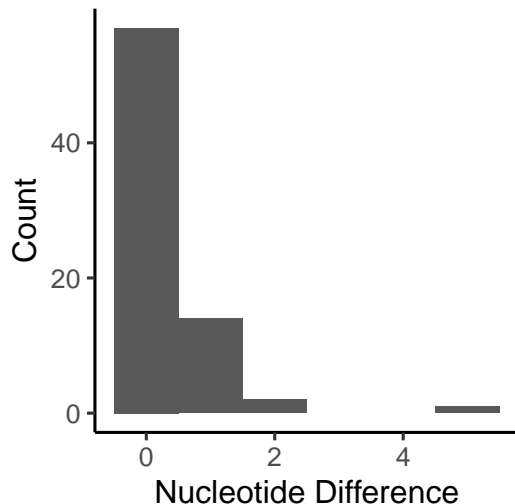


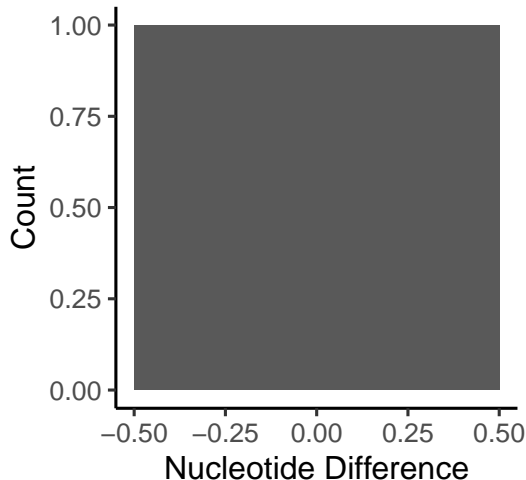
IGHV1-2*04

74 sequences assigned
57 (77%) exact matches, in which:
57 unique CDR3
6 unique J



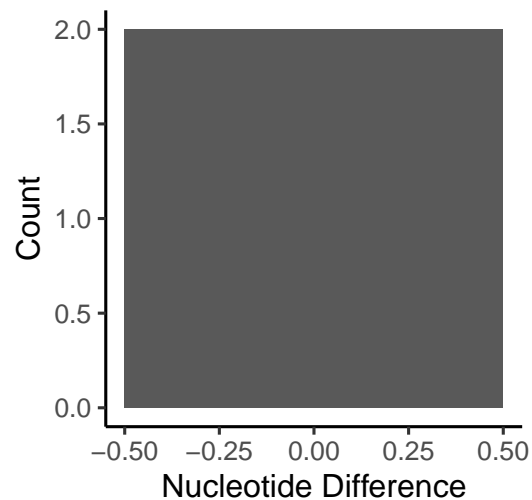
IGHV1-8*01

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



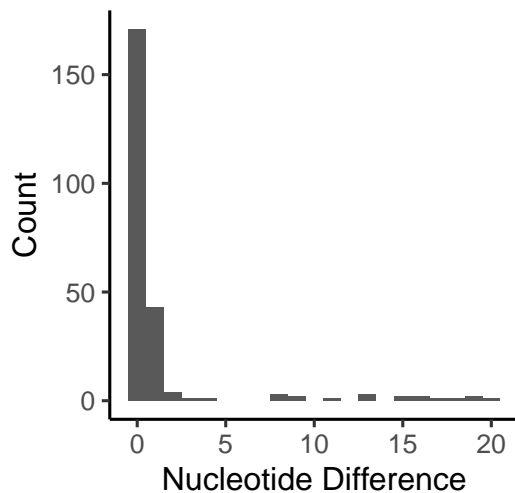
IGHV1-45*02

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



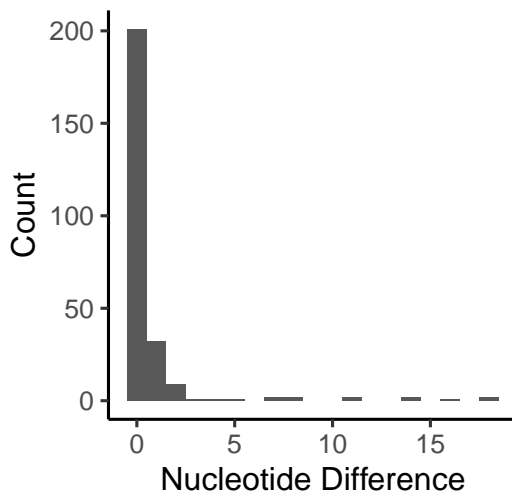
IGHV1-2*06

241 sequences assigned
171 (71%) exact matches, in which:
171 unique CDR3
6 unique J



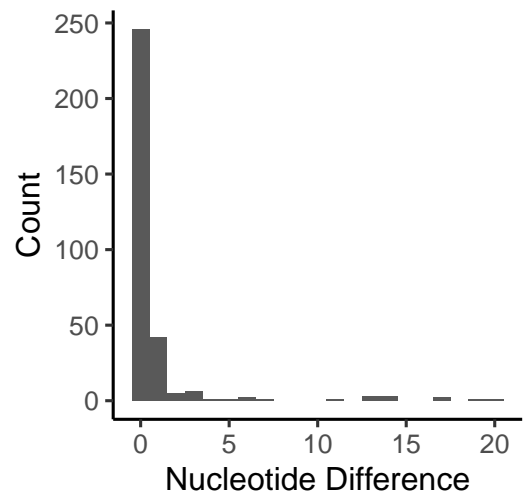
IGHV1-18*01

261 sequences assigned
201 (77%) exact matches, in which:
201 unique CDR3
7 unique J



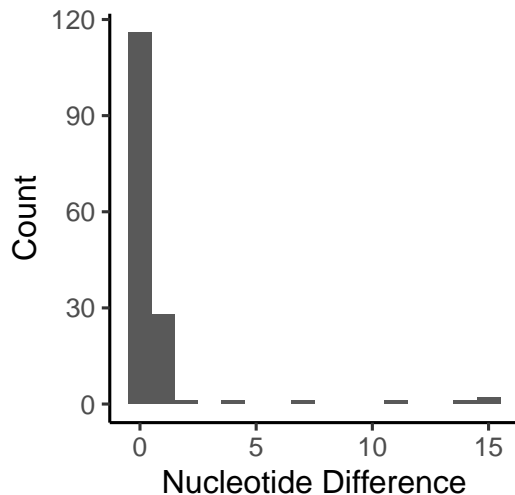
IGHV1-46*01

319 sequences assigned
246 (77.1%) exact matches, in which:
245 unique CDR3
7 unique J



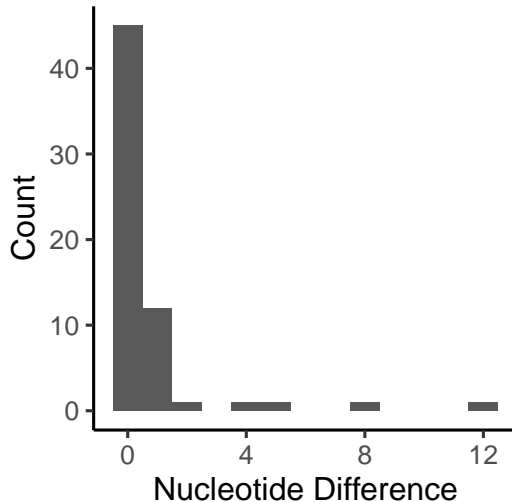
IGHV1-3*01

152 sequences assigned
116 (76.3%) exact matches, in which:
116 unique CDR3
6 unique J



