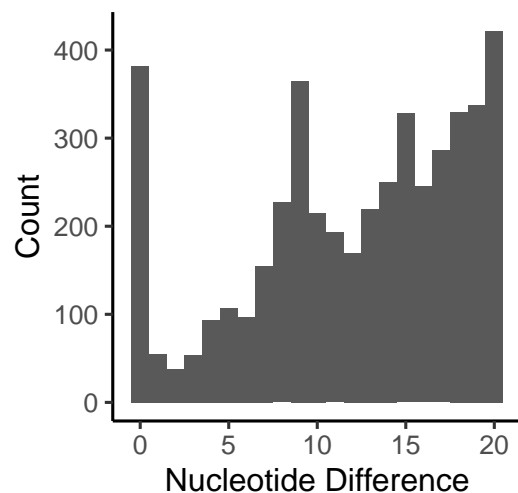


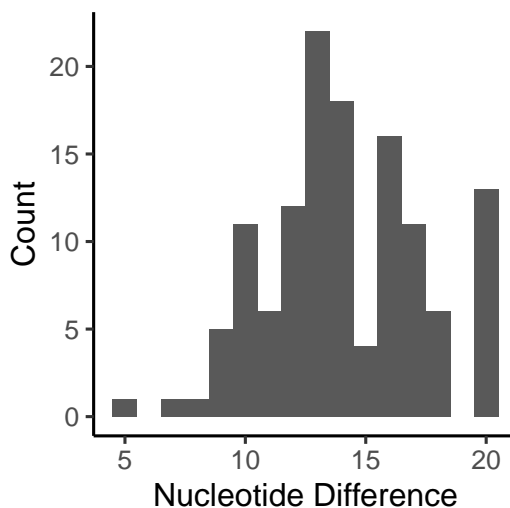
IGHV1-2*02

6891 sequences assigned
382 (5.5%) exact matches, in which:
353 unique CDR3
6 unique J



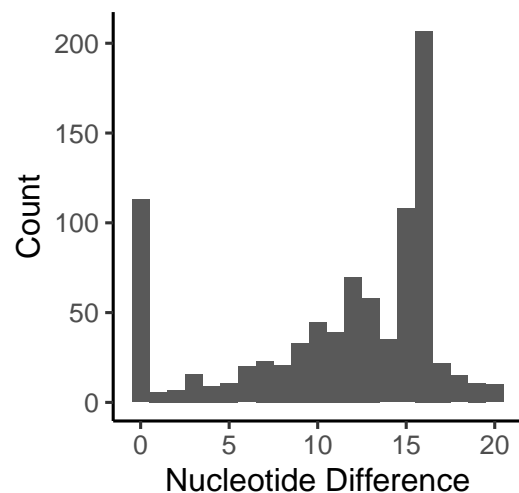
IGHV1-3*04

185 sequences assigned
No exact matches.



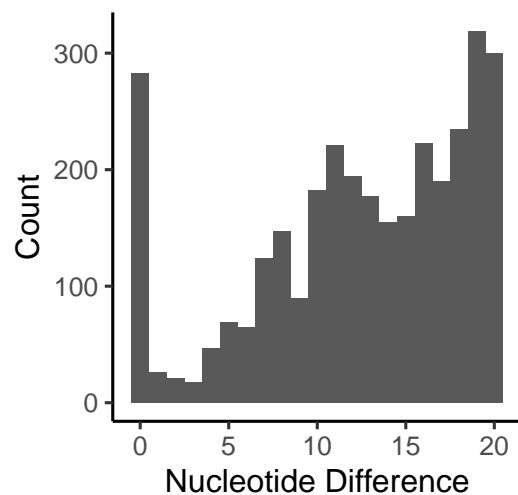
IGHV1-24*01

983 sequences assigned
113 (11.5%) exact matches, in which:
109 unique CDR3
6 unique J



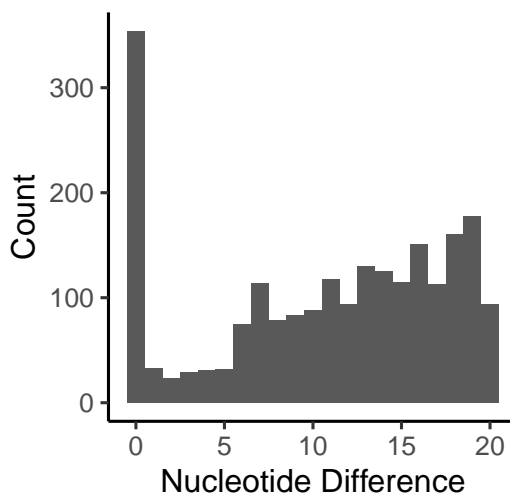
IGHV1-2*06

5384 sequences assigned
283 (5.3%) exact matches, in which:
268 unique CDR3
6 unique J



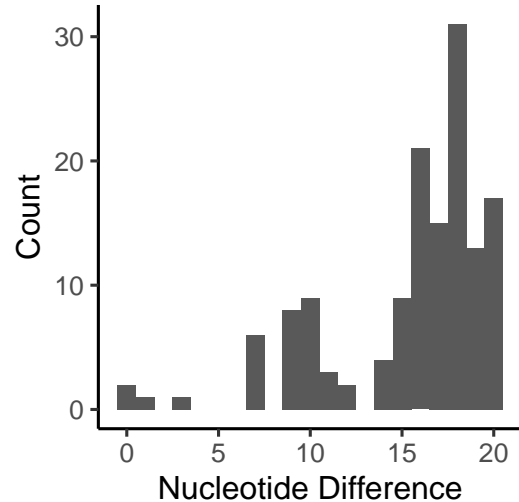
IGHV1-8*01

3327 sequences assigned
354 (10.6%) exact matches, in which:
332 unique CDR3
7 unique J



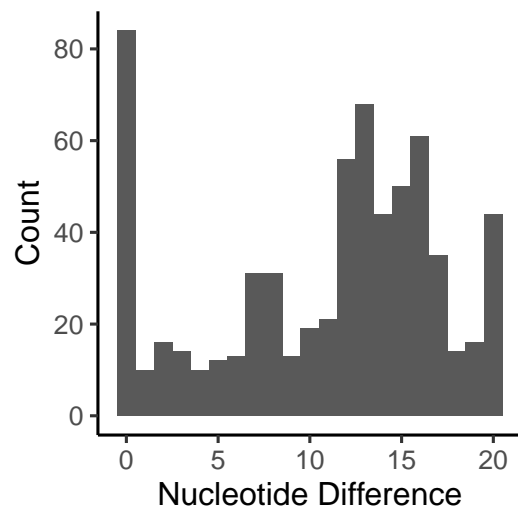
IGHV1-45*02

409 sequences assigned
2 (0.5%) exact matches, in which:
2 unique CDR3
2 unique J



