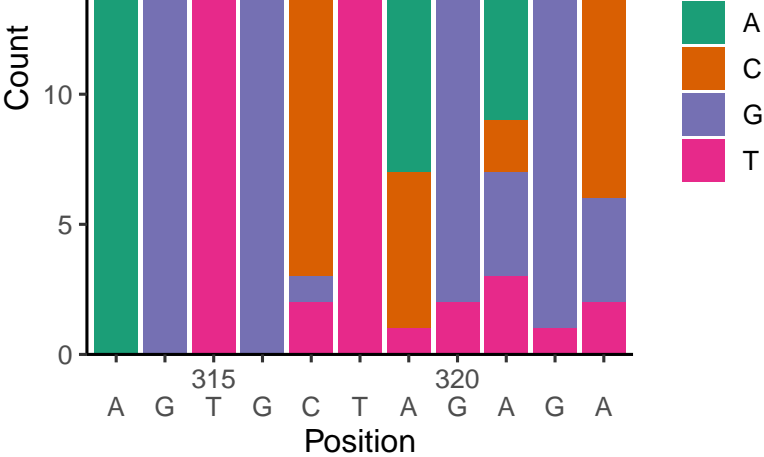


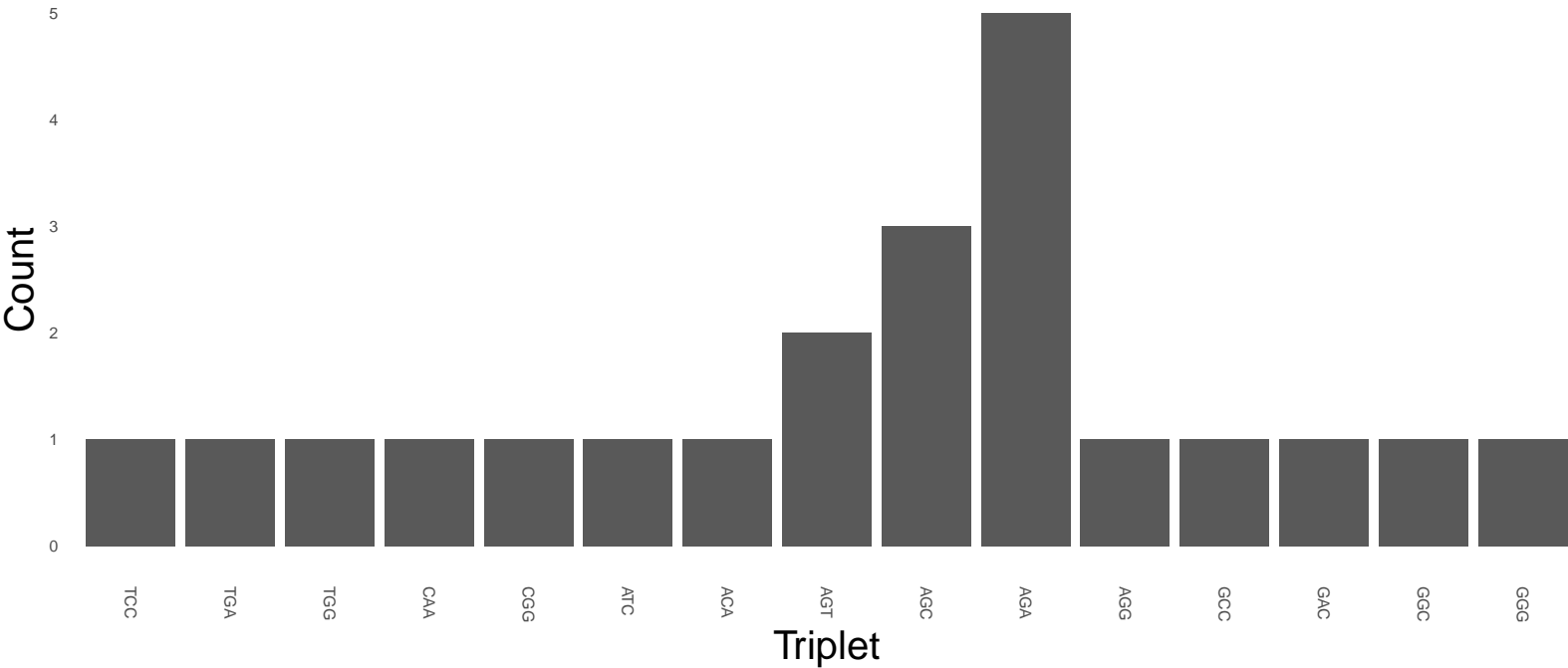
Gene TRBV20-1*02_319AGAGA323



Gene TRBV20-1*02_319AGAGA323

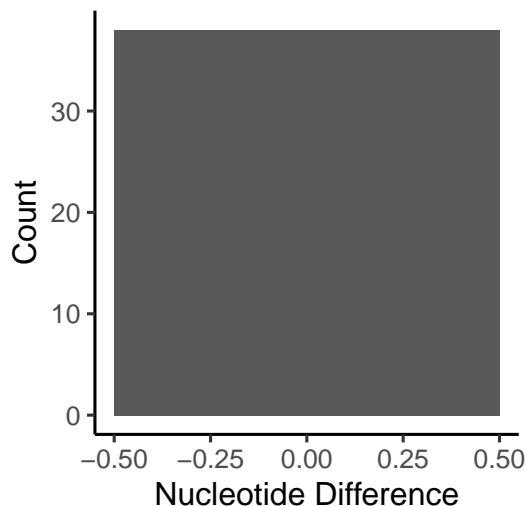


TRBV20-1*02_319AGAGA323- Final 3 nucleotides as a triplet



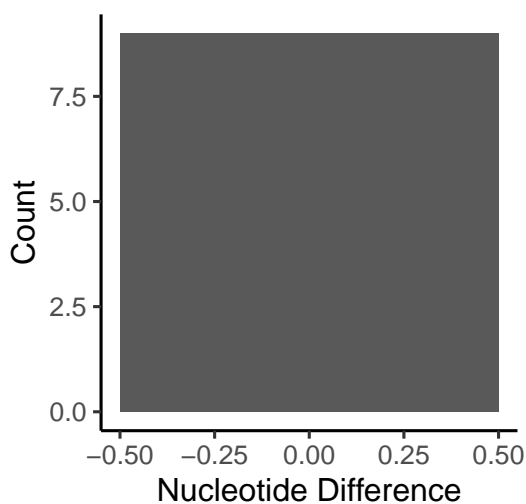
TRBV20–1*01

38 sequences assigned
38 (100%) exact matches, in which:
38 unique CDR3
12 unique J



