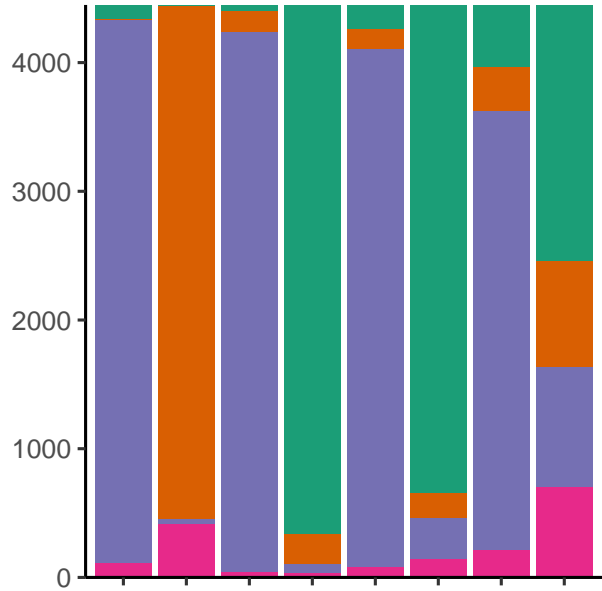


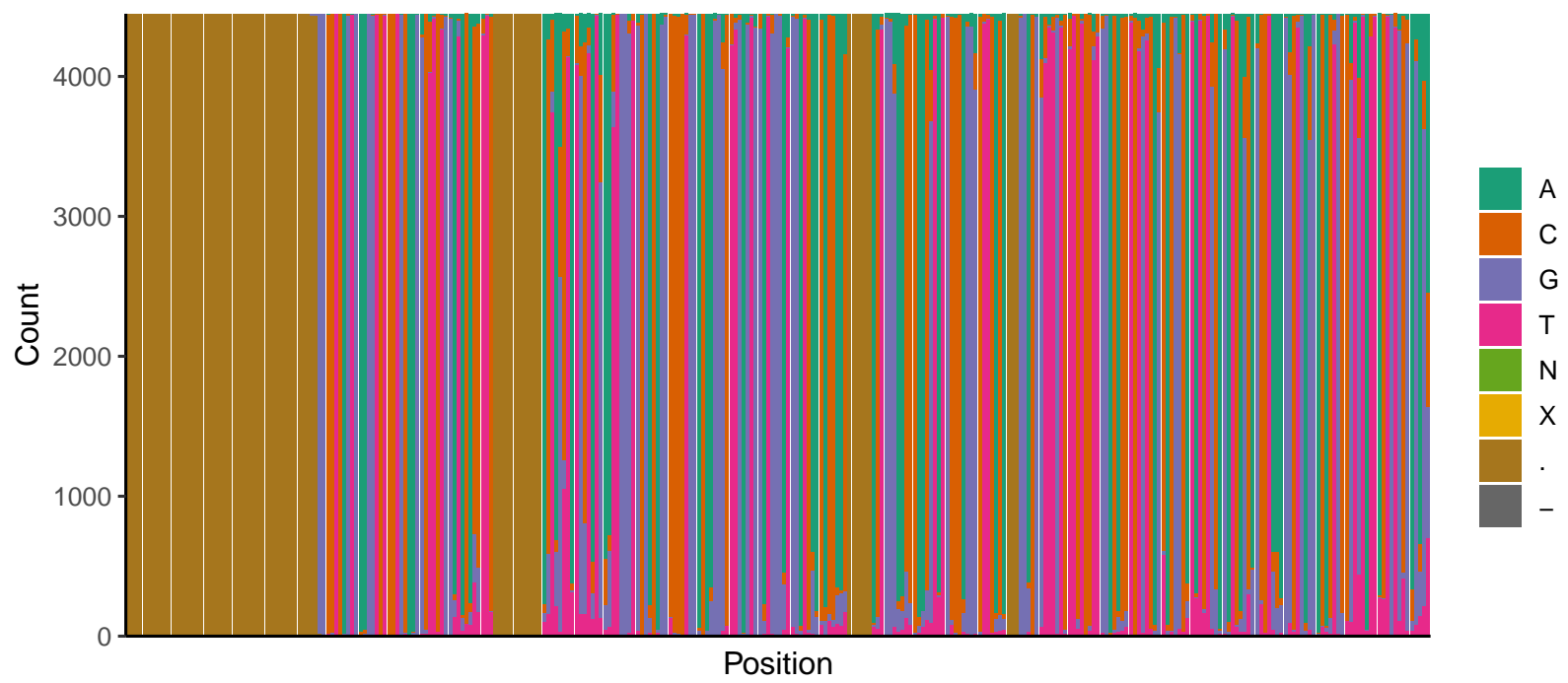
Gene IGHV7-4-1*02_T63C

Count

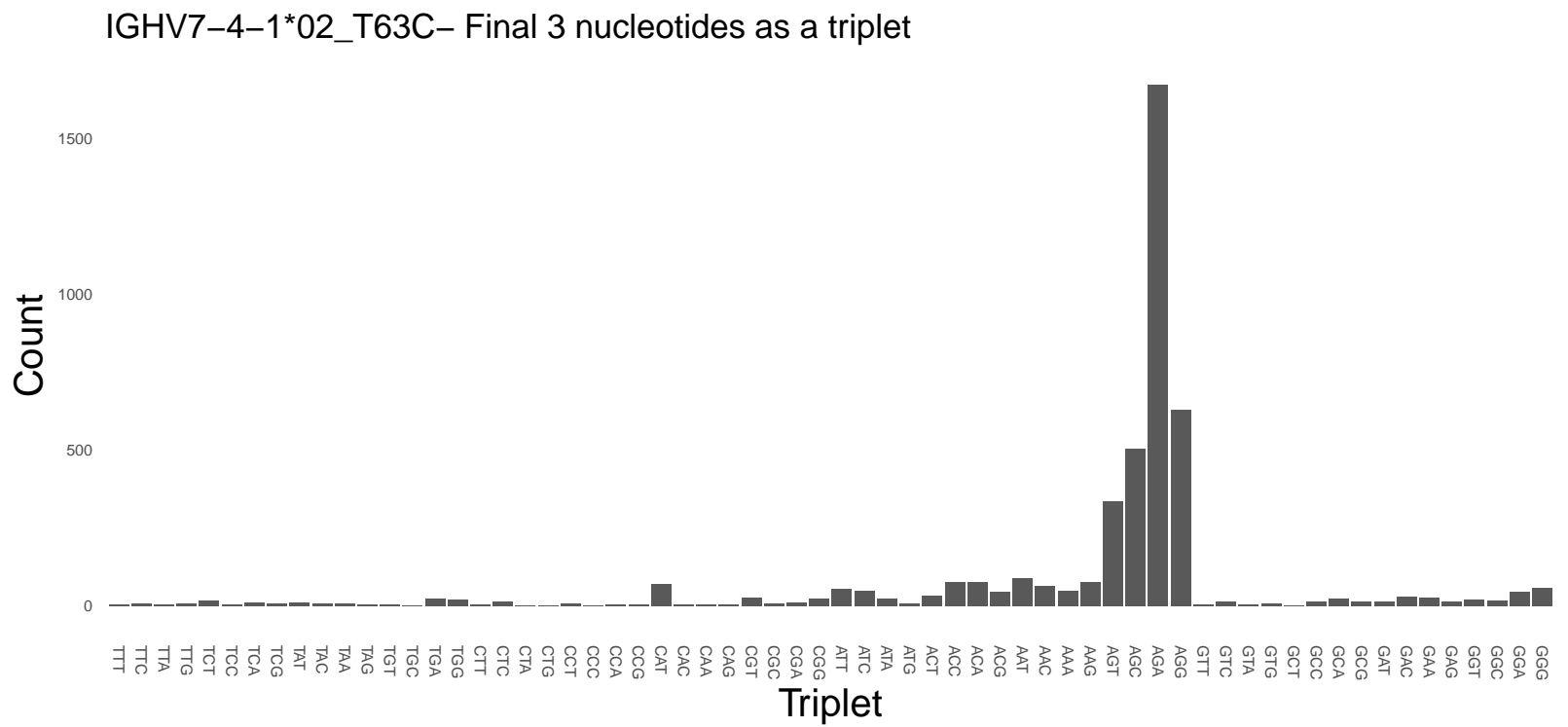


Position

Gene IGHV7-4-1*02_T63C

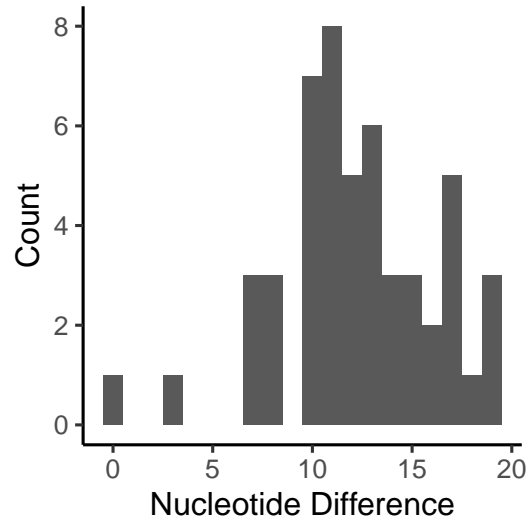


IGHV7-4-1*02_T63C- Final 3 nucleotides as a triplet



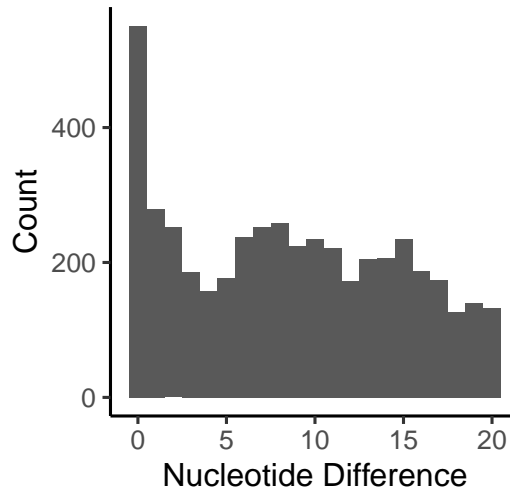
IGHV1-2*02

53 sequences assigned
1 (1.9%) exact matches, in which:
1 unique CDR3
1 unique J



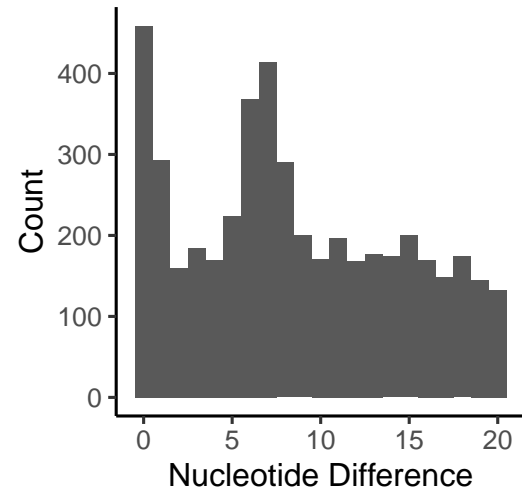
IGHV1-3*01_05

5776 sequences assigned
551 (9.5%) exact matches, in which:
543 unique CDR3
7 unique J



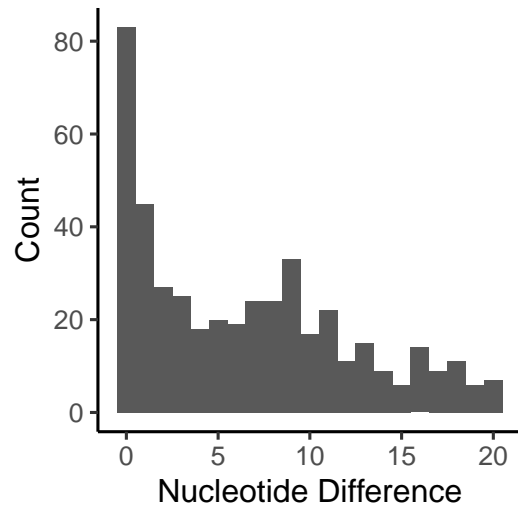
IGHV1-18*01

5518 sequences assigned
459 (8.3%) exact matches, in which:
449 unique CDR3
7 unique J



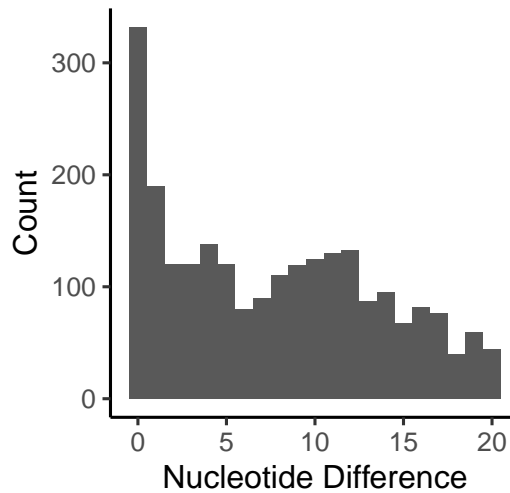
IGHV1-2*04

477 sequences assigned
83 (17.4%) exact matches, in which:
82 unique CDR3
6 unique J



