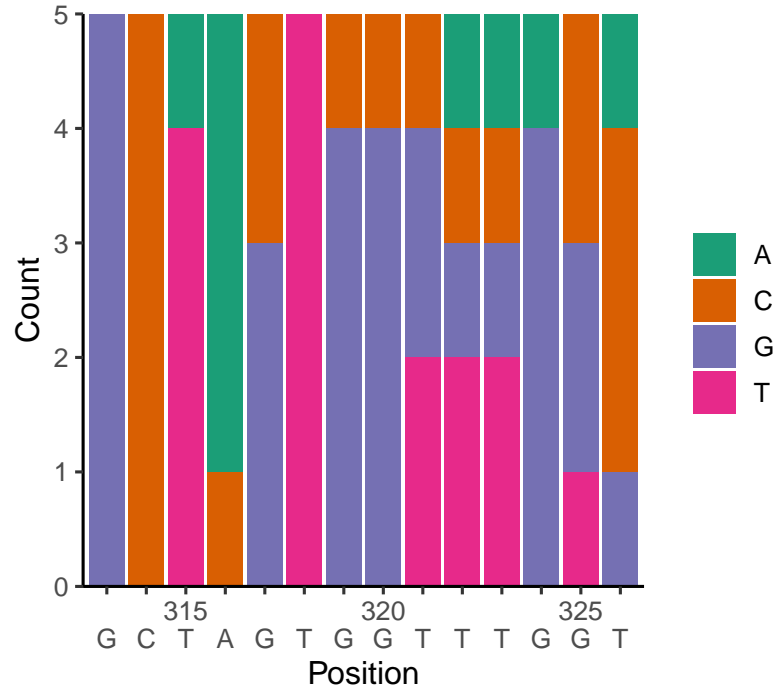
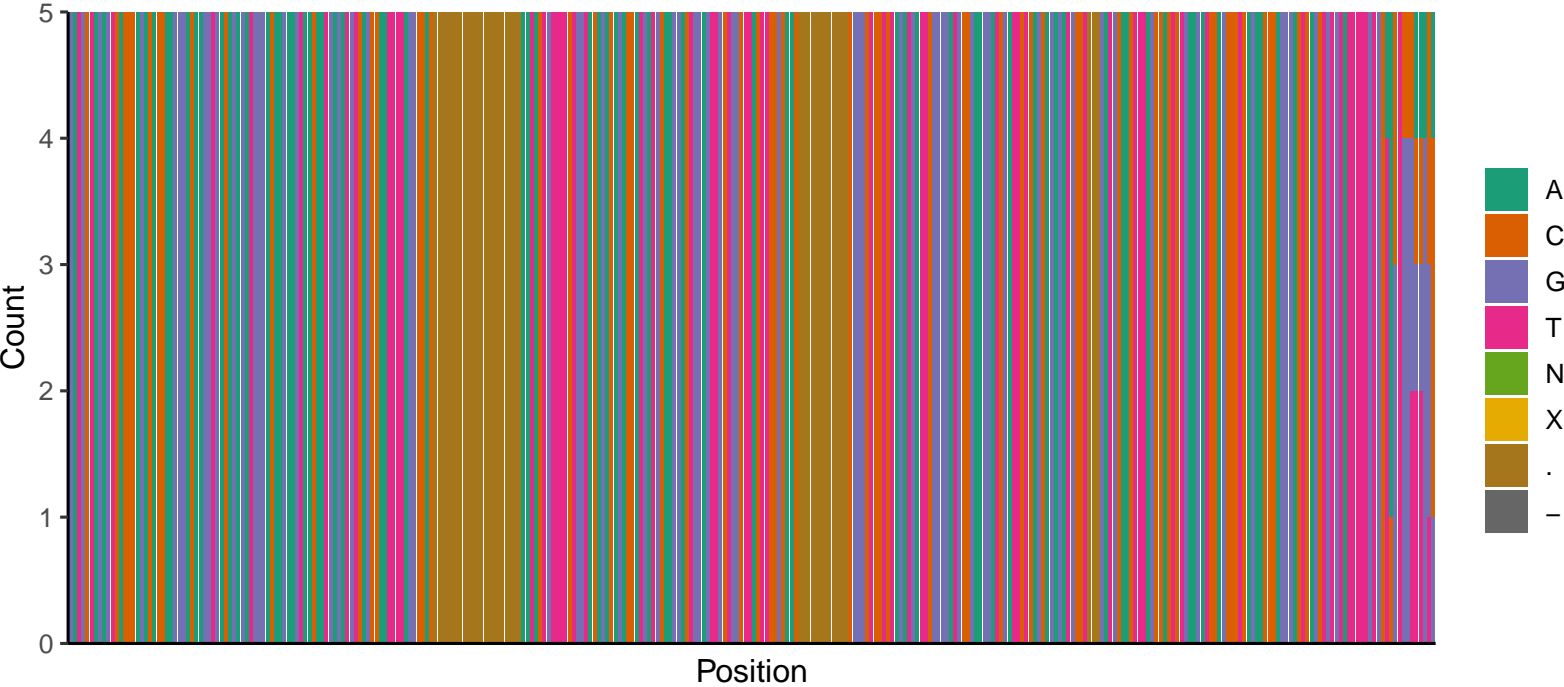