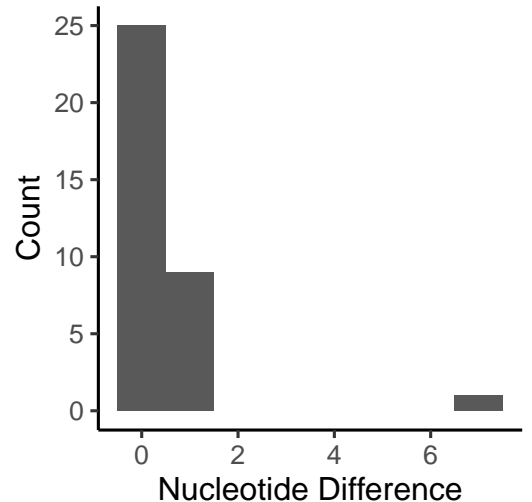
IGHV1-24*01

62 sequences assigned
45 (72.6%) exact matches, in which:
45 unique CDR3
7 unique J



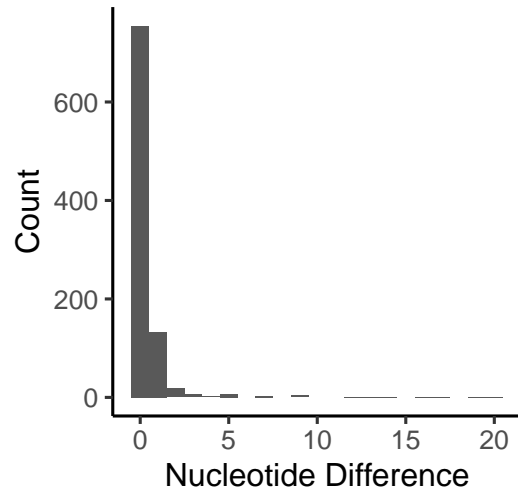
IGHV1-58*01

35 sequences assigned
25 (71.4%) exact matches, in which:
25 unique CDR3
5 unique J



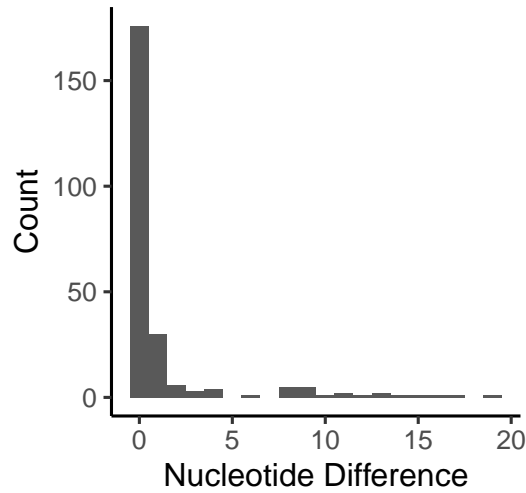
IGHV1-69*01

953 sequences assigned
755 (79.2%) exact matches, in which:
751 unique CDR3
7 unique J



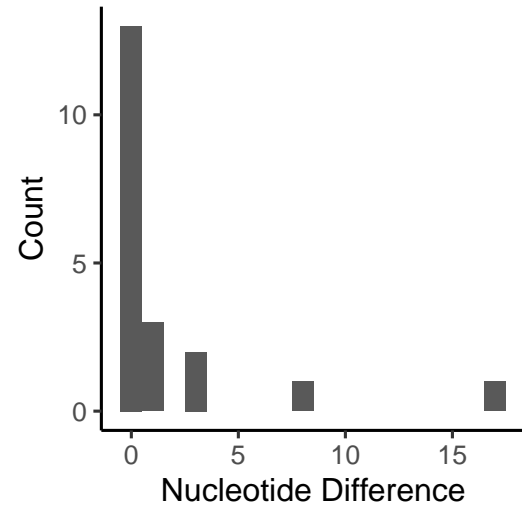
IGHV2-5*02

242 sequences assigned
176 (72.7%) exact matches, in which:
176 unique CDR3
7 unique J



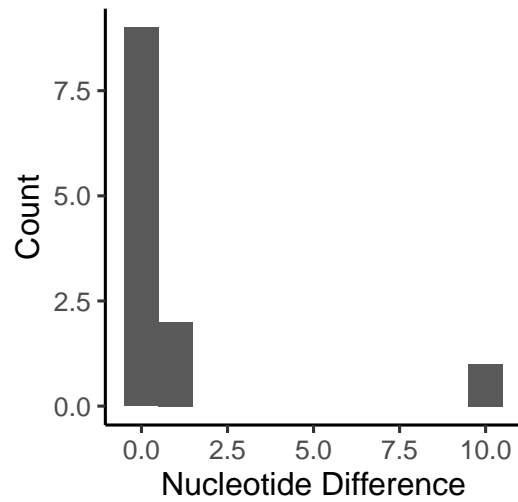
IGHV2-70*04

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



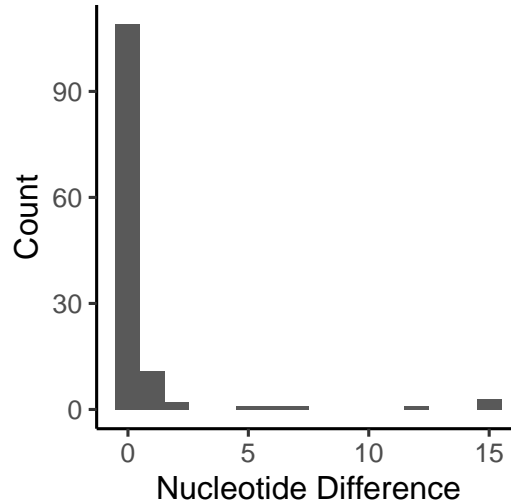
IGHV1-69-2*01

12 sequences assigned
9 (75%) exact matches, in which:
9 unique CDR3
4 unique J



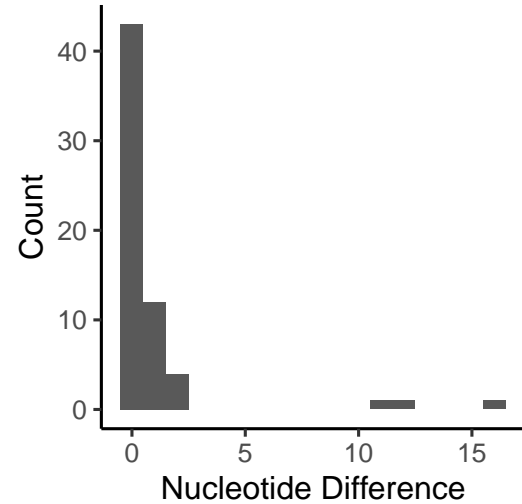
IGHV2-26*01

129 sequences assigned
109 (84.5%) exact matches, in which:
109 unique CDR3
7 unique J



