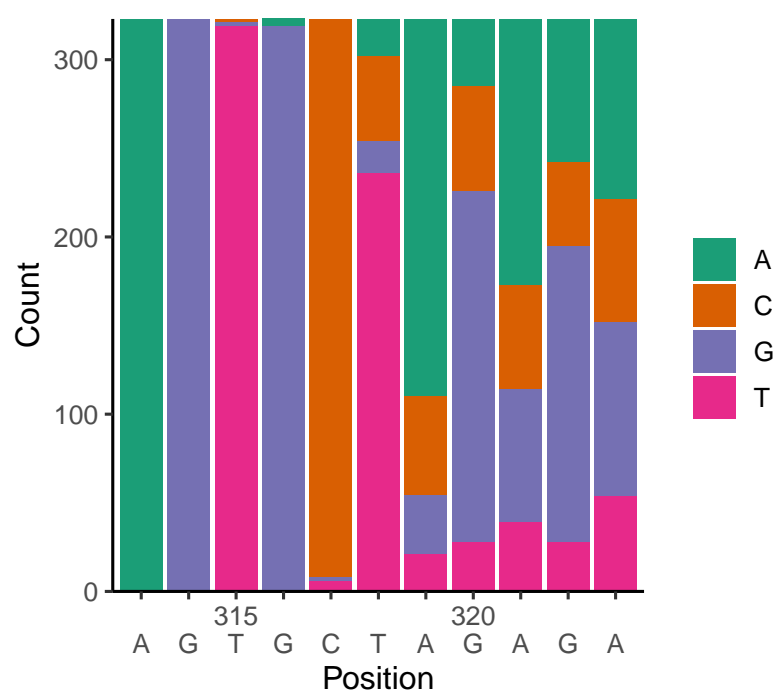
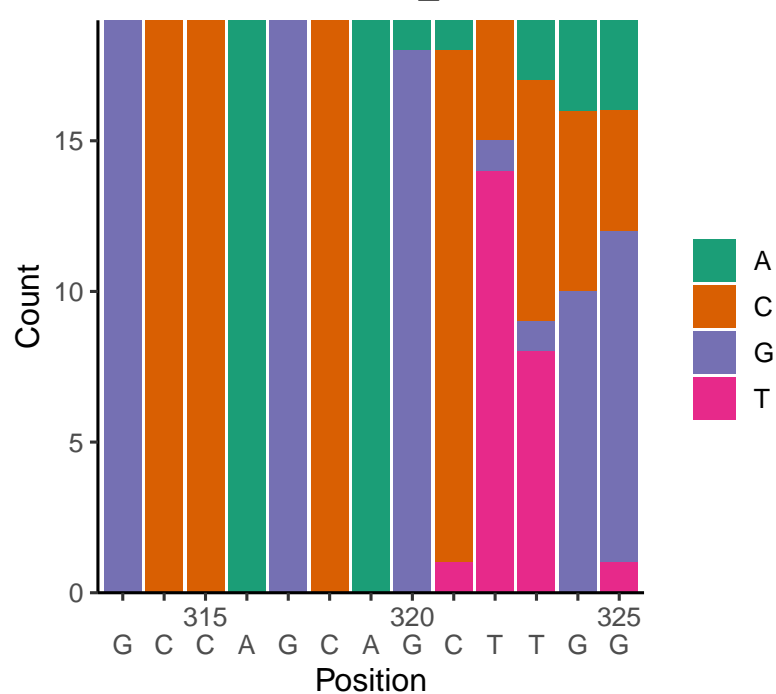