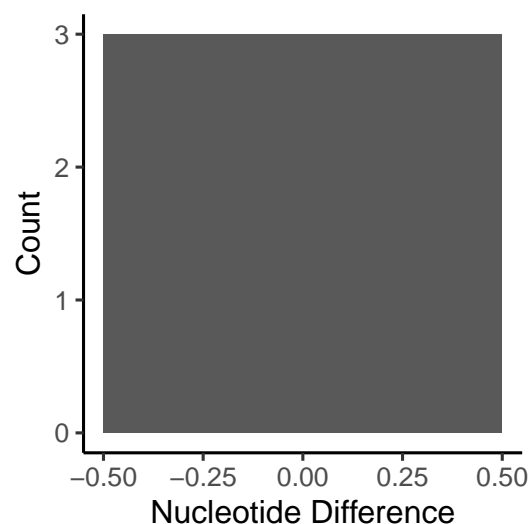
TRBV10–3*01

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



TRBV14*01

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



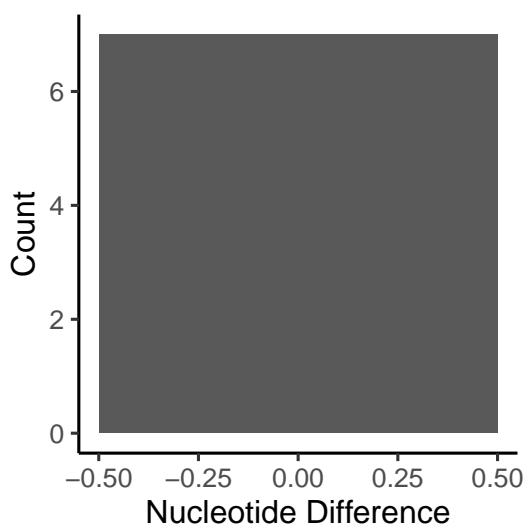
TRBV10–1*01

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



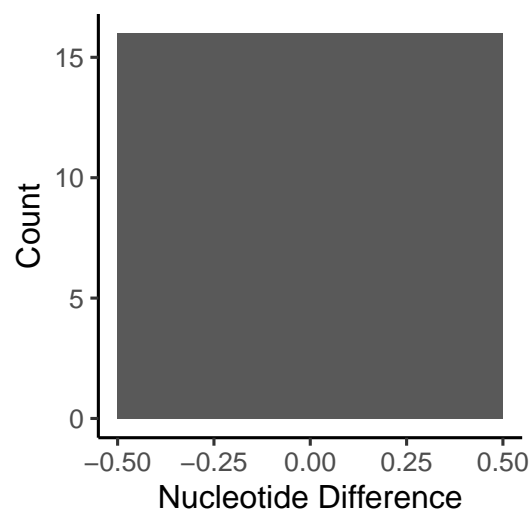
TRBV10–3*02

7 sequences assigned
7 (100%) exact matches, in which:
7 unique CDR3
7 unique J



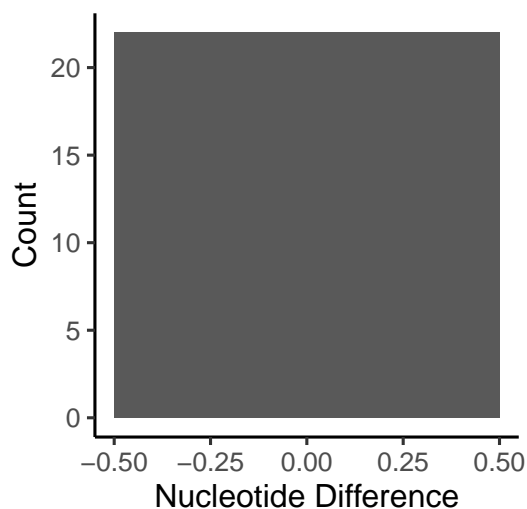
TRBV15*02

16 sequences assigned
16 (100%) exact matches, in which:
16 unique CDR3
7 unique J



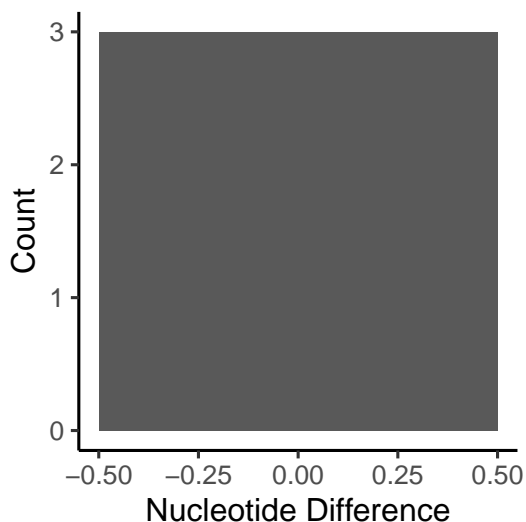
TRBV20–1*02_319AGAG/

22 sequences assigned
22 (100%) exact matches, in which:
22 unique CDR3
10 unique J



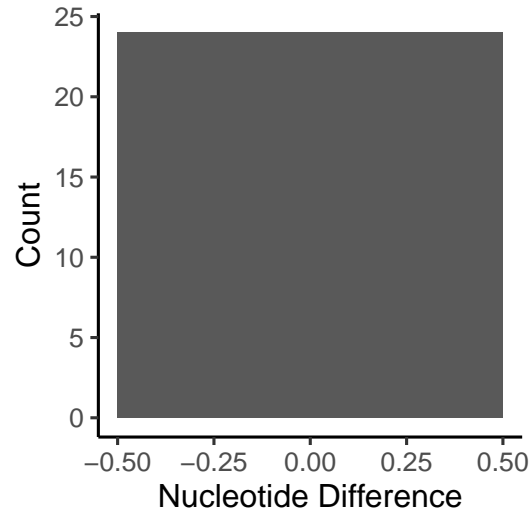
TRBV13*01

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



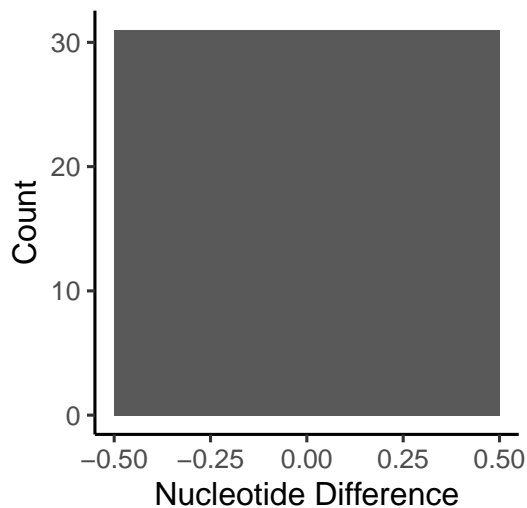
TRBV18*01

24 sequences assigned
24 (100%) exact matches, in which:
24 unique CDR3
11 unique J