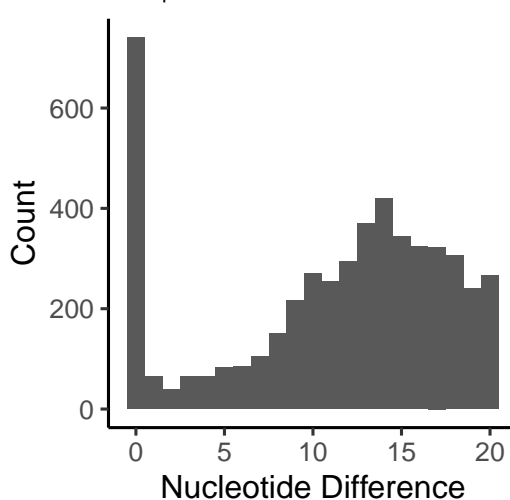
IGHV1-3*01

1014 sequences assigned
84 (8.3%) exact matches, in which:
81 unique CDR3
6 unique J



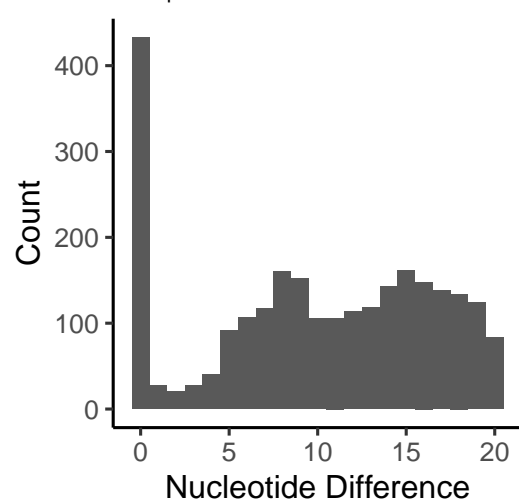
IGHV1-18*01

7573 sequences assigned
741 (9.8%) exact matches, in which:
691 unique CDR3
6 unique J



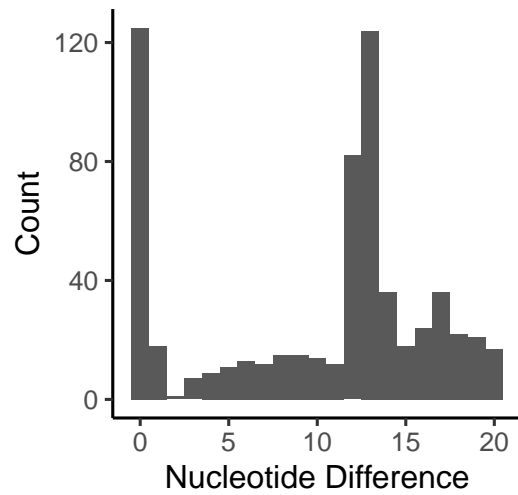
IGHV1-46*03

3883 sequences assigned
433 (11.2%) exact matches, in which:
396 unique CDR3
6 unique J



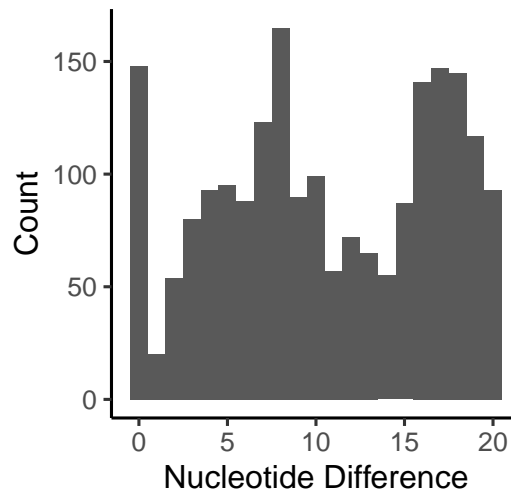
IGHV1–58*02

717 sequences assigned
125 (17.4%) exact matches, in which:
110 unique CDR3
5 unique J



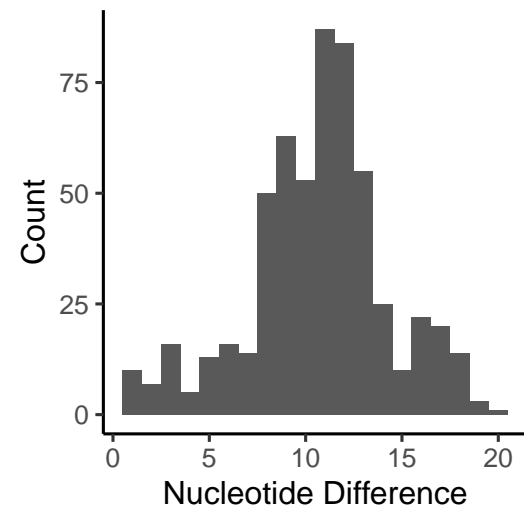
IGHV2–5*01

2334 sequences assigned
148 (6.3%) exact matches, in which:
127 unique CDR3
5 unique J



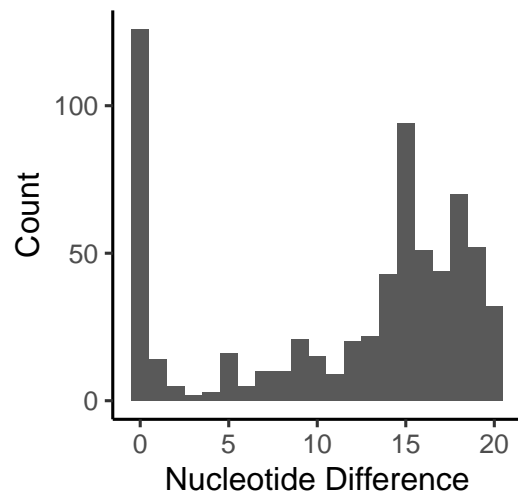
IGHV2–70*17

594 sequences assigned
No exact matches.



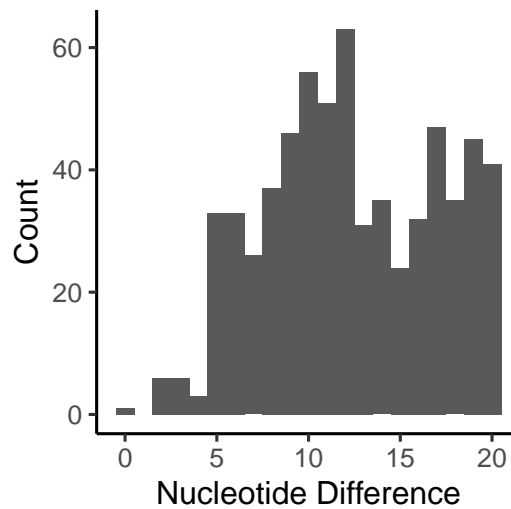
IGHV1–69*02

2215 sequences assigned
126 (5.7%) exact matches, in which:
122 unique CDR3
5 unique J