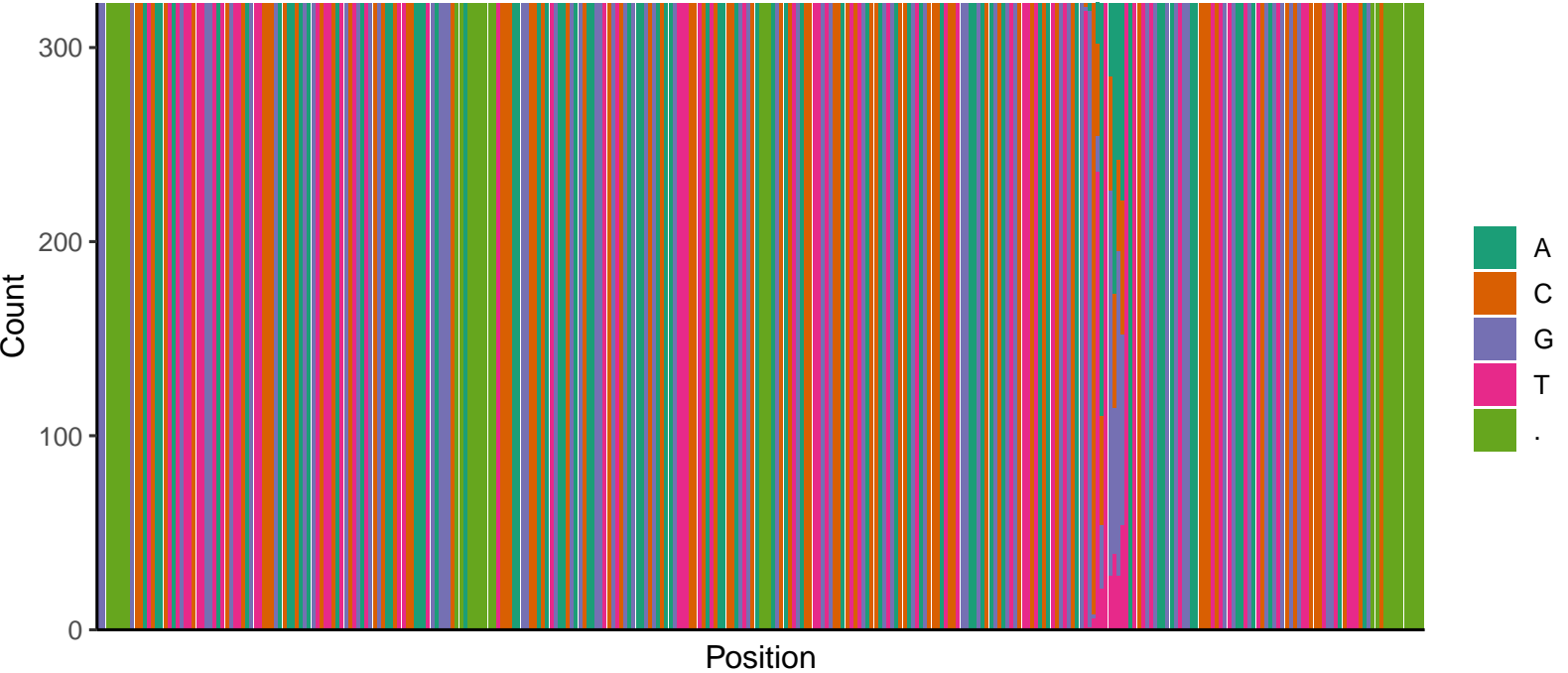
Gene TRBV20-1\*02\_319AGAGA323



Gene TRBV5-5\*02\_322TTGG325



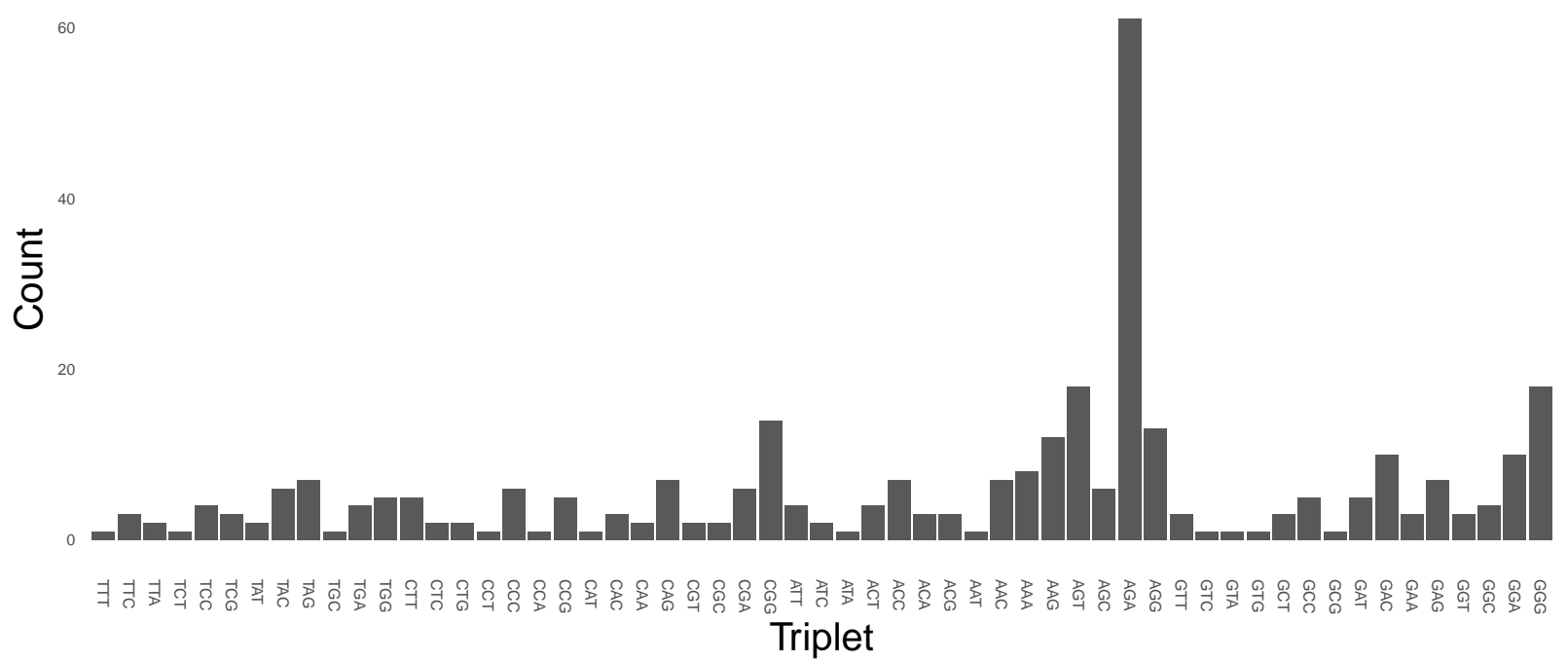
Gene TRBV20-1\*02\_319AGAGA323



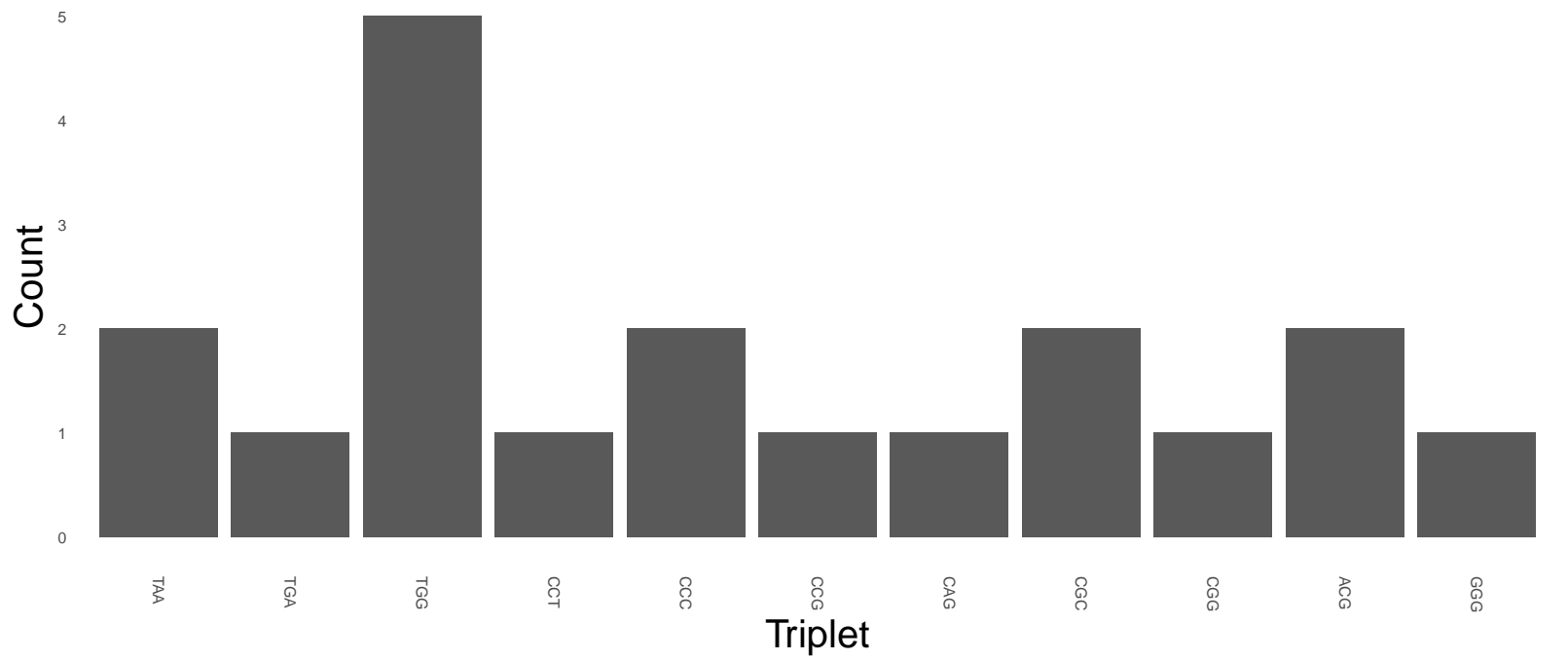
Gene TRBV5-5\*02\_322TTGG325



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

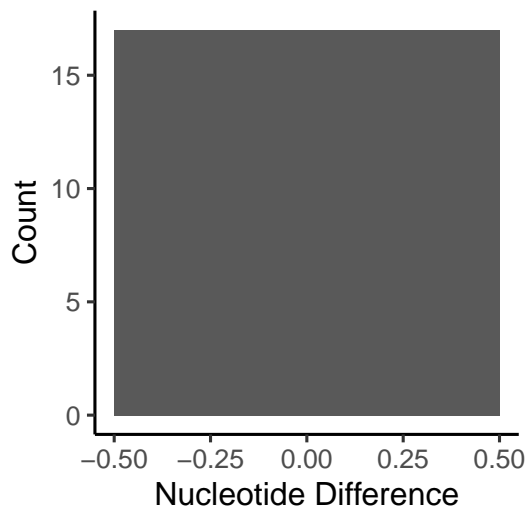


TRBV5–5\*02\_322TTGG325– Final 3 nucleotides as a triplet



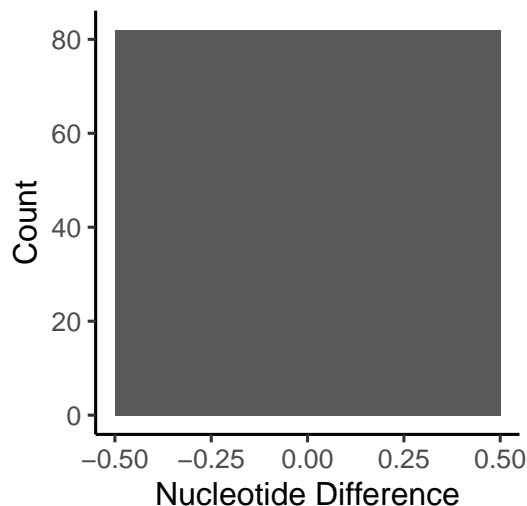
### TRBV10–1\*01

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



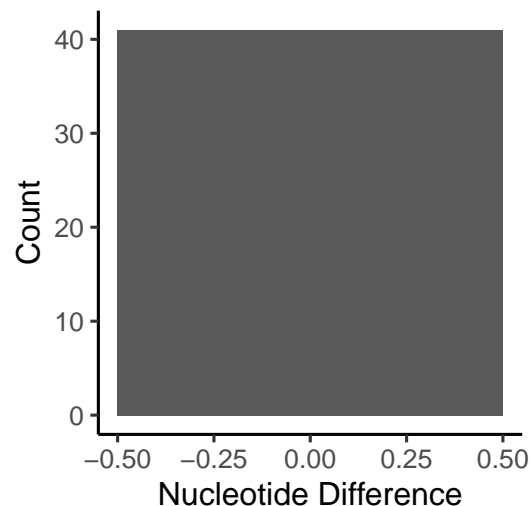
### TRBV10–3\*01

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



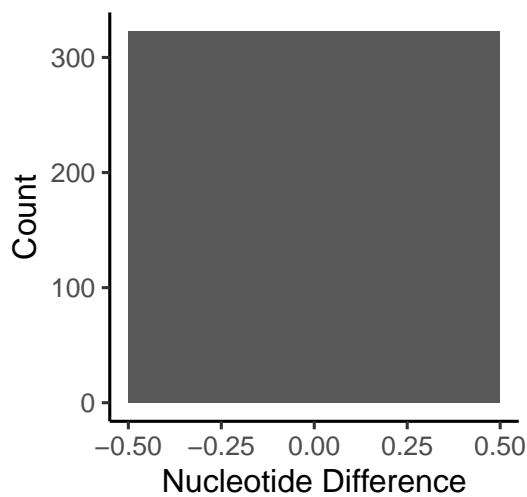
### TRBV15\*02

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



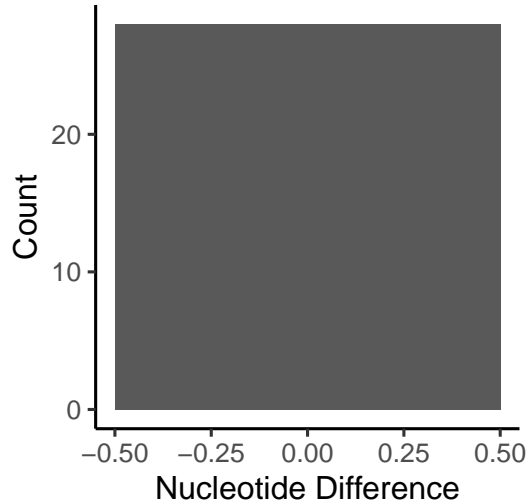
### TRBV20–1\*02\_319AGAG

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