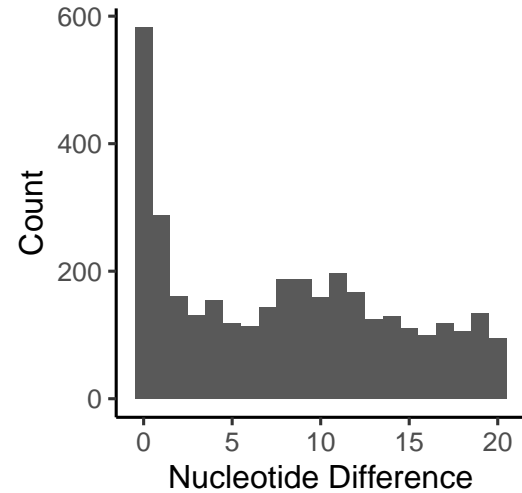
IGHV1-8*01

2756 sequences assigned
332 (12%) exact matches, in which:
321 unique CDR3
7 unique J



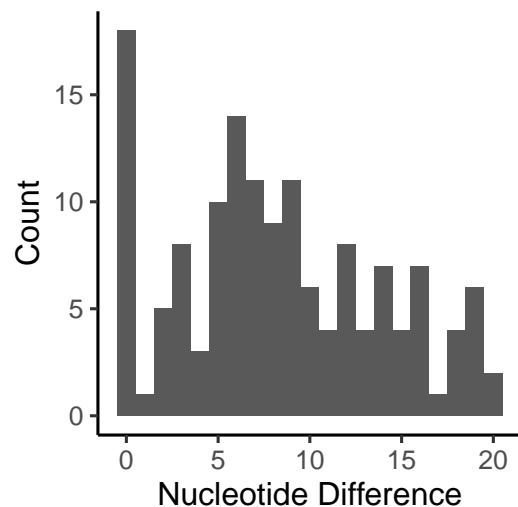
IGHV1-18*04

4296 sequences assigned
583 (13.6%) exact matches, in which:
567 unique CDR3
7 unique J



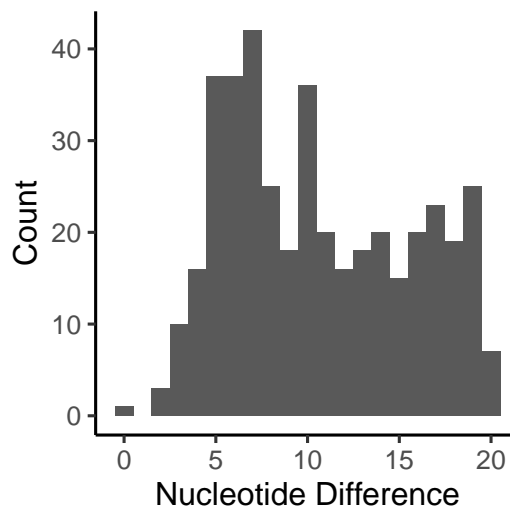
IGHV1-2*05

168 sequences assigned
18 (10.7%) exact matches, in which:
18 unique CDR3
4 unique J



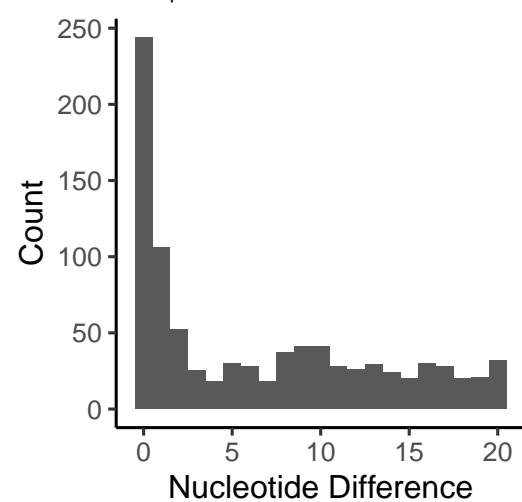
IGHV1-8*02

483 sequences assigned
1 (0.2%) exact matches, in which:
1 unique CDR3
1 unique J



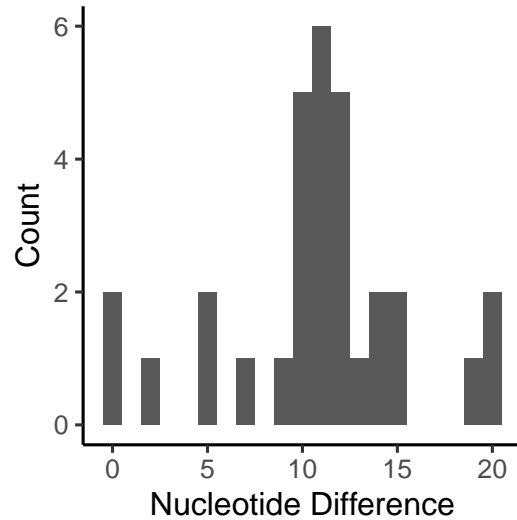
IGHV1-24*01

1121 sequences assigned
244 (21.8%) exact matches, in which:
243 unique CDR3
7 unique J



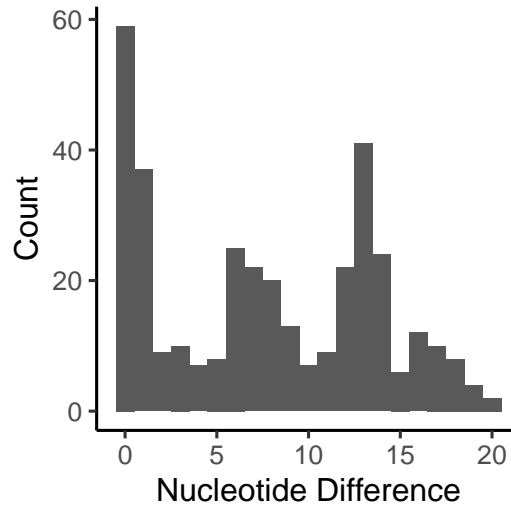
IGHV1-45*02

37 sequences assigned
2 (5.4%) exact matches, in which:
2 unique CDR3
1 unique J



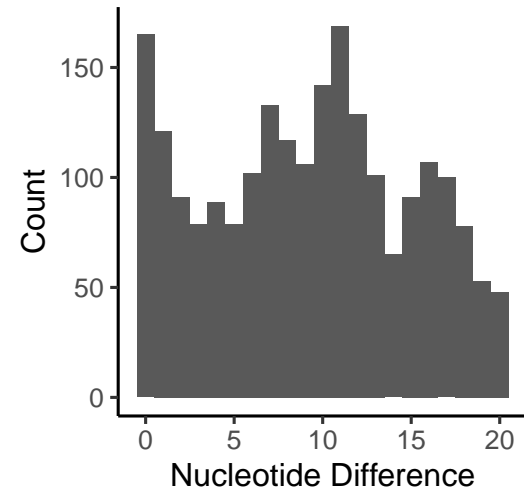
IGHV1-58*01_03

395 sequences assigned
59 (14.9%) exact matches, in which:
59 unique CDR3
7 unique J



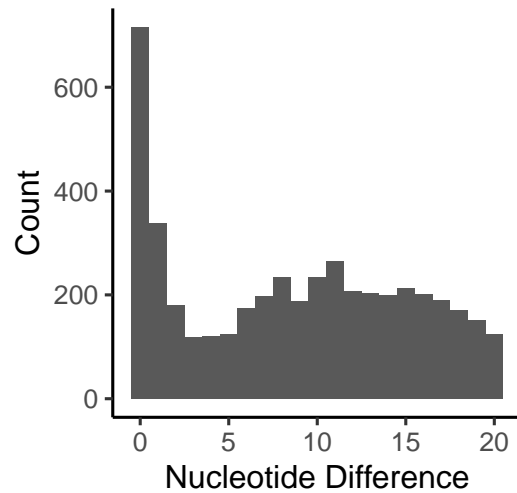
IGHV2-5*02

2444 sequences assigned
165 (6.8%) exact matches, in which:
160 unique CDR3
6 unique J



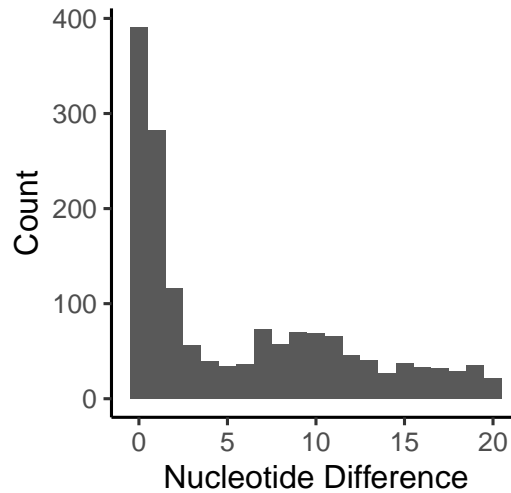
IGHV1-46*01

5908 sequences assigned
716 (12.1%) exact matches, in which:
708 unique CDR3
7 unique J



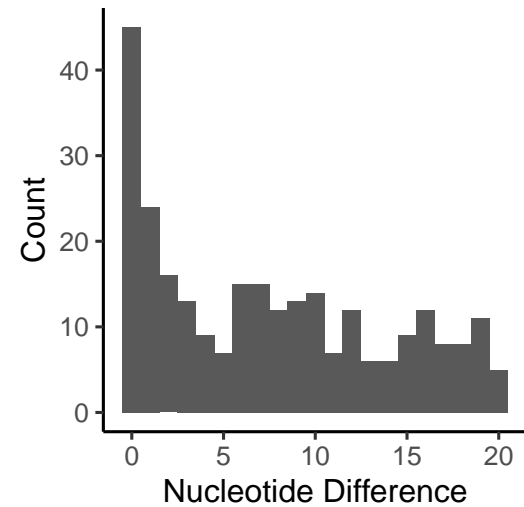
IGHV1-69*10

1972 sequences assigned
391 (19.8%) exact matches, in which:
385 unique CDR3
7 unique J



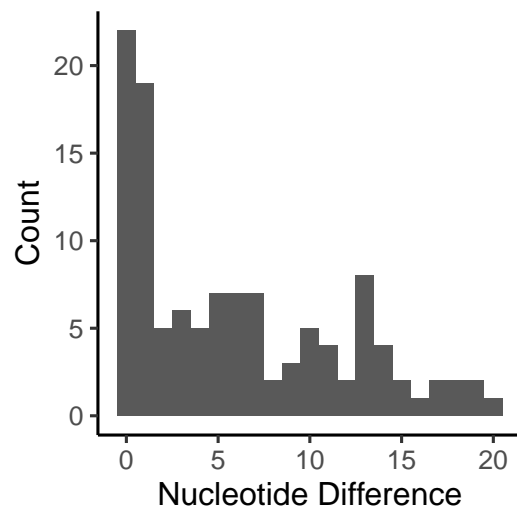
IGHV2-26*01

309 sequences assigned
45 (14.6%) exact matches, in which:
44 unique CDR3
6 unique J



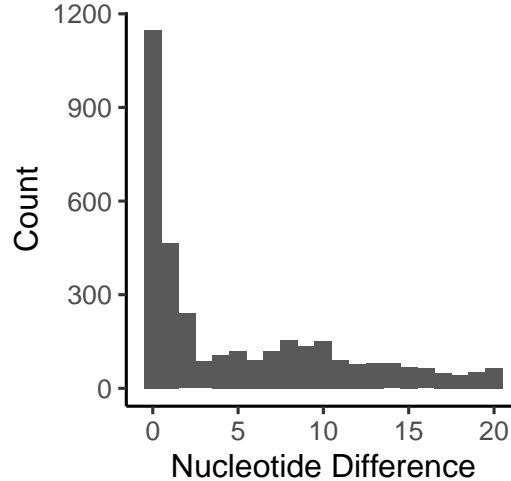
IGHV1-58*02

127 sequences assigned
22 (17.3%) exact matches, in which:
22 unique CDR3
5 unique J



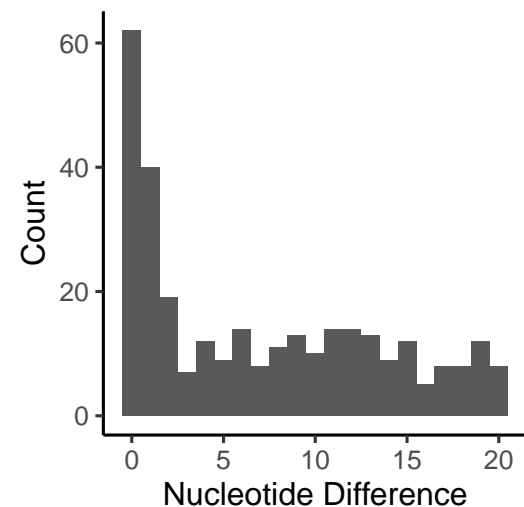
IGHV1-69*01_12_13

4021 sequences assigned
1148 (28.6%) exact matches, in which:
1138 unique CDR3
7 unique J



