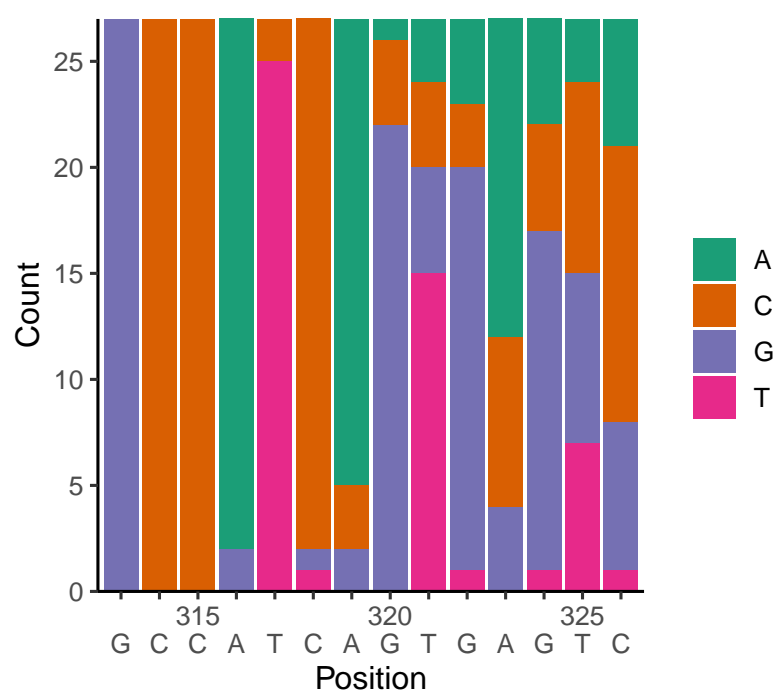
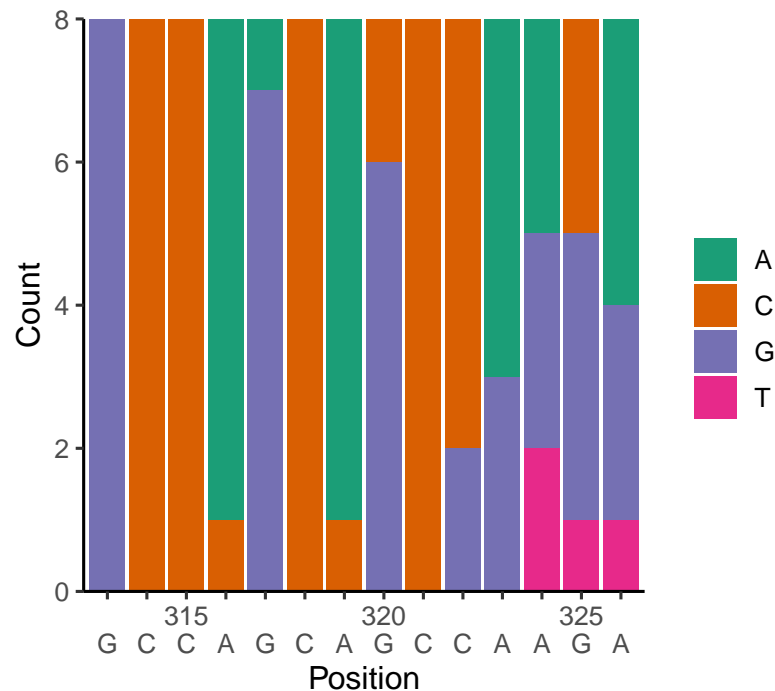


