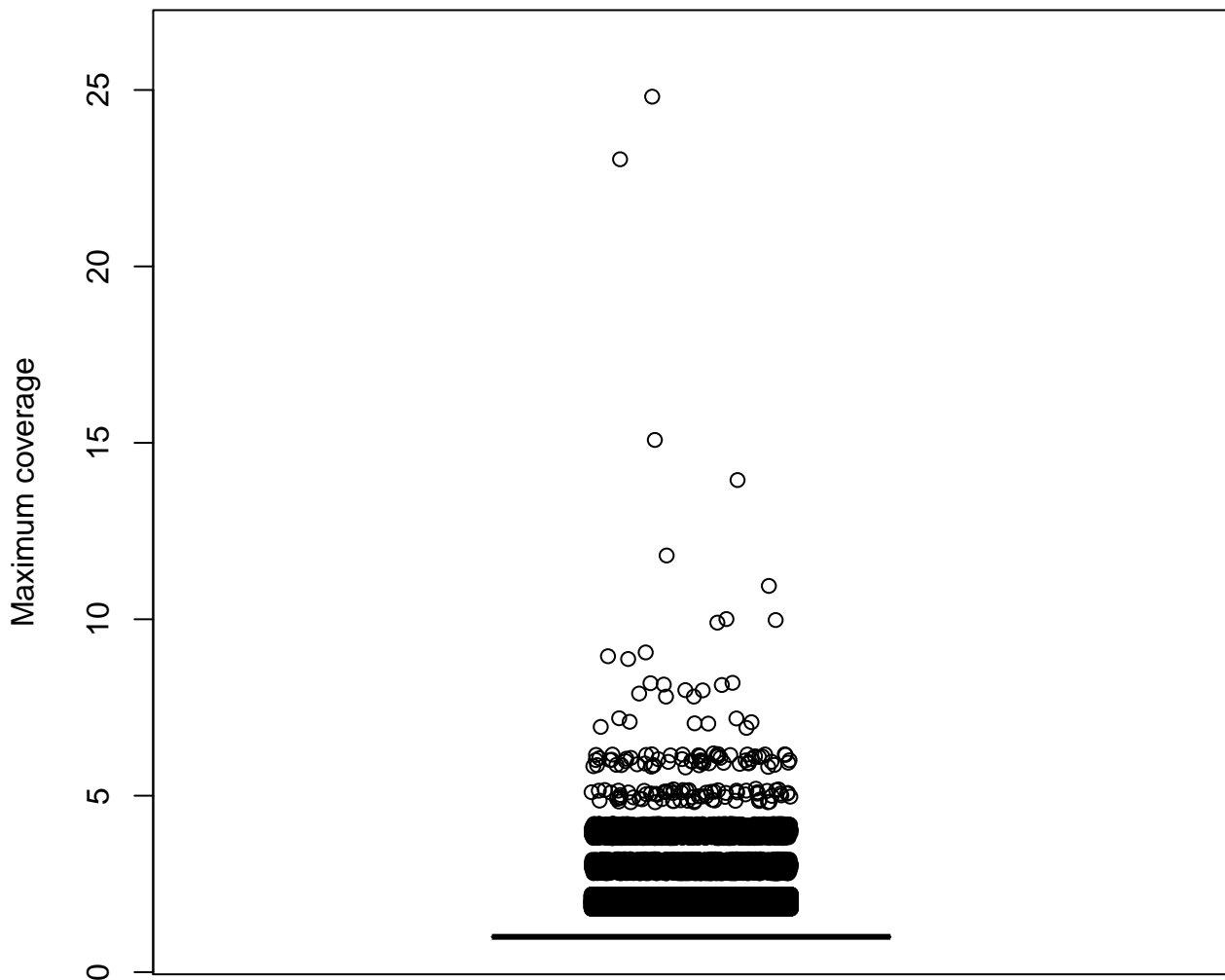
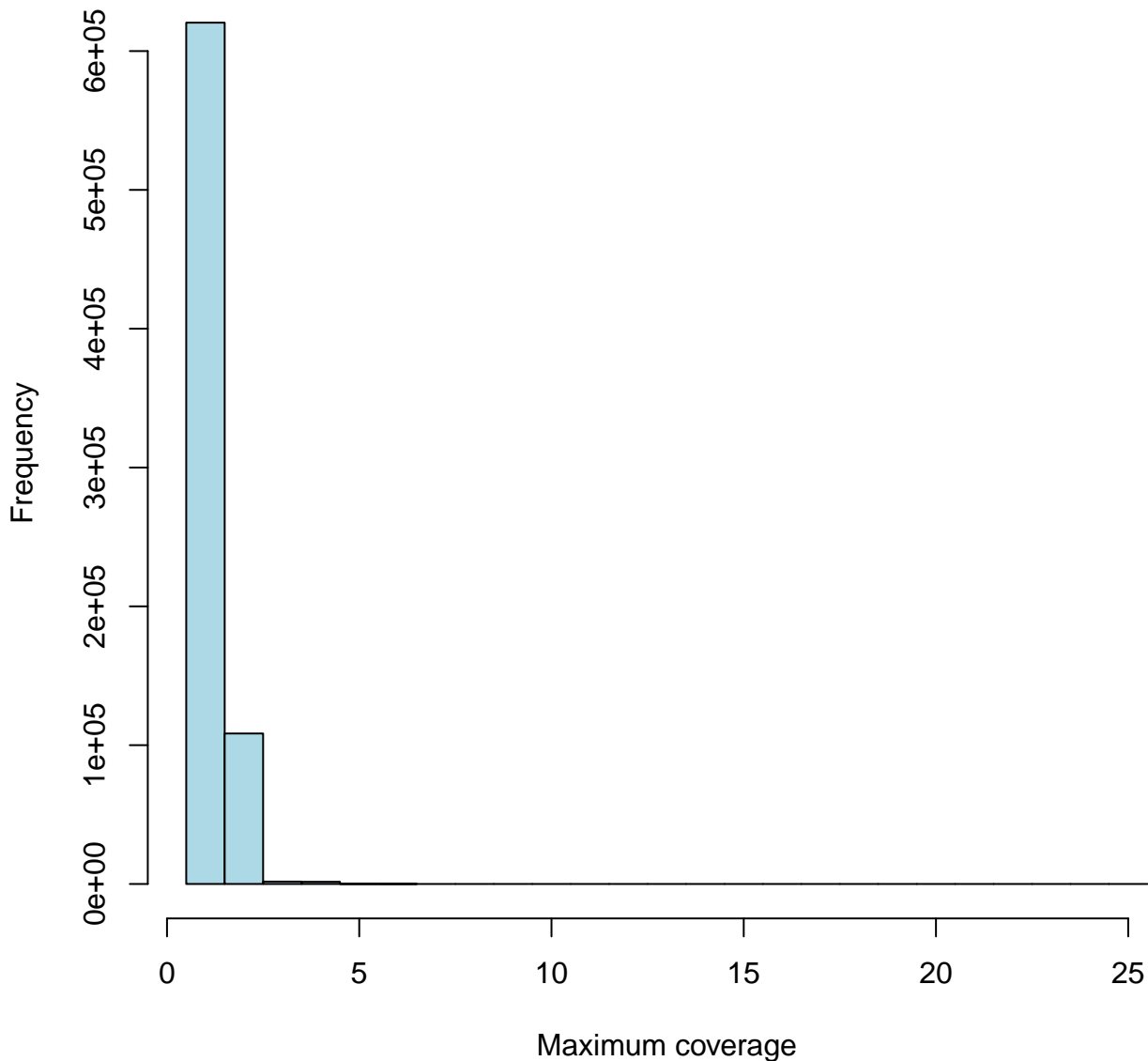


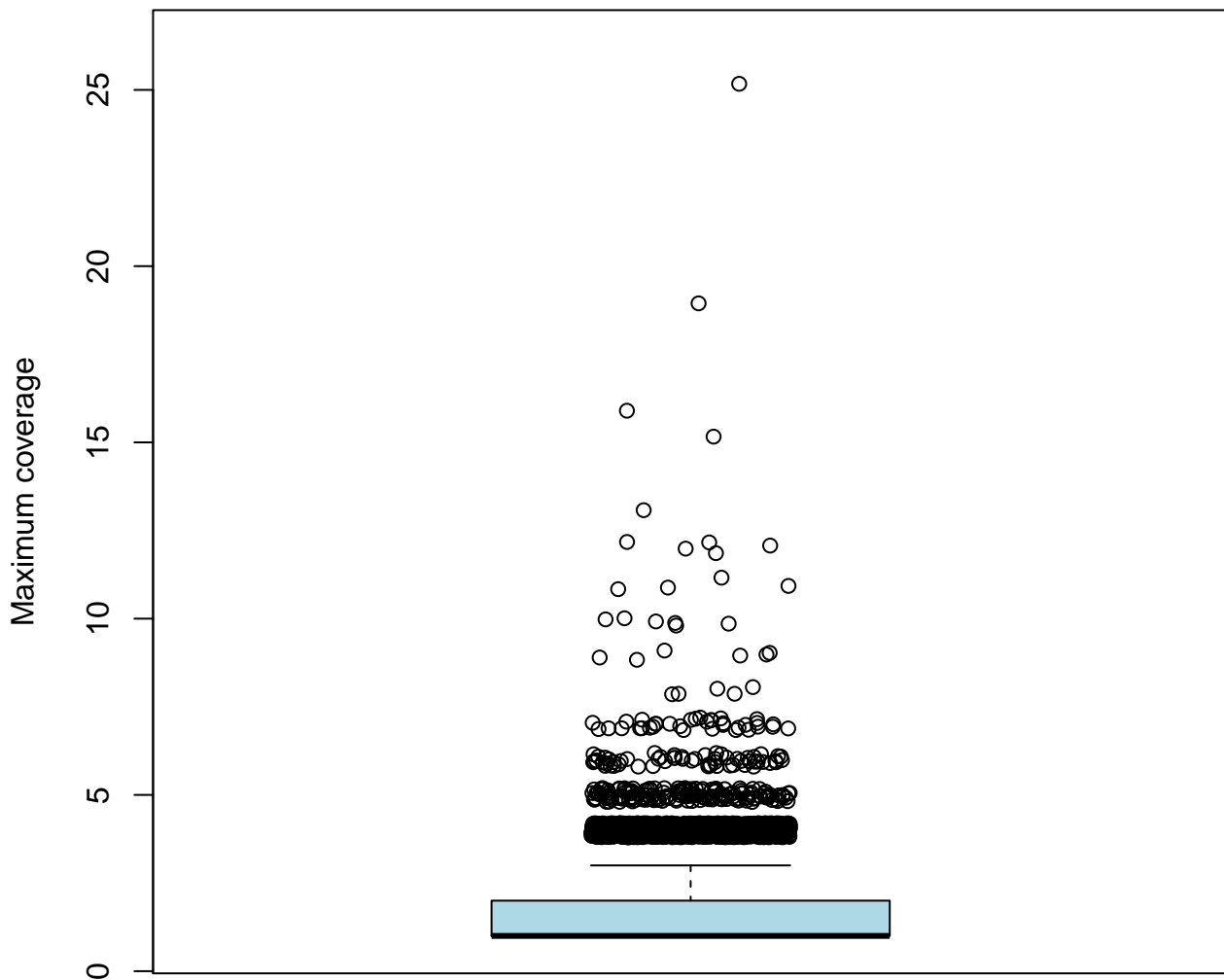
**jx in Intropolis but not "annotated" by knownGene:
1 samples; n jx: 732314**



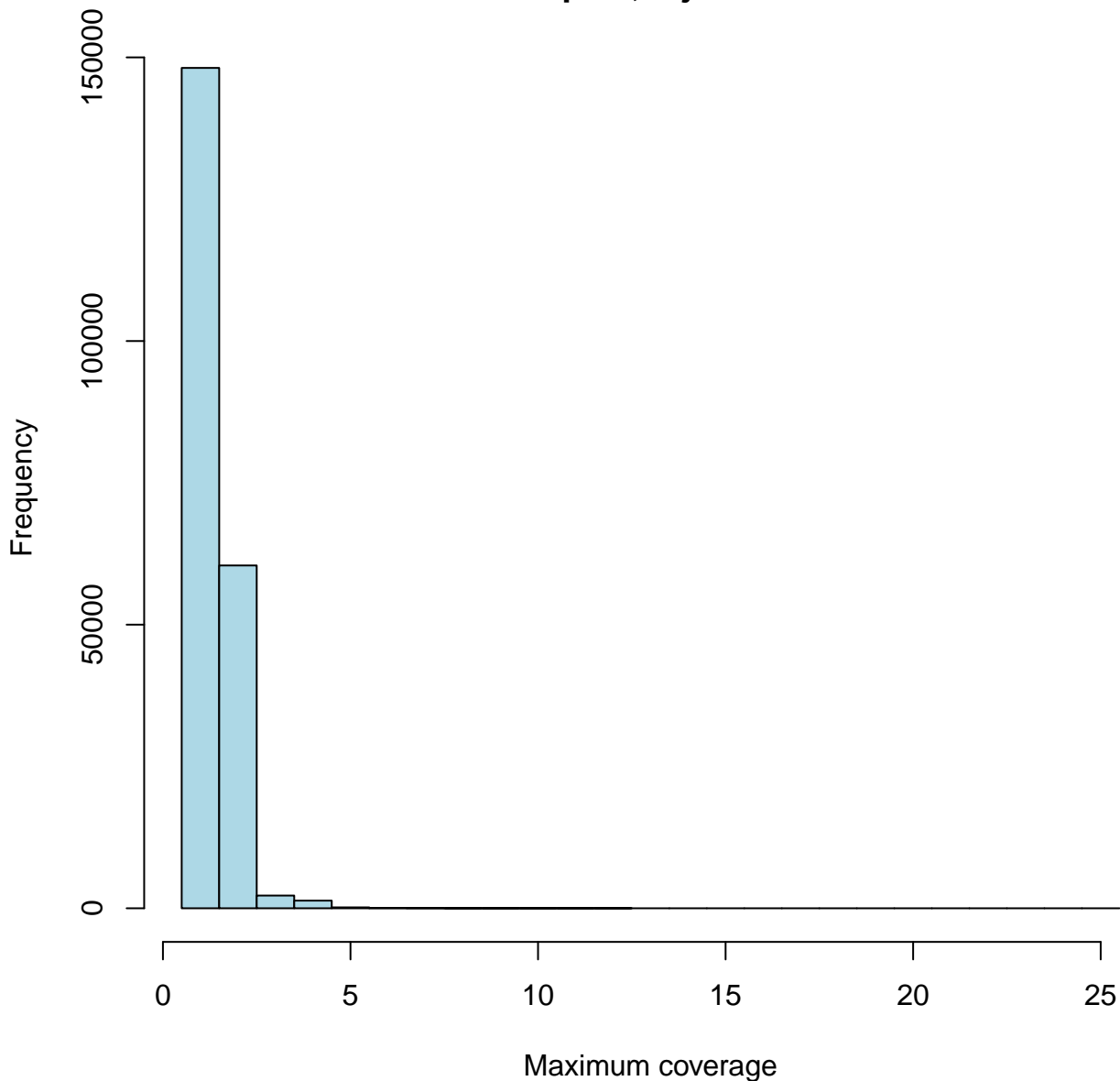
**jx in Intropolis but not "annotated" by knownGene:
1 samples; n jx: 732314**



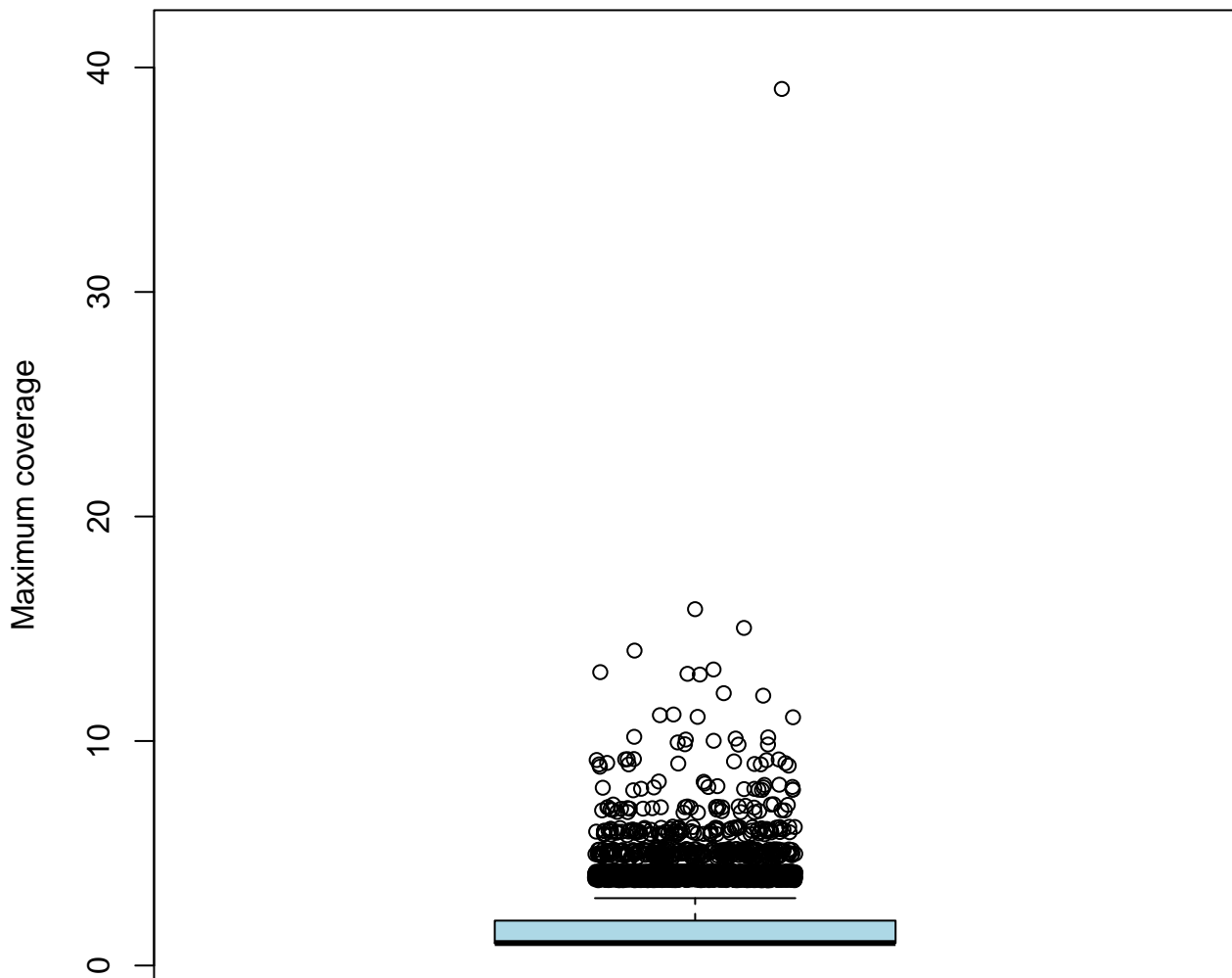
**jx in Intropolis but not "annotated" by knownGene:
2 samples; n jx: 212456**



