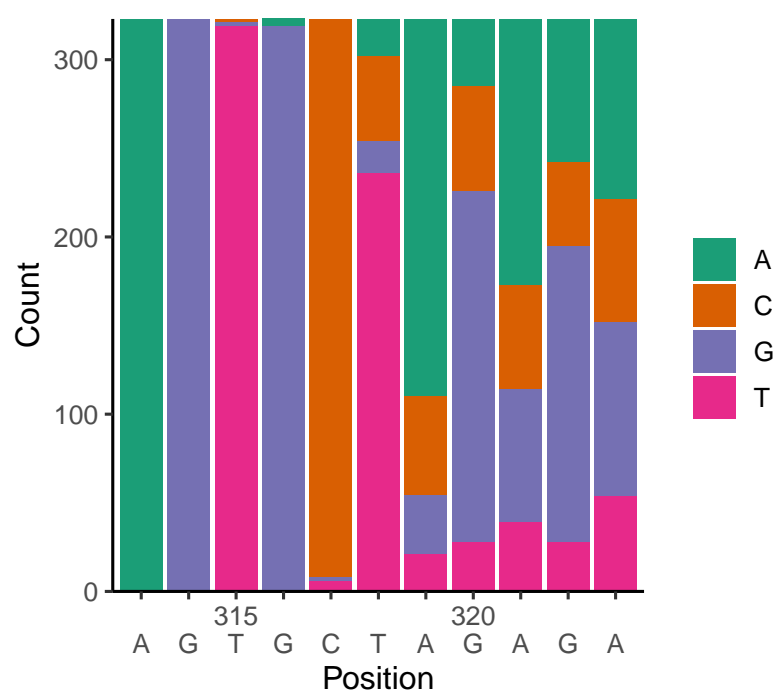
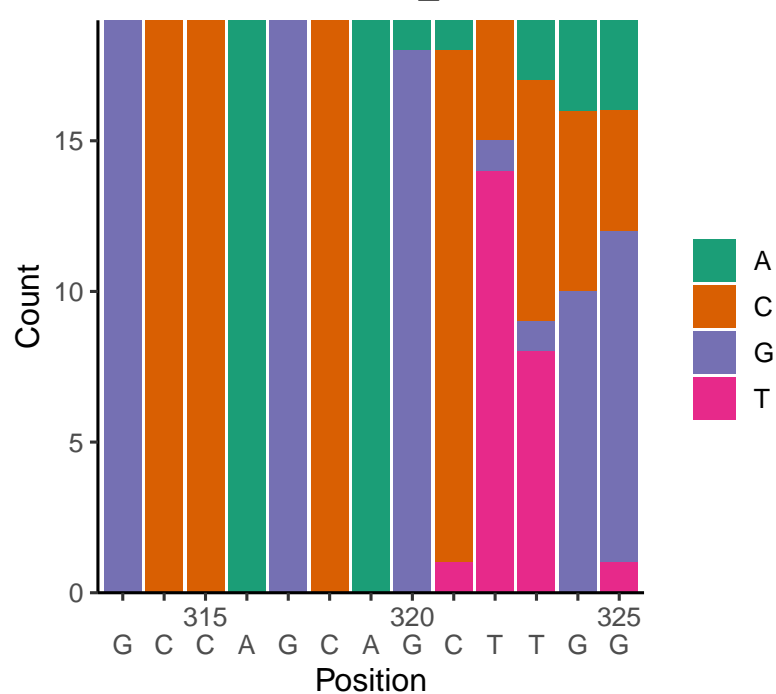