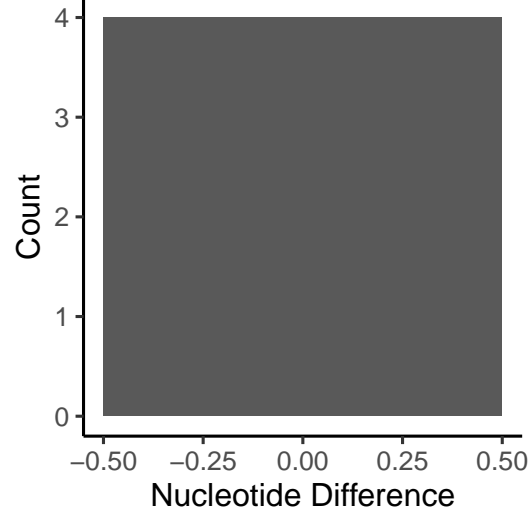
### TRBV13\*01

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



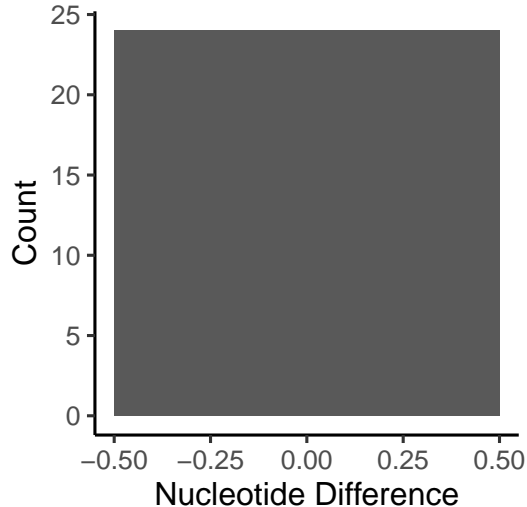
### TRBV16\*01

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



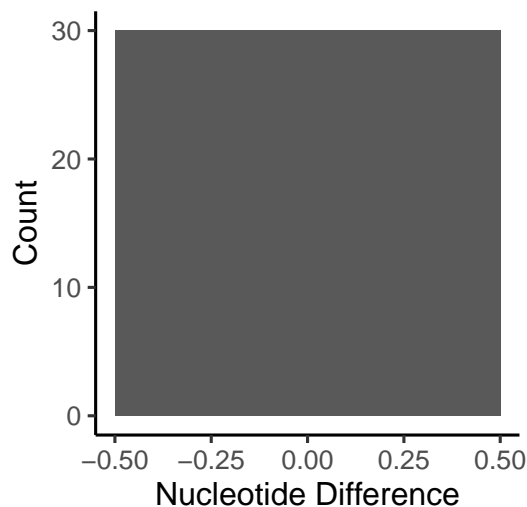
### TRBV10–2\*01

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



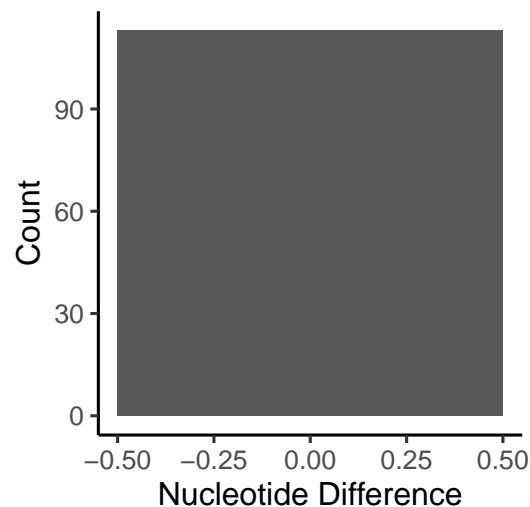
### TRBV14\*01

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



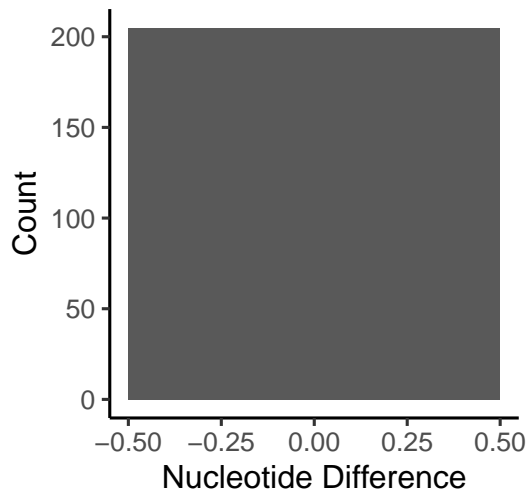
### TRBV18\*01

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



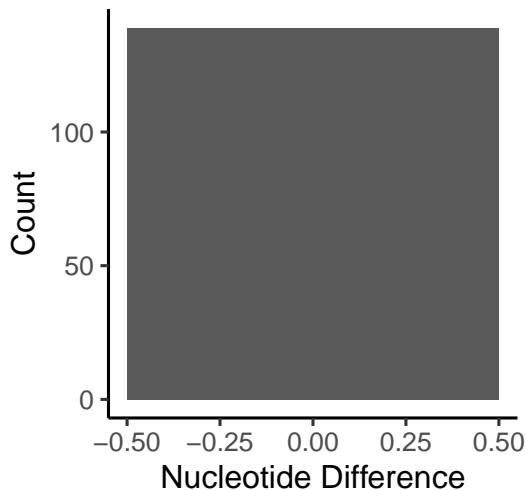
### TRBV19\*01

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



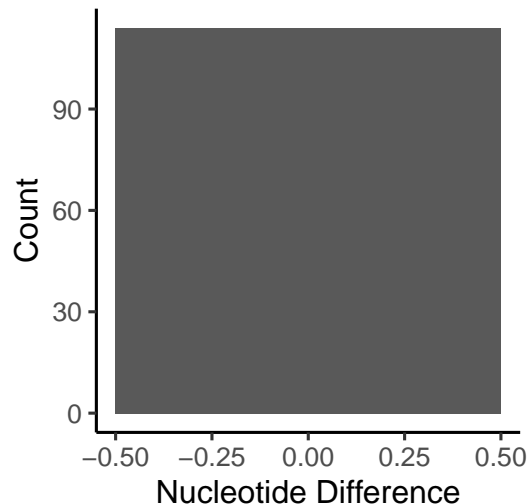
### TRBV28\*01

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



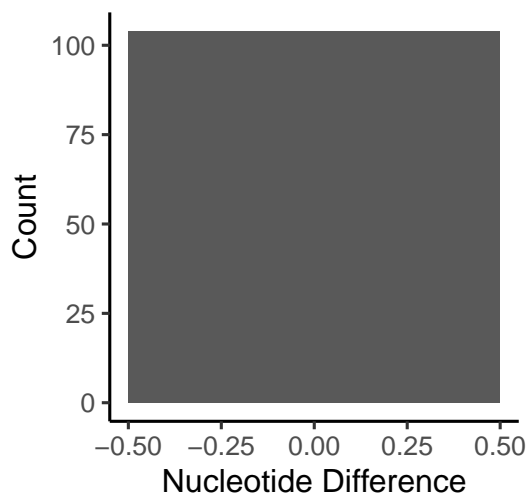
### TRBV11-2\*01

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