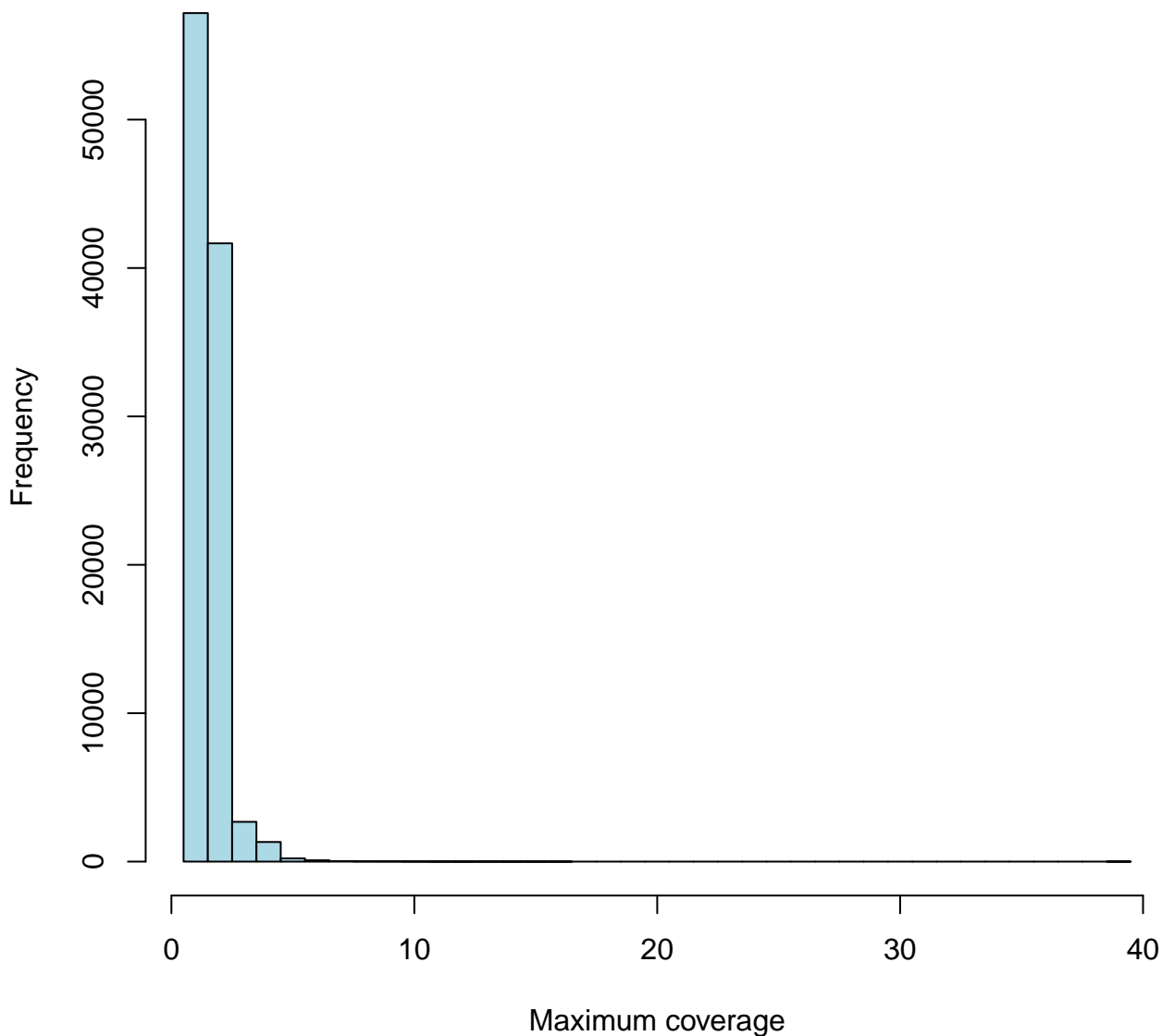
**jx in Intropolis but not "annotated" by knownGene:
2 samples; n jx: 212456**



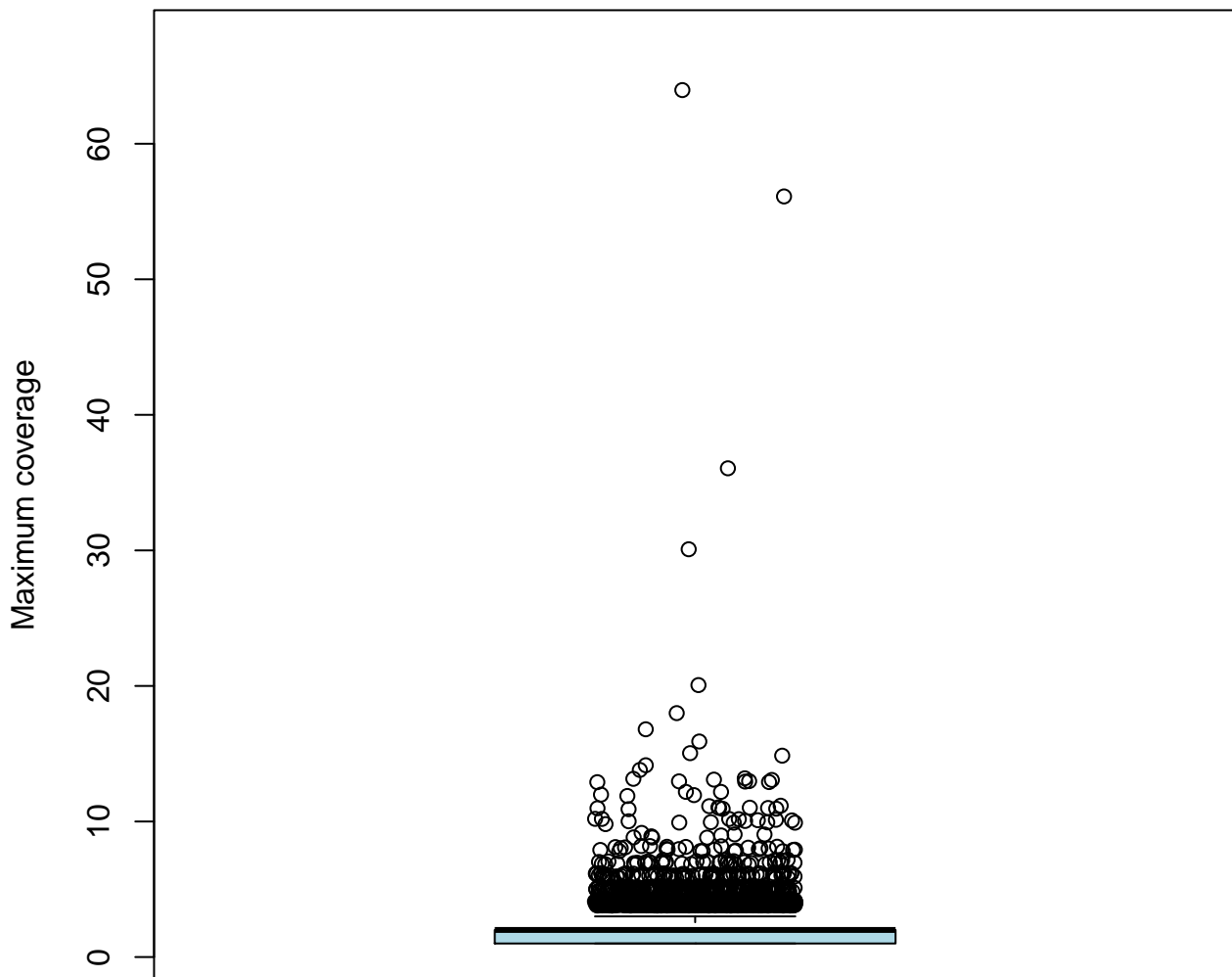
**jx in Intropolis but not "annotated" by knownGene:
3 samples; n jx: 103243**



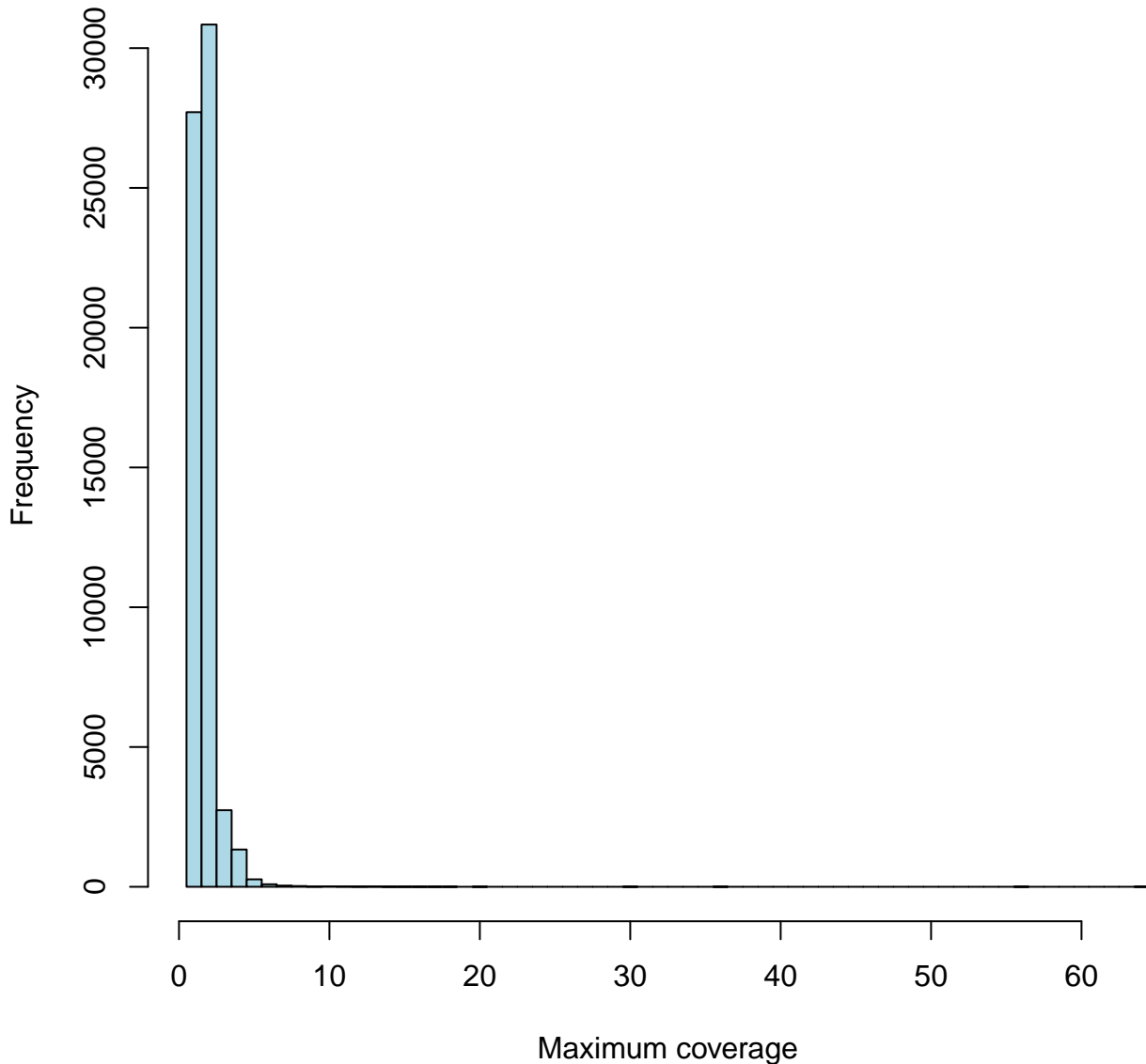
**jx in Intropolis but not "annotated" by knownGene:
3 samples; n jx: 103243**



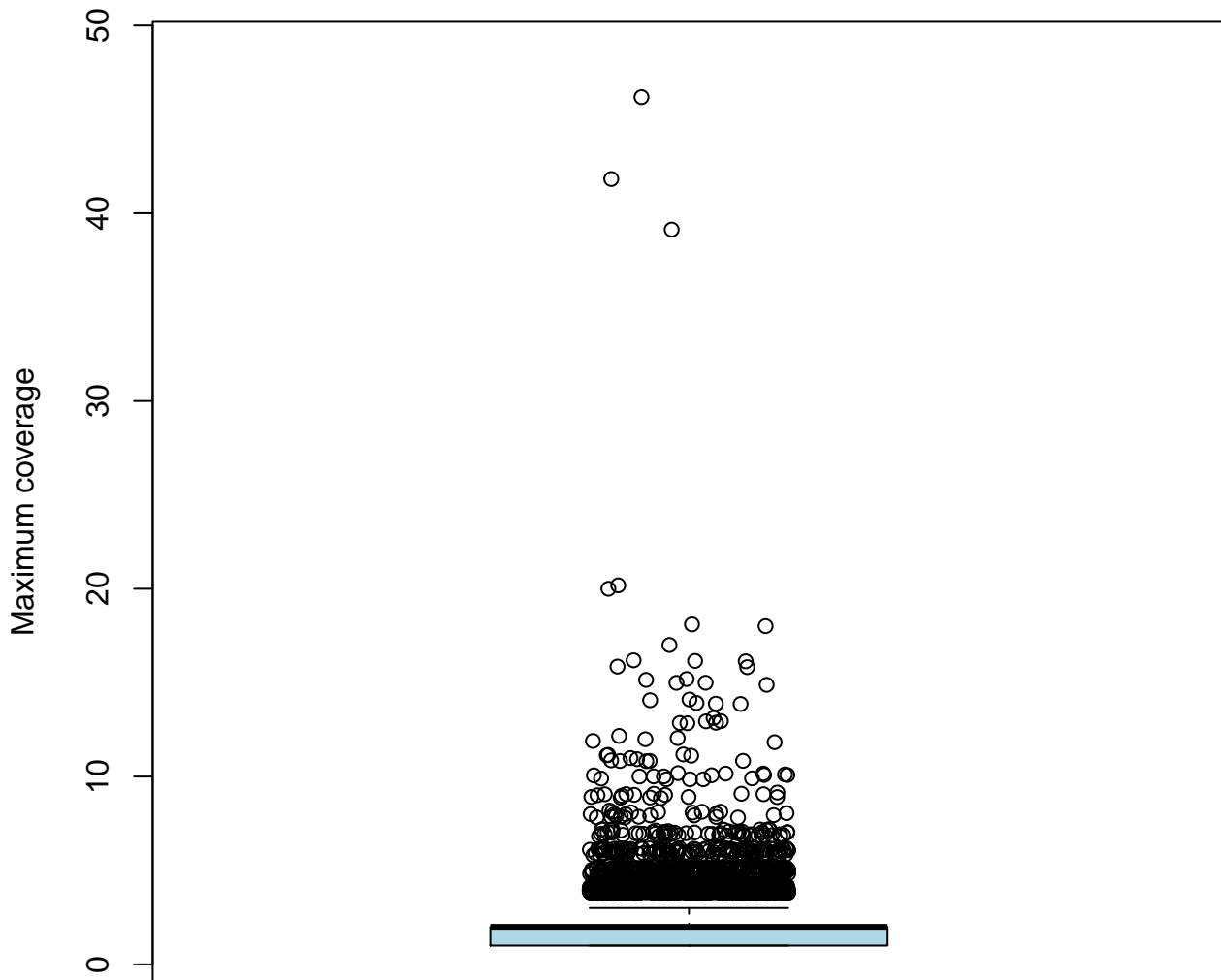
**jx in Intropolis but not "annotated" by knownGene:
4 samples; n jx: 63109**



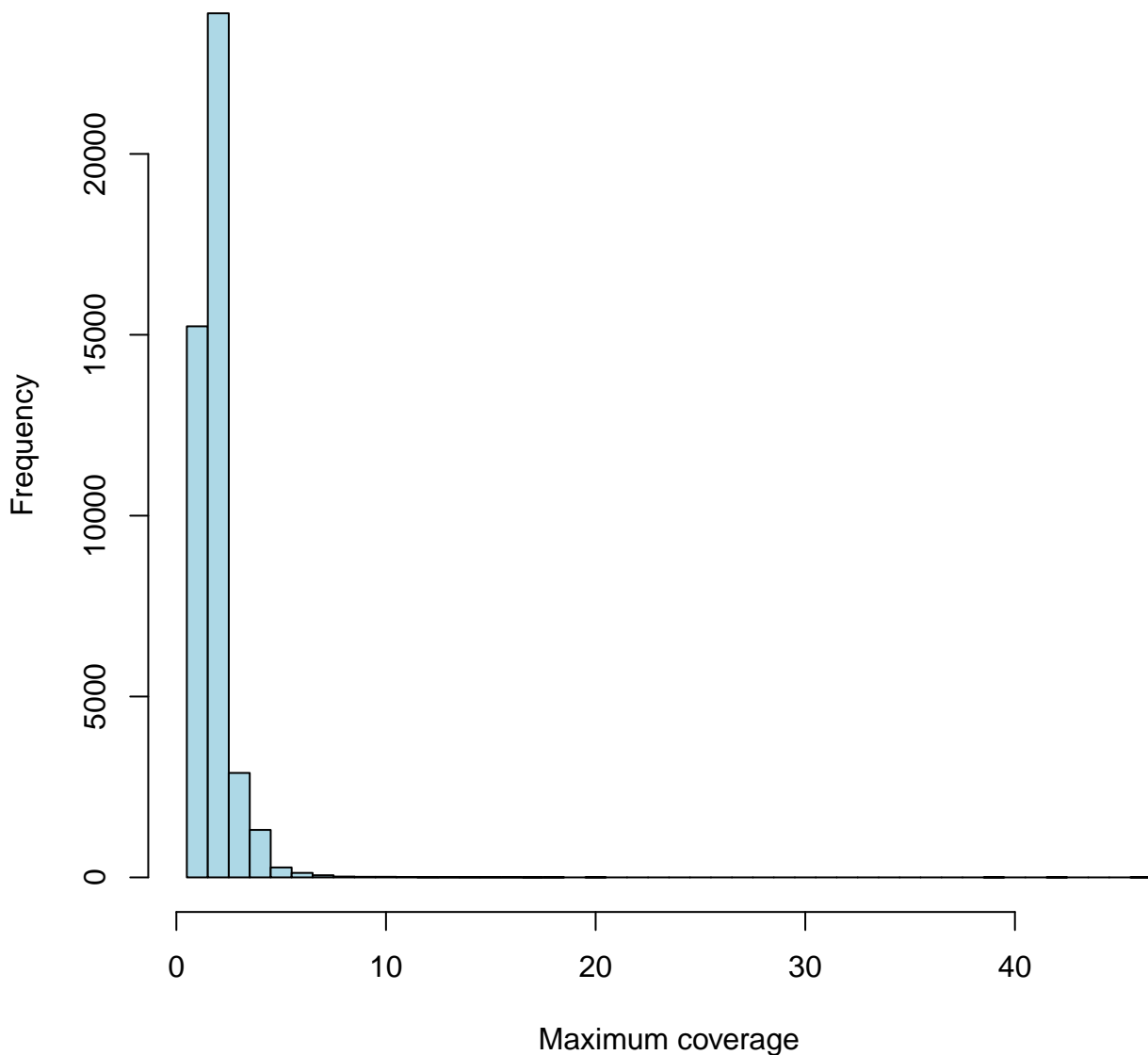
**jx in Intropolis but not "annotated" by knownGene:
4 samples; n jx: 63109**



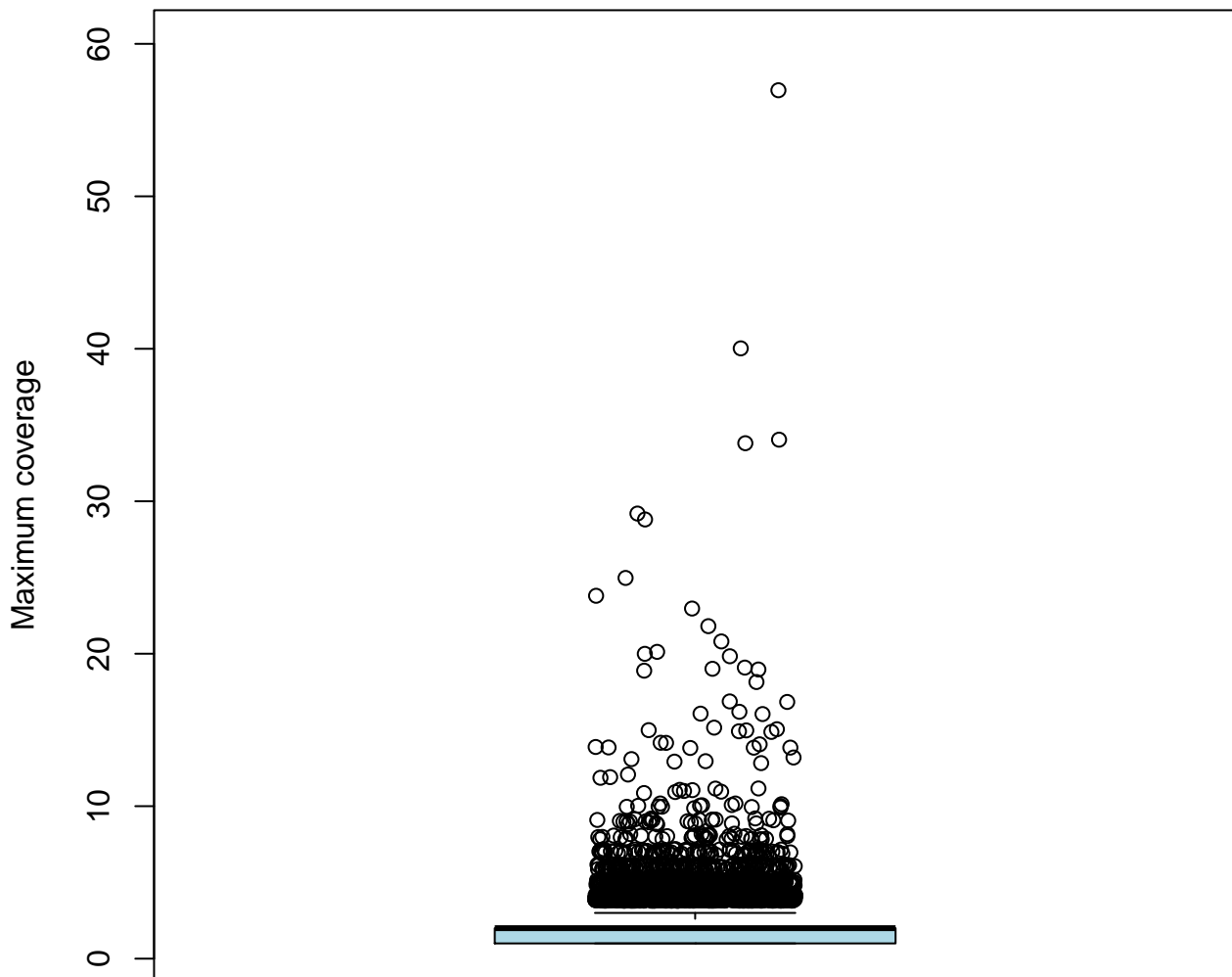
**jx in Intropolis but not "annotated" by knownGene:
5 samples; n jx: 43882**