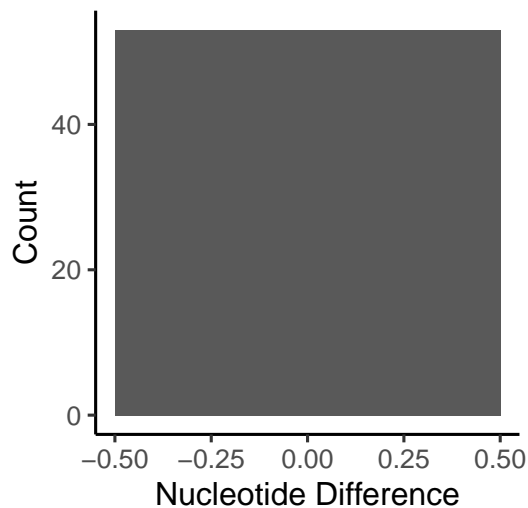
TRBV19*01

31 sequences assigned
31 (100%) exact matches, in which:
31 unique CDR3
10 unique J



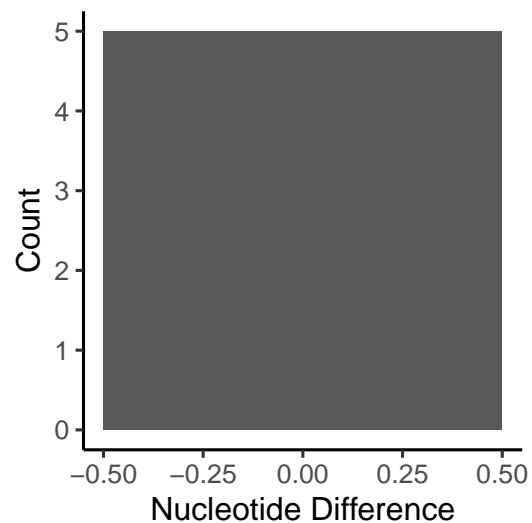
TRBV28*01

53 sequences assigned
53 (100%) exact matches, in which:
53 unique CDR3
13 unique J



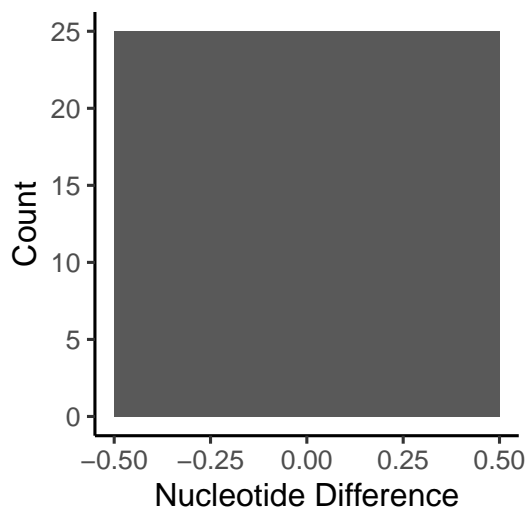
TRBV11-1*01

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



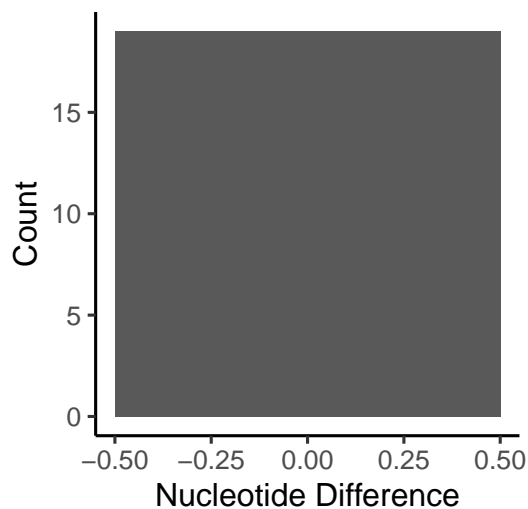
TRBV2*01

25 sequences assigned
25 (100%) exact matches, in which:
25 unique CDR3
9 unique J



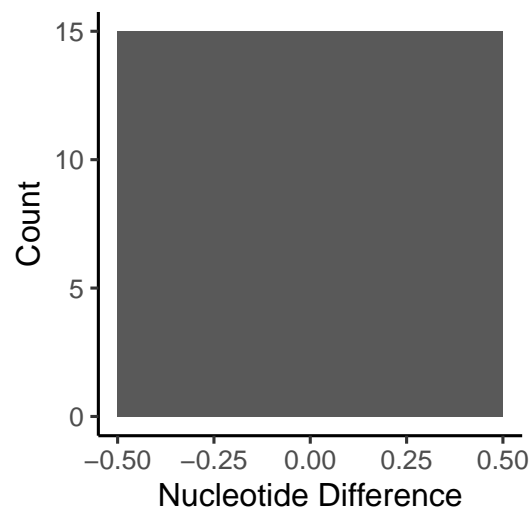
TRBV30*01

19 sequences assigned
19 (100%) exact matches, in which:
19 unique CDR3
10 unique J



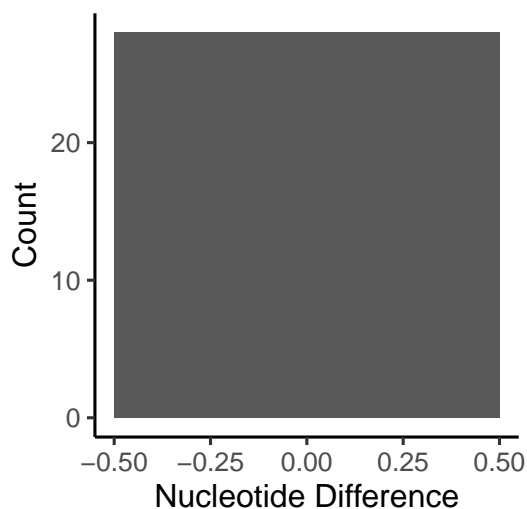
TRBV11-2*01

15 sequences assigned
15 (100%) exact matches, in which:
15 unique CDR3
6 unique J



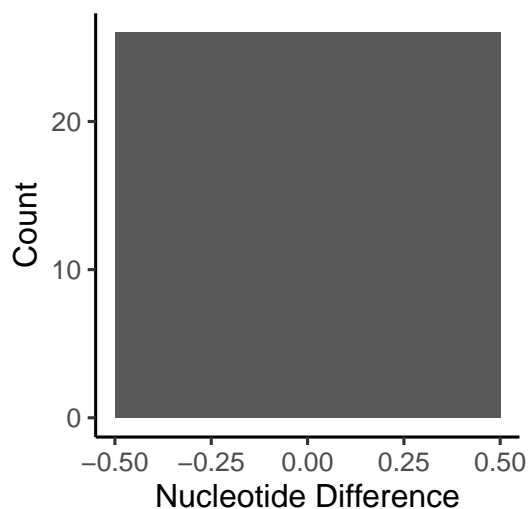
TRBV27*01

28 sequences assigned
28 (100%) exact matches, in which:
28 unique CDR3
9 unique J