Gene TRBV20-1*02_319AGAGA323



Gene TRBV5-5*02_322TTGG325



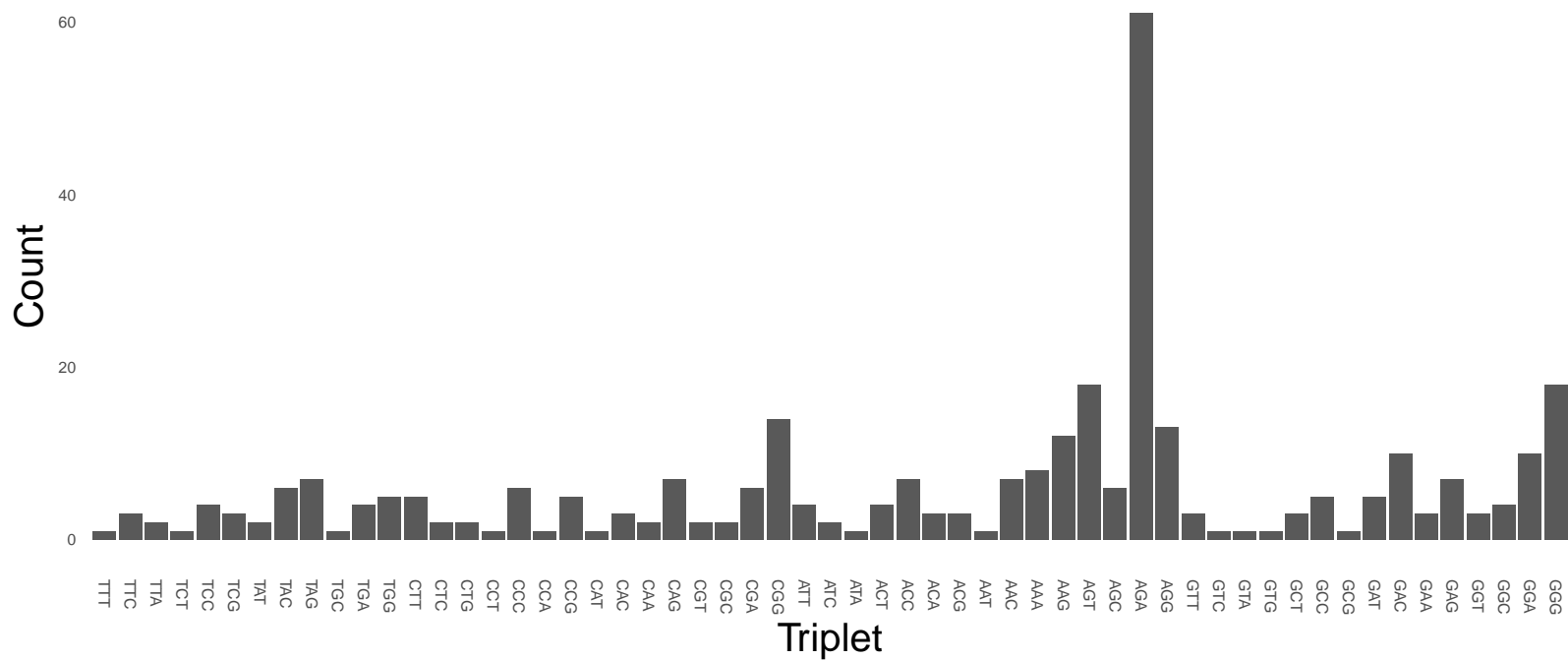
Gene TRBV20-1*02_319AGAGA323



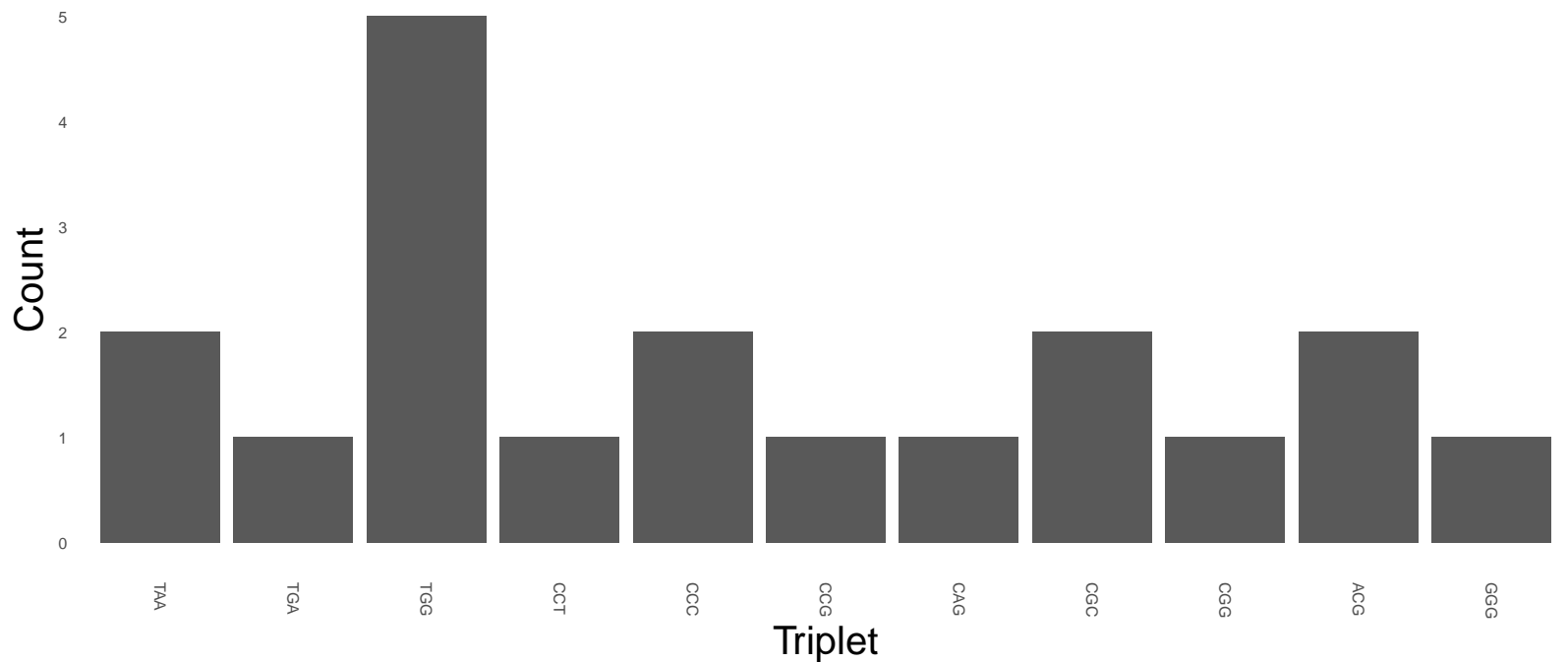
Gene TRBV5-5*02_322TTGG325



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

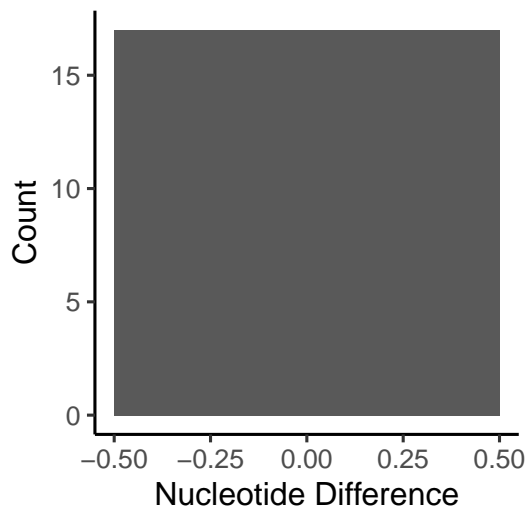


TRBV5–5*02_322TTGG325– Final 3 nucleotides as a triplet



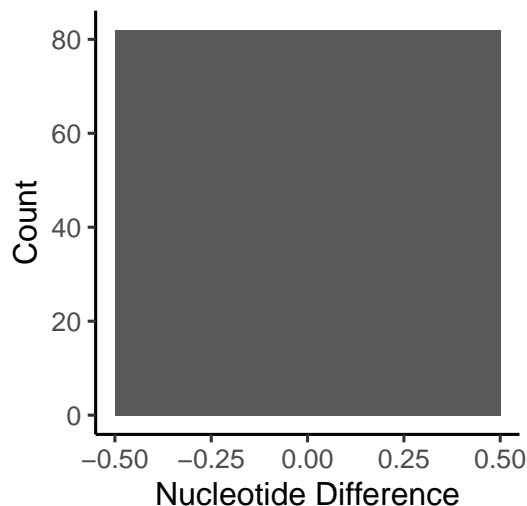
TRBV10–1*01

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



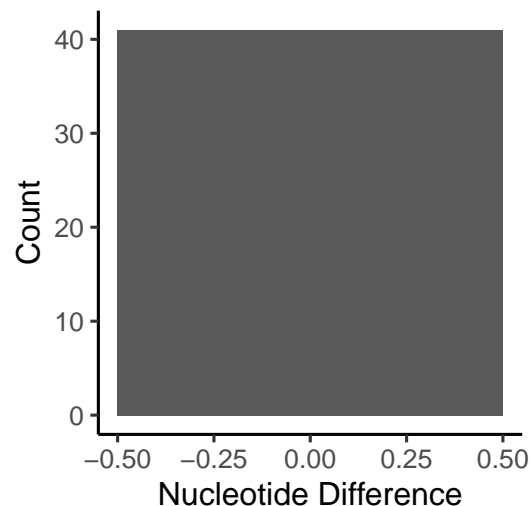
TRBV10–3*01

82 sequences assigned
82 (100%) exact matches, in which:
82 unique CDR3
11 unique J



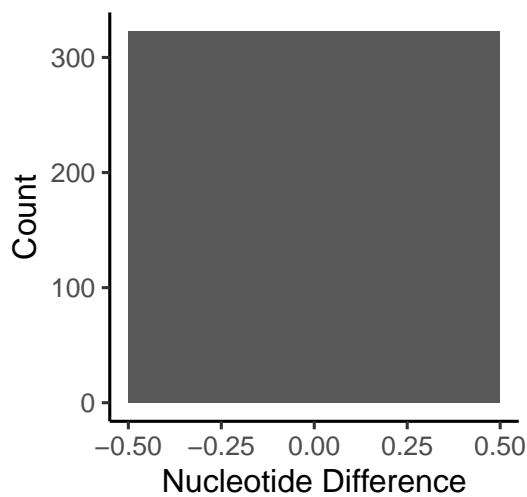
TRBV15*02

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



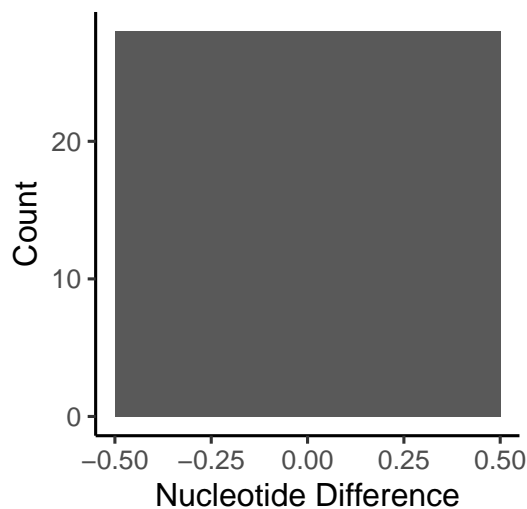
TRBV20–1*02_319AGAG

323 sequences assigned
323 (100%) exact matches, in which:
323 unique CDR3
12 unique J



