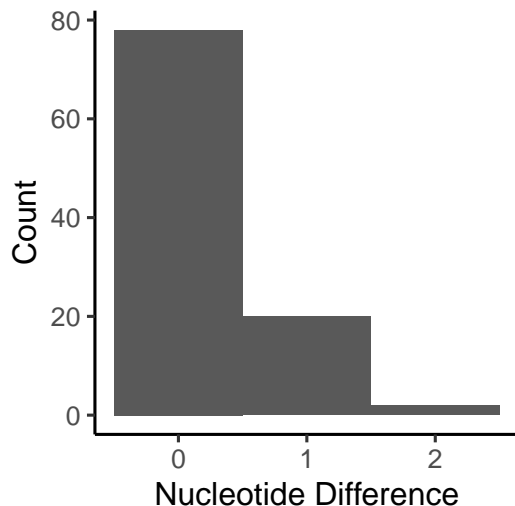


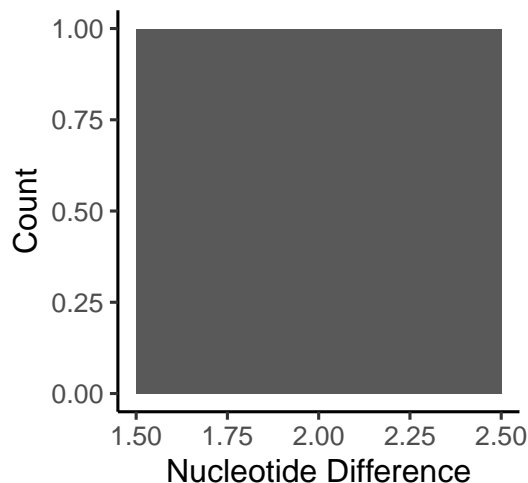
### TRBV20–1\*bp01

100 sequences assigned  
78 (78%) exact matches, in which:  
72 unique CDR3  
12 unique J



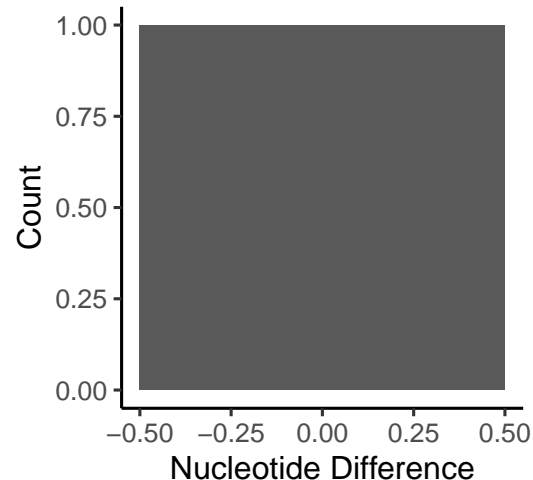
### TRBV10–2\*bp02

1 sequences assigned  
No exact matches.



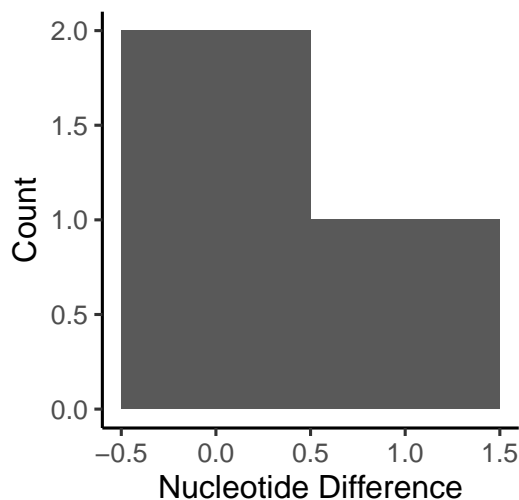
### TRBV14\*bp01

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



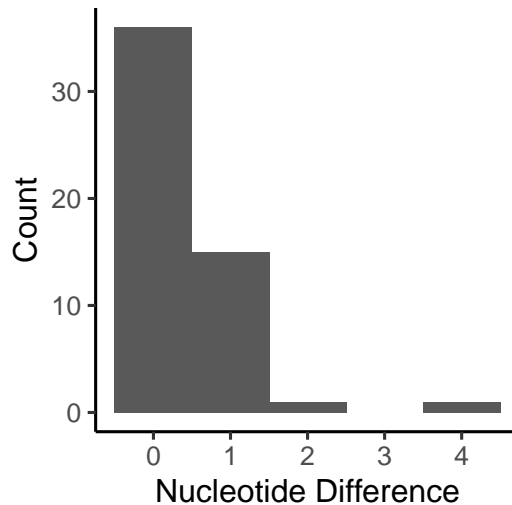
### TRBV10–1\*bp01

3 sequences assigned  
2 (66.7%) exact matches, in which:  
1 unique CDR3  
1 unique J



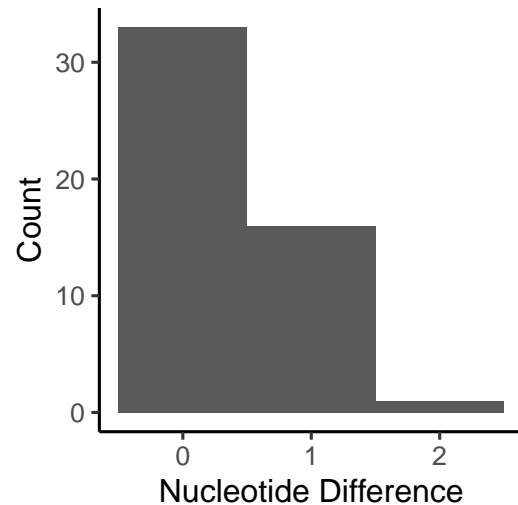
### TRBV10–3\*bp01

53 sequences assigned  
36 (67.9%) exact matches, in which:  
31 unique CDR3  
10 unique J