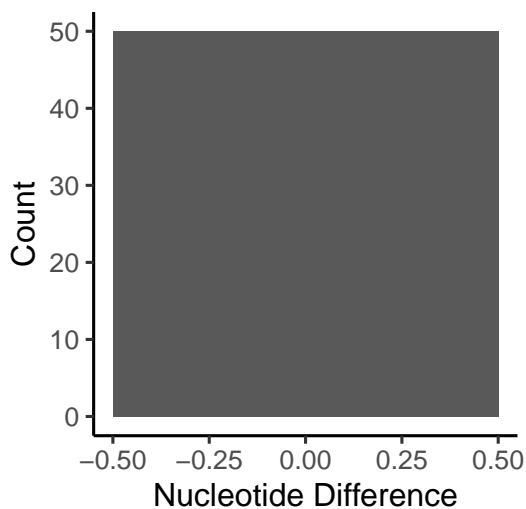
### TRBV2\*01

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



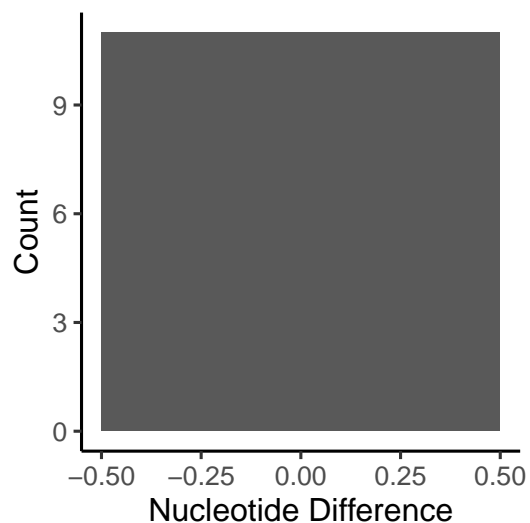
### TRBV30\*01

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



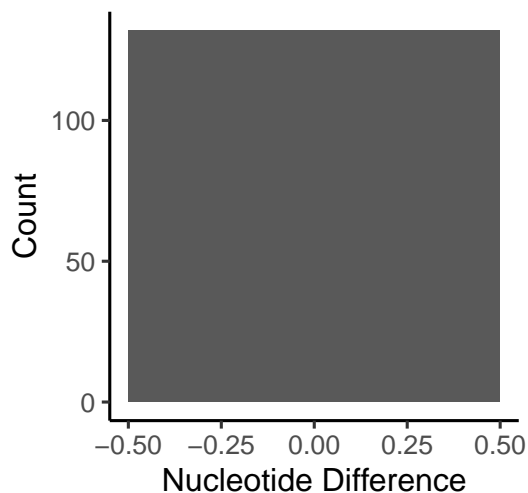
### TRBV11-3\*01

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



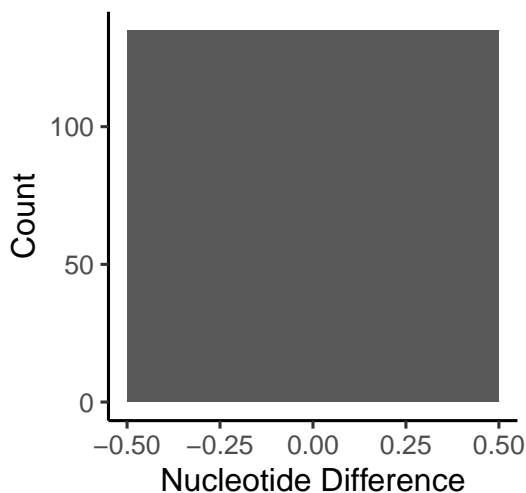
### TRBV27\*01

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



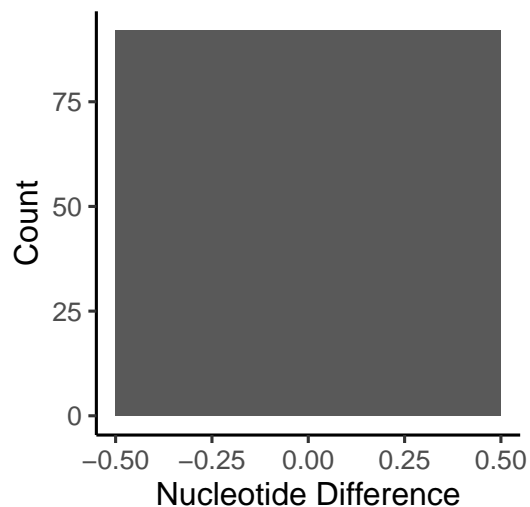
### TRBV9\*01

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



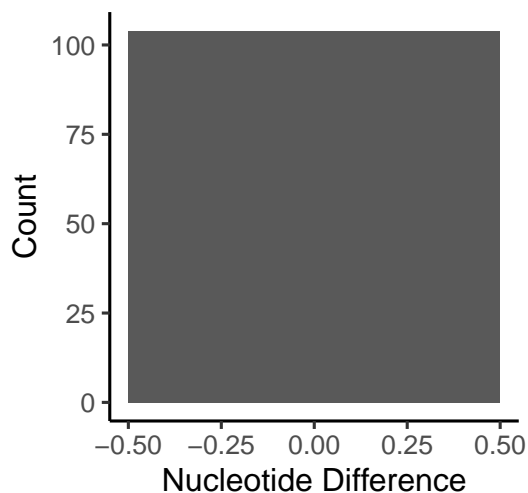
### TRBV12-3\*01

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



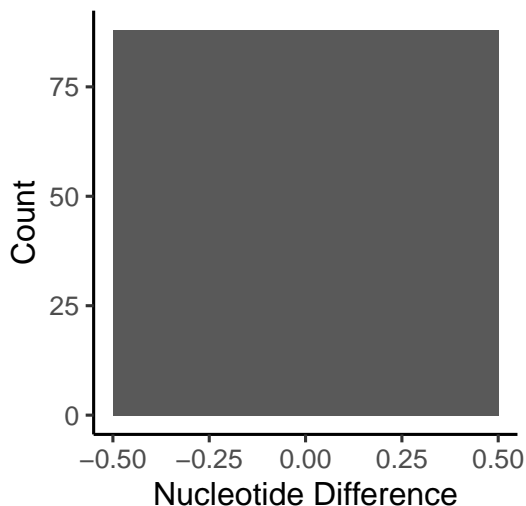
### TRBV12–4\*01

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



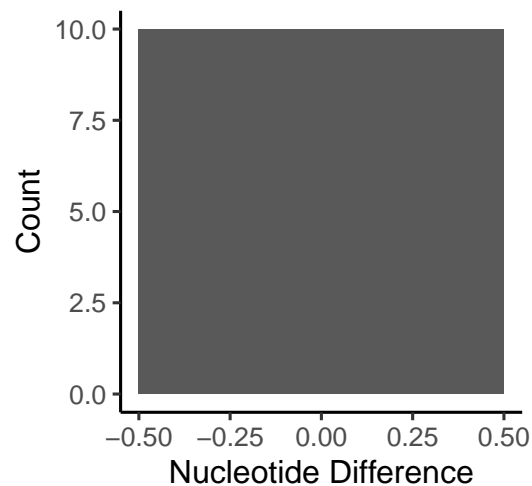