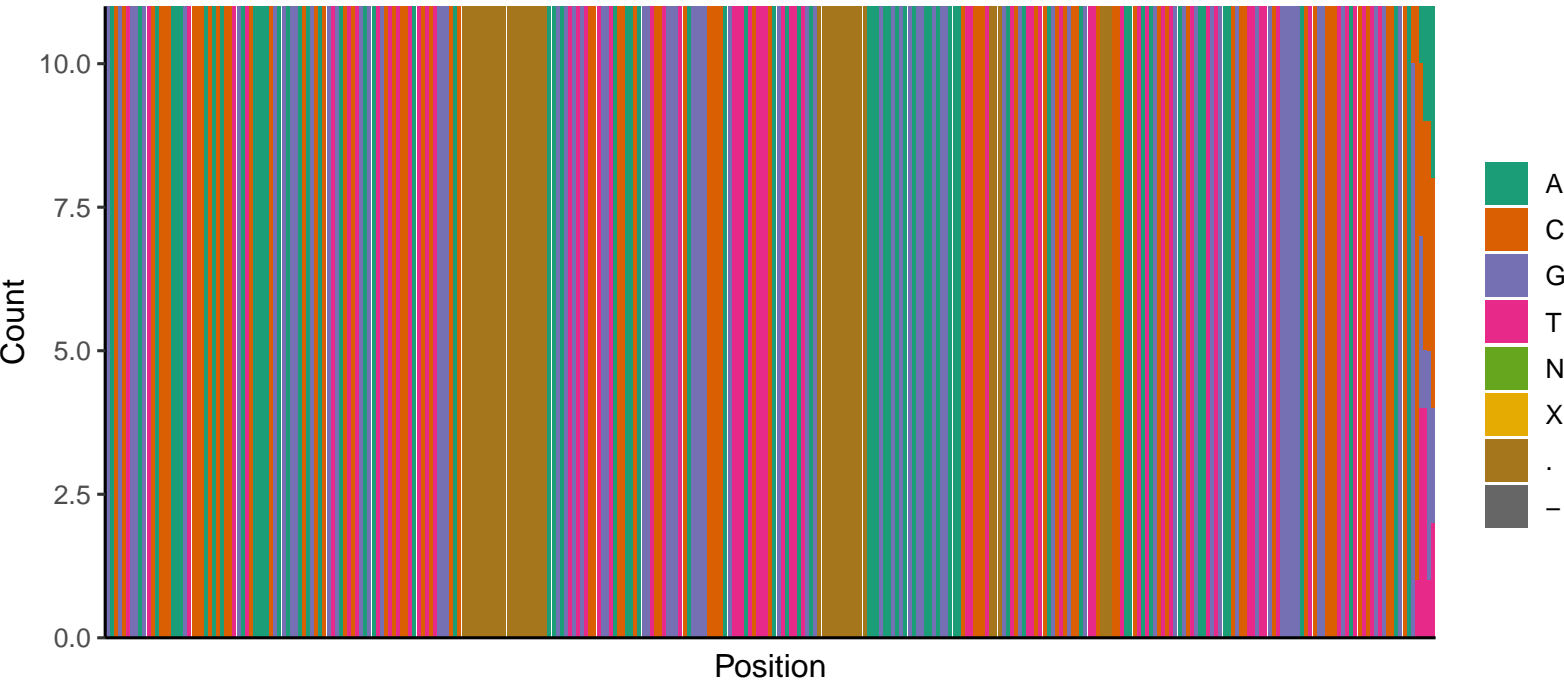
Gene TRBV12-5*01_C28G



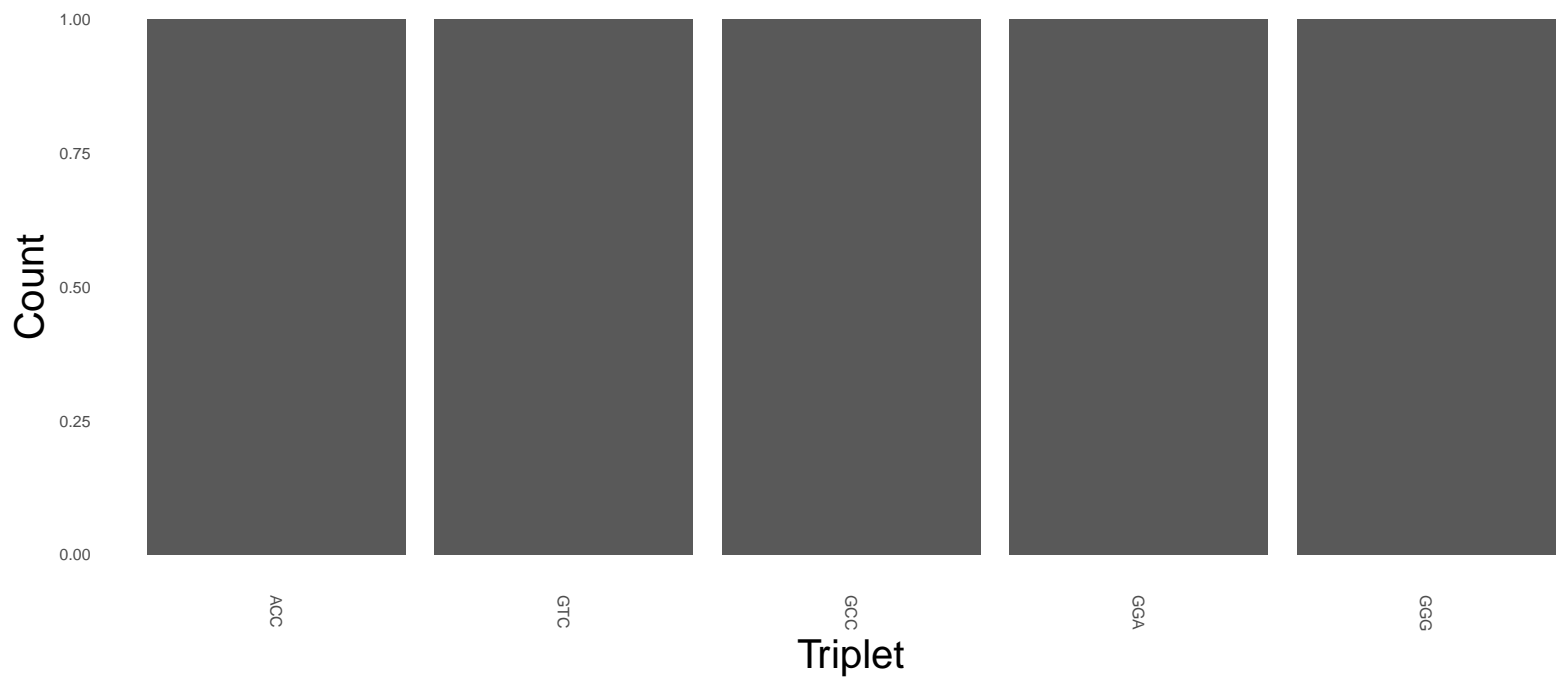
Gene TRBV12-5*01_C28G



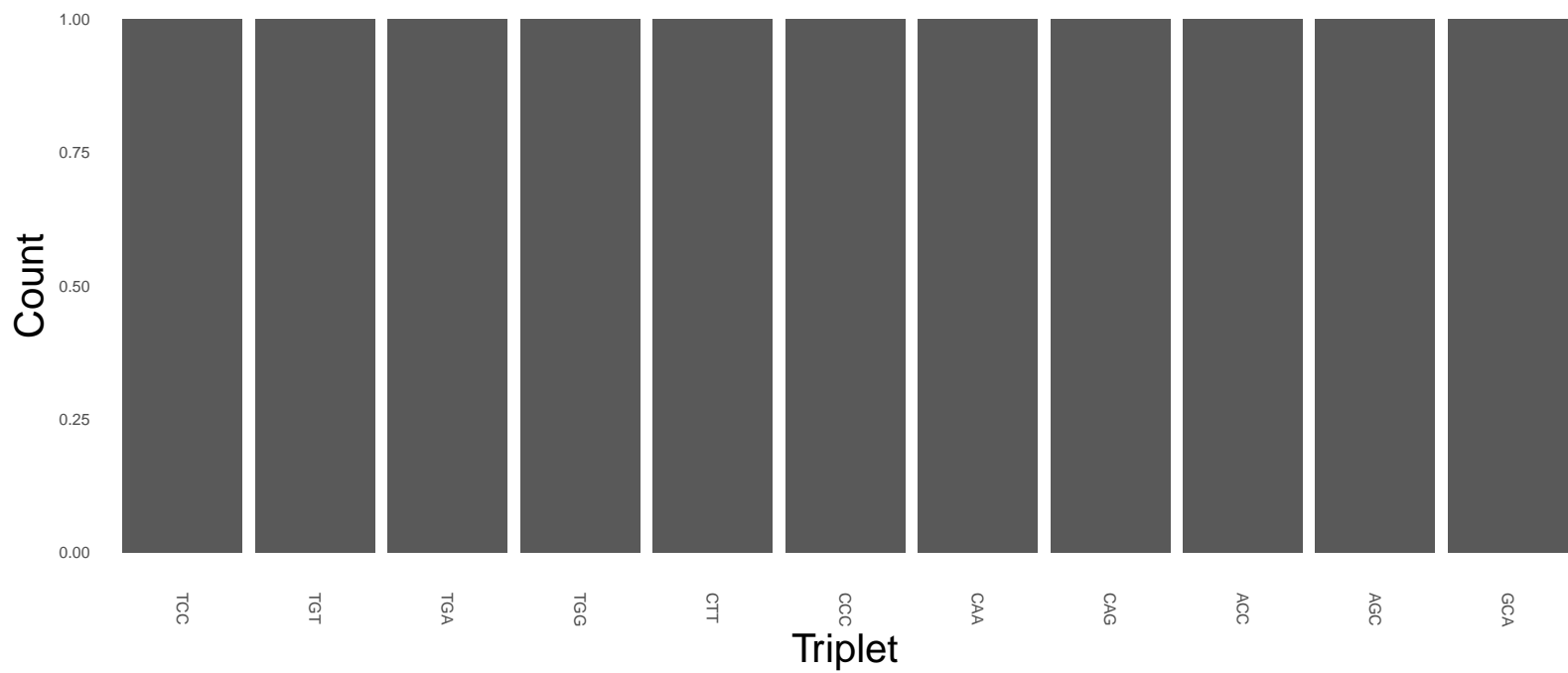
Gene TRBV5-5*02_322TTGG325



TRBV12–5*01_C28G– Final 3 nucleotides as a triplet

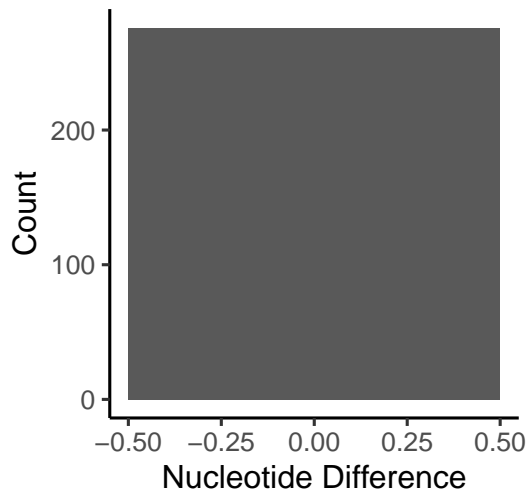


TRBV5–5*02_322TTGG325– Final 3 nucleotides as a triplet



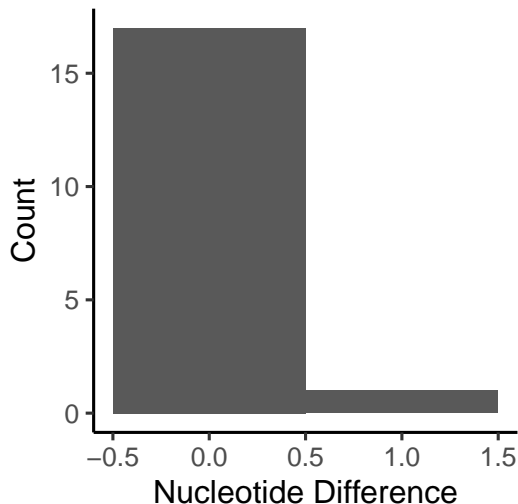
TRBV20–1*01

276 sequences assigned
276 (100%) exact matches, in which:
275 unique CDR3
13 unique J



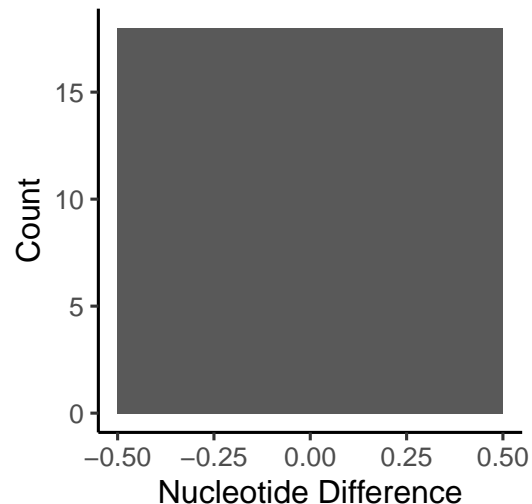
TRBV10–2*01

18 sequences assigned
17 (94.4%) exact matches, in which:
17 unique CDR3
10 unique J