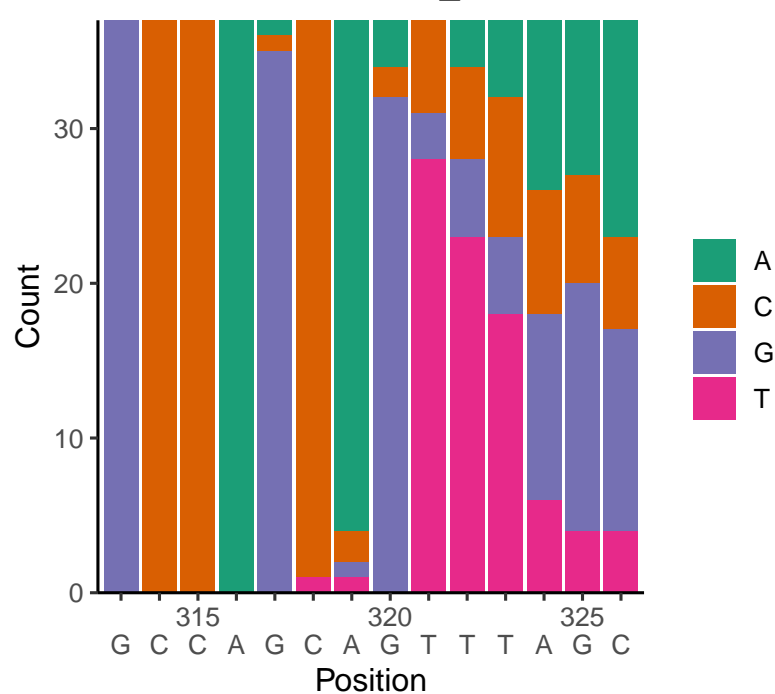
Gene TRBV10-3*03_313GCCATCAGTGA



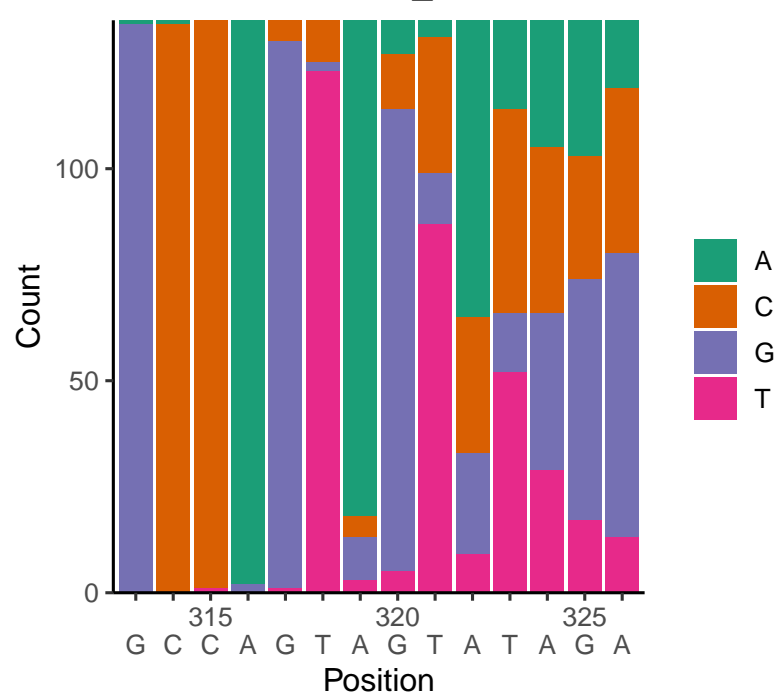
Gene TRBV14*02_322CAAGA326



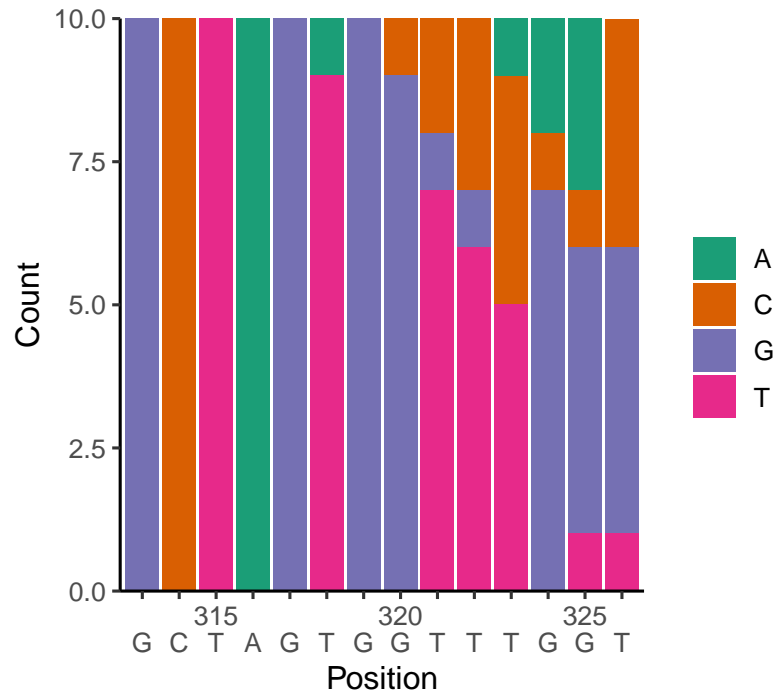
Gene TRBV12-4*01_C87T



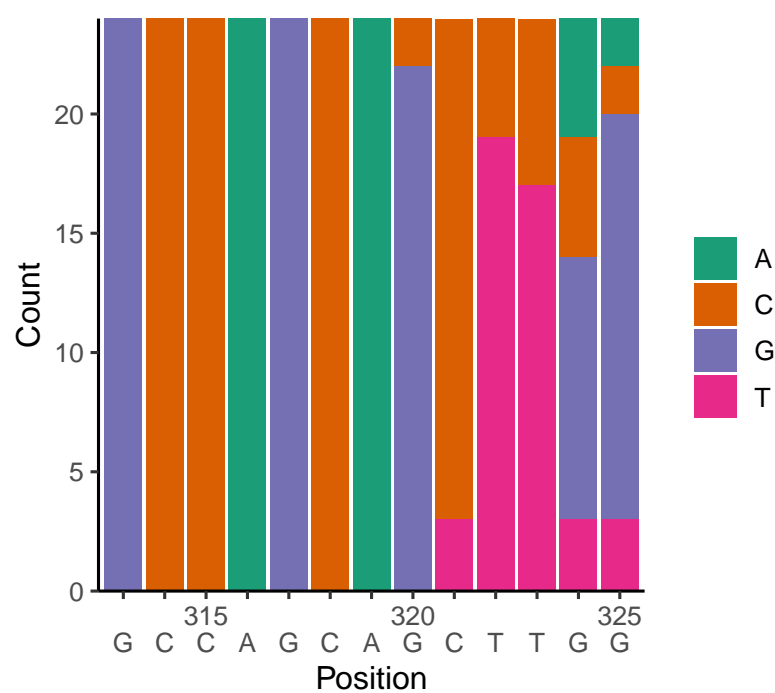
Gene TRBV19*01_A24G



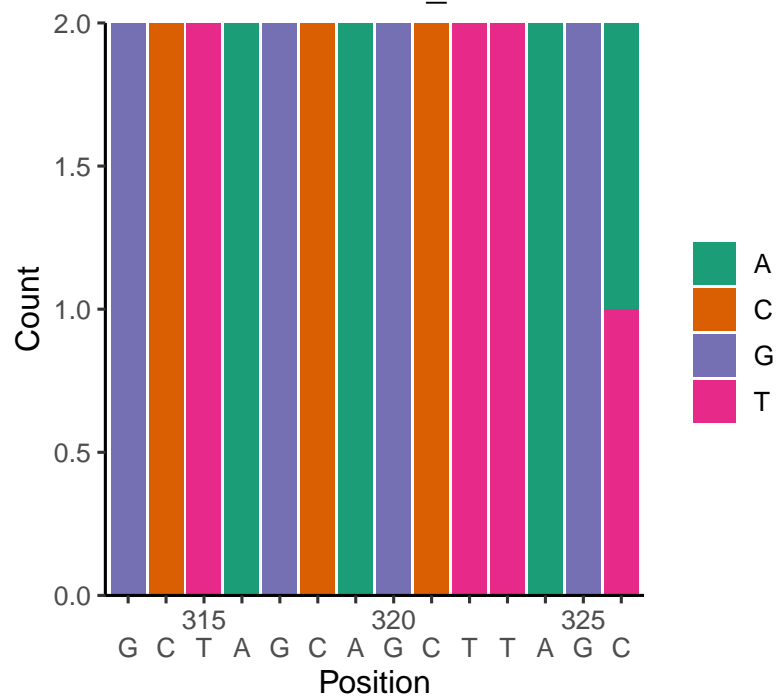
Gene TRBV12-5*01_C28G



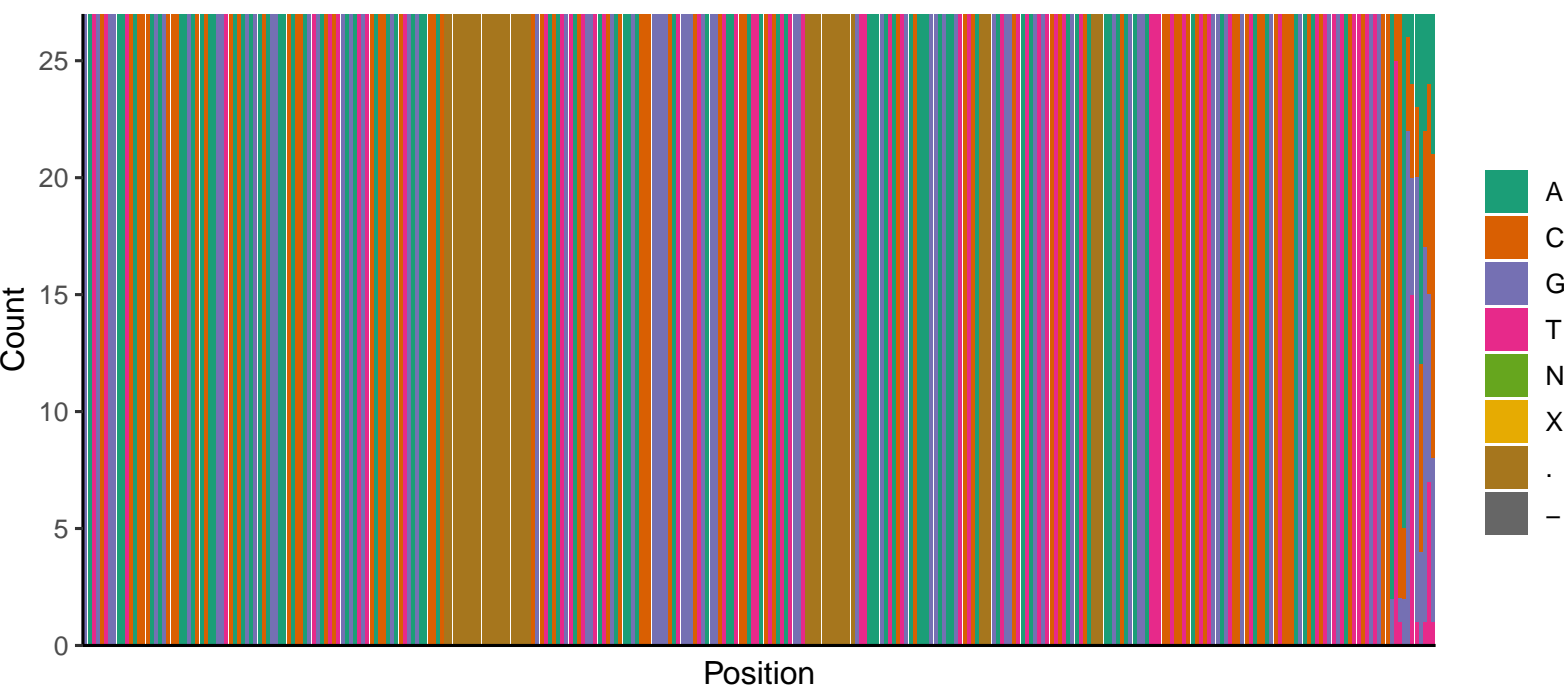
Gene TRBV5-5*02_322TTGG325



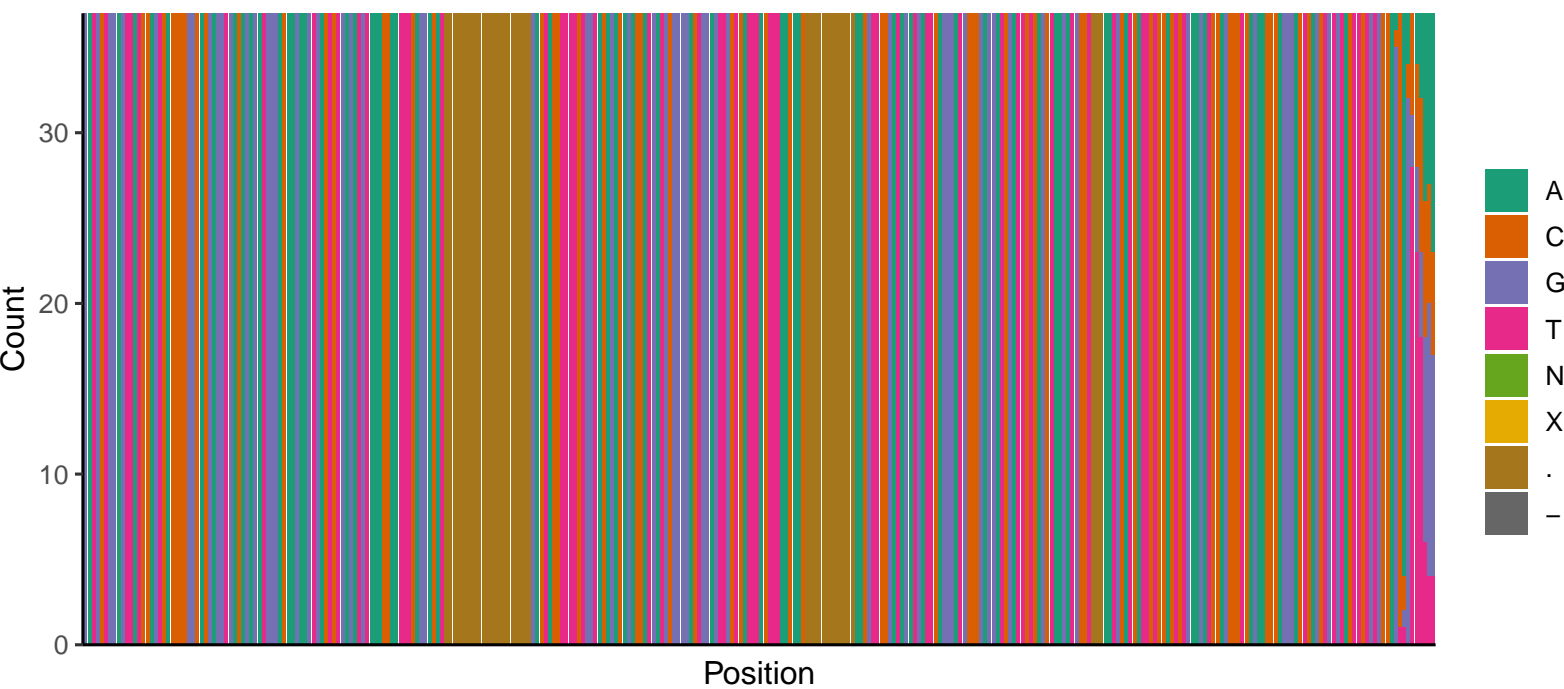
Gene TRBV7-7*01_C315T



Gene TRBV10-3*03_313GCCATCAGTGAGTC326



Gene TRBV12-4*01_C87T



