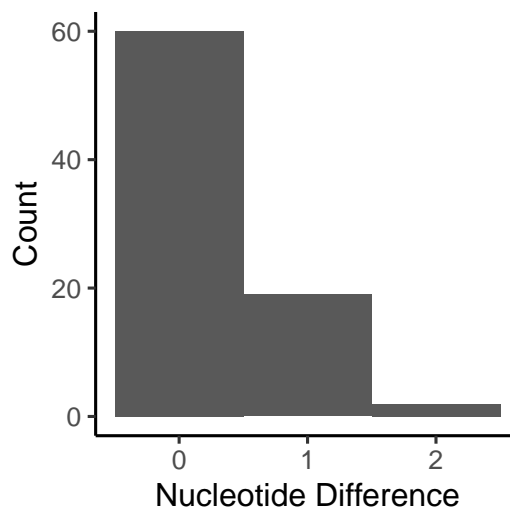
### TRBV10-1\*bp01

228 sequences assigned  
111 (48.7%) exact matches, in which:  
91 unique CDR3  
13 unique J



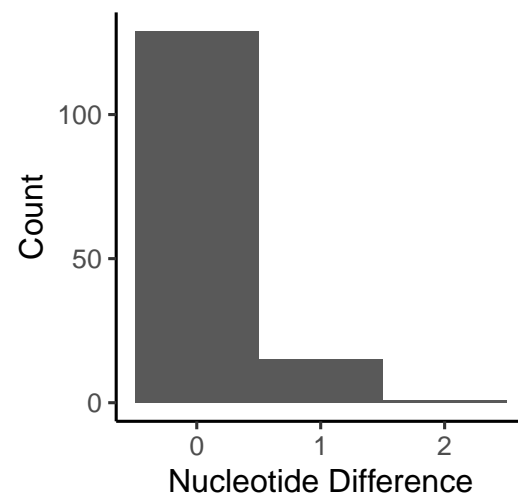
### TRBV10-2\*bp02

81 sequences assigned  
60 (74.1%) exact matches, in which:  
50 unique CDR3  
14 unique J



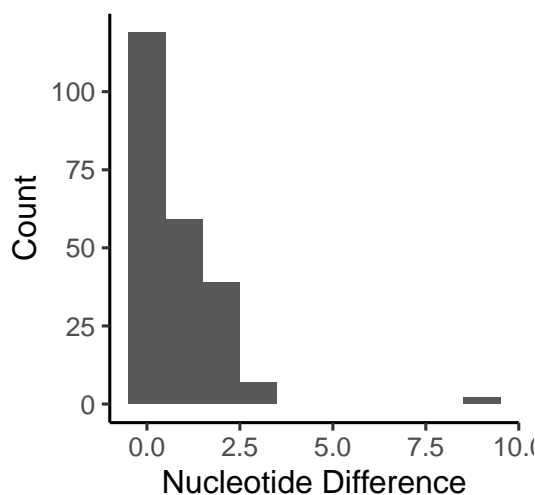
### TRBV14\*bp01

145 sequences assigned  
129 (89%) exact matches, in which:  
108 unique CDR3  
13 unique J



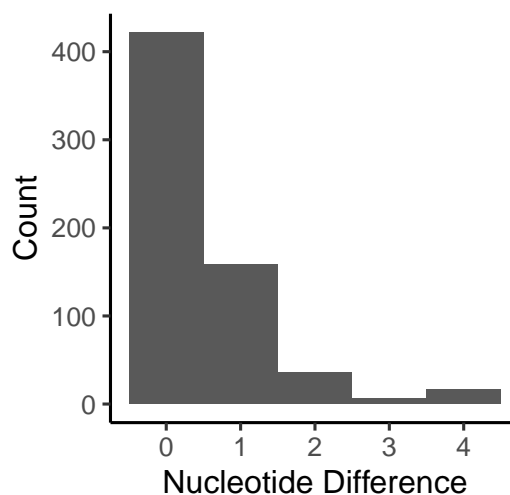
### TRBV10-1\*bp02

226 sequences assigned  
119 (52.7%) exact matches, in which:  
101 unique CDR3  
13 unique J



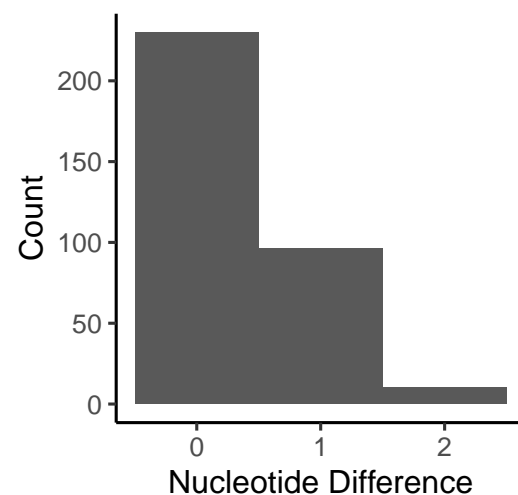
### TRBV10-3\*bp01

638 sequences assigned  
422 (66.1%) exact matches, in which:  
367 unique CDR3  
14 unique J



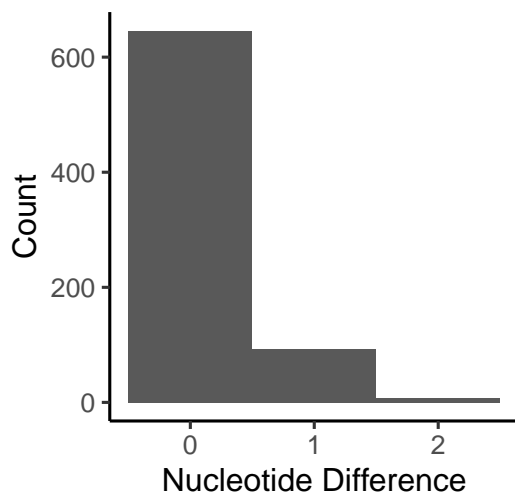
### TRBV15\*bp02

336 sequences assigned  
230 (68.5%) exact matches, in which:  
158 unique CDR3  
14 unique J



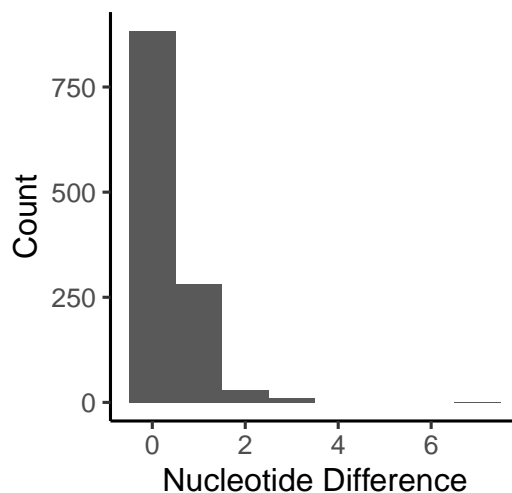
### TRBV18\*bp01

747 sequences assigned  
646 (86.5%) exact matches, in which:  
425 unique CDR3  
14 unique J



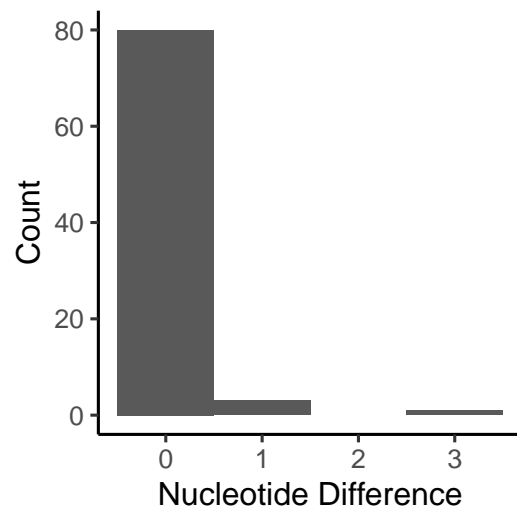
### TRBV27\*bp01

1208 sequences assigned  
884 (73.2%) exact matches, in which:  
759 unique CDR3  
14 unique J