### TRBV4–1\*01

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



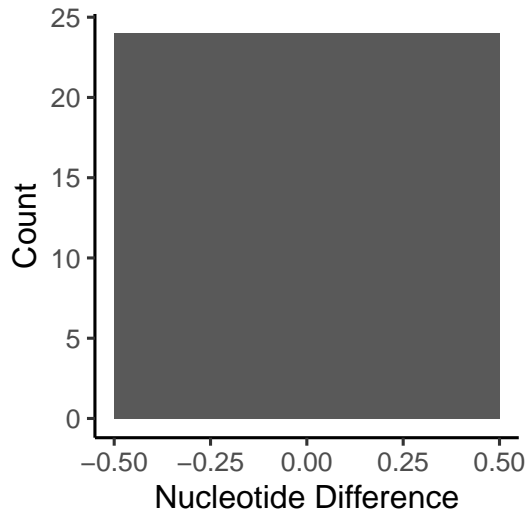
### TRBV4–2\*01

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



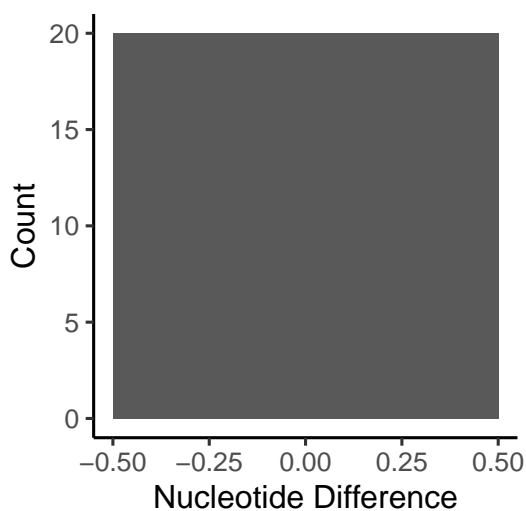
### TRBV12–5\*01

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



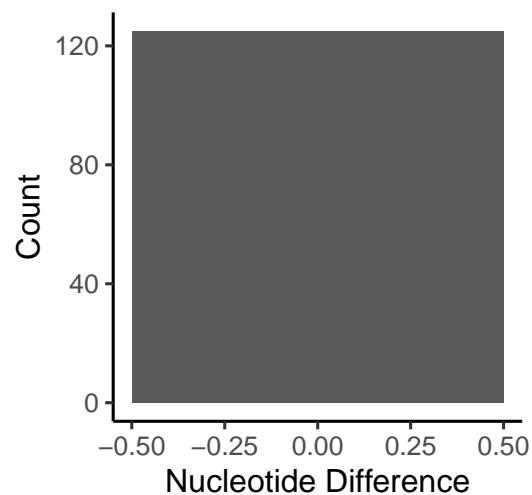
### TRBV24–1\*01

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



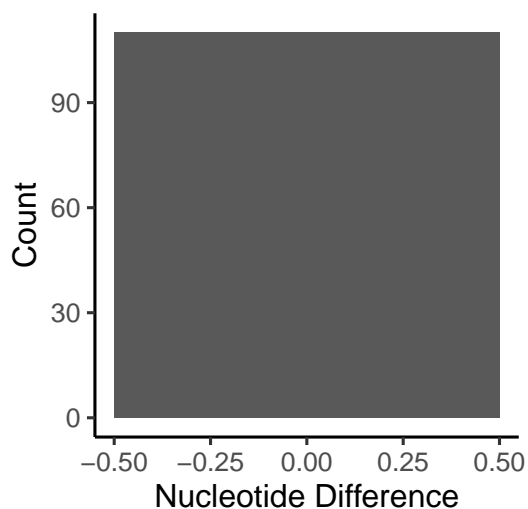
### TRBV4–3\*01

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



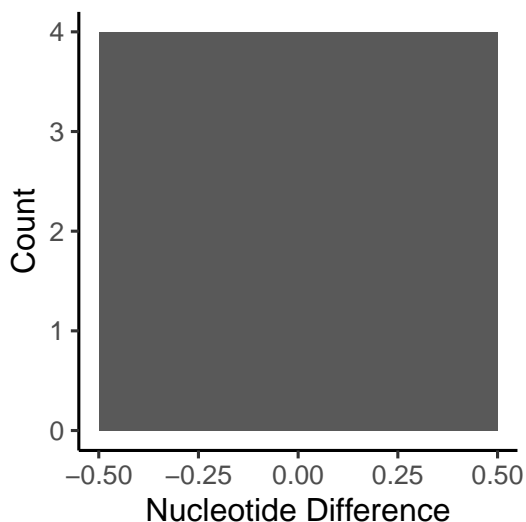
### TRBV3–1\*01

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



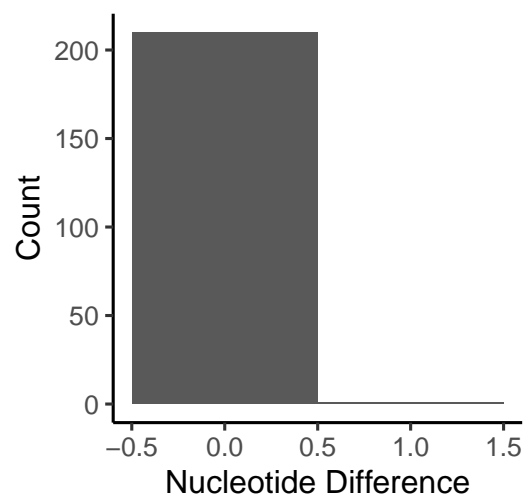
### TRBV24–1\*02

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



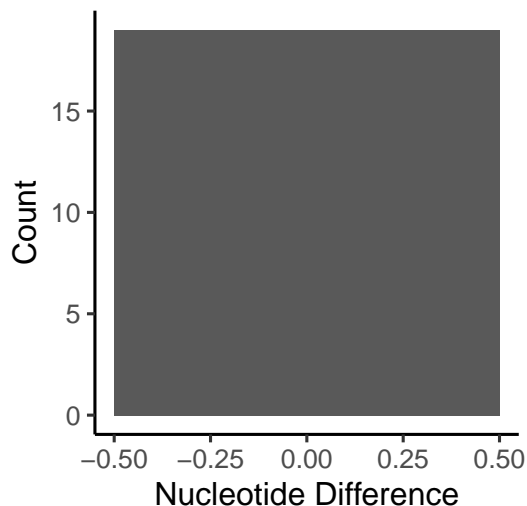
### TRBV5–1\*01

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



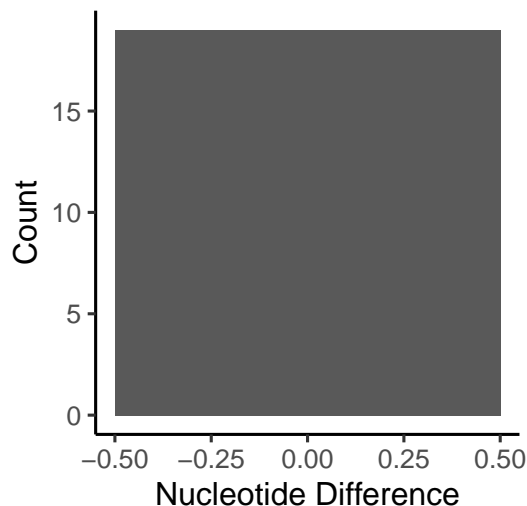