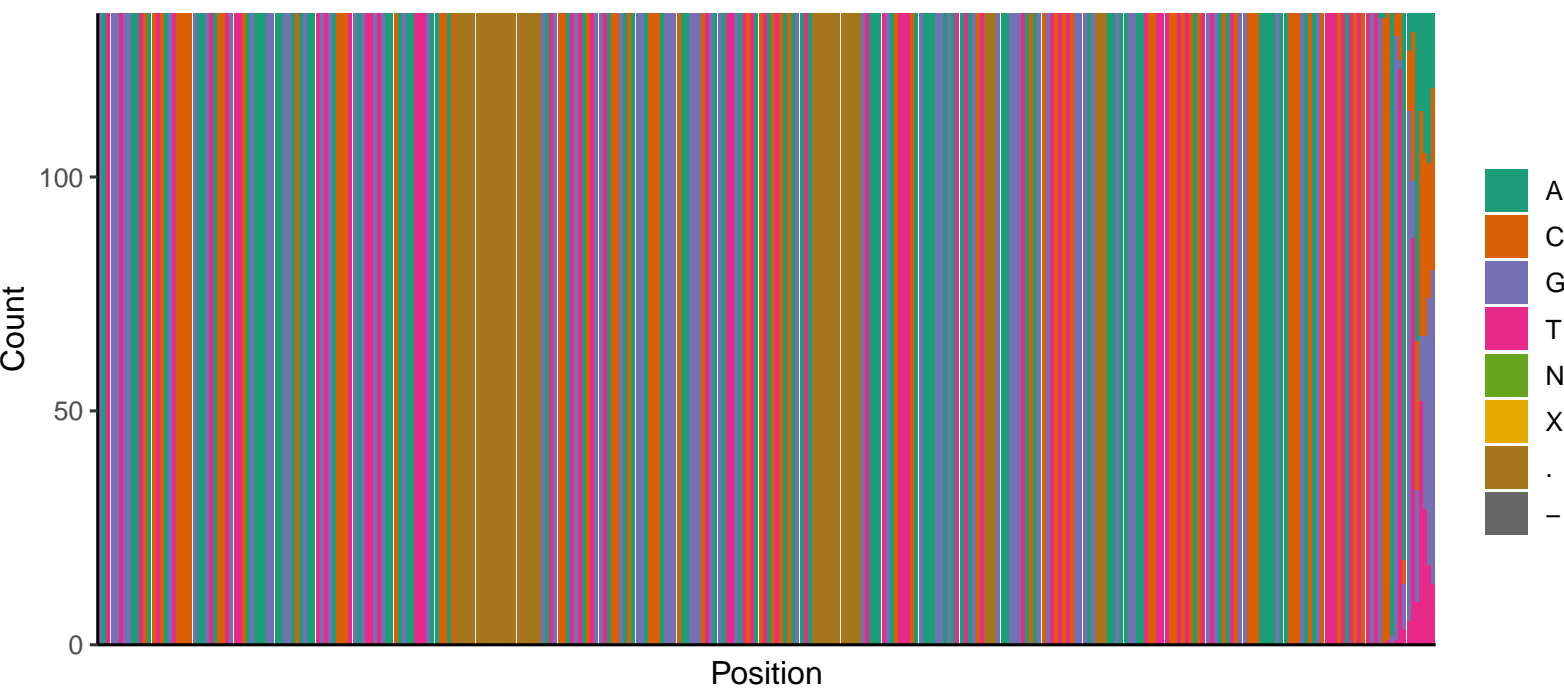
Gene TRBV12-5*01_C28G



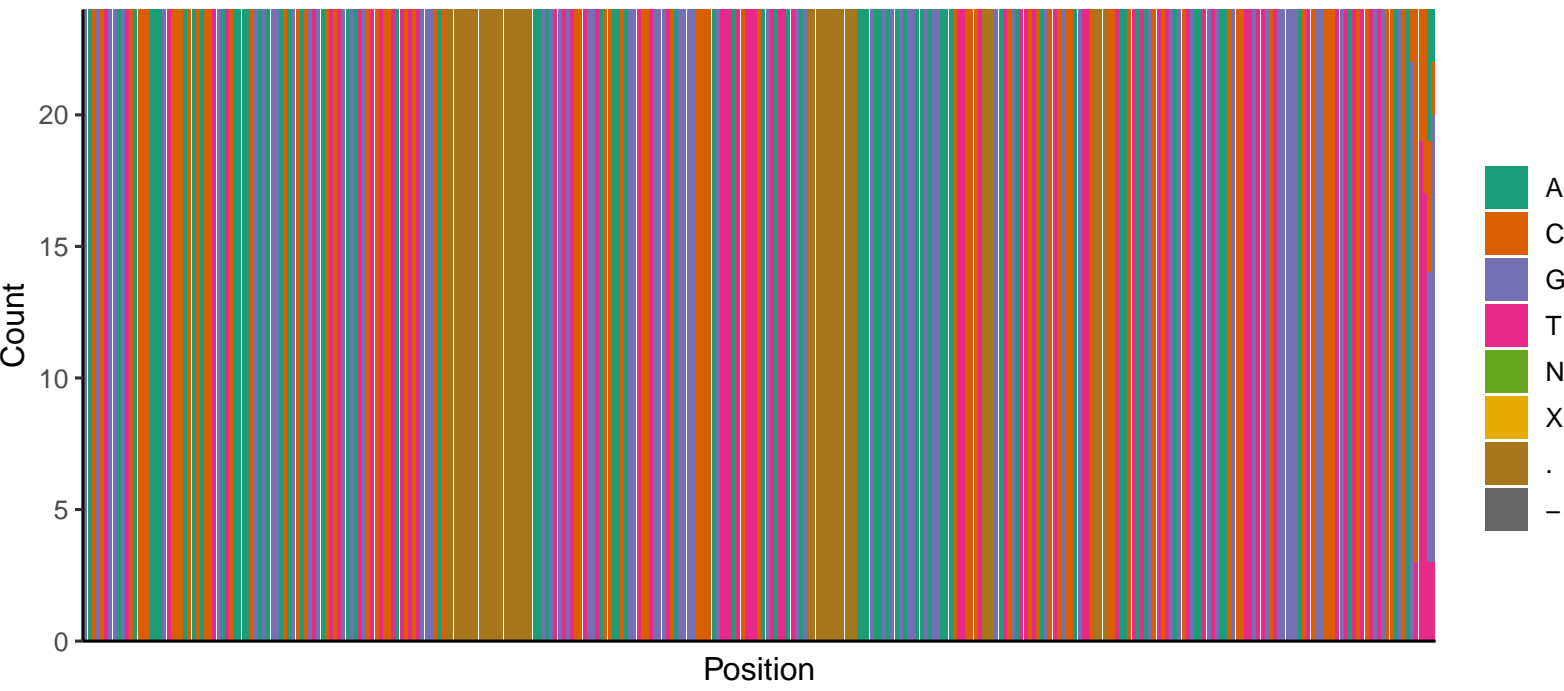
Gene TRBV14*02_322CAAGA326



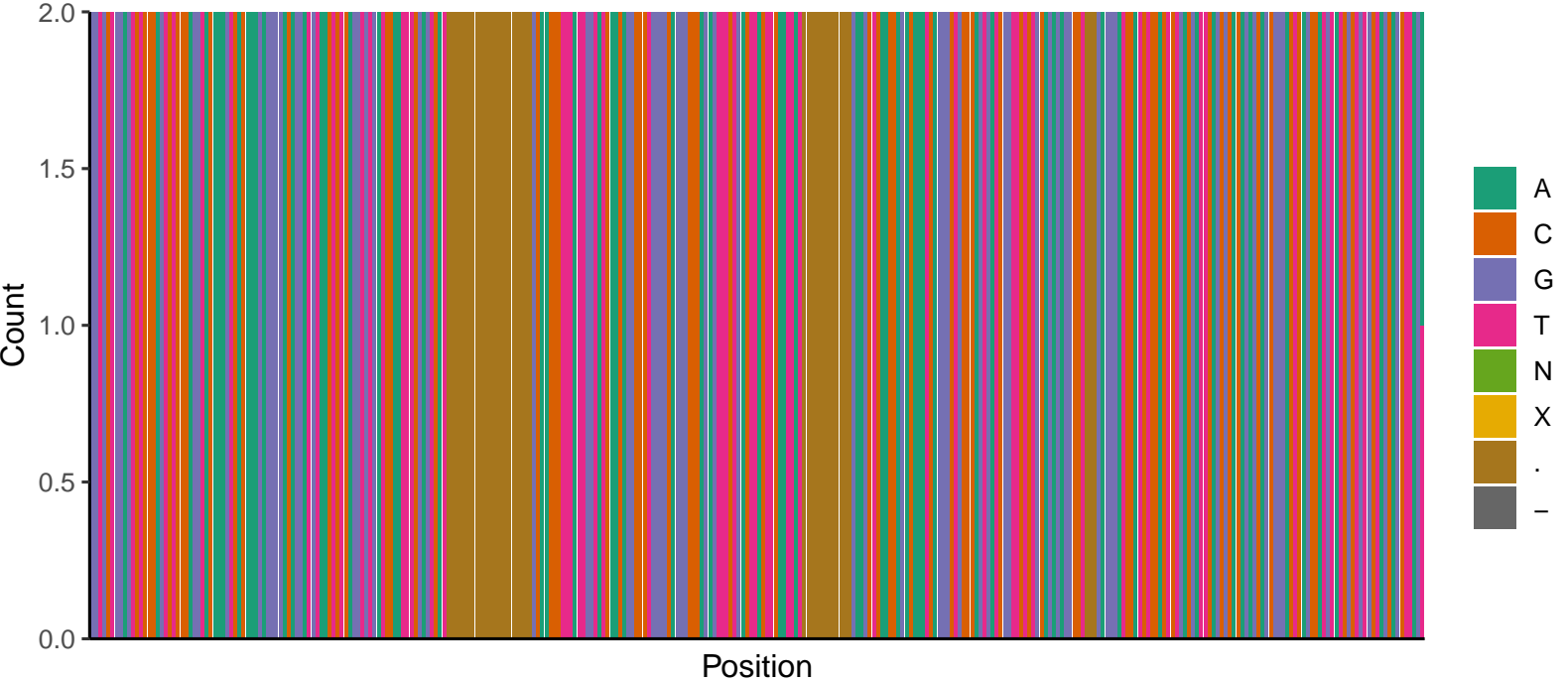
Gene TRBV19*01_A24G



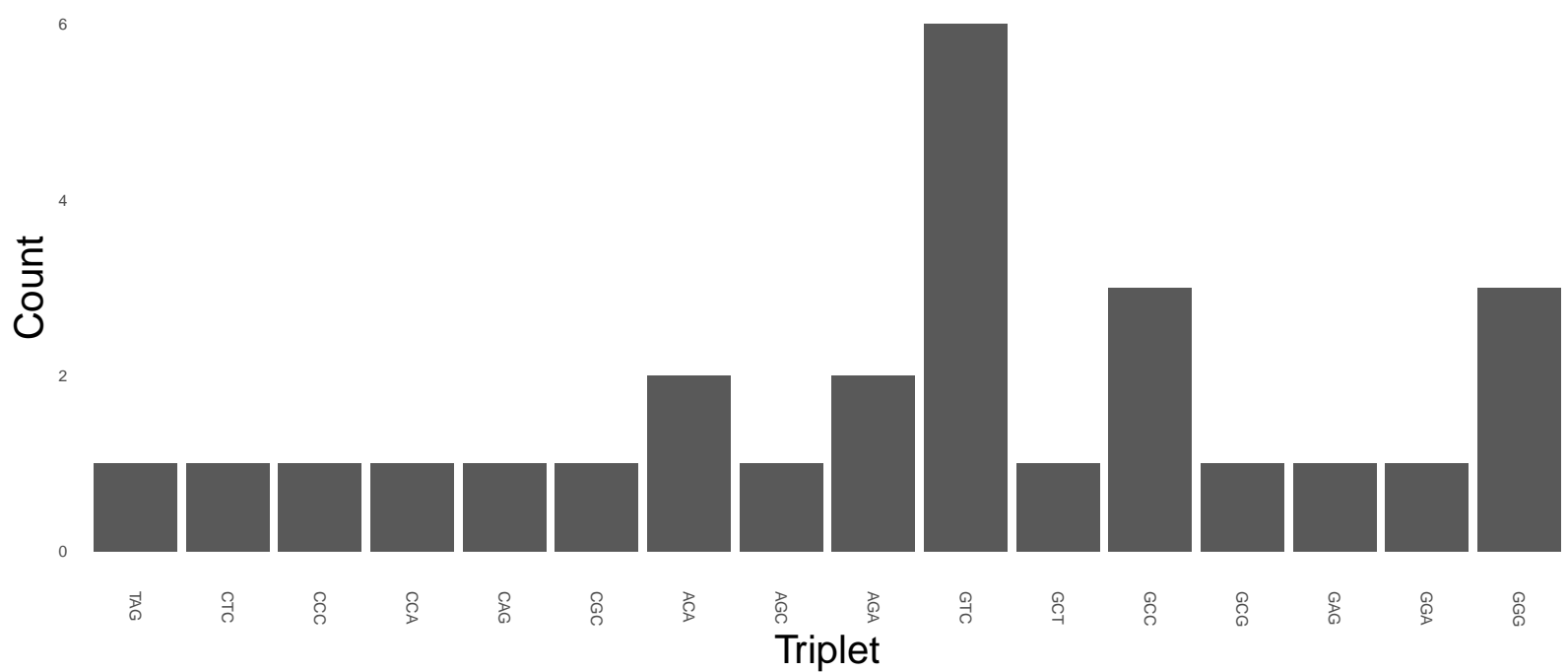
Gene TRBV5-5*02_322TTGG325



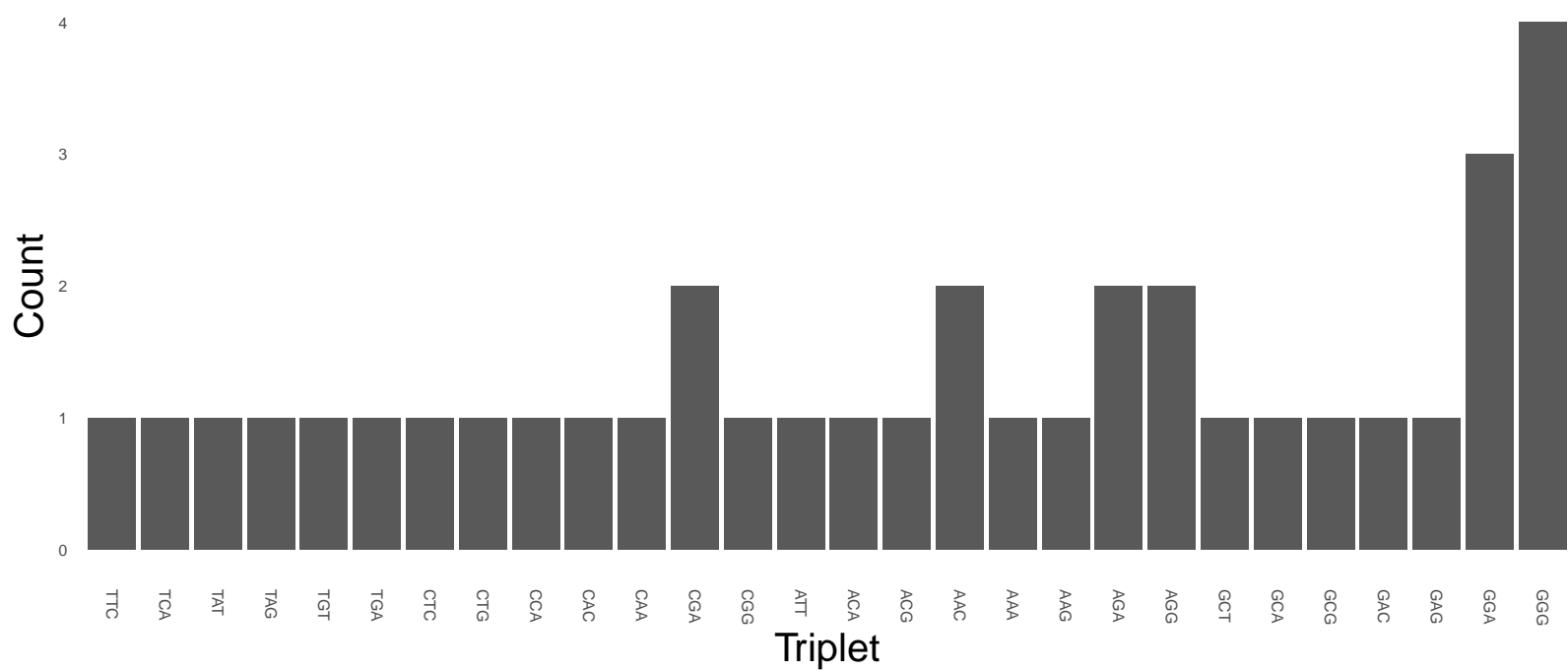
Gene TRBV7-7*01_C315T



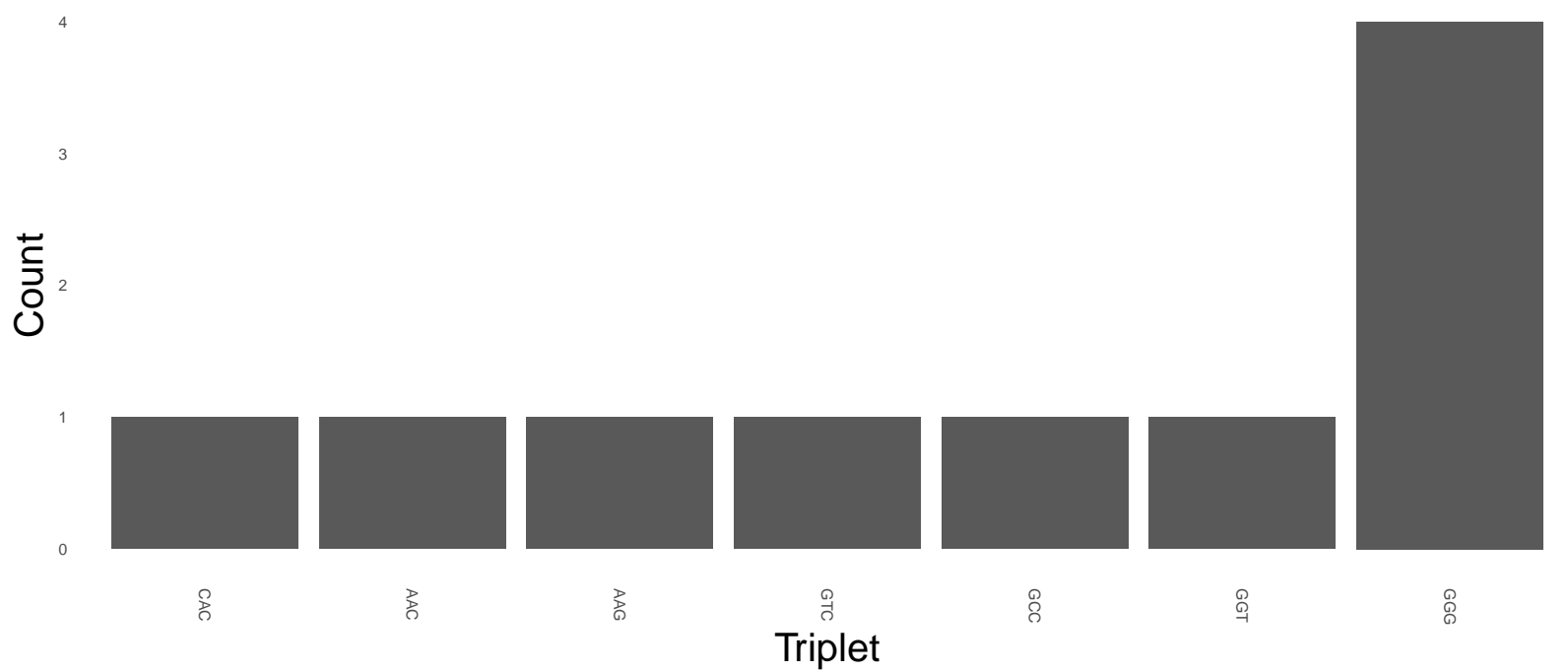
TRBV10–3*03_313GCCATCAGTGAGTC326– Final 3 nucleotides as a triplet



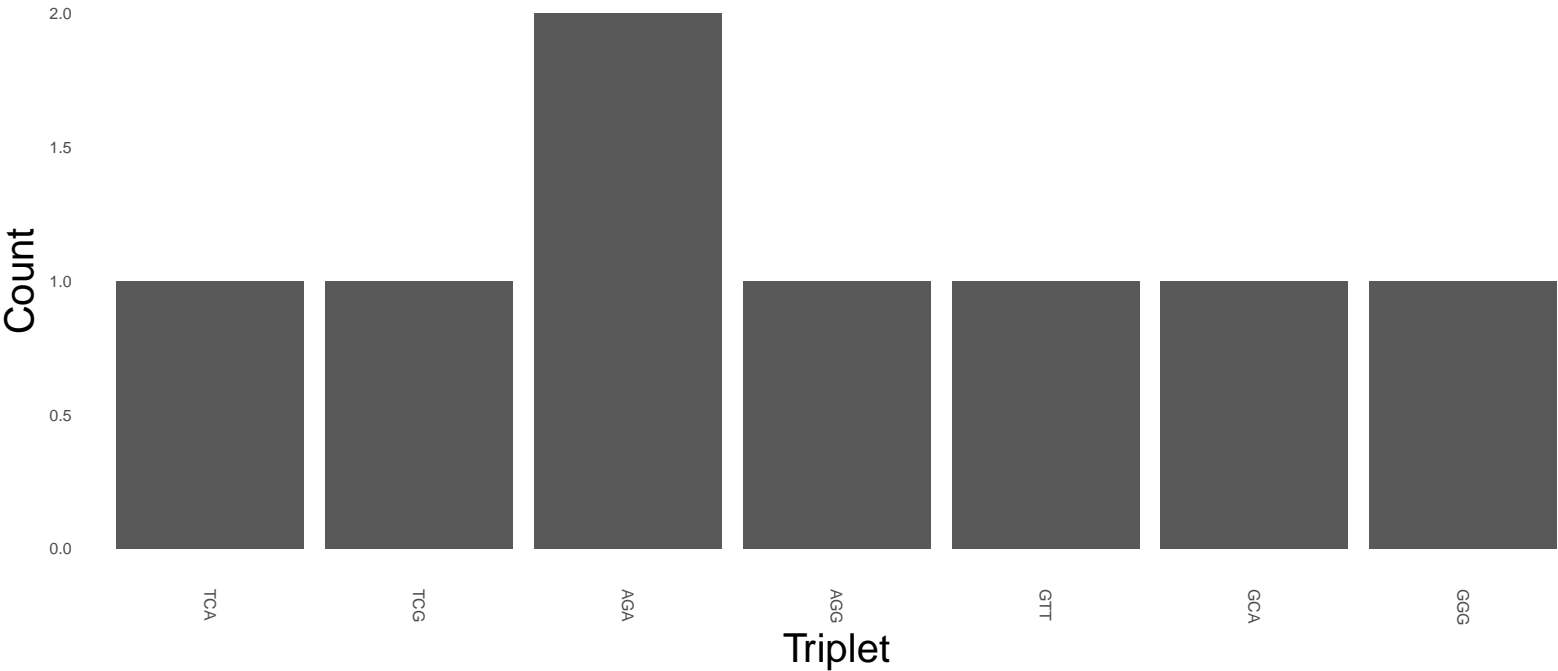
TRBV12–4*01_C87T– Final 3 nucleotides as a triplet



