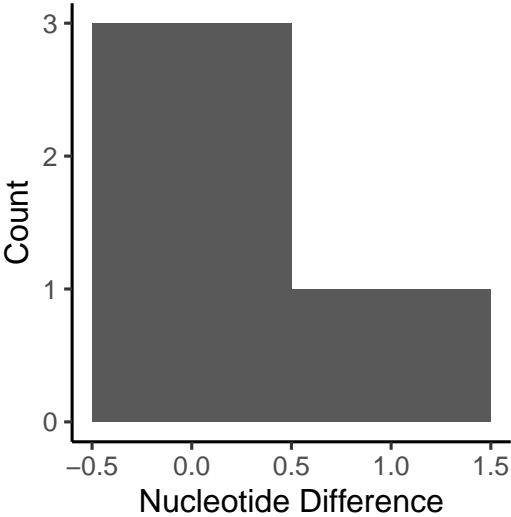


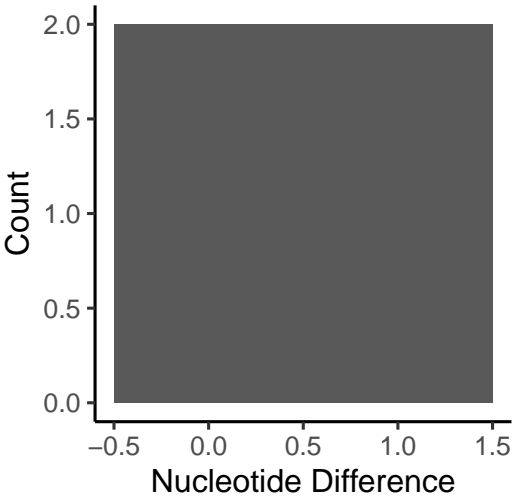
TRBV10–1\*bp01

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



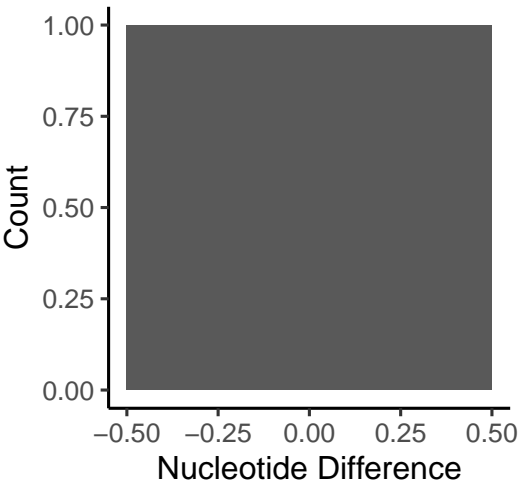
TRBV10–3\*bp01

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



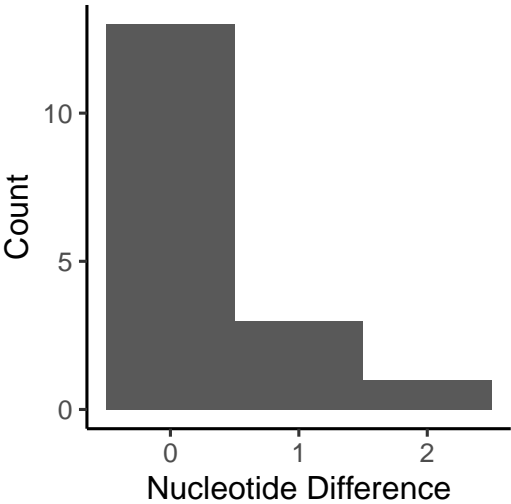
TRBV18\*bp01

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



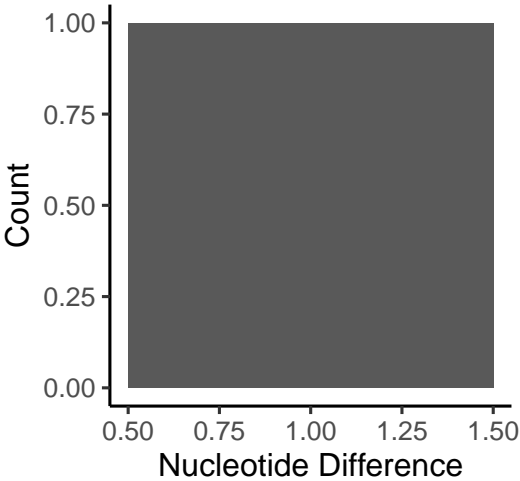
TRBV20–1\*bp01

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



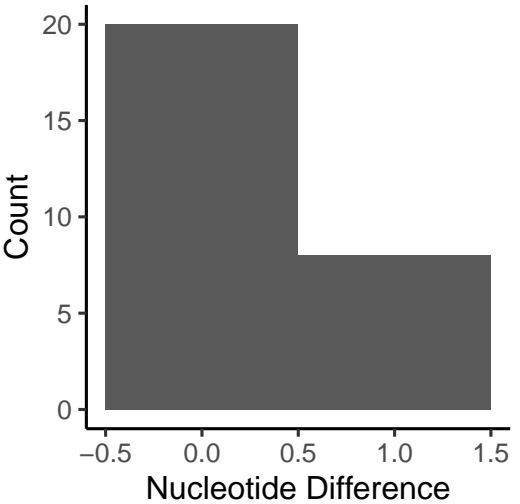
TRBV13\*bp01

1 sequences assigned  
No exact matches.



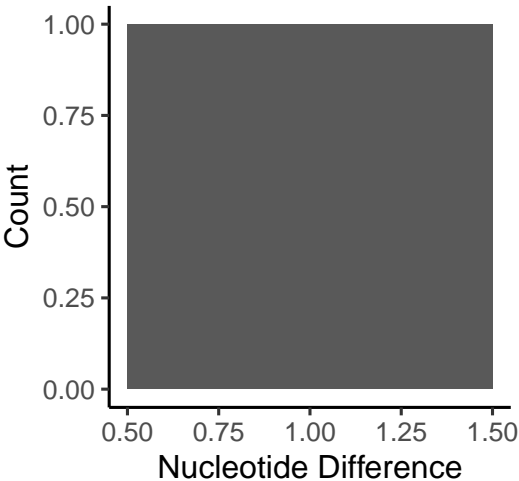
TRBV19\*bp01

28 sequences assigned  
20 (71.4%) exact matches, in which:  
18 unique CDR3  
7 unique J



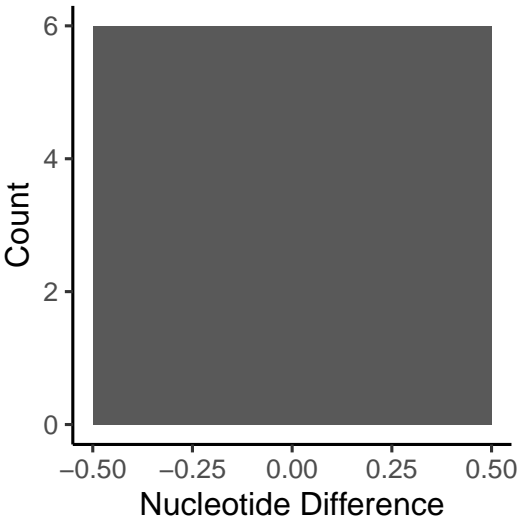
TRBV10–2\*bp01

1 sequences assigned  
No exact matches.