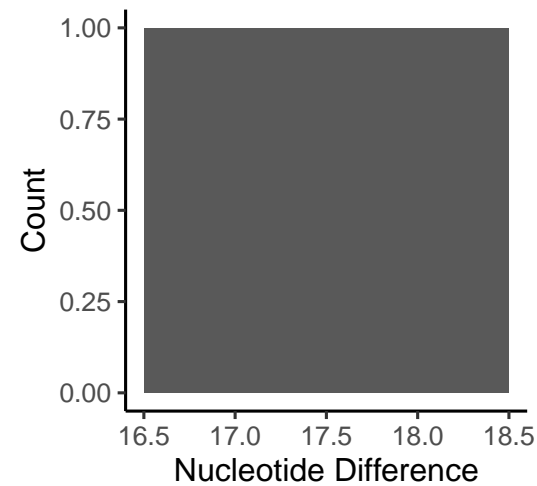
IGHV2–5*02

832 sequences assigned
1 (0.1%) exact matches, in which:
1 unique CDR3
1 unique J



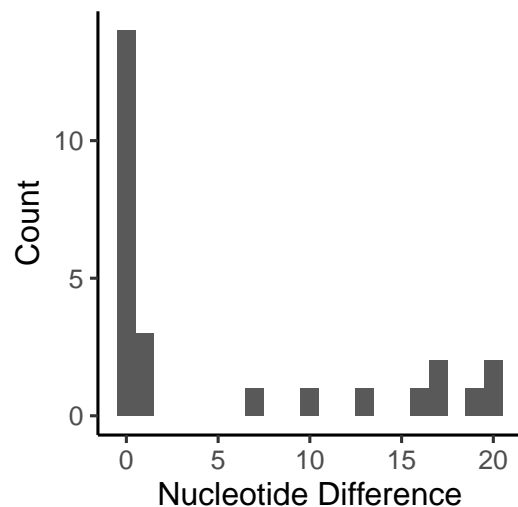
IGHV2–70D*04

5 sequences assigned
No exact matches.



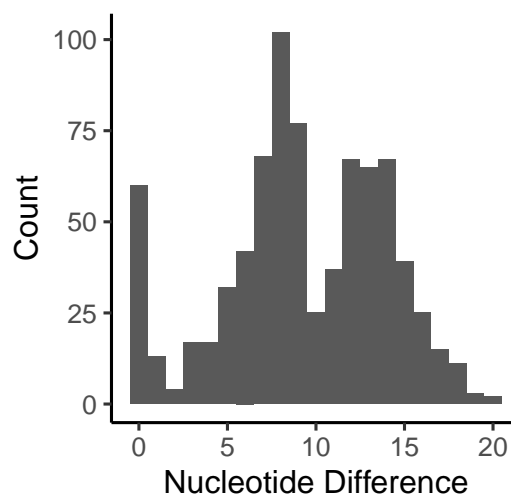
IGHV1–69D*01

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



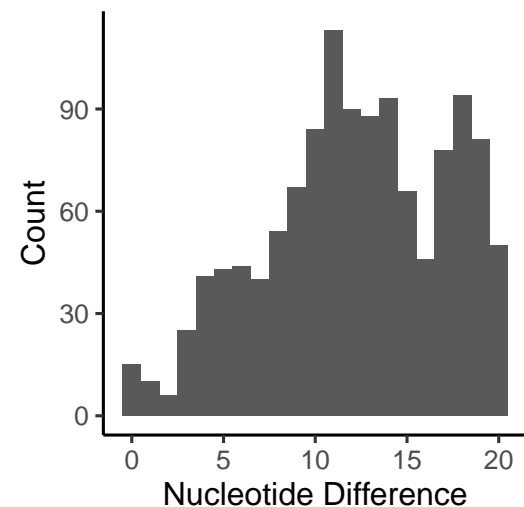
IGHV2–26*01

805 sequences assigned
60 (7.5%) exact matches, in which:
57 unique CDR3
5 unique J



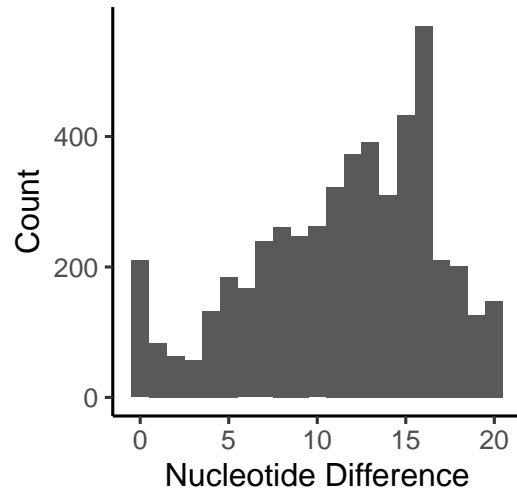
IGHV3–7*01

1559 sequences assigned
15 (1%) exact matches, in which:
12 unique CDR3
3 unique J



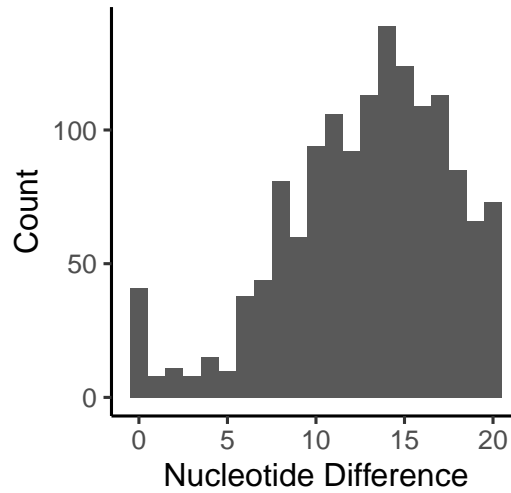
IGHV3-9*01

6127 sequences assigned
210 (3.4%) exact matches, in which:
181 unique CDR3
6 unique J



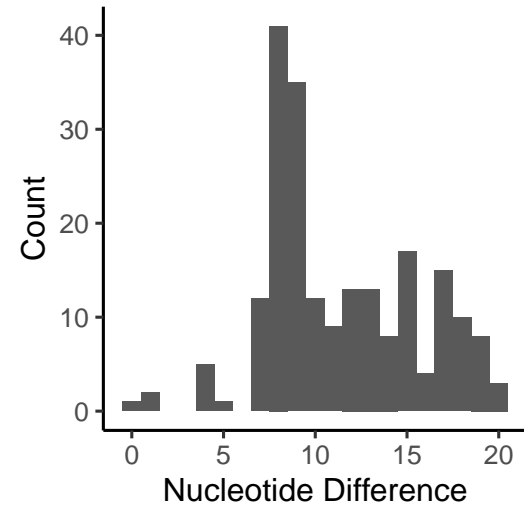
IGHV3-15*01

1932 sequences assigned
41 (2.1%) exact matches, in which:
36 unique CDR3
4 unique J