### TRBV25–1\*01

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



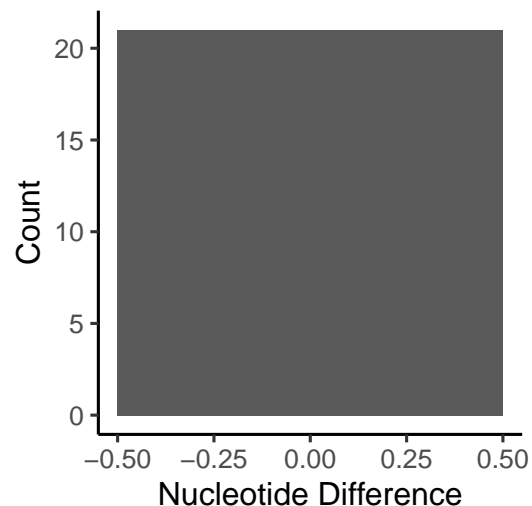
### TRBV5–5\*02\_322TTGG32

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



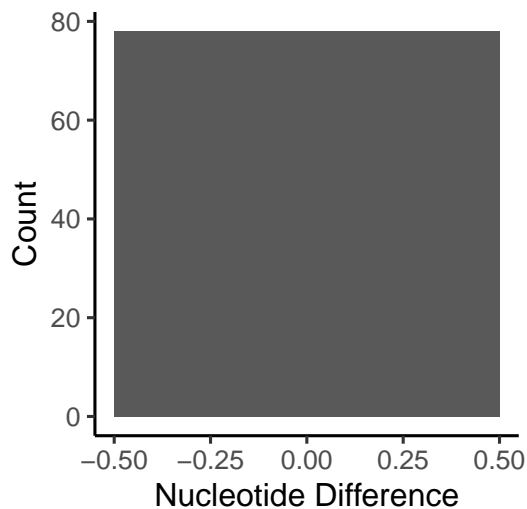
### TRBV5–8\*01

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



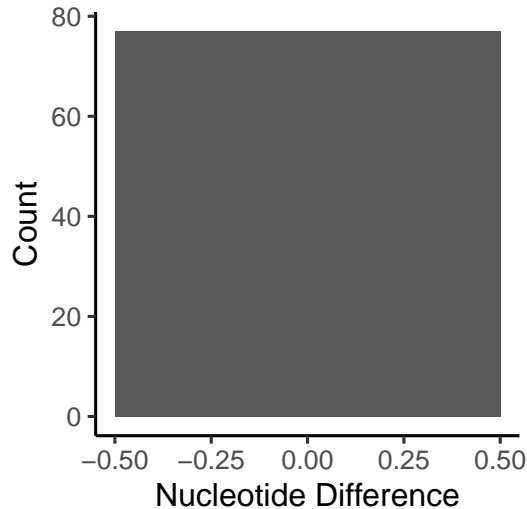
### TRBV5–4\*01

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



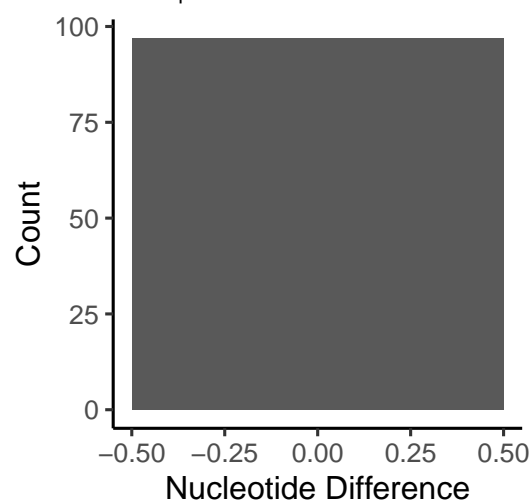
### TRBV5–6\*01

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



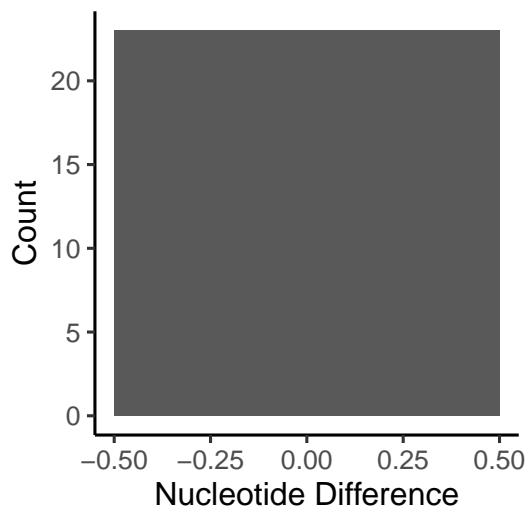
### TRBV6–1\*01

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



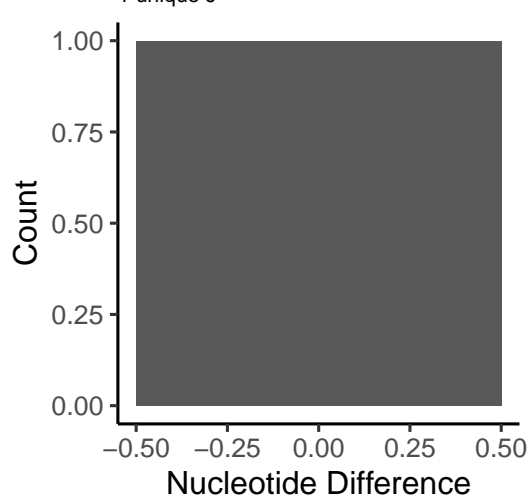
### TRBV5–5\*01

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



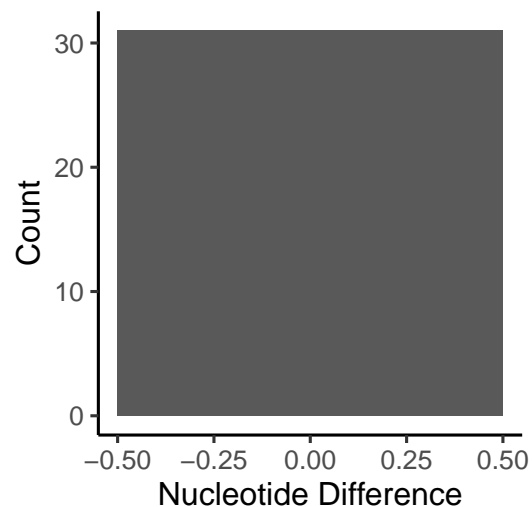
### TRBV5–7\*01

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