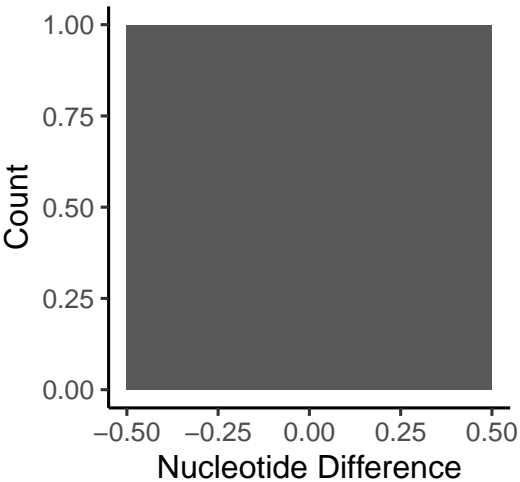
TRBV15\*bp02

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



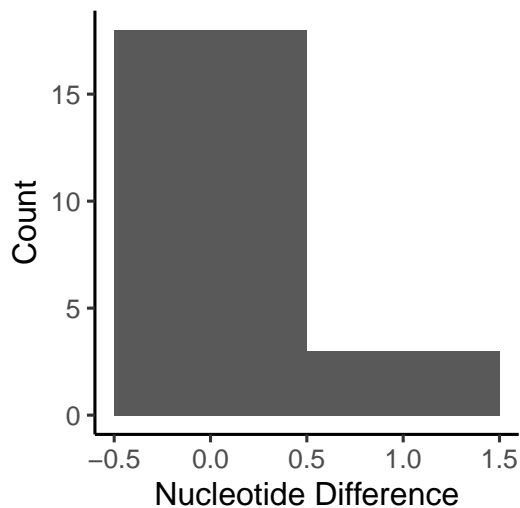
TRBV2\*bp01

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



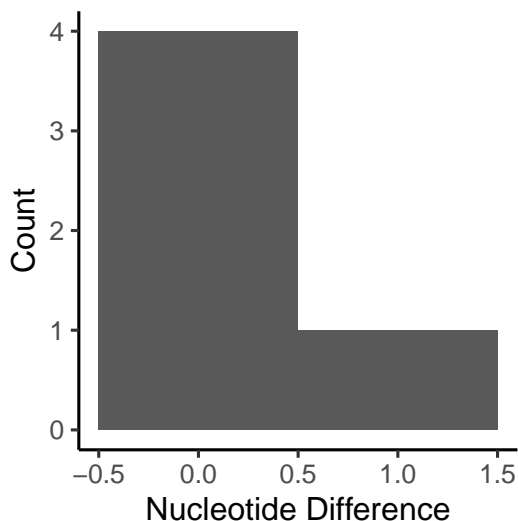
### TRBV27\*bp01

21 sequences assigned  
18 (85.7%) exact matches, in which:  
16 unique CDR3  
6 unique J