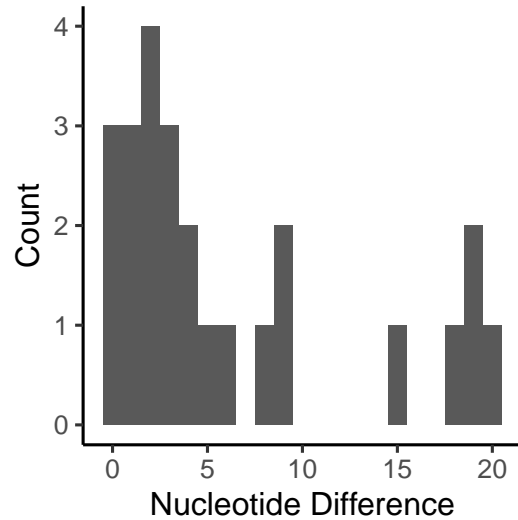
IGHV2-70*01

344 sequences assigned
62 (18%) exact matches, in which:
62 unique CDR3
5 unique J



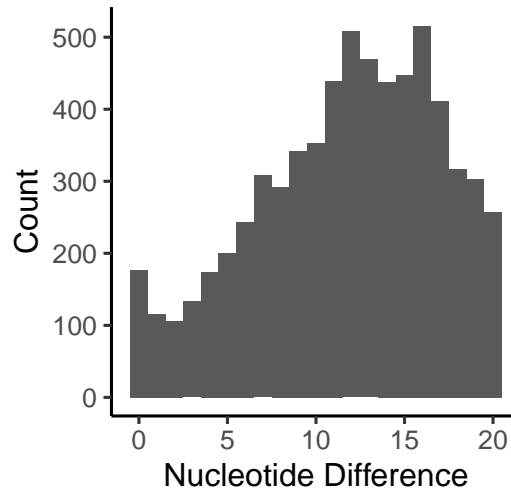
IGHV2-70*12

29 sequences assigned
3 (10.3%) exact matches, in which:
3 unique CDR3
2 unique J



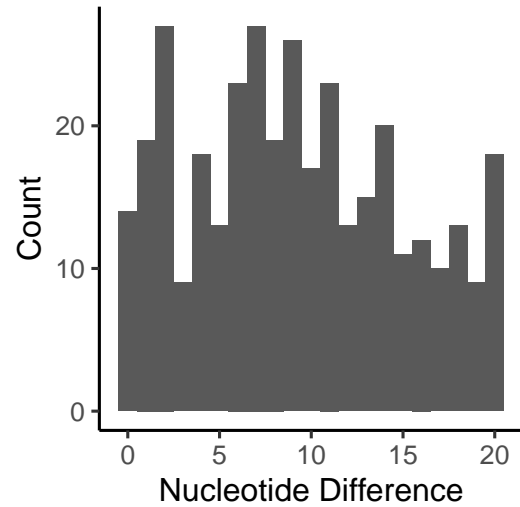
IGHV3-7*03

8422 sequences assigned
177 (2.1%) exact matches, in which:
176 unique CDR3
7 unique J



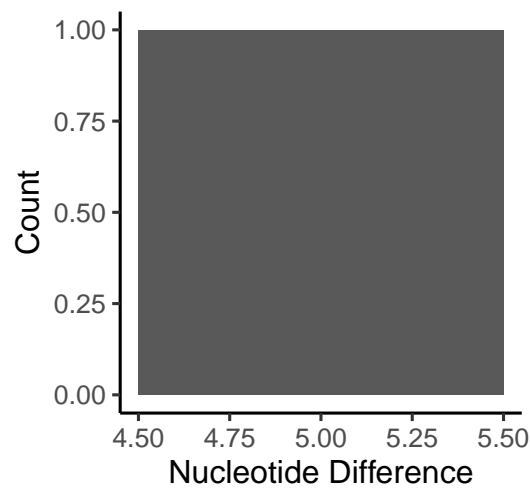
IGHV3-11*04

417 sequences assigned
14 (3.4%) exact matches, in which:
14 unique CDR3
4 unique J



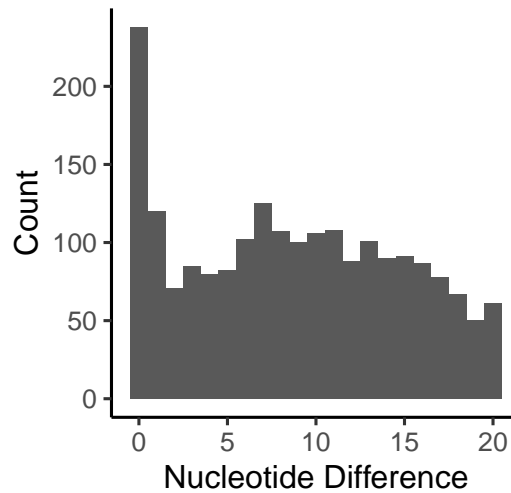
IGHV2-70D*14

1 sequences assigned
No exact matches.



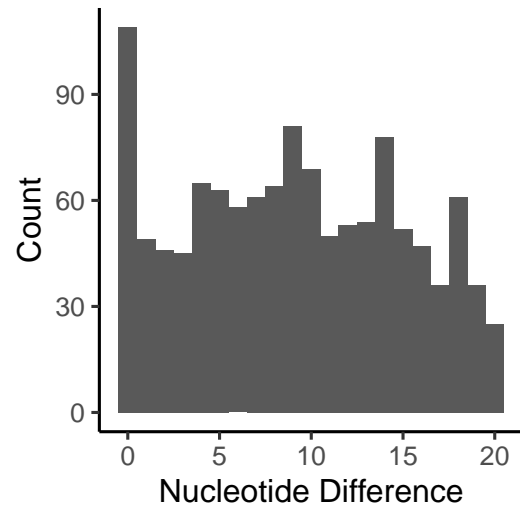
IGHV3-9*01

2347 sequences assigned
238 (10.1%) exact matches, in which:
236 unique CDR3
7 unique J



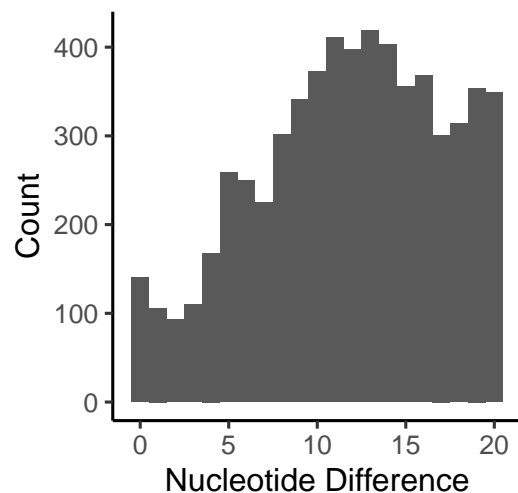
IGHV3-11*06

1490 sequences assigned
109 (7.3%) exact matches, in which:
108 unique CDR3
6 unique J



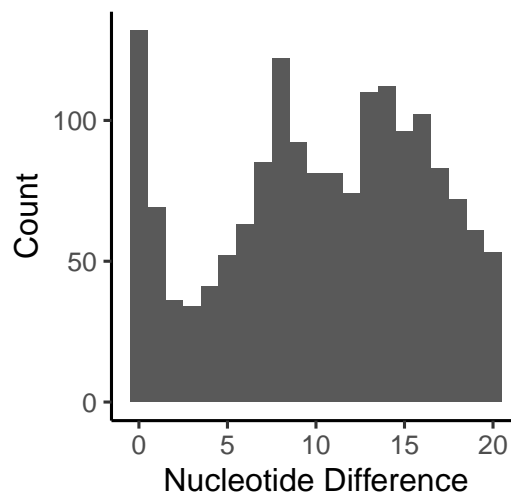
IGHV3-7*01

8162 sequences assigned
140 (1.7%) exact matches, in which:
138 unique CDR3
7 unique J



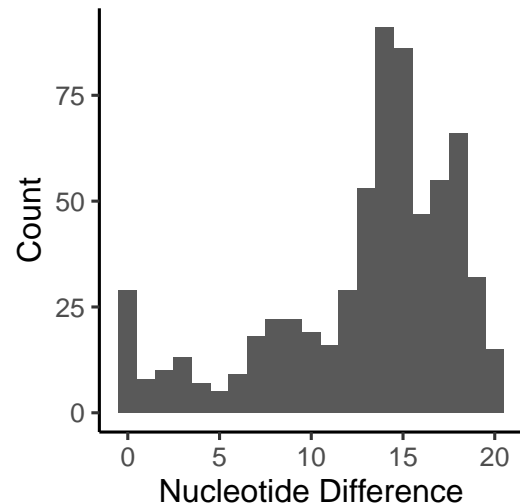
IGHV3-11*01

2373 sequences assigned
132 (5.6%) exact matches, in which:
130 unique CDR3
7 unique J



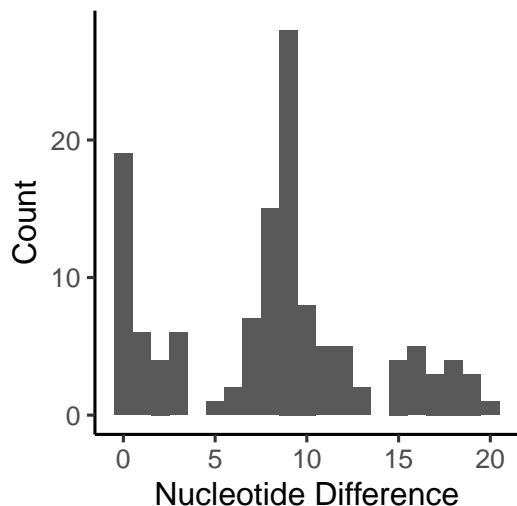
IGHV3-13*01

790 sequences assigned
29 (3.7%) exact matches, in which:
28 unique CDR3
7 unique J



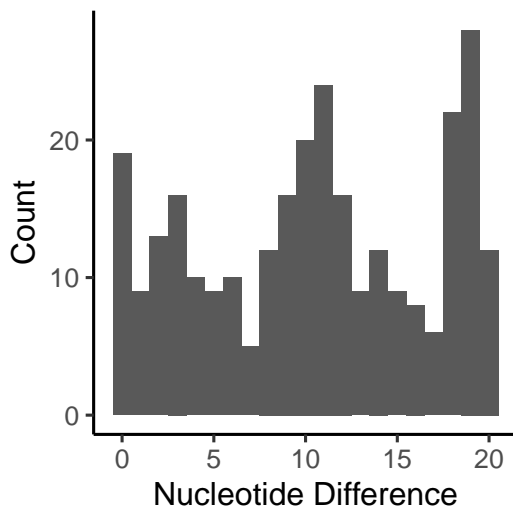
IGHV3-13*05

133 sequences assigned
19 (14.3%) exact matches, in which:
19 unique CDR3
5 unique J



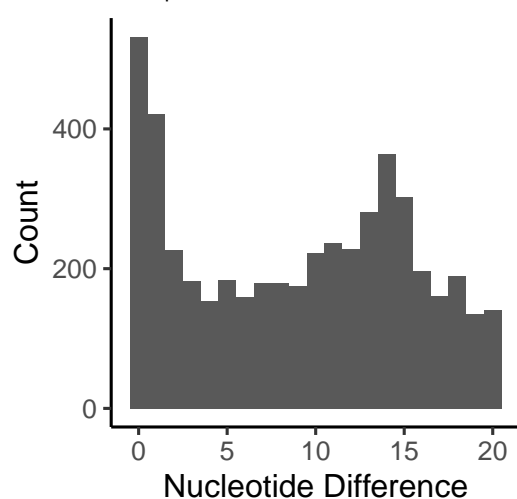
IGHV3-20*01_02

317 sequences assigned
19 (6%) exact matches, in which:
19 unique CDR3
5 unique J



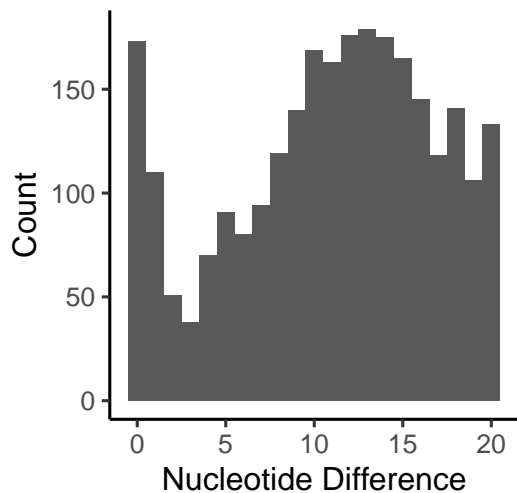
IGHV3-21*01_02

5903 sequences assigned
532 (9%) exact matches, in which:
499 unique CDR3
7 unique J



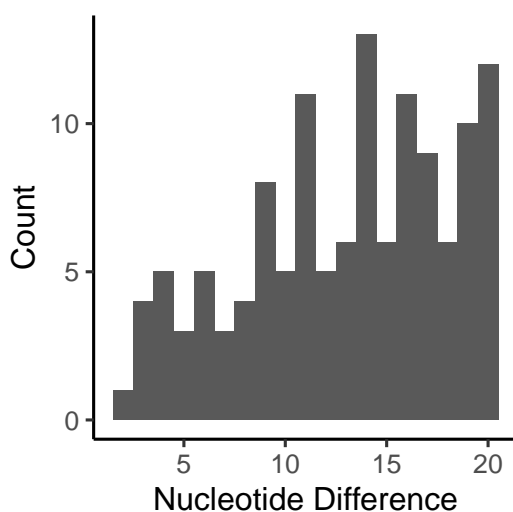
IGHV3-15*01_02

3692 sequences assigned
173 (4.7%) exact matches, in which:
170 unique CDR3
7 unique J



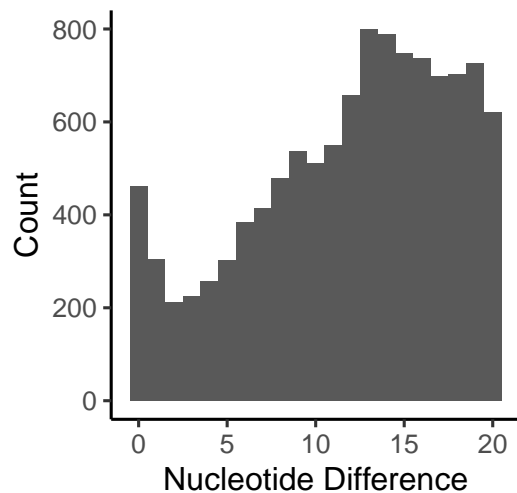
IGHV3-20*03_04

201 sequences assigned
No exact matches.