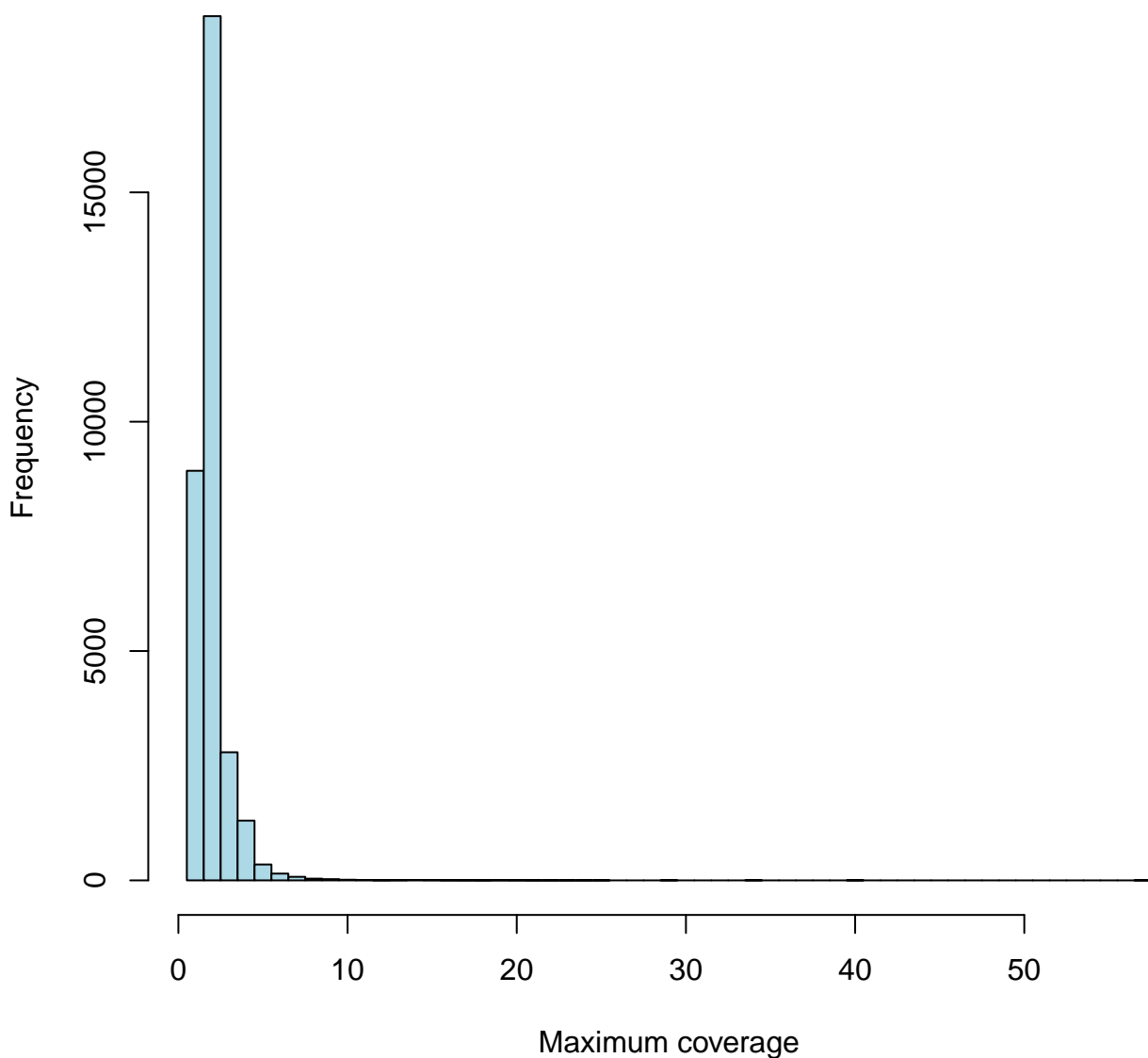
**jx in Intropolis but not "annotated" by knownGene:
5 samples; n jx: 43882**



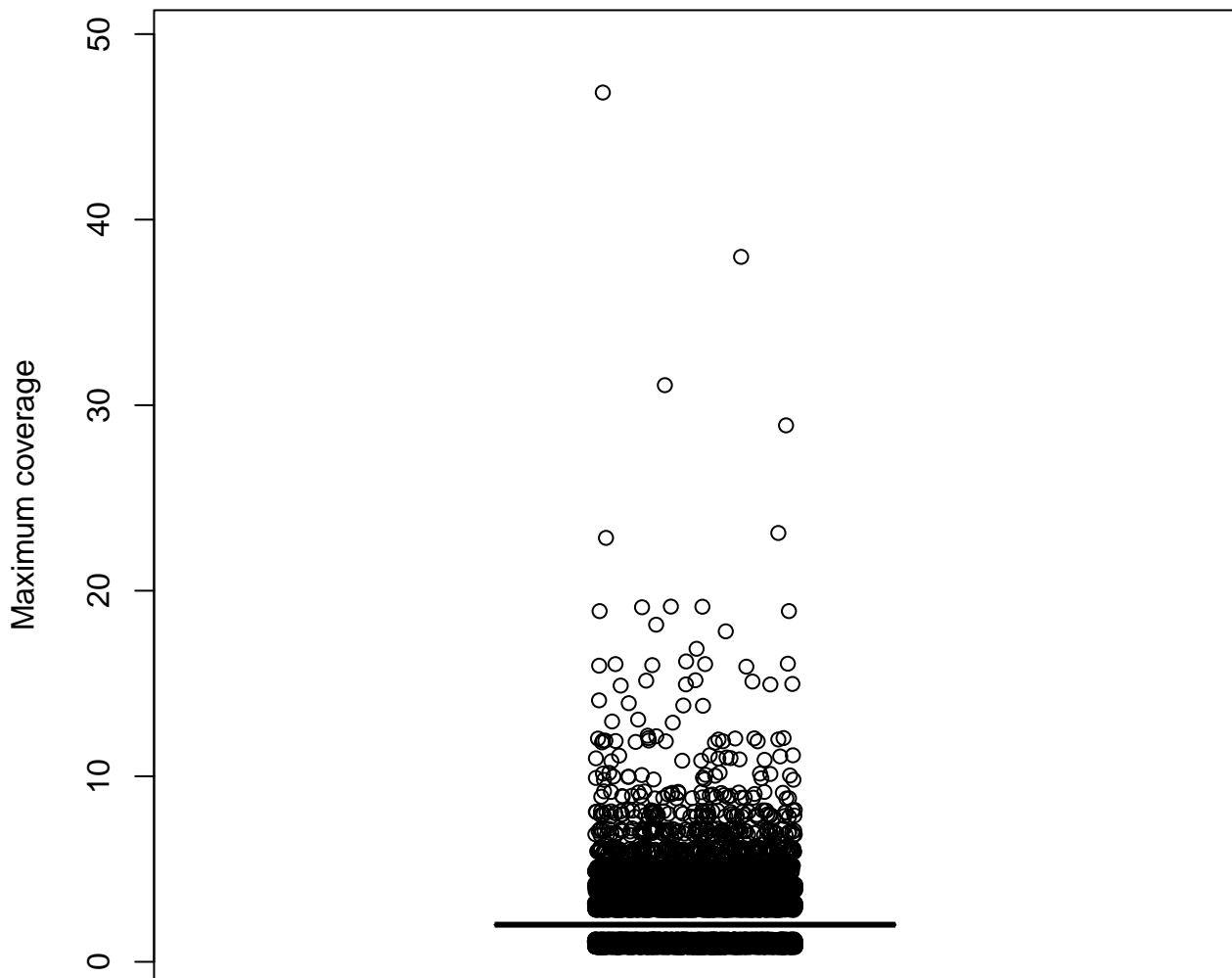
**jx in Intropolis but not "annotated" by knownGene:
6 samples; n jx: 32572**



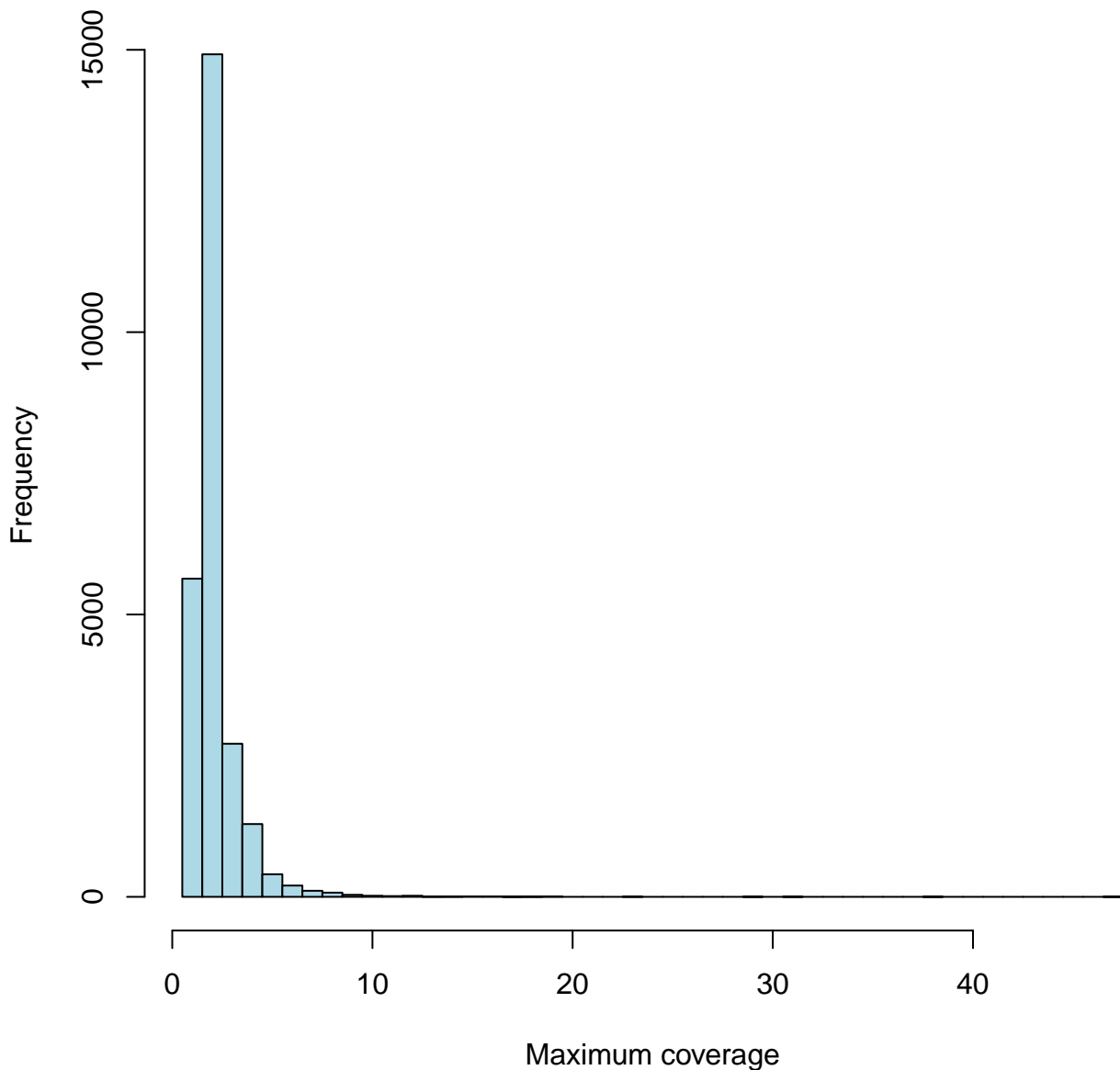
**jx in Intropolis but not "annotated" by knownGene:
6 samples; n jx: 32572**



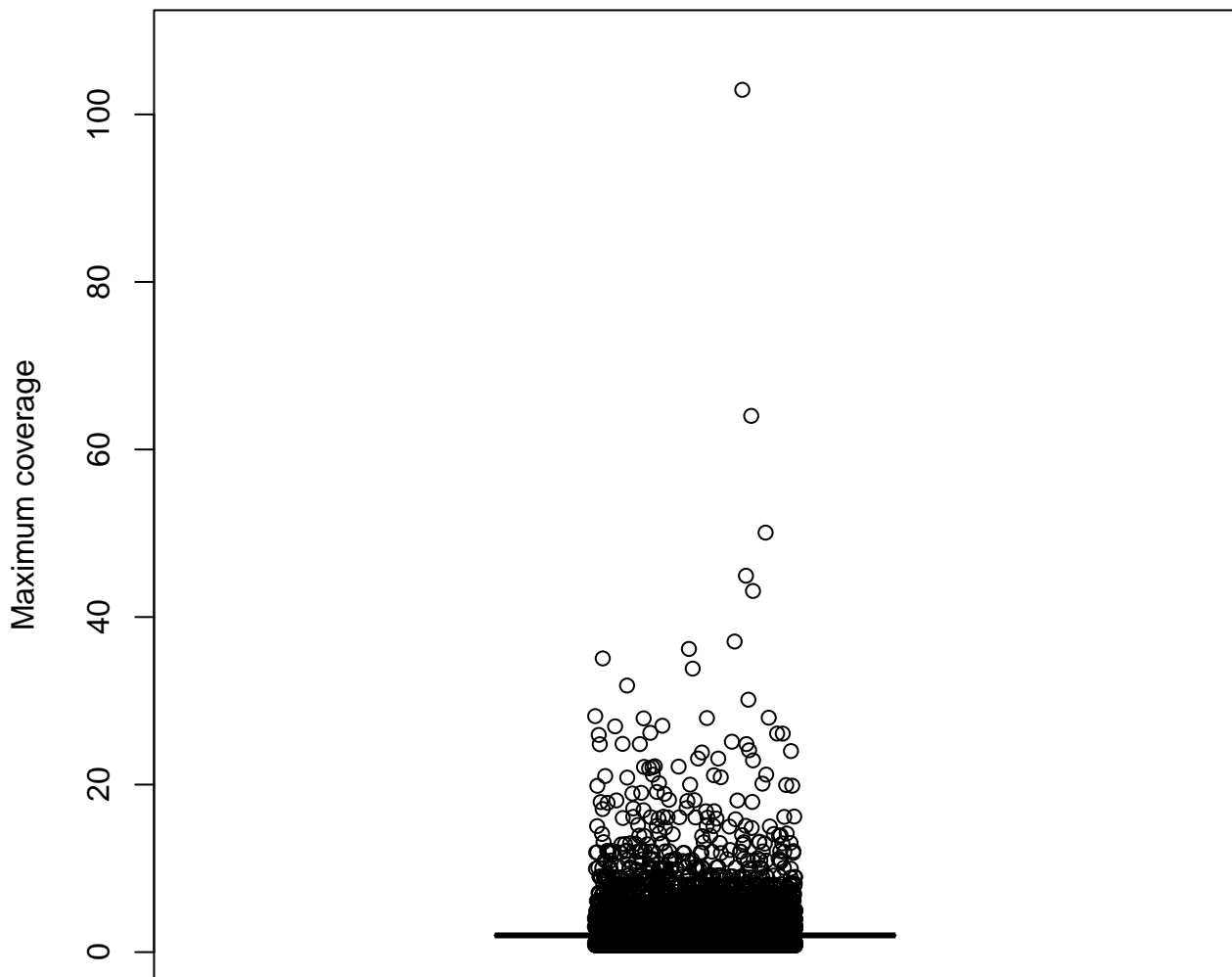
**jx in Intropolis but not "annotated" by knownGene:
7 samples; n jx: 25457**



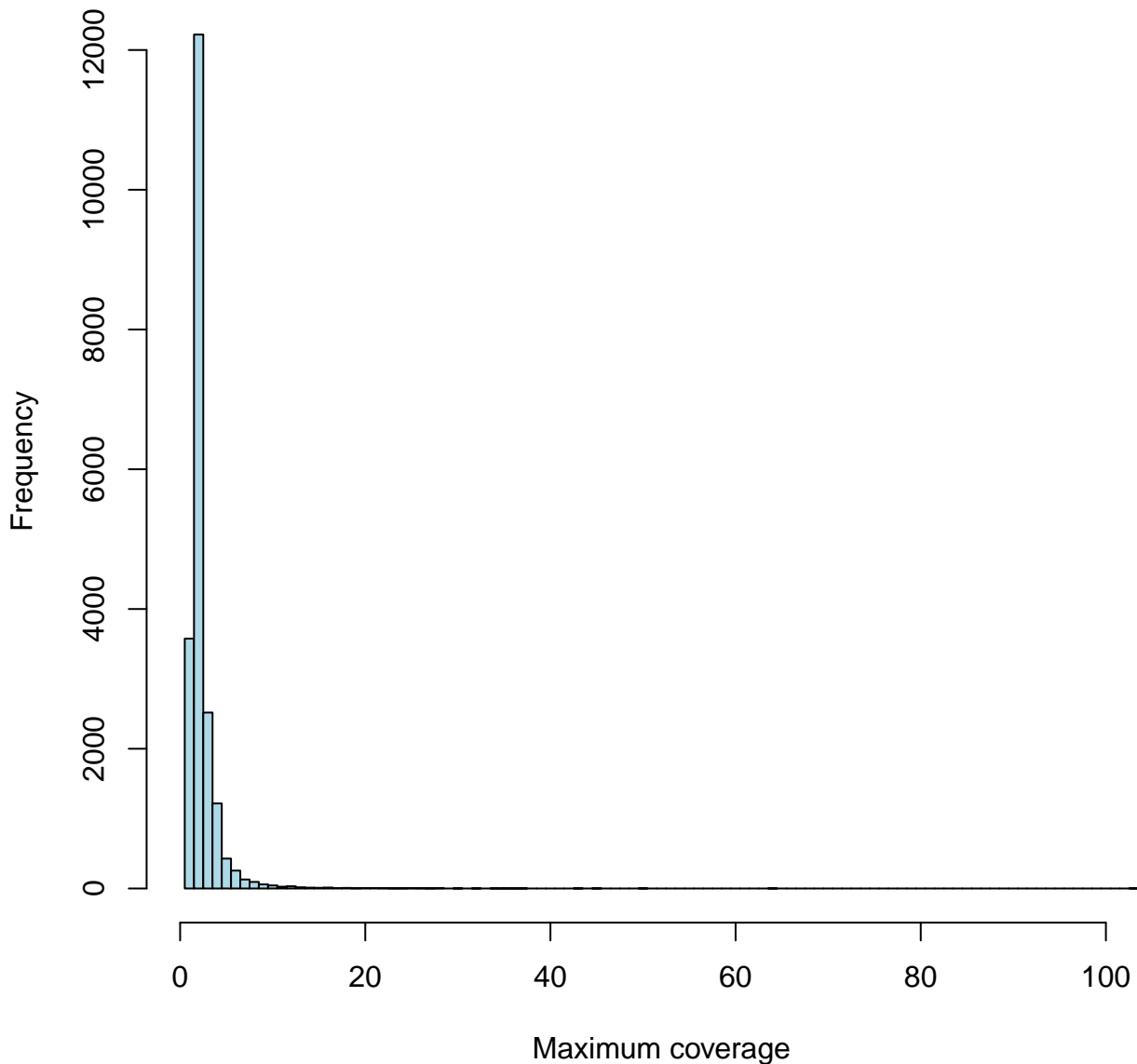
**jx in Intropolis but not "annotated" by knownGene:
7 samples; n jx: 25457**