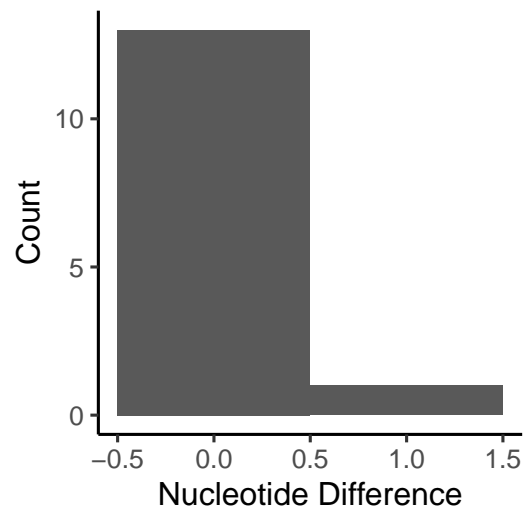
### TRBV9\*bp01

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



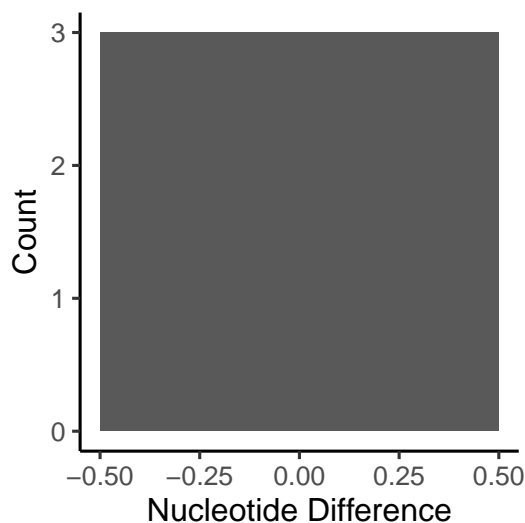
### TRBV12-34\*bp01

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



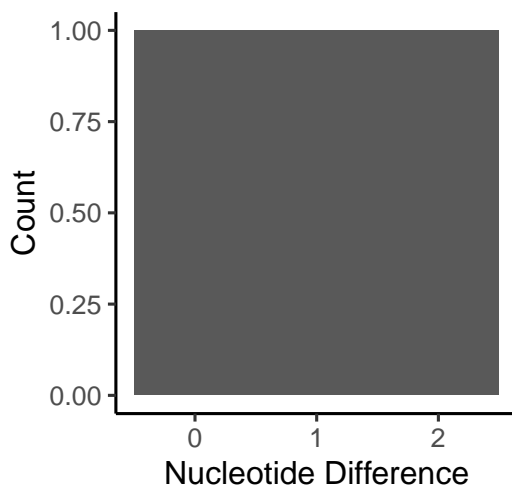
### TRBV28\*bp01

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



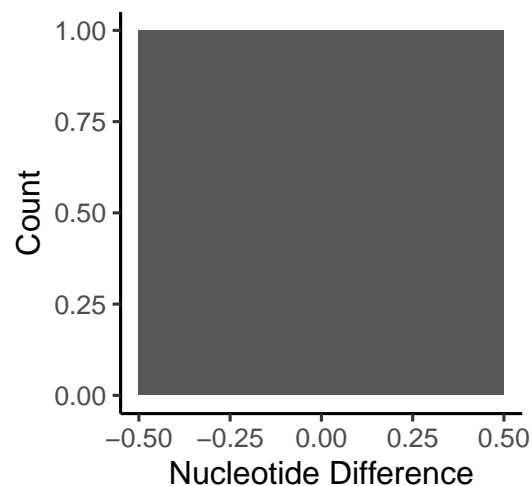
### TRBV11-2\*bp01

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