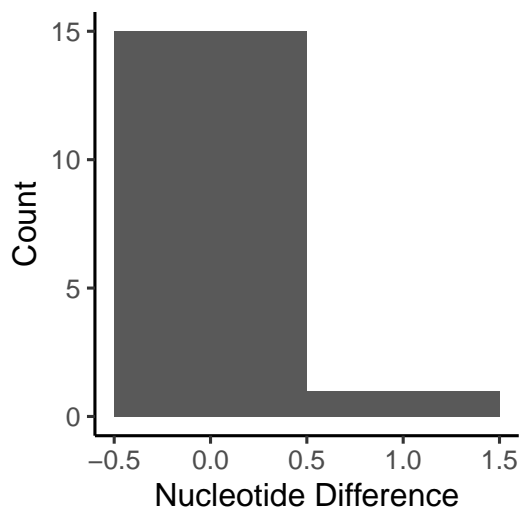
TRBV13*01

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



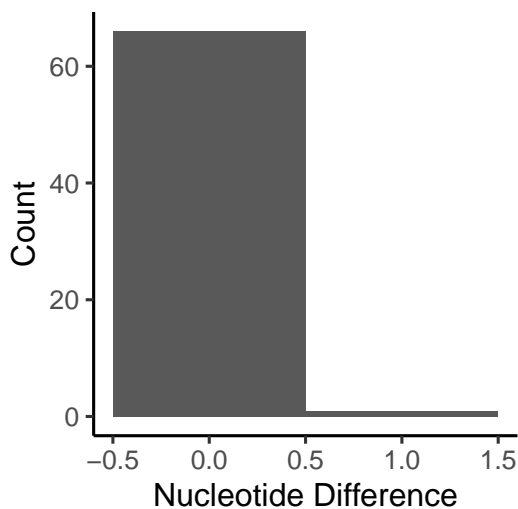
TRBV10–1*01

16 sequences assigned
15 (93.8%) exact matches, in which:
13 unique CDR3
6 unique J



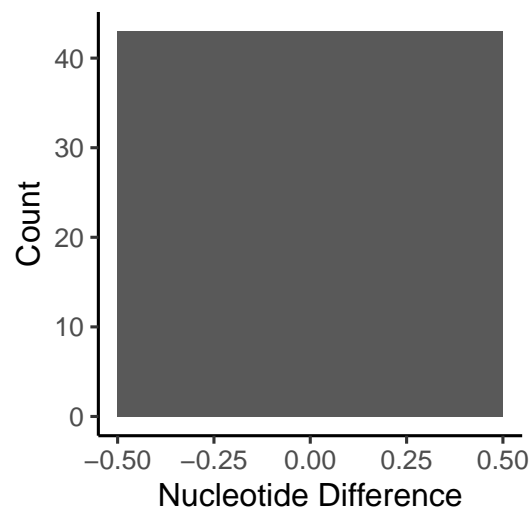
TRBV10–3*01

67 sequences assigned
66 (98.5%) exact matches, in which:
66 unique CDR3
12 unique J



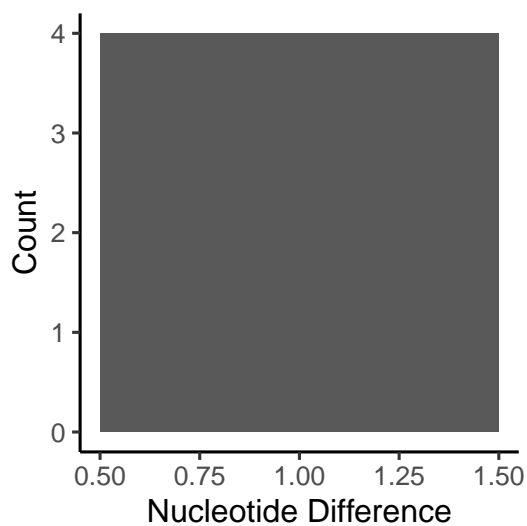
TRBV14*01

43 sequences assigned
43 (100%) exact matches, in which:
42 unique CDR3
10 unique J



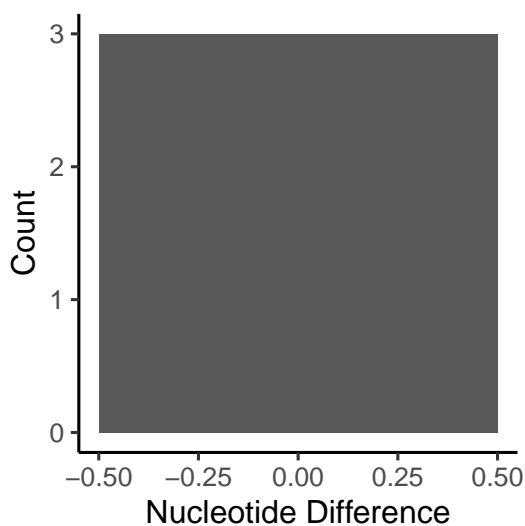
TRBV10–1*02

4 sequences assigned
No exact matches.



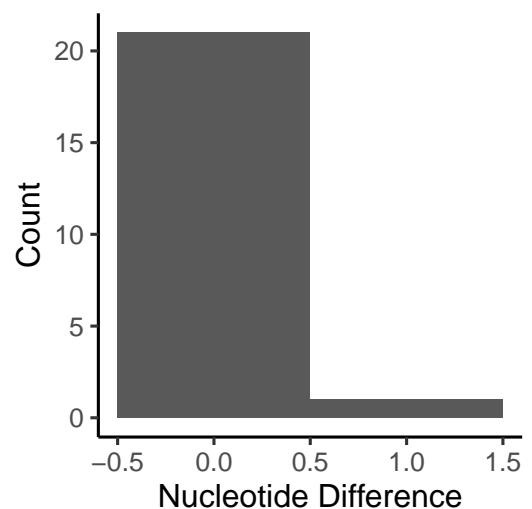
TRBV1*01

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



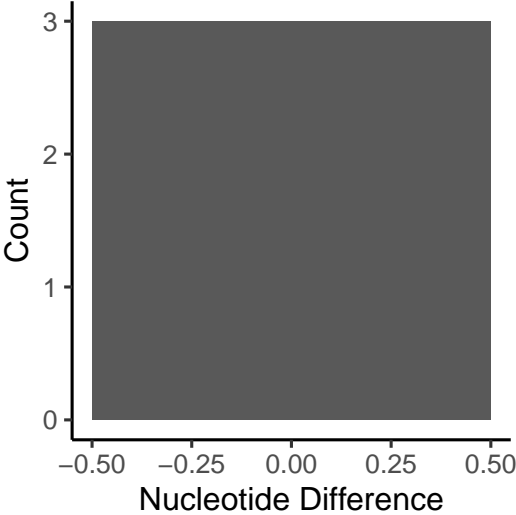
TRBV15*02

22 sequences assigned
21 (95.5%) exact matches, in which:
21 unique CDR3
10 unique J



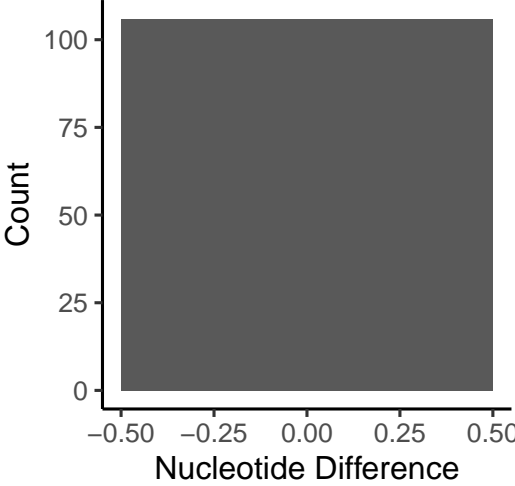
TRBV16*01

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



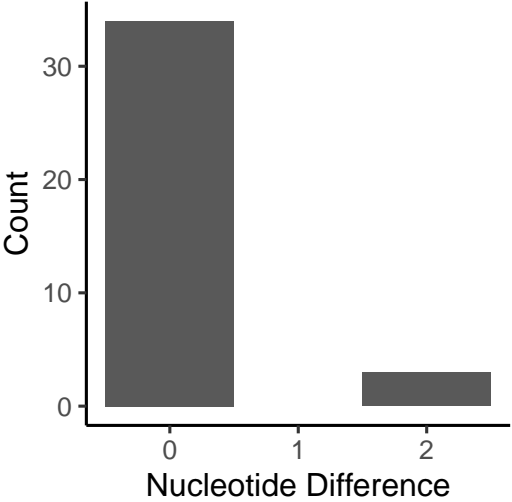
TRBV2*01

106 sequences assigned
106 (100%) exact matches, in which:
105 unique CDR3
13 unique J



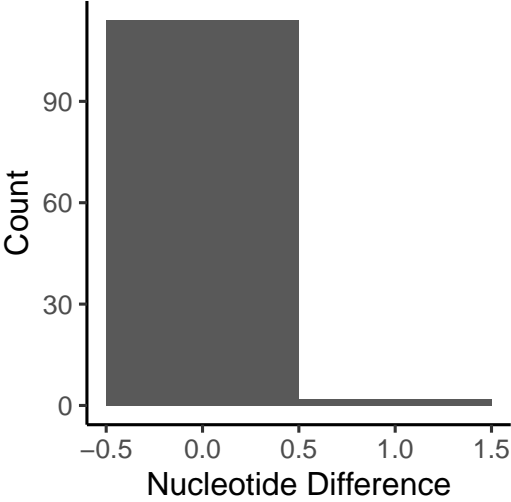
TRBV30*02

39 sequences assigned
34 (87.2%) exact matches, in which:
34 unique CDR3
11 unique J



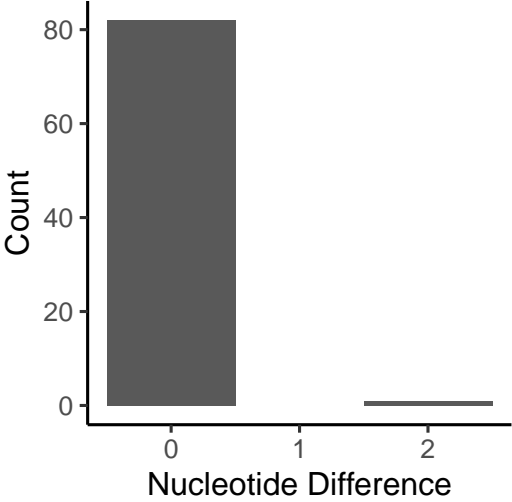
TRBV18*01

116 sequences assigned
114 (98.3%) exact matches, in which:
109 unique CDR3
13 unique J