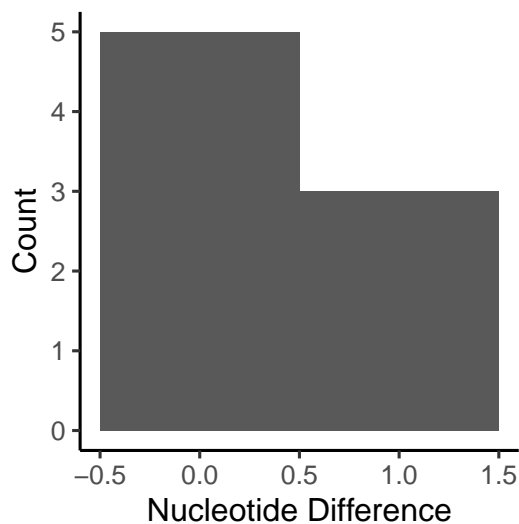
### TRBV15\*bp02

50 sequences assigned  
33 (66%) exact matches, in which:  
28 unique CDR3  
10 unique J



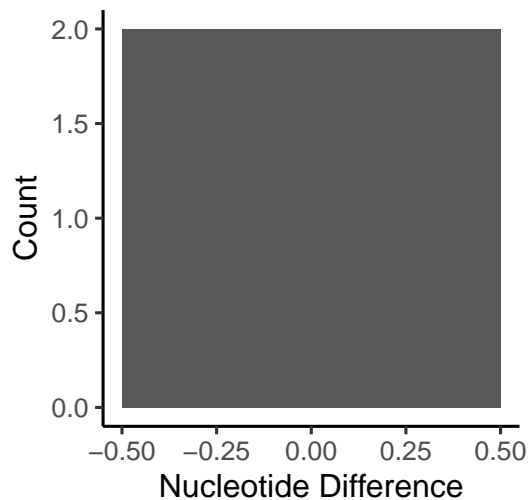
### TRBV10–1\*bp02

8 sequences assigned  
5 (62.5%) exact matches, in which:  
5 unique CDR3  
4 unique J



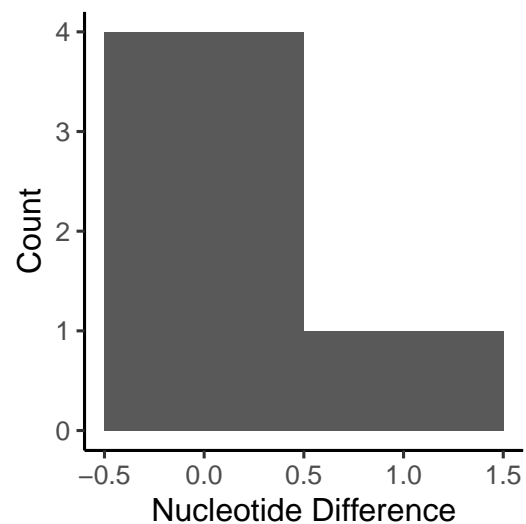
### TRBV13\*bp01

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



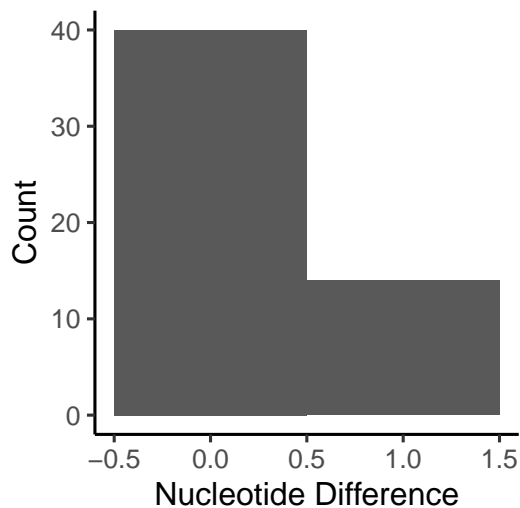
### TRBV18\*bp01

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



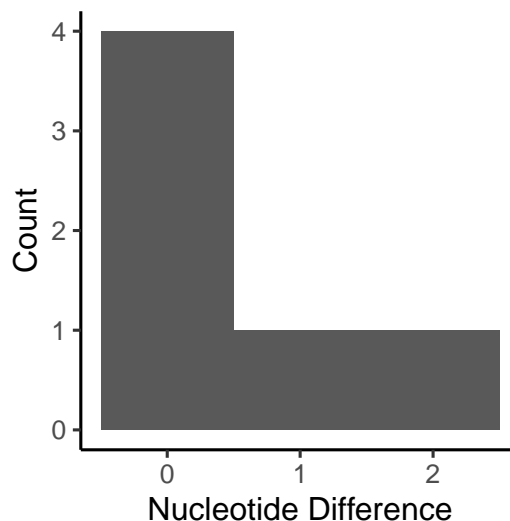
### TRBV19\*bp01

54 sequences assigned  
40 (74.1%) exact matches, in which:  
36 unique CDR3  
11 unique J