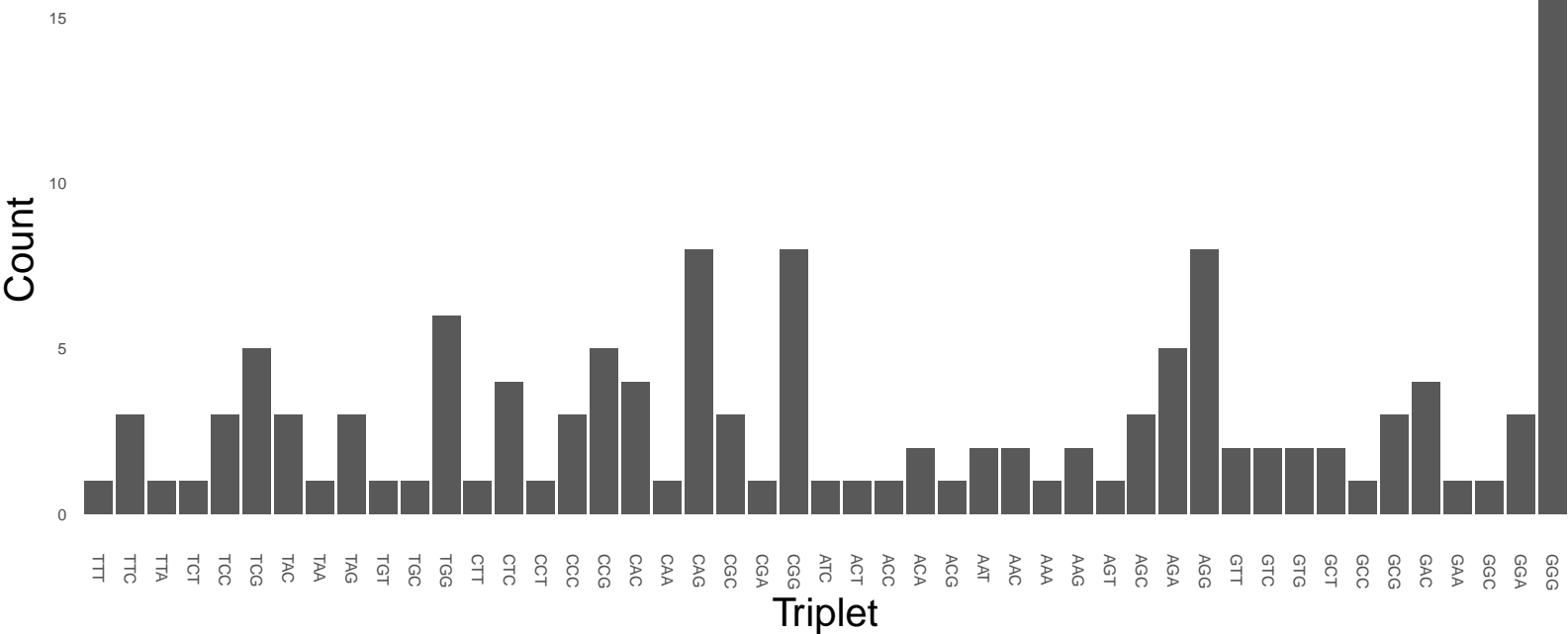
TRBV12–5*01_C28G– Final 3 nucleotides as a triplet



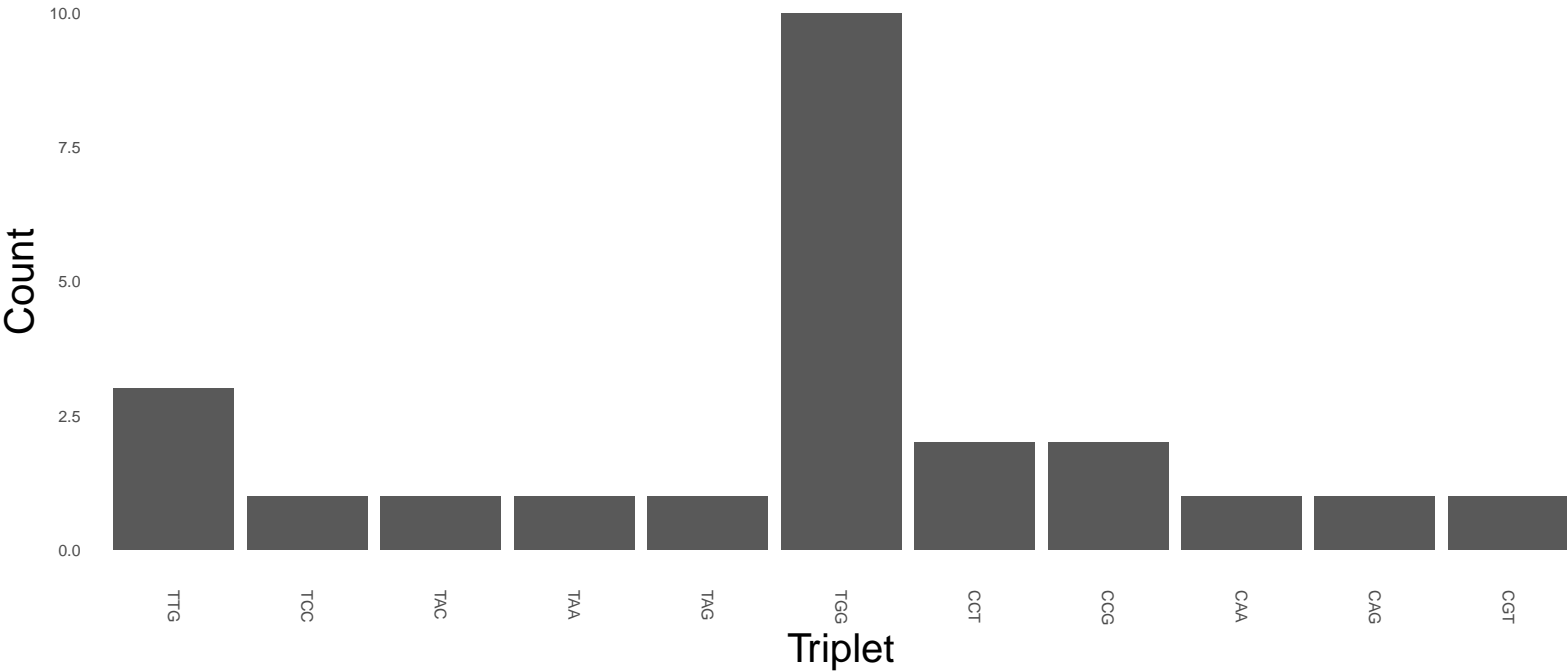
TRBV14*02_322CAAGA326– Final 3 nucleotides as a triplet



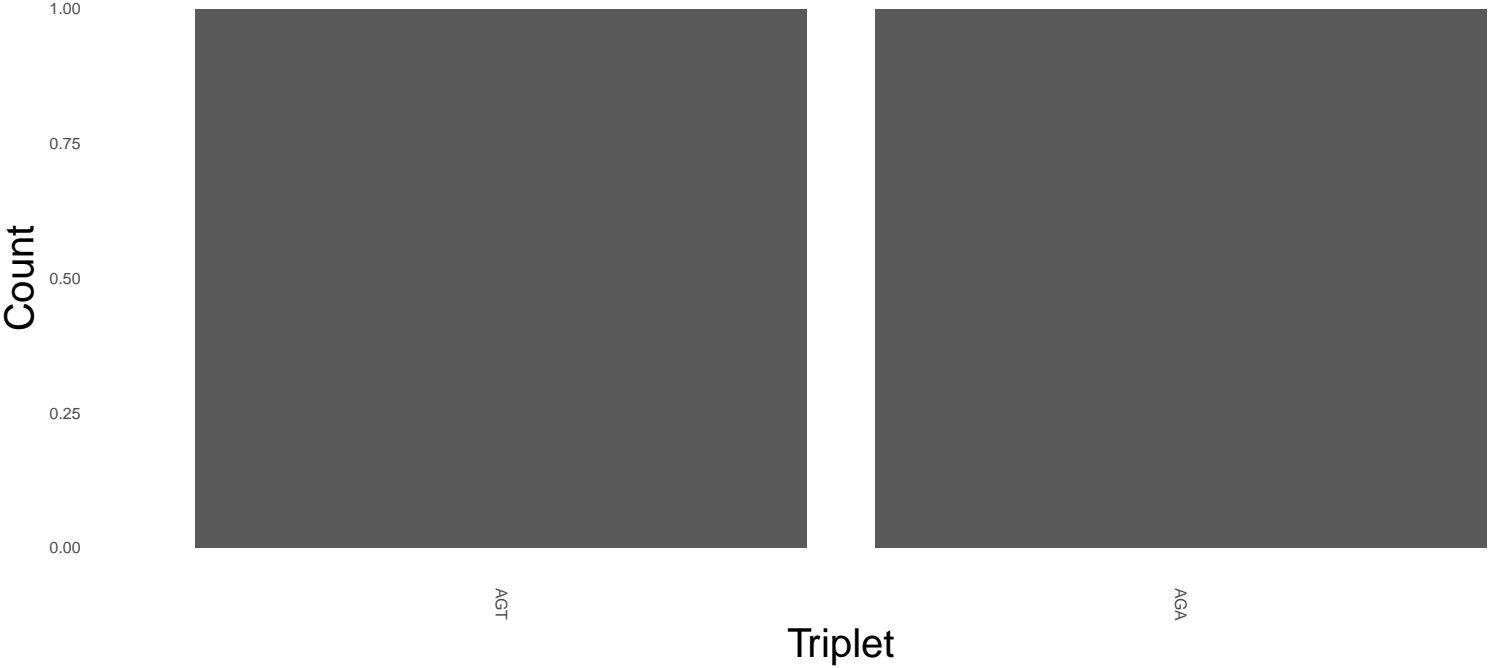
TRBV19*01_A24G– Final 3 nucleotides as a triplet



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

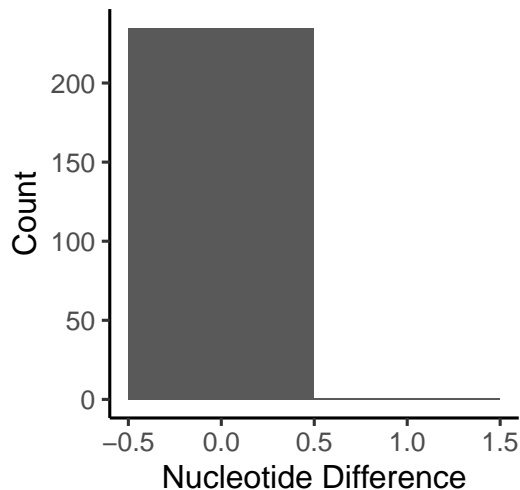


TRBV7-7*01_C315T- Final 3 nucleotides as a triplet



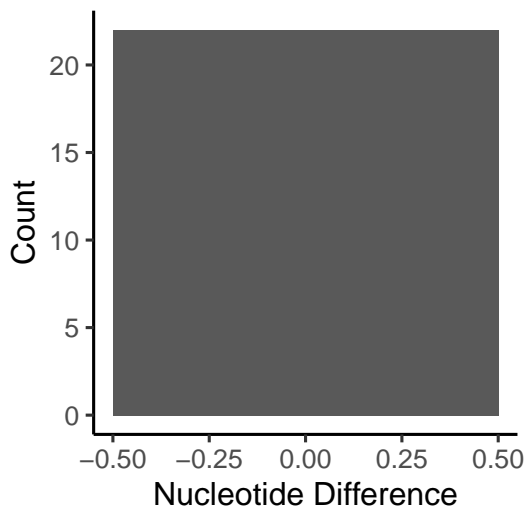
TRBV20–1*01

236 sequences assigned
235 (99.6%) exact matches, in which:
235 unique CDR3
13 unique J



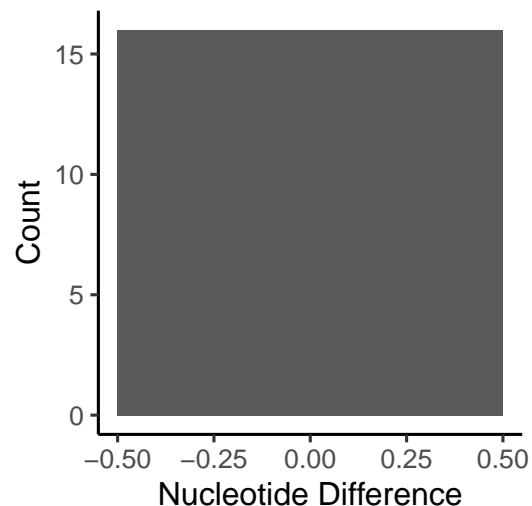
TRBV10–2*01

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



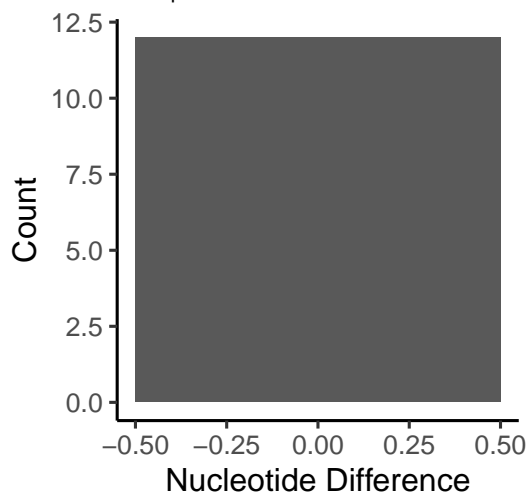
TRBV13*01

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



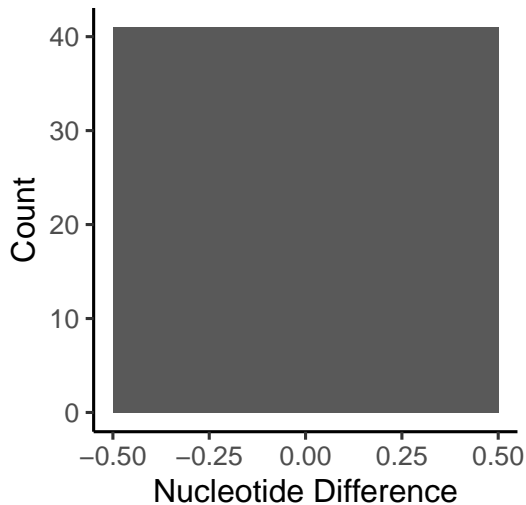
TRBV10–1*01

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



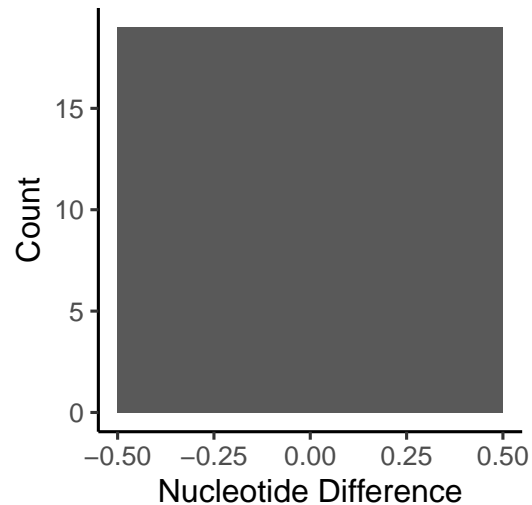
TRBV10–3*02

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



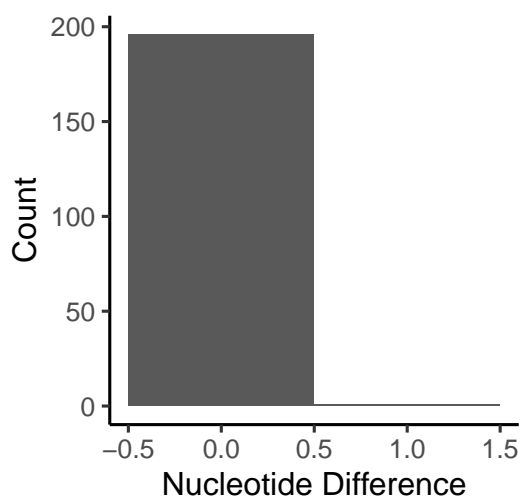
TRBV14*01

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



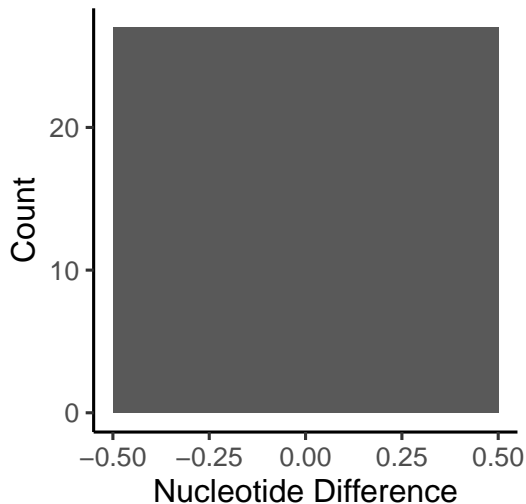
TRBV20–1*05

197 sequences assigned
196 (99.5%) exact matches, in which:
196 unique CDR3
12 unique J