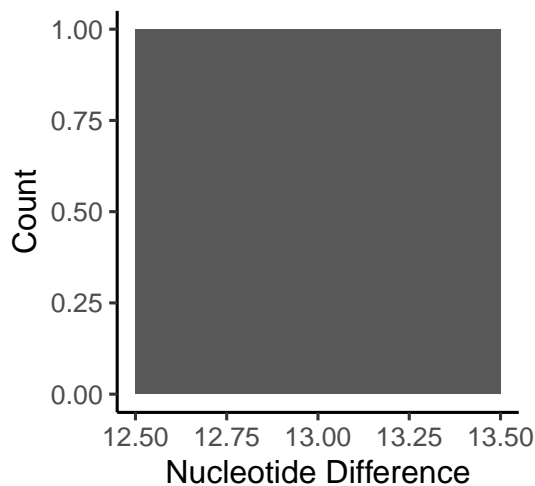
IGHV3-23*01_04

16228 sequences assigned
461 (2.8%) exact matches, in which:
458 unique CDR3
7 unique J



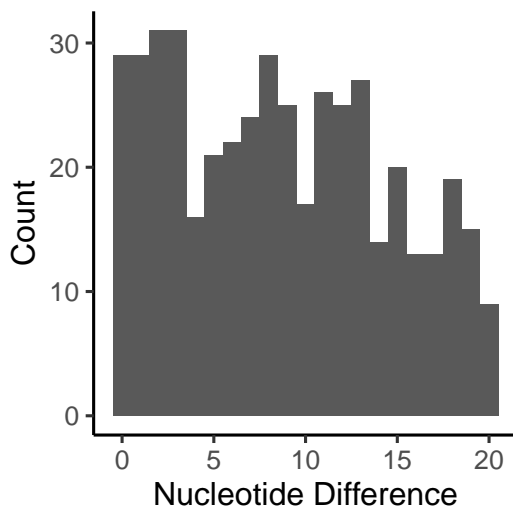
IGHV3-19*01

1 sequences assigned
No exact matches.



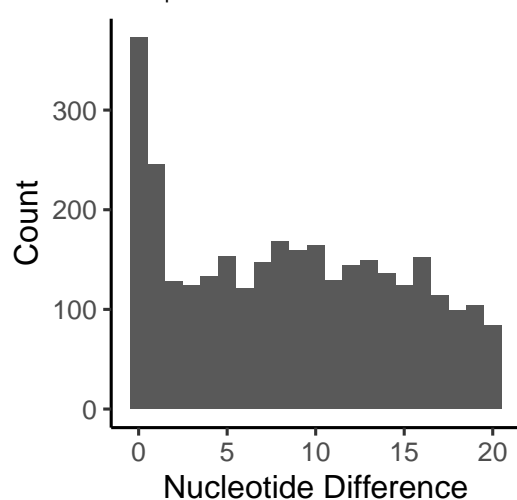
IGHV3-21*04

564 sequences assigned
29 (5.1%) exact matches, in which:
29 unique CDR3
5 unique J



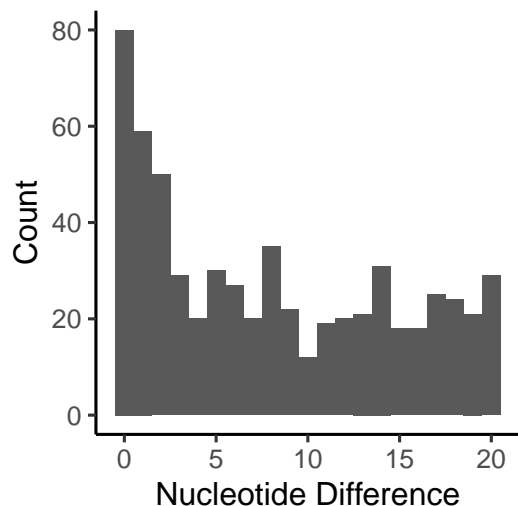
IGHV3-30-3*01

3996 sequences assigned
373 (9.3%) exact matches, in which:
368 unique CDR3
7 unique J



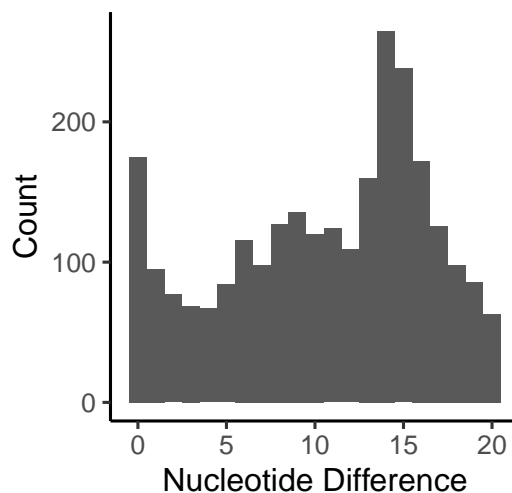
IGHV3-30*03

873 sequences assigned
80 (9.2%) exact matches, in which:
80 unique CDR3
7 unique J



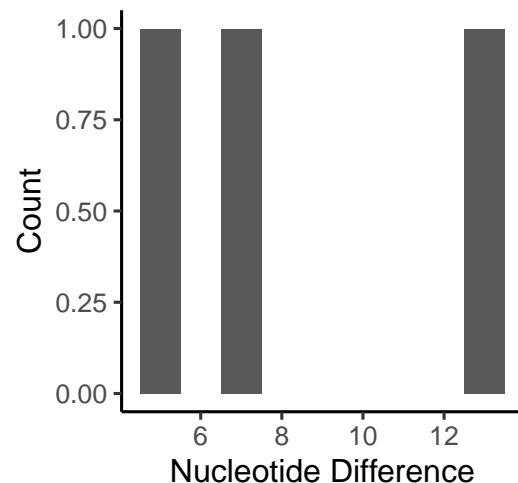
IGHV3-33*01

3166 sequences assigned
175 (5.5%) exact matches, in which:
173 unique CDR3
6 unique J



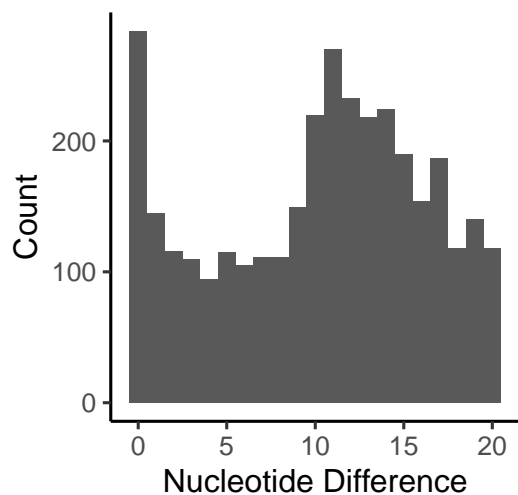
IGHV3-35*01

3 sequences assigned
No exact matches.



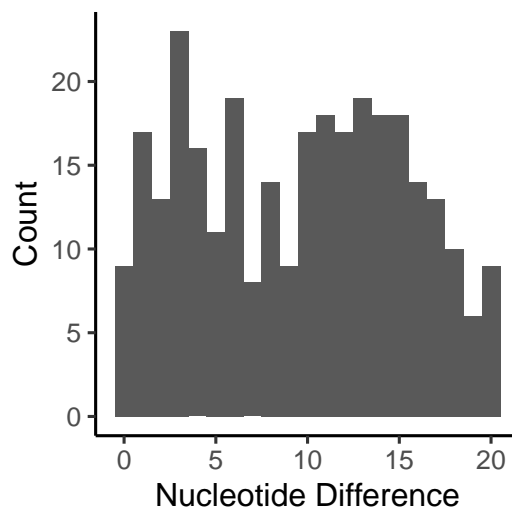
IGHV3-30*04

4269 sequences assigned
284 (6.7%) exact matches, in which:
283 unique CDR3
7 unique J



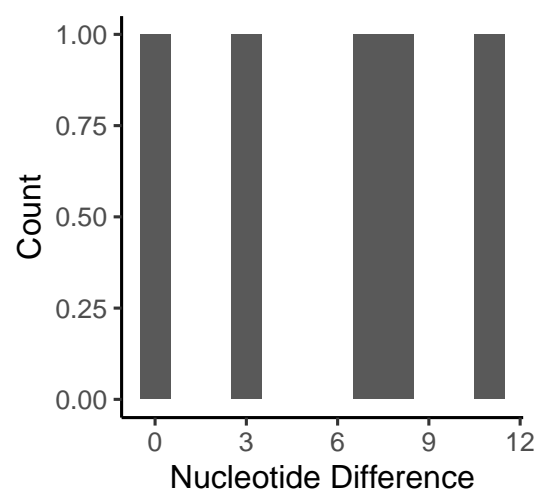
IGHV3-33*06

396 sequences assigned
9 (2.3%) exact matches, in which:
9 unique CDR3
3 unique J



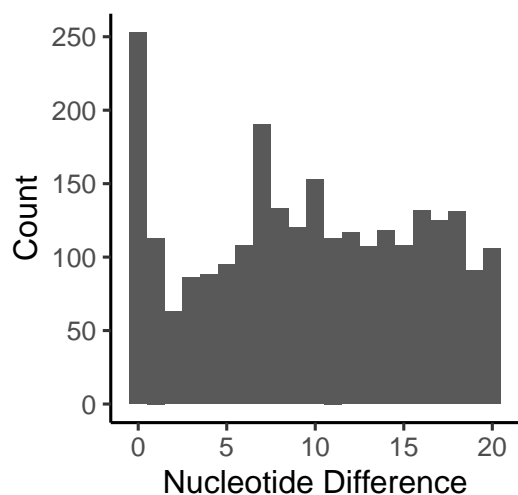
IGHV3-38-3*01

5 sequences assigned
1 (20%) exact matches, in which:
1 unique CDR3
1 unique J



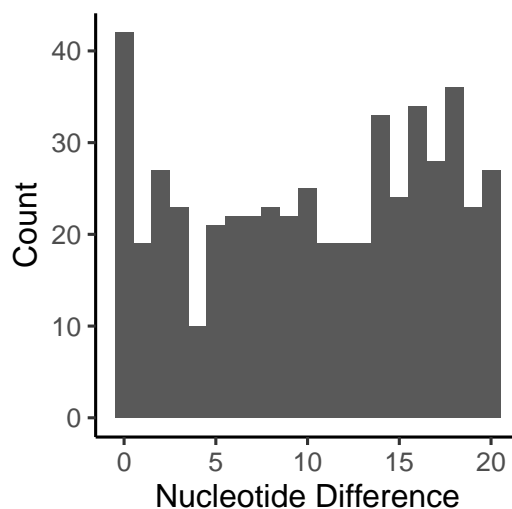
IGHV3-30*18

3208 sequences assigned
253 (7.9%) exact matches, in which:
250 unique CDR3
6 unique J



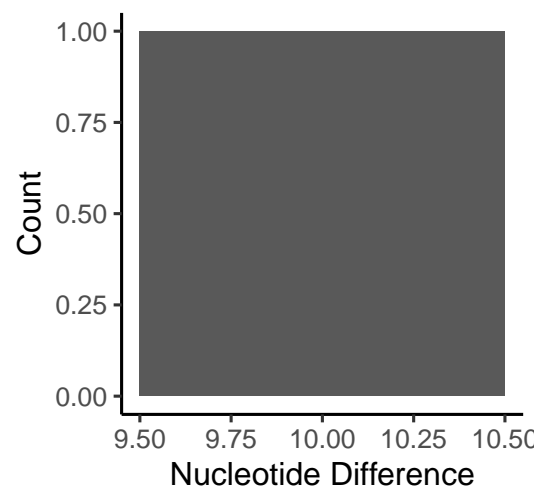
IGHV3-33*08

689 sequences assigned
42 (6.1%) exact matches, in which:
41 unique CDR3
5 unique J



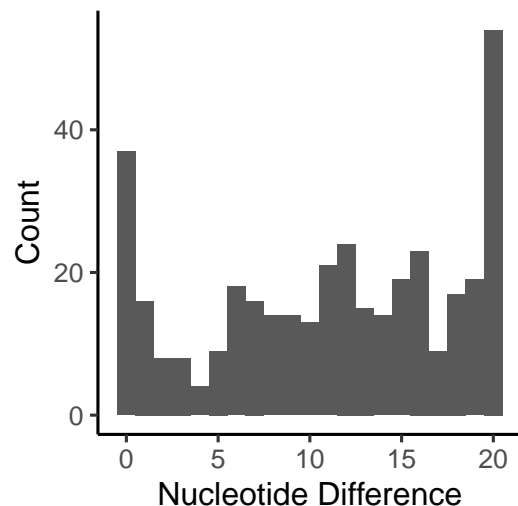
IGHV3-38*03

1 sequences assigned
No exact matches.