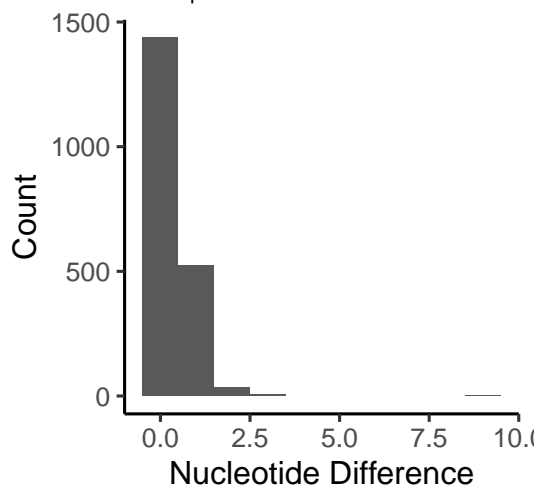
### TRBV9\*bp01

84 sequences assigned  
80 (95.2%) exact matches, in which:  
68 unique CDR3  
12 unique J



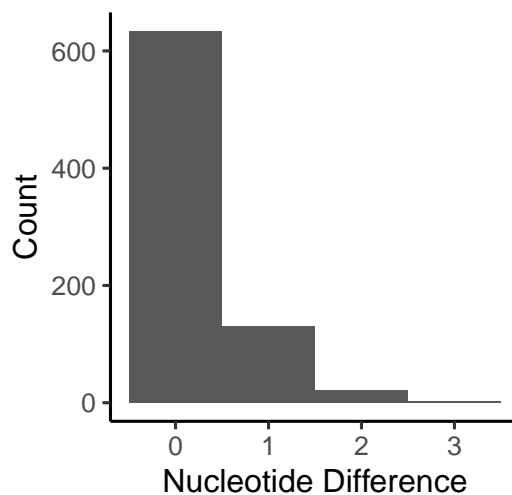
### TRBV19\*bp01

2000 sequences assigned  
1437 (71.8%) exact matches, in which:  
1015 unique CDR3  
14 unique J



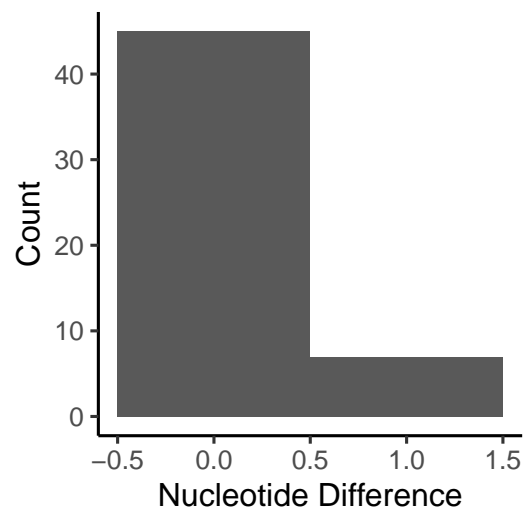
### TRBV28\*bp01

789 sequences assigned  
634 (80.4%) exact matches, in which:  
556 unique CDR3  
14 unique J



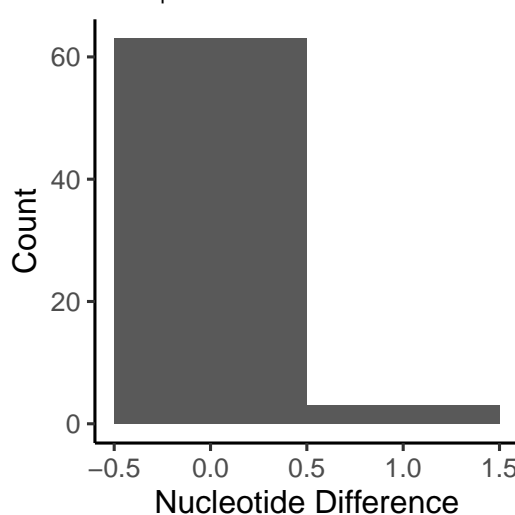
### TRBV9\*bp02

52 sequences assigned  
45 (86.5%) exact matches, in which:  
44 unique CDR3  
11 unique J



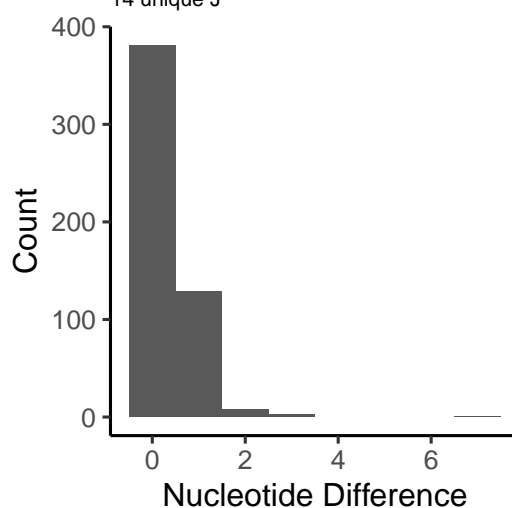
### TRBV2\*bp01

67 sequences assigned  
63 (94%) exact matches, in which:  
62 unique CDR3  
14 unique J



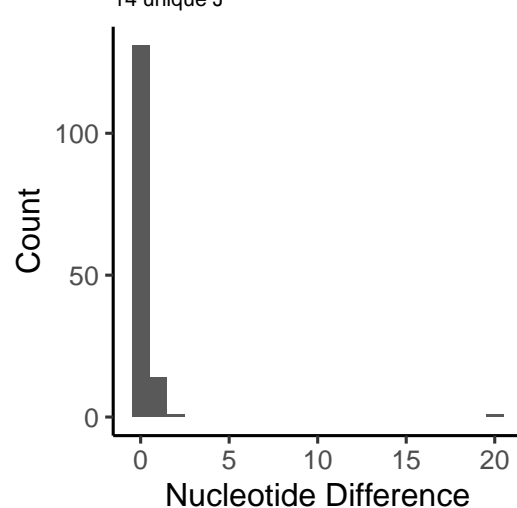
### TRBV30\*bp01

522 sequences assigned  
381 (73%) exact matches, in which:  
217 unique CDR3  
14 unique J



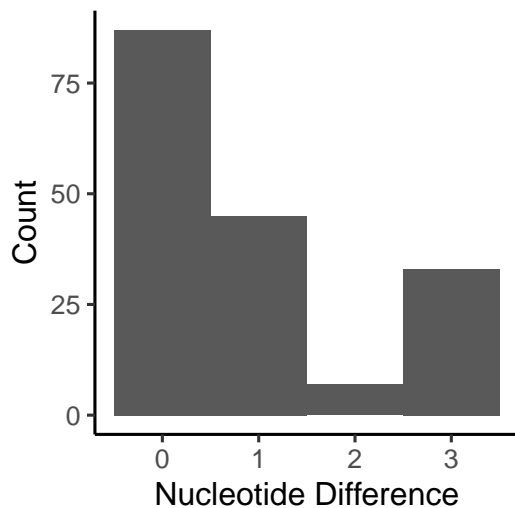
### TRBV21-1\*bp01

147 sequences assigned  
131 (89.1%) exact matches, in which:  
119 unique CDR3  
14 unique J