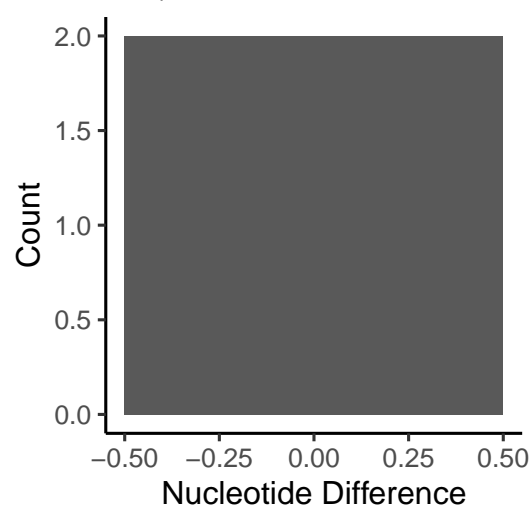
TRBV9*01

26 sequences assigned
26 (100%) exact matches, in which:
26 unique CDR3
10 unique J



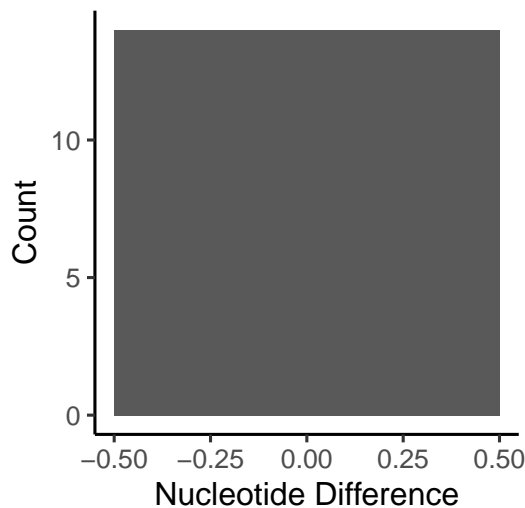
TRBV11-3*01

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



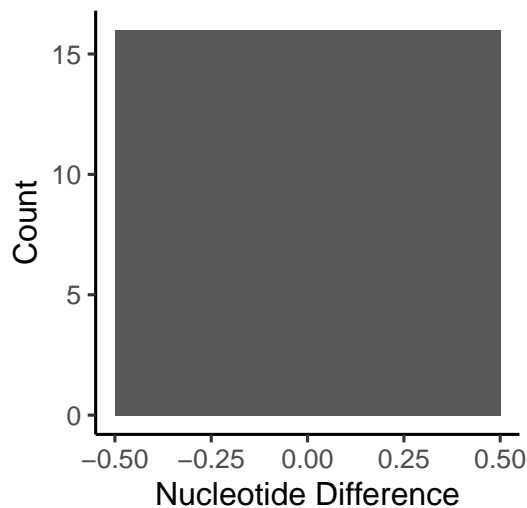
TRBV12-3*01

14 sequences assigned
14 (100%) exact matches, in which:
14 unique CDR3
7 unique J



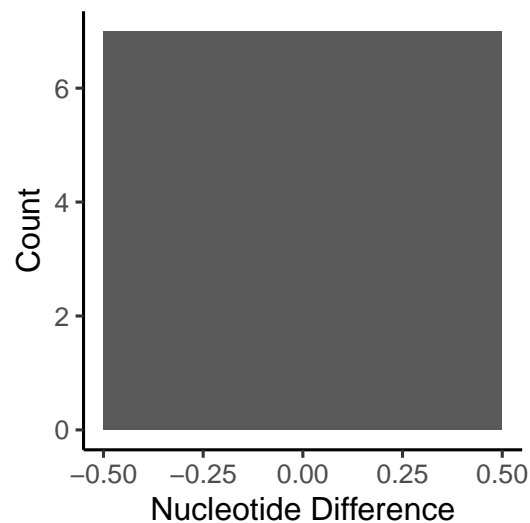
TRBV3-1*01

16 sequences assigned
16 (100%) exact matches, in which:
16 unique CDR3
9 unique J



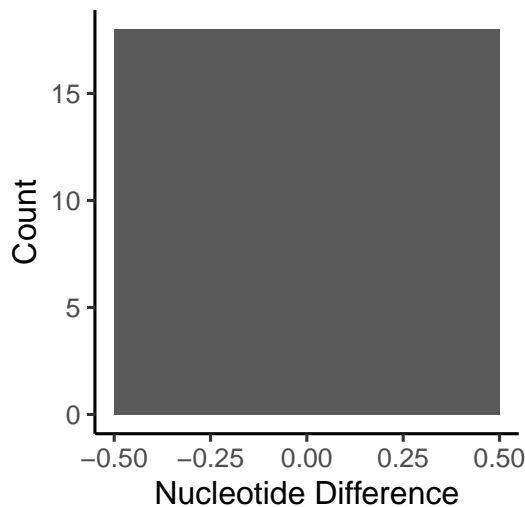
TRBV4-2*01

7 sequences assigned
7 (100%) exact matches, in which:
7 unique CDR3
4 unique J



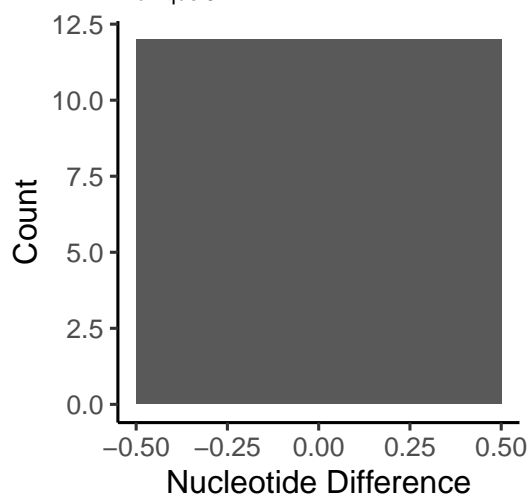
TRBV12-4*01

18 sequences assigned
18 (100%) exact matches, in which:
18 unique CDR3
7 unique J