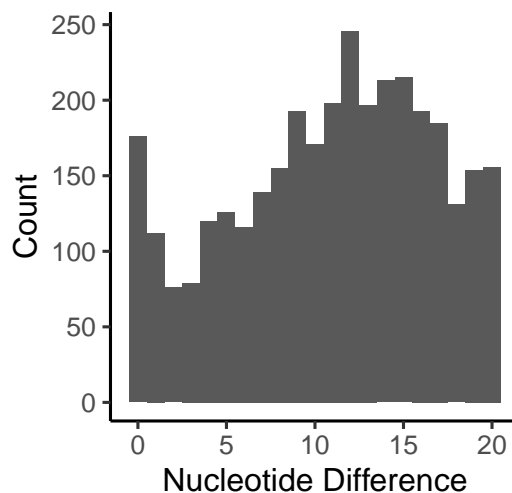
IGHV3-43*01

581 sequences assigned
37 (6.4%) exact matches, in which:
37 unique CDR3
4 unique J



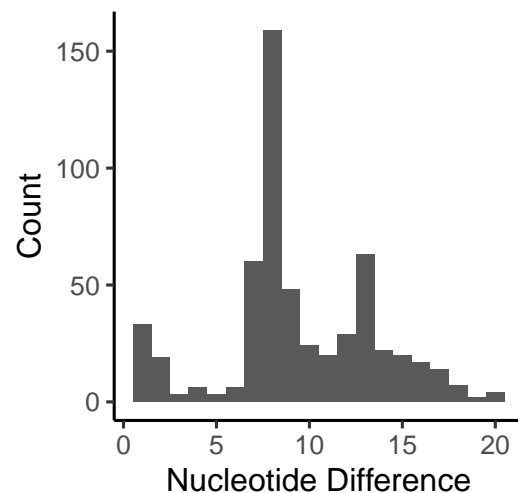
IGHV3-48*04

4424 sequences assigned
176 (4%) exact matches, in which:
172 unique CDR3
7 unique J



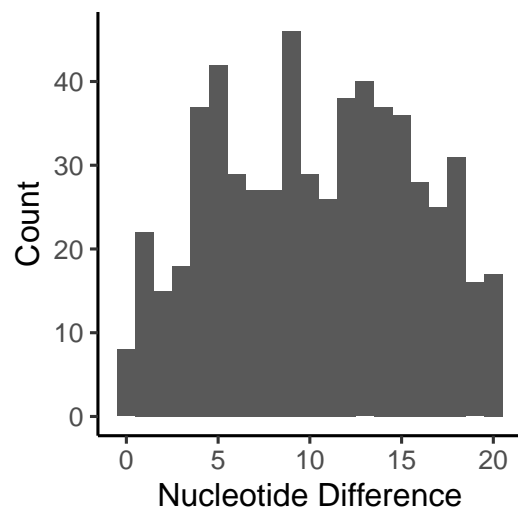
IGHV3-53*05

596 sequences assigned
No exact matches.



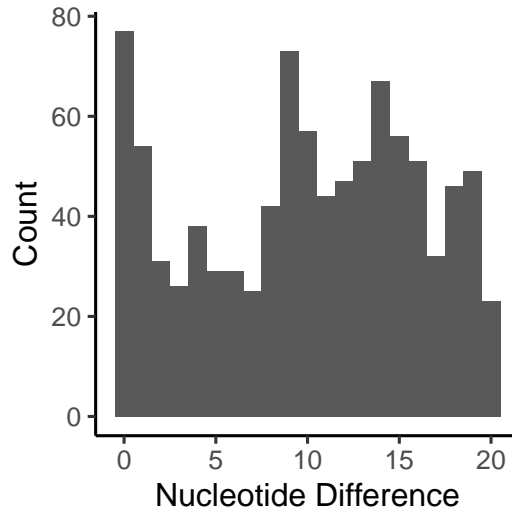
IGHV3-48*01

721 sequences assigned
8 (1.1%) exact matches, in which:
8 unique CDR3
3 unique J



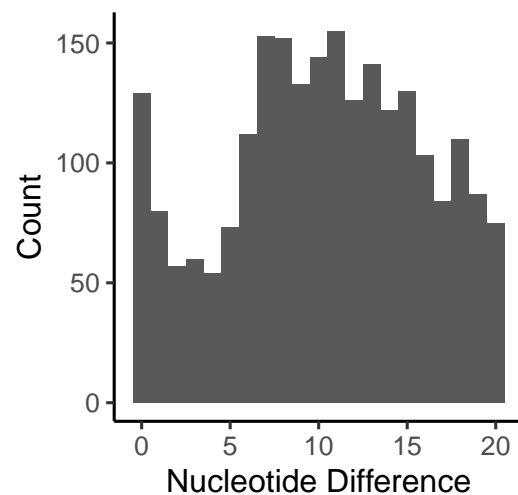
IGHV3-49*03_05

1205 sequences assigned
77 (6.4%) exact matches, in which:
73 unique CDR3
7 unique J



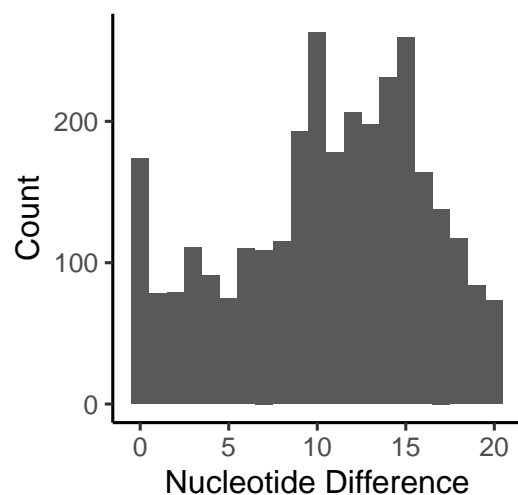
IGHV3-53*01_02

3054 sequences assigned
129 (4.2%) exact matches, in which:
126 unique CDR3
7 unique J



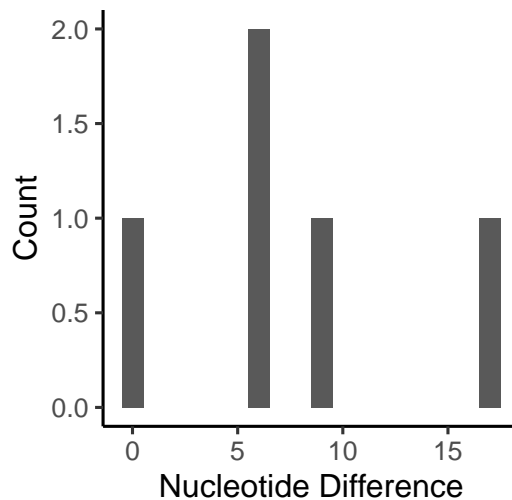
IGHV3-48*02

3580 sequences assigned
174 (4.9%) exact matches, in which:
170 unique CDR3
6 unique J



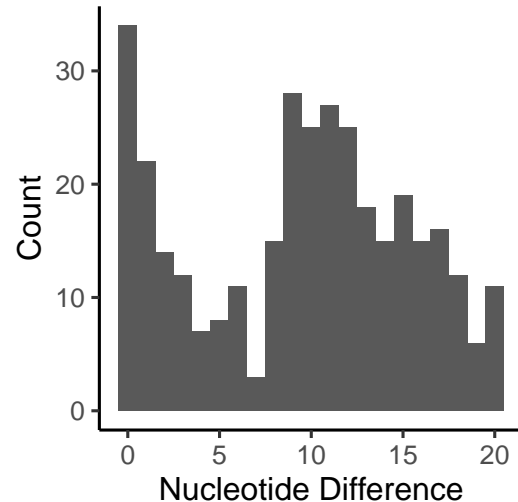
IGHV3-52*01_03

6 sequences assigned
1 (16.7%) exact matches, in which:
1 unique CDR3
1 unique J



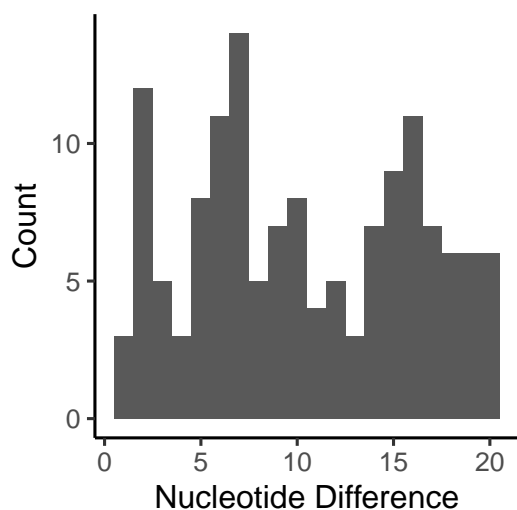
IGHV3-64*01

474 sequences assigned
34 (7.2%) exact matches, in which:
32 unique CDR3
4 unique J



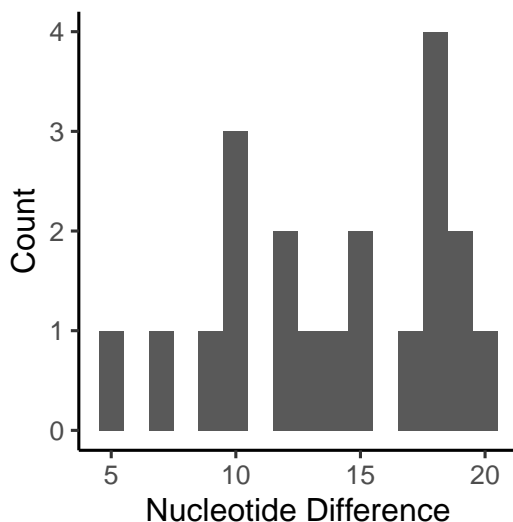
IGHV3-64*04

197 sequences assigned
No exact matches.



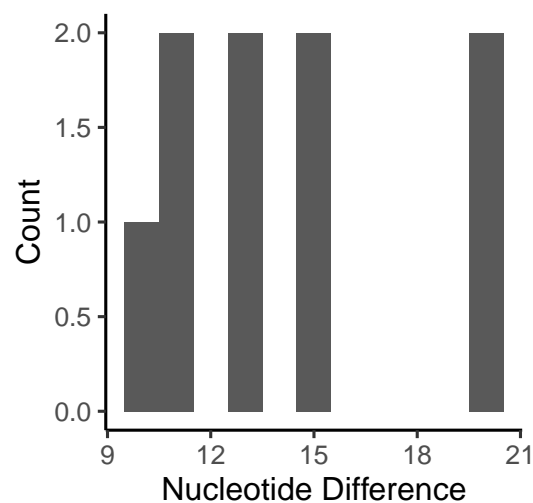
IGHV3-66*02

28 sequences assigned
No exact matches.



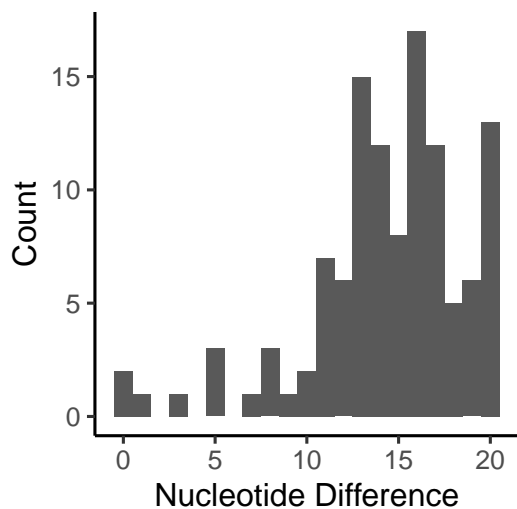
IGHV3-69-1*02

36 sequences assigned
No exact matches.