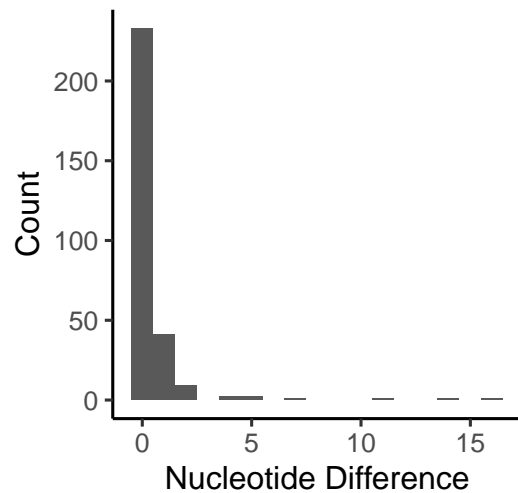
IGHV3-7*01

64 sequences assigned
43 (67.2%) exact matches, in which:
43 unique CDR3
6 unique J



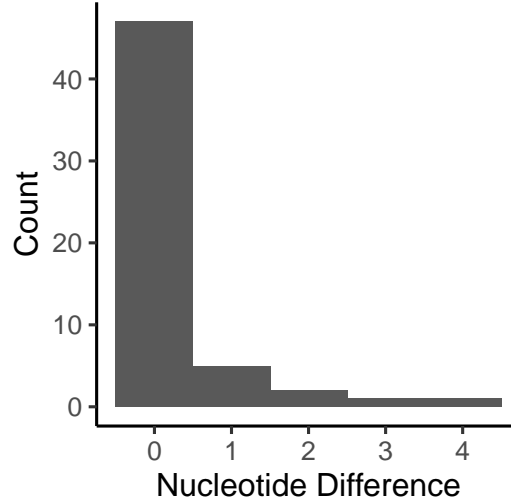
IGHV1-69*06

296 sequences assigned
233 (78.7%) exact matches, in which:
233 unique CDR3
6 unique J



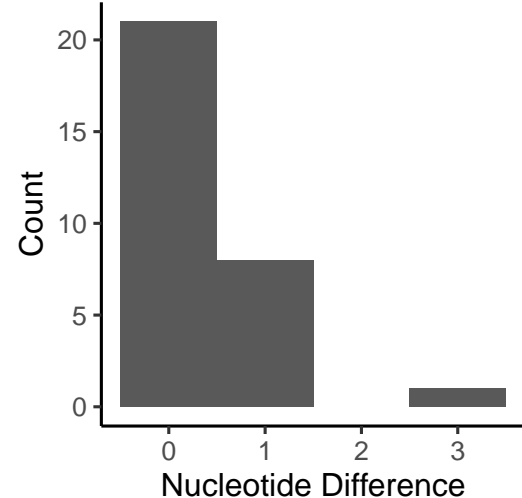
IGHV2-70*01

56 sequences assigned
47 (83.9%) exact matches, in which:
47 unique CDR3
5 unique J



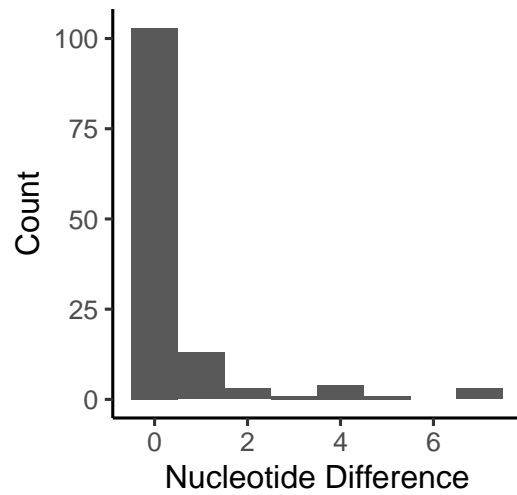
IGHV3-7*04

30 sequences assigned
21 (70%) exact matches, in which:
21 unique CDR3
4 unique J



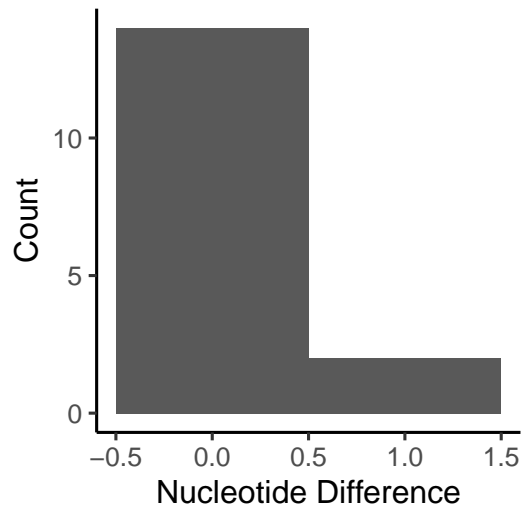
IGHV3-11*05

128 sequences assigned
103 (80.5%) exact matches, in which:
103 unique CDR3
5 unique J



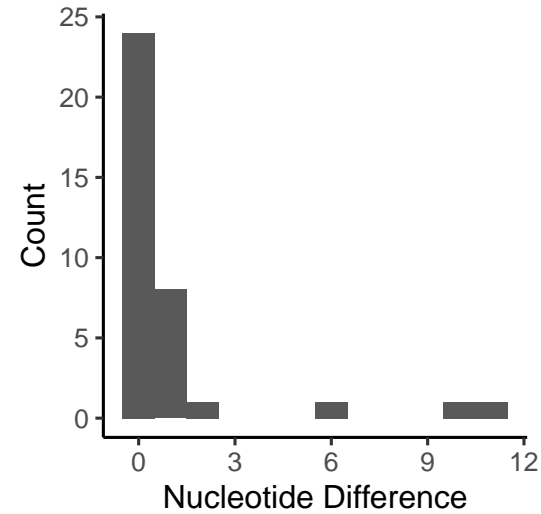
IGHV3-13*05

16 sequences assigned
14 (87.5%) exact matches, in which:
14 unique CDR3
4 unique J



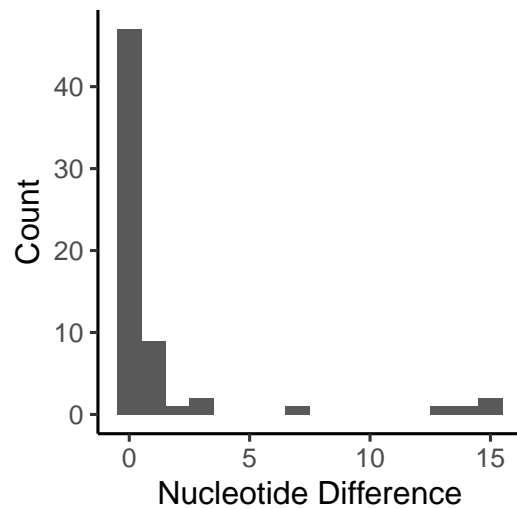
IGHV3-20*04

36 sequences assigned
24 (66.7%) exact matches, in which:
24 unique CDR3
5 unique J