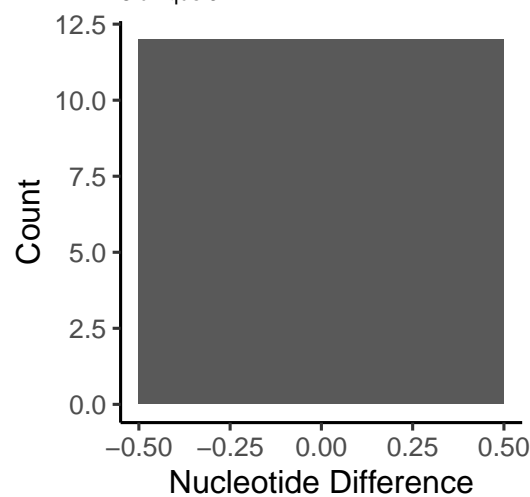
TRBV24-1*01

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



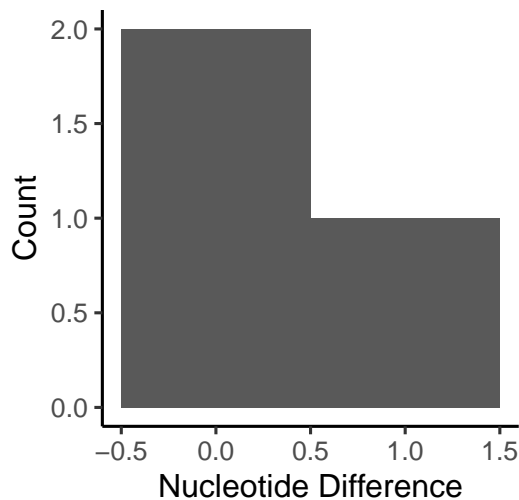
TRBV4-3*01

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



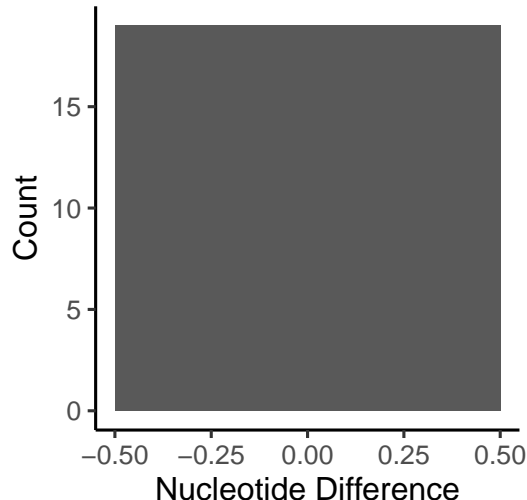
TRBV12-5*01

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



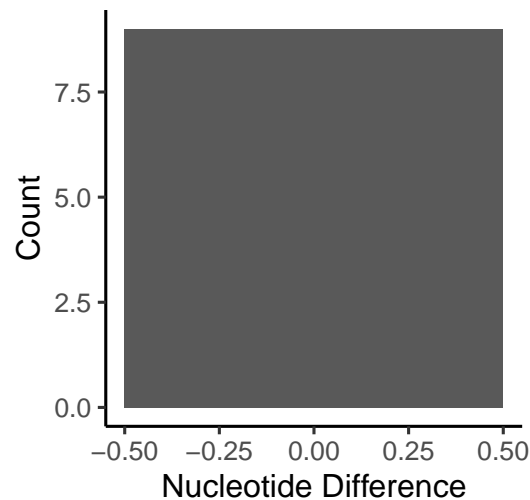
TRBV4-1*01

19 sequences assigned
19 (100%) exact matches, in which:
19 unique CDR3
8 unique J



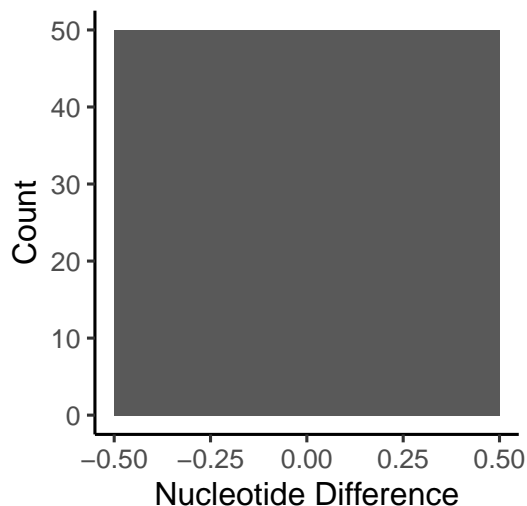
TRBV25-1*01

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



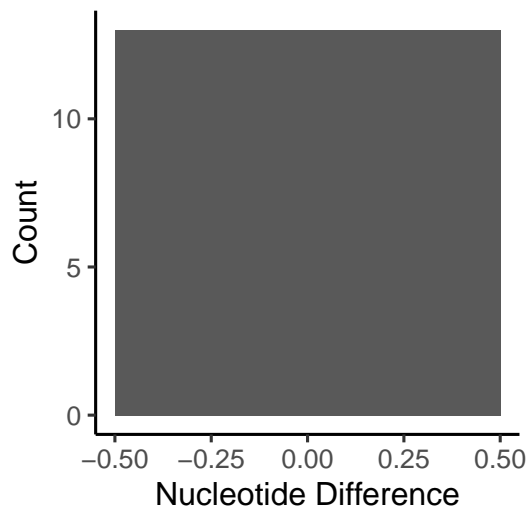
TRBV5-1*01

50 sequences assigned
50 (100%) exact matches, in which:
50 unique CDR3
11 unique J



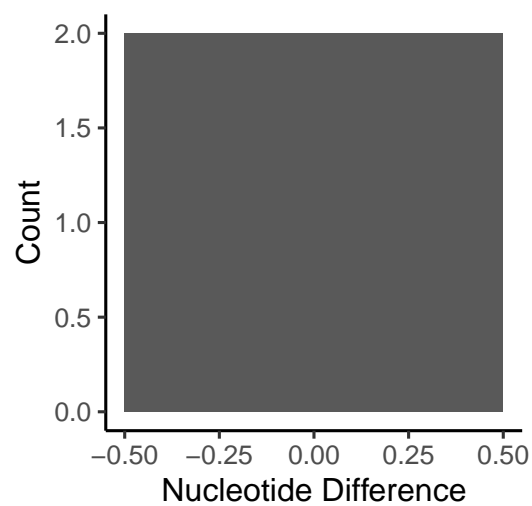
TRBV5-6*01

13 sequences assigned
13 (100%) exact matches, in which:
13 unique CDR3
8 unique J