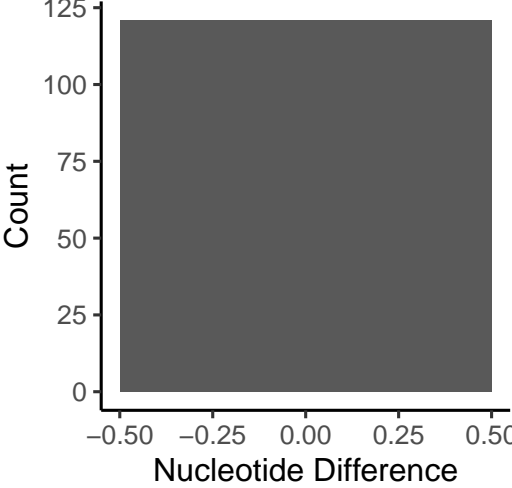
TRBV27*01

83 sequences assigned
82 (98.8%) exact matches, in which:
82 unique CDR3
13 unique J



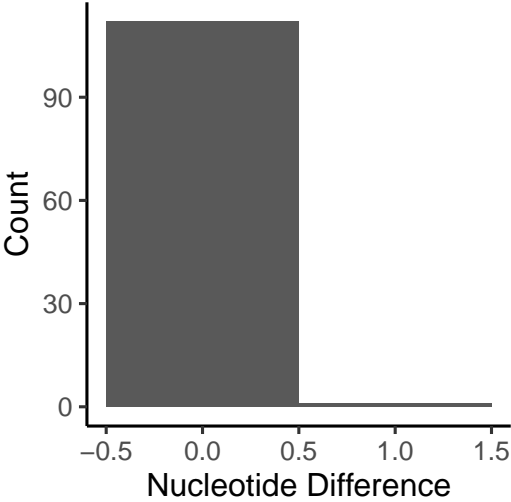
TRBV9*01

121 sequences assigned
121 (100%) exact matches, in which:
121 unique CDR3
13 unique J



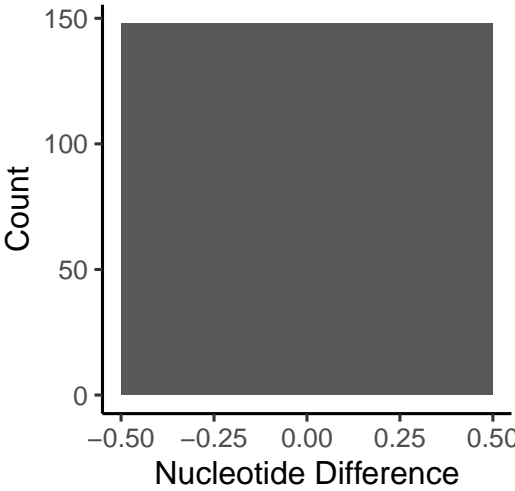
TRBV19*01

113 sequences assigned
112 (99.1%) exact matches, in which:
112 unique CDR3
13 unique J



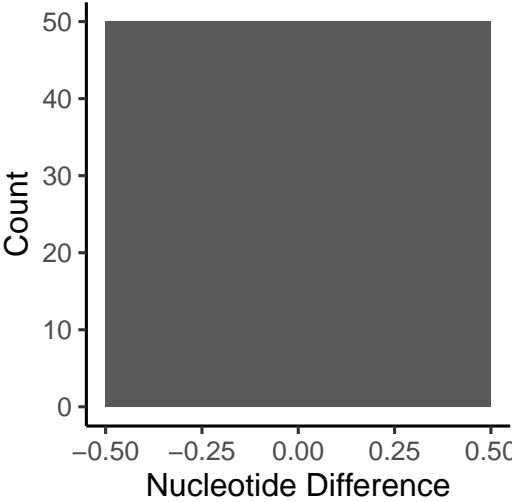
TRBV28*01

148 sequences assigned
148 (100%) exact matches, in which:
146 unique CDR3
12 unique J



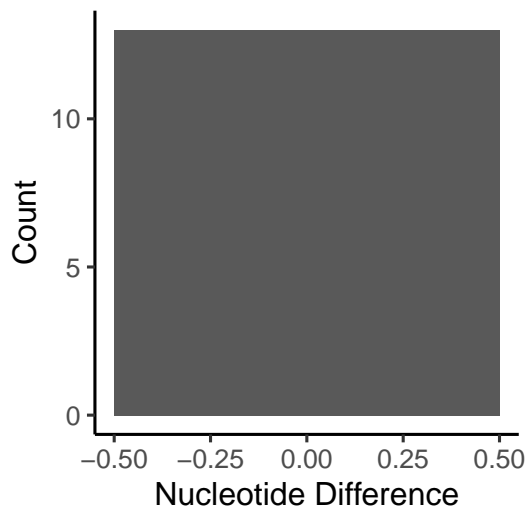
TRBV21-1*01

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



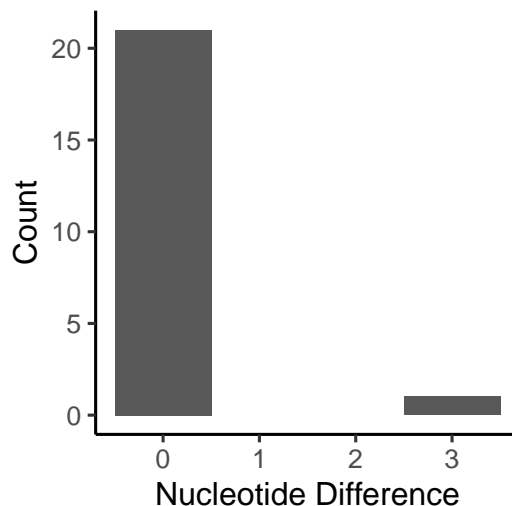
TRBV11-1*01

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



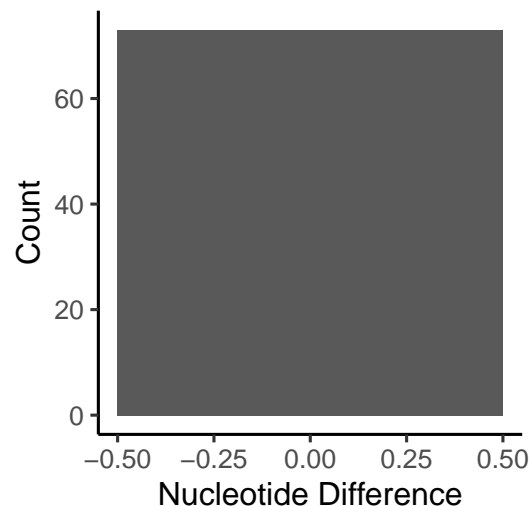
TRBV11-3*01

22 sequences assigned
21 (95.5%) exact matches, in which:
21 unique CDR3
9 unique J



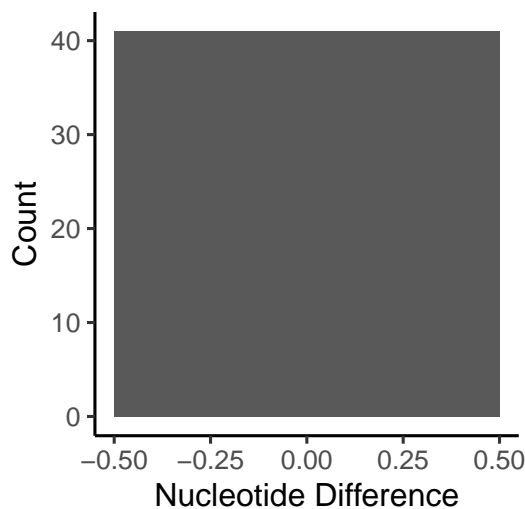
TRBV12-3*01

73 sequences assigned
73 (100%) exact matches, in which:
73 unique CDR3
13 unique J



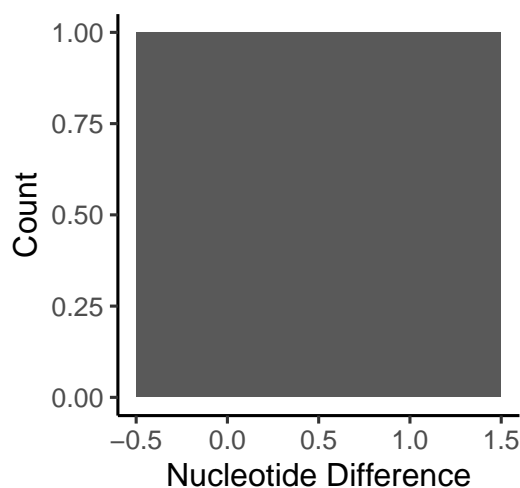
TRBV11-2*01

41 sequences assigned
41 (100%) exact matches, in which:
41 unique CDR3
11 unique J