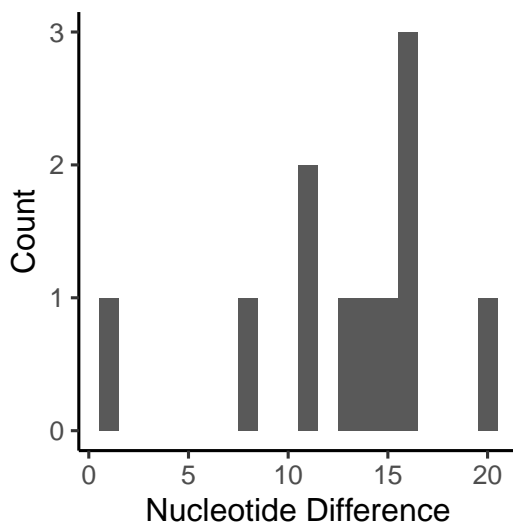
IGHV3-64*02_07

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



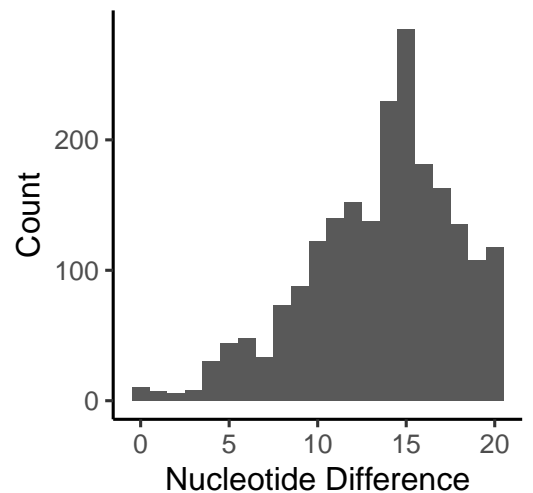
IGHV3-66*03

15 sequences assigned
No exact matches.



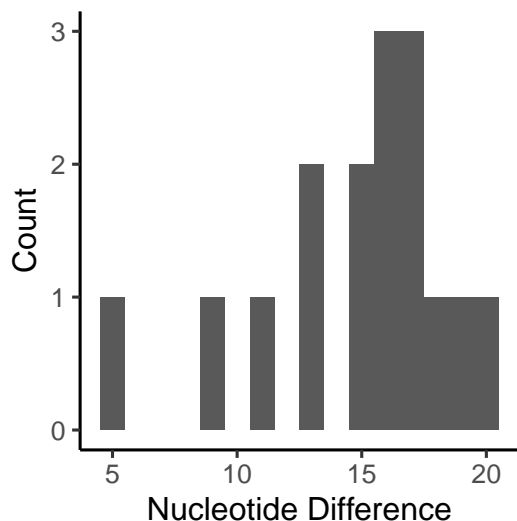
IGHV3-72*01

2896 sequences assigned
10 (0.3%) exact matches, in which:
10 unique CDR3
4 unique J



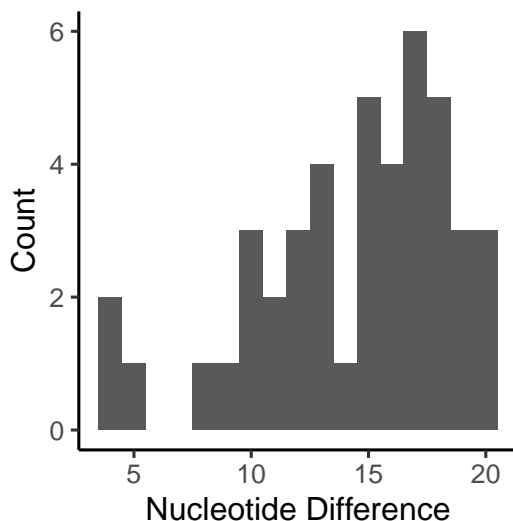
IGHV3-66*01

21 sequences assigned
No exact matches.



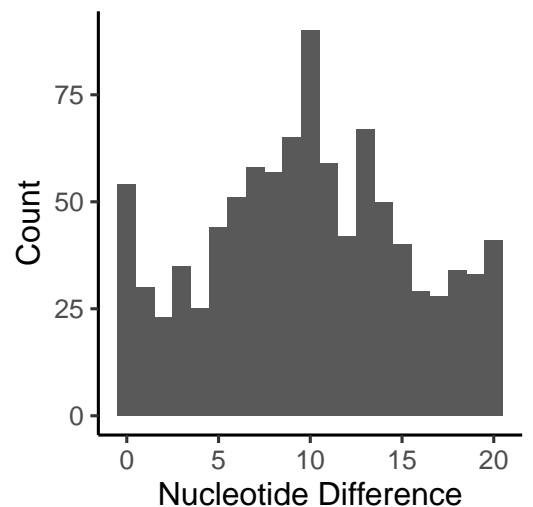
IGHV3-69-1*01

95 sequences assigned
No exact matches.



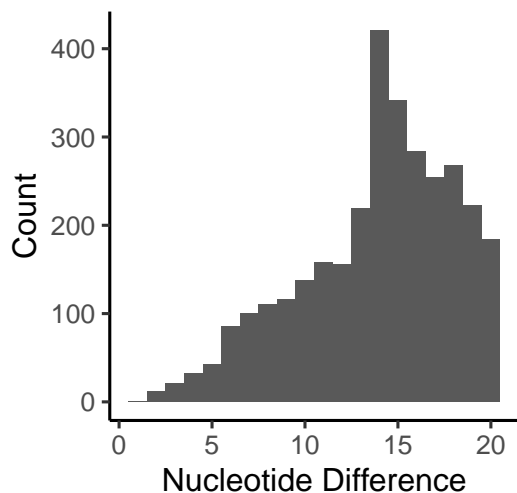
IGHV3-73*01_02

1125 sequences assigned
54 (4.8%) exact matches, in which:
54 unique CDR3
7 unique J



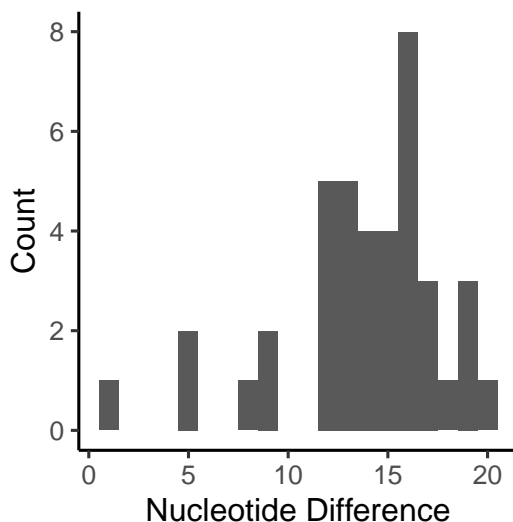
IGHV3-74*03

4513 sequences assigned
No exact matches.



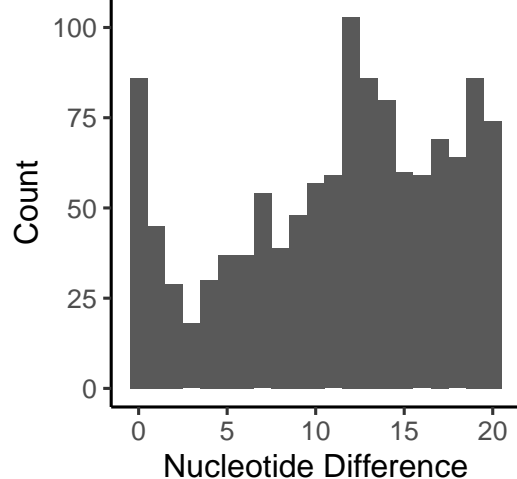
IGHV3-43D*04

72 sequences assigned
No exact matches.



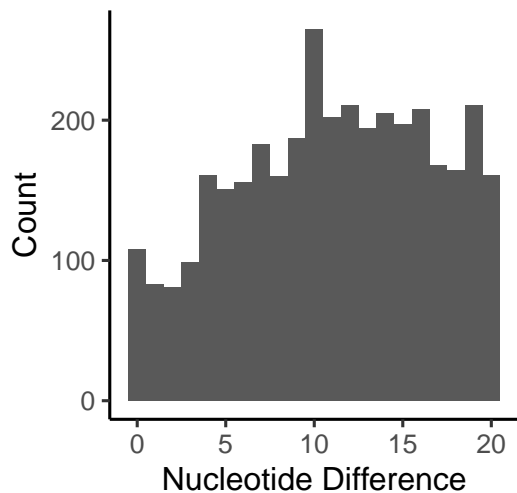
IGHV4-4*02_03

1749 sequences assigned
86 (4.9%) exact matches, in which:
86 unique CDR3
7 unique J



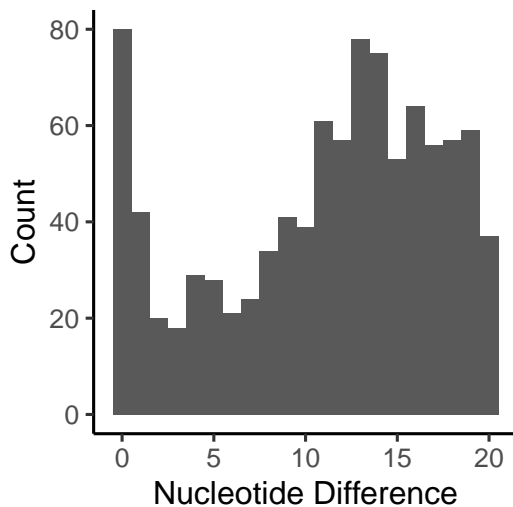
IGHV3-74*01_02

4208 sequences assigned
108 (2.6%) exact matches, in which:
102 unique CDR3
6 unique J



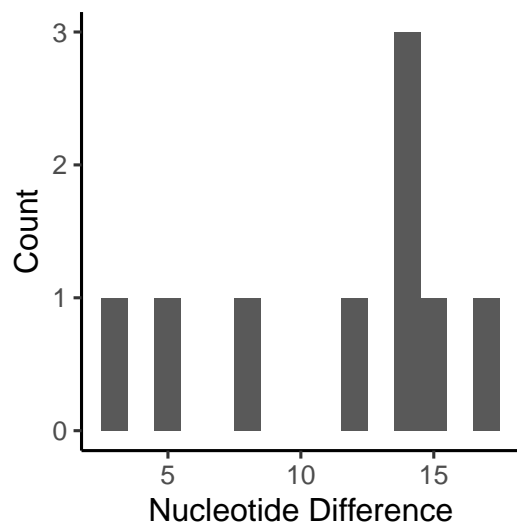
IGHV3-64D*06

1499 sequences assigned
80 (5.3%) exact matches, in which:
80 unique CDR3
7 unique J



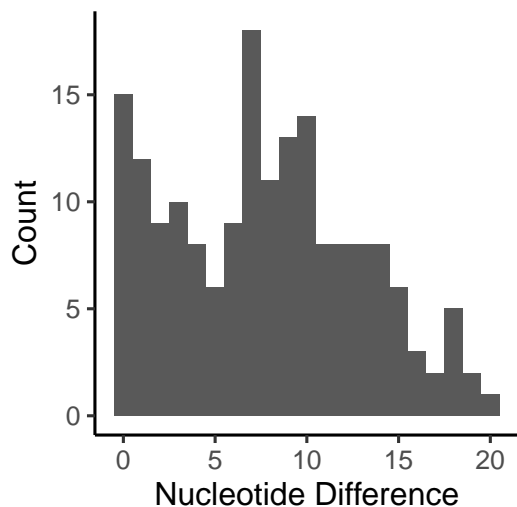
IGHV4-28*03

11 sequences assigned
No exact matches.



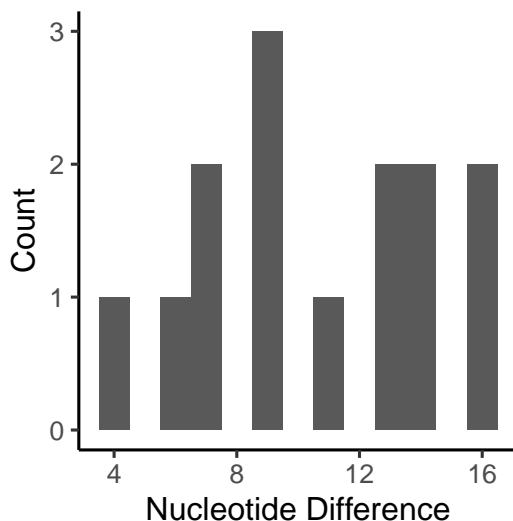
IGHV3-43D*03

187 sequences assigned
15 (8%) exact matches, in which:
14 unique CDR3
2 unique J



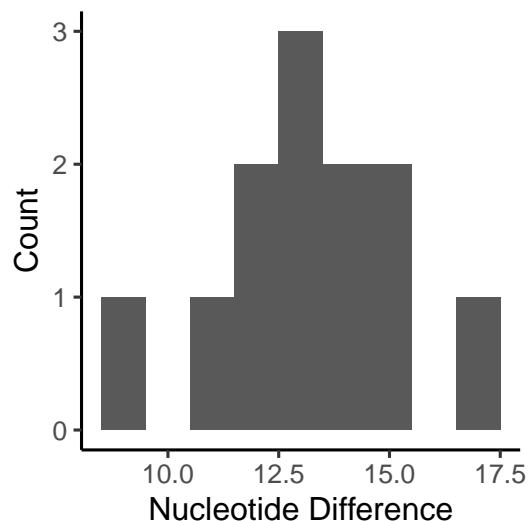
IGHV3-NL1*01

29 sequences assigned
No exact matches.