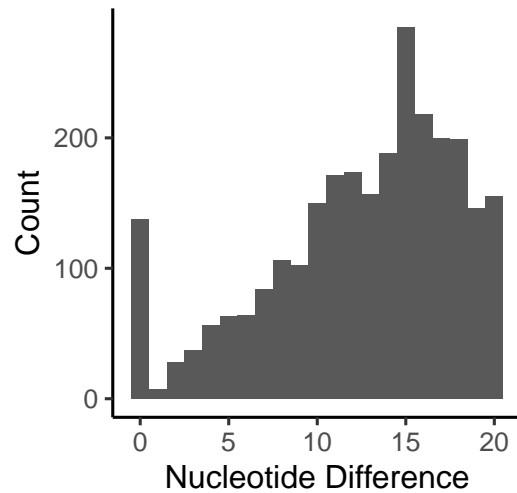
IGHV3-21*06

317 sequences assigned
1 (0.3%) exact matches, in which:
1 unique CDR3
1 unique J



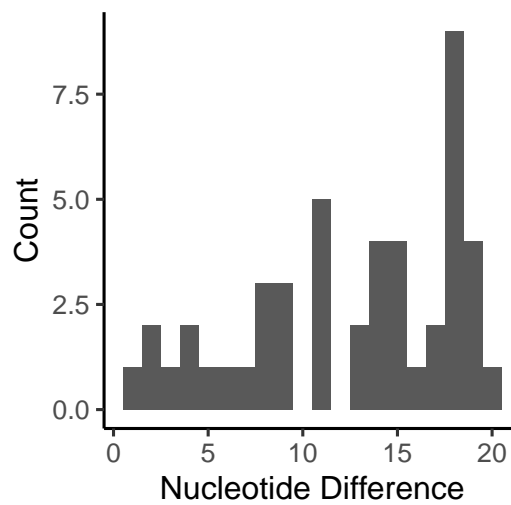
IGHV3-11*01

3817 sequences assigned
138 (3.6%) exact matches, in which:
125 unique CDR3
6 unique J



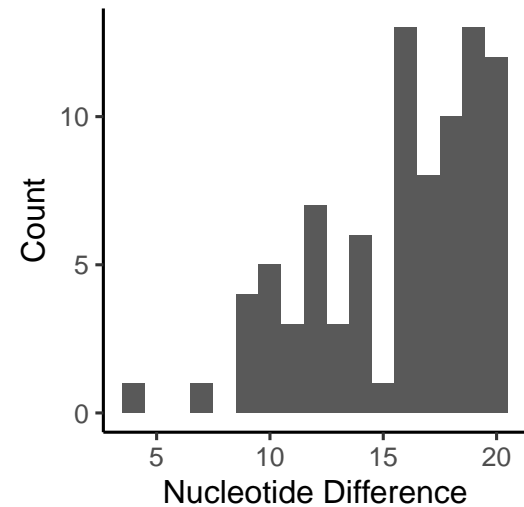
IGHV3-20*01

65 sequences assigned
No exact matches.



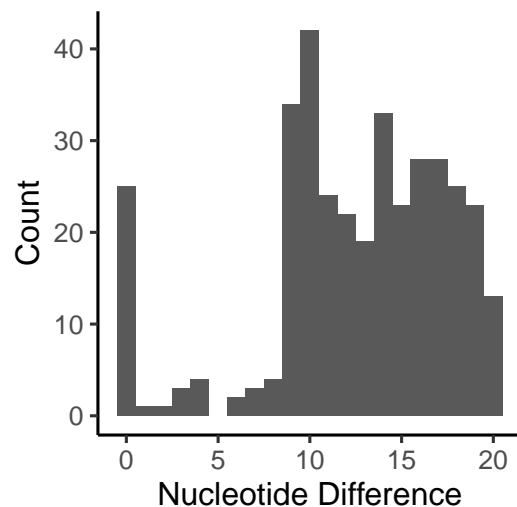
IGHV3-30-5*01

264 sequences assigned
No exact matches.



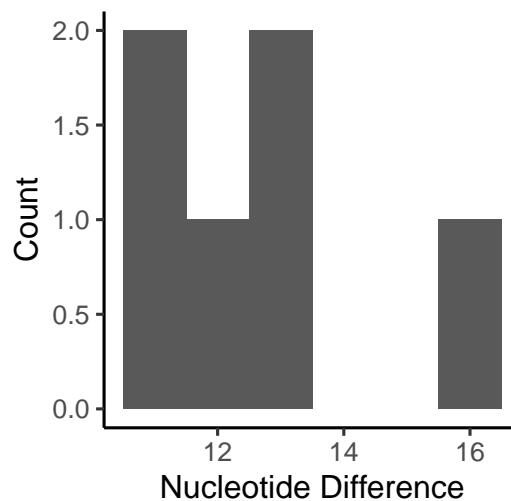
IGHV3-13*01

469 sequences assigned
25 (5.3%) exact matches, in which:
24 unique CDR3
5 unique J



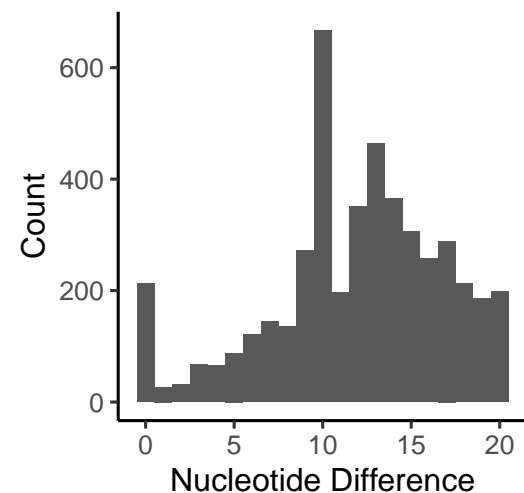
IGHV3-20*04

8 sequences assigned
No exact matches.



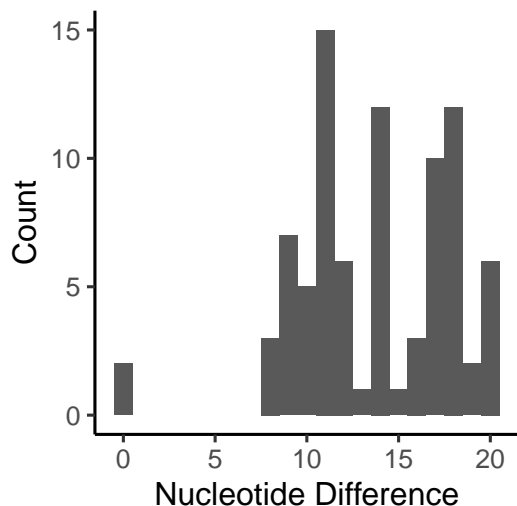
IGHV3-33*01

5870 sequences assigned
212 (3.6%) exact matches, in which:
191 unique CDR3
6 unique J