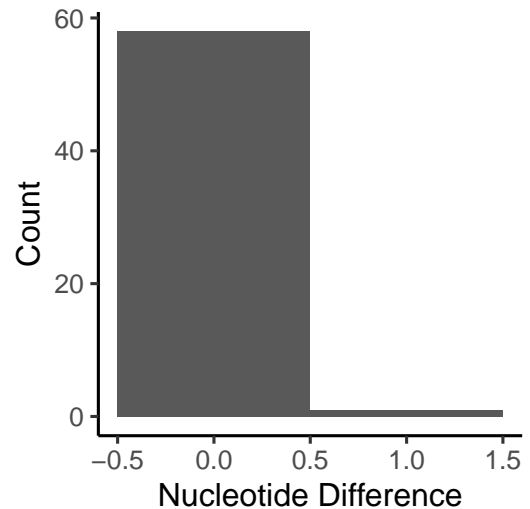
TRBV12-1*01

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



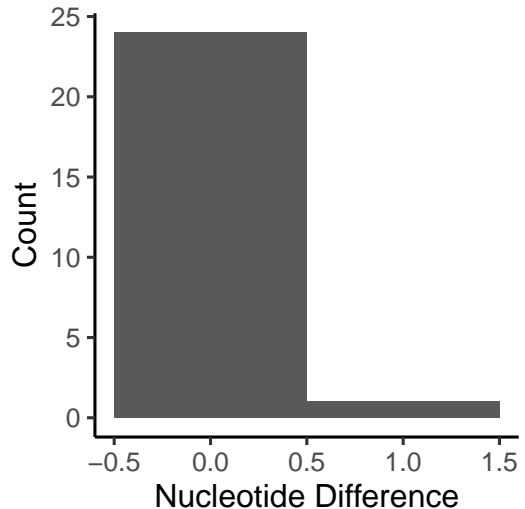
TRBV12-4*01

59 sequences assigned
58 (98.3%) exact matches, in which:
58 unique CDR3
12 unique J



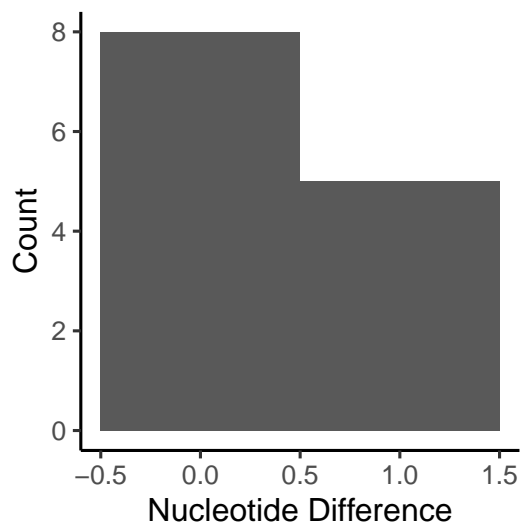
TRBV11-2*03

25 sequences assigned
24 (96%) exact matches, in which:
22 unique CDR3
10 unique J



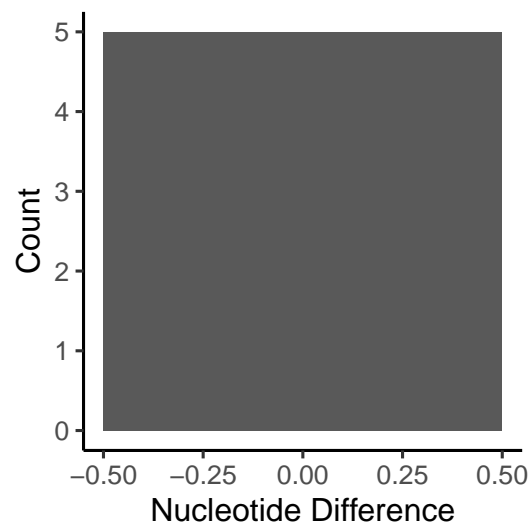
TRBV12-2*01

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



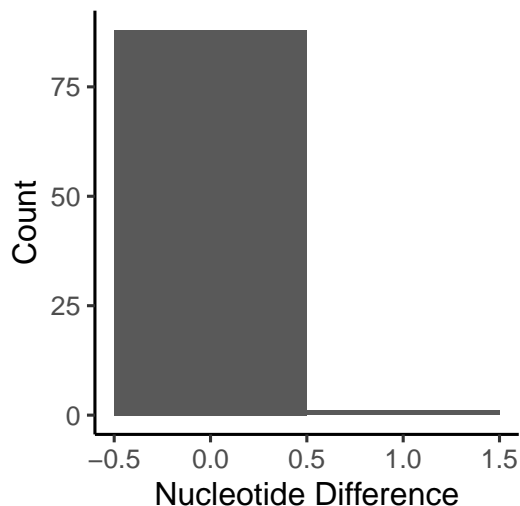
TRBV12-5*01_C28G

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



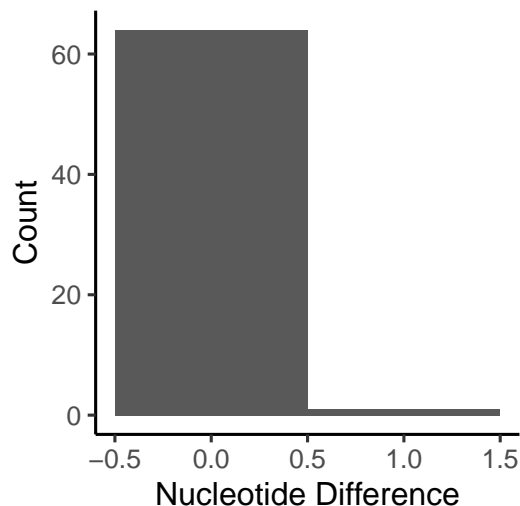
TRBV3-1*01

89 sequences assigned
88 (98.9%) exact matches, in which:
88 unique CDR3
11 unique J



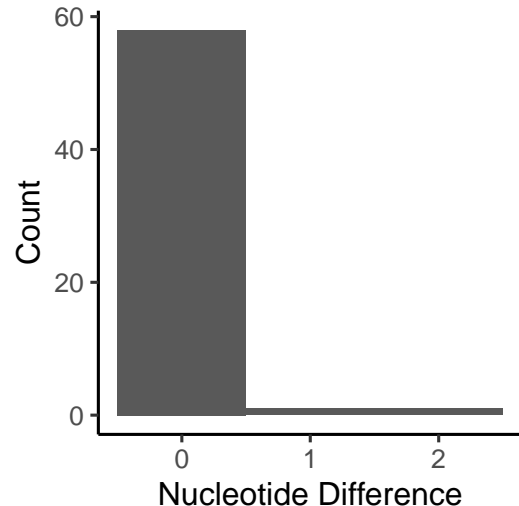
TRBV4-1*01

65 sequences assigned
64 (98.5%) exact matches, in which:
64 unique CDR3
13 unique J



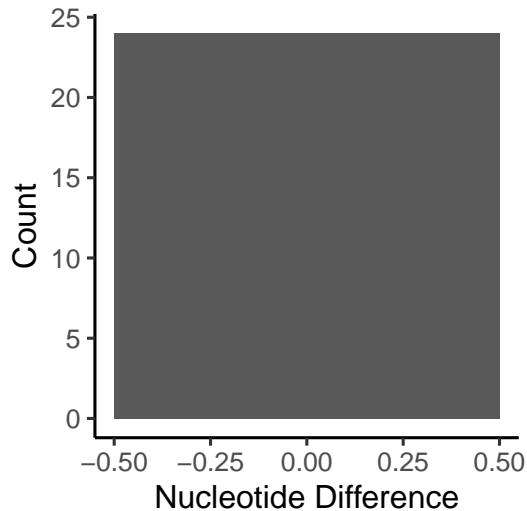
TRBV4-3*01

60 sequences assigned
58 (96.7%) exact matches, in which:
58 unique CDR3
10 unique J



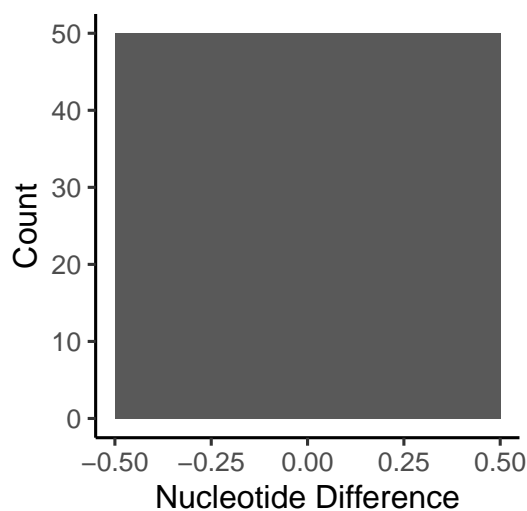
TRBV23-1*01

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



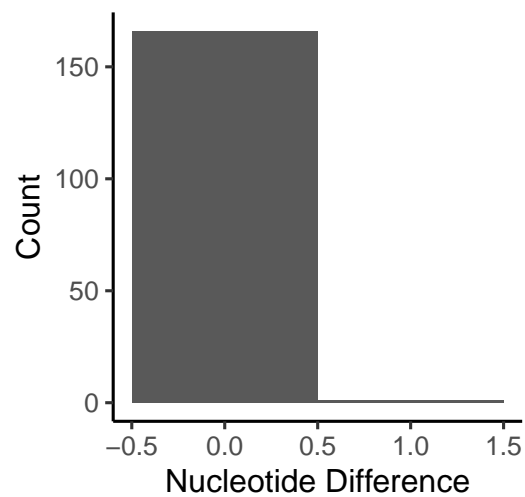
TRBV24-1*01

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



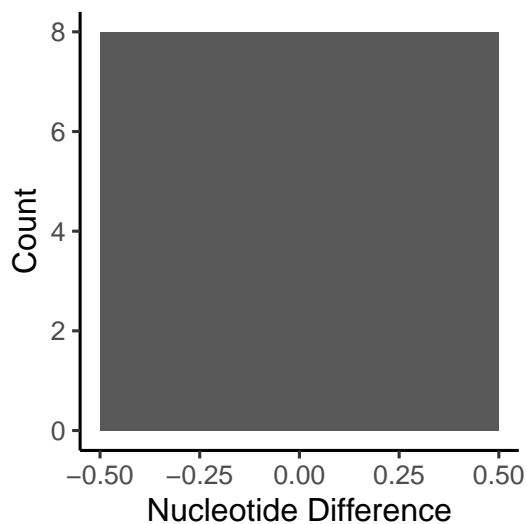
TRBV5-1*01

167 sequences assigned
166 (99.4%) exact matches, in which:
166 unique CDR3
13 unique J