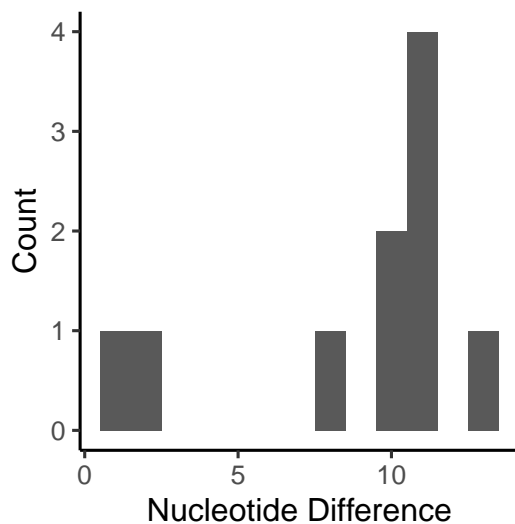
IGHV4-28*01_07

13 sequences assigned
No exact matches.



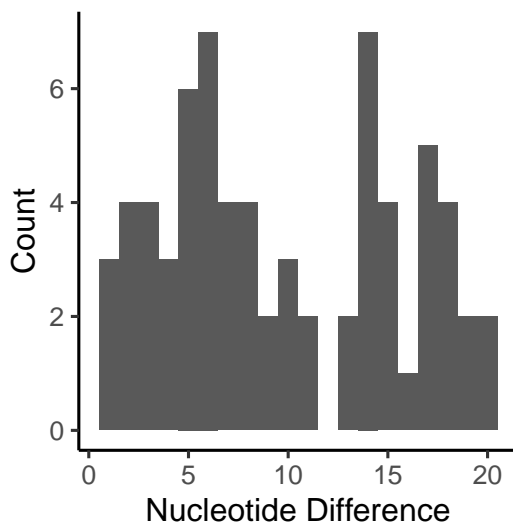
IGHV4-28*02_05

10 sequences assigned
No exact matches.



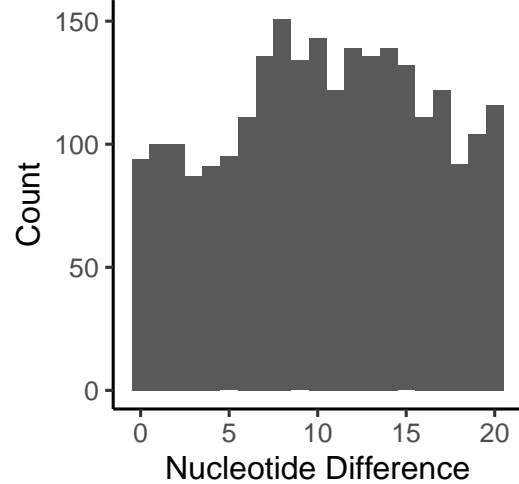
IGHV4-30-4*08

103 sequences assigned
No exact matches.



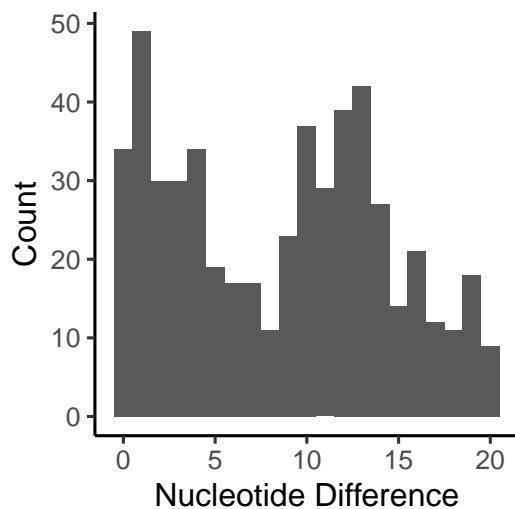
IGHV4-38-2*02

3490 sequences assigned
94 (2.7%) exact matches, in which:
92 unique CDR3
7 unique J



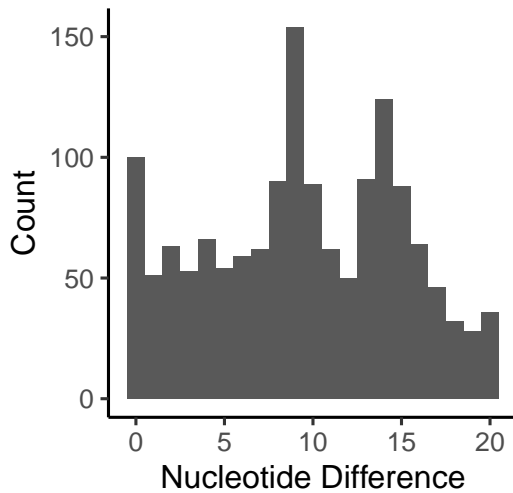
IGHV4-30-2*01

711 sequences assigned
34 (4.8%) exact matches, in which:
34 unique CDR3
5 unique J



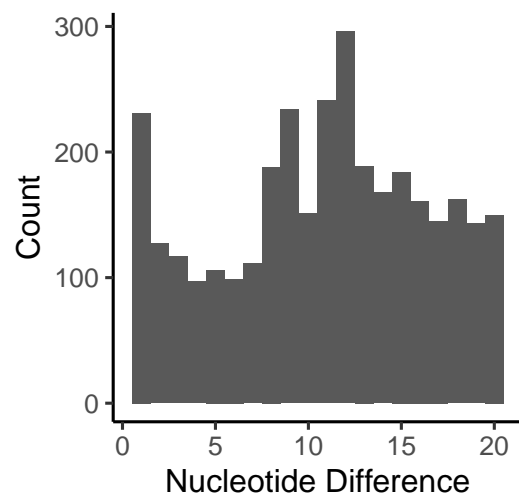
IGHV4-31*02

1838 sequences assigned
100 (5.4%) exact matches, in which:
99 unique CDR3
6 unique J



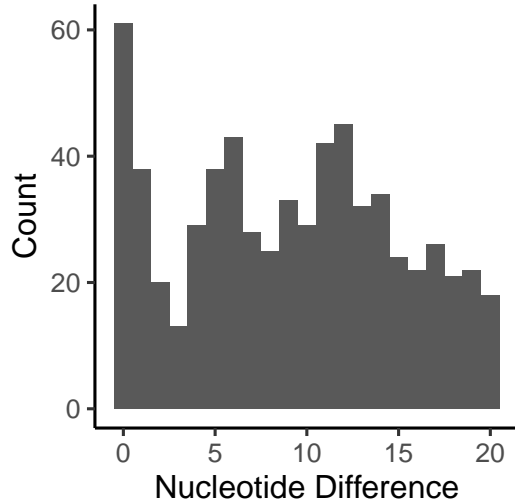
IGHV4-39*02

5385 sequences assigned
No exact matches.



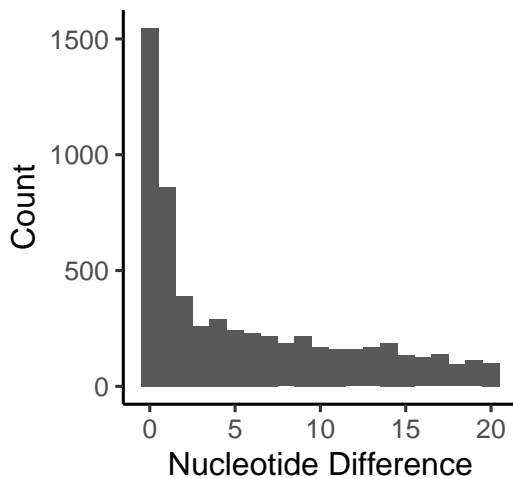
IGHV4-30-4*01

905 sequences assigned
61 (6.7%) exact matches, in which:
60 unique CDR3
7 unique J



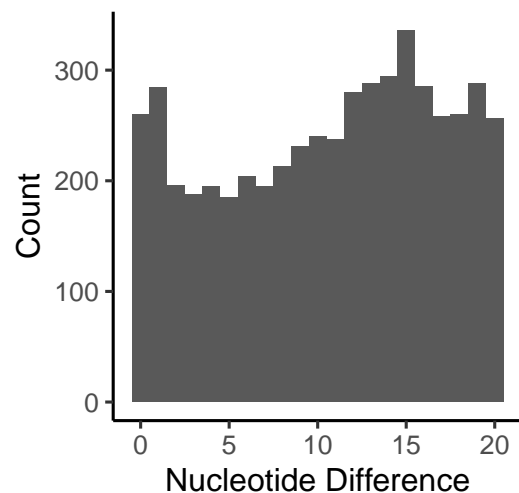
IGHV4-34*01_02

6879 sequences assigned
1547 (22.5%) exact matches, in which:
1507 unique CDR3
7 unique J



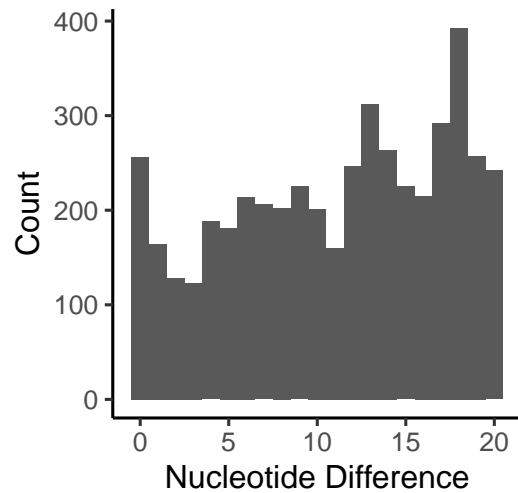
IGHV4-39*07

7302 sequences assigned
260 (3.6%) exact matches, in which:
257 unique CDR3
7 unique J



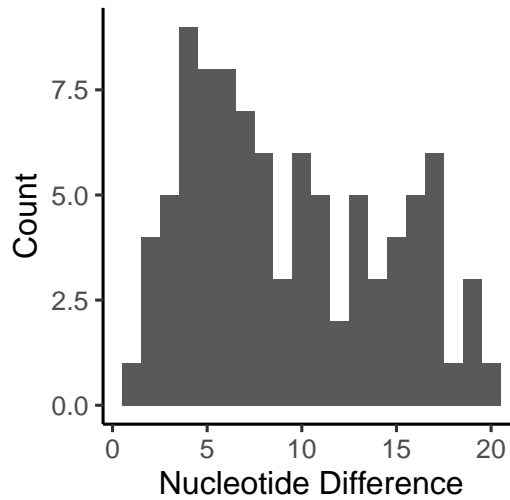
IGHV4–59*08

6279 sequences assigned
256 (4.1%) exact matches, in which:
251 unique CDR3
7 unique J



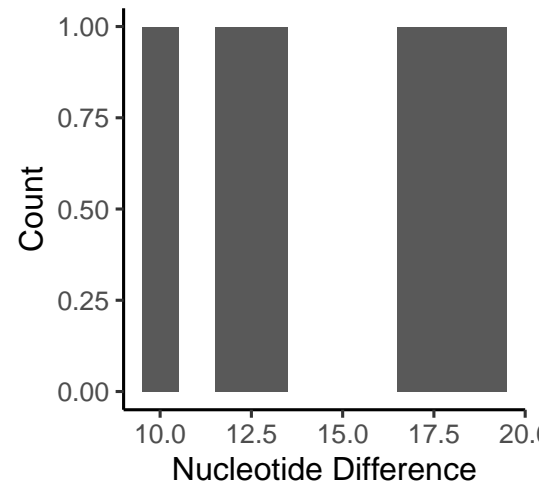
IGHV4–61*05

113 sequences assigned
No exact matches.



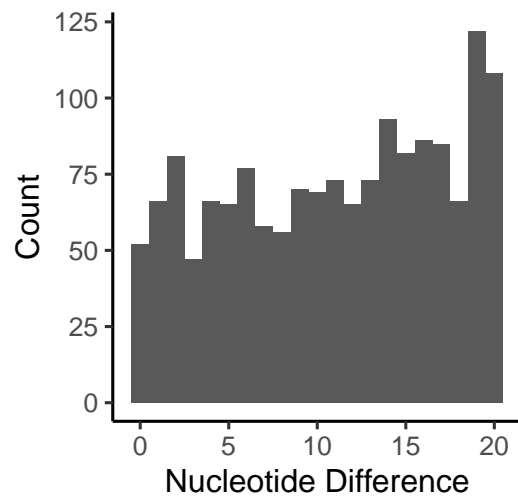
IGHV4/OR15–8*01

14 sequences assigned
No exact matches.



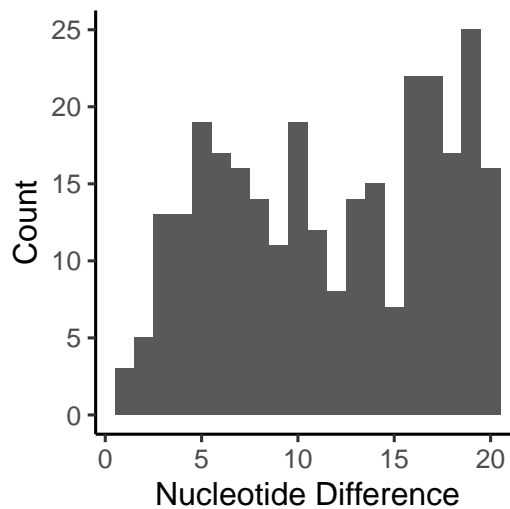
IGHV4–59*12

2633 sequences assigned
52 (2%) exact matches, in which:
52 unique CDR3
6 unique J



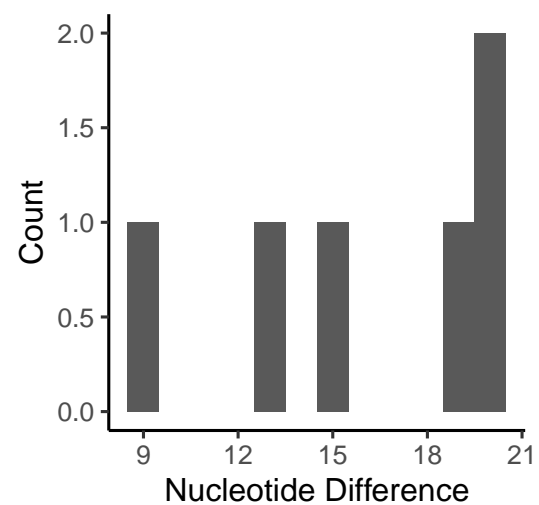
IGHV4–61*08

477 sequences assigned
No exact matches.