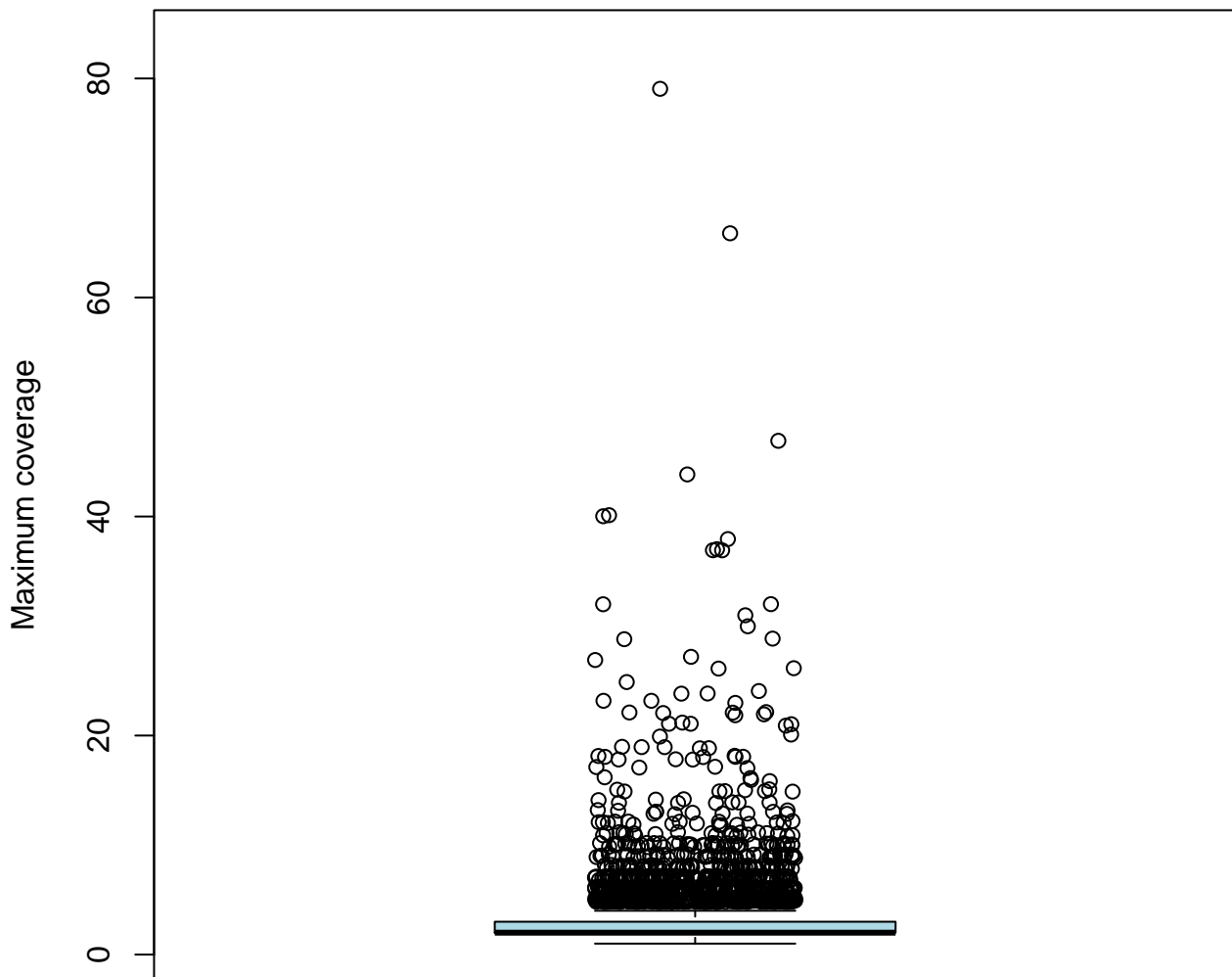
**jx in Intropolis but not "annotated" by knownGene:
8 samples; n jx: 20719**



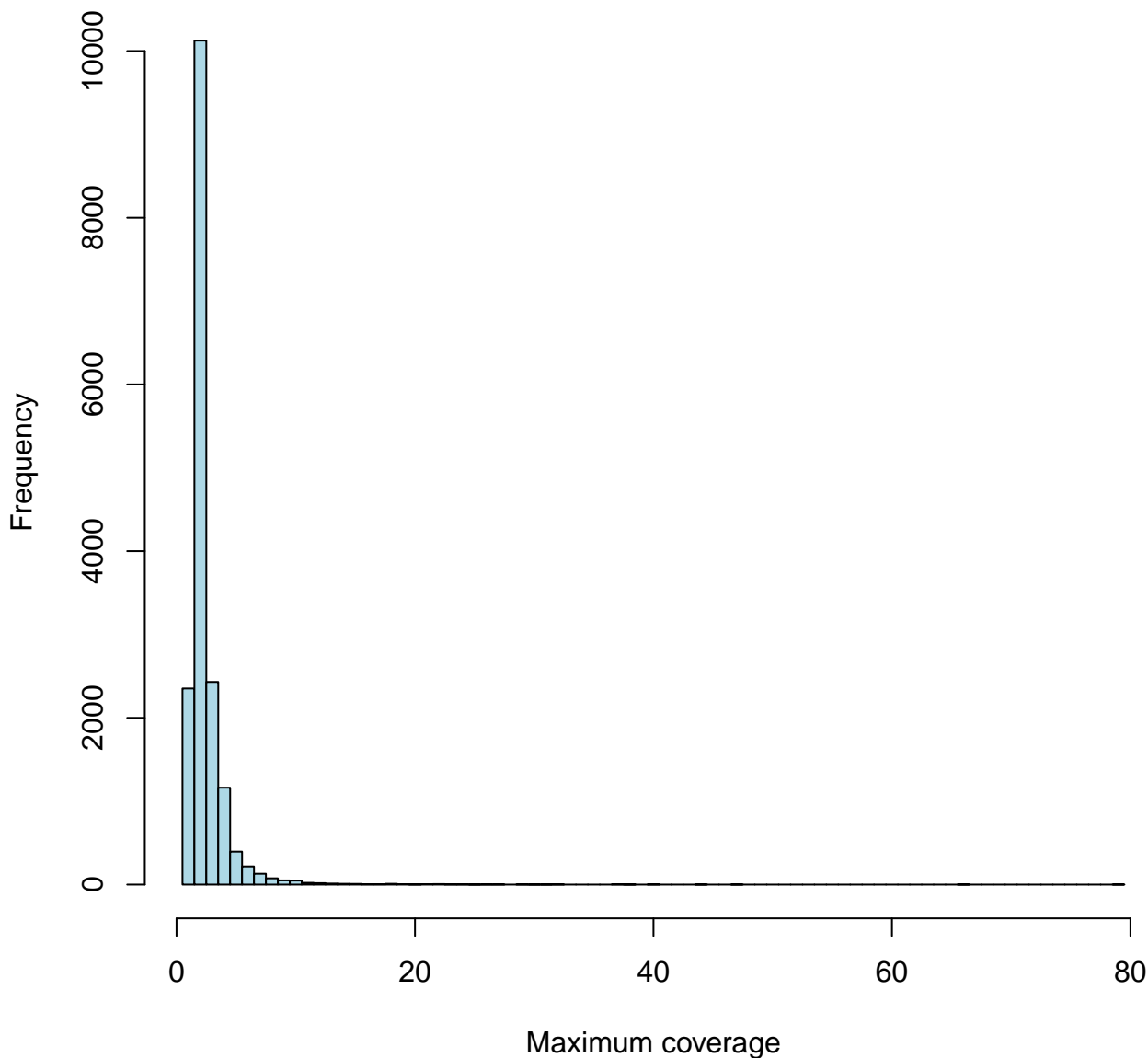
**jx in Intropolis but not "annotated" by knownGene:
8 samples; n jx: 20719**



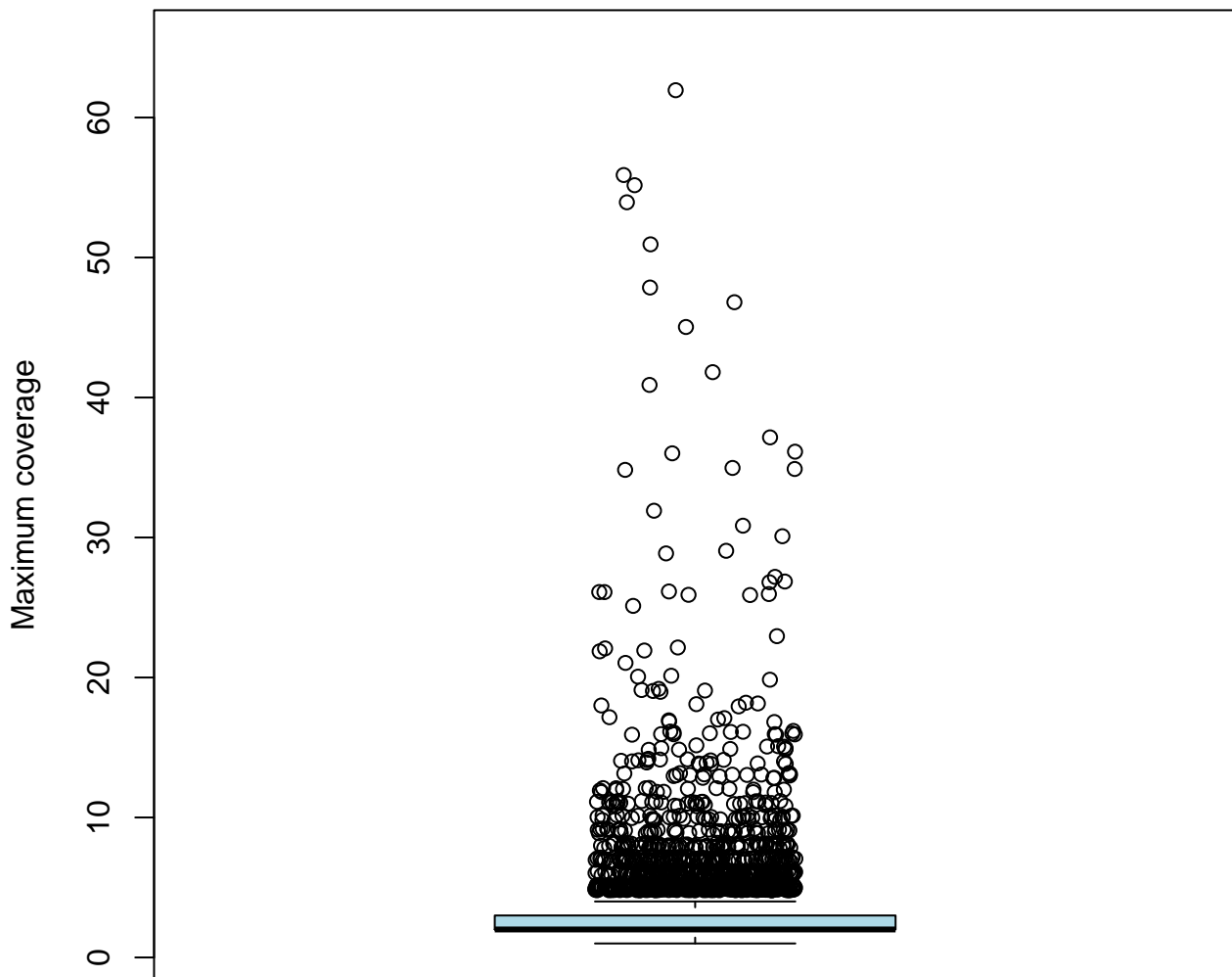
**jx in Intropolis but not "annotated" by knownGene:
9 samples; n jx: 17115**



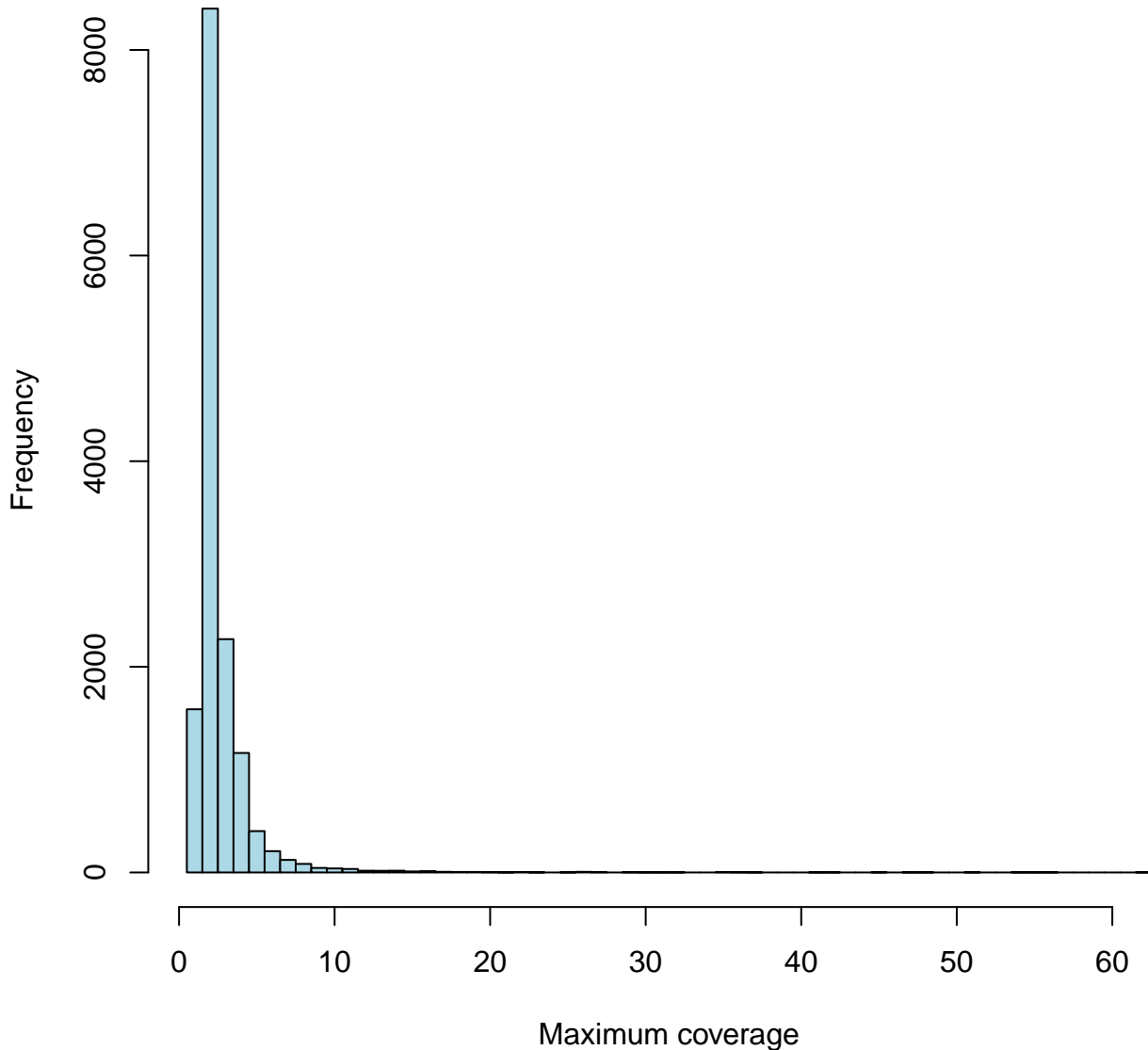
**jx in Intropolis but not "annotated" by knownGene:
9 samples; n jx: 17115**



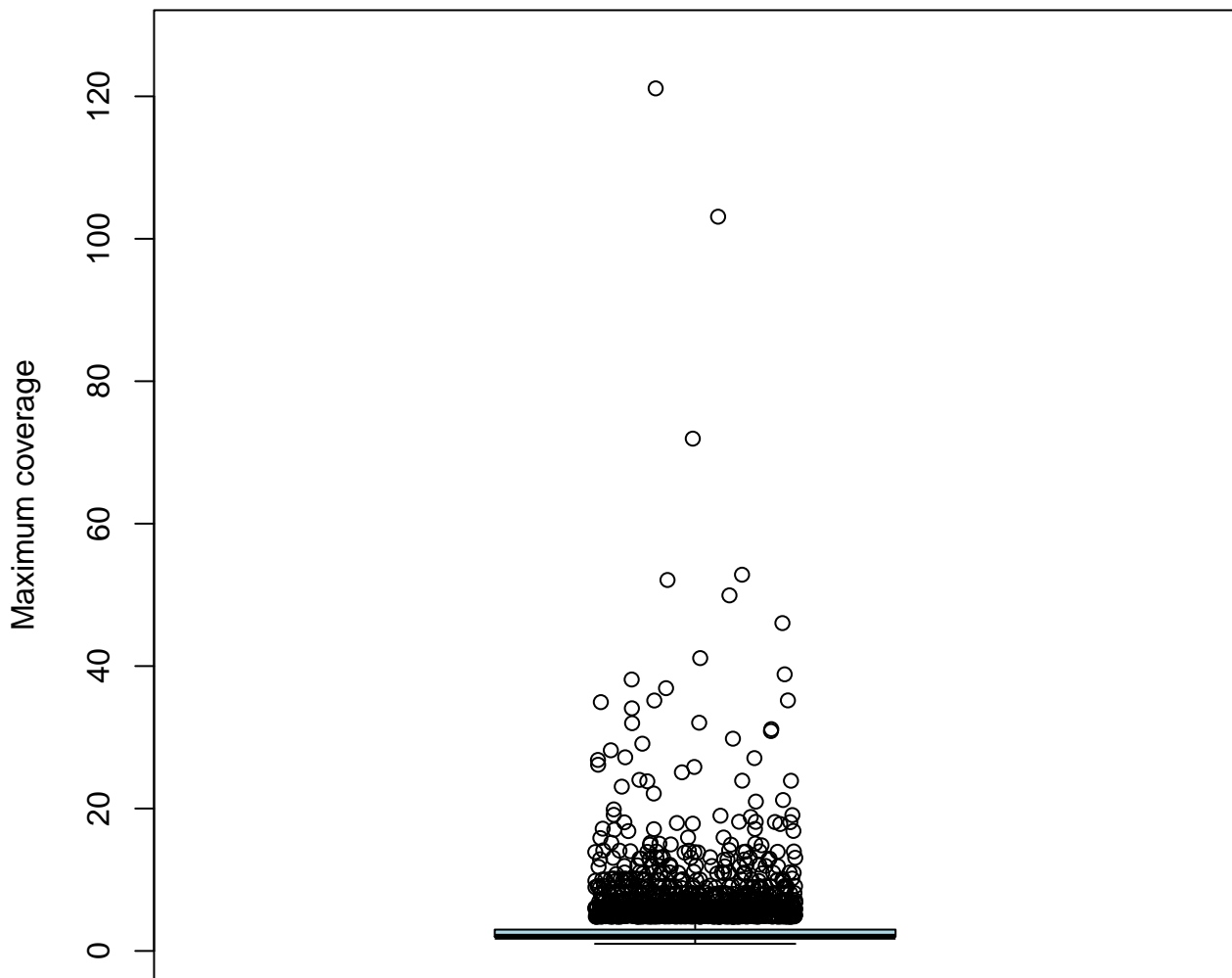
**jx in Intropolis but not "annotated" by knownGene:
10 samples; n jx: 14480**



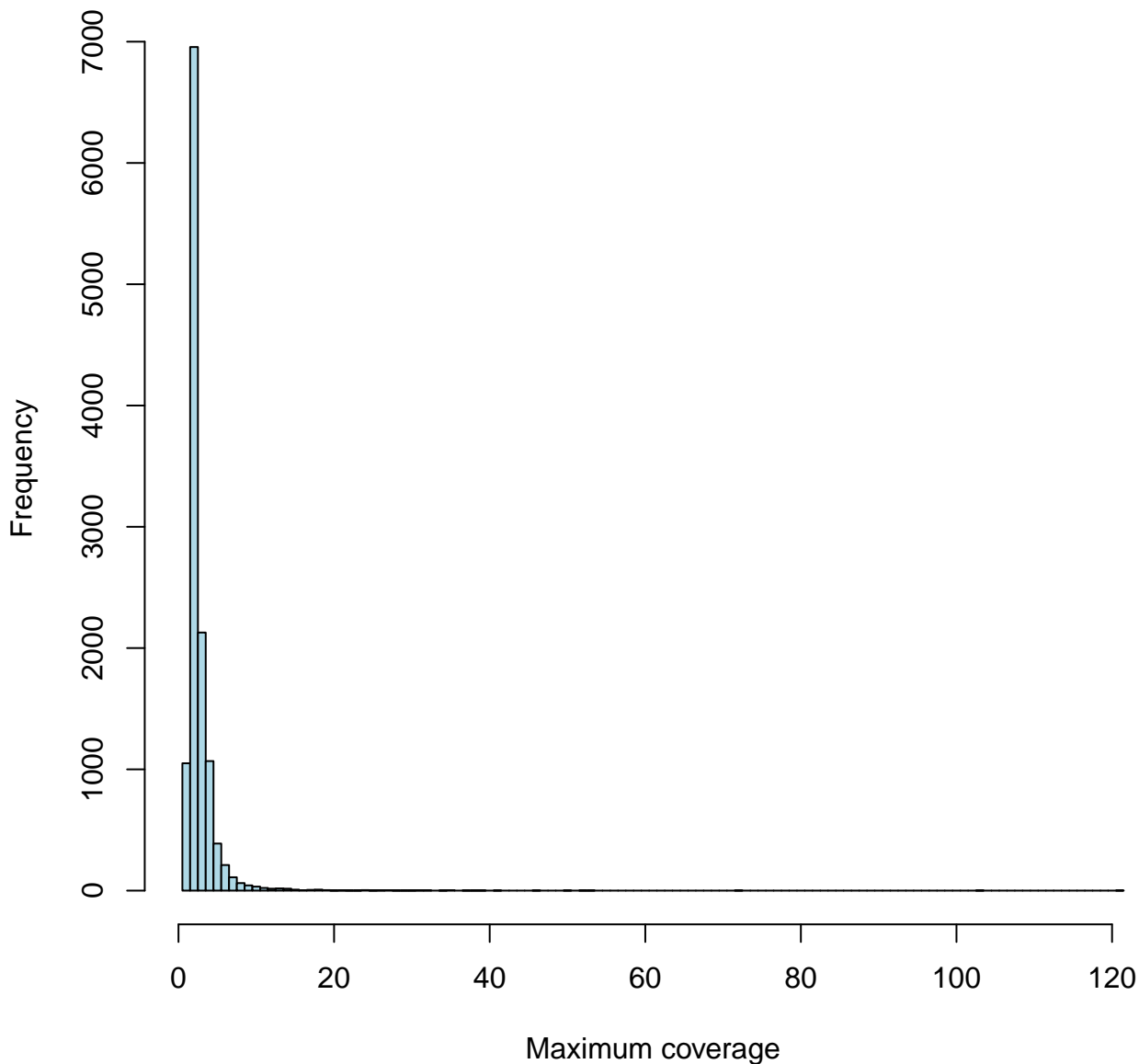
**jx in Intropolis but not "annotated" by knownGene:
10 samples; n jx: 14480**