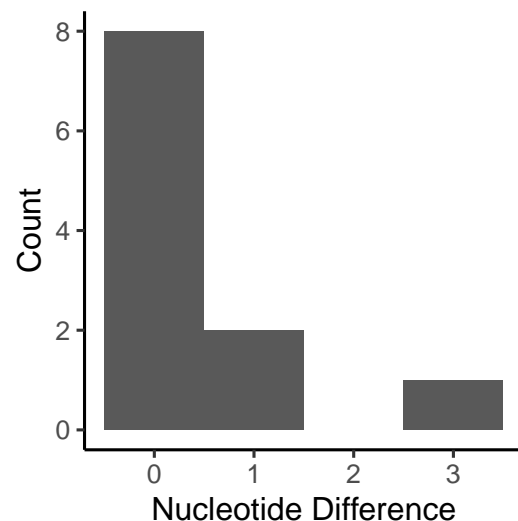
### TRBV28\*bp01

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



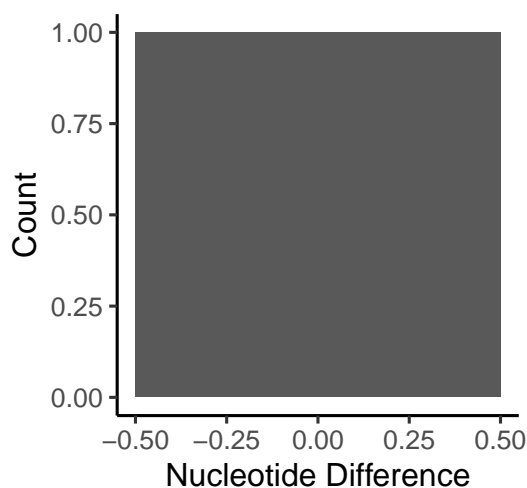
### TRBV11-1\*bp01

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



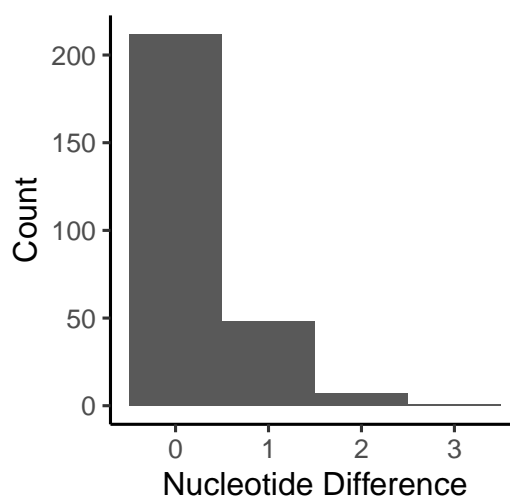
### TRBV2\*bp01

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



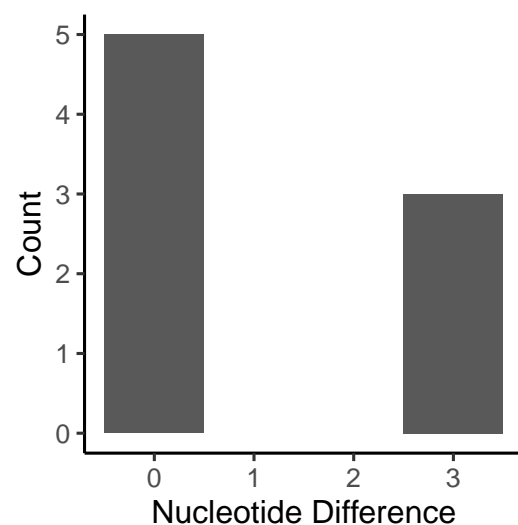
### TRBV30\*bp01

268 sequences assigned  
212 (79.1%) exact matches, in which:  
124 unique CDR3  
13 unique J



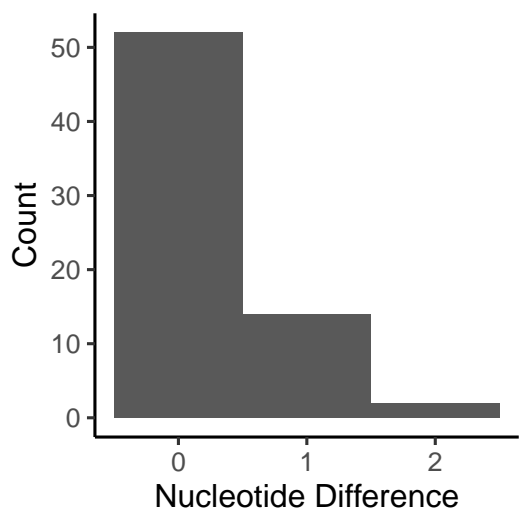
### TRBV11-2\*bp01

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



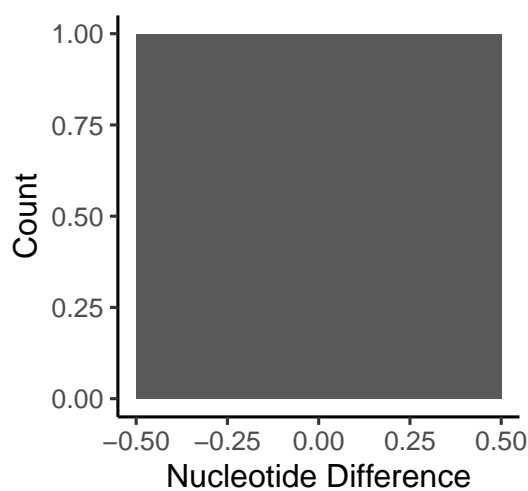
### TRBV27\*bp01

68 sequences assigned  
52 (76.5%) exact matches, in which:  
46 unique CDR3  
9 unique J