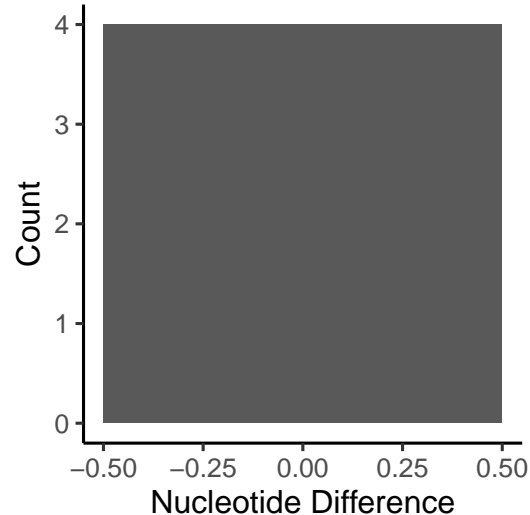
TRBV13*01

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



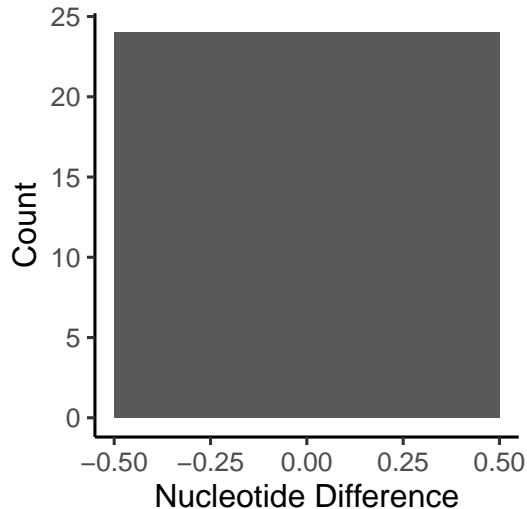
TRBV16*01

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



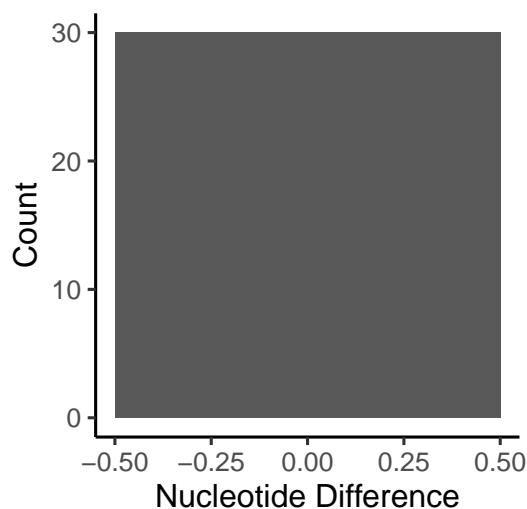
TRBV10–2*01

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



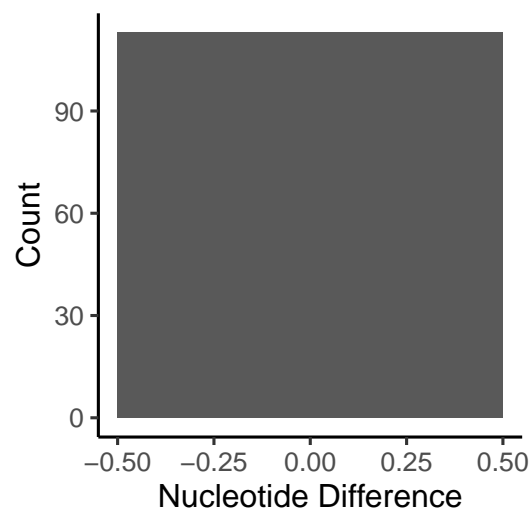
TRBV14*01

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



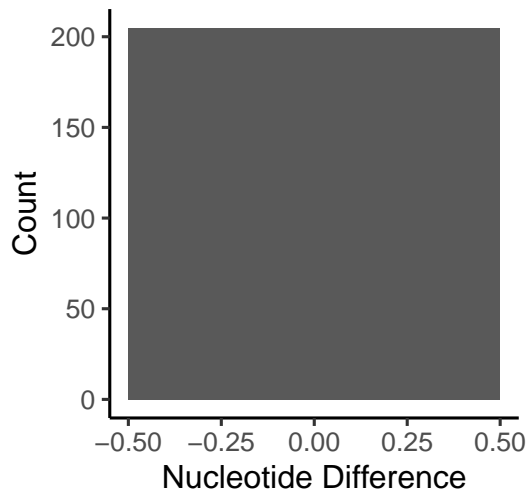
TRBV18*01

113 sequences assigned
113 (100%) exact matches, in which:
113 unique CDR3
13 unique J



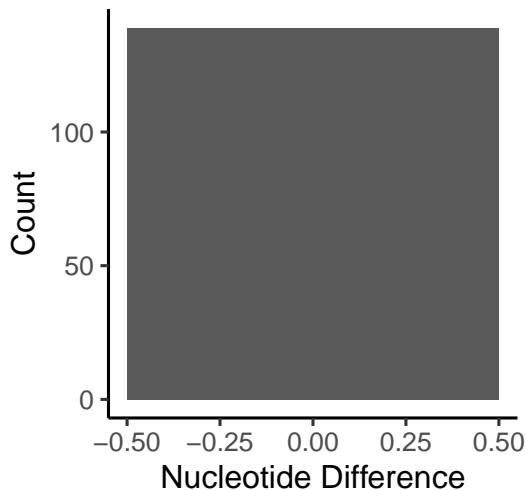
TRBV19*01

205 sequences assigned
205 (100%) exact matches, in which:
205 unique CDR3
13 unique J



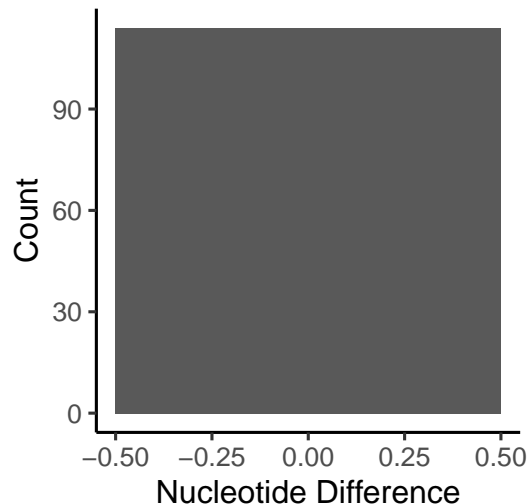
TRBV28*01

139 sequences assigned
139 (100%) exact matches, in which:
139 unique CDR3
13 unique J



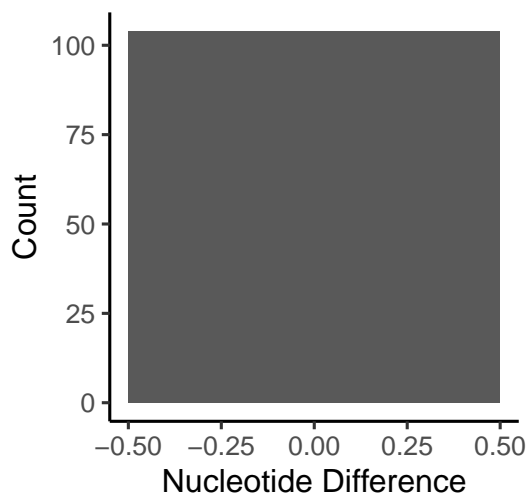
TRBV11-2*01

114 sequences assigned
114 (100%) exact matches, in which:
114 unique CDR3
13 unique J