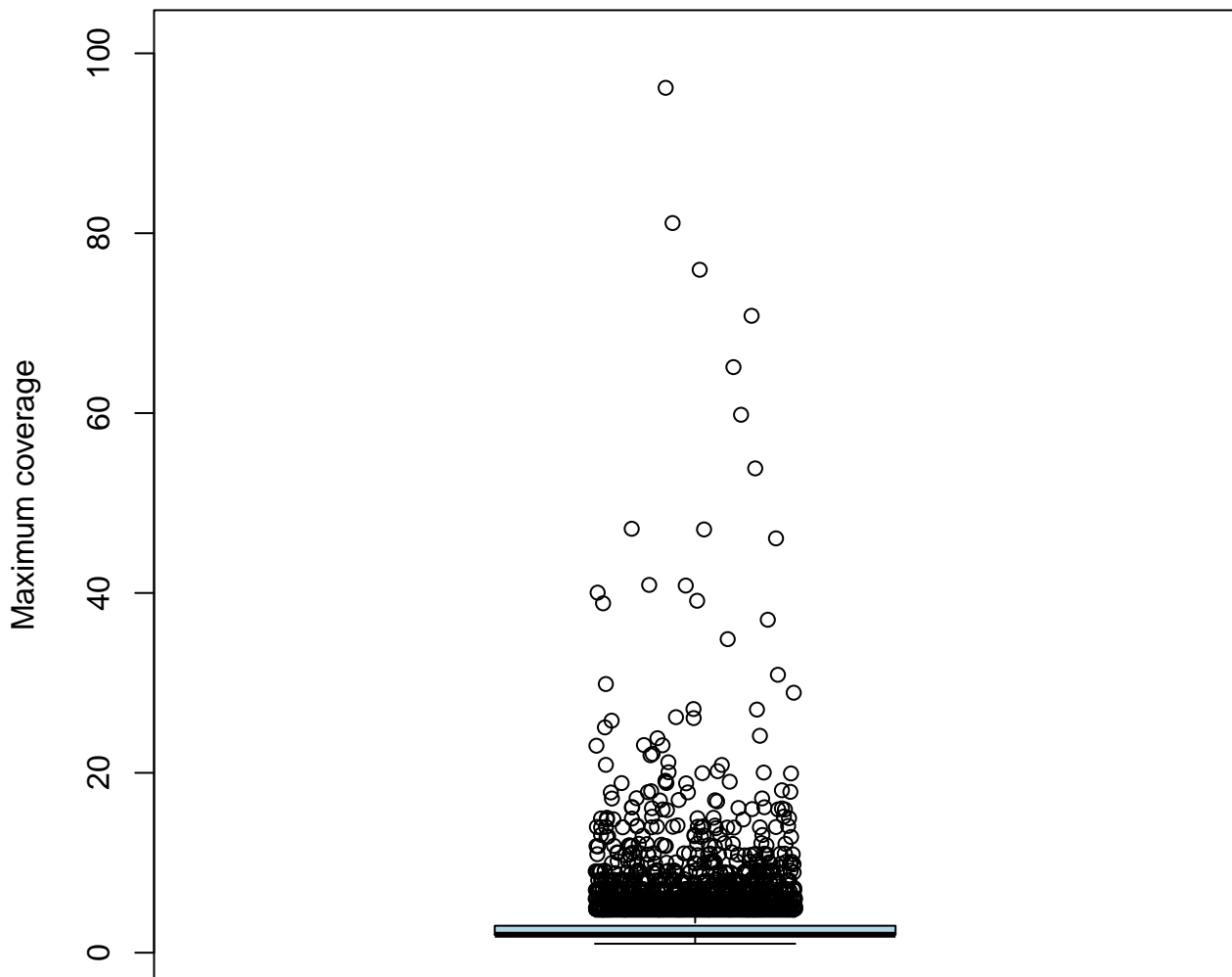
**jx in Intropolis but not "annotated" by knownGene:
11 samples; n jx: 12187**



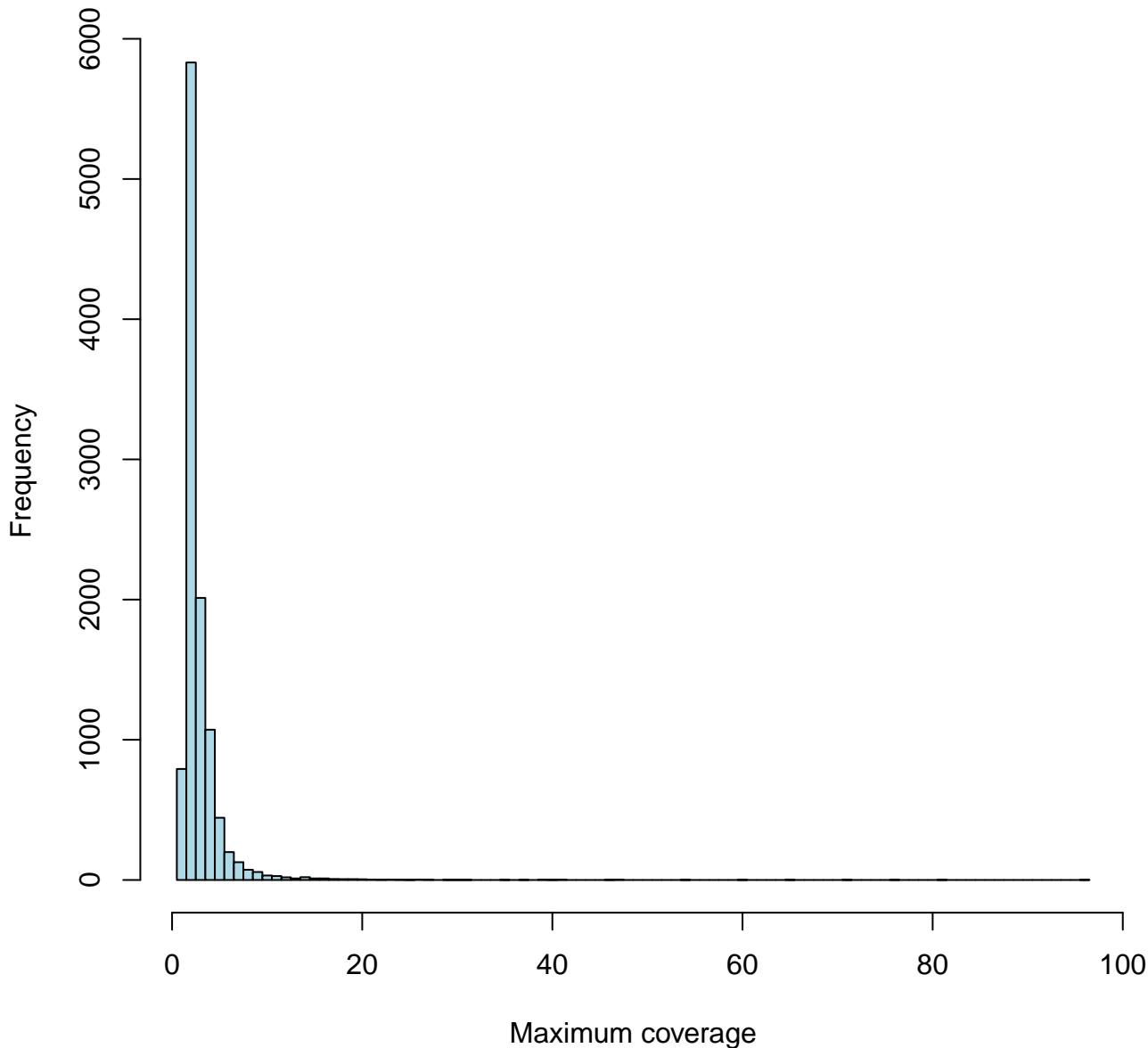
**jx in Intropolis but not "annotated" by knownGene:
11 samples; n jx: 12187**



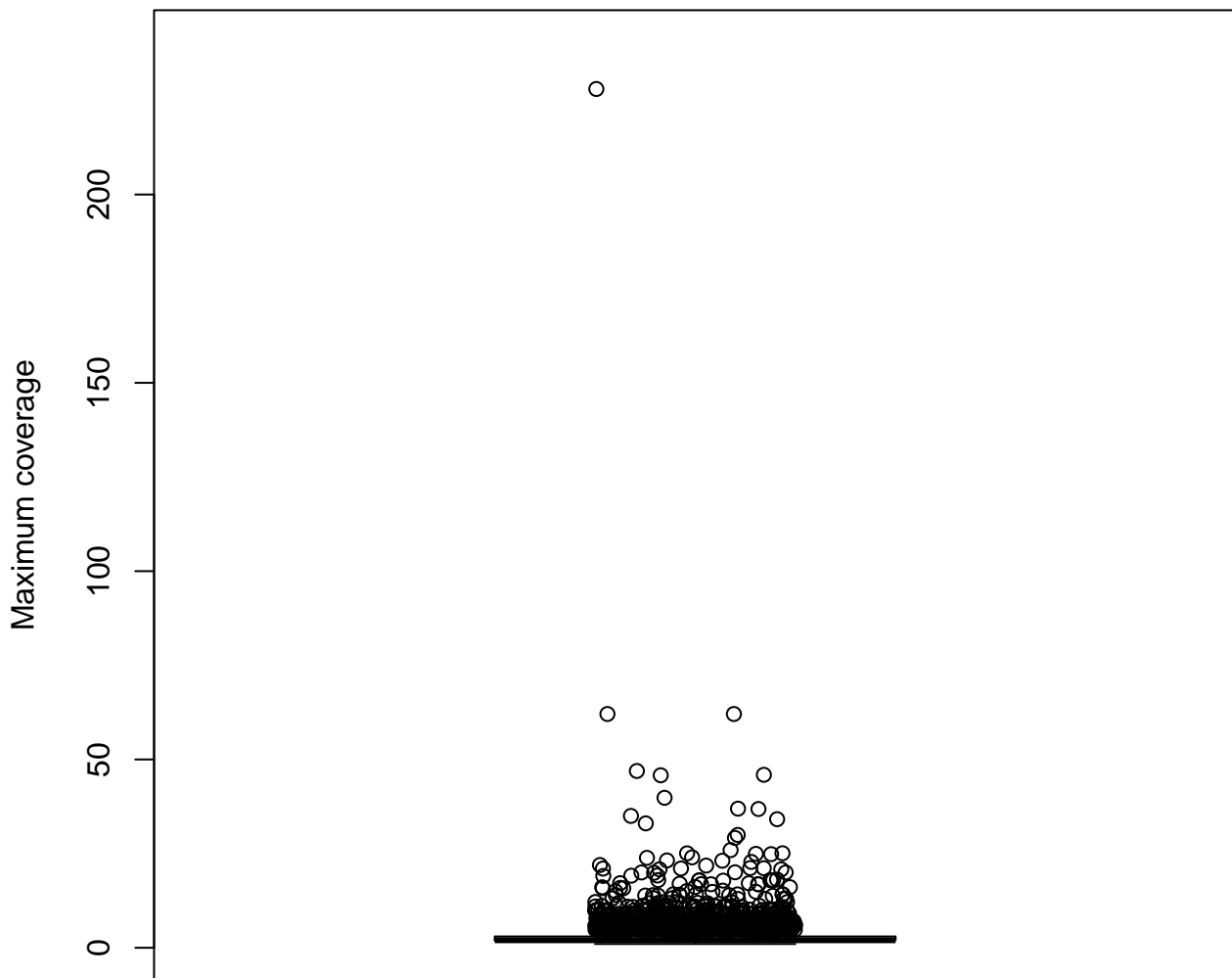
**jx in Intropolis but not "annotated" by knownGene:
12 samples; n jx: 10799**



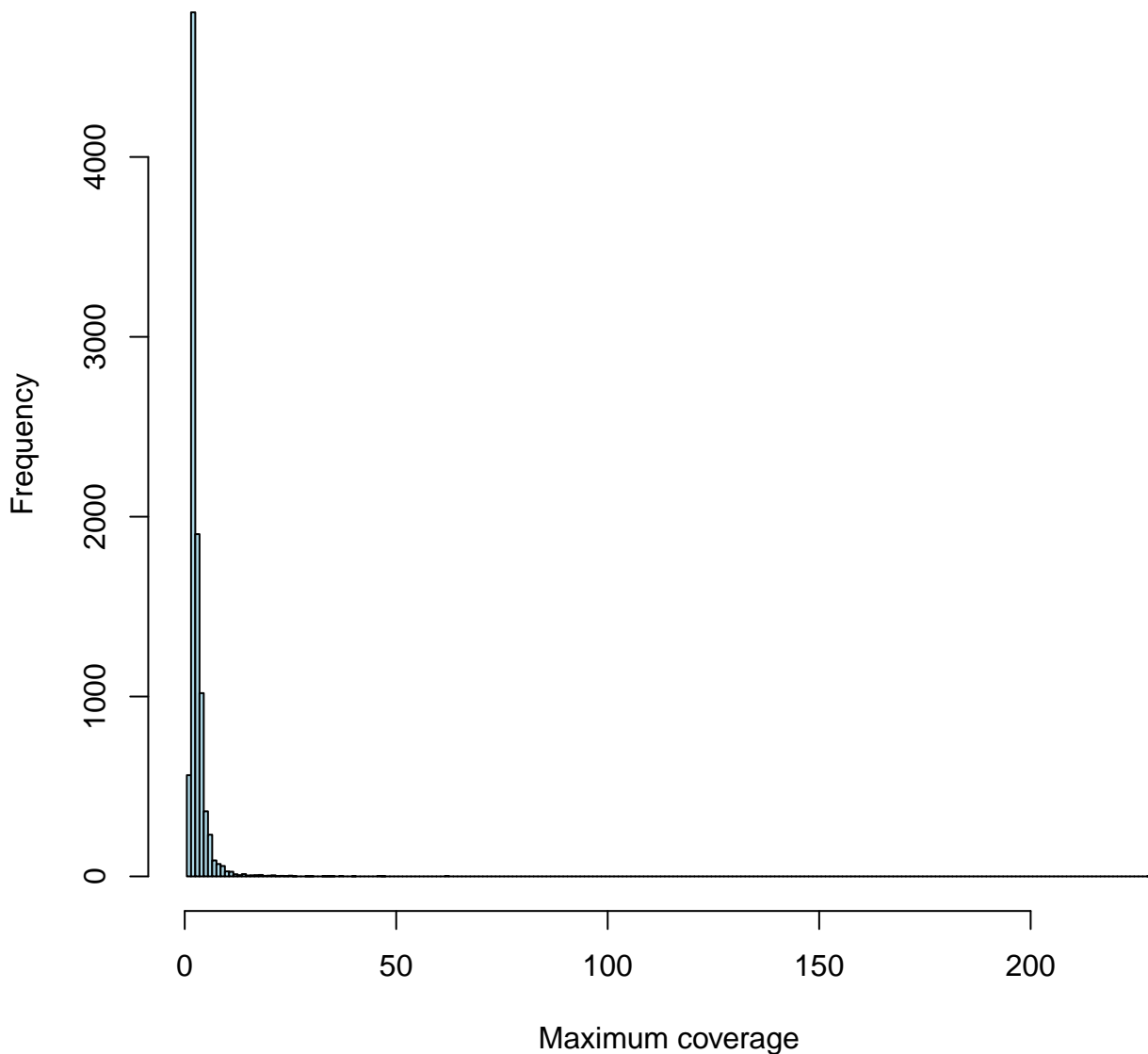
**jx in Intropolis but not "annotated" by knownGene:
12 samples; n jx: 10799**



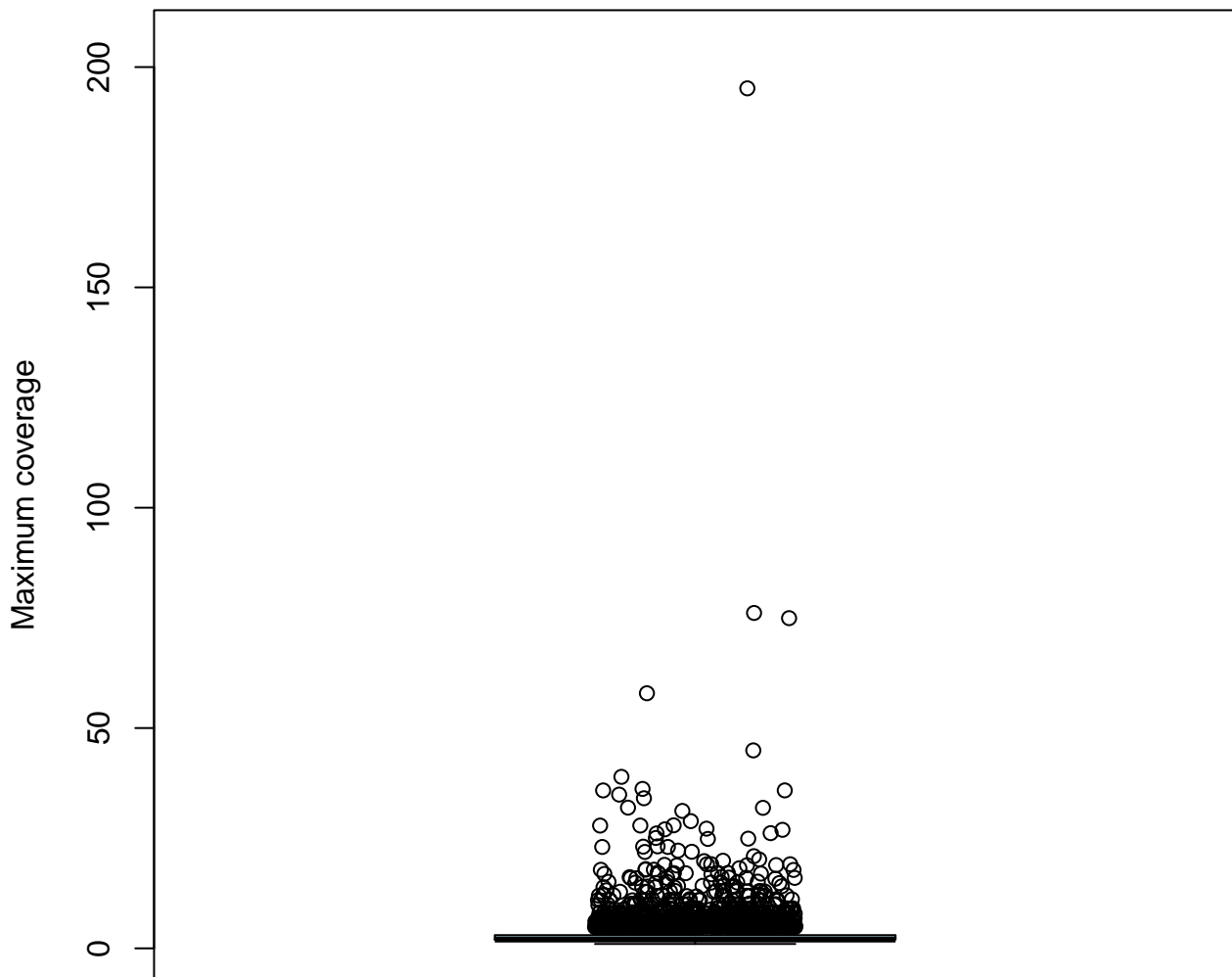
**jx in Intropolis but not "annotated" by knownGene:
13 samples; n jx: 9250**