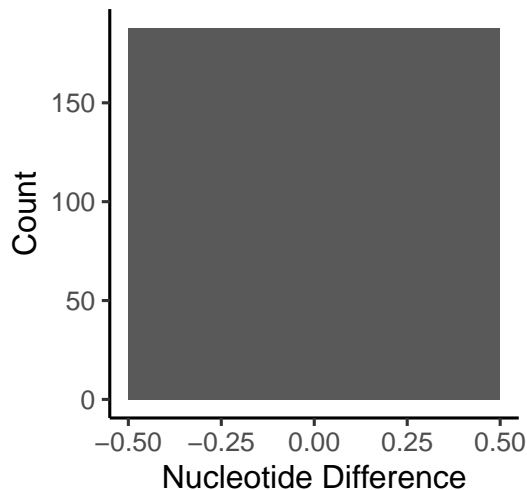
### TRBV6–4\*01

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



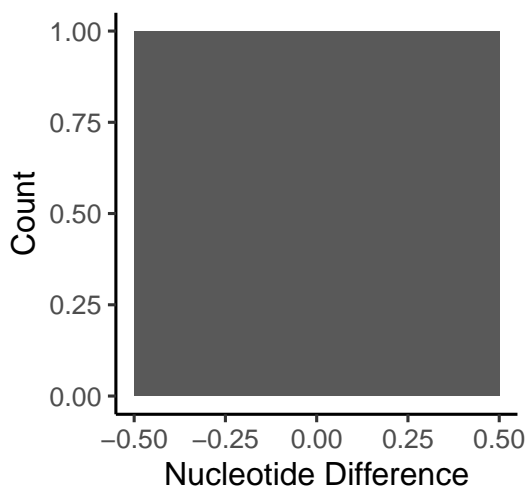
### TRBV6–5\*01

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



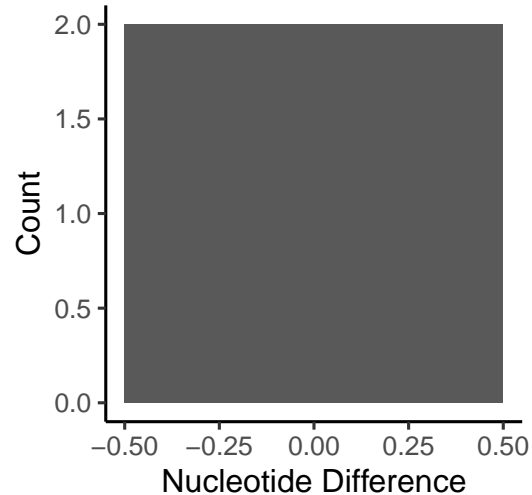
### TRBV6–8\*01

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



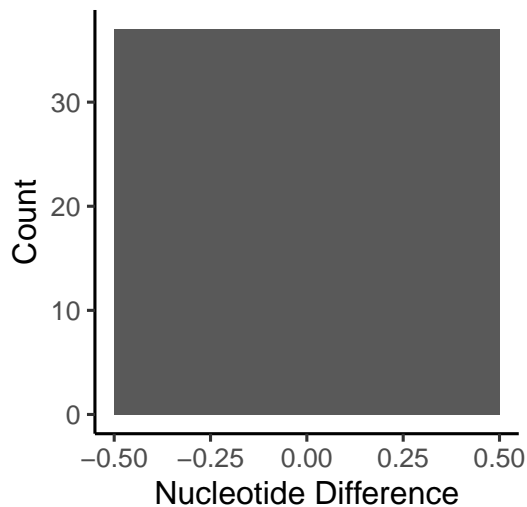
### TRBV7–4\*01

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



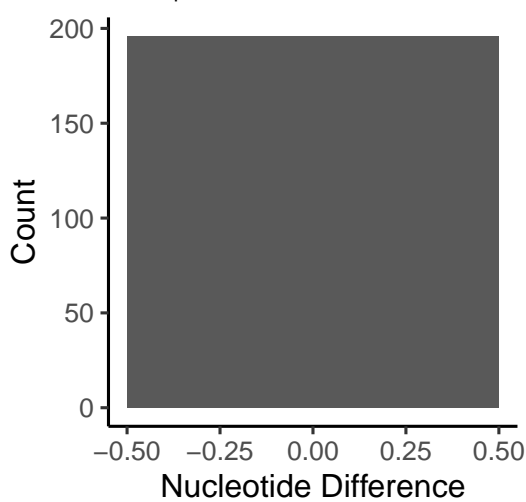
### TRBV6–6\*01

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



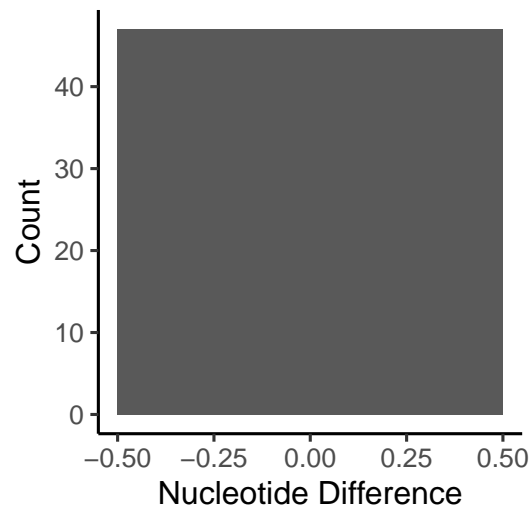
### TRBV6–23\*01

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



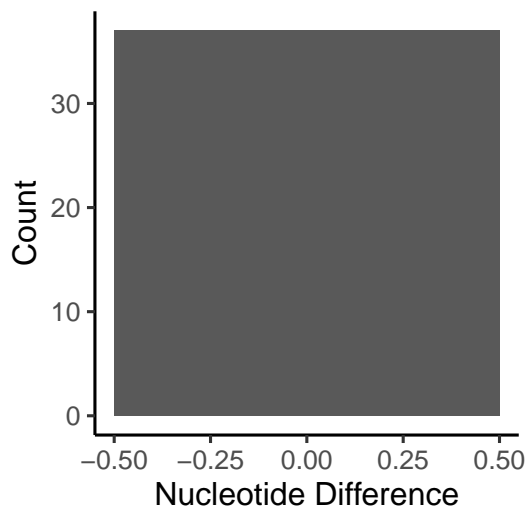
### TRBV7–6\*01

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



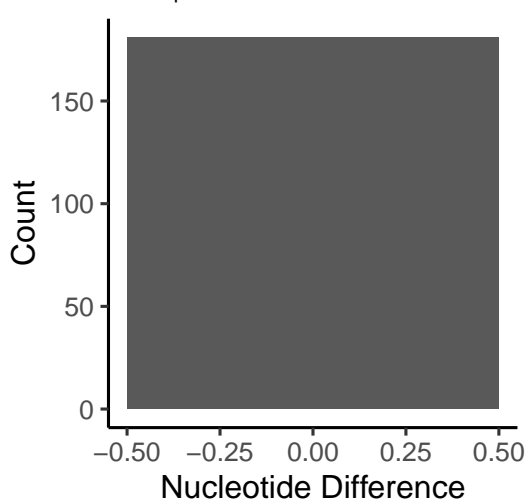
### TRBV6–6\*02

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