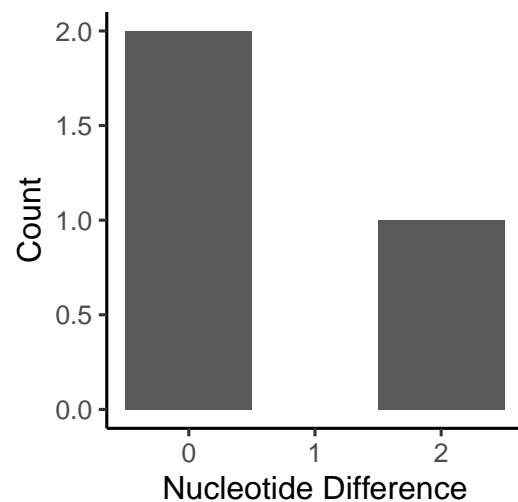
### TRBV9\*bp01

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



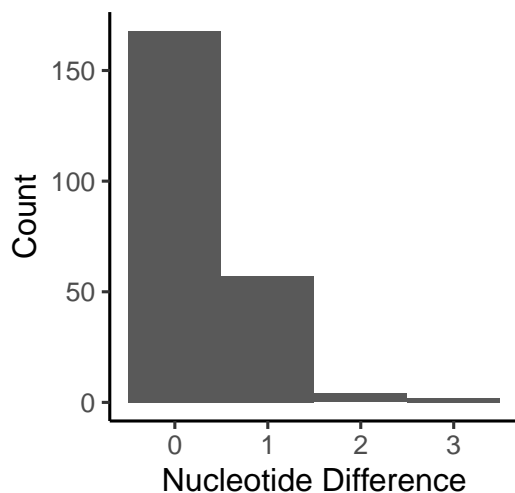
### TRBV12-5\*bp01

3 sequences assigned  
2 (66.7%) exact matches, in which:  
2 unique CDR3  
2 unique J



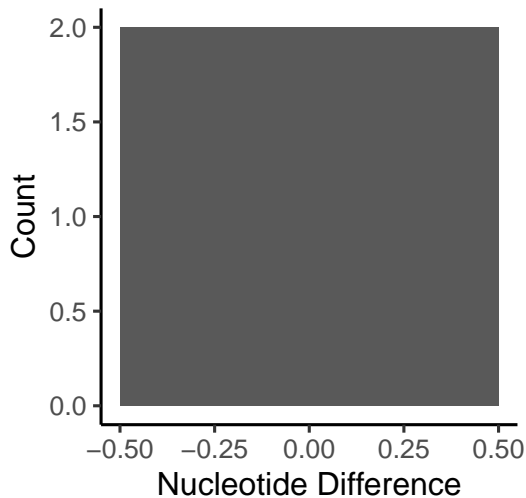
### TRBV12–34\*bp01

231 sequences assigned  
168 (72.7%) exact matches, in which:  
129 unique CDR3  
14 unique J



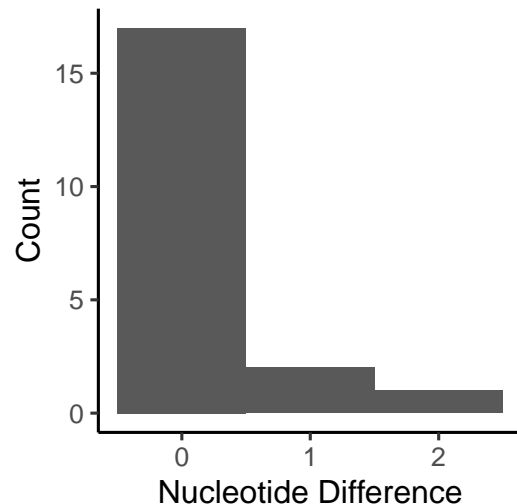
### TRBV25–1\*bp01

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



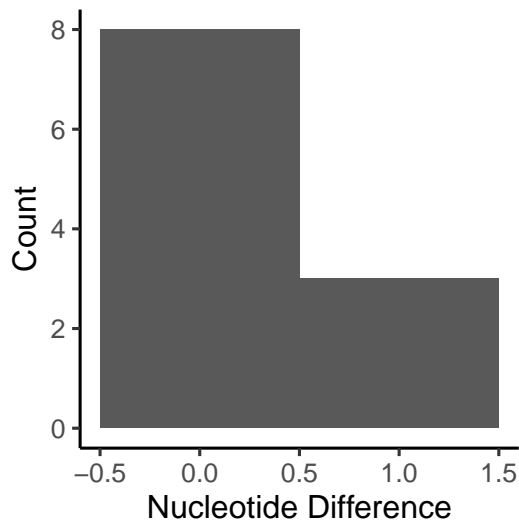
### TRBV5–4\*bp01

20 sequences assigned  
17 (85%) exact matches, in which:  
17 unique CDR3  
8 unique J