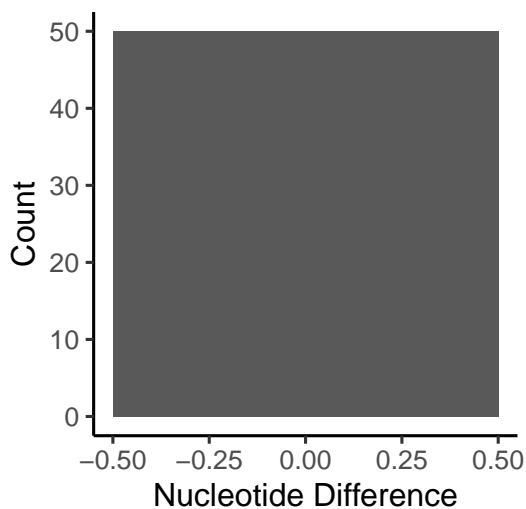
TRBV2*01

104 sequences assigned
104 (100%) exact matches, in which:
104 unique CDR3
11 unique J



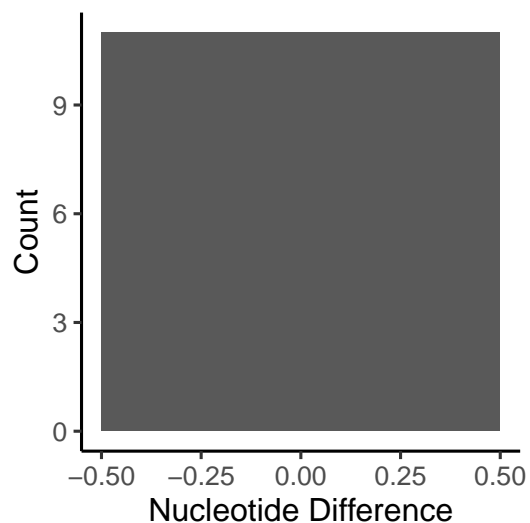
TRBV30*01

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



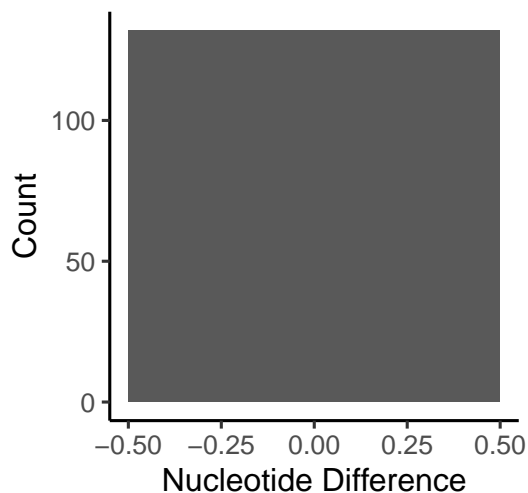
TRBV11-3*01

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



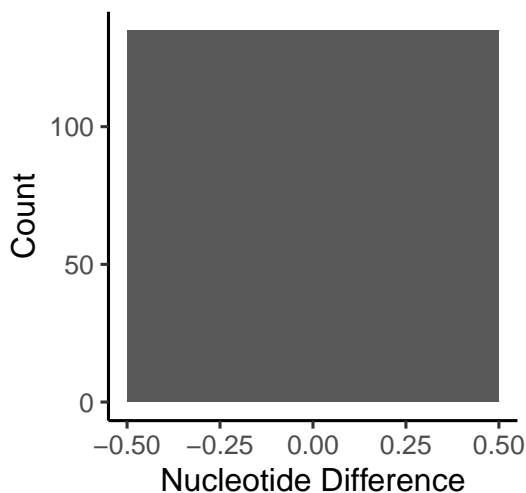
TRBV27*01

132 sequences assigned
132 (100%) exact matches, in which:
132 unique CDR3
13 unique J



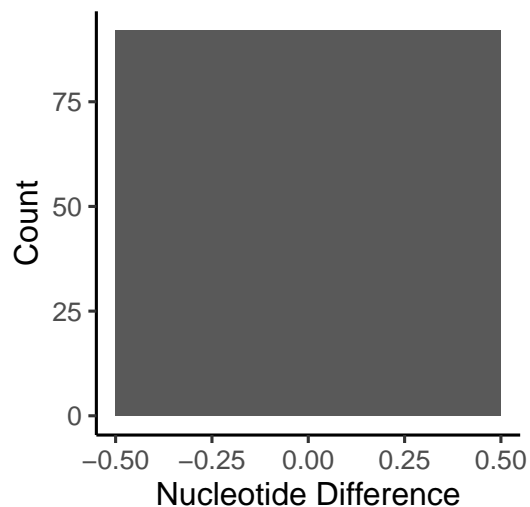
TRBV9*01

135 sequences assigned
135 (100%) exact matches, in which:
135 unique CDR3
12 unique J



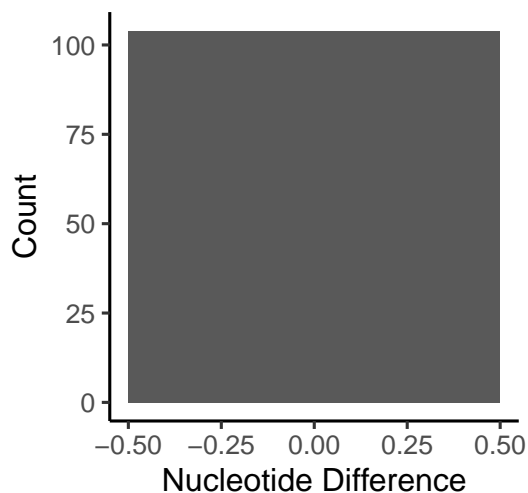
TRBV12-3*01

92 sequences assigned
92 (100%) exact matches, in which:
92 unique CDR3
13 unique J



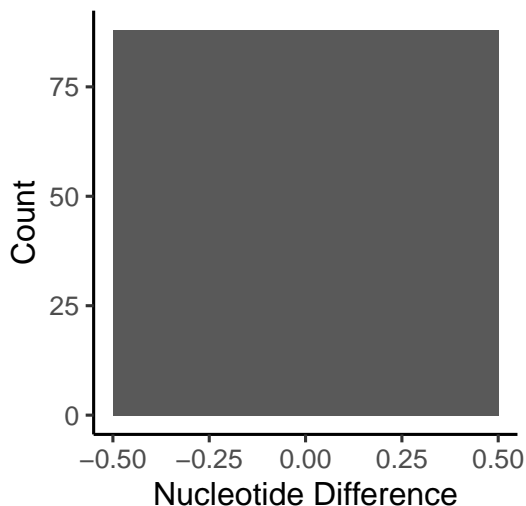
TRBV12–4*01

104 sequences assigned
104 (100%) exact matches, in which:
104 unique CDR3
12 unique J



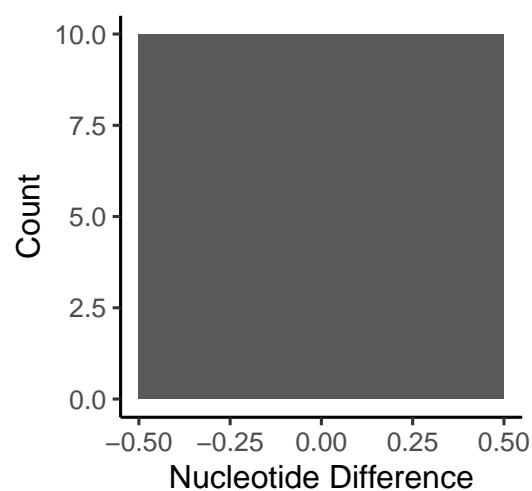
TRBV4–1*01

88 sequences assigned
88 (100%) exact matches, in which:
88 unique CDR3
11 unique J