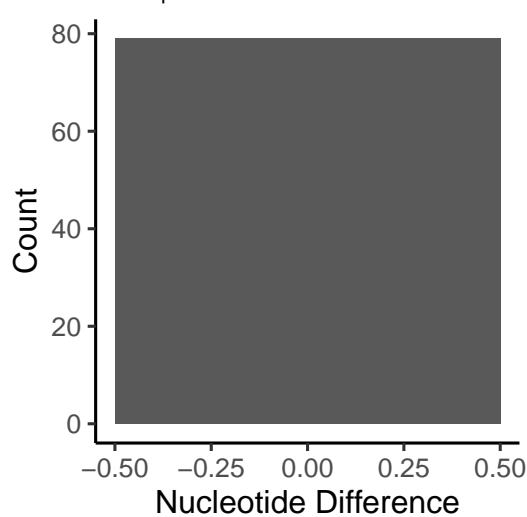
TRBV3-2*03

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



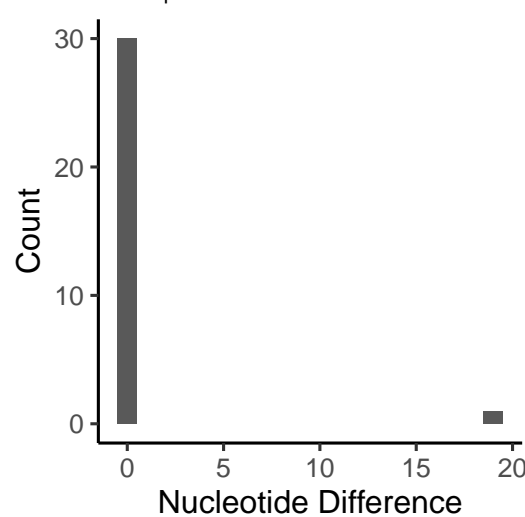
TRBV4-2*01

79 sequences assigned
79 (100%) exact matches, in which:
79 unique CDR3
12 unique J



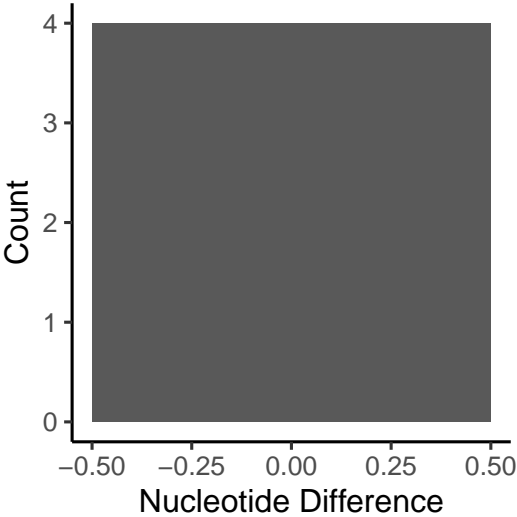
TRBV25-1*01

31 sequences assigned
30 (96.8%) exact matches, in which:
30 unique CDR3
13 unique J



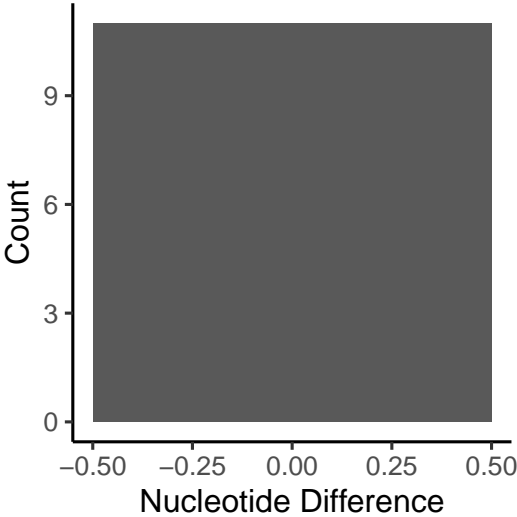
TRBV5-3*01

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



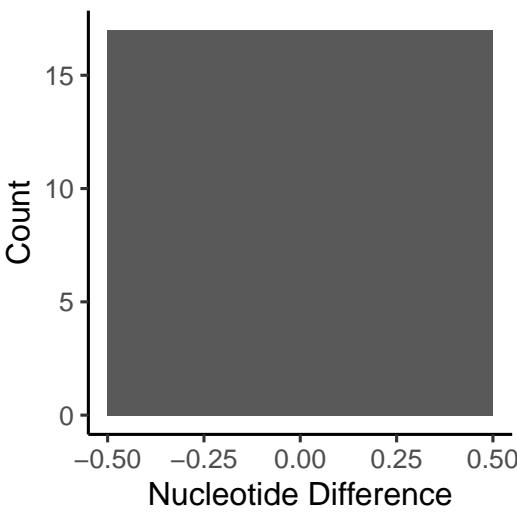
TRBV5-5*02_322TTGG32!

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



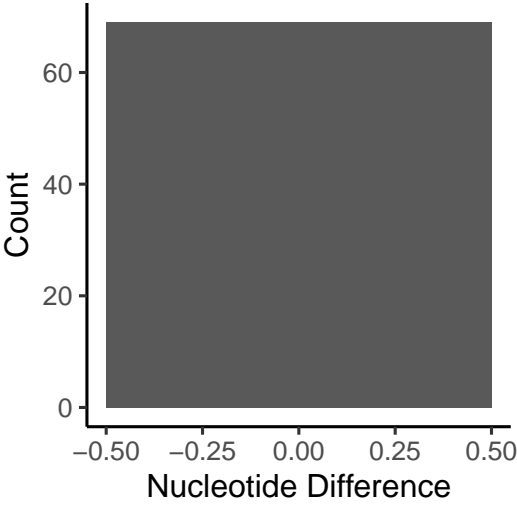
TRBV5-8*01

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



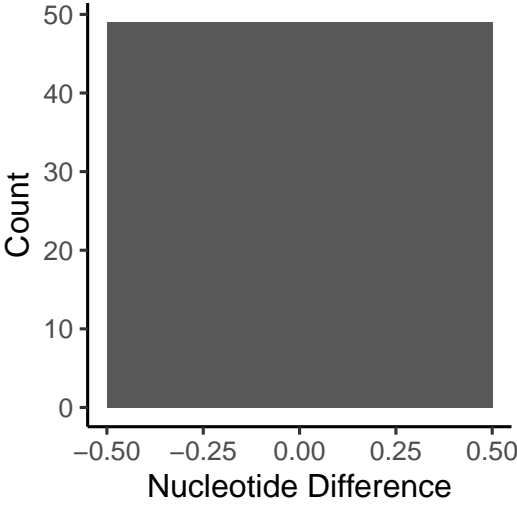
TRBV5-4*01

69 sequences assigned
69 (100%) exact matches, in which:
69 unique CDR3
12 unique J



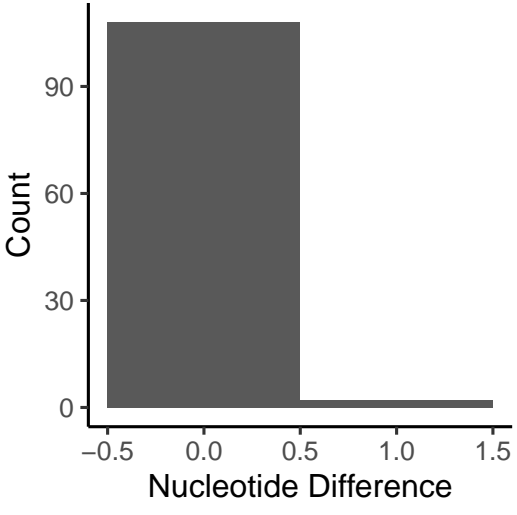
TRBV5-6*01

49 sequences assigned
49 (100%) exact matches, in which:
49 unique CDR3
11 unique J



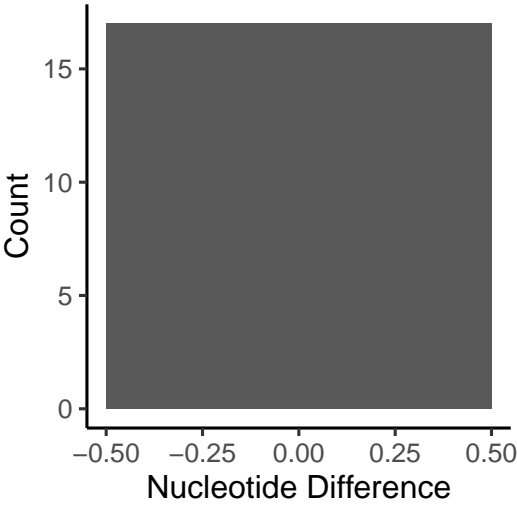
TRBV6-1*01

110 sequences assigned
108 (98.2%) exact matches, in which:
107 unique CDR3
12 unique J