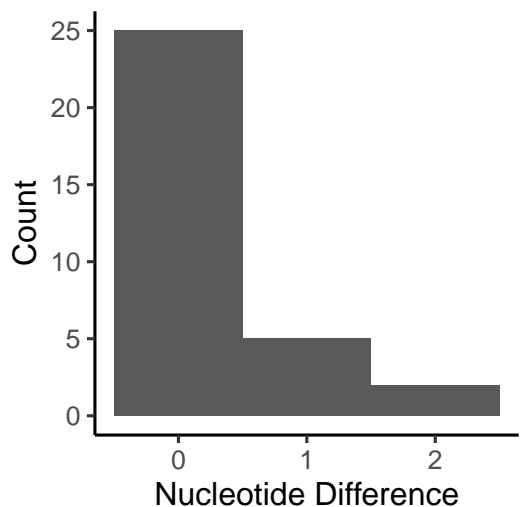
### TRBV23-1\*bp01

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



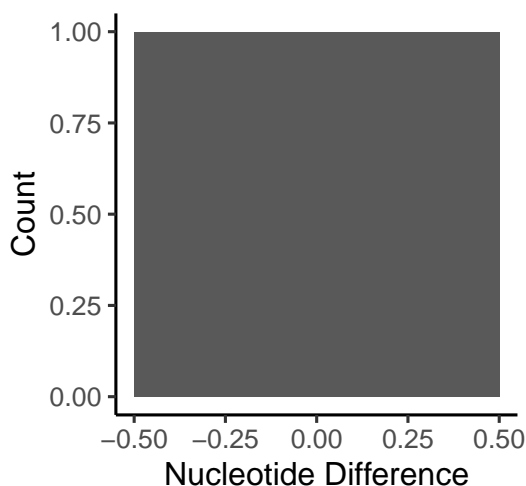
### TRBV30\*bp01

32 sequences assigned  
25 (78.1%) exact matches, in which:  
17 unique CDR3  
6 unique J



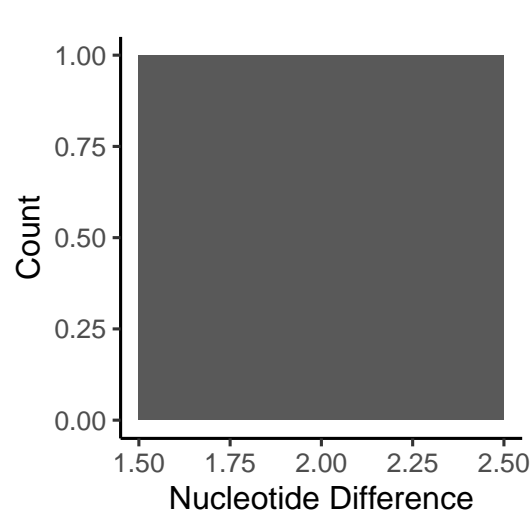
### TRBV11-3\*bp01

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



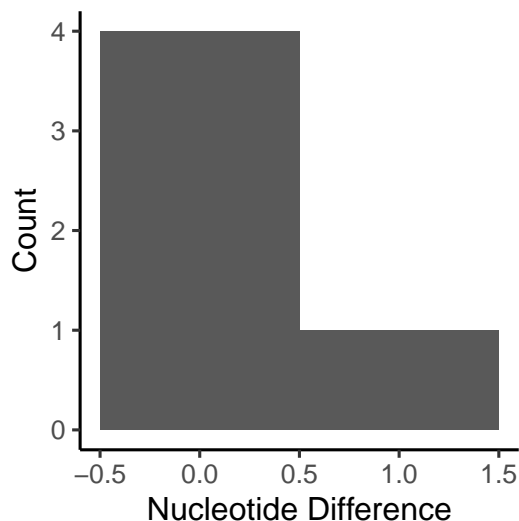
### TRBV24-1\*bp01

1 sequences assigned  
No exact matches.