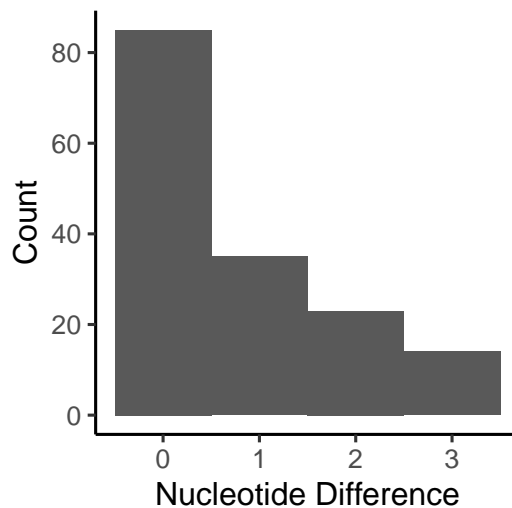
### TRBV11–1\*bp01

172 sequences assigned  
87 (50.6%) exact matches, in which:  
69 unique CDR3  
12 unique J



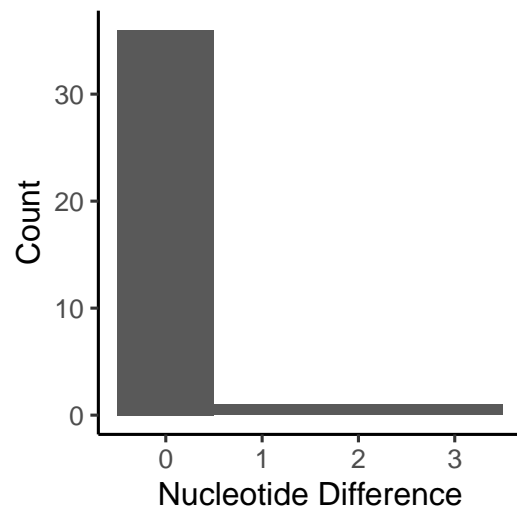
### TRBV11–3\*bp01

157 sequences assigned  
85 (54.1%) exact matches, in which:  
74 unique CDR3  
14 unique J



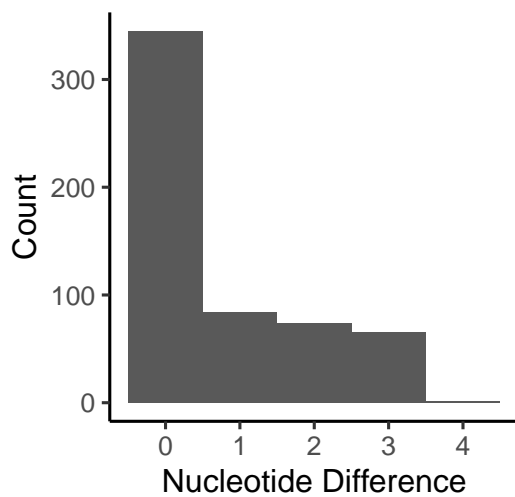
### TRBV23–1\*bp01

39 sequences assigned  
36 (92.3%) exact matches, in which:  
34 unique CDR3  
10 unique J



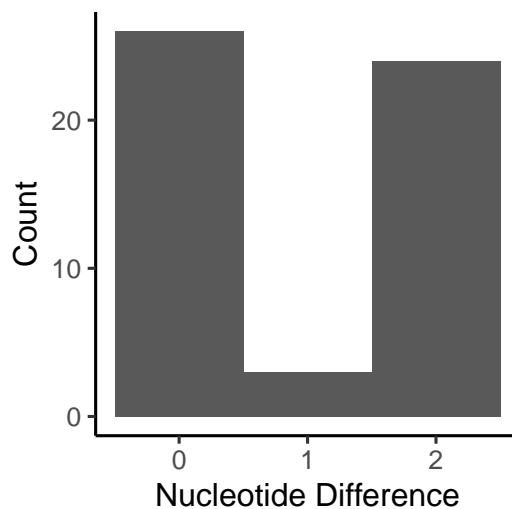
### TRBV11–2\*bp01

569 sequences assigned  
345 (60.6%) exact matches, in which:  
291 unique CDR3  
14 unique J



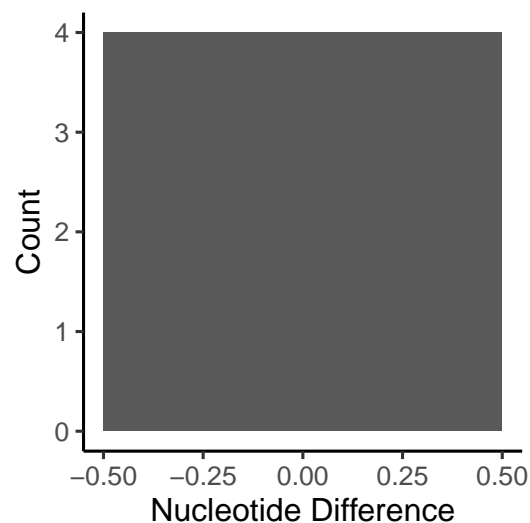
### TRBV12–5\*bp01

53 sequences assigned  
26 (49.1%) exact matches, in which:  
23 unique CDR3  
10 unique J



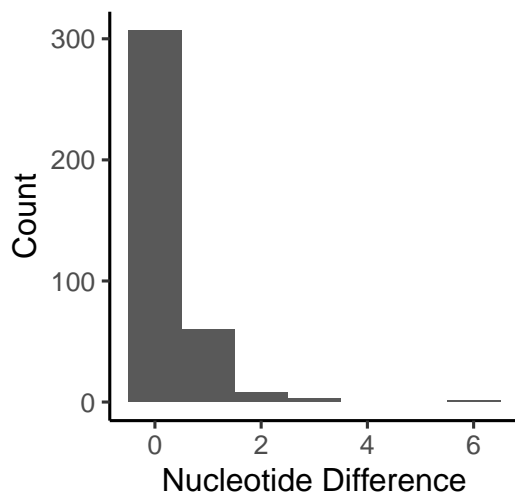
### TRBV3–1\*bp01

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



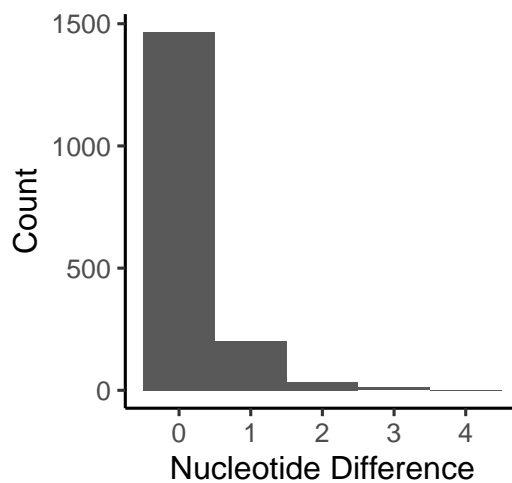
### TRBV11–2\*bp03

379 sequences assigned  
307 (81%) exact matches, in which:  
264 unique CDR3  
14 unique J



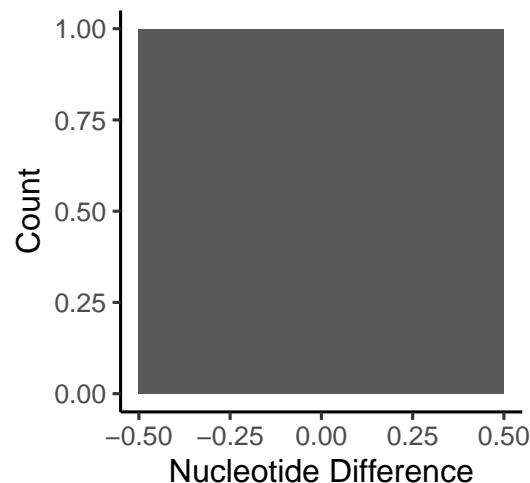
### TRBV12–34\*bp01

1716 sequences assigned  
1466 (85.4%) exact matches, in which:  
920 unique CDR3  
14 unique J