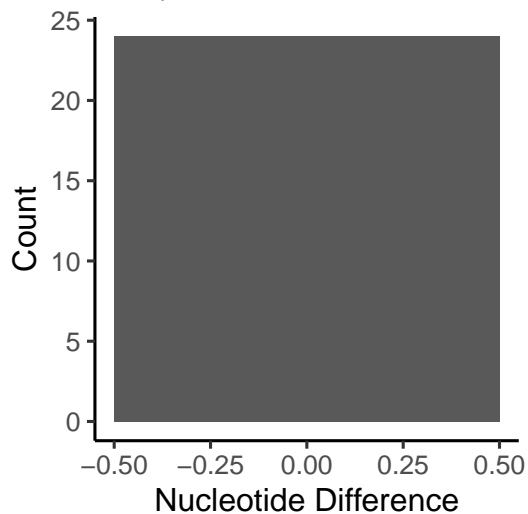
TRBV4–2*01

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



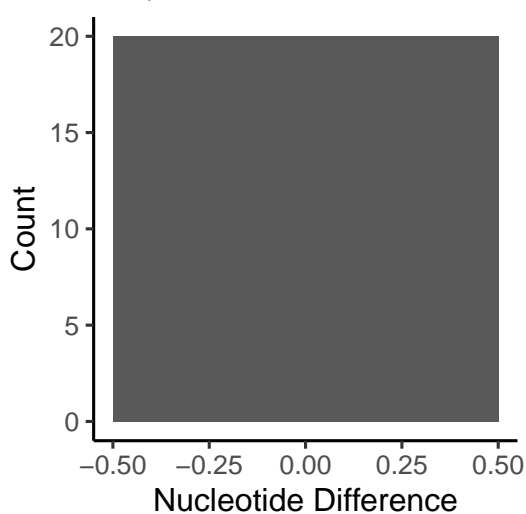
TRBV12–5*01

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



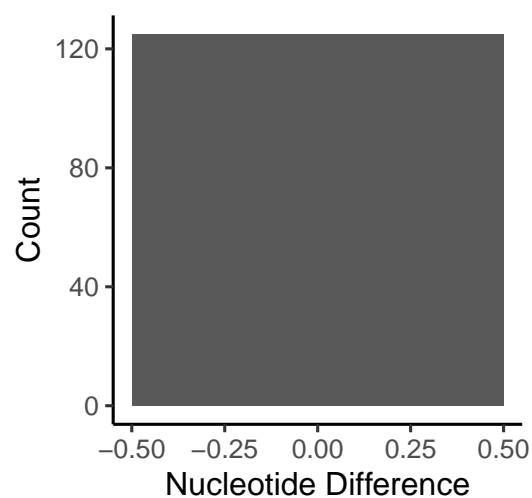
TRBV24–1*01

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



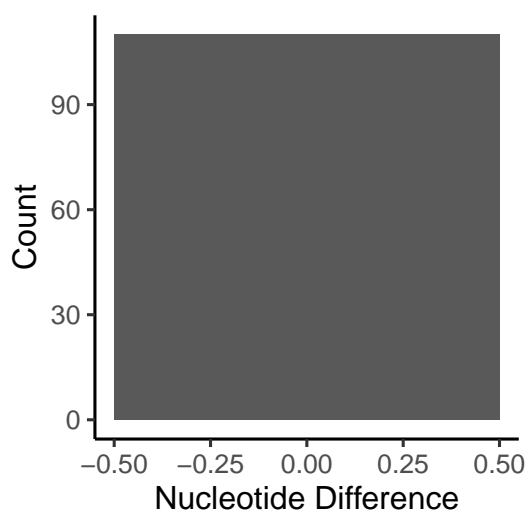
TRBV4–3*01

125 sequences assigned
125 (100%) exact matches, in which:
125 unique CDR3
13 unique J



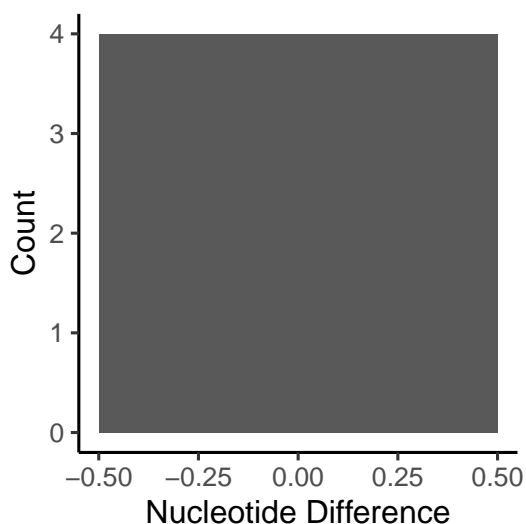
TRBV3–1*01

110 sequences assigned
110 (100%) exact matches, in which:
110 unique CDR3
13 unique J



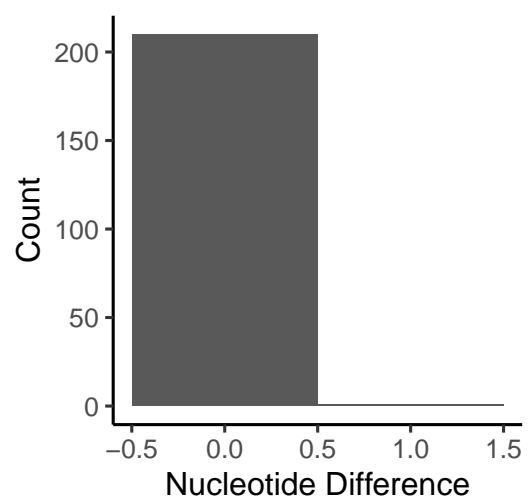
TRBV24–1*02

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



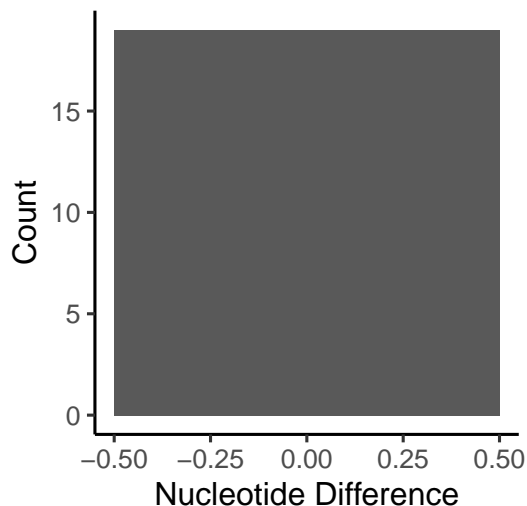
TRBV5–1*01

211 sequences assigned
210 (99.5%) exact matches, in which:
210 unique CDR3
13 unique J