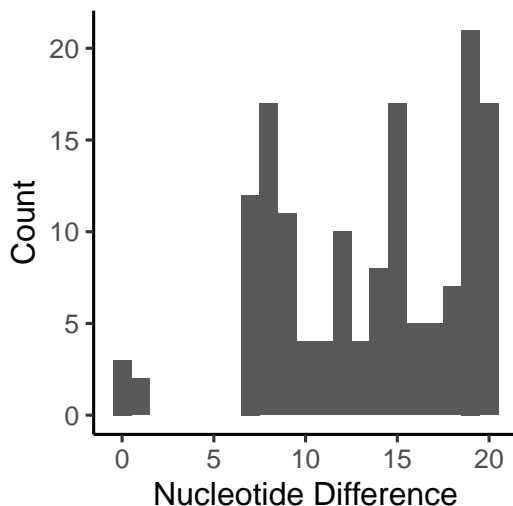
IGHV3-43*01

105 sequences assigned
2 (1.9%) exact matches, in which:
2 unique CDR3
1 unique J



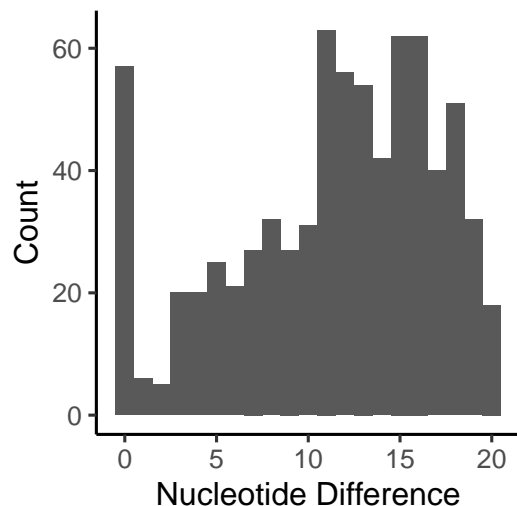
IGHV3-53*04

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



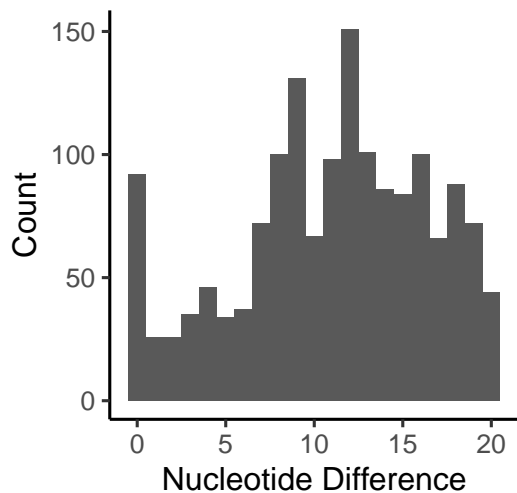
IGHV3-66*01

2119 sequences assigned
57 (2.7%) exact matches, in which:
52 unique CDR3
4 unique J



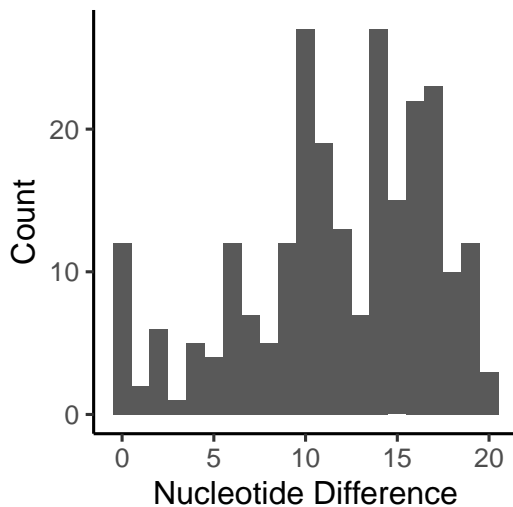
IGHV3-48*01

2032 sequences assigned
92 (4.5%) exact matches, in which:
72 unique CDR3
5 unique J



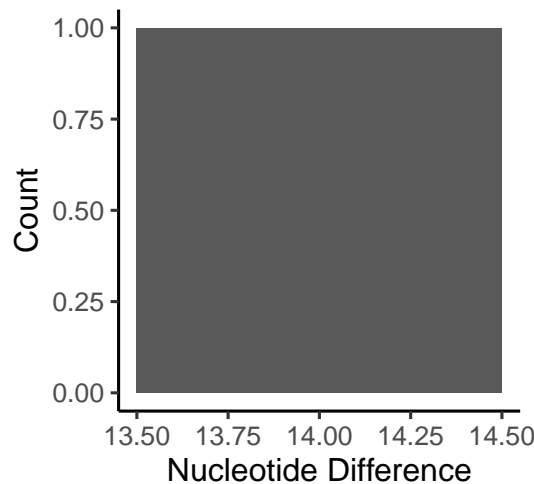
IGHV3-64*01

295 sequences assigned
12 (4.1%) exact matches, in which:
11 unique CDR3
4 unique J



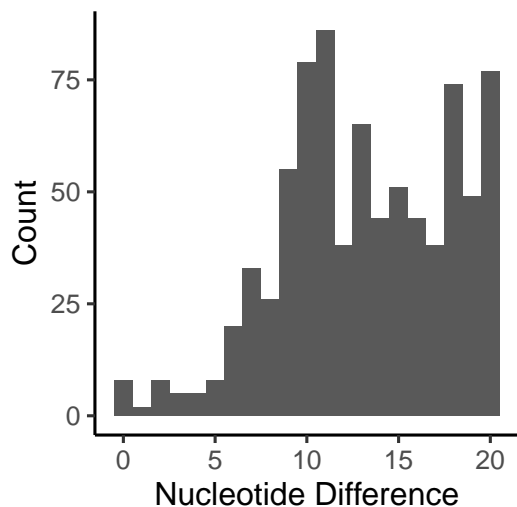
IGHV3-69-1*02

2 sequences assigned
No exact matches.



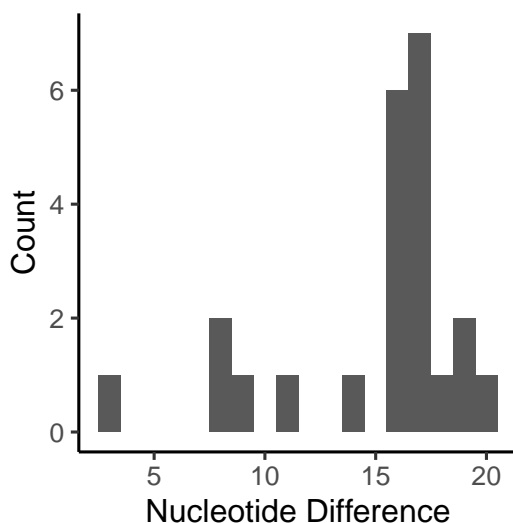
IGHV3-49*03

1421 sequences assigned
8 (0.6%) exact matches, in which:
7 unique CDR3
2 unique J



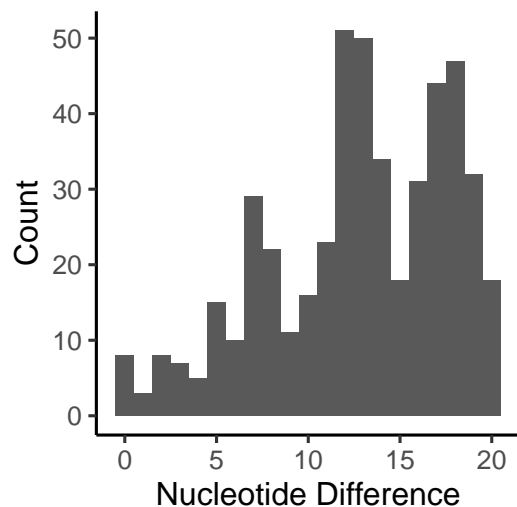
IGHV3-64*07

79 sequences assigned
No exact matches.