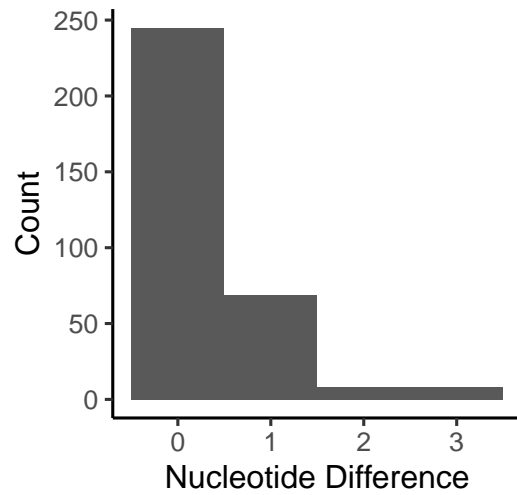
### TRBV3–2\*bp02

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



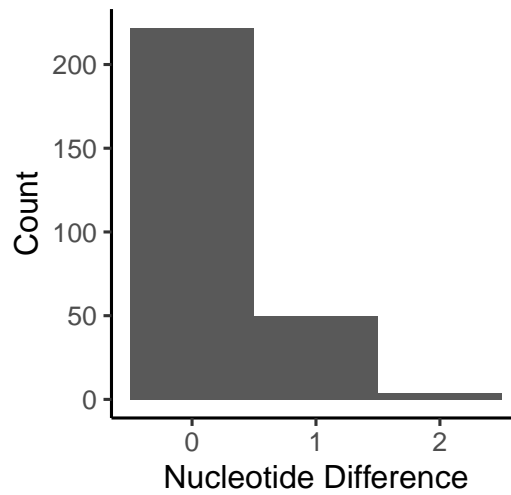
### TRBV24–1\*bp01

330 sequences assigned  
245 (74.2%) exact matches, in which:  
195 unique CDR3  
14 unique J



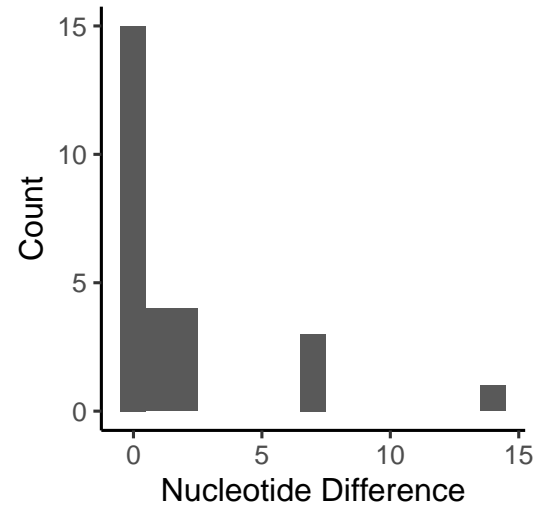
### TRBV4–3\*bp01

276 sequences assigned  
222 (80.4%) exact matches, in which:  
186 unique CDR3  
13 unique J



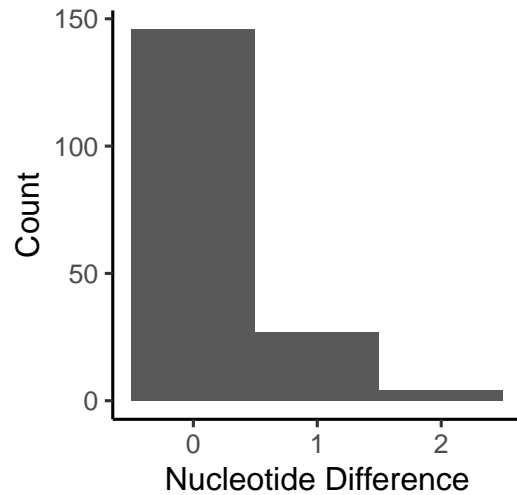
### TRBV5–3\*bp01

27 sequences assigned  
15 (55.6%) exact matches, in which:  
12 unique CDR3  
6 unique J



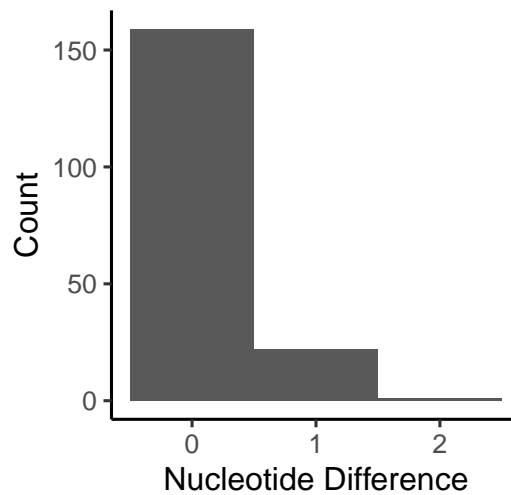
### TRBV4–1\*bp01

177 sequences assigned  
146 (82.5%) exact matches, in which:  
131 unique CDR3  
14 unique J



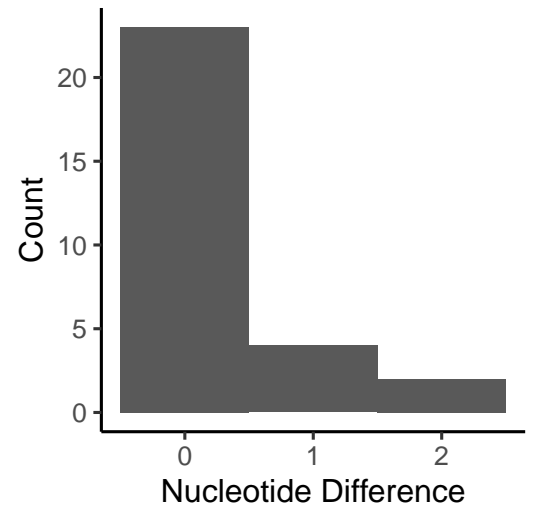
### TRBV25–1\*bp01

182 sequences assigned  
159 (87.4%) exact matches, in which:  
108 unique CDR3  
13 unique J



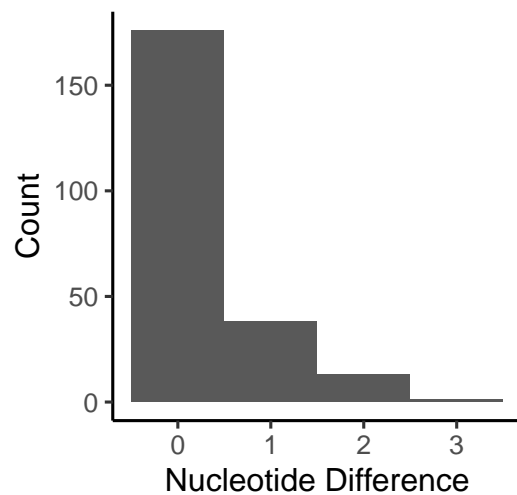
### TRBV5–3\*bp02

29 sequences assigned  
23 (79.3%) exact matches, in which:  
19 unique CDR3  
9 unique J



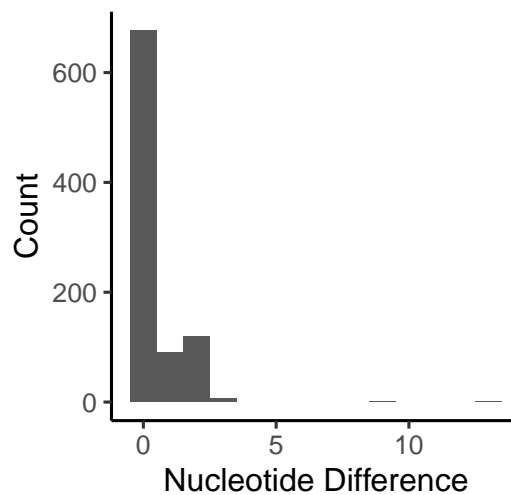
### TRBV4–2\*bp01

228 sequences assigned  
176 (77.2%) exact matches, in which:  
164 unique CDR3  
13 unique J