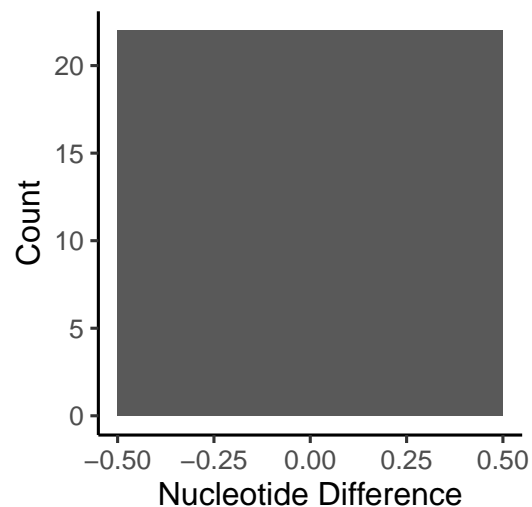
### TRBV7–2\*01

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



### TRBV7–7\*01

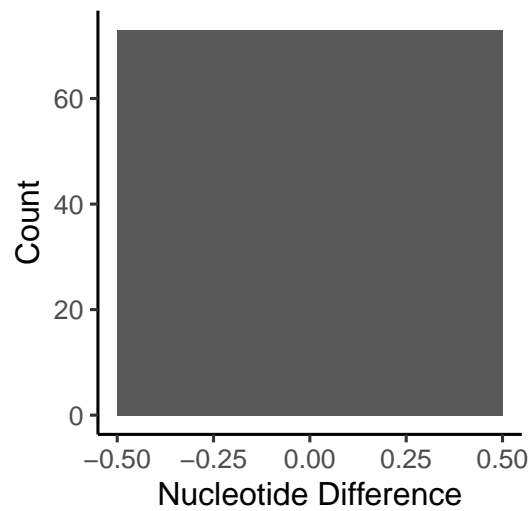
22 sequences assigned  
22 (100%) exact matches, in which:  
22 unique CDR3  
8 unique J





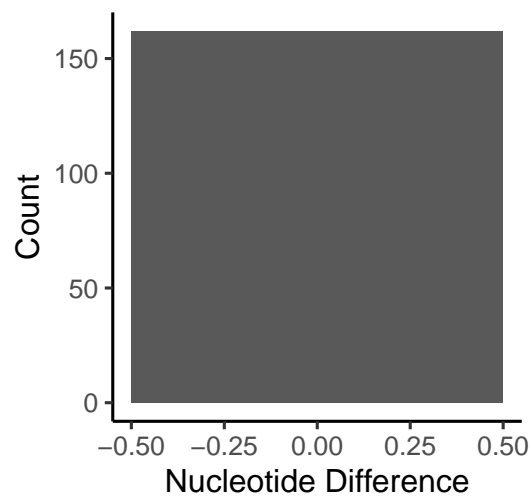
## TRBV7-8\*01

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



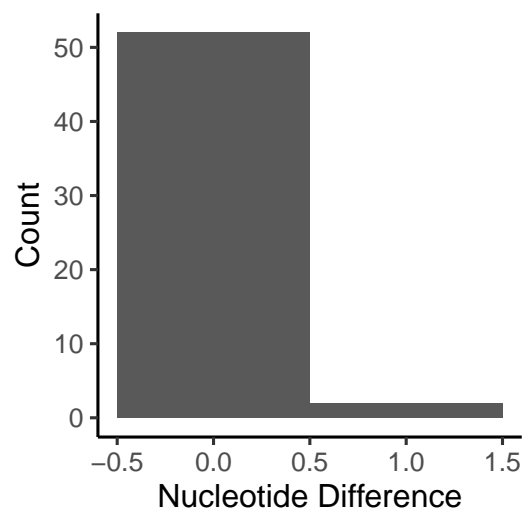
## TRBV7-9\*01

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

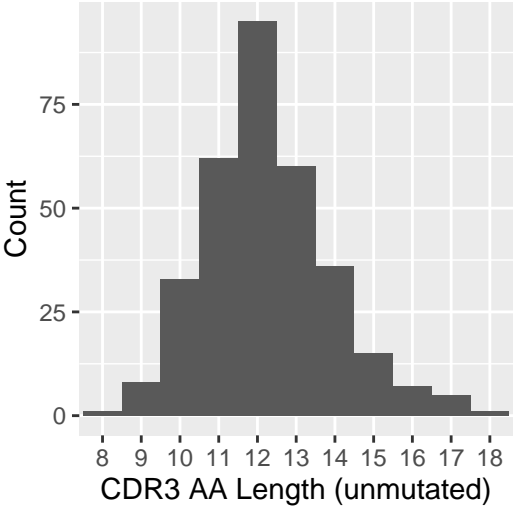


## TRBV29-1\*01

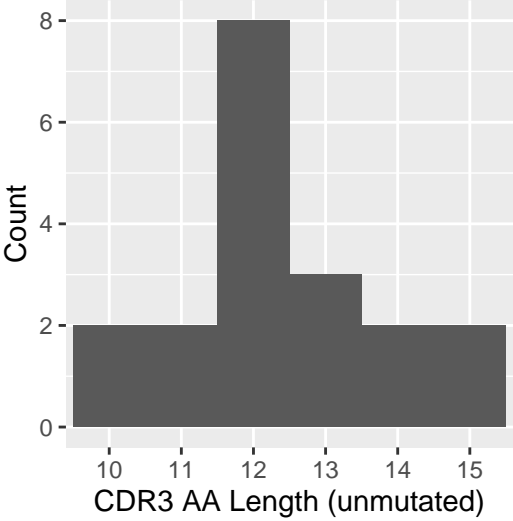
54 sequences assigned  
52 (96.3%) exact matches, in which:  
52 unique CDR3  
11 unique J

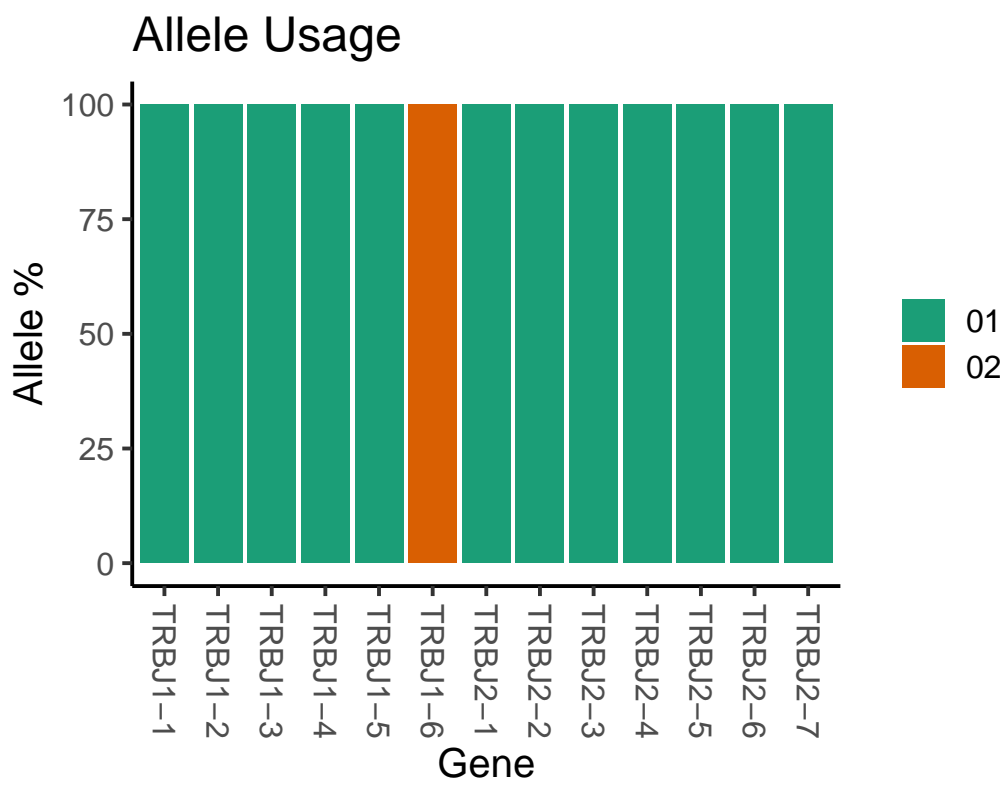


TRBV20-1\*02\_319AGAGA323



TRBV5-5\*02\_322TTGG325





Warning – no inferred sequences found.

Warning – no inferred sequences found.