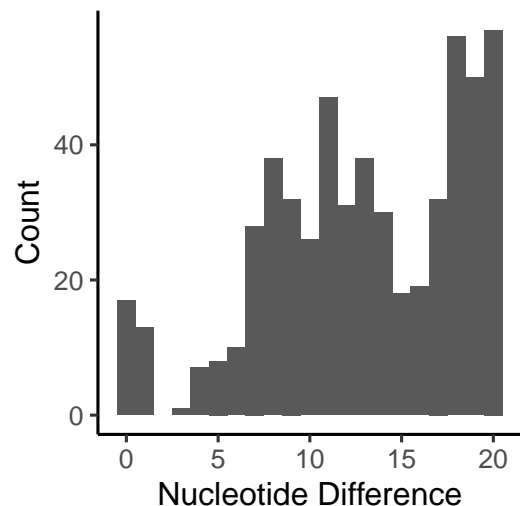
IGHV3-72*01

776 sequences assigned
8 (1%) exact matches, in which:
8 unique CDR3
2 unique J



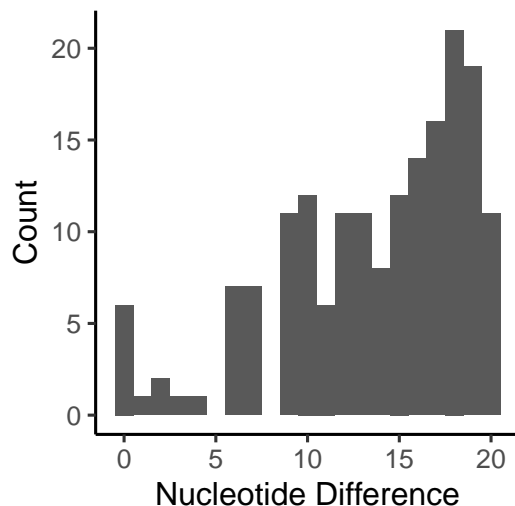
IGHV3-73*01

965 sequences assigned
17 (1.8%) exact matches, in which:
16 unique CDR3
4 unique J



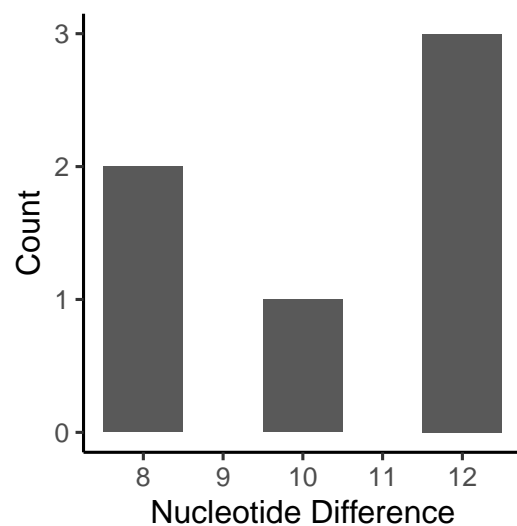
IGHV4-4*02

327 sequences assigned
6 (1.8%) exact matches, in which:
4 unique CDR3
3 unique J



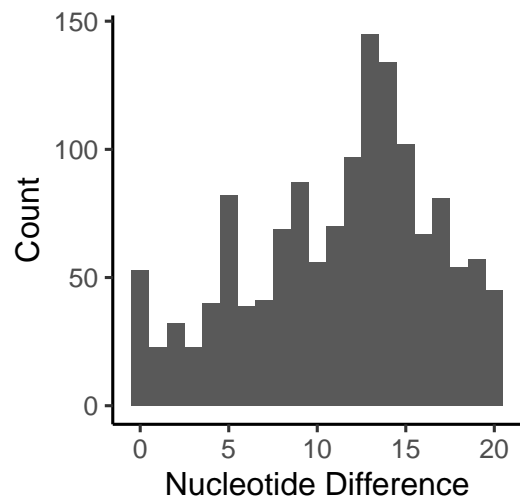
IGHV4-28*05

6 sequences assigned
No exact matches.



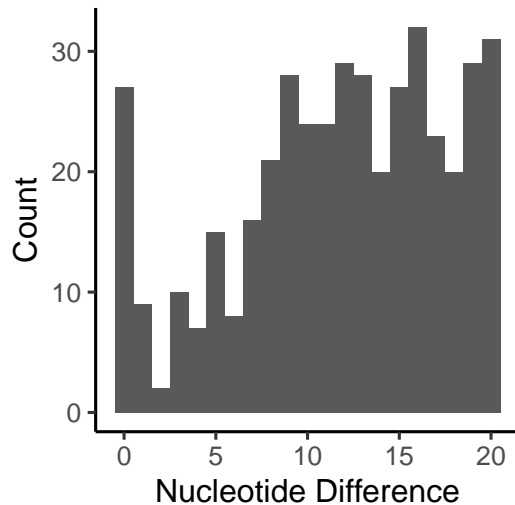
IGHV3-74*01

1667 sequences assigned
53 (3.2%) exact matches, in which:
36 unique CDR3
5 unique J



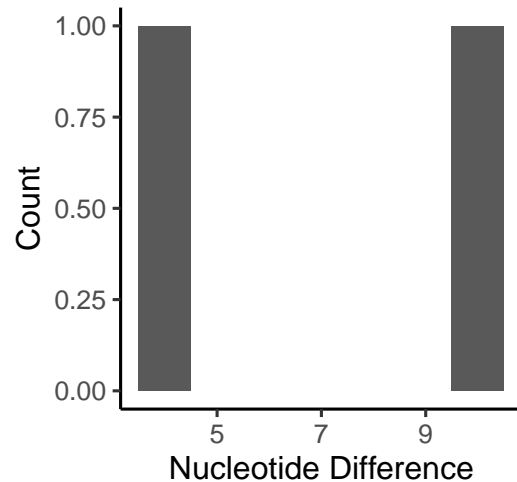
IGHV4-4*07

635 sequences assigned
27 (4.3%) exact matches, in which:
27 unique CDR3
6 unique J



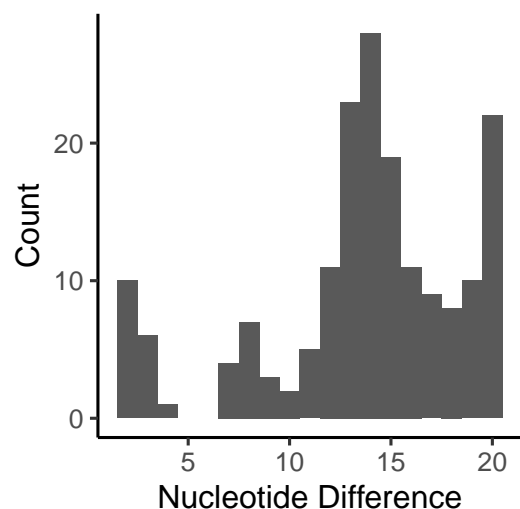
IGHV4-30-4*01

2 sequences assigned
No exact matches.