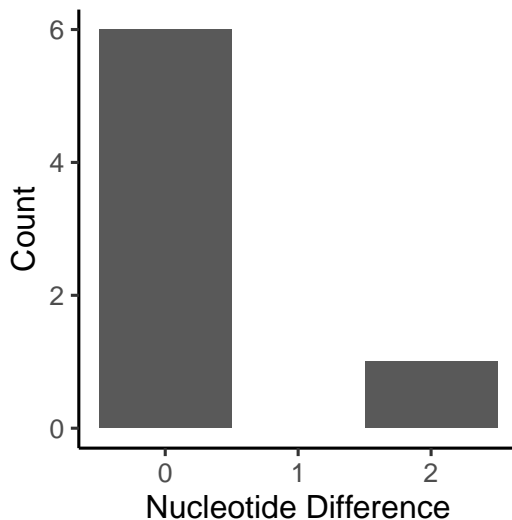
### TRBV24–1\*bp01

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



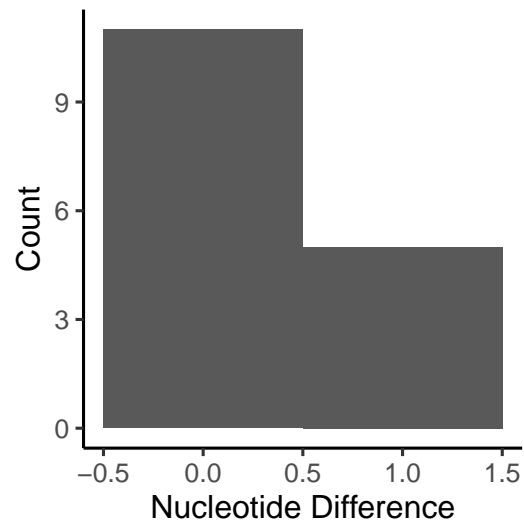
### TRBV5–1\*bp01

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



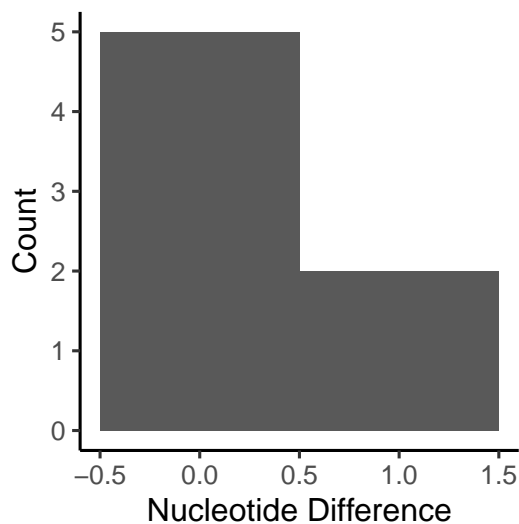
### TRBV5–5\*bp01

16 sequences assigned  
11 (68.8%) exact matches, in which:  
11 unique CDR3  
7 unique J



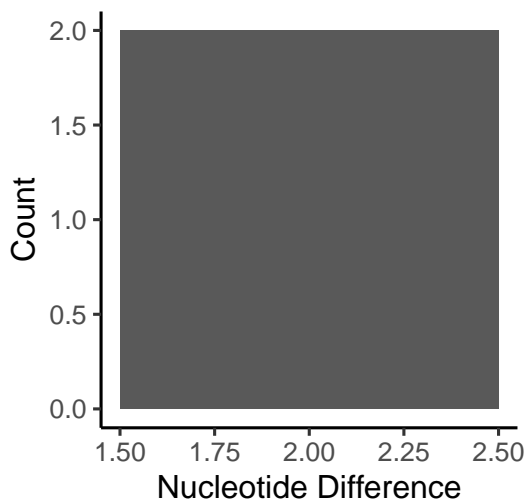
### TRBV4–3\*bp01

7 sequences assigned  
5 (71.4%) exact matches, in which:  
5 unique CDR3  
3 unique J



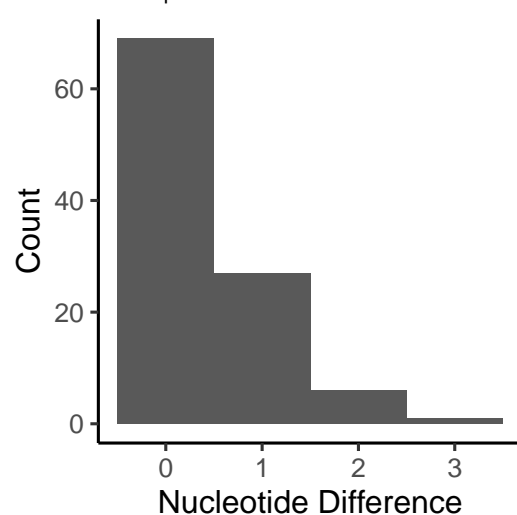
### TRBV5–3\*bp01

2 sequences assigned  
No exact matches.