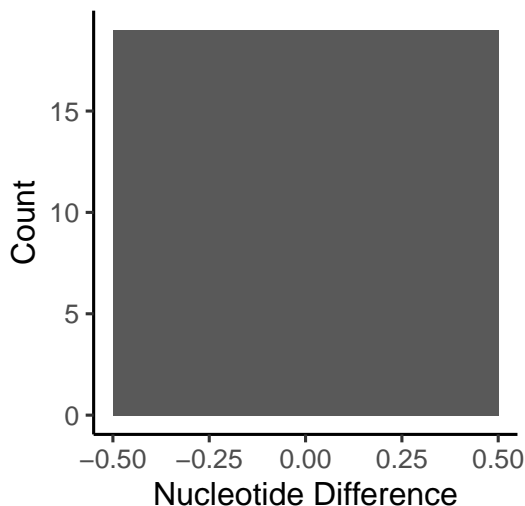
TRBV25–1*01

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



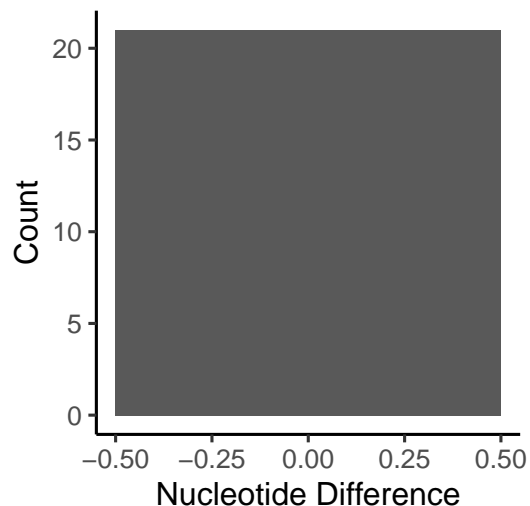
TRBV5–5*02_322TTGG32

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



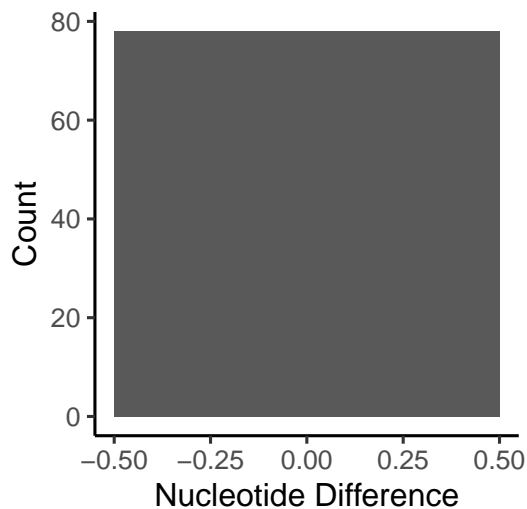
TRBV5–8*01

21 sequences assigned
21 (100%) exact matches, in which:
21 unique CDR3
10 unique J



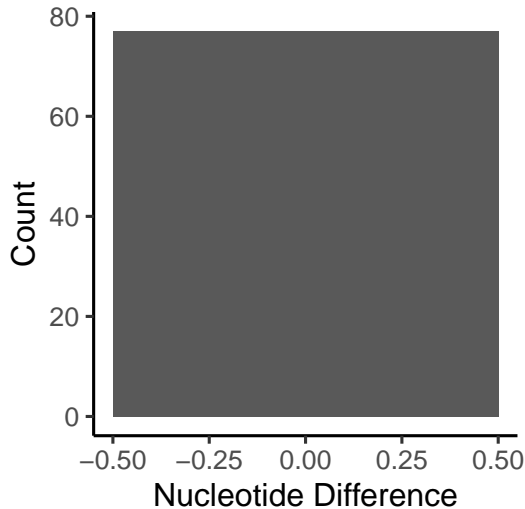
TRBV5–4*01

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



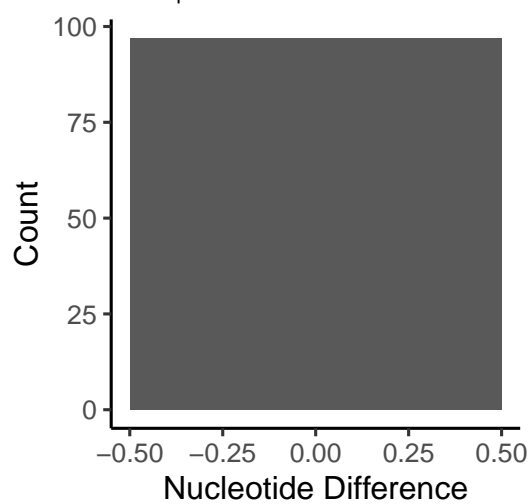
TRBV5–6*01

77 sequences assigned
77 (100%) exact matches, in which:
77 unique CDR3
13 unique J



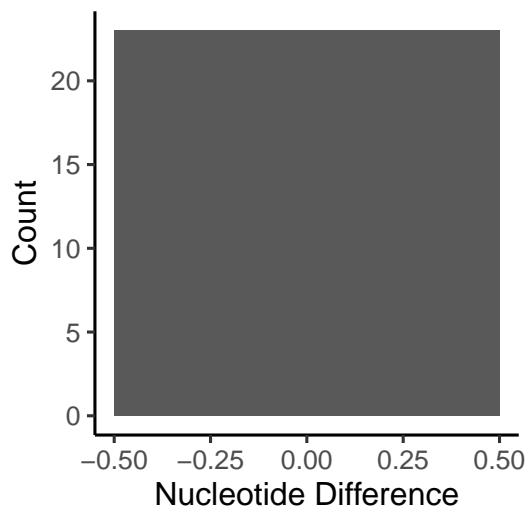
TRBV6–1*01

97 sequences assigned
97 (100%) exact matches, in which:
97 unique CDR3
13 unique J



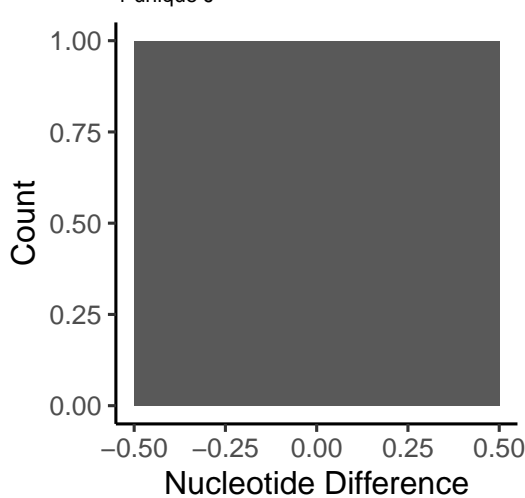
TRBV5–5*01

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



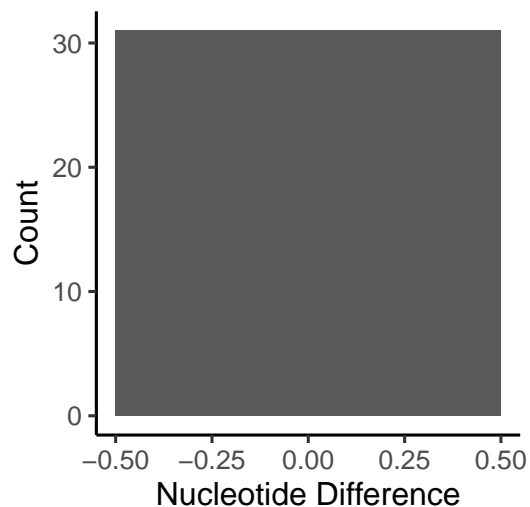
TRBV5–7*01

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



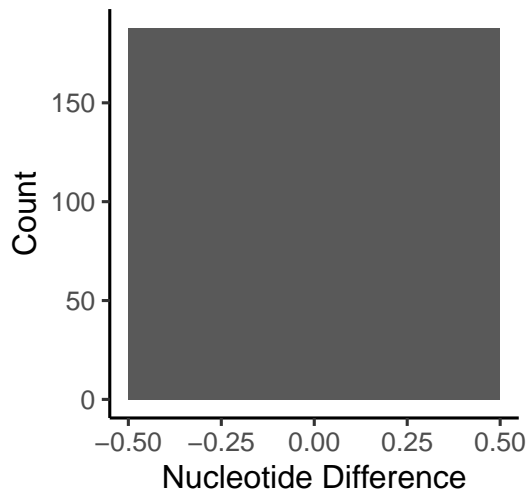
TRBV6–4*01

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



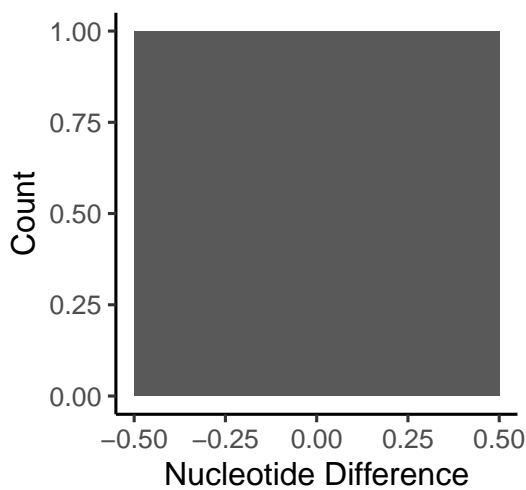
TRBV6–5*01

188 sequences assigned
188 (100%) exact matches, in which:
188 unique CDR3
13 unique J