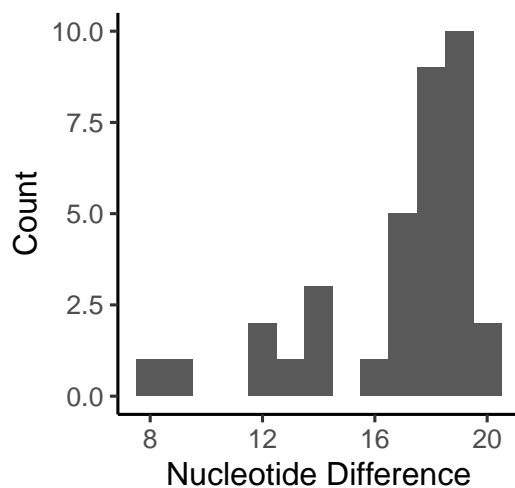
IGHV3-74*03

270 sequences assigned
No exact matches.



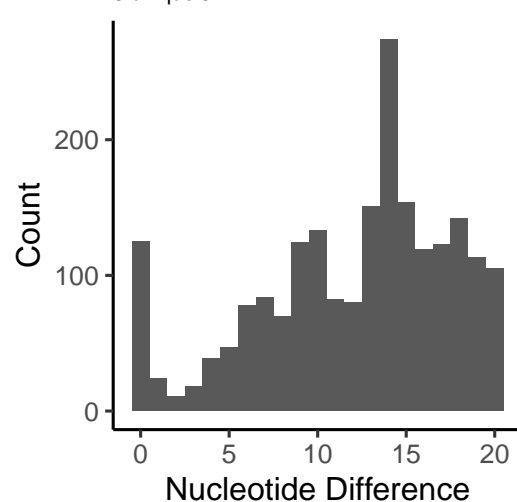
IGHV4-4*08

131 sequences assigned
No exact matches.



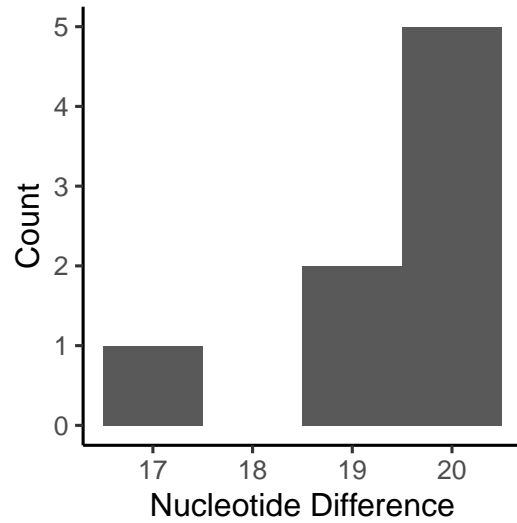
IGHV4-31*03

2782 sequences assigned
125 (4.5%) exact matches, in which:
109 unique CDR3
5 unique J



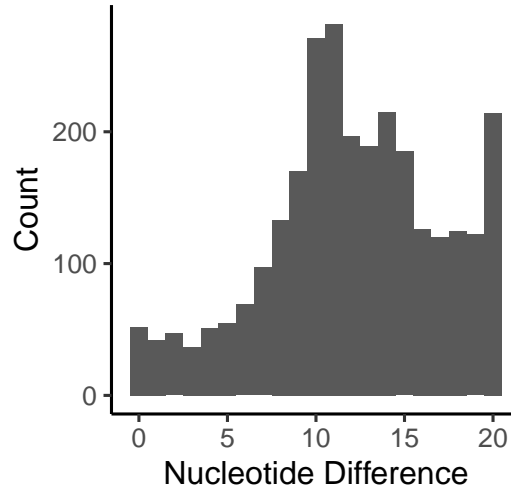
IGHV4-38-2*02

9 sequences assigned
No exact matches.



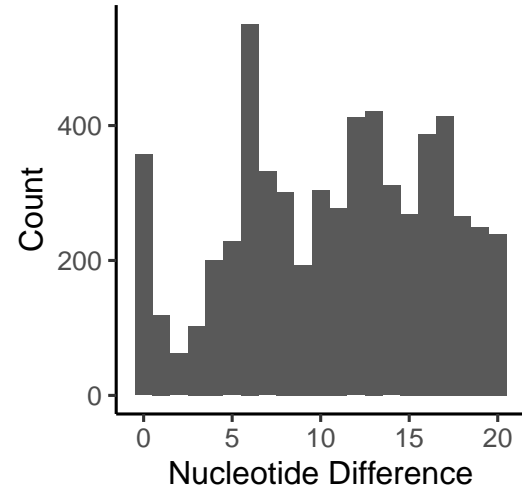
IGHV4-59*08

3915 sequences assigned
52 (1.3%) exact matches, in which:
39 unique CDR3
5 unique J



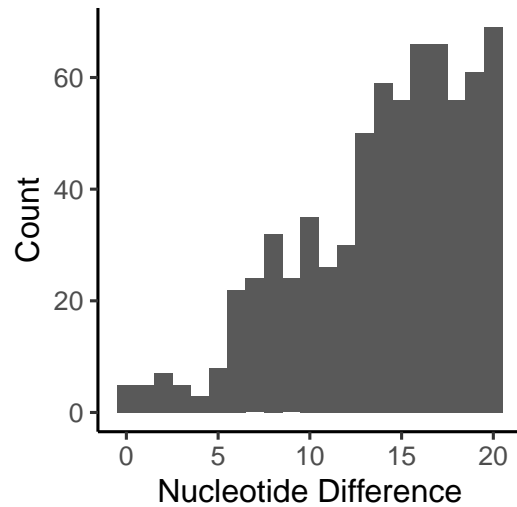
IGHV5-51*03

7324 sequences assigned
357 (4.9%) exact matches, in which:
297 unique CDR3
6 unique J



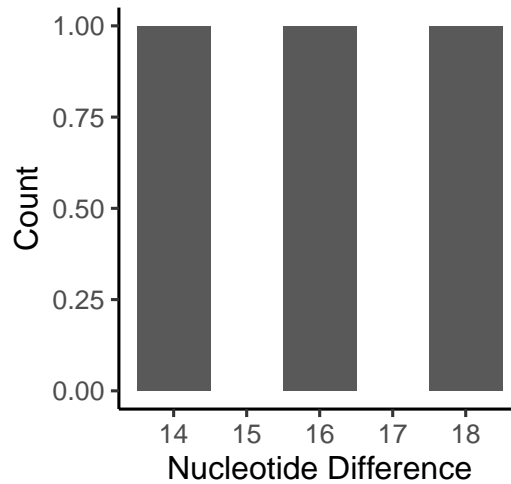
IGHV4-39*01

1591 sequences assigned
5 (0.3%) exact matches, in which:
5 unique CDR3
3 unique J