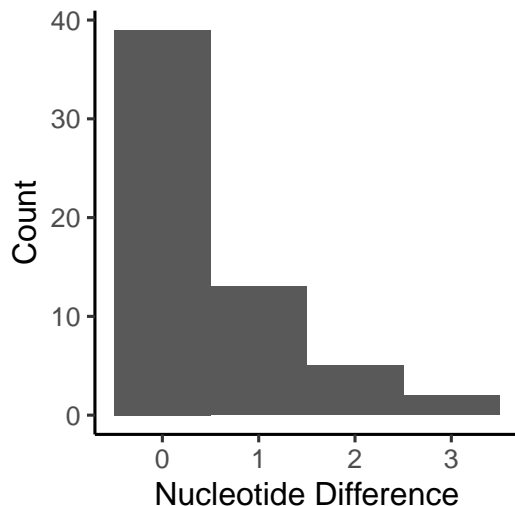
### TRBV5–6\*bp01

103 sequences assigned  
69 (67%) exact matches, in which:  
57 unique CDR3  
10 unique J



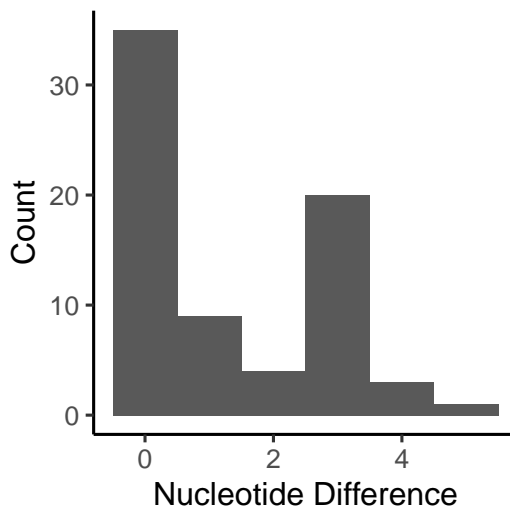
### TRBV6-1\*bp01

59 sequences assigned  
39 (66.1%) exact matches, in which:  
35 unique CDR3  
10 unique J



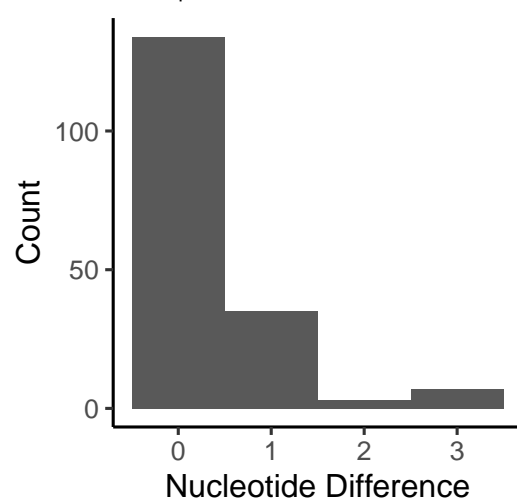
### TRBV6-6\*bp01

72 sequences assigned  
35 (48.6%) exact matches, in which:  
33 unique CDR3  
9 unique J



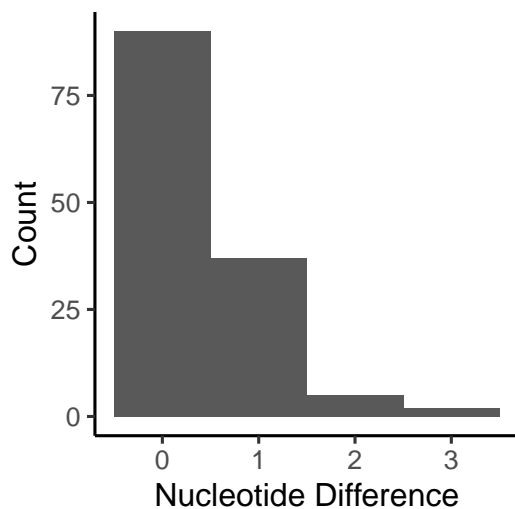
### TRBV7-2\*bp01

179 sequences assigned  
134 (74.9%) exact matches, in which:  
107 unique CDR3  
14 unique J



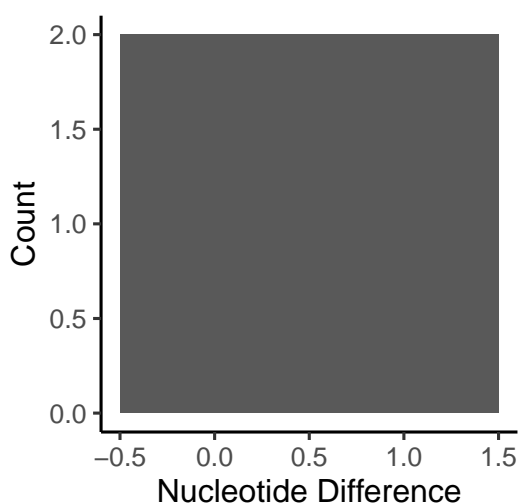
### TRBV6-4\*bp01

134 sequences assigned  
90 (67.2%) exact matches, in which:  
64 unique CDR3  
11 unique J