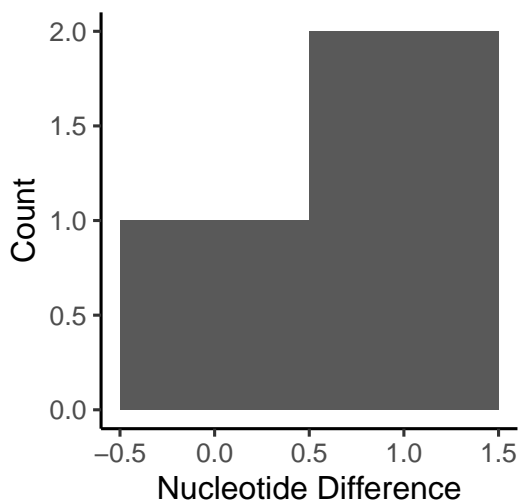
### TRBV4-2\*bp01

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



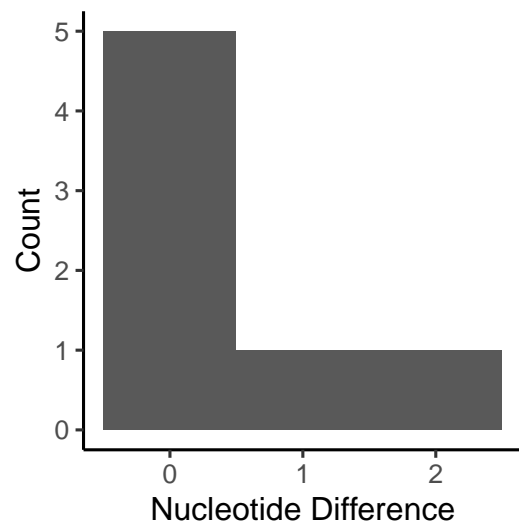
### TRBV5-4\*bp01

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



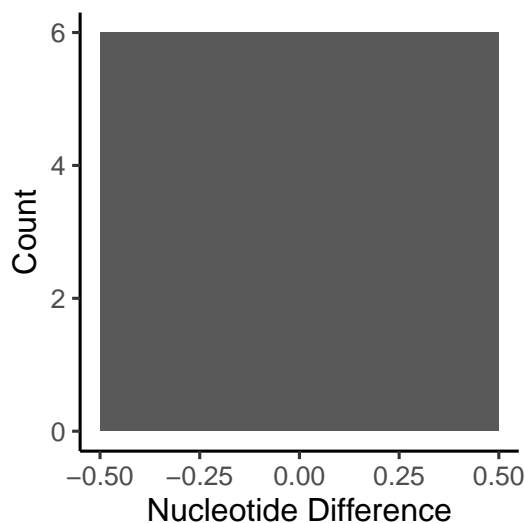
### TRBV6-1\*bp01

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



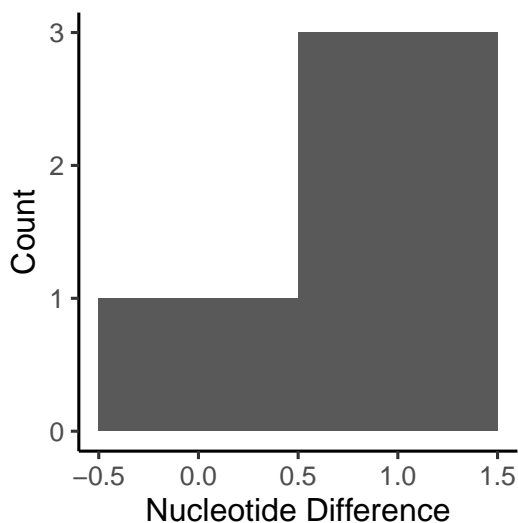
### TRBV4-3\*bp01

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



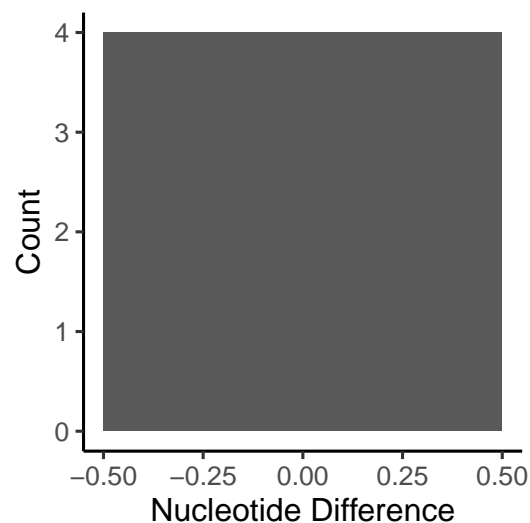
### TRBV5-5\*bp01

4 sequences assigned  
1 (25%) exact matches, in which:  
1 unique CDR3  
1 unique J