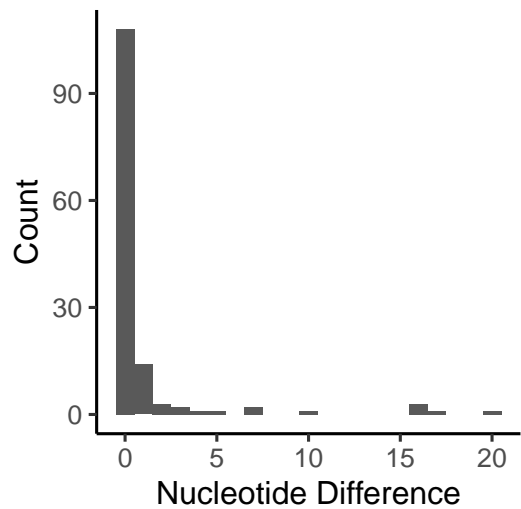
IGHV3-11*06

64 sequences assigned
47 (73.4%) exact matches, in which:
47 unique CDR3
5 unique J



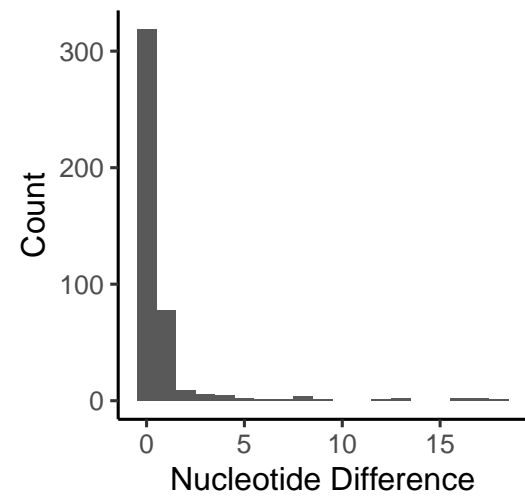
IGHV3-15*01

140 sequences assigned
108 (77.1%) exact matches, in which:
108 unique CDR3
7 unique J



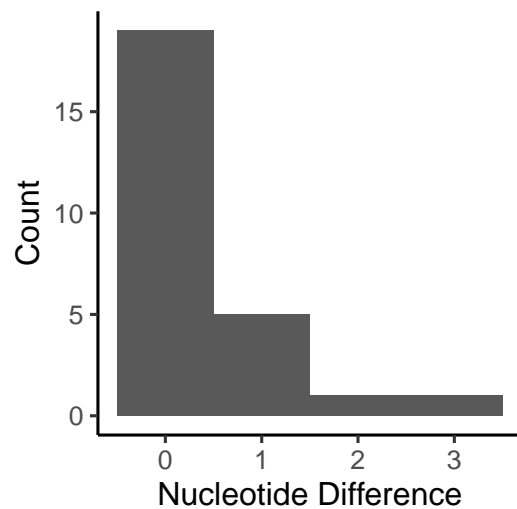
IGHV3-21*01

436 sequences assigned
319 (73.2%) exact matches, in which:
317 unique CDR3
7 unique J



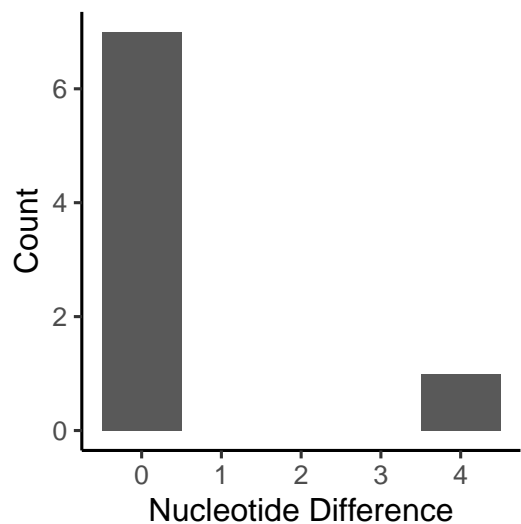
IGHV3-13*04

26 sequences assigned
19 (73.1%) exact matches, in which:
19 unique CDR3
4 unique J



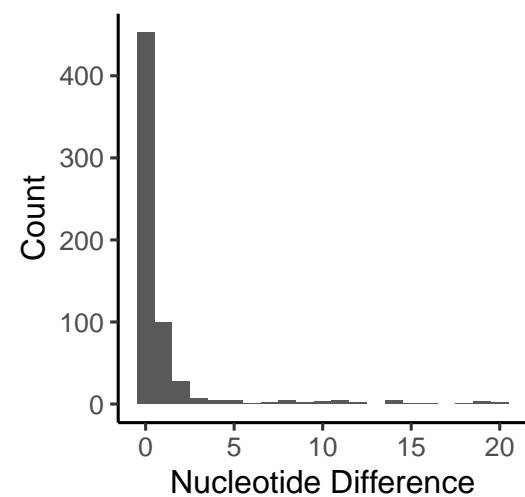
IGHV3-20*01

8 sequences assigned
7 (87.5%) exact matches, in which:
7 unique CDR3
3 unique J



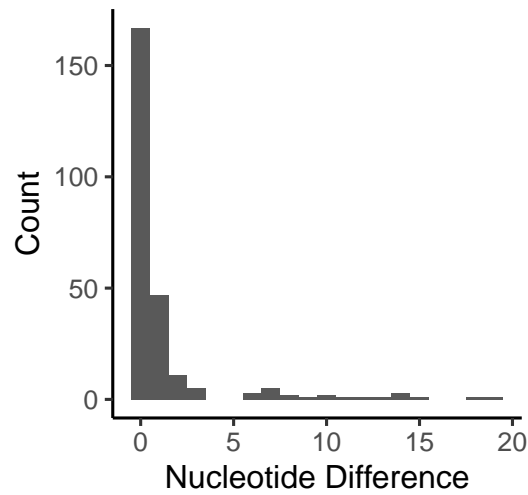
IGHV3-23*01

628 sequences assigned
453 (72.1%) exact matches, in which:
453 unique CDR3
7 unique J



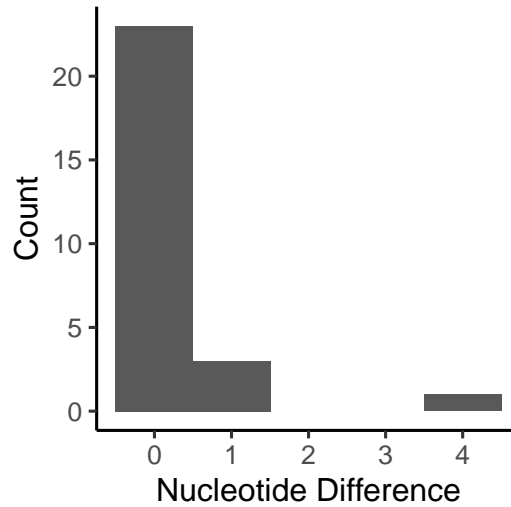
IGHV3-30-3*01

254 sequences assigned
167 (65.7%) exact matches, in which:
166 unique CDR3
6 unique J



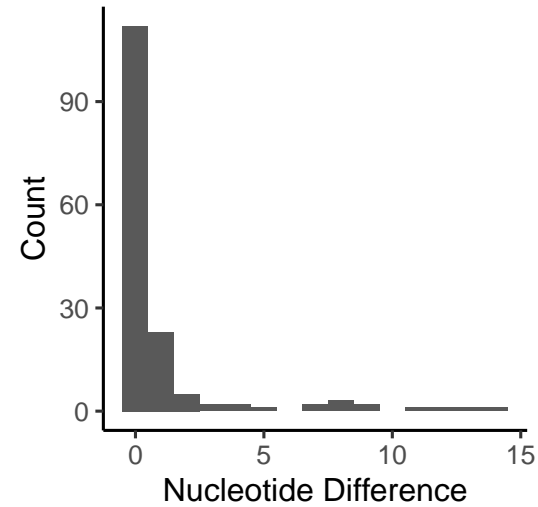
IGHV3-43*01

27 sequences assigned
23 (85.2%) exact matches, in which:
23 unique CDR3
5 unique J