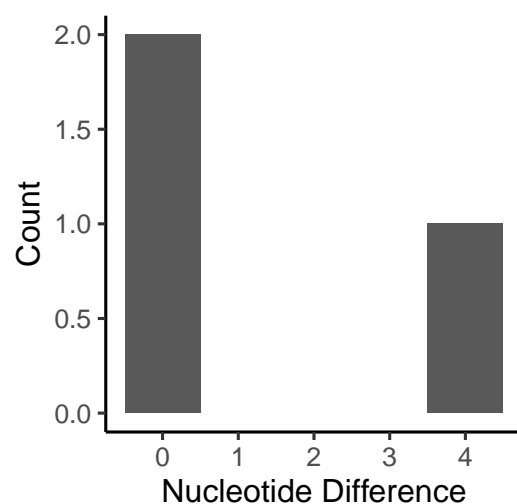
### TRBV6-9\*bp01

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



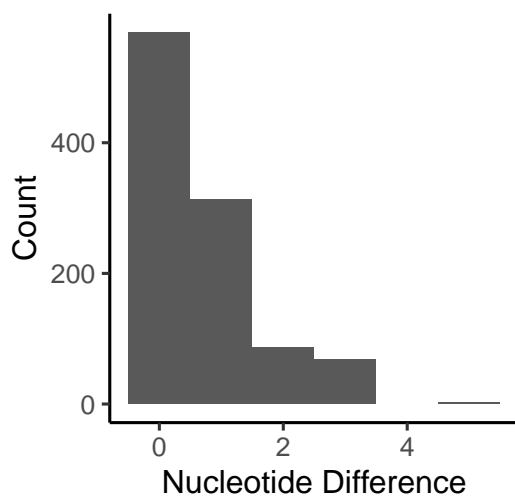
### TRBV7-3\*bp02

3 sequences assigned  
2 (66.7%) exact matches, in which:  
2 unique CDR3  
2 unique J



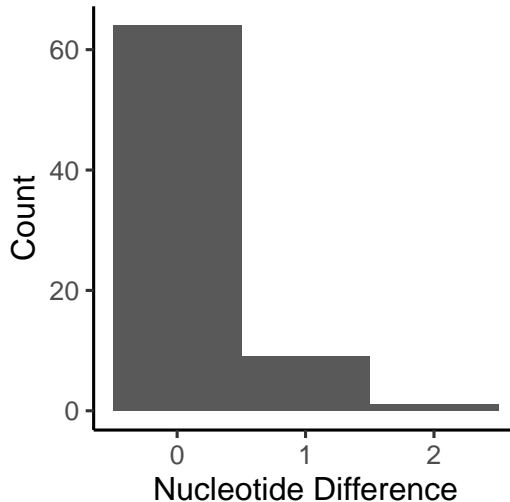
### TRBV6-5\*bp01

1039 sequences assigned  
569 (54.8%) exact matches, in which:  
429 unique CDR3  
14 unique J



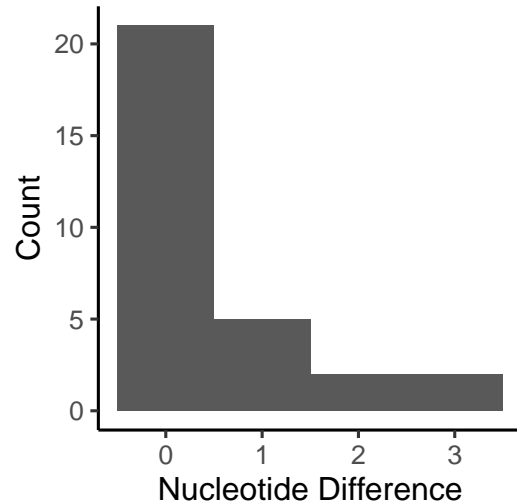
### TRBV6-23\*bp01

74 sequences assigned  
64 (86.5%) exact matches, in which:  
62 unique CDR3  
10 unique J



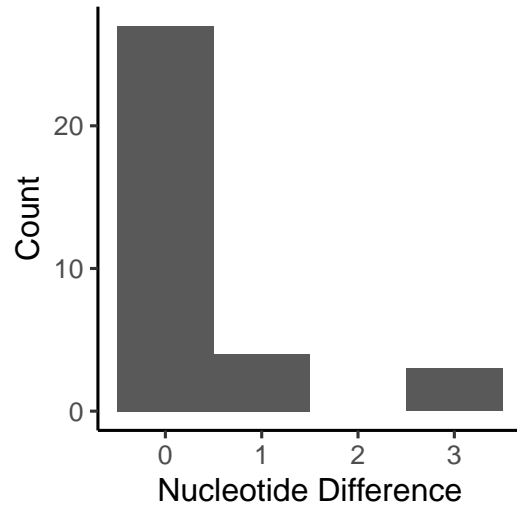
### TRBV7-4\*bp01

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



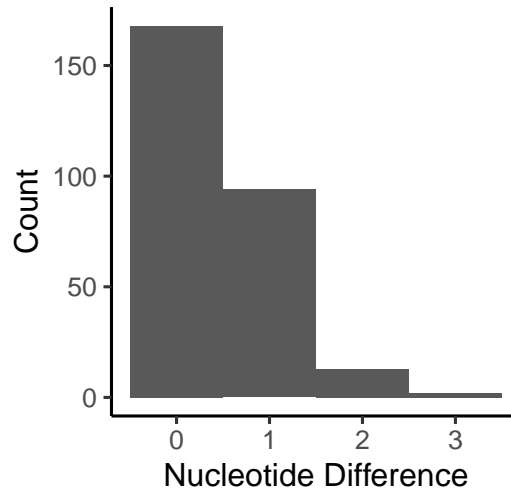
### TRBV7-6\*bp01

34 sequences assigned  
27 (79.4%) exact matches, in which:  
24 unique CDR3  
7 unique J