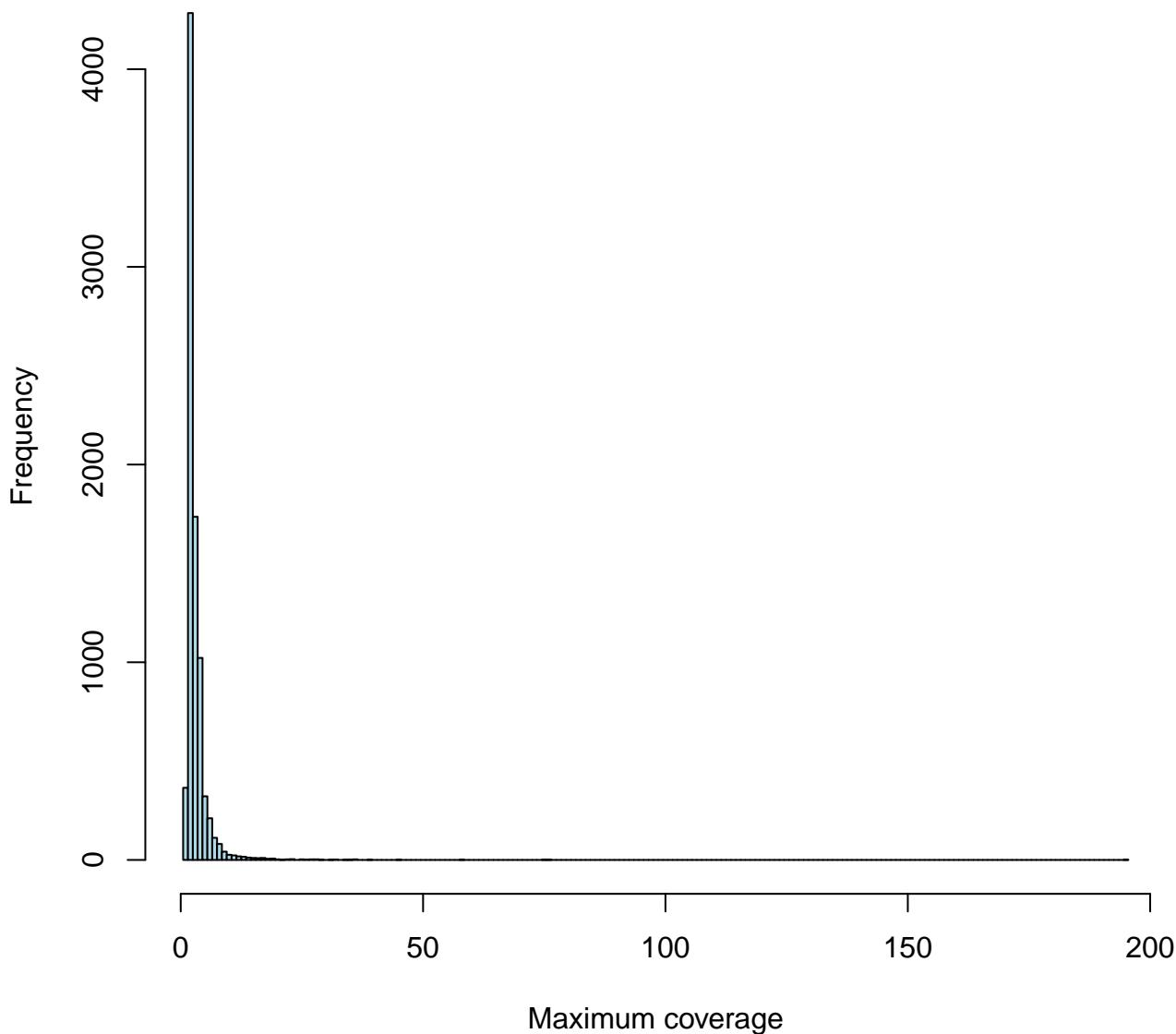
**jx in Intropolis but not "annotated" by knownGene:
13 samples; n jx: 9250**



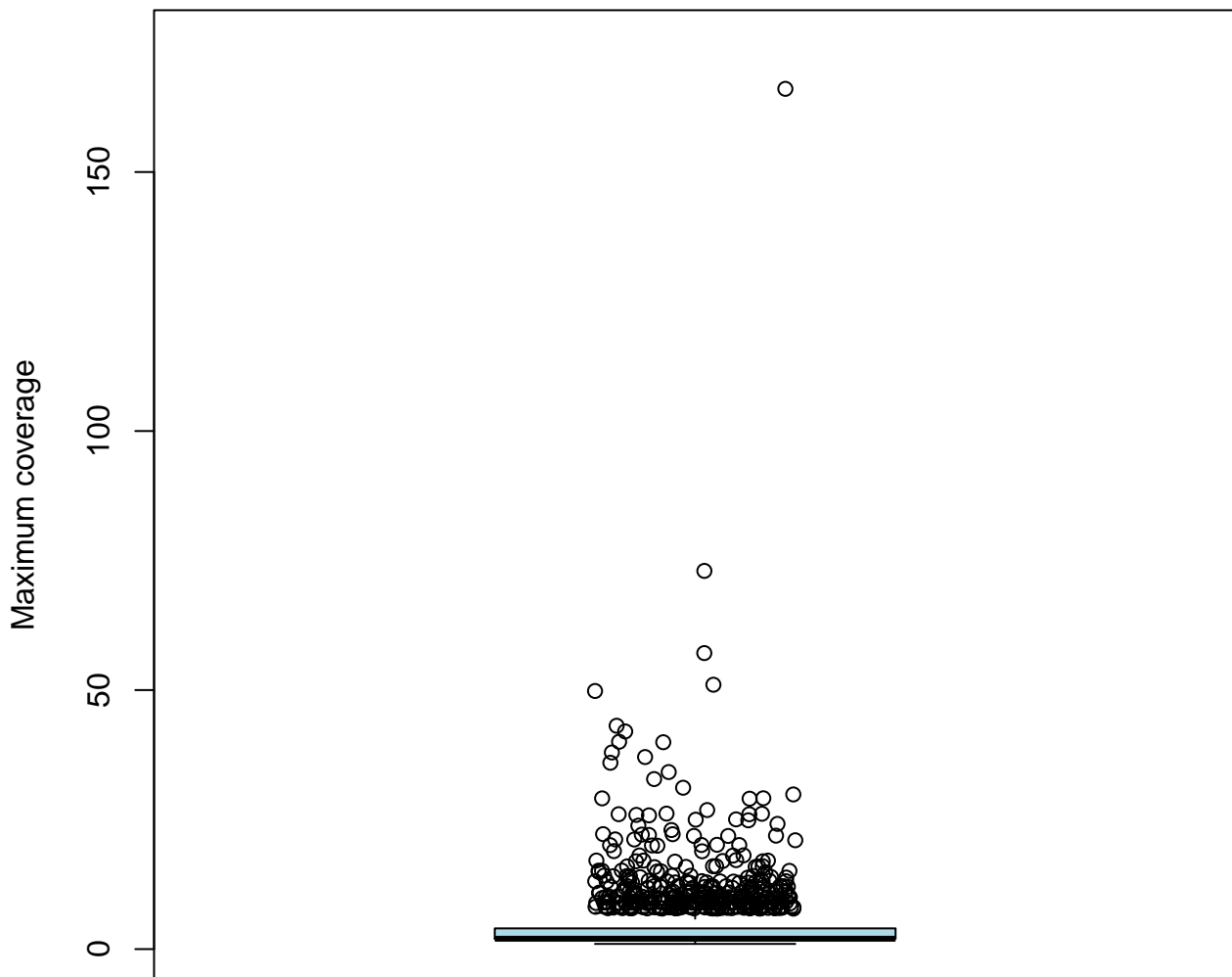
**jx in Intropolis but not "annotated" by knownGene:
14 samples; n jx: 8349**



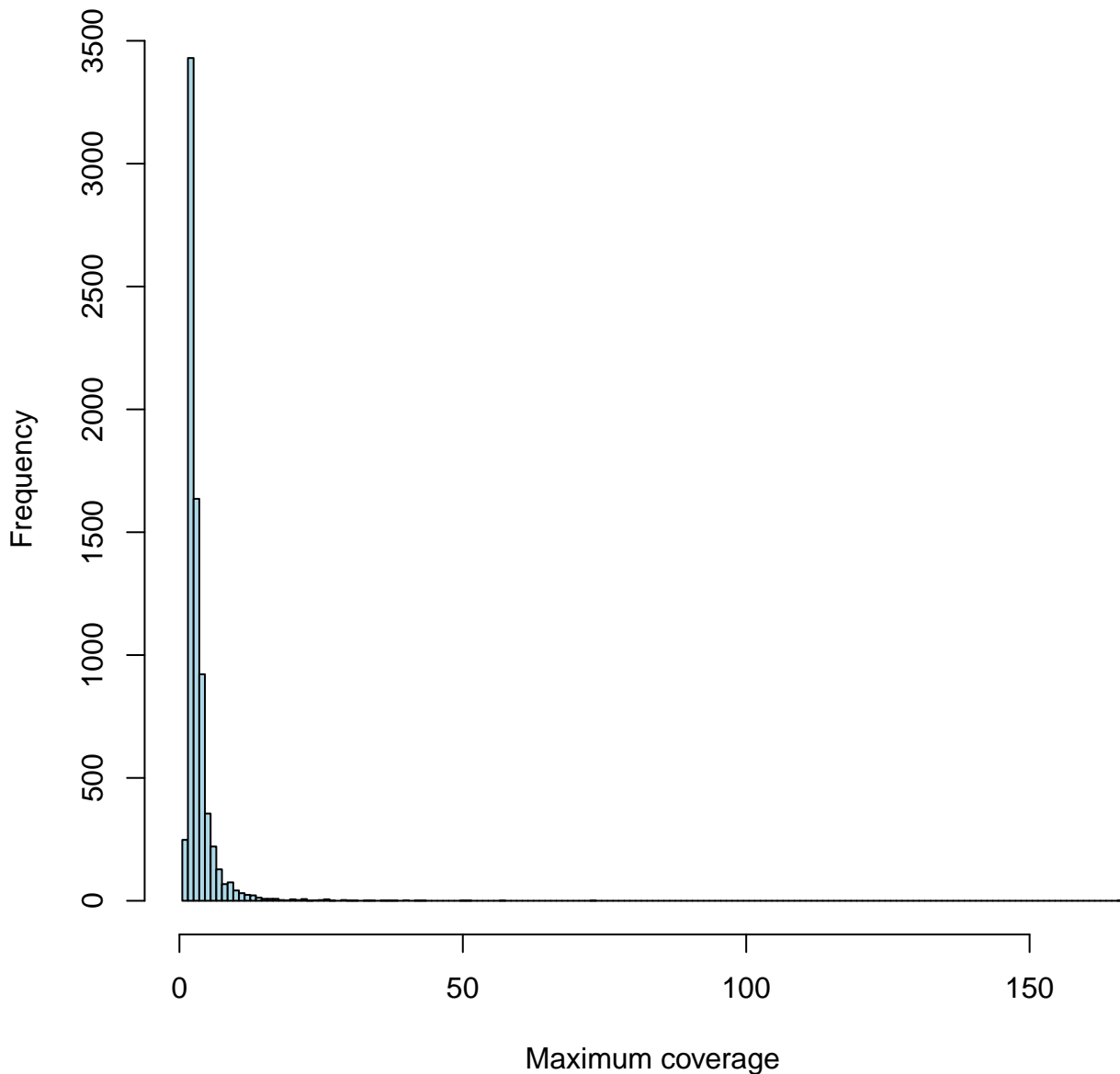
**jx in Intropolis but not "annotated" by knownGene:
14 samples; n jx: 8349**



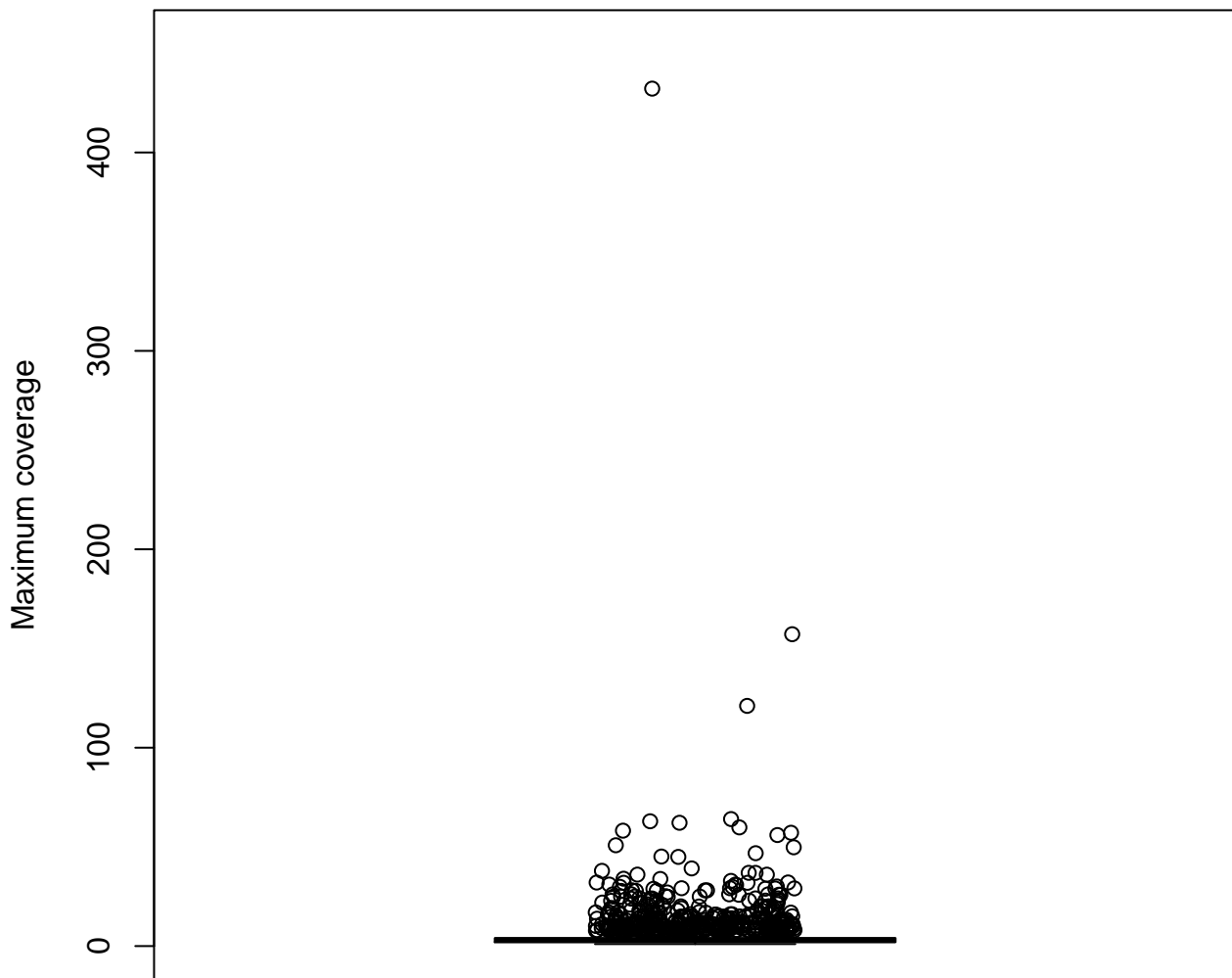
**jx in Intropolis but not "annotated" by knownGene:
15 samples; n jx: 7292**



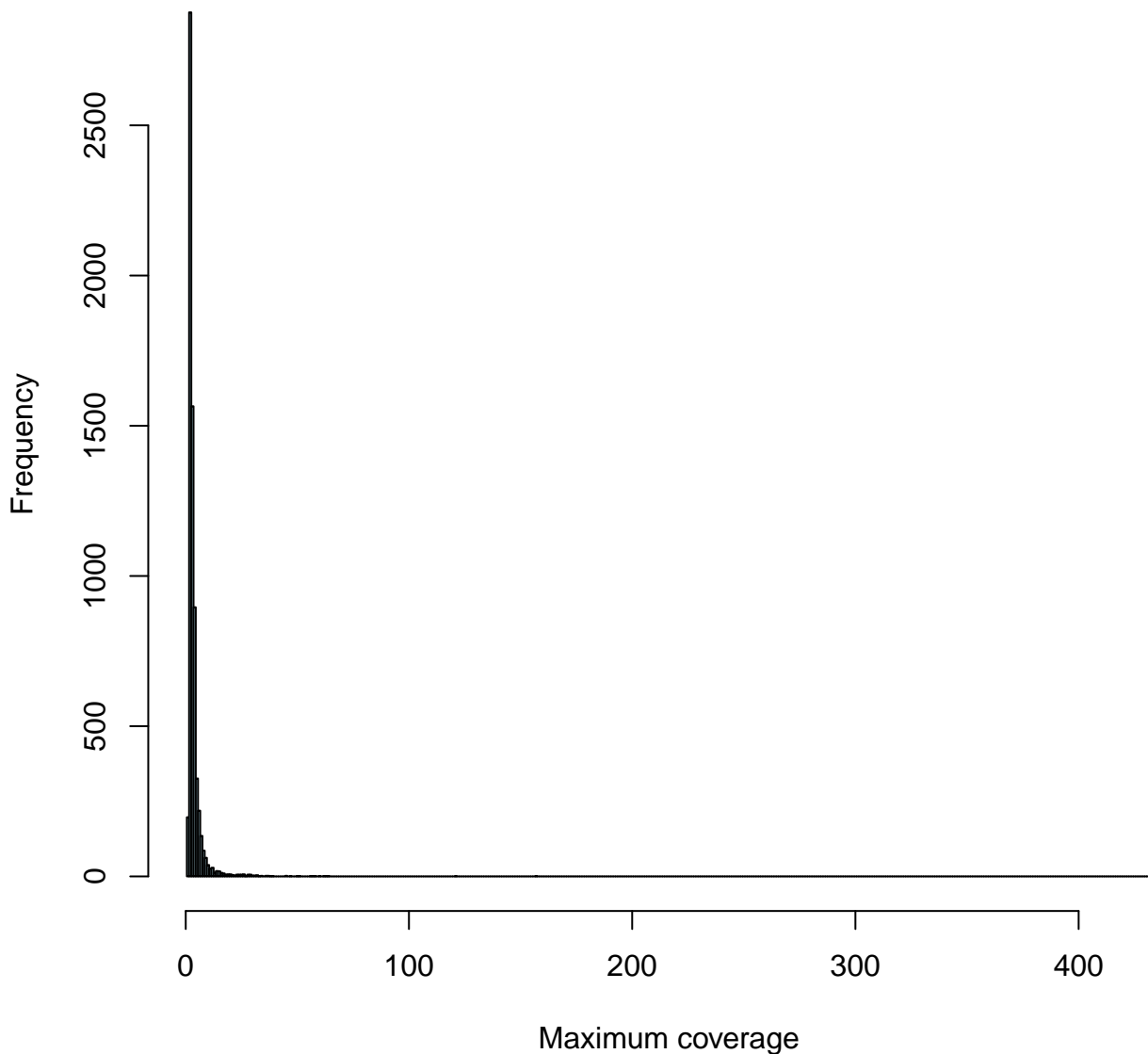
**jx in Intropolis but not "annotated" by knownGene:
15 samples; n jx: 7292**