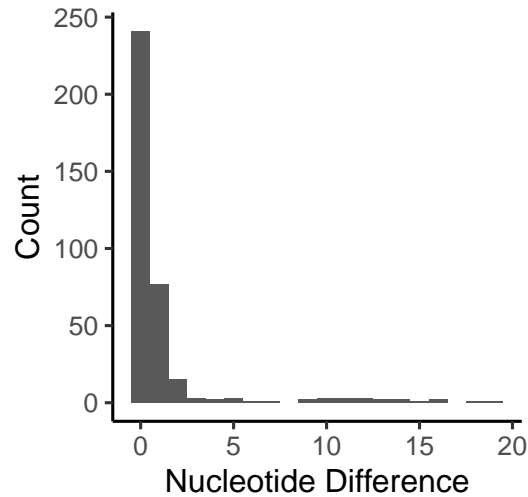
IGHV3-53*01

157 sequences assigned
112 (71.3%) exact matches, in which:
112 unique CDR3
6 unique J



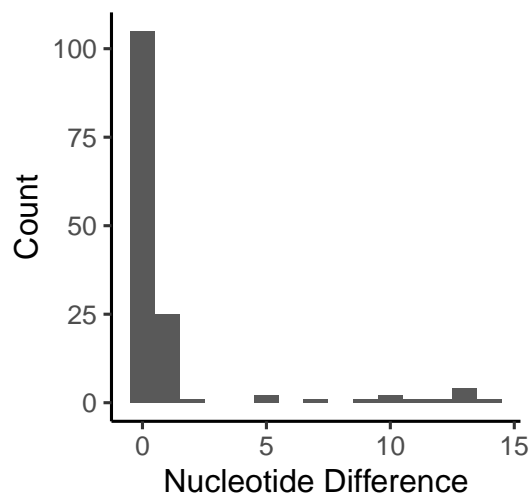
IGHV3-30*18

365 sequences assigned
241 (66%) exact matches, in which:
241 unique CDR3
7 unique J



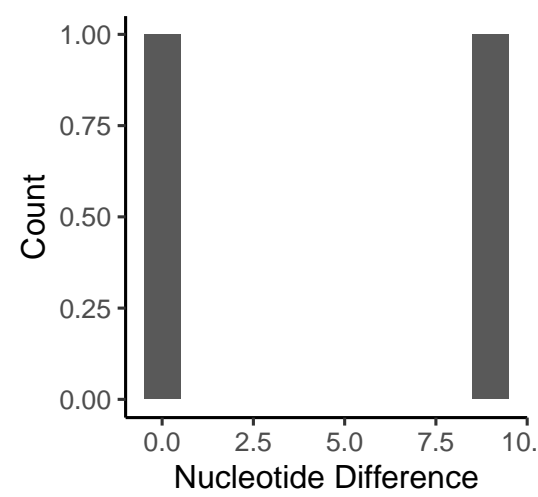
IGHV3-48*02

146 sequences assigned
105 (71.9%) exact matches, in which:
105 unique CDR3
7 unique J



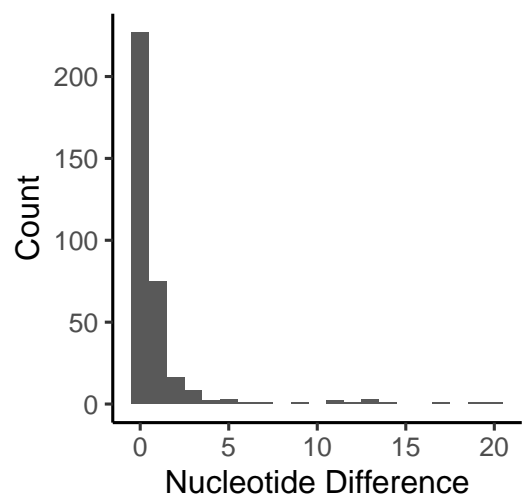
IGHV3-64*02

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



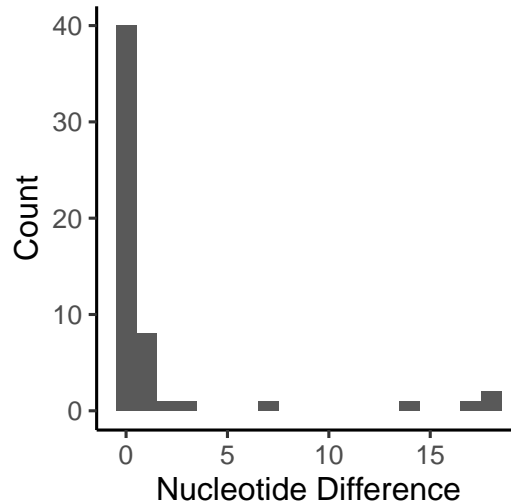
IGHV3-33*01

349 sequences assigned
227 (65%) exact matches, in which:
226 unique CDR3
7 unique J



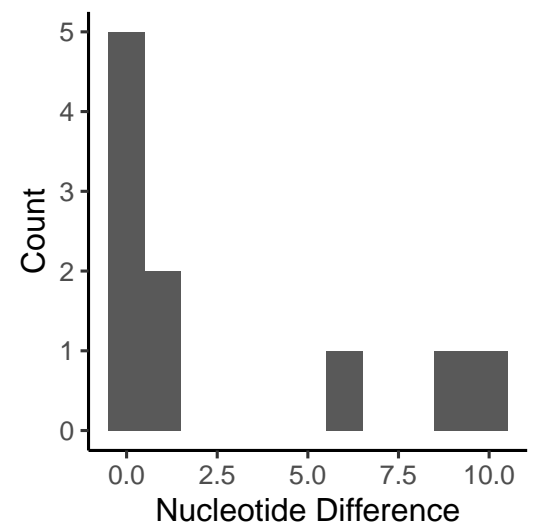
IGHV3-49*05

55 sequences assigned
40 (72.7%) exact matches, in which:
40 unique CDR3
7 unique J



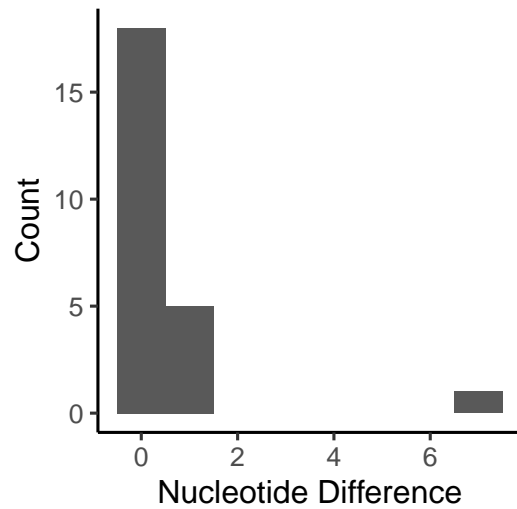
IGHV3-72*01

11 sequences assigned
5 (45.5%) exact matches, in which:
5 unique CDR3
3 unique J