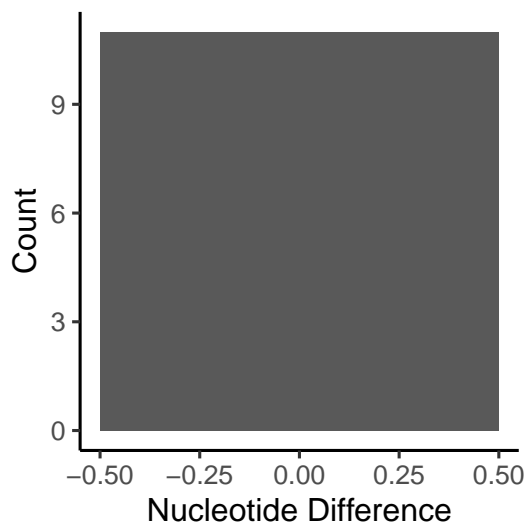
### TRBV6-4\*bp01

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



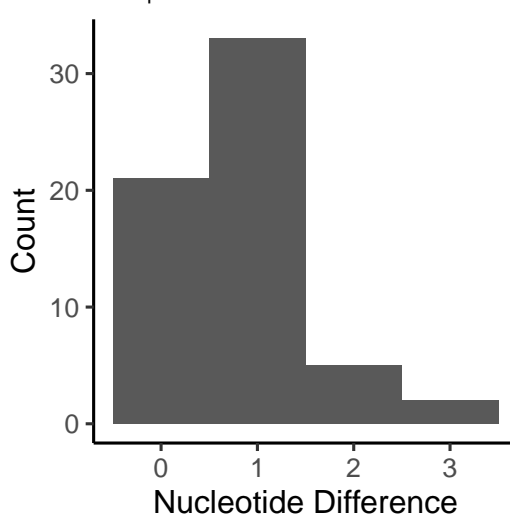
### TRBV5-1\*bp01

11 sequences assigned  
11 (100%) exact matches, in which:  
9 unique CDR3  
4 unique J



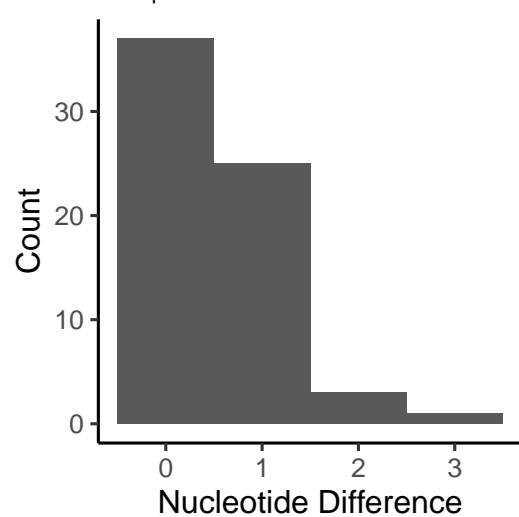
### TRBV5-6\*bp01

61 sequences assigned  
21 (34.4%) exact matches, in which:  
12 unique CDR3  
5 unique J



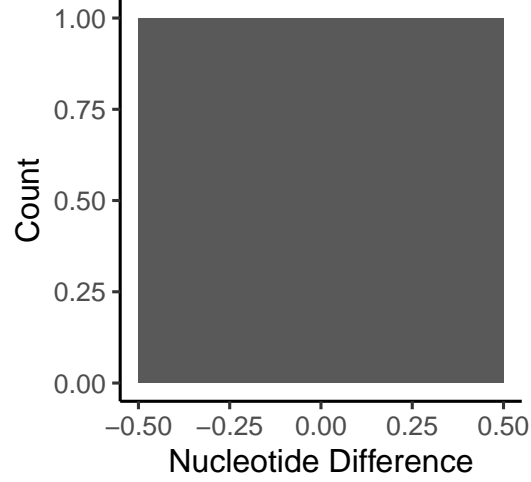
### TRBV6-5\*bp01

66 sequences assigned  
37 (56.1%) exact matches, in which:  
27 unique CDR3  
7 unique J