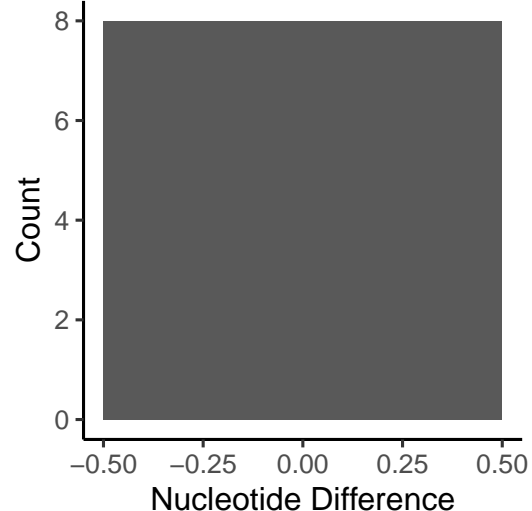
TRBV10–3*03_313GCCA1

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



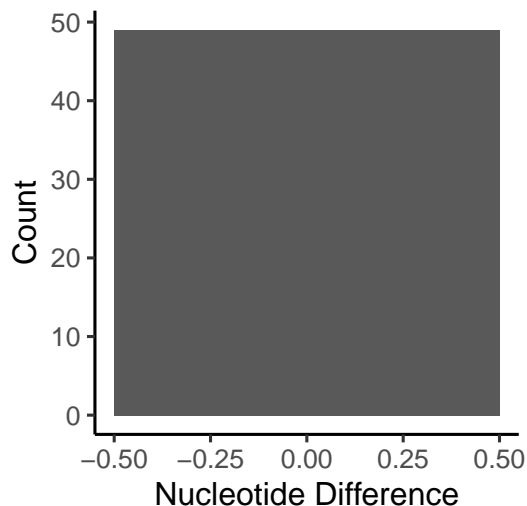
TRBV14*02_322CAAGA32

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



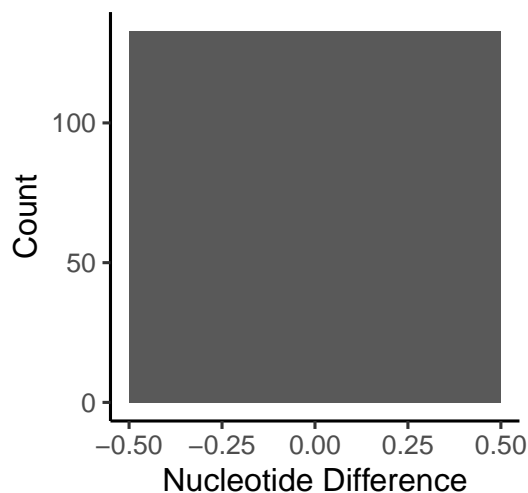
TRBV15*02

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



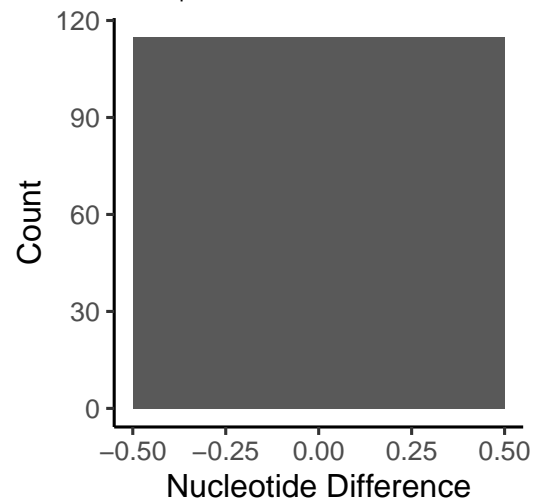
TRBV19*01

133 sequences assigned
133 (100%) exact matches, in which:
133 unique CDR3
13 unique J



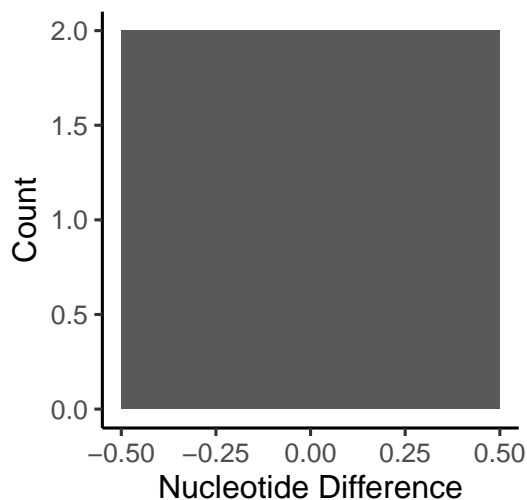
TRBV27*01

115 sequences assigned
115 (100%) exact matches, in which:
115 unique CDR3
12 unique J



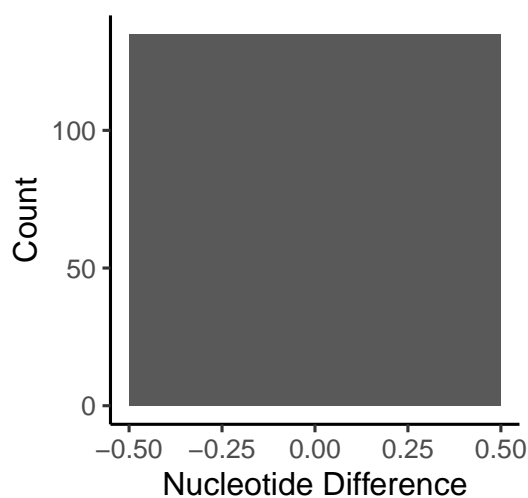
TRBV16*01

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



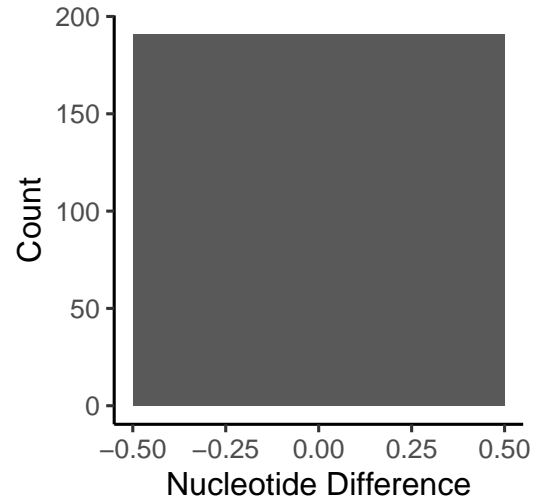
TRBV19*01_A24G

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



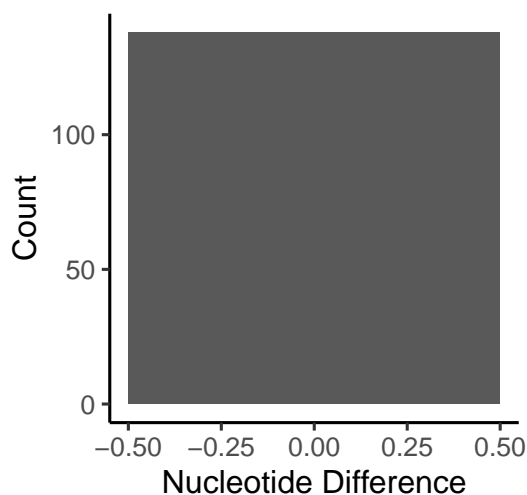
TRBV28*01

191 sequences assigned
191 (100%) exact matches, in which:
191 unique CDR3
13 unique J



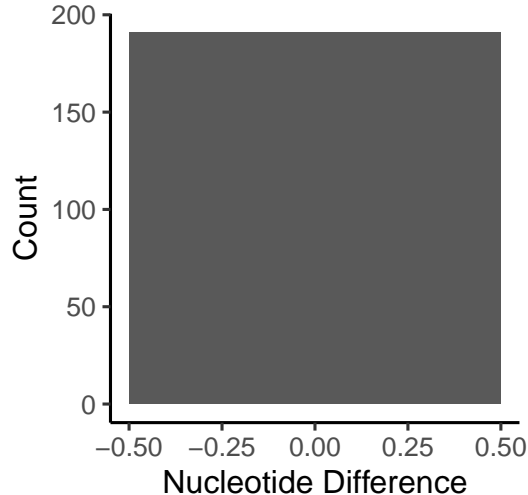
TRBV18*01

138 sequences assigned
138 (100%) exact matches, in which:
138 unique CDR3
13 unique J



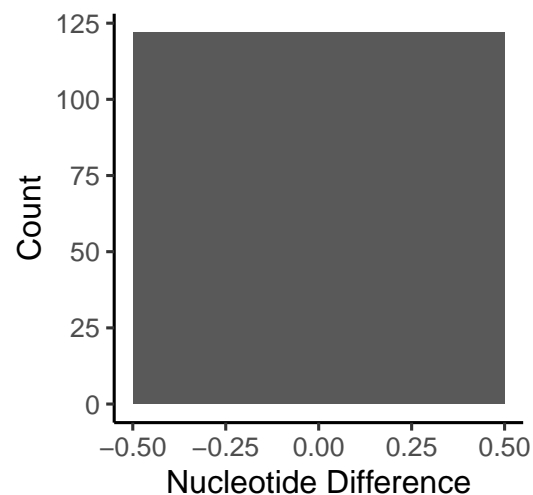
TRBV2*01

191 sequences assigned
191 (100%) exact matches, in which:
191 unique CDR3
13 unique J



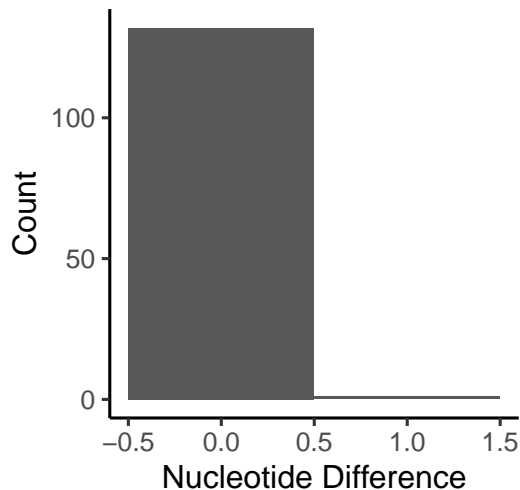
TRBV30*01

122 sequences assigned
122 (100%) exact matches, in which:
122 unique CDR3
13 unique J



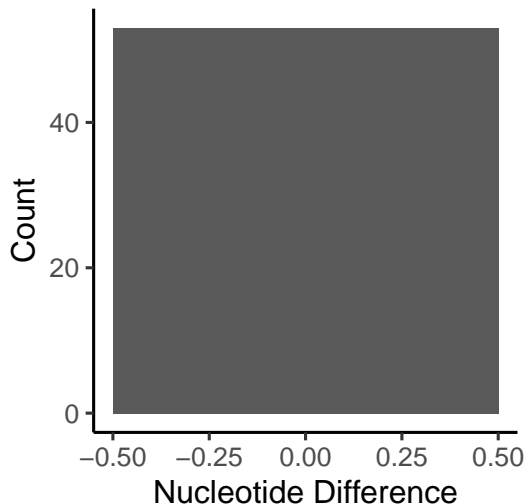
TRBV9*01

133 sequences assigned
132 (99.2%) exact matches, in which:
132 unique CDR3
12 unique J