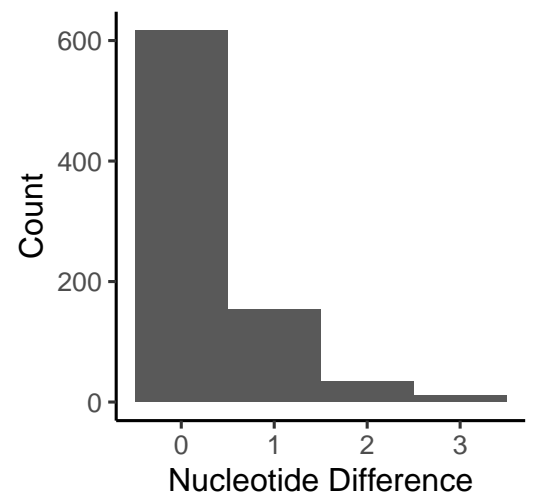
### TRBV5–1\*bp01

897 sequences assigned  
677 (75.5%) exact matches, in which:  
561 unique CDR3  
14 unique J



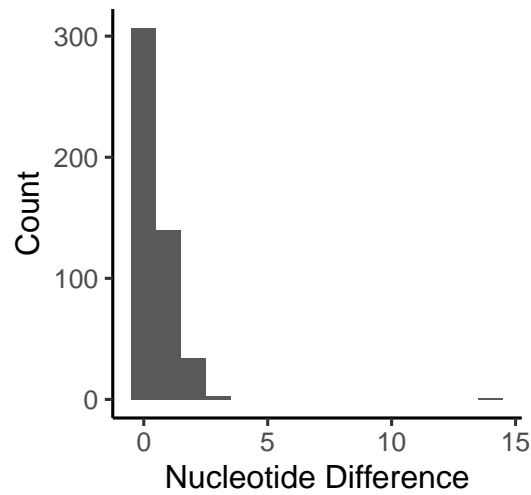
### TRBV5–4\*bp01

816 sequences assigned  
617 (75.6%) exact matches, in which:  
468 unique CDR3  
14 unique J



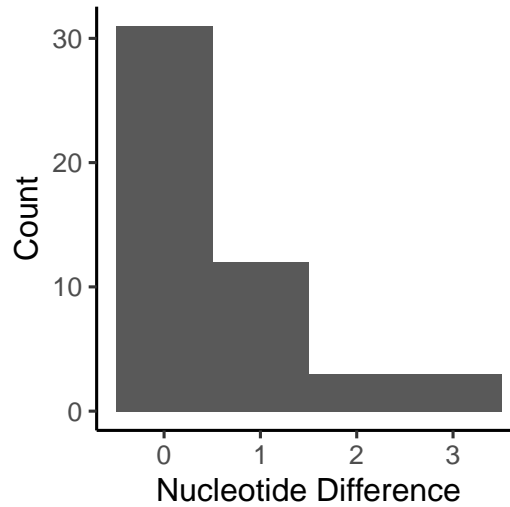
### TRBV5-5\*bp01

485 sequences assigned  
307 (63.3%) exact matches, in which:  
216 unique CDR3  
14 unique J



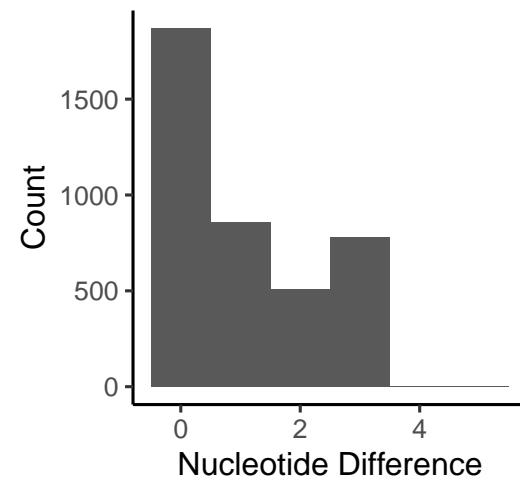
### TRBV5-8\*bp01

49 sequences assigned  
31 (63.3%) exact matches, in which:  
29 unique CDR3  
12 unique J



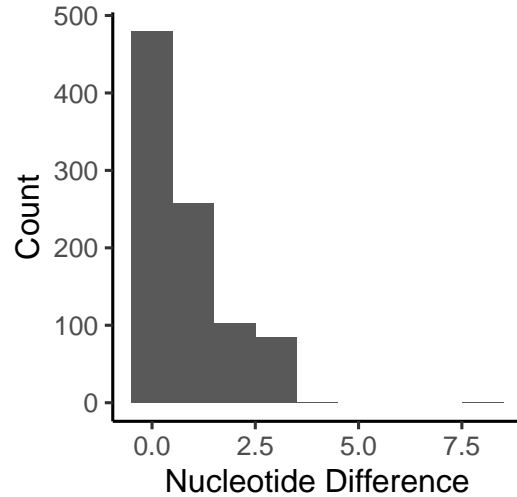
### TRBV6-5\*bp01

4016 sequences assigned  
1869 (46.5%) exact matches, in which:  
1526 unique CDR3  
14 unique J



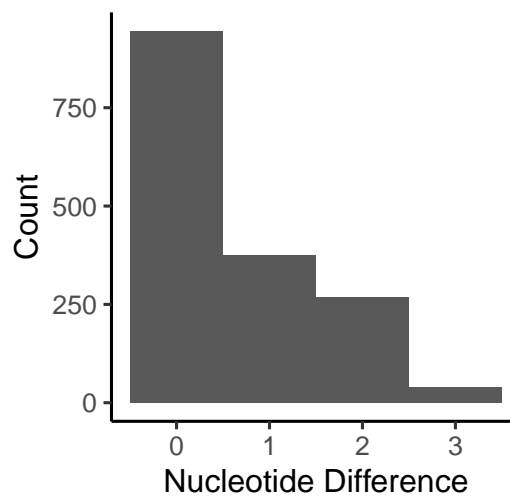
### TRBV5-6\*bp01

926 sequences assigned  
480 (51.8%) exact matches, in which:  
341 unique CDR3  
14 unique J



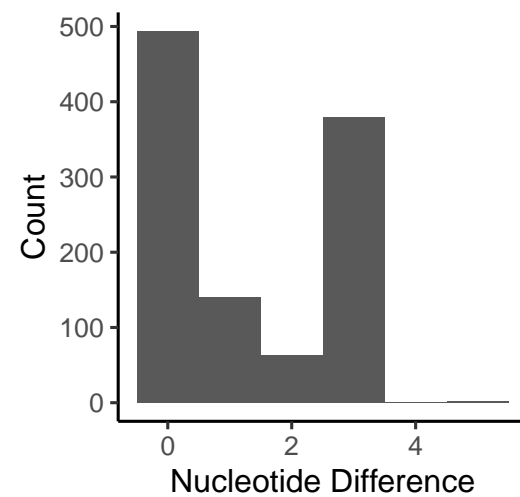
### TRBV6-1\*bp01

1629 sequences assigned  
945 (58%) exact matches, in which:  
820 unique CDR3  
14 unique J