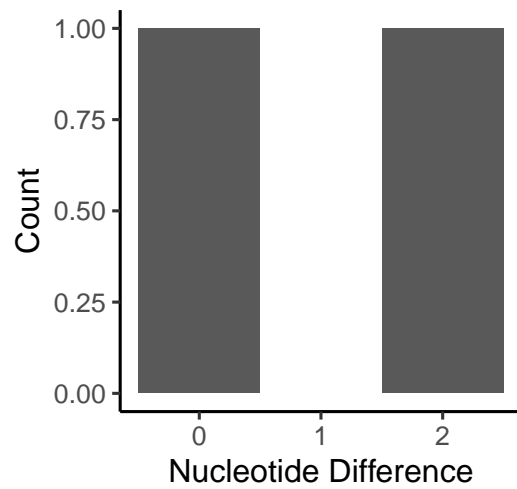
### TRBV29-1\*bp01

277 sequences assigned  
168 (60.6%) exact matches, in which:  
115 unique CDR3  
13 unique J



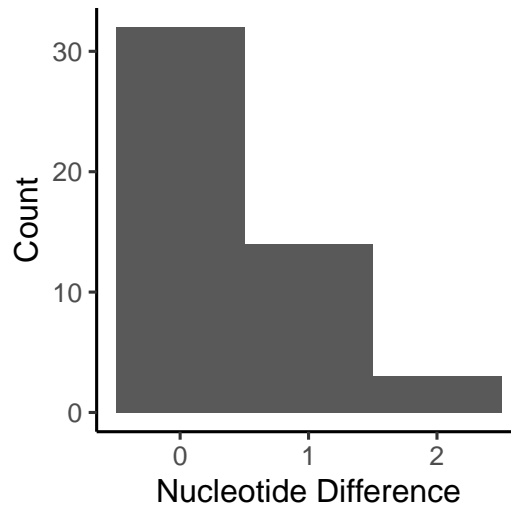
### TRBV7-7\*bp01

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



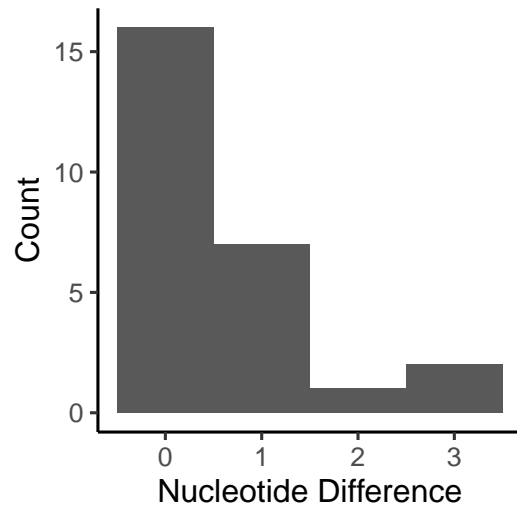
### TRBV29-1\*bp03

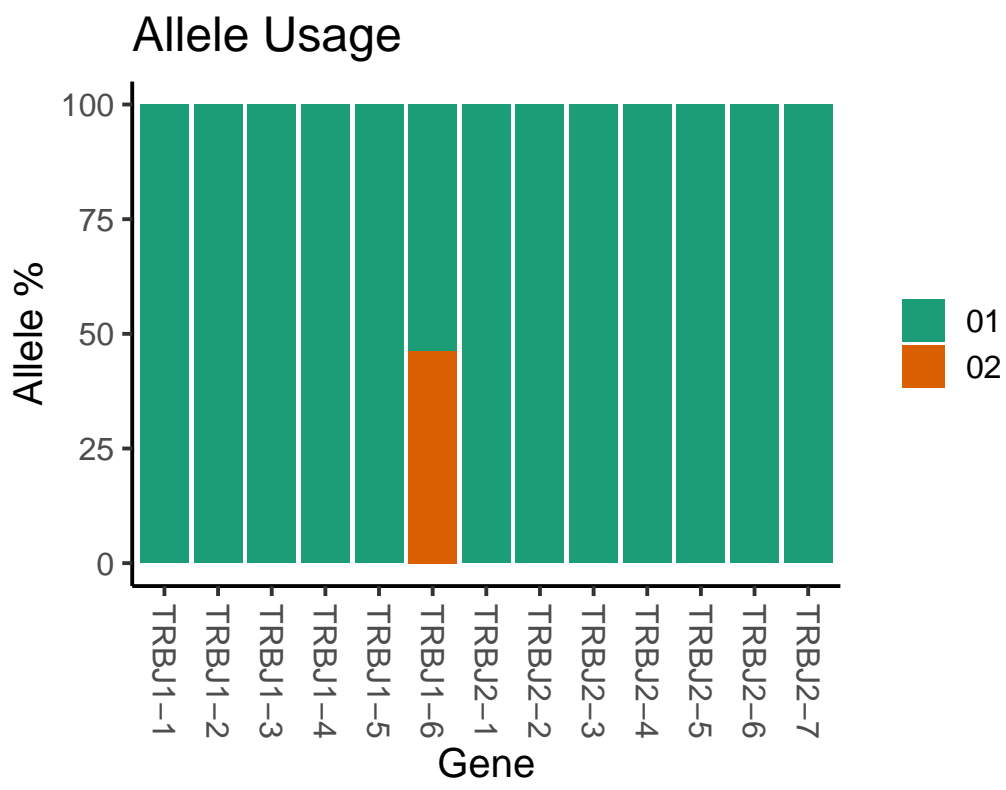
49 sequences assigned  
32 (65.3%) exact matches, in which:  
19 unique CDR3  
11 unique J



### TRBV7-9\*bp01

26 sequences assigned  
16 (61.5%) exact matches, in which:  
16 unique CDR3  
7 unique J





Sequence Count by TRBJ1–6 allele usage