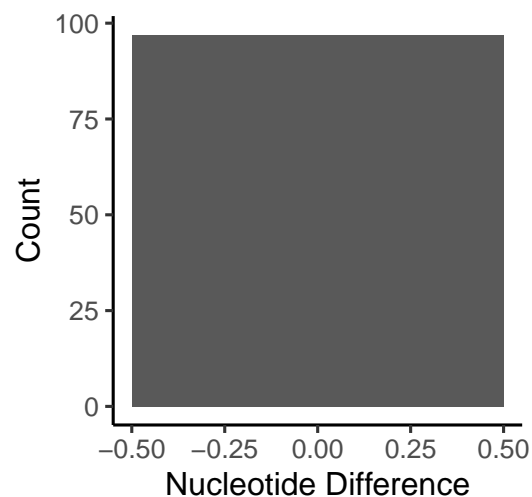
TRBV11-2*03

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



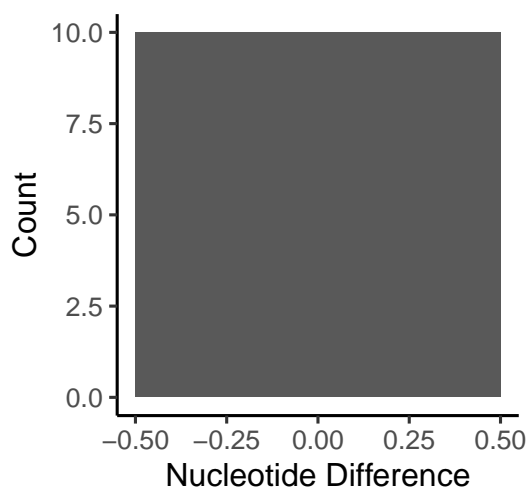
TRBV12-3*01

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



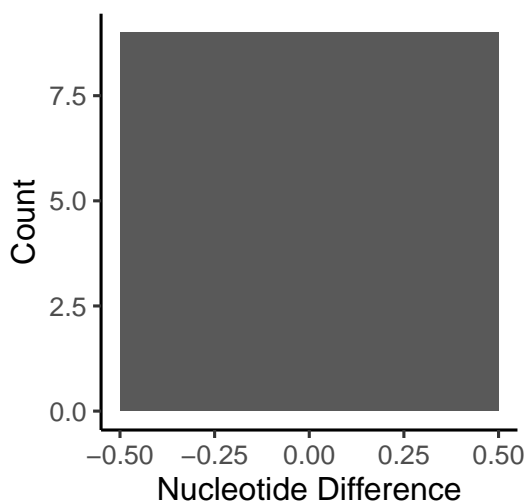
TRBV11-1*01

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



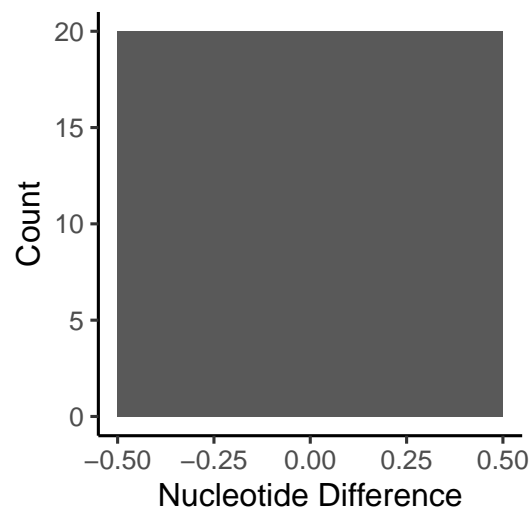
TRBV11-3*01

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



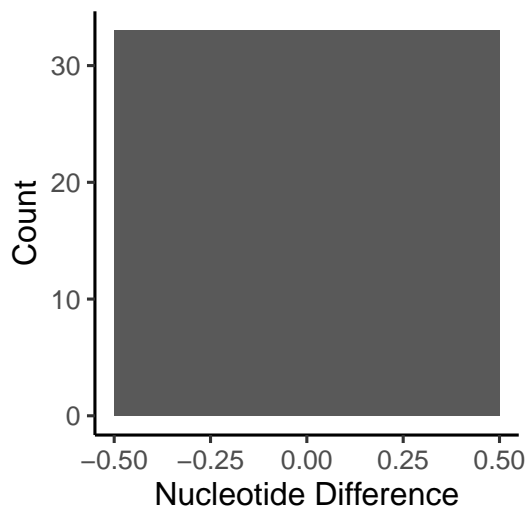
TRBV12-4*01

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



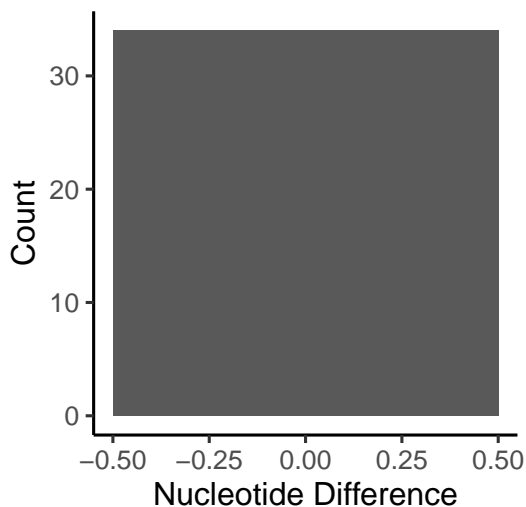
TRBV11-2*01

33 sequences assigned
33 (100%) exact matches, in which:
33 unique CDR3
10 unique J



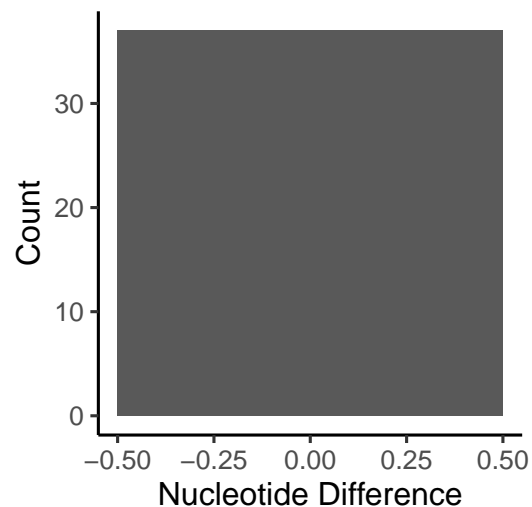
TRBV11-3*04

34 sequences assigned
34 (100%) exact matches, in which:
34 unique CDR3
9 unique J



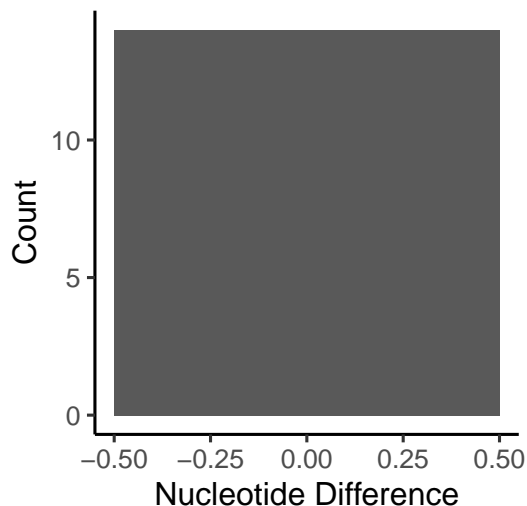
TRBV12-4*01_C87T

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



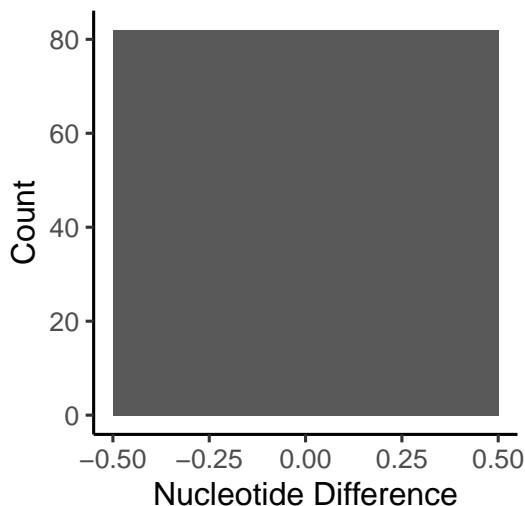
TRBV12–5*01

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



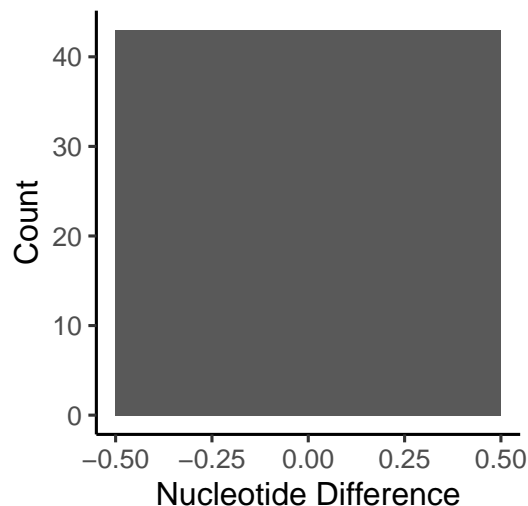
TRBV24–1*01

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



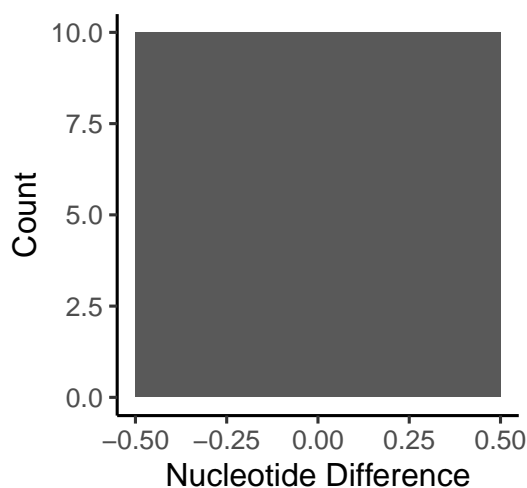
TRBV4–3*01

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



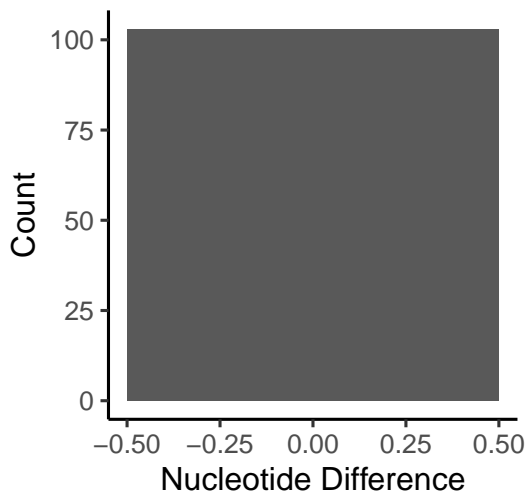
TRBV12–5*01_C28G

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



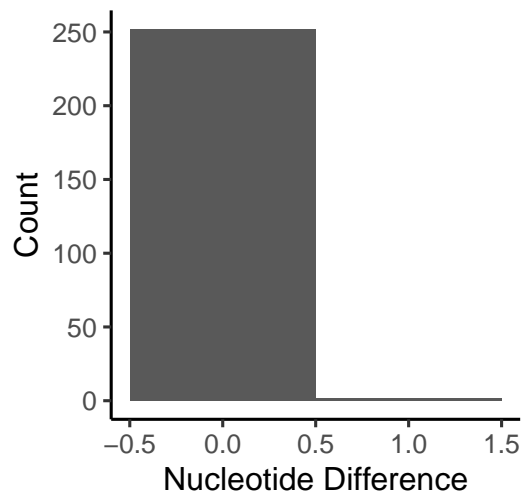
TRBV4–1*01

103 sequences assigned
103 (100%) exact matches, in which:
103 unique CDR3
12 unique J



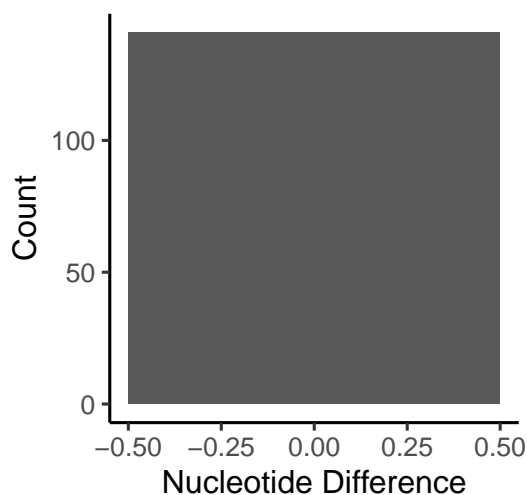
TRBV5–1*01

254 sequences assigned
252 (99.2%) exact matches, in which:
252 unique CDR3
13 unique J



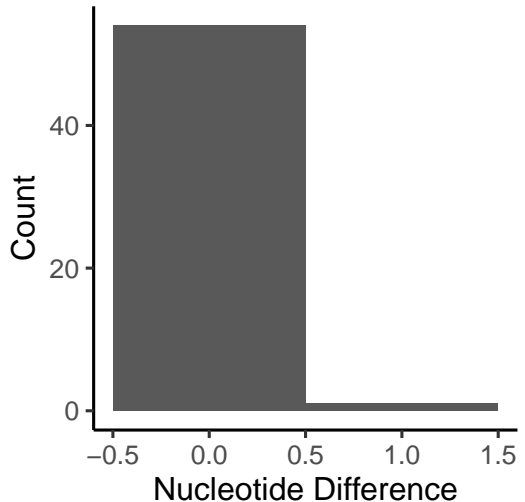
TRBV3–1*01

141 sequences assigned
141 (100%) exact matches, in which:
141 unique CDR3
12 unique J



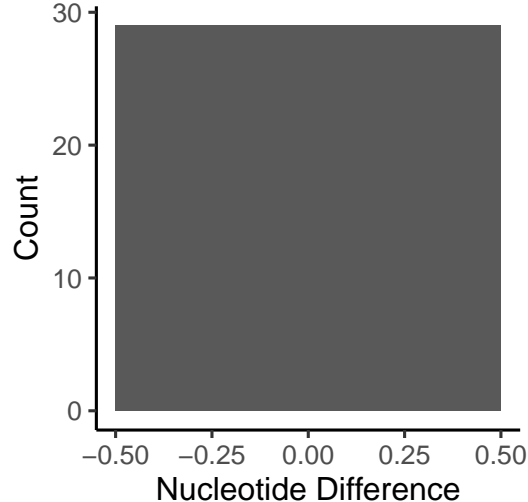
TRBV4–2*01

55 sequences assigned
54 (98.2%) exact matches, in which:
54 unique CDR3
10 unique J



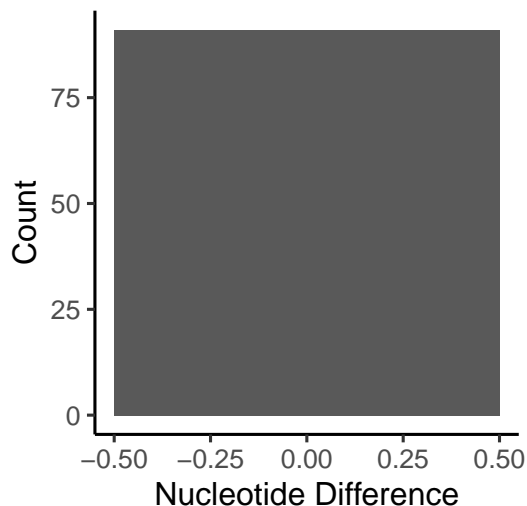
TRBV25–1*01

29 sequences assigned
29 (100%) exact matches, in which:
29 unique CDR3
9 unique J