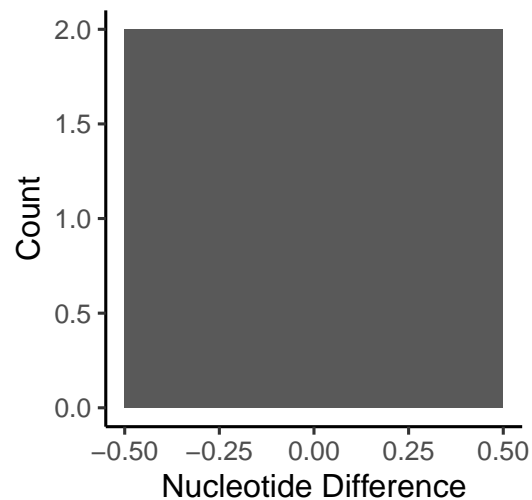
TRBV6–8*01

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



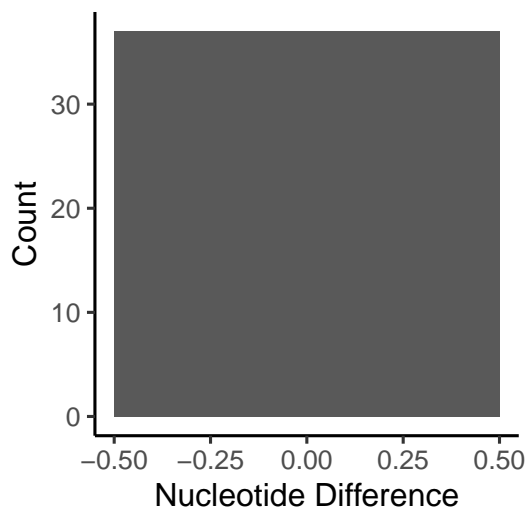
TRBV7–4*01

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



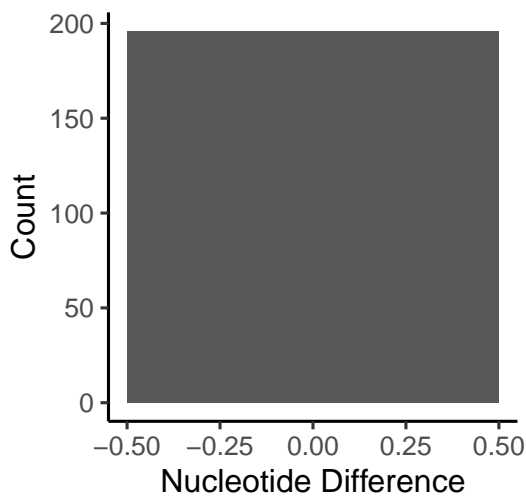
TRBV6–6*01

37 sequences assigned
37 (100%) exact matches, in which:
37 unique CDR3
12 unique J



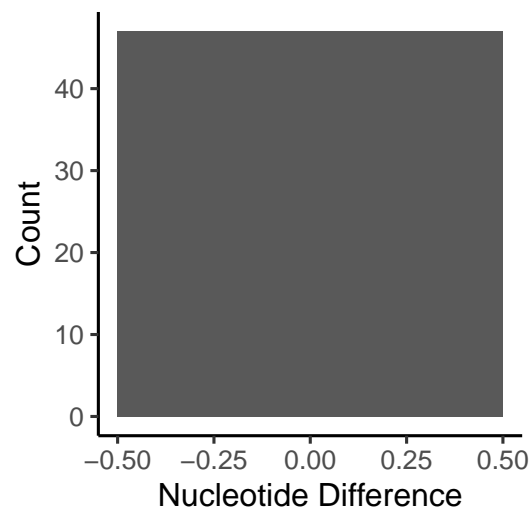
TRBV6–23*01

196 sequences assigned
196 (100%) exact matches, in which:
196 unique CDR3
13 unique J



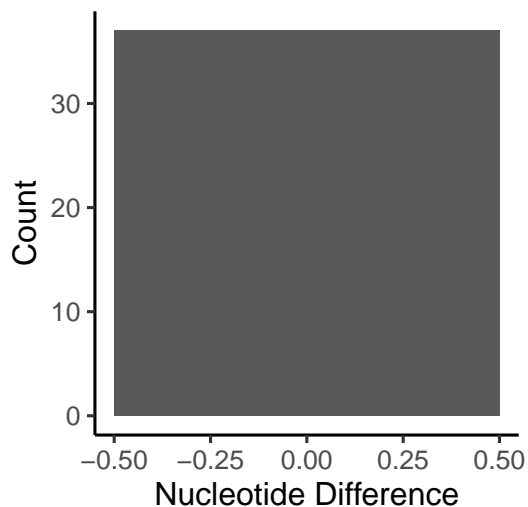
TRBV7–6*01

47 sequences assigned
47 (100%) exact matches, in which:
47 unique CDR3
13 unique J



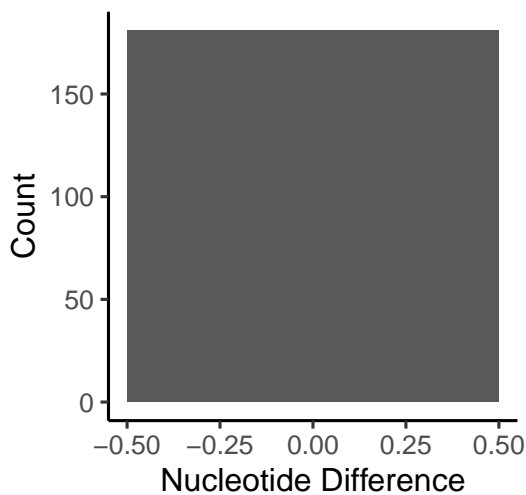
TRBV6–6*02

37 sequences assigned
37 (100%) exact matches, in which:
37 unique CDR3
11 unique J



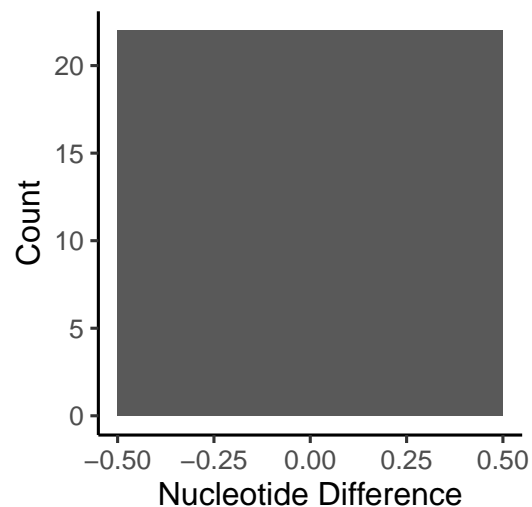
TRBV7–2*01

181 sequences assigned
181 (100%) exact matches, in which:
181 unique CDR3
13 unique J