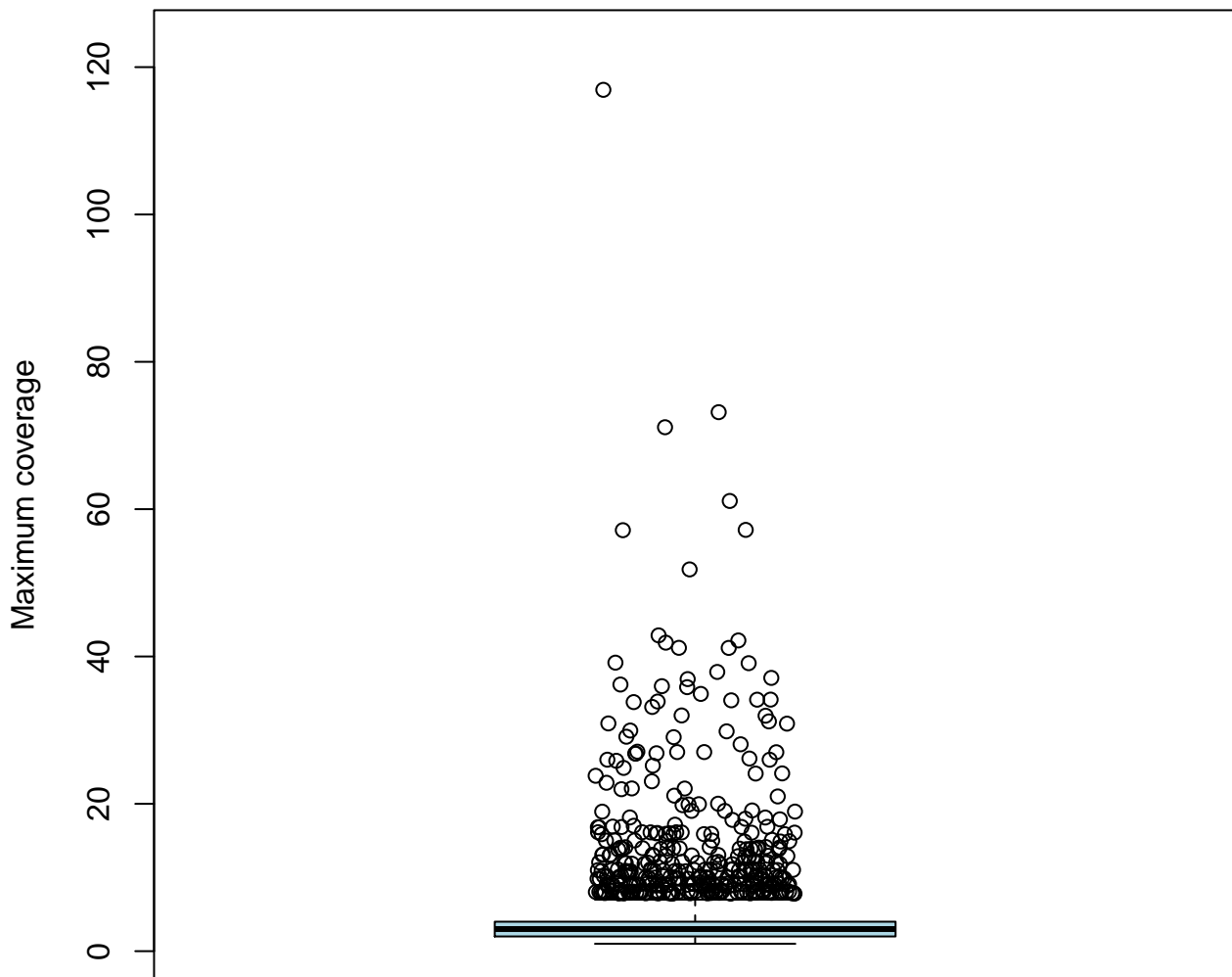
**jx in Intropolis but not "annotated" by knownGene:
16 samples; n jx: 6625**



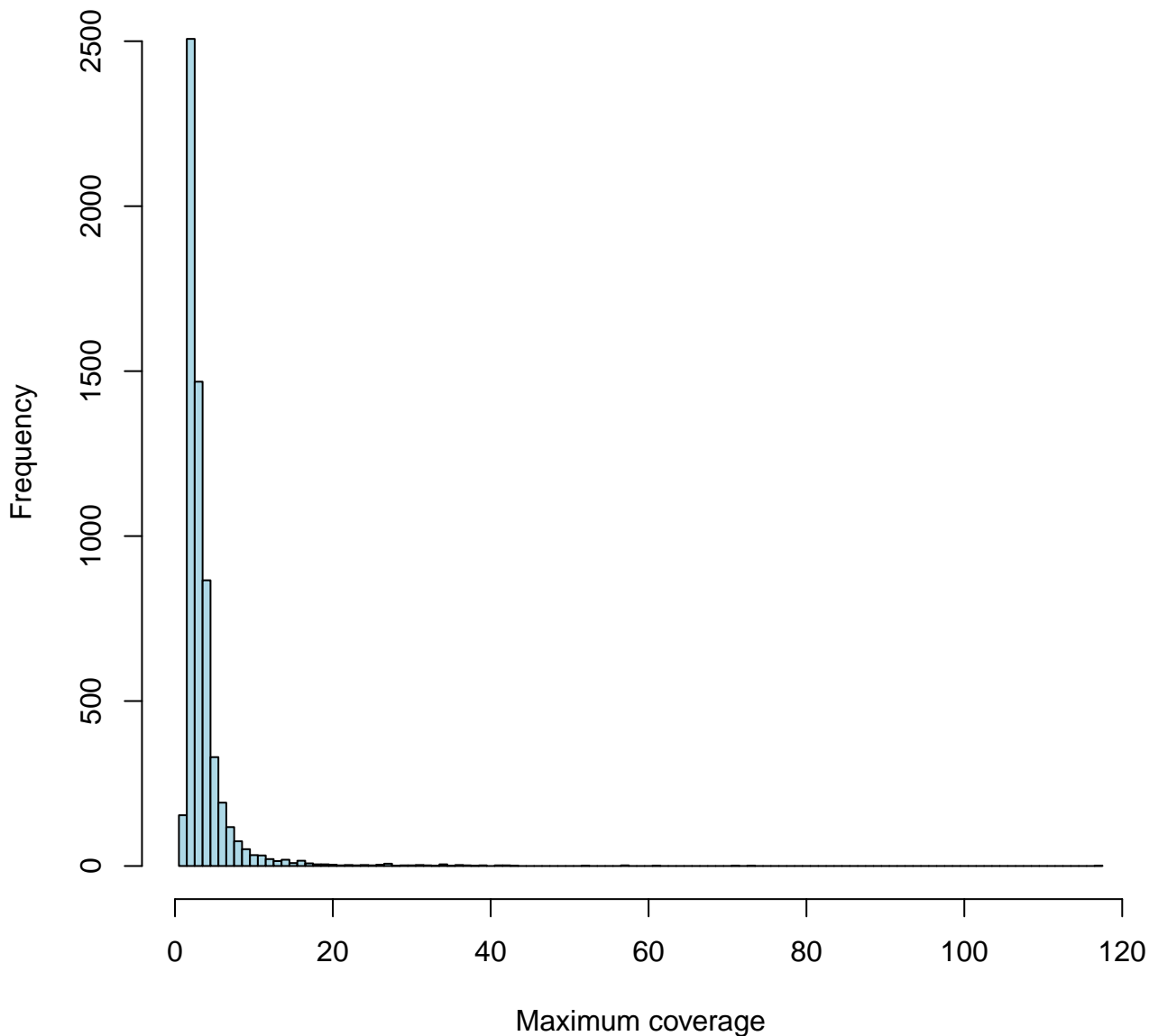
**jx in Intropolis but not "annotated" by knownGene:
16 samples; n jx: 6625**



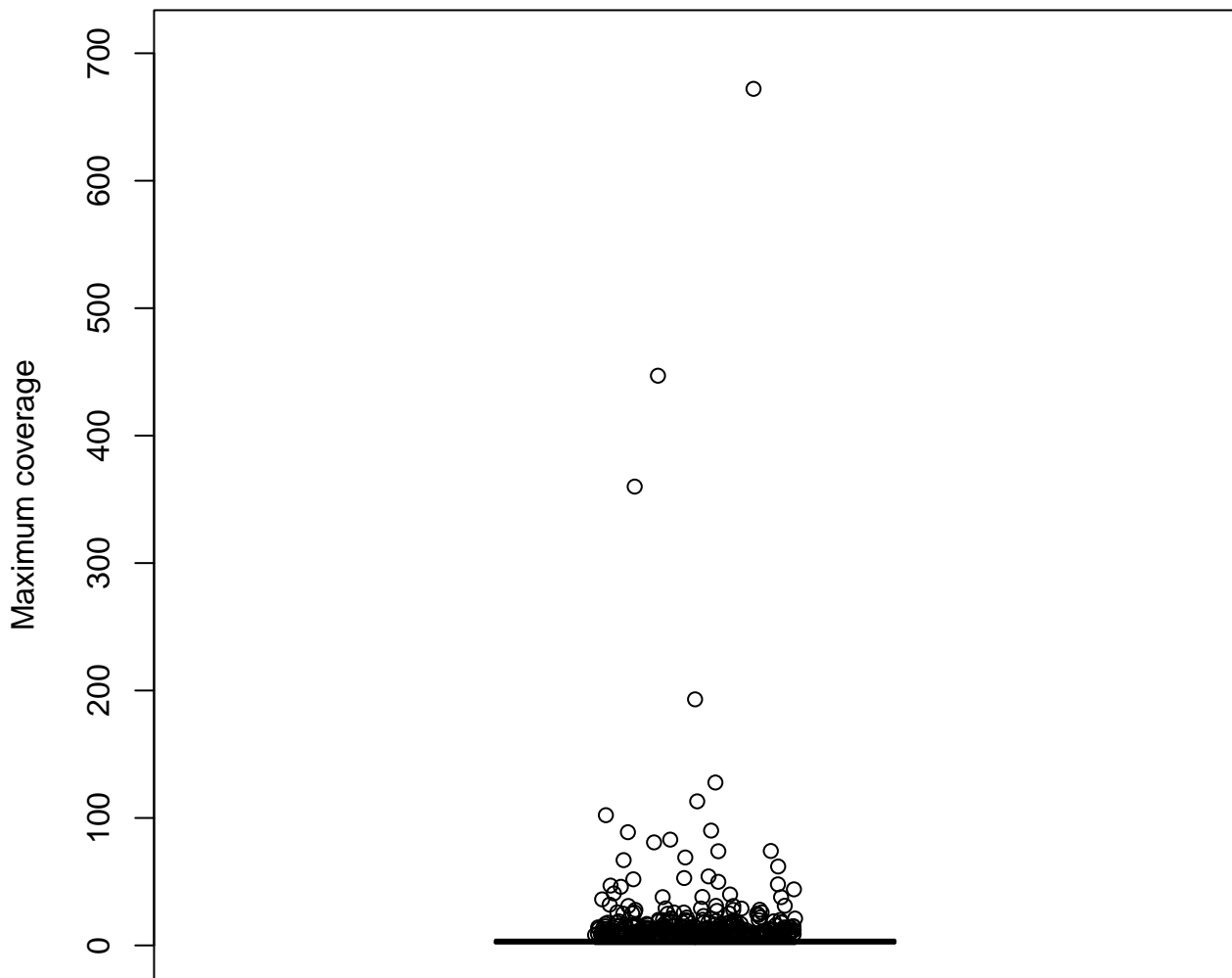
**jx in Intropolis but not "annotated" by knownGene:
17 samples; n jx: 5988**



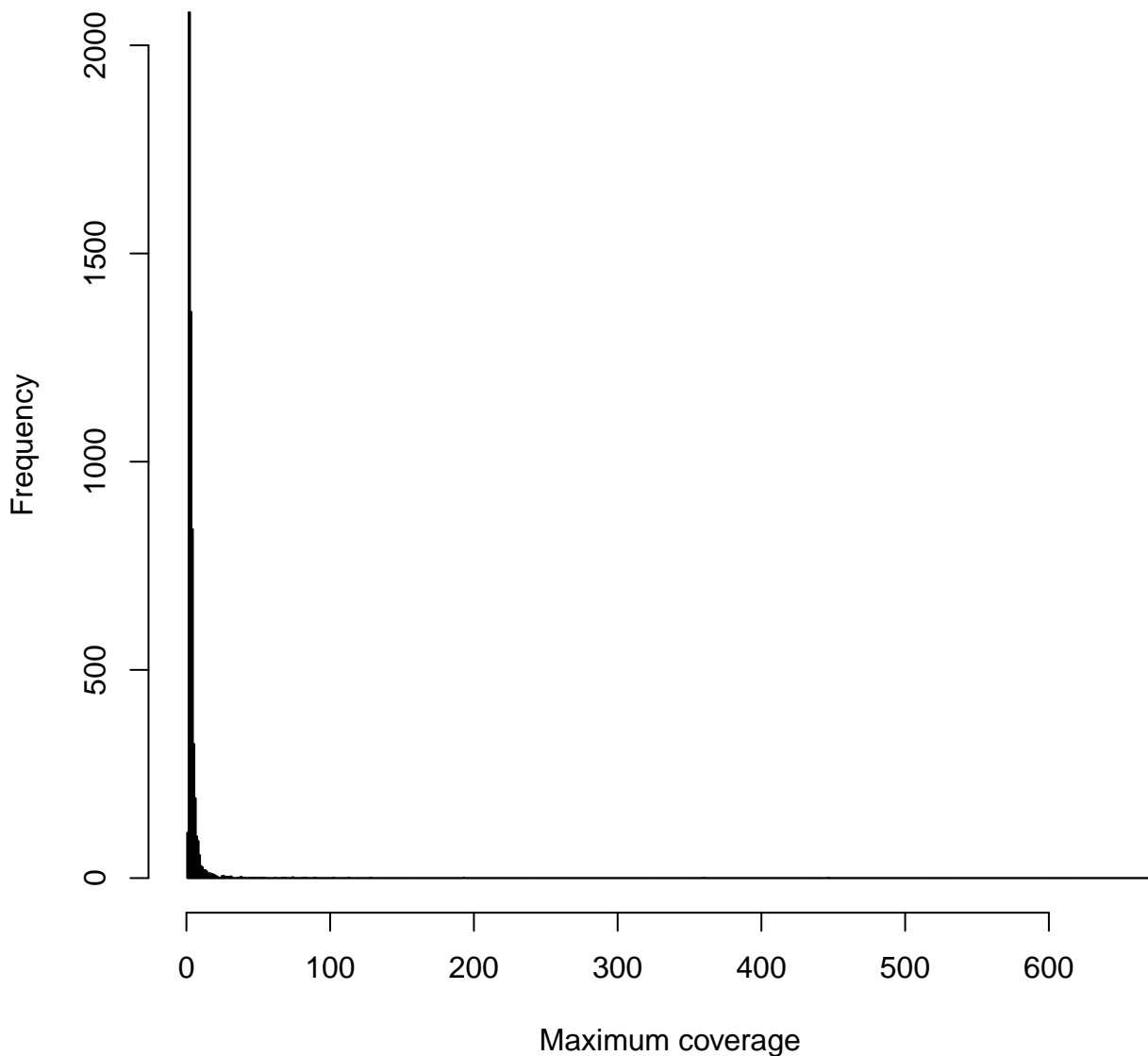
**jx in Intropolis but not "annotated" by knownGene:
17 samples; n jx: 5988**



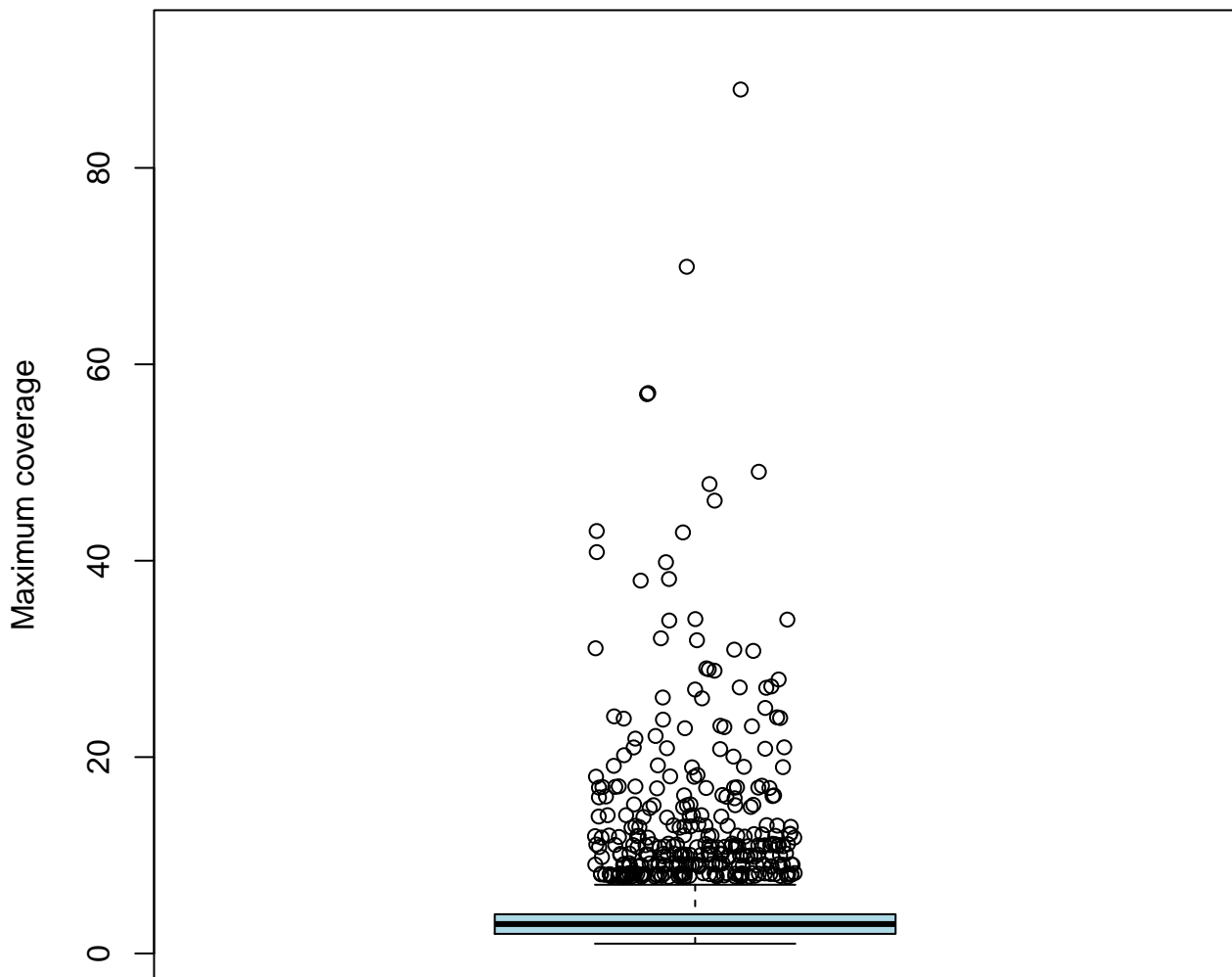
**jx in Intropolis but not "annotated" by knownGene:
18 samples; n jx: 5360**



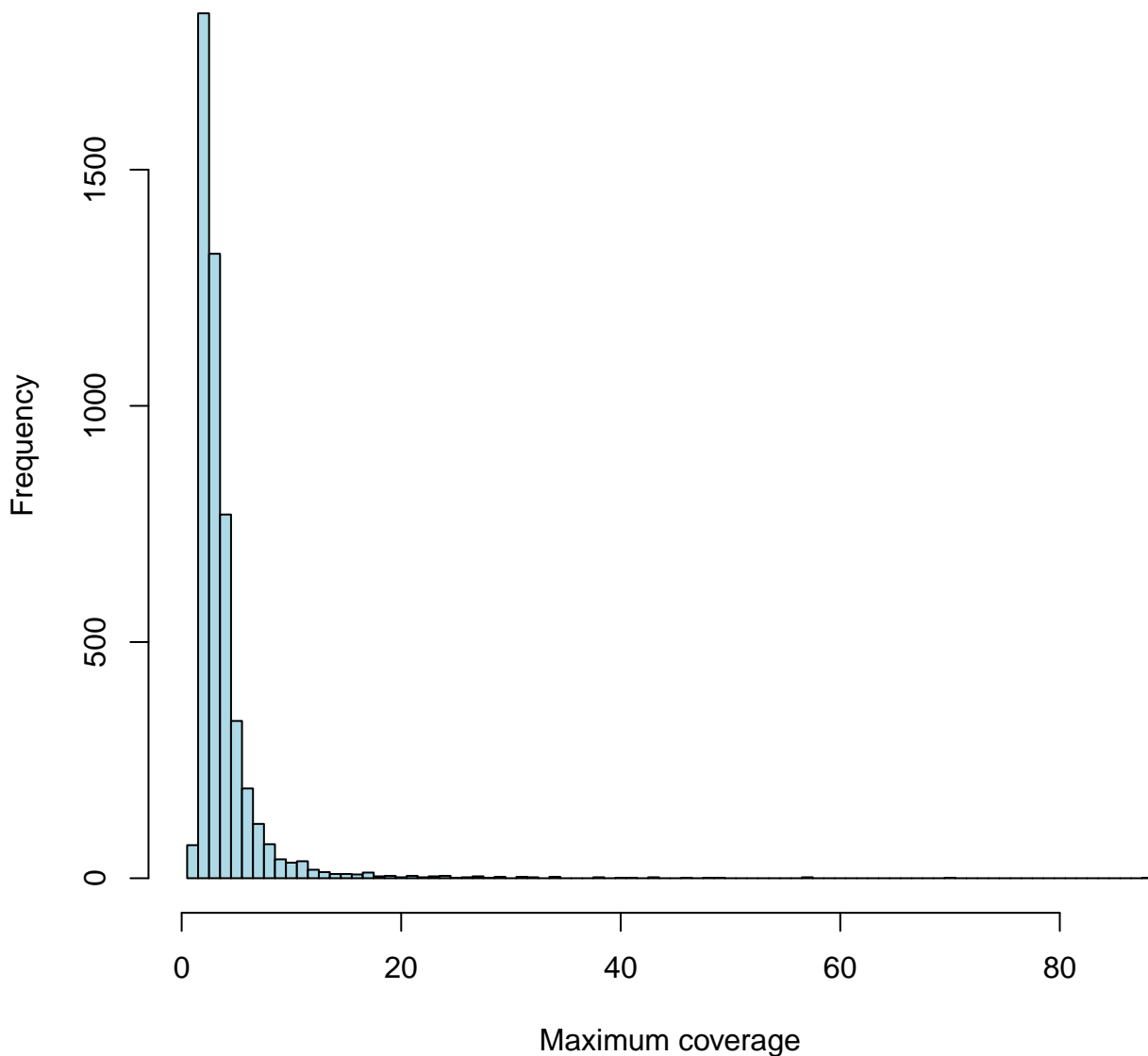
**jx in Intropolis but not "annotated" by knownGene:
18 samples; n jx: 5360**