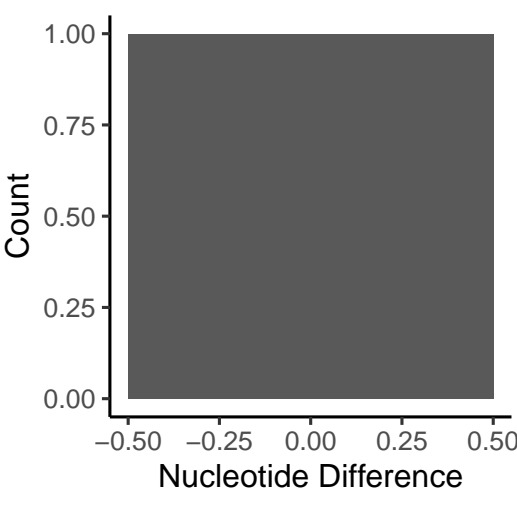
TRBV5-5*01

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



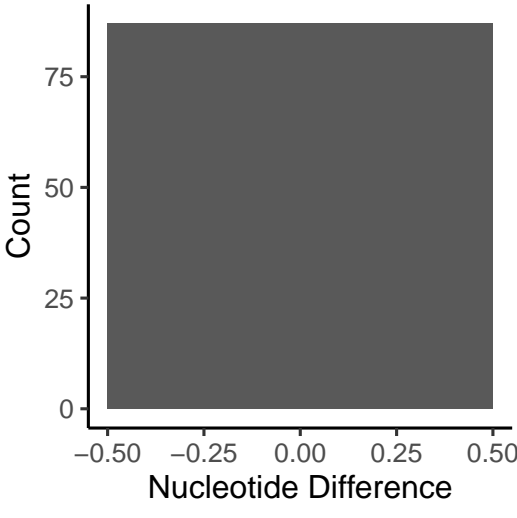
TRBV5-7*01

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



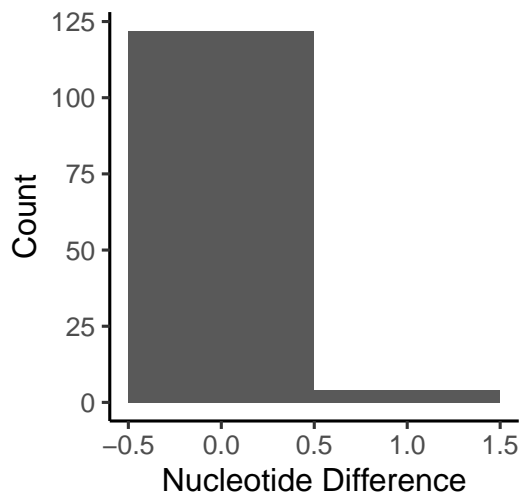
TRBV6-4*01

87 sequences assigned
87 (100%) exact matches, in which:
86 unique CDR3
13 unique J



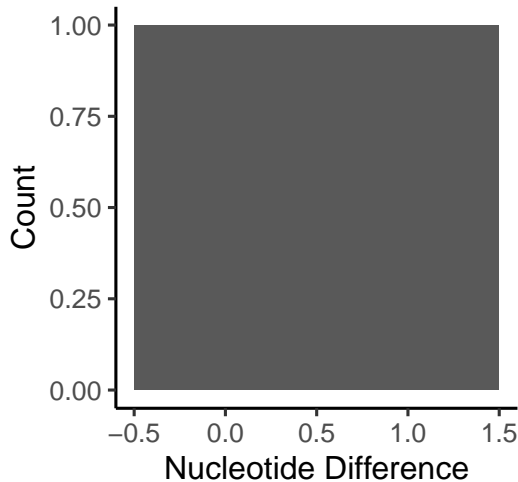
TRBV6–5*01

126 sequences assigned
122 (96.8%) exact matches, in which:
121 unique CDR3
13 unique J



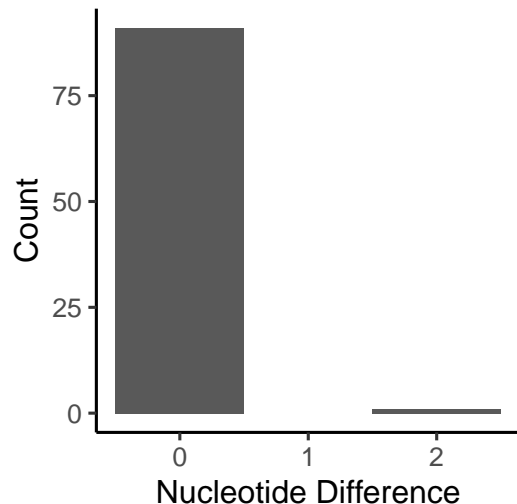
TRBV6–7*01

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



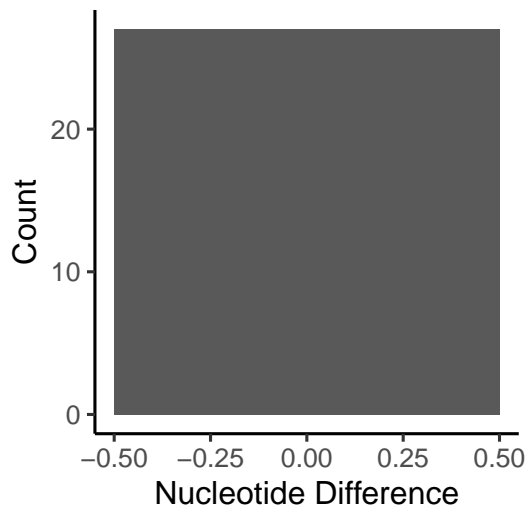
TRBV6–23*01

92 sequences assigned
91 (98.9%) exact matches, in which:
90 unique CDR3
12 unique J



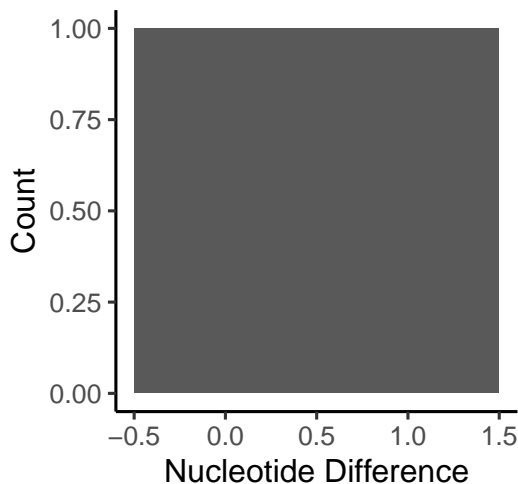
TRBV6–6*01

27 sequences assigned
27 (100%) exact matches, in which:
27 unique CDR3
9 unique J



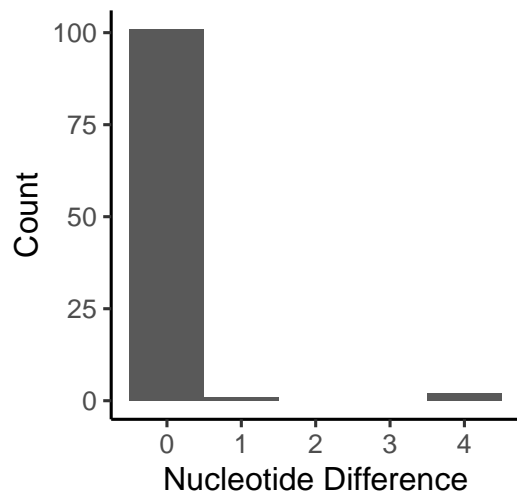
TRBV6–8*01

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



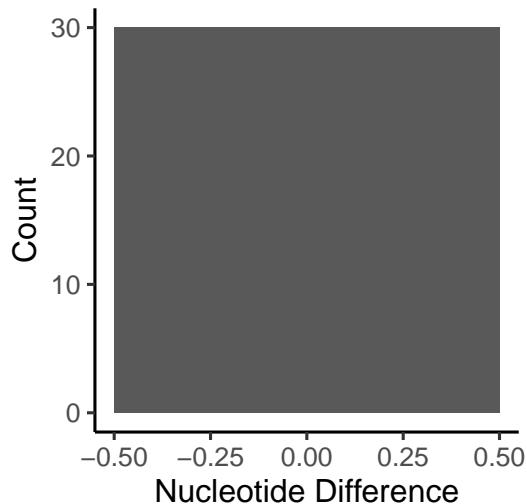
TRBV7–2*01

104 sequences assigned
101 (97.1%) exact matches, in which:
101 unique CDR3
13 unique J



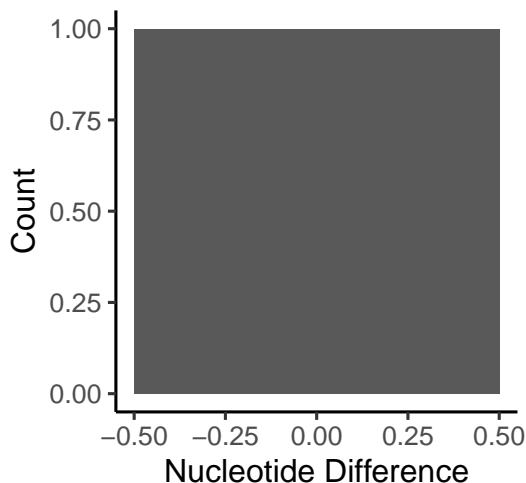
TRBV6–6*02

30 sequences assigned
30 (100%) exact matches, in which:
30 unique CDR3
8 unique J