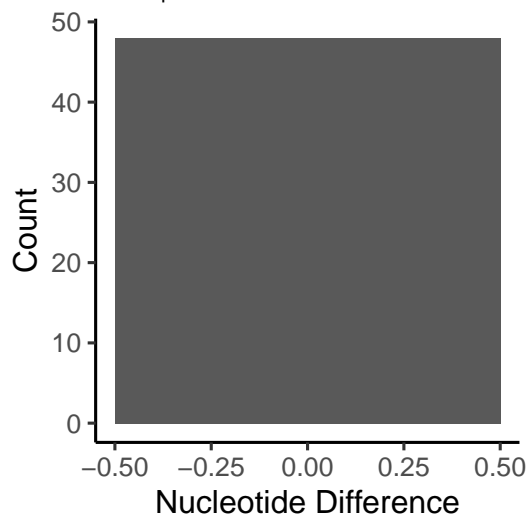
TRBV5-4*01

91 sequences assigned
91 (100%) exact matches, in which:
91 unique CDR3
12 unique J



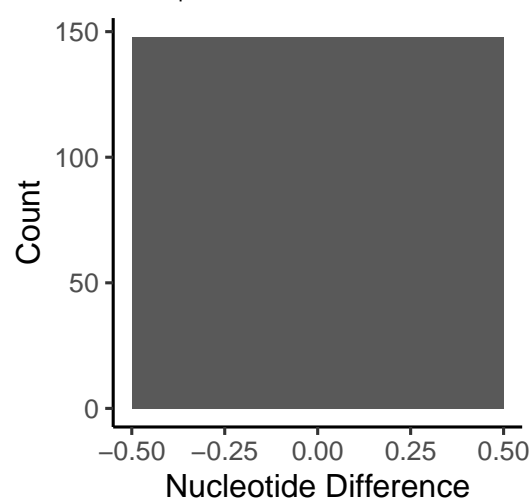
TRBV5-6*01

48 sequences assigned
48 (100%) exact matches, in which:
48 unique CDR3
11 unique J



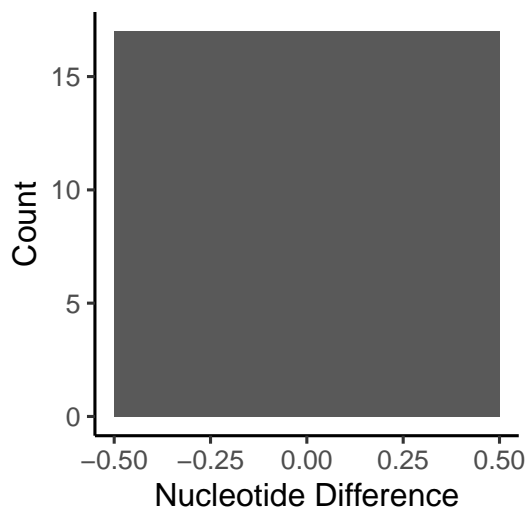
TRBV6-1*01

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



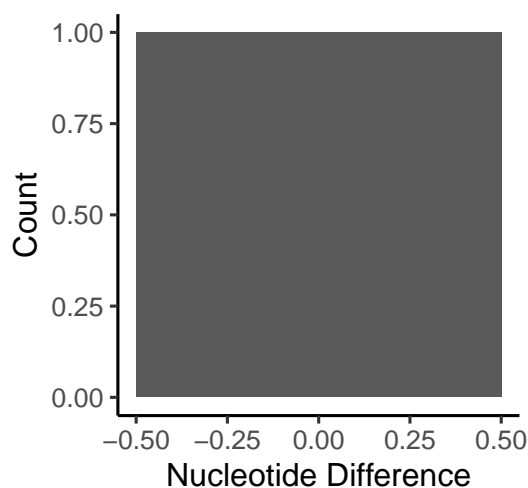
TRBV5-5*01

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



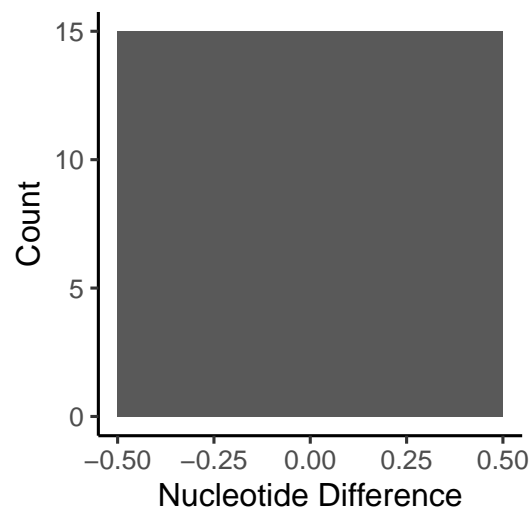
TRBV5-7*01

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



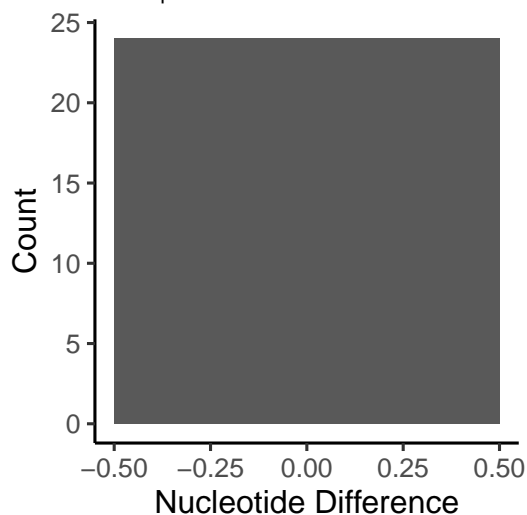
TRBV6-4*01

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



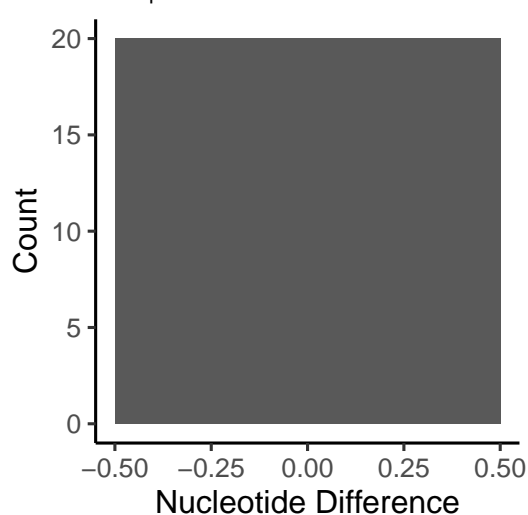
TRBV5-5*02_322TTGG32

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



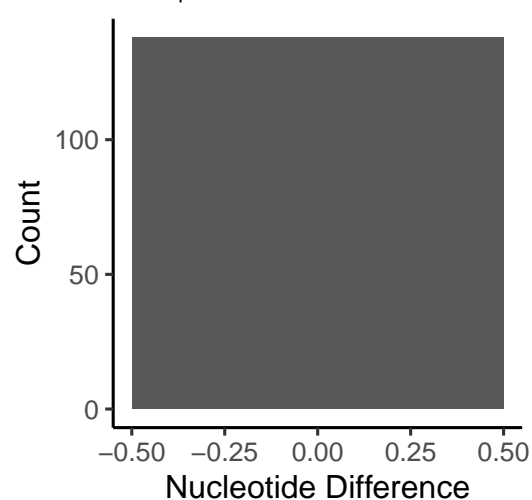
TRBV5-8*01

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



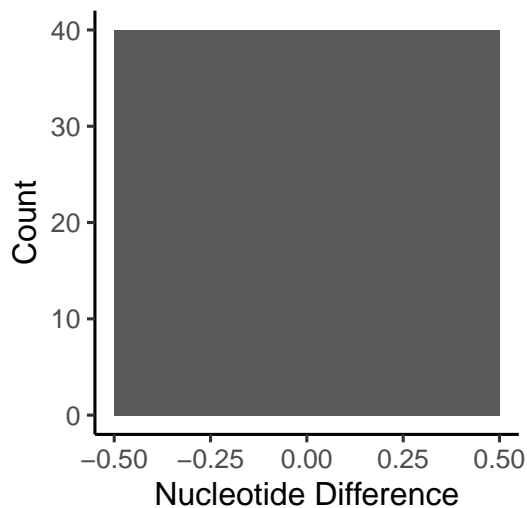
TRBV6-5*01

138 sequences assigned
138 (100%) exact matches, in which:
138 unique CDR3
13 unique J



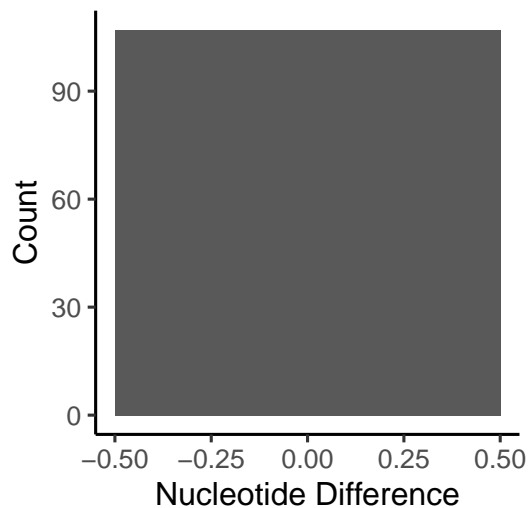
TRBV6-6*01

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



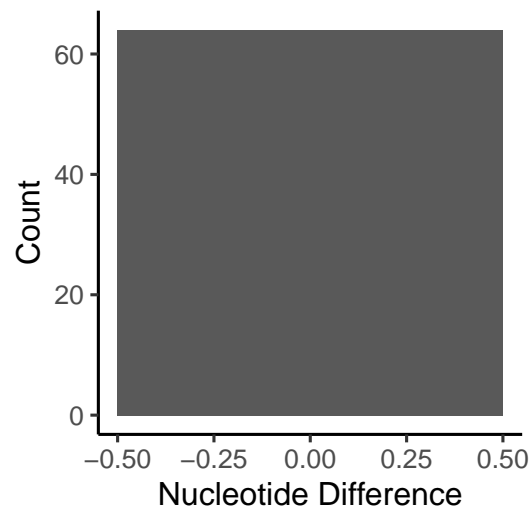
TRBV6-23*01

107 sequences assigned
107 (100%) exact matches, in which:
107 unique CDR3
12 unique J



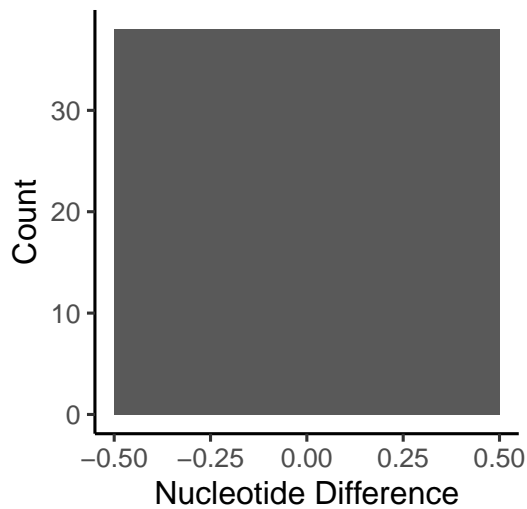
TRBV7-3*01

64 sequences assigned
64 (100%) exact matches, in which:
64 unique CDR3
11 unique J



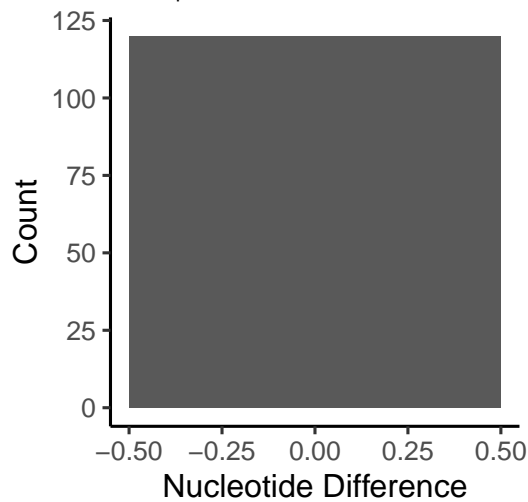
TRBV6-6*02

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



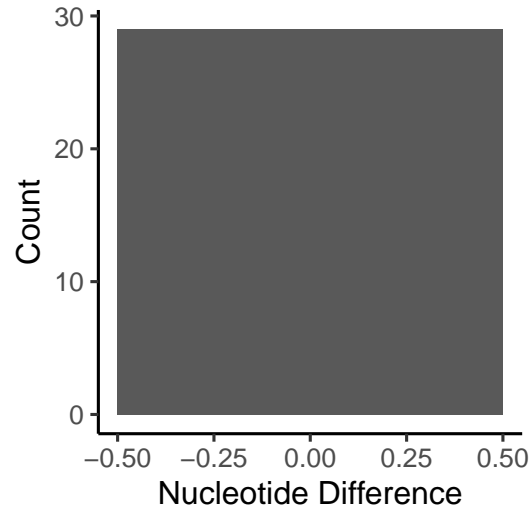
TRBV7-2*01

120 sequences assigned
120 (100%) exact matches, in which:
120 unique CDR3
13 unique J



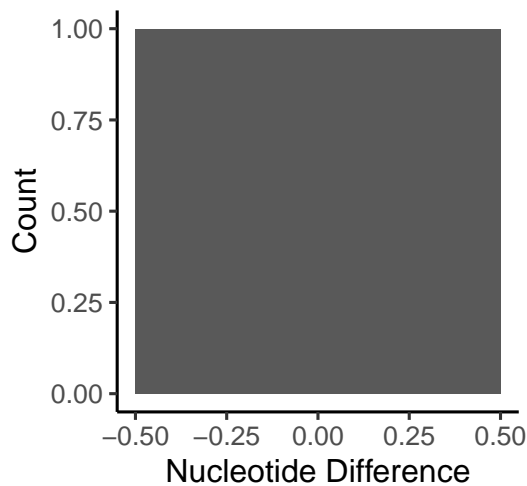
TRBV7-6*01

29 sequences assigned
29 (100%) exact matches, in which:
29 unique CDR3
8 unique J