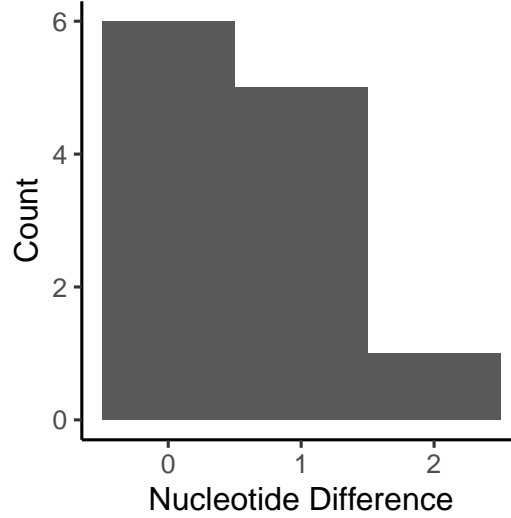
### TRBV6–6\*bp01

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



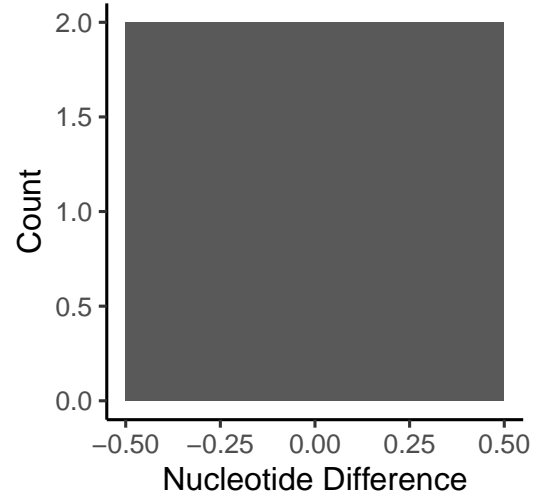
### TRBV6–23\*bp01

12 sequences assigned  
6 (50%) exact matches, in which:  
6 unique CDR3  
5 unique J



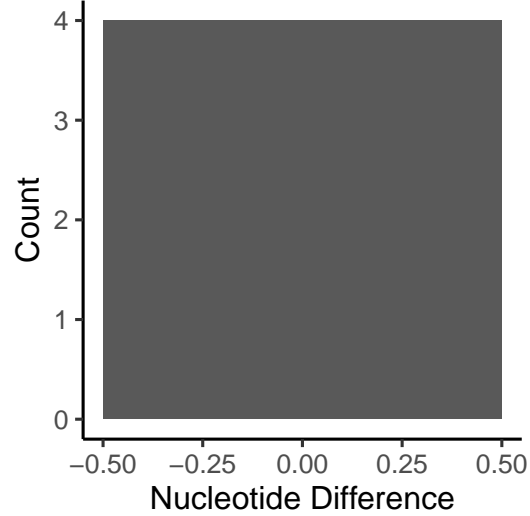
### TRBV7–3\*bp02

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



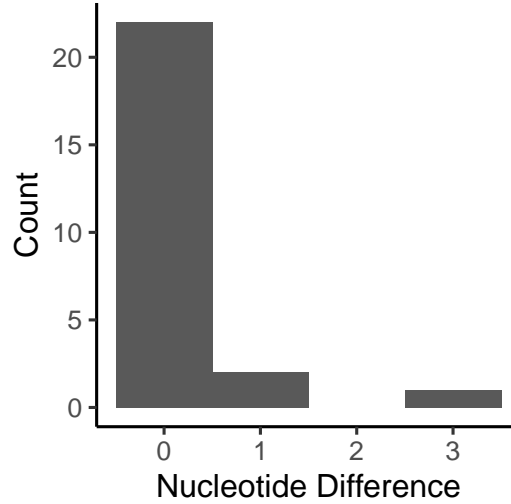
### TRBV6–6\*bp02

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



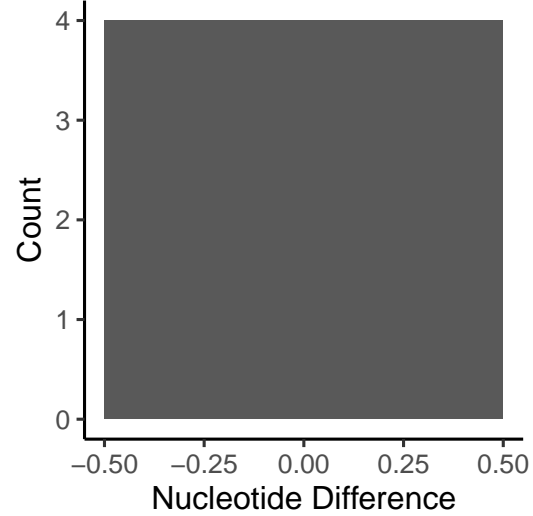
### TRBV7–2\*bp01

25 sequences assigned  
22 (88%) exact matches, in which:  
18 unique CDR3  
8 unique J