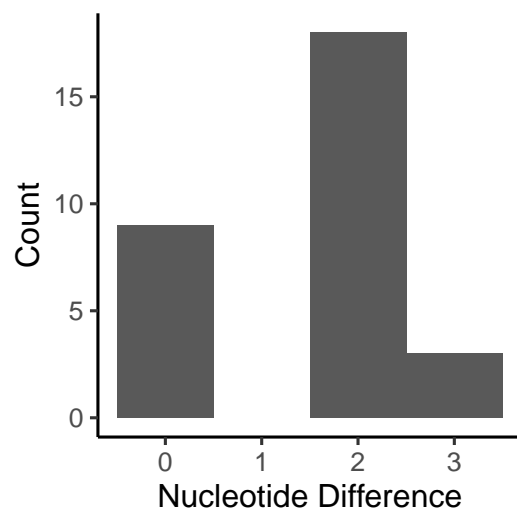
### TRBV6-6\*bp02

1079 sequences assigned  
494 (45.8%) exact matches, in which:  
427 unique CDR3  
14 unique J



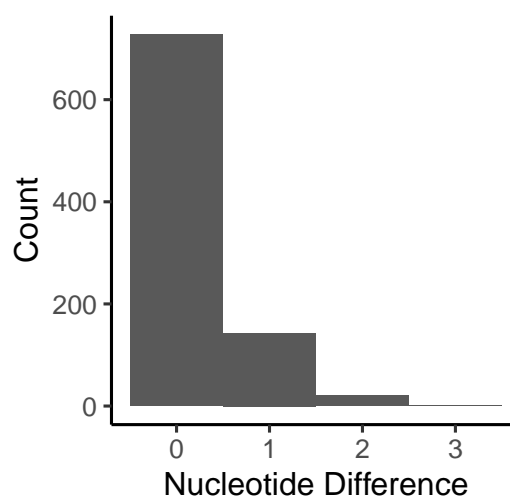
### TRBV5-7\*bp01

30 sequences assigned  
9 (30%) exact matches, in which:  
7 unique CDR3  
6 unique J



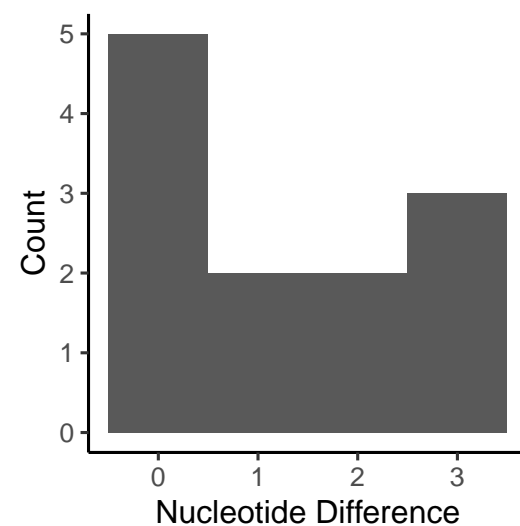
### TRBV6-4\*bp01

893 sequences assigned  
728 (81.5%) exact matches, in which:  
593 unique CDR3  
14 unique J



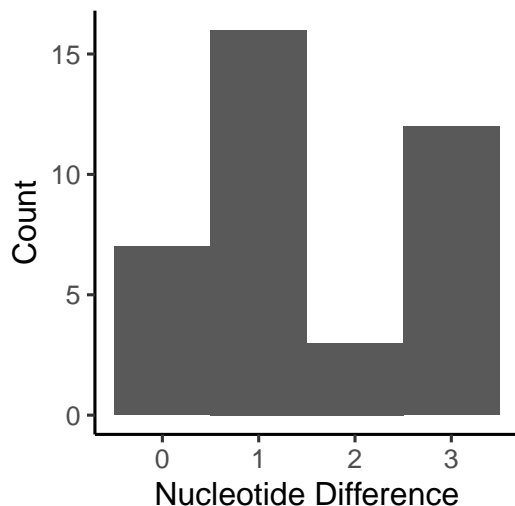
### TRBV6-7\*bp01

12 sequences assigned  
5 (41.7%) exact matches, in which:  
4 unique CDR3  
4 unique J



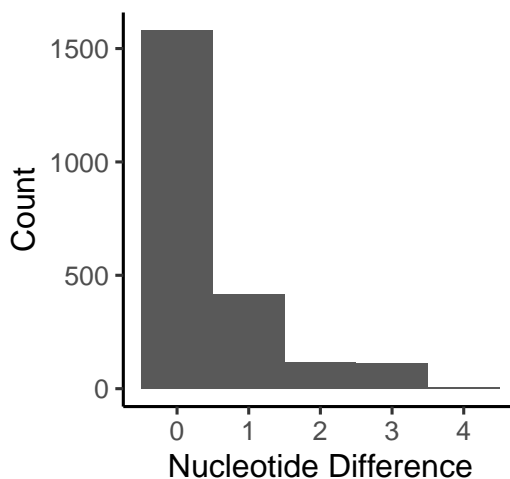
### TRBV6-8\*bp01

38 sequences assigned  
7 (18.4%) exact matches, in which:  
6 unique CDR3  
5 unique J



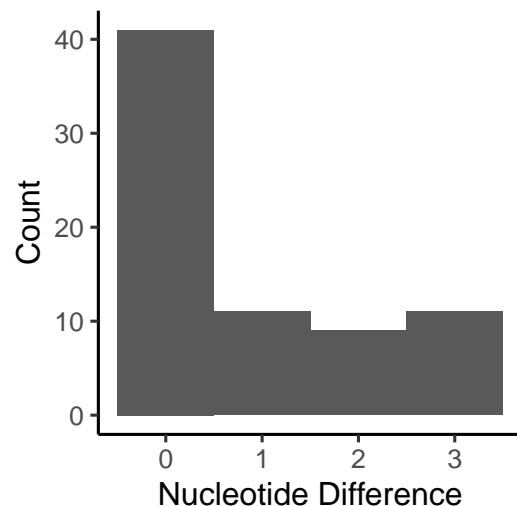
### TRBV7-2\*bp02

2234 sequences assigned  
1580 (70.7%) exact matches, in which:  
1008 unique CDR3  
14 unique J



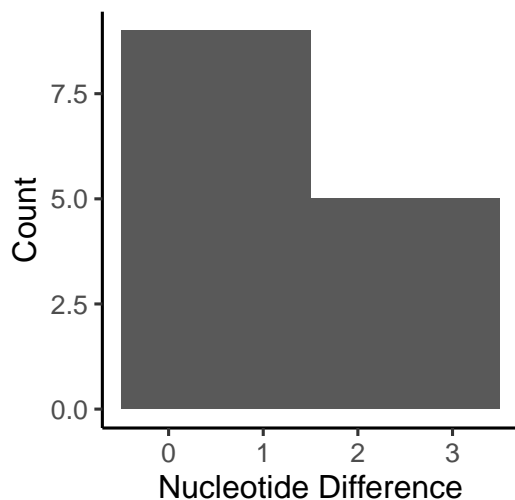
### TRBV7-4\*bp01

72 sequences assigned  
41 (56.9%) exact matches, in which:  
31 unique CDR3  
12 unique J



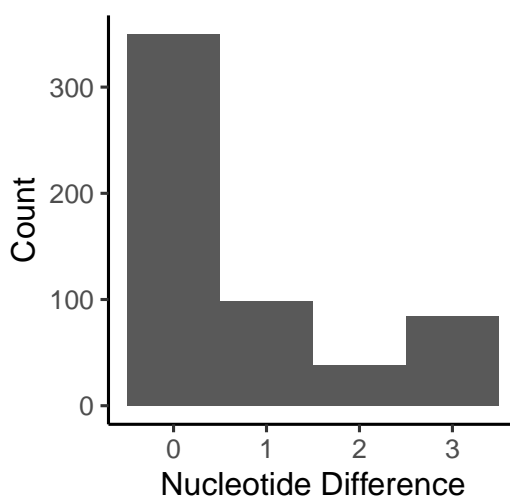
### TRBV6-9\*bp01

28 sequences assigned  
9 (32.1%) exact matches, in which:  
8 unique CDR3  
6 unique J