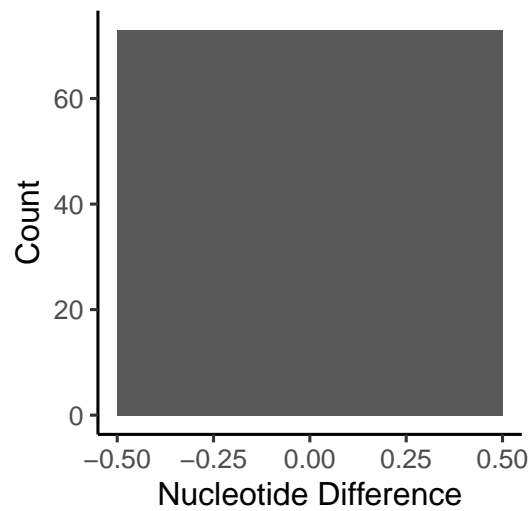
TRBV7–7*01

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



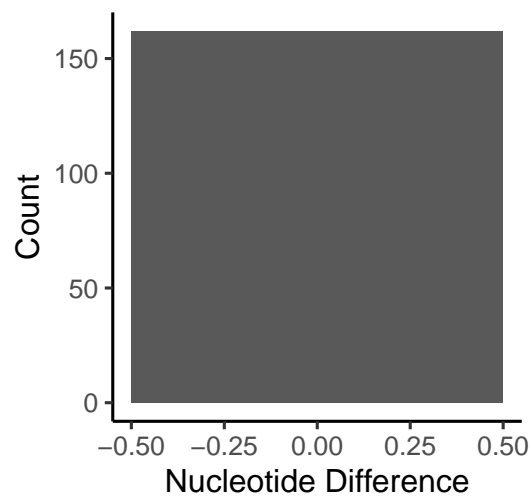
TRBV7-8*01

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



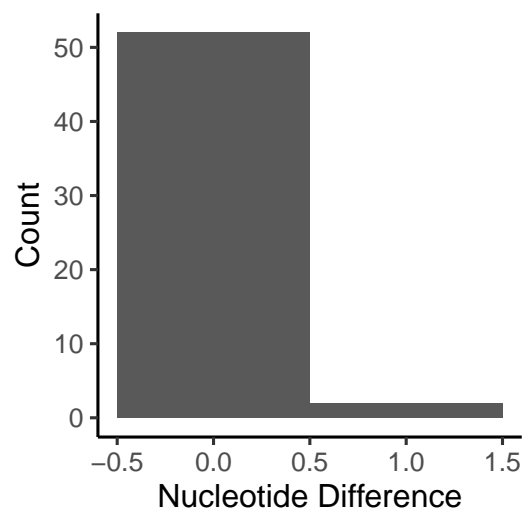
TRBV7-9*01

162 sequences assigned
162 (100%) exact matches, in which:
162 unique CDR3
13 unique J

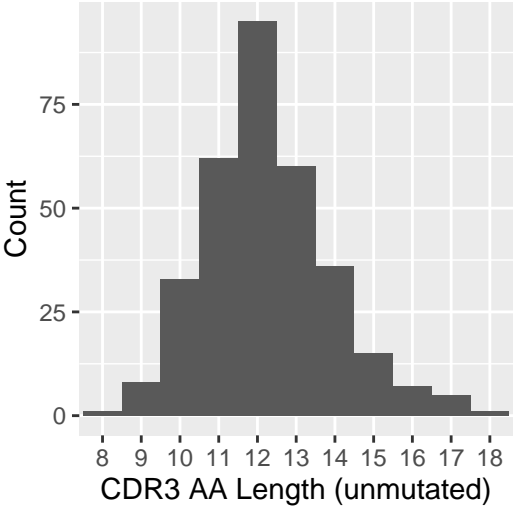


TRBV29-1*01

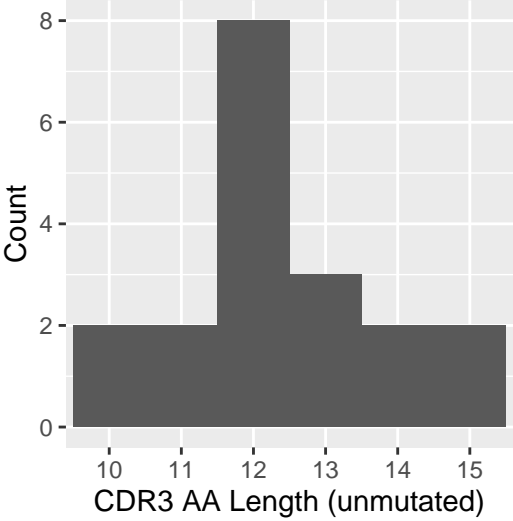
54 sequences assigned
52 (96.3%) exact matches, in which:
52 unique CDR3
11 unique J

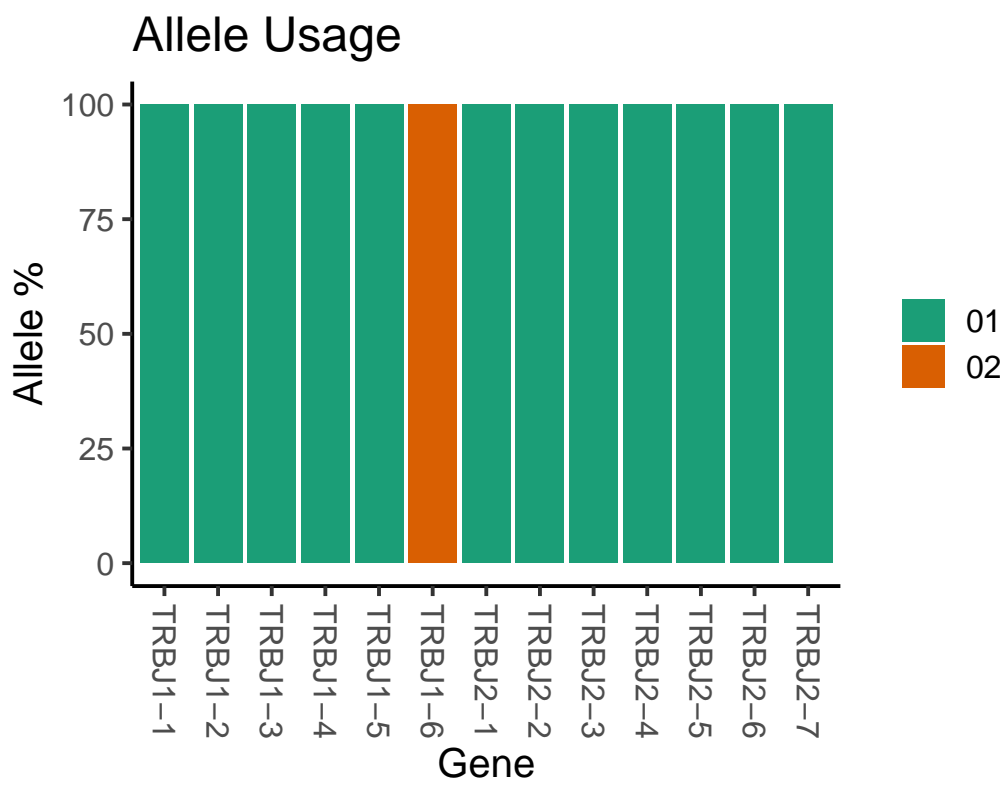


TRBV20-1*02_319AGAGA323



TRBV5-5*02_322TTGG325





Warning – no inferred sequences found.

Warning – no inferred sequences found.