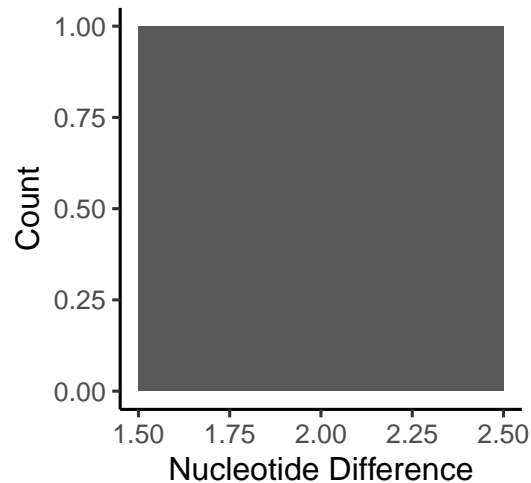
### TRBV7–6\*bp01

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



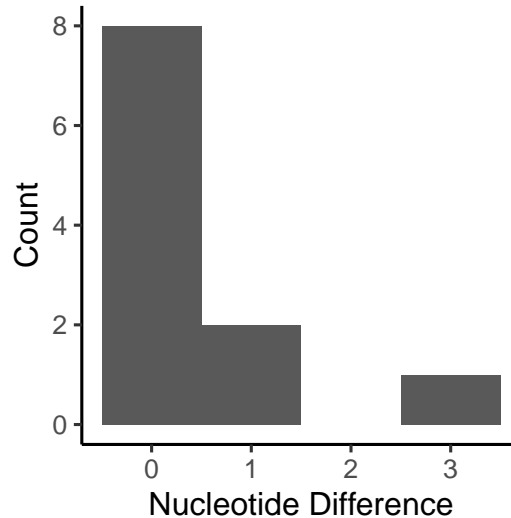
### TRBV6–9\*bp01

1 sequences assigned  
No exact matches.



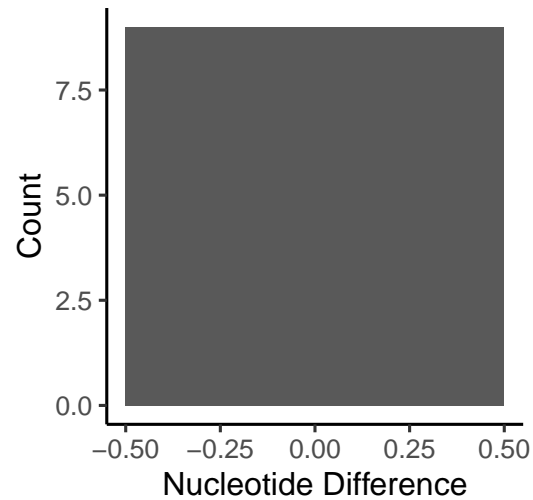
### TRBV7–3\*bp01

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



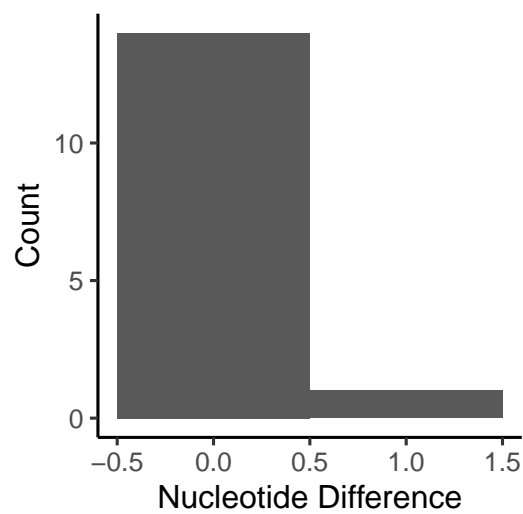
### TRBV7–9\*bp01

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



## TRBV29–1\*bp01

15 sequences assigned  
14 (93.3%) exact matches, in which:  
14 unique CDR3  
8 unique J



## TRBV29–1\*bp03

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