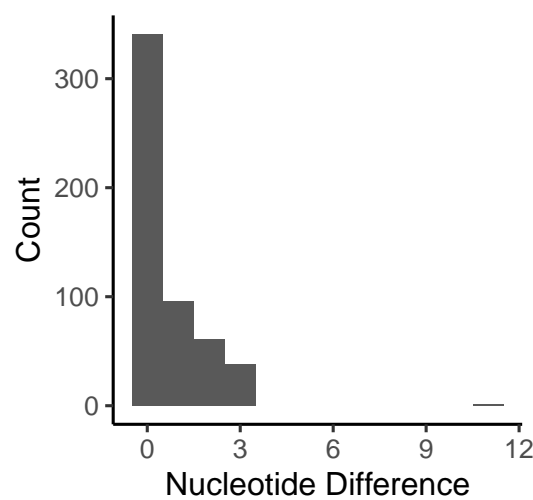
### TRBV7-3\*bp01

570 sequences assigned  
350 (61.4%) exact matches, in which:  
272 unique CDR3  
14 unique J



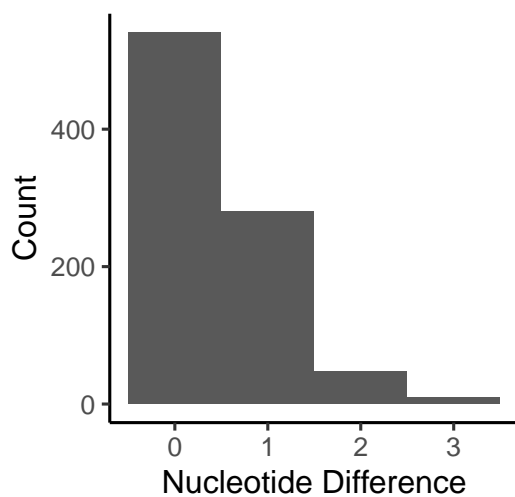
### TRBV7-6\*bp01

537 sequences assigned  
341 (63.5%) exact matches, in which:  
228 unique CDR3  
13 unique J



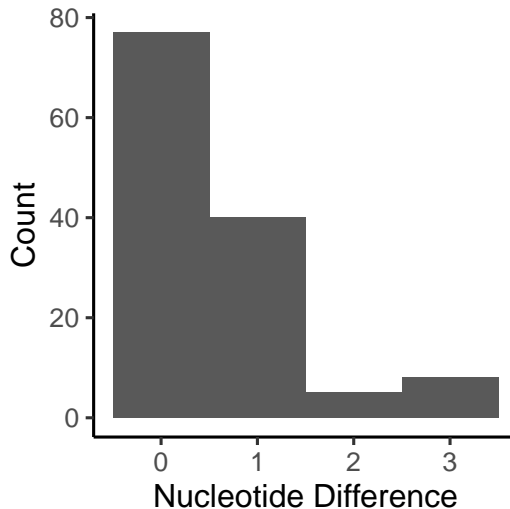
### TRBV6-23\*bp01

877 sequences assigned  
541 (61.7%) exact matches, in which:  
475 unique CDR3  
14 unique J



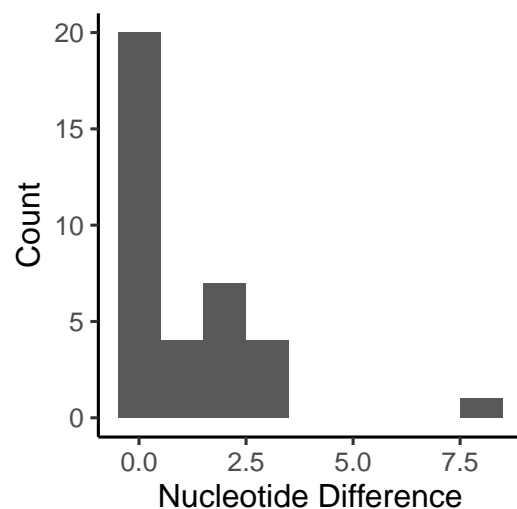
### TRBV7-3\*bp03

130 sequences assigned  
77 (59.2%) exact matches, in which:  
53 unique CDR3  
13 unique J



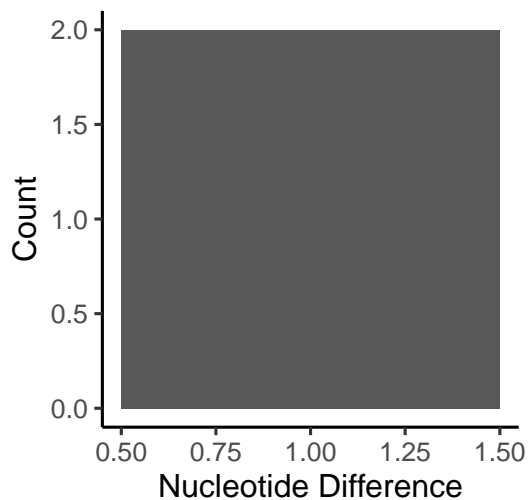
### TRBV7-7\*bp01

36 sequences assigned  
20 (55.6%) exact matches, in which:  
19 unique CDR3  
9 unique J



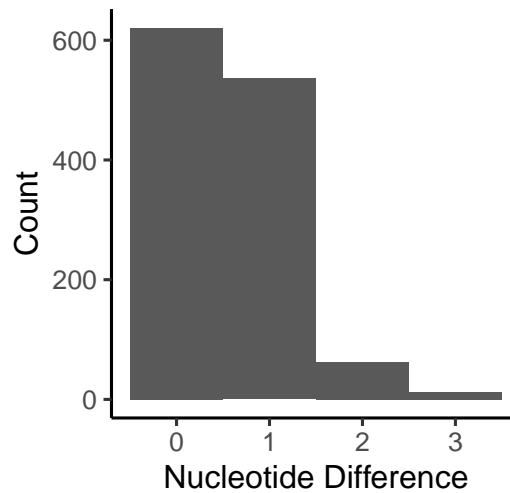
### TRBV7–8\*bp01

2 sequences assigned  
No exact matches.



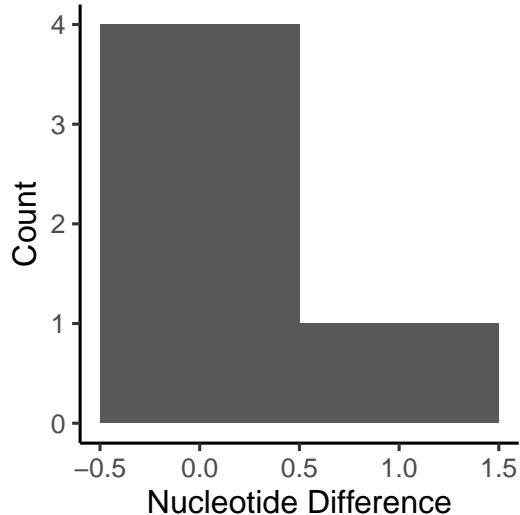
### TRBV29–1\*bp01

1232 sequences assigned  
621 (50.4%) exact matches, in which:  
460 unique CDR3  
14 unique J