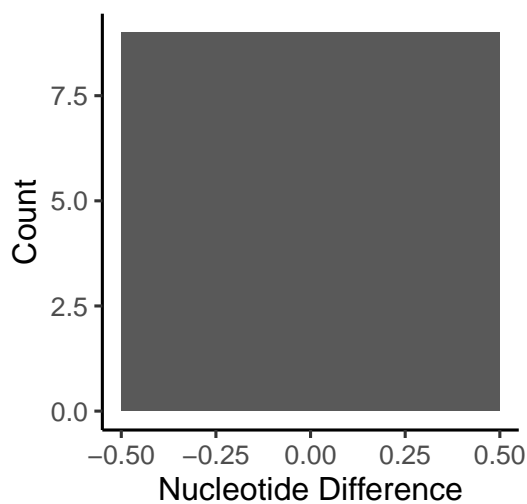
TRBV6-4*01

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



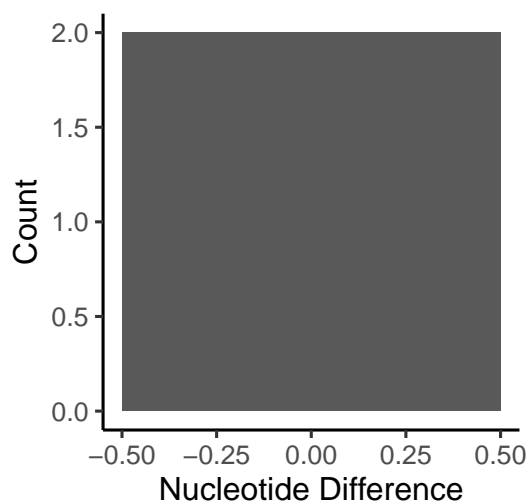
TRBV5-4*01

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



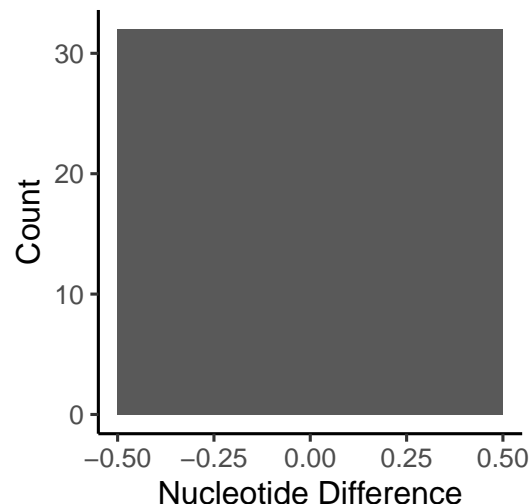
TRBV5-8*01

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



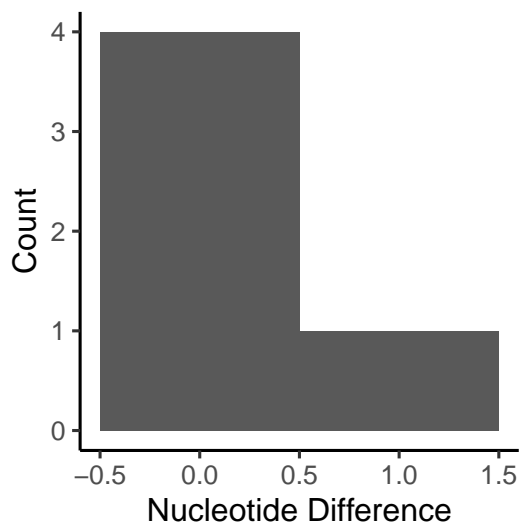
TRBV6-5*01

32 sequences assigned
32 (100%) exact matches, in which:
32 unique CDR3
13 unique J



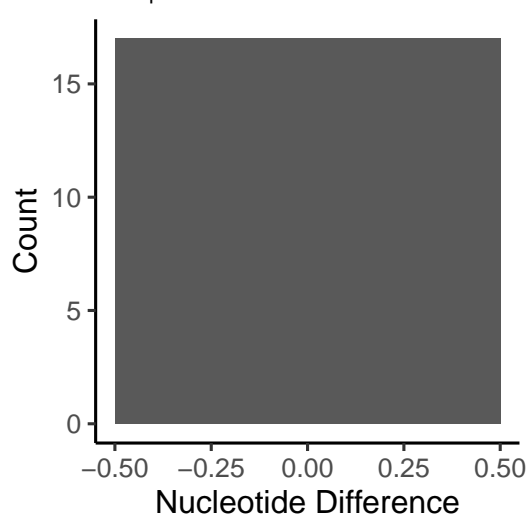
TRBV5-5*01

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



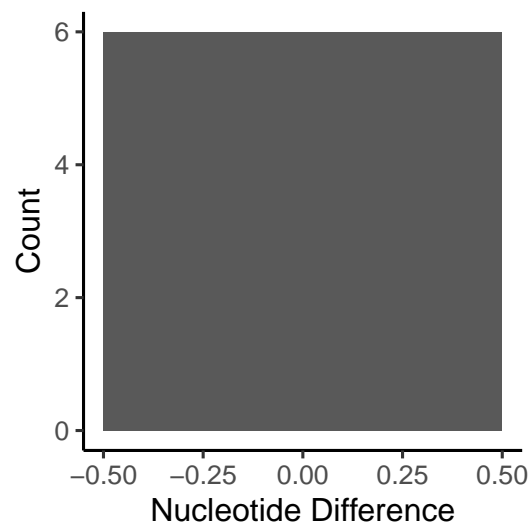
TRBV6-1*01

17 sequences assigned
17 (100%) exact matches, in which:
17 unique CDR3
7 unique J