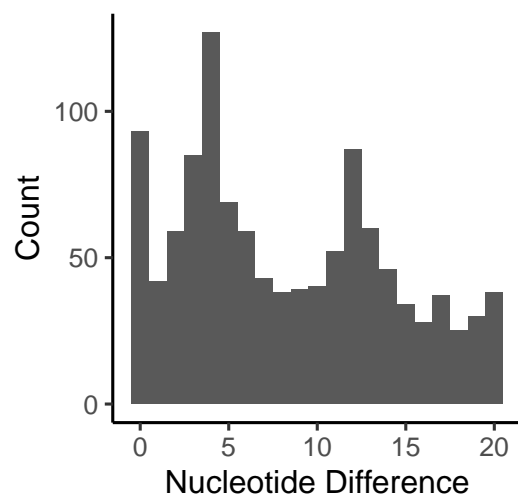
IGHV4/OR15–8*02

32 sequences assigned
No exact matches.



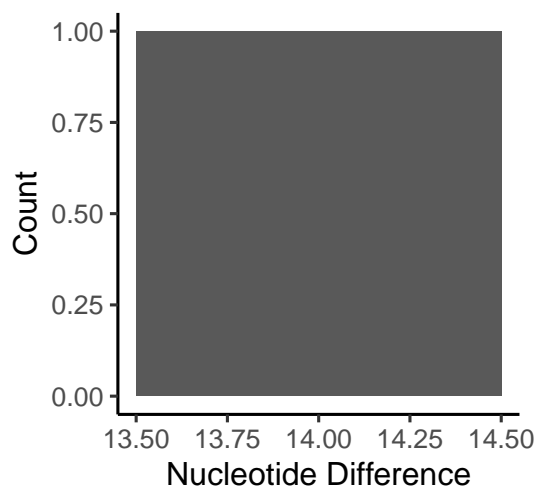
IGHV4–61*01

1547 sequences assigned
93 (6%) exact matches, in which:
91 unique CDR3
7 unique J



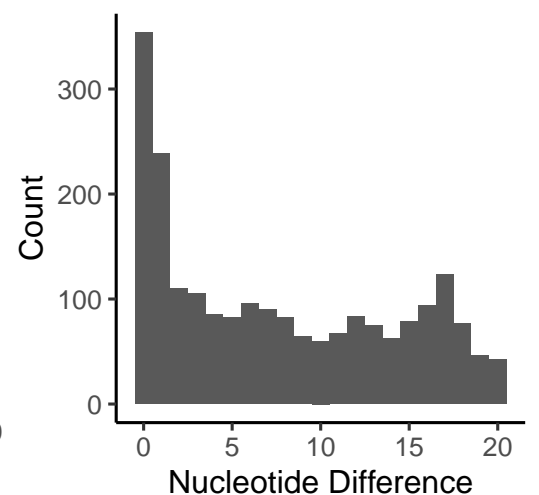
IGHV1/OR15–1*04

1 sequences assigned
No exact matches.



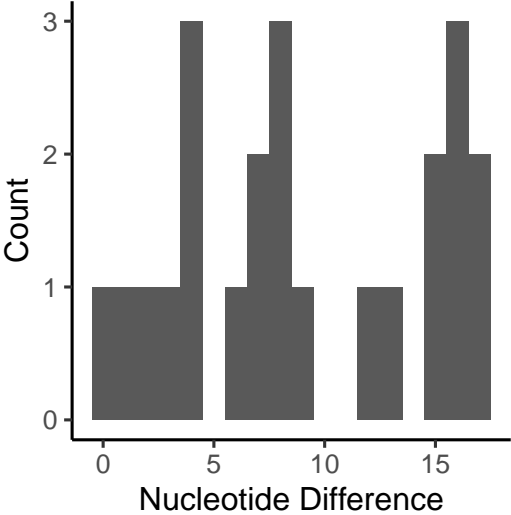
IGHV5–10–1*01_03

2366 sequences assigned
354 (15%) exact matches, in which:
350 unique CDR3
7 unique J



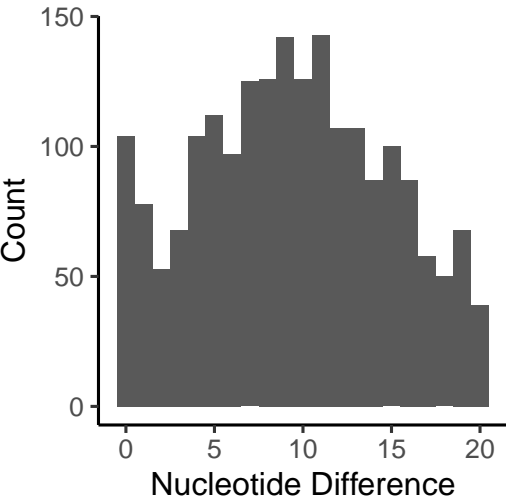
IGHV5-51*04

29 sequences assigned
1 (3.4%) exact matches, in which:
1 unique CDR3
1 unique J



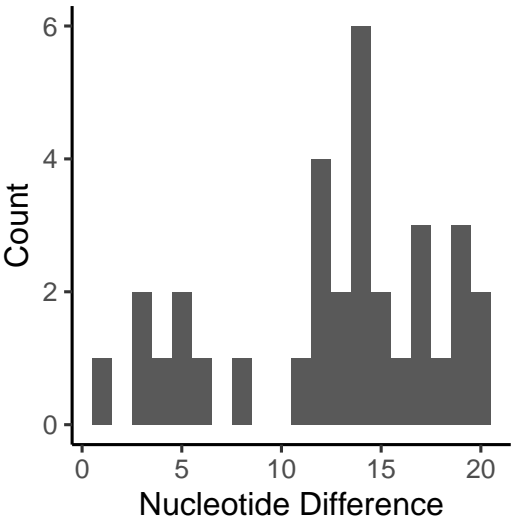
IGHV6-1*01_02

2116 sequences assigned
104 (4.9%) exact matches, in which:
104 unique CDR3
7 unique J



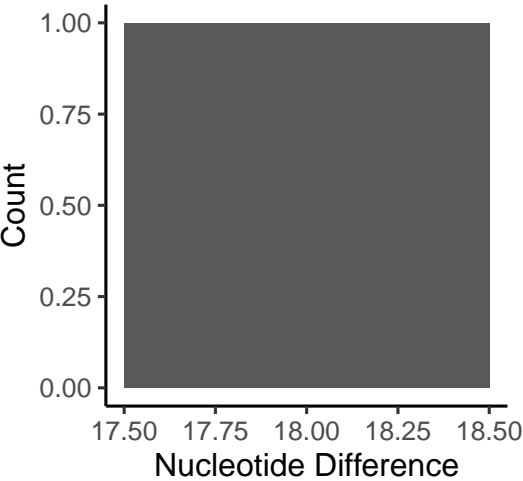
IGHV5-51*06

41 sequences assigned
No exact matches.



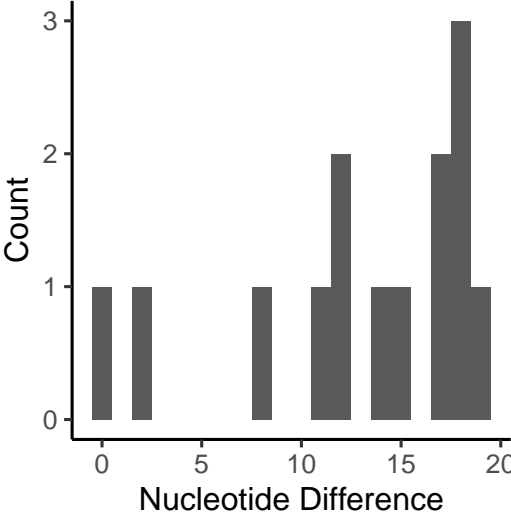
IGHV3/OR16-6*02

1 sequences assigned
No exact matches.



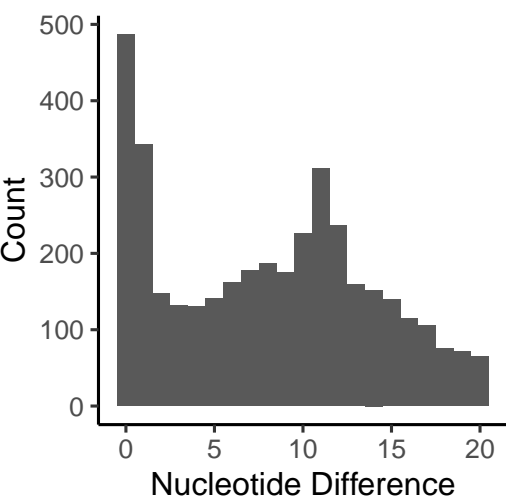
IGHV5-51*07

15 sequences assigned
1 (6.7%) exact matches, in which:
1 unique CDR3
1 unique J

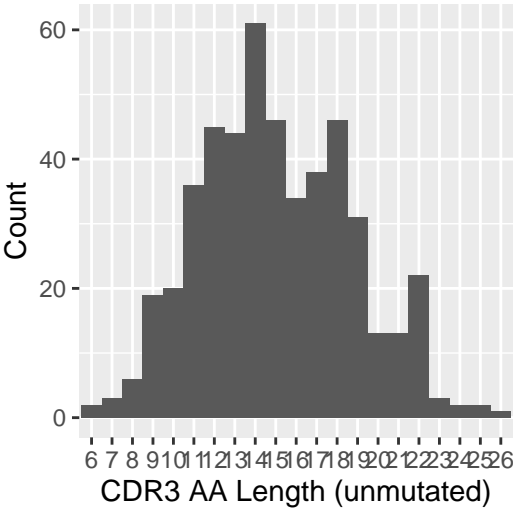


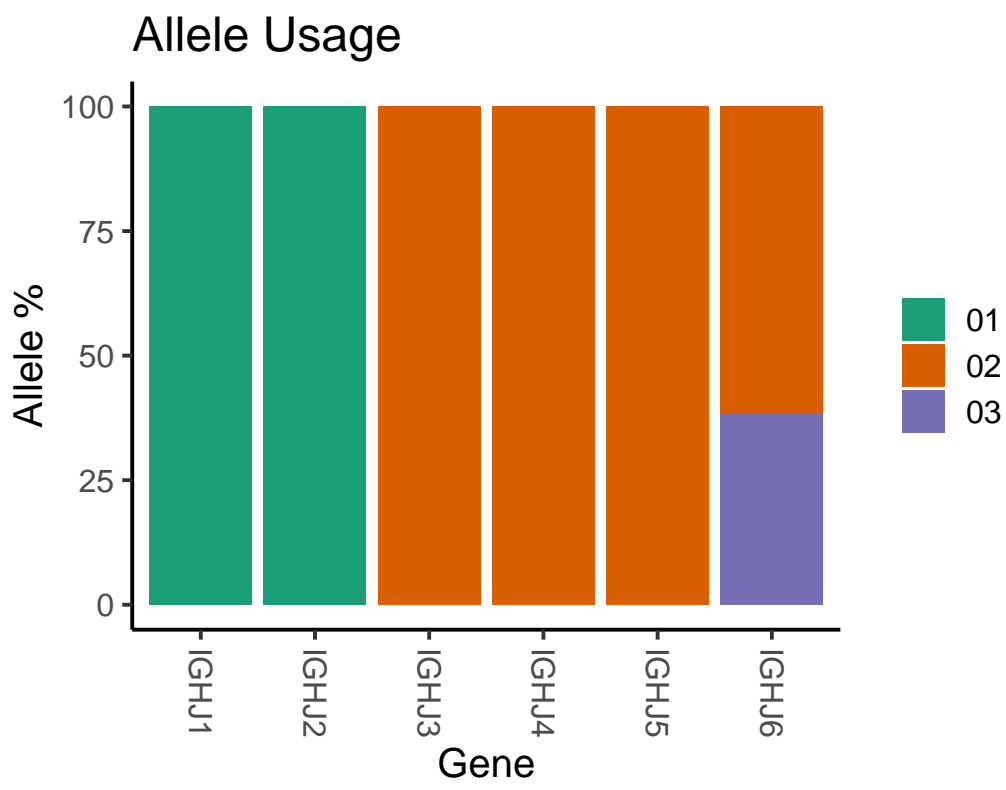
IGHV7-4-1*02_T63C

4448 sequences assigned
487 (10.9%) exact matches, in which:
478 unique CDR3
7 unique J



IGHV7-4-1*02_T63C





Sequence Count byIGHJ6 allele usage