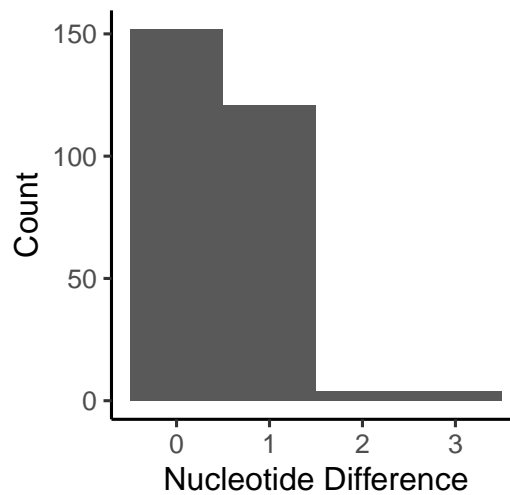
### TRBV7–8\*bp02

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



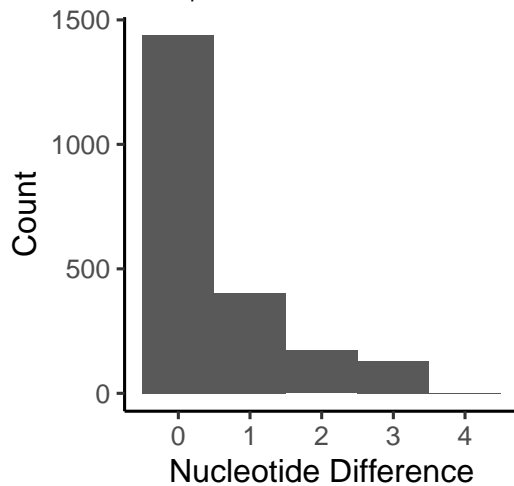
### TRBV29–1\*bp03

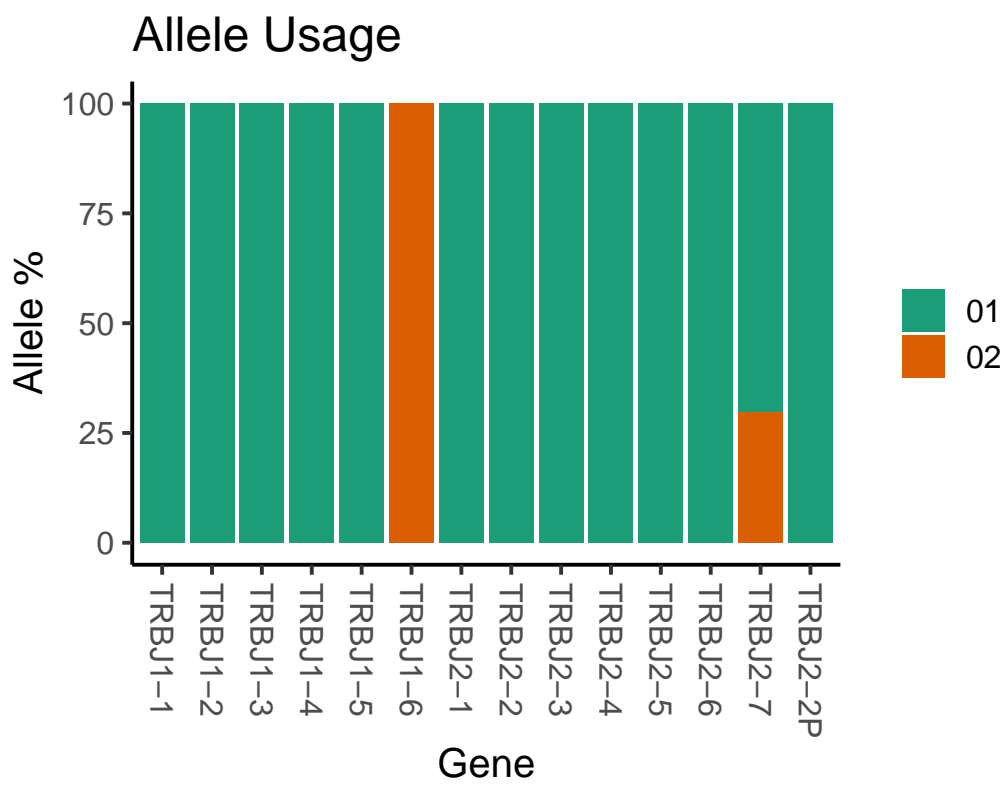
281 sequences assigned  
152 (54.1%) exact matches, in which:  
119 unique CDR3  
12 unique J



### TRBV7–9\*bp01

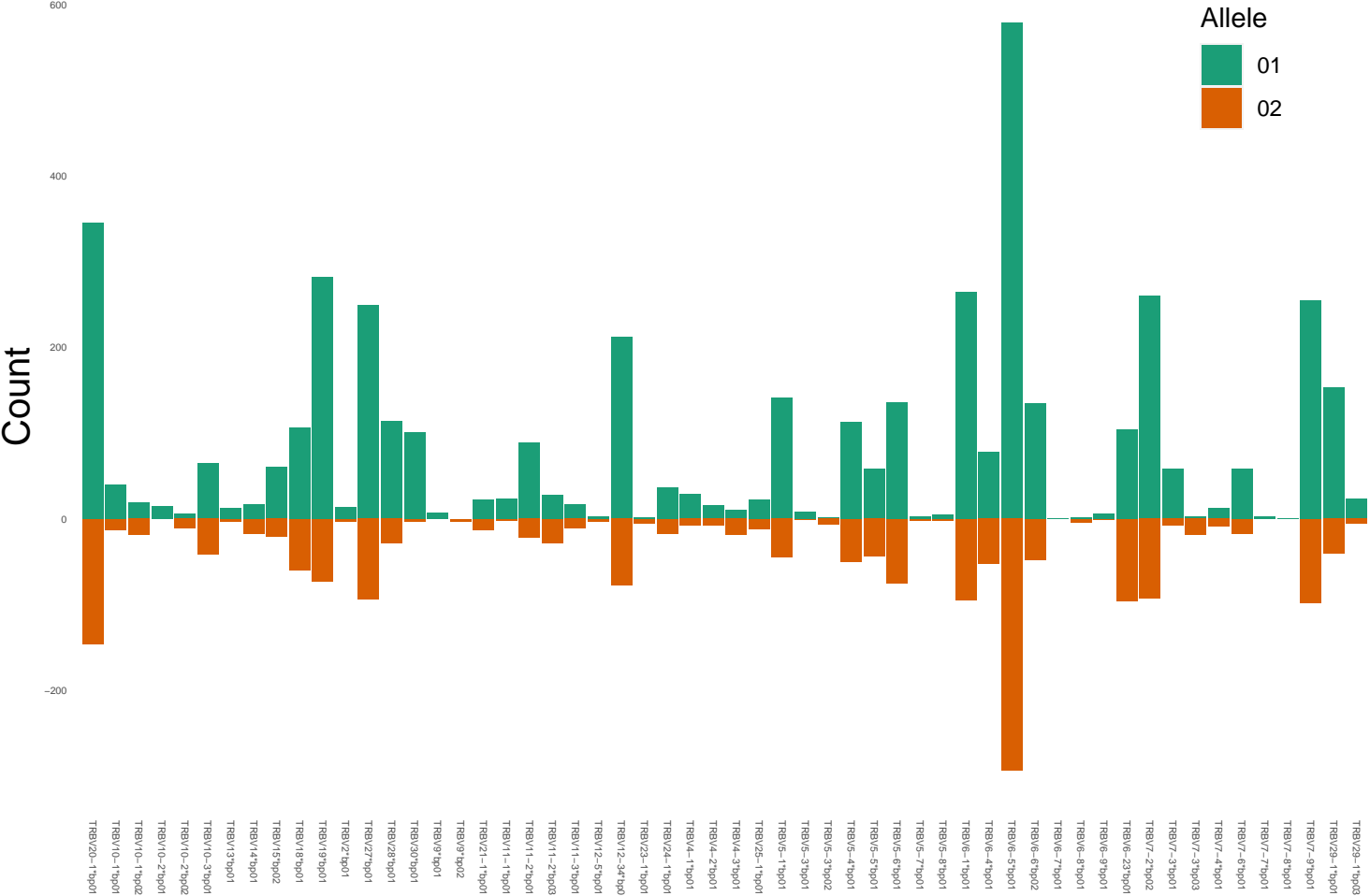
2143 sequences assigned  
1439 (67.1%) exact matches, in which:  
939 unique CDR3  
14 unique J







Sequence Count by TRBJ2-7 allele usage



Warning – no inferred sequences found.