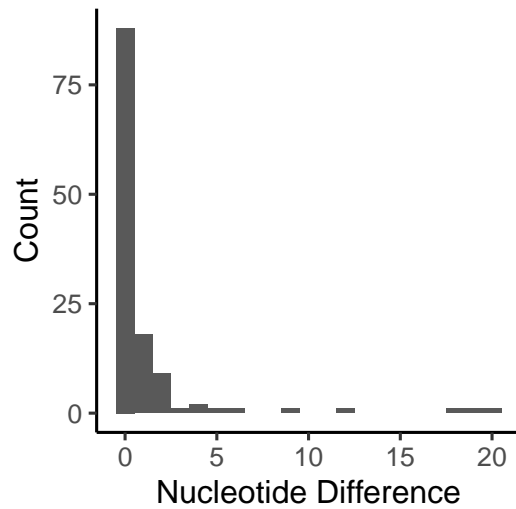
IGHV3-73*02

24 sequences assigned
18 (75%) exact matches, in which:
18 unique CDR3
5 unique J



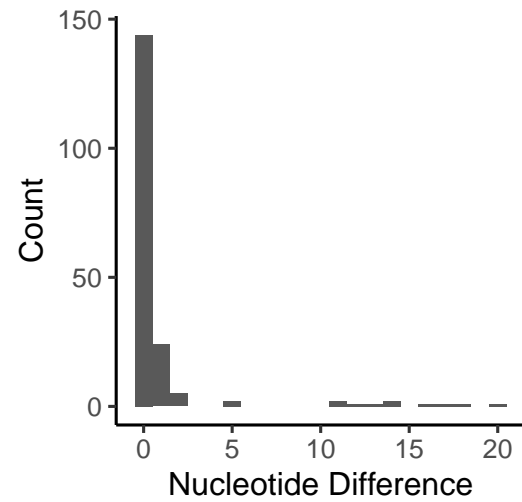
IGHV4-4*02

128 sequences assigned
88 (68.8%) exact matches, in which:
87 unique CDR3
7 unique J



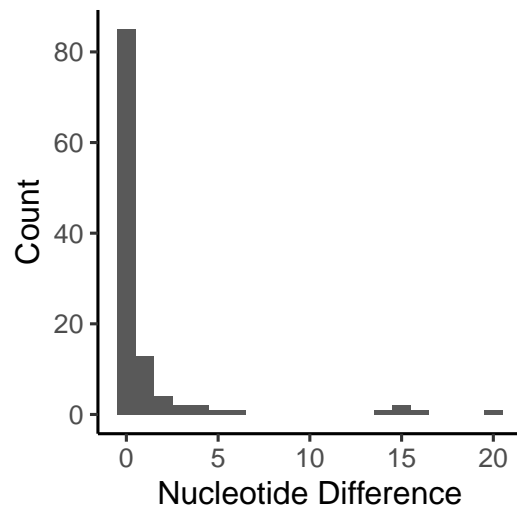
IGHV4-31*03

186 sequences assigned
144 (77.4%) exact matches, in which:
144 unique CDR3
6 unique J



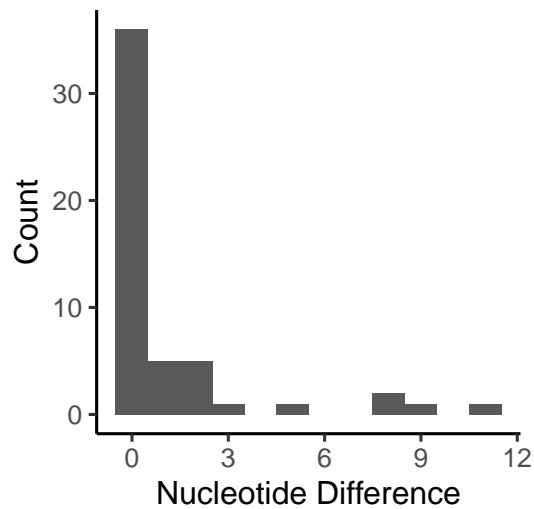
IGHV3-74*01

114 sequences assigned
85 (74.6%) exact matches, in which:
85 unique CDR3
6 unique J



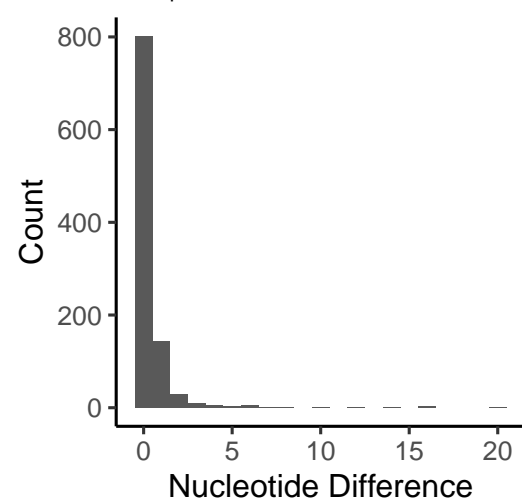
IGHV4-30-4*01

52 sequences assigned
36 (69.2%) exact matches, in which:
36 unique CDR3
6 unique J



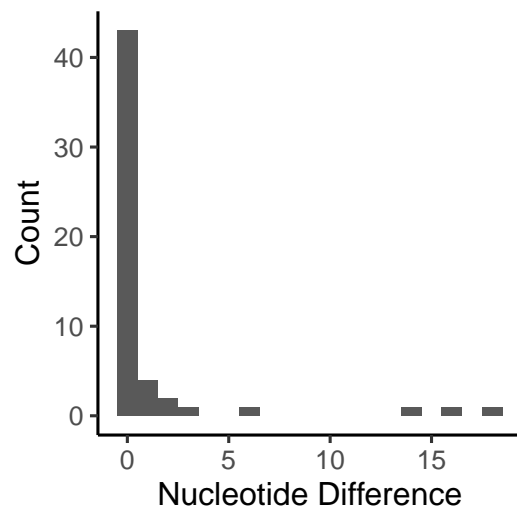
IGHV4-34*01

1014 sequences assigned
802 (79.1%) exact matches, in which:
794 unique CDR3
7 unique J



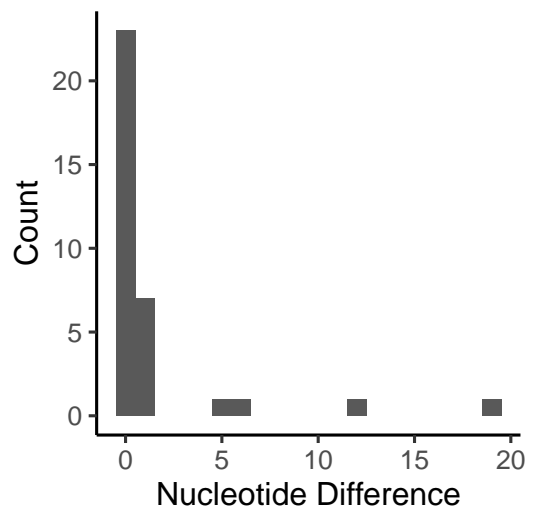
IGHV3-64D*06

56 sequences assigned
43 (76.8%) exact matches, in which:
42 unique CDR3
7 unique J