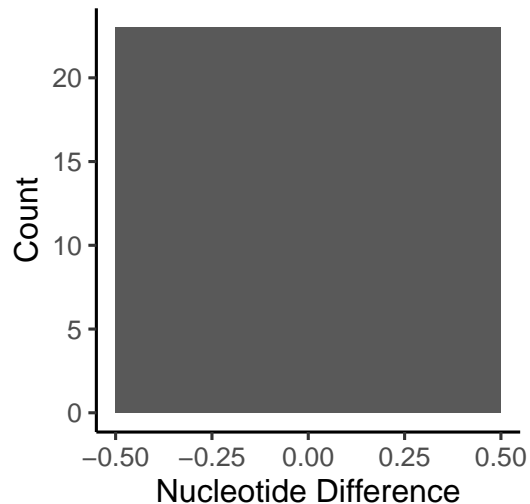
TRBV6–9*01

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



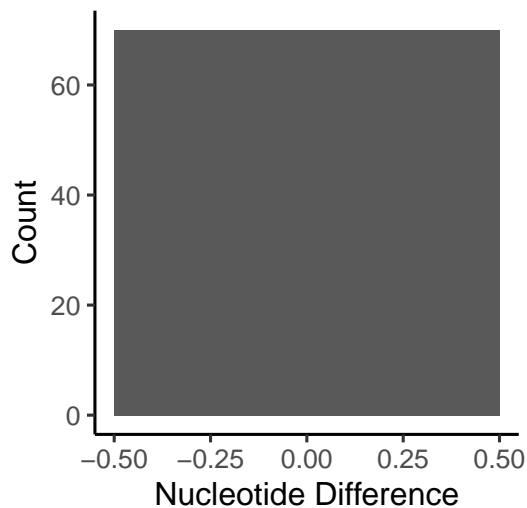
TRBV7–2*02

23 sequences assigned
23 (100%) exact matches, in which:
23 unique CDR3
11 unique J



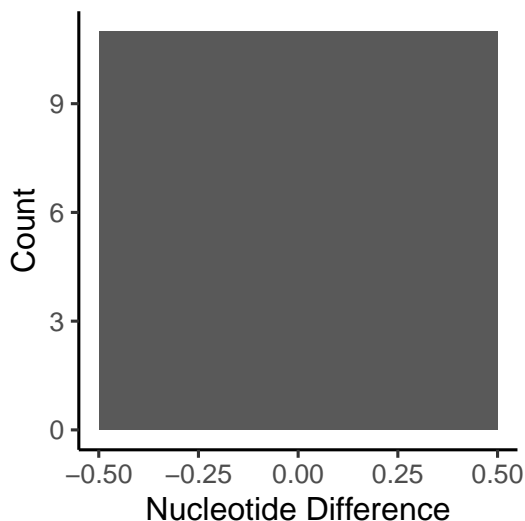
TRBV7-3*01

70 sequences assigned
70 (100%) exact matches, in which:
68 unique CDR3
12 unique J



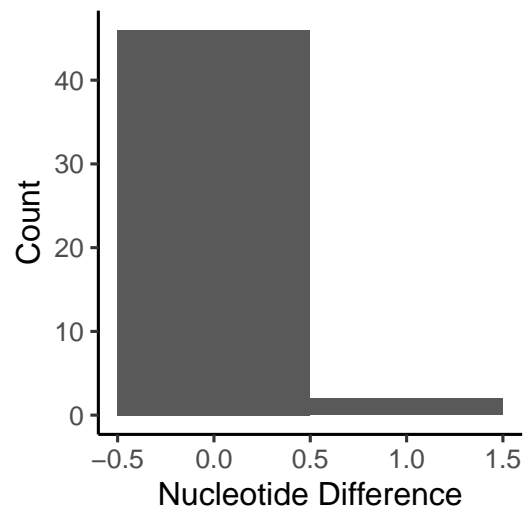
TRBV7-7*01

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



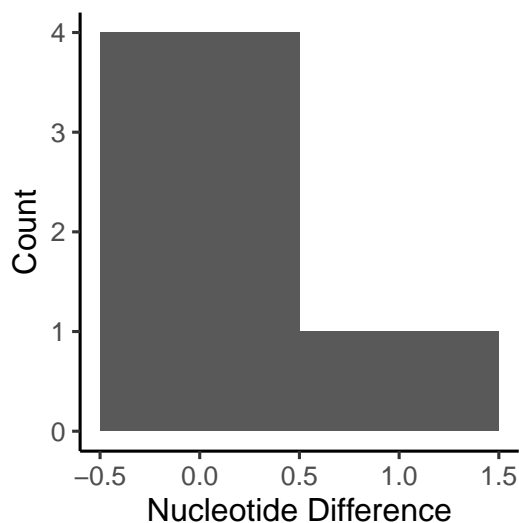
TRBV29-1*01

48 sequences assigned
46 (95.8%) exact matches, in which:
46 unique CDR3
10 unique J



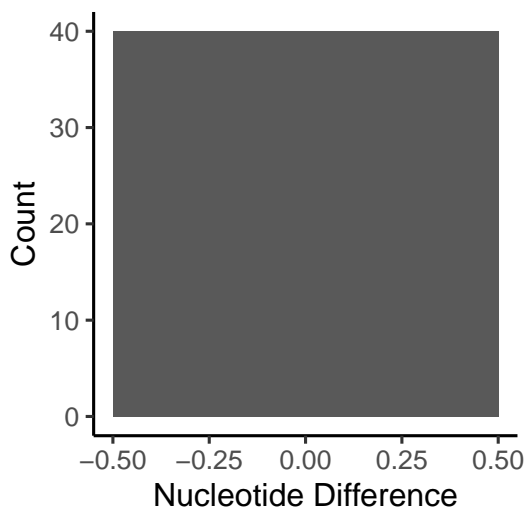
TRBV7-4*01

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



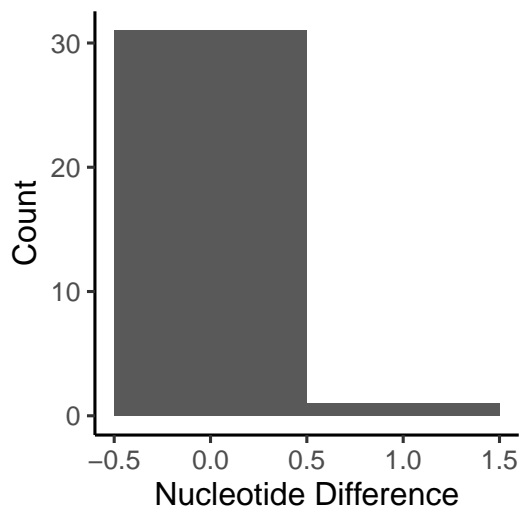
TRBV7-8*01

40 sequences assigned
40 (100%) exact matches, in which:
40 unique CDR3
13 unique J



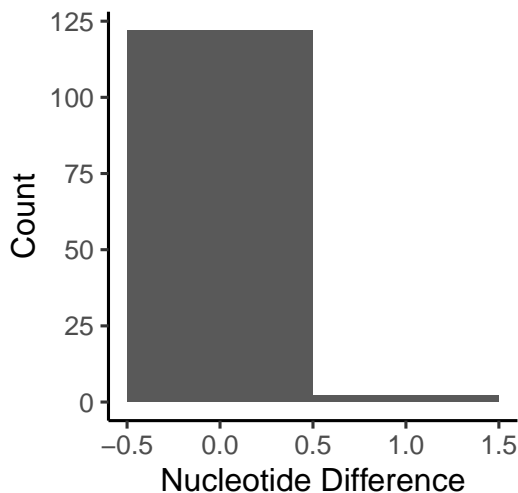
TRBV7-6*01

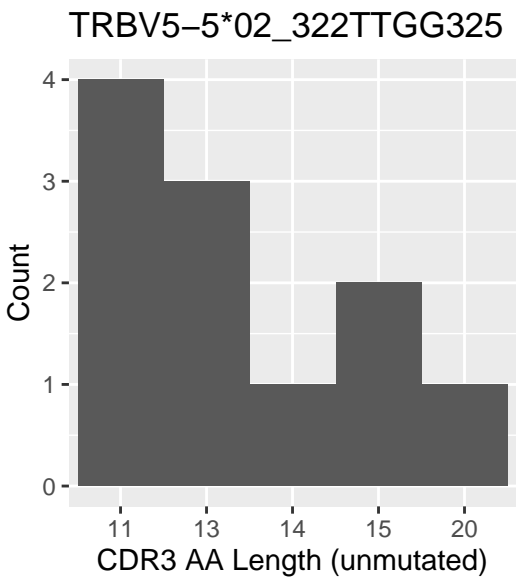
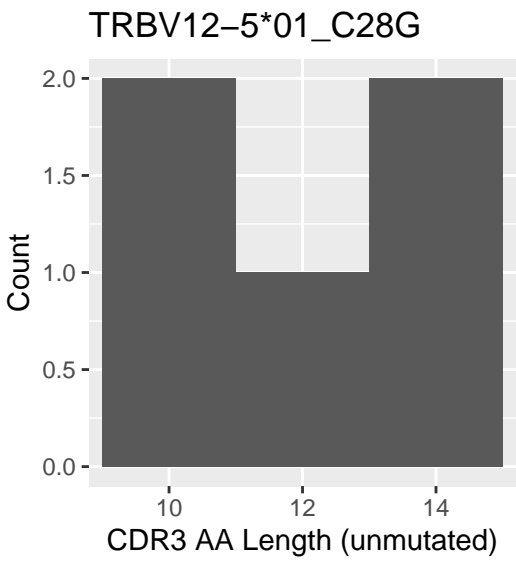
32 sequences assigned
31 (96.9%) exact matches, in which:
30 unique CDR3
12 unique J



TRBV7-9*01

124 sequences assigned
122 (98.4%) exact matches, in which:
121 unique CDR3
13 unique J







Warning – no inferred sequences found.

Warning – no inferred sequences found.