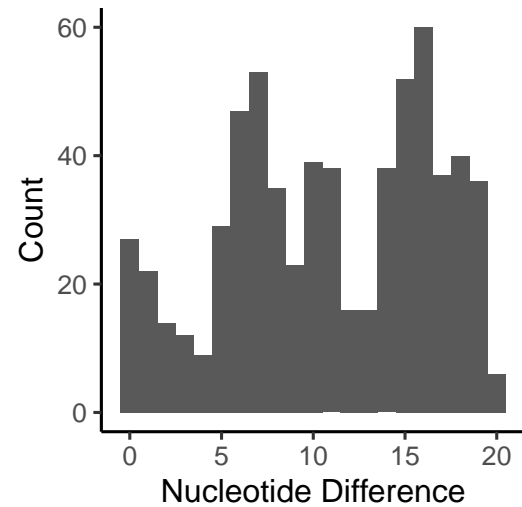
IGHV4-61*02

3 sequences assigned
No exact matches.



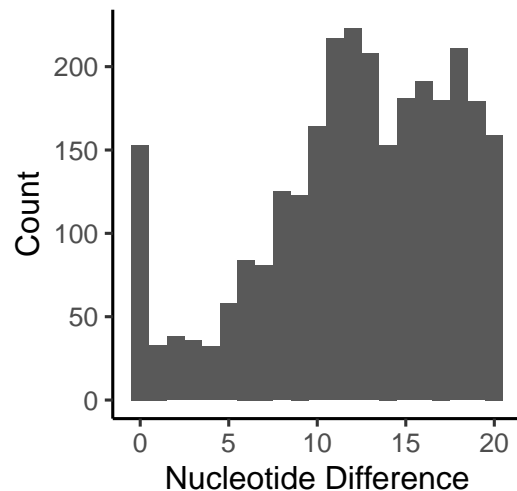
IGHV6-1*01

708 sequences assigned
27 (3.8%) exact matches, in which:
15 unique CDR3
4 unique J



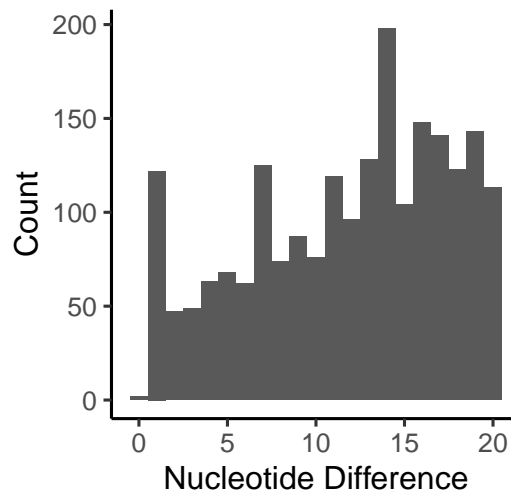
IGHV4-59*01

4020 sequences assigned
153 (3.8%) exact matches, in which:
132 unique CDR3
5 unique J



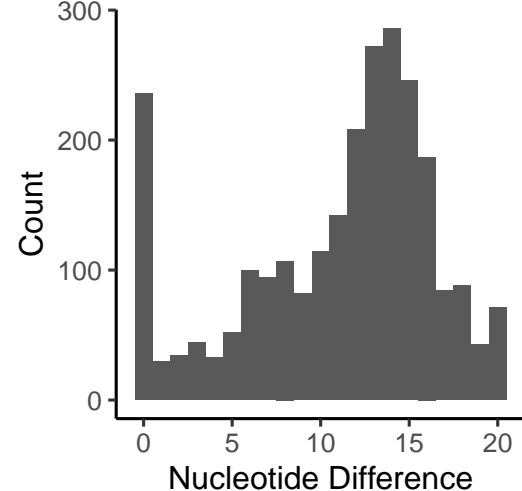
IGHV5-51*01

2732 sequences assigned
2 (0.1%) exact matches, in which:
2 unique CDR3
2 unique J

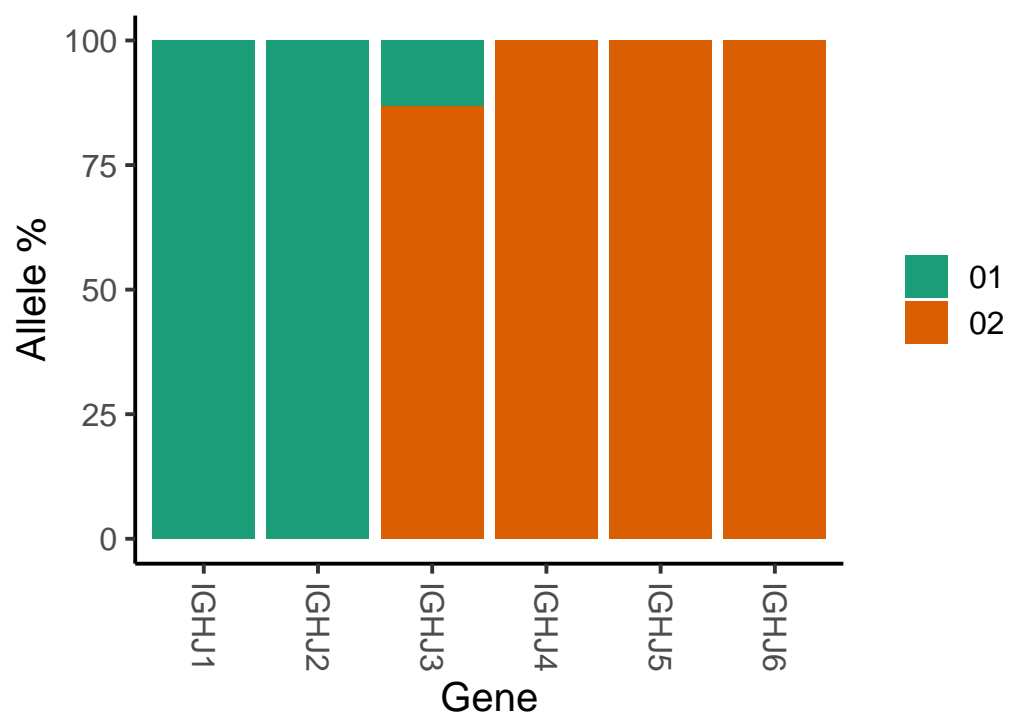


IGHV7-4-1*02

3232 sequences assigned
236 (7.3%) exact matches, in which:
211 unique CDR3
5 unique J



Allele Usage



Warning – no inferred sequences found.