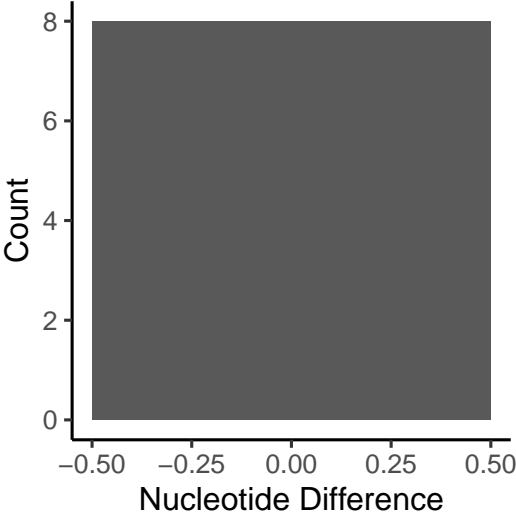
TRBV6-6*01

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



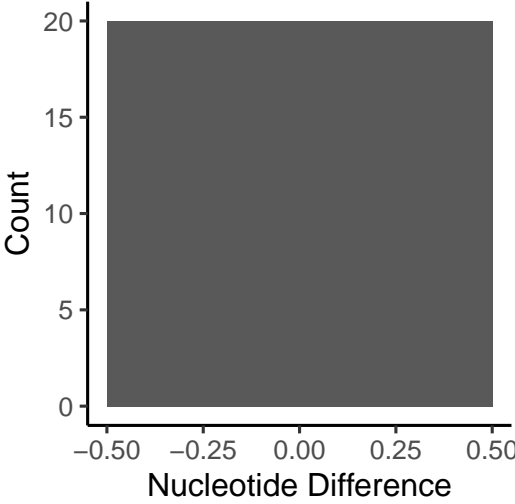
TRBV6-6*02

8 sequences assigned
8 (100%) exact matches, in which:
8 unique CDR3
7 unique J



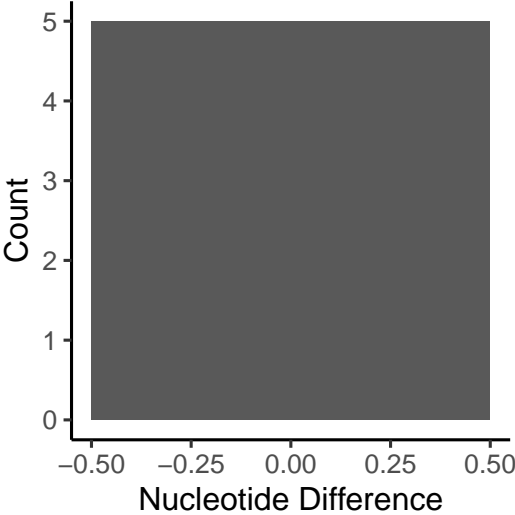
TRBV7-2*01

20 sequences assigned
20 (100%) exact matches, in which:
20 unique CDR3
9 unique J



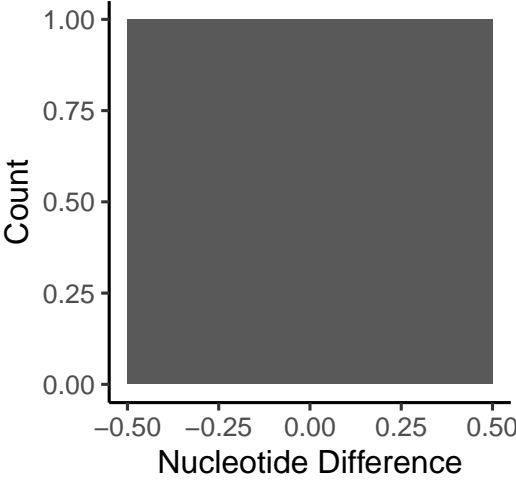
TRBV7-6*01

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



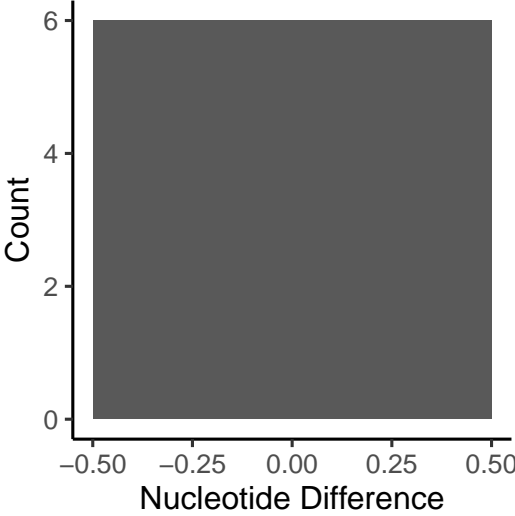
TRBV6-7*01

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



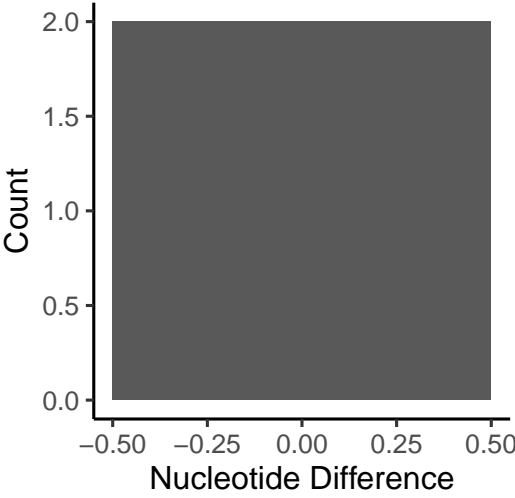
TRBV7-2*02

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



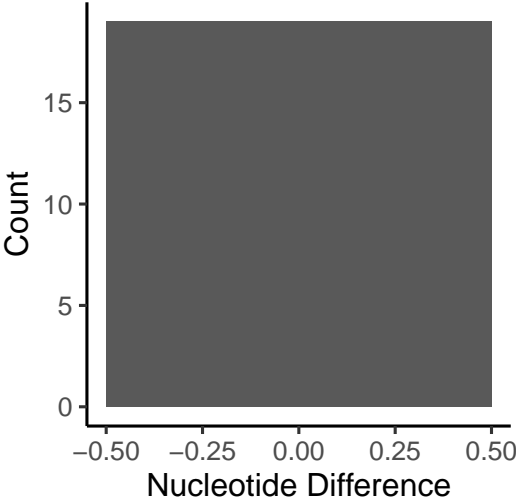
TRBV7-7*01

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



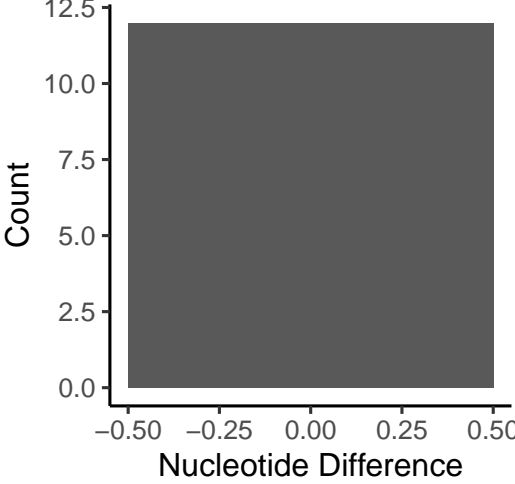
TRBV6-23*01

19 sequences assigned
19 (100%) exact matches, in which:
19 unique CDR3
10 unique J