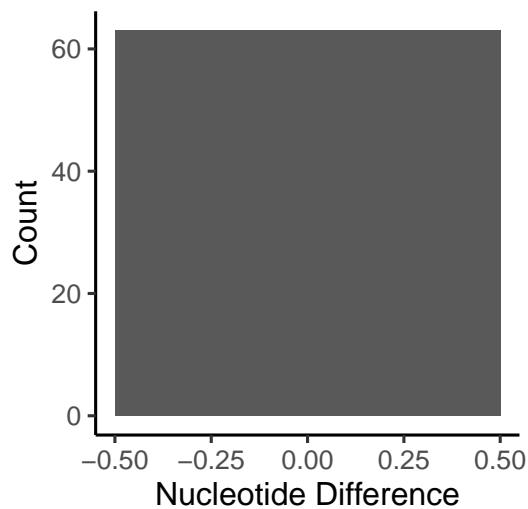
TRBV6-7*01

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



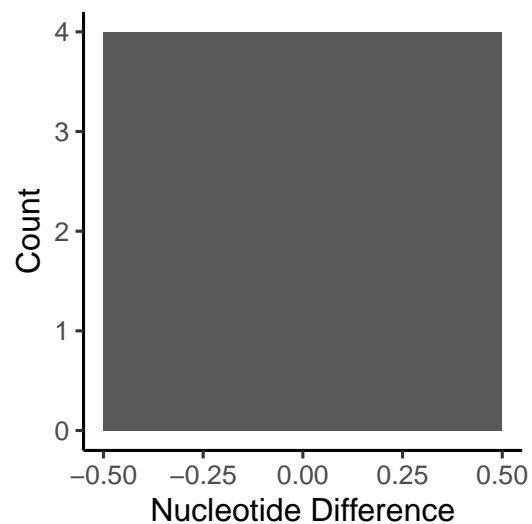
TRBV7-2*02

63 sequences assigned
63 (100%) exact matches, in which:
63 unique CDR3
12 unique J



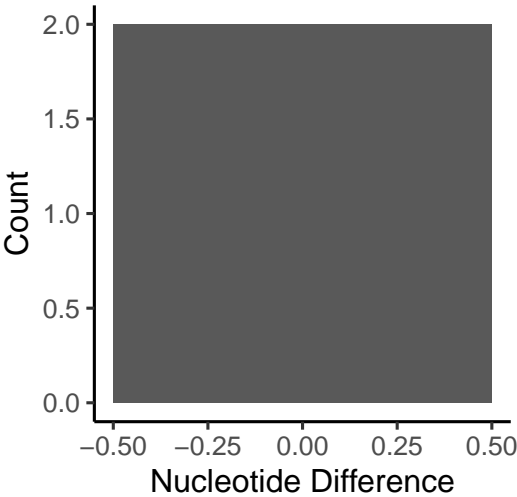
TRBV7-7*01

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



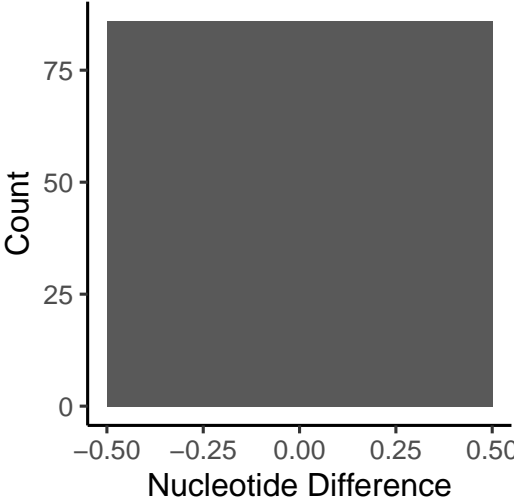
TRBV7-7*01_C315T

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



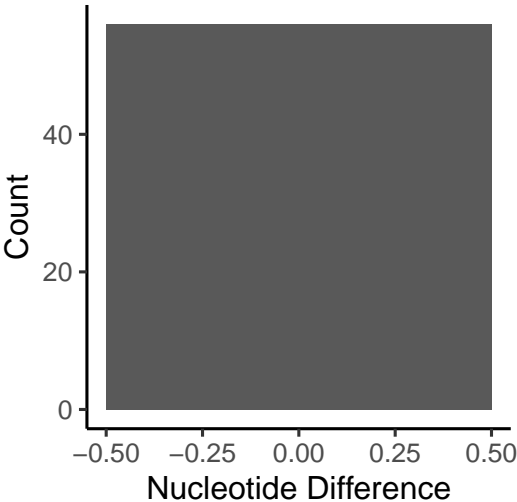
TRBV29-1*01

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



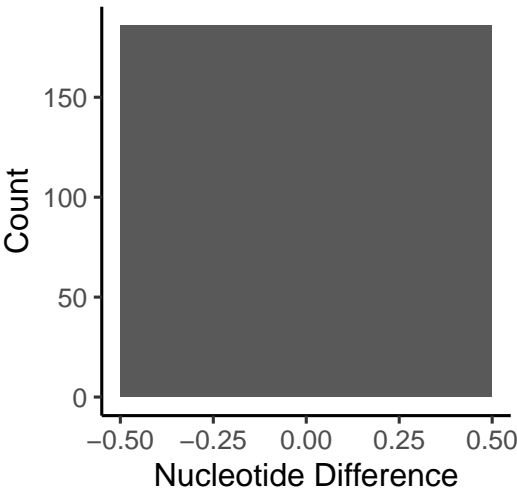
TRBV7-8*01

56 sequences assigned
56 (100%) exact matches, in which:
56 unique CDR3
10 unique J

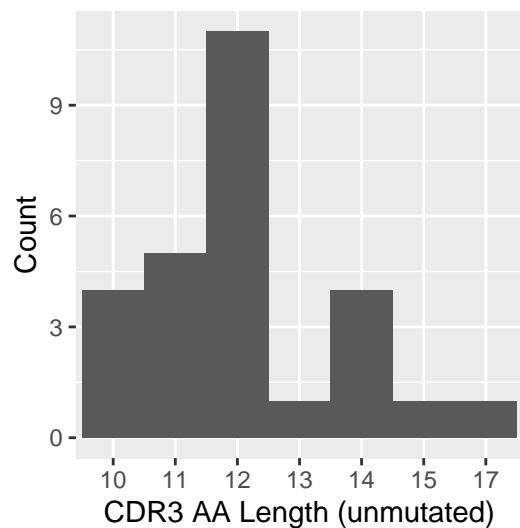


TRBV7-9*03

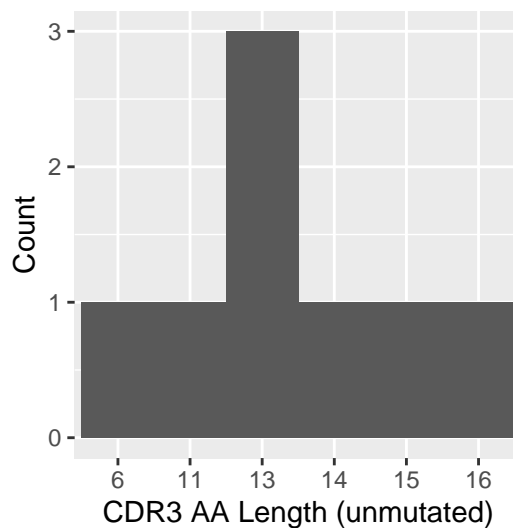
186 sequences assigned
186 (100%) exact matches, in which:
186 unique CDR3
13 unique J



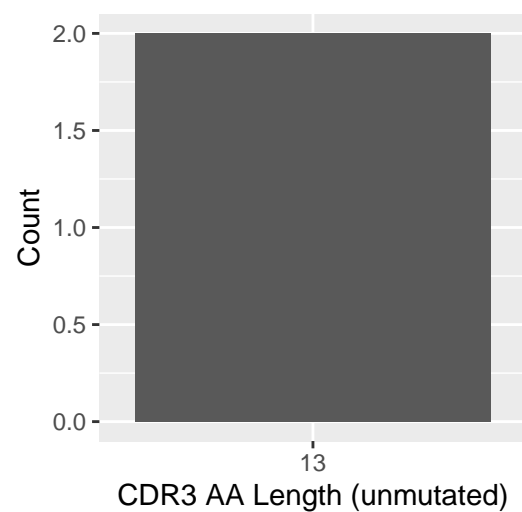
TRBV10-3*03_313GCCATCA



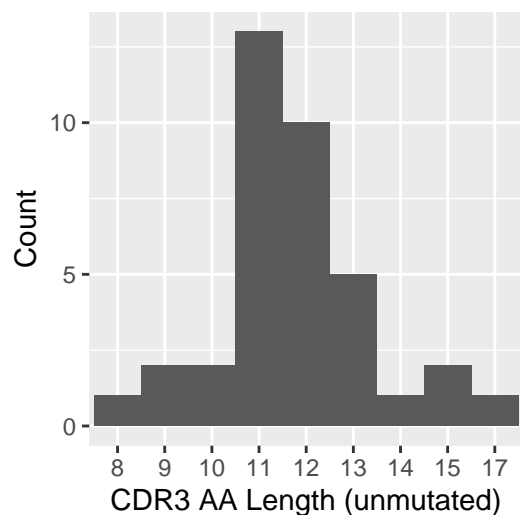
TRBV14*02_322CAAGA326



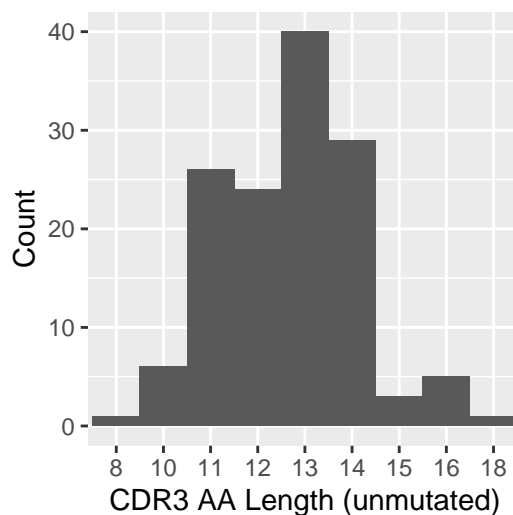
TRBV7-7*01_C315T



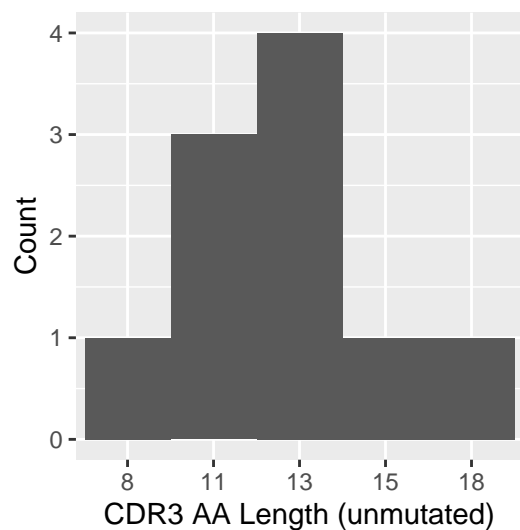
TRBV12-4*01_C87T



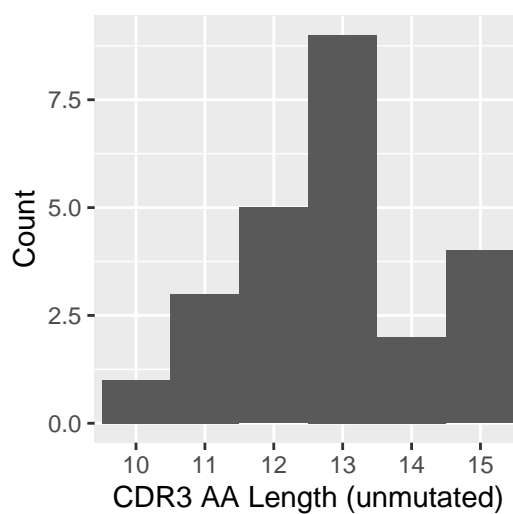
TRBV19*01_A24G

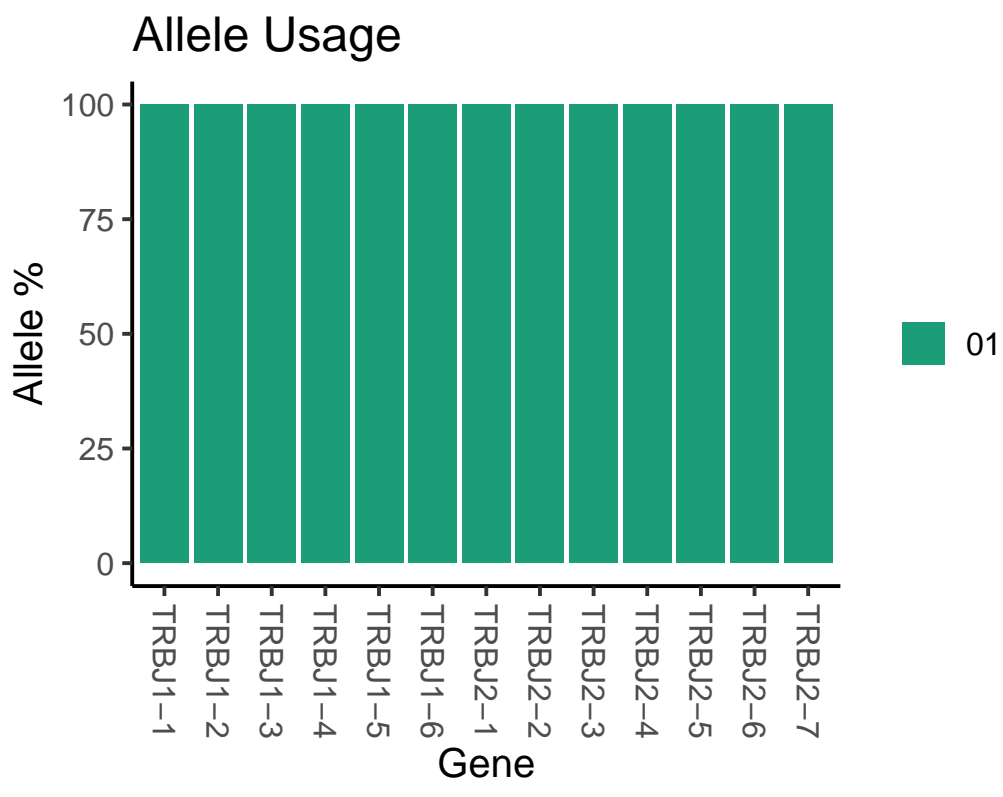


TRBV12-5*01_C28G



TRBV5-5*02_322TTGG325





Warning – no inferred sequences found.

Warning – no inferred sequences found.