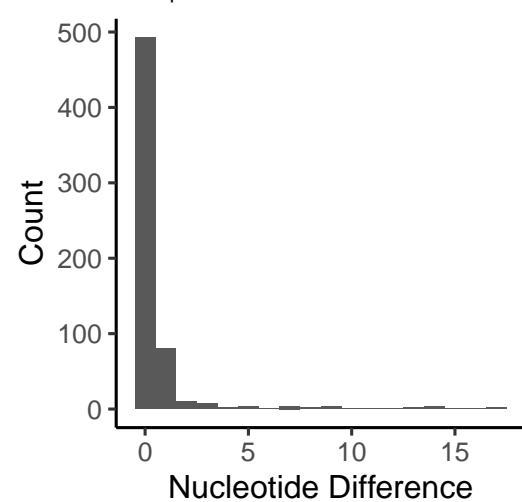
IGHV4-30-2*01

36 sequences assigned
23 (63.9%) exact matches, in which:
23 unique CDR3
7 unique J



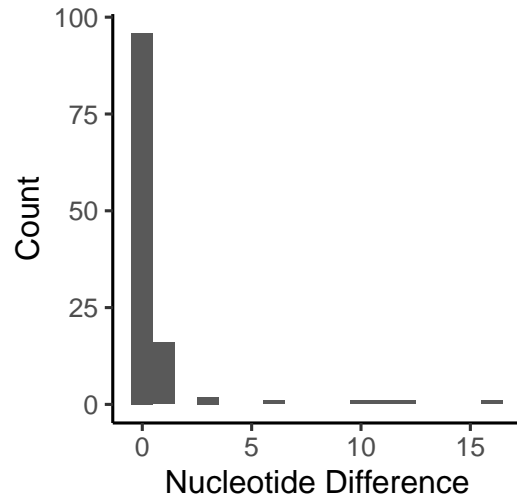
IGHV4-39*01

624 sequences assigned
493 (79%) exact matches, in which:
489 unique CDR3
7 unique J



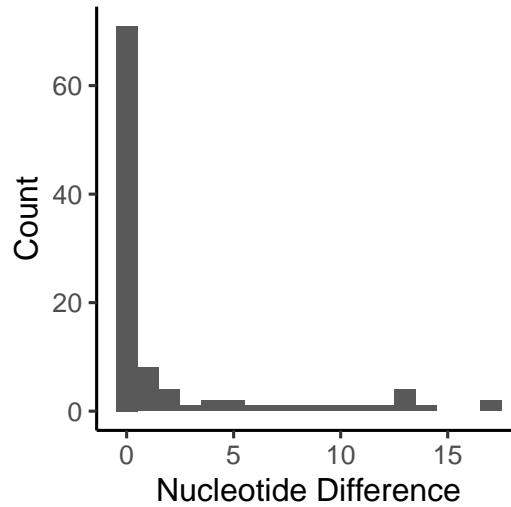
IGHV4-59*01

125 sequences assigned
96 (76.8%) exact matches, in which:
96 unique CDR3
7 unique J



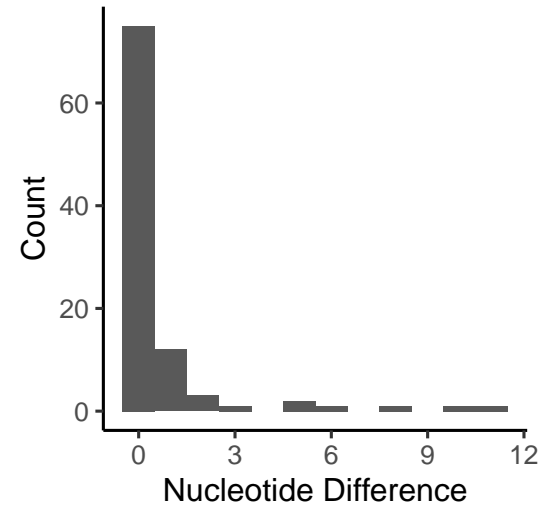
IGHV5-10-1*03

103 sequences assigned
71 (68.9%) exact matches, in which:
71 unique CDR3
6 unique J



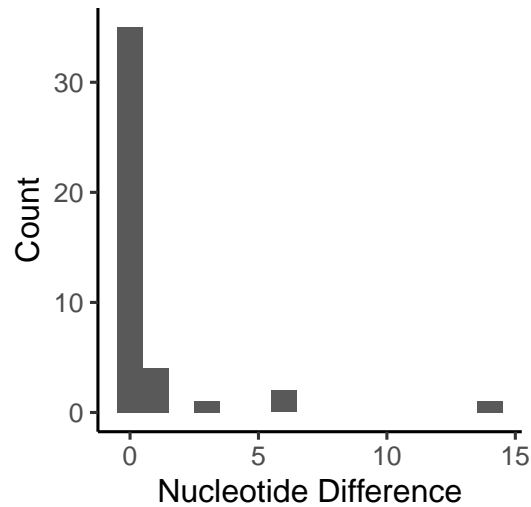
IGHV7-4-1*02

98 sequences assigned
75 (76.5%) exact matches, in which:
75 unique CDR3
6 unique J



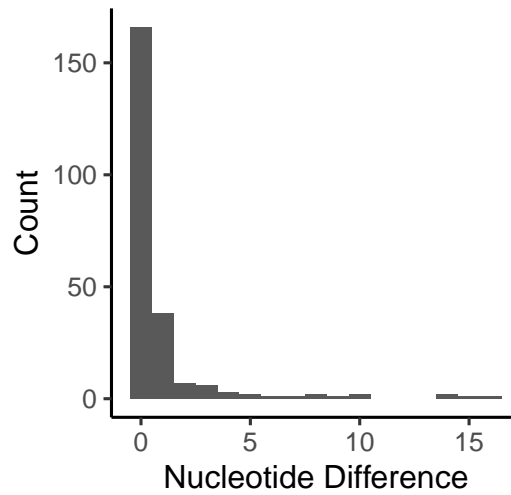
IGHV4-61*01

44 sequences assigned
35 (79.5%) exact matches, in which:
35 unique CDR3
6 unique J



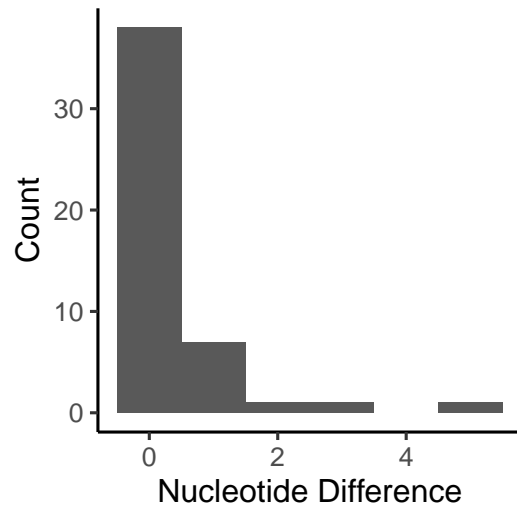
IGHV5-51*01

233 sequences assigned
166 (71.2%) exact matches, in which:
166 unique CDR3
7 unique J