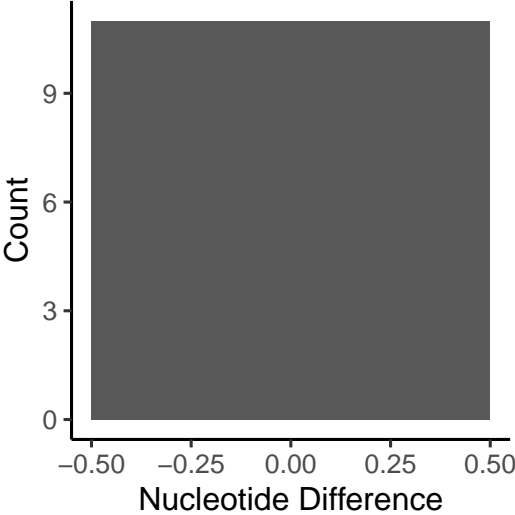
TRBV7-3*01

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



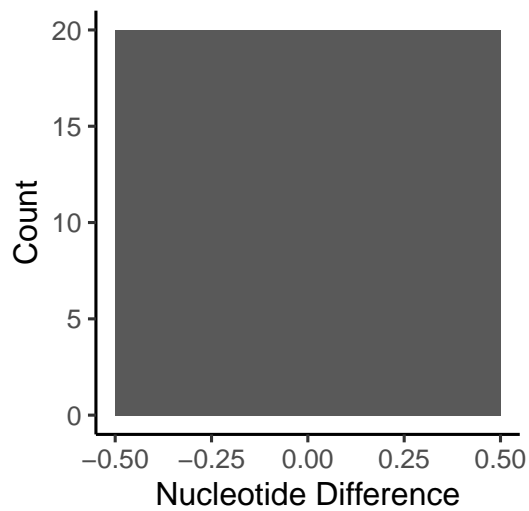
TRBV7-8*01

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



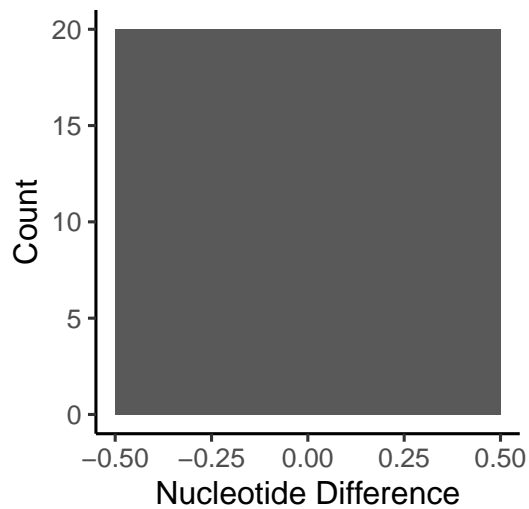
TRBV7–9*01

20 sequences assigned
20 (100%) exact matches, in which:
20 unique CDR3
8 unique J



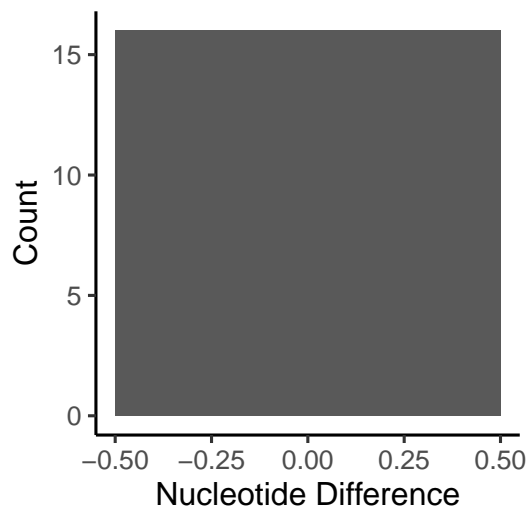
TRBV7–9*03

20 sequences assigned
20 (100%) exact matches, in which:
20 unique CDR3
7 unique J

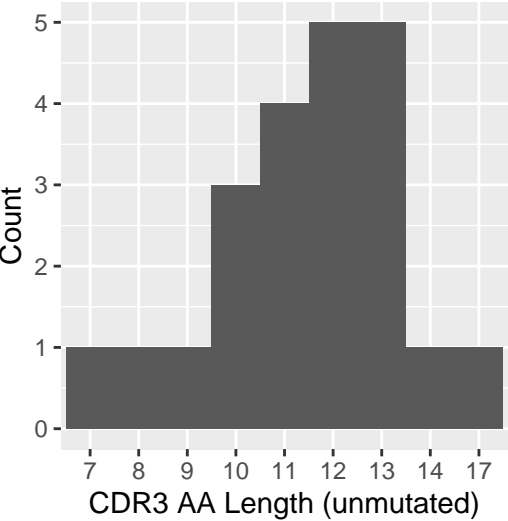


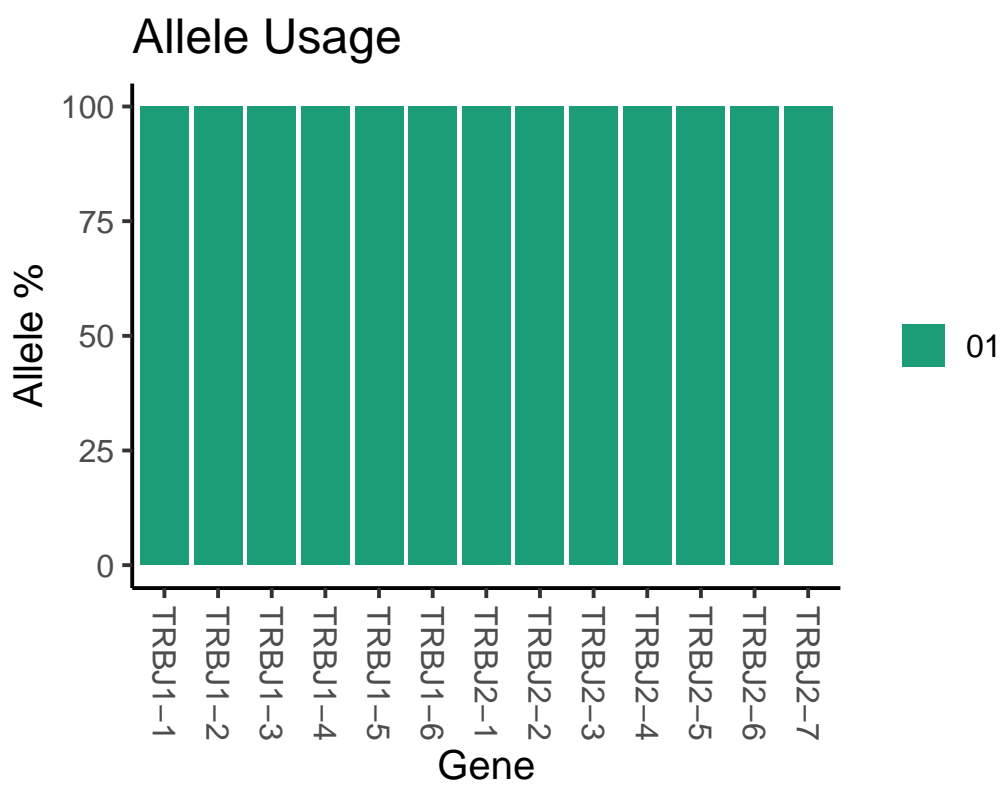
TRBV29–1*01

16 sequences assigned
16 (100%) exact matches, in which:
16 unique CDR3
6 unique J



TRBV20-1*02_319AGAGA323





Warning – no inferred sequences found.

Warning – no inferred sequences found.