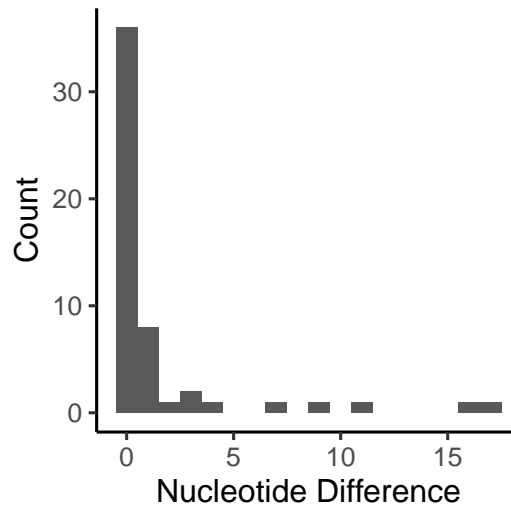
IGHV5-10-1*01

48 sequences assigned
38 (79.2%) exact matches, in which:
38 unique CDR3
6 unique J

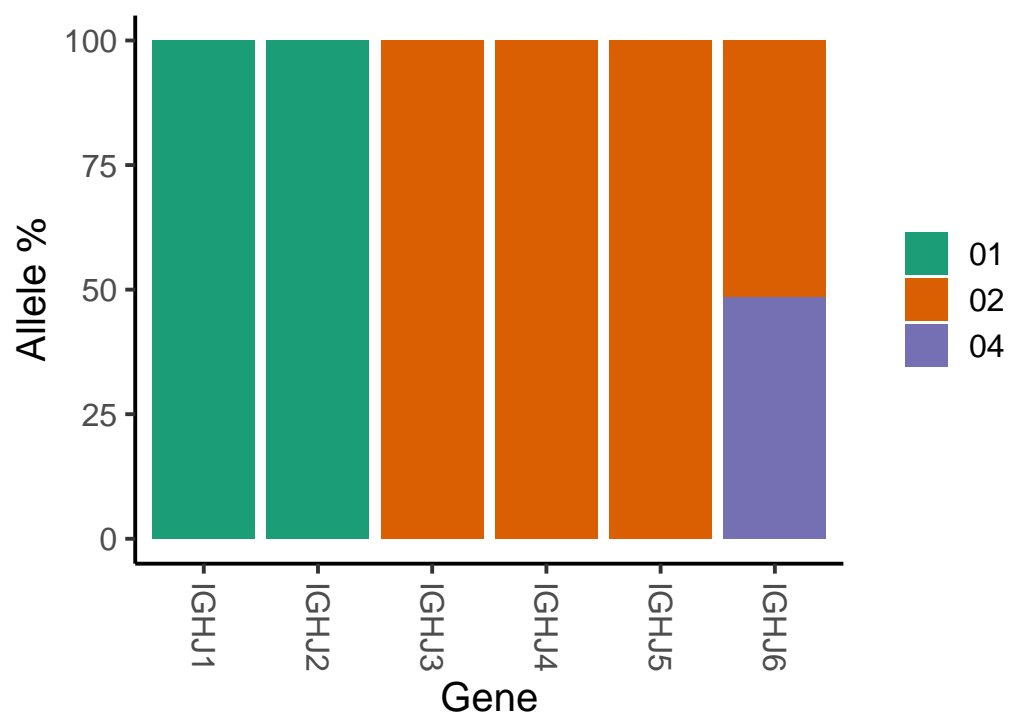


IGHV6-1*01

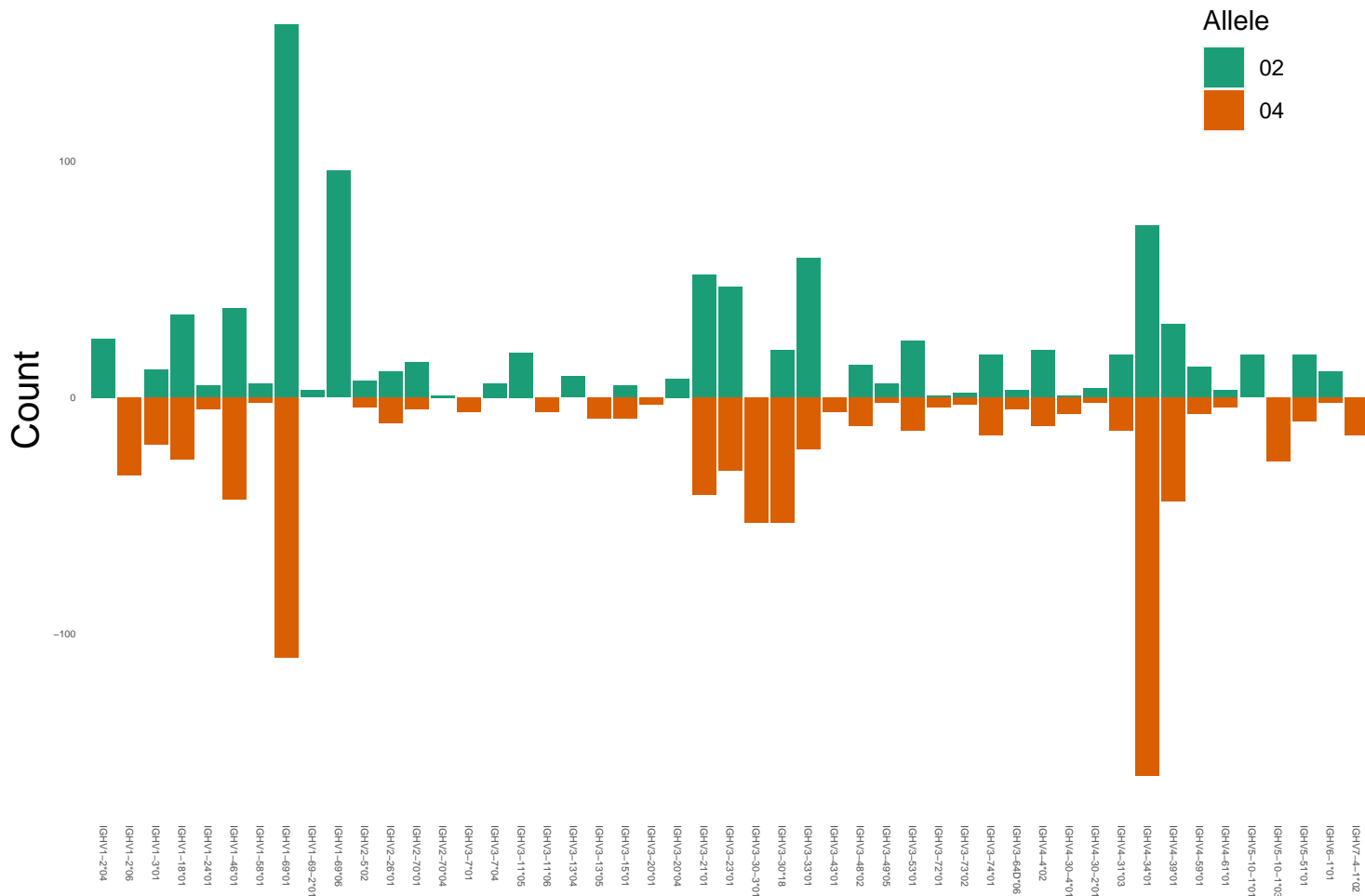
54 sequences assigned
36 (66.7%) exact matches, in which:
36 unique CDR3
6 unique J



Allele Usage



Sequence Count by IGHJ6 allele usage



Warning – no inferred sequences found.