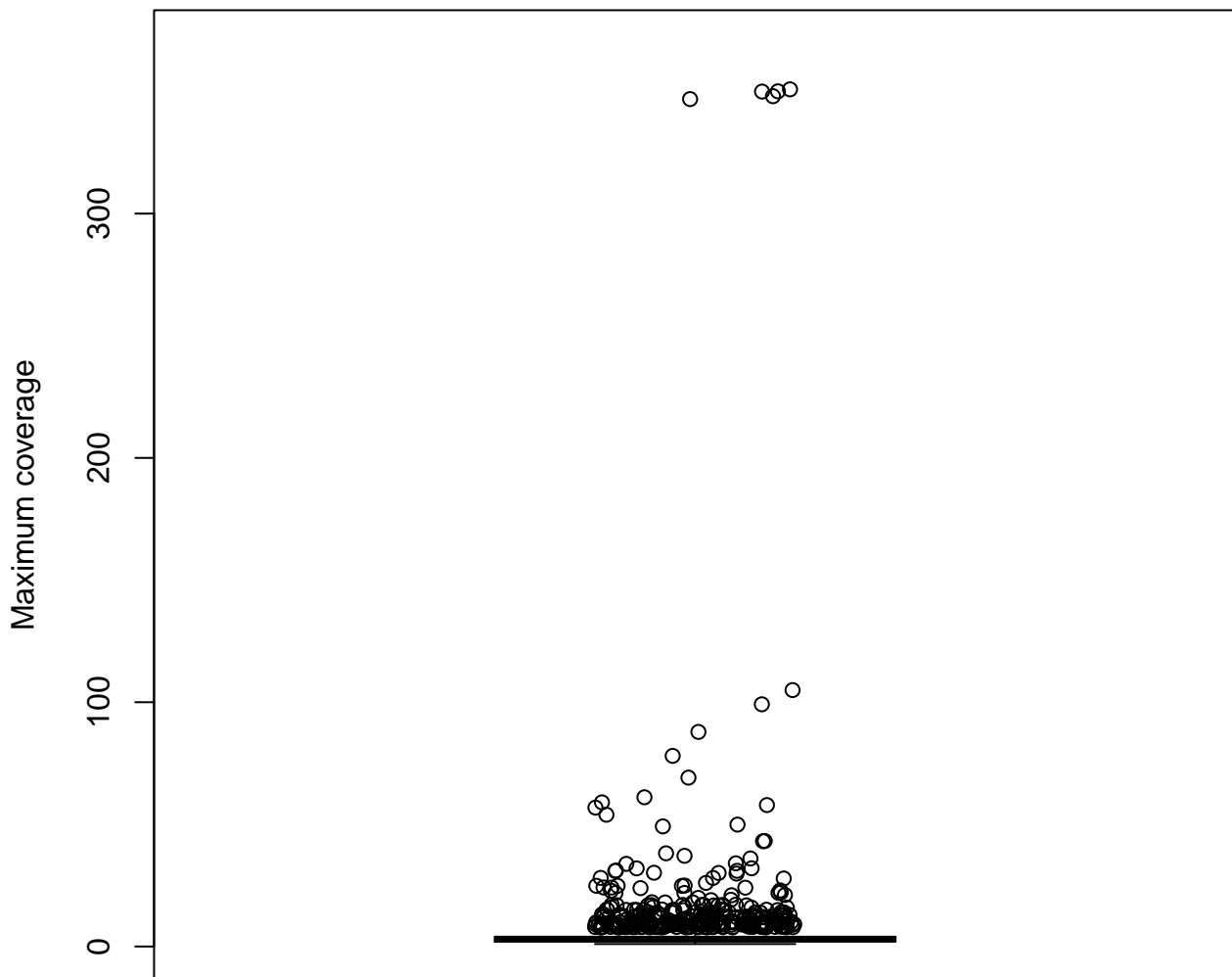
**jx in Intropolis but not "annotated" by knownGene:
19 samples; n jx: 4940**



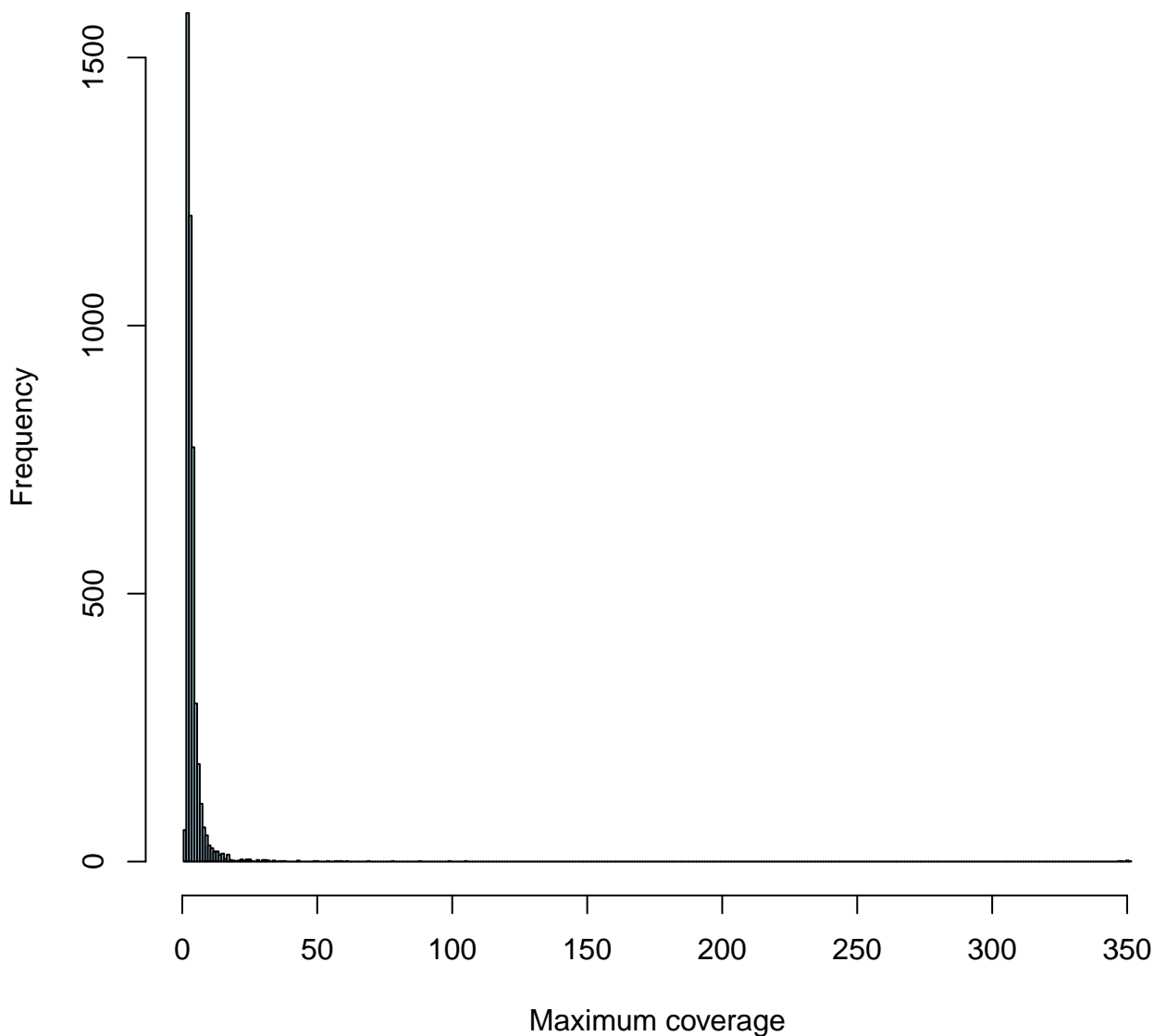
**jx in Intropolis but not "annotated" by knownGene:
19 samples; n jx: 4940**



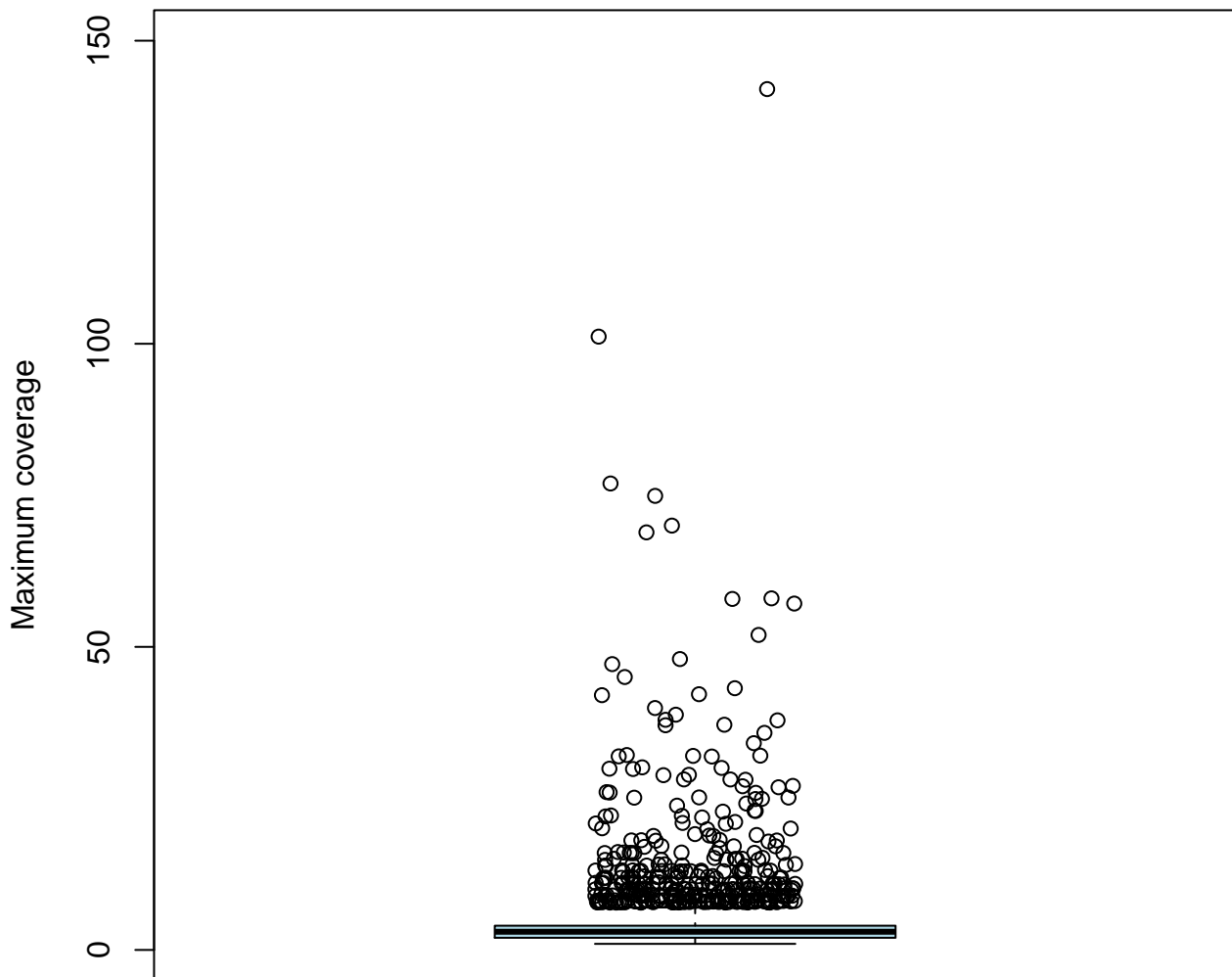
**jx in Intropolis but not "annotated" by knownGene:
20 samples; n jx: 4515**



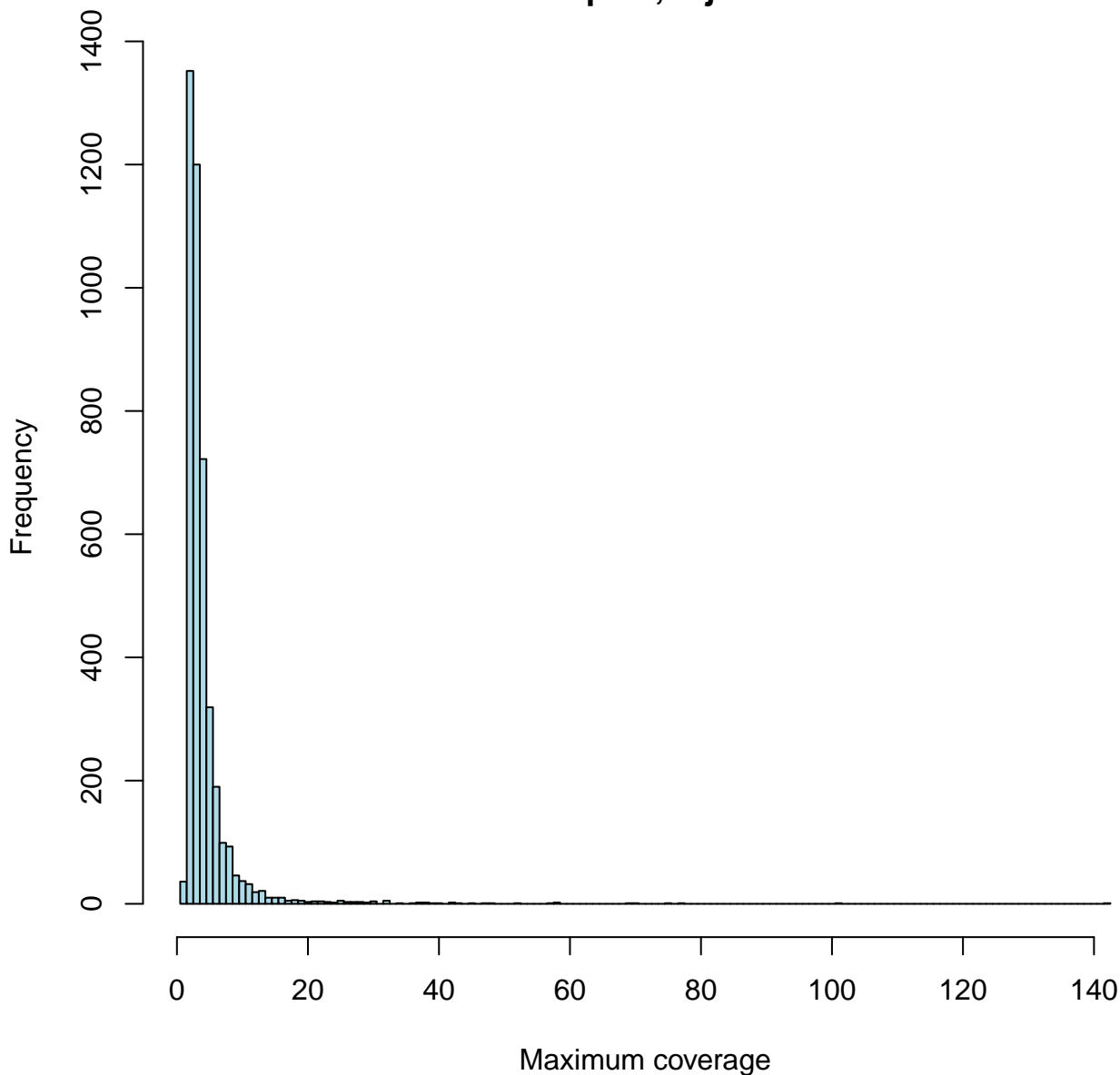
**jx in Intropolis but not "annotated" by knownGene:
20 samples; n jx: 4515**



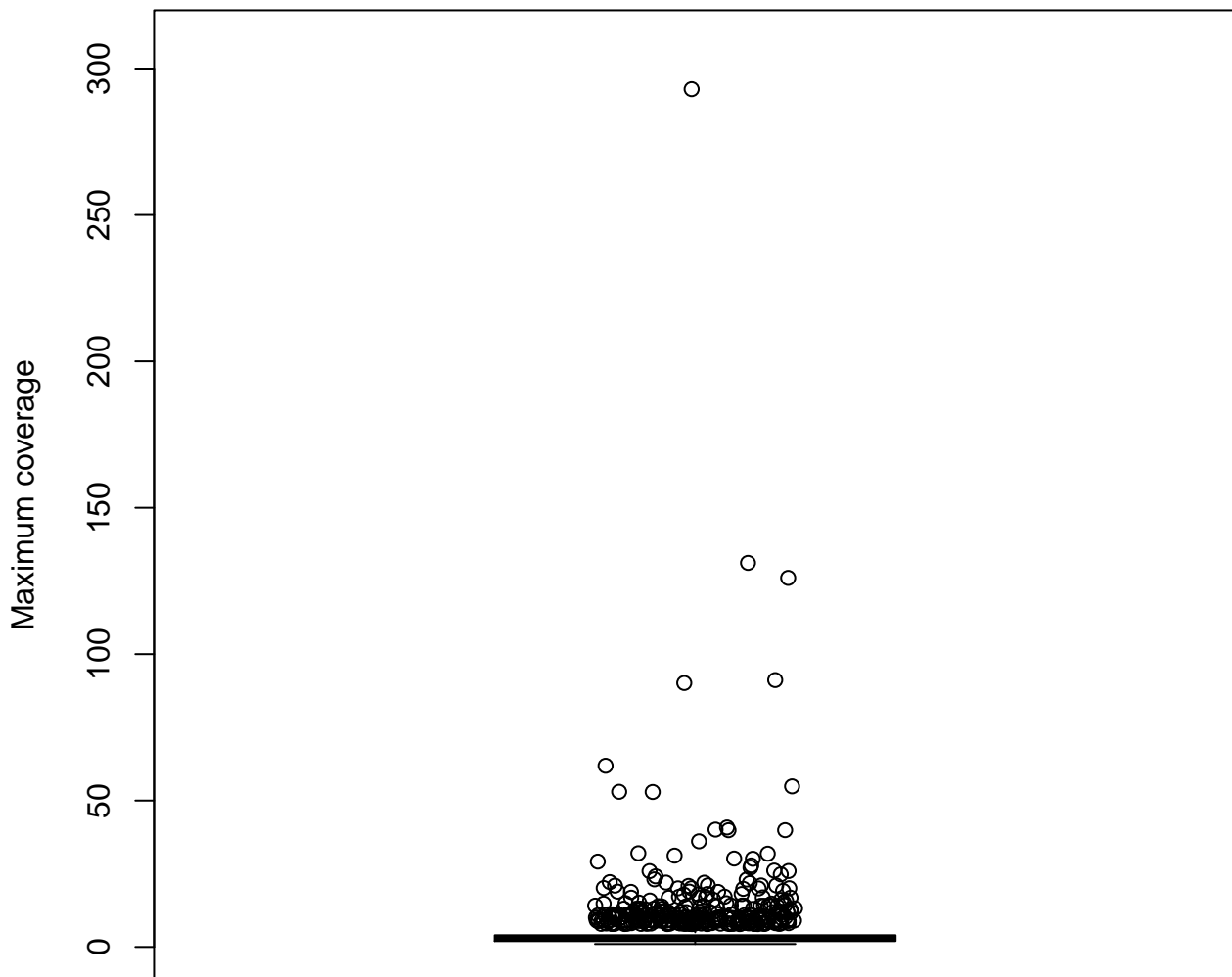
**jx in Intropolis but not "annotated" by knownGene:
21 samples; n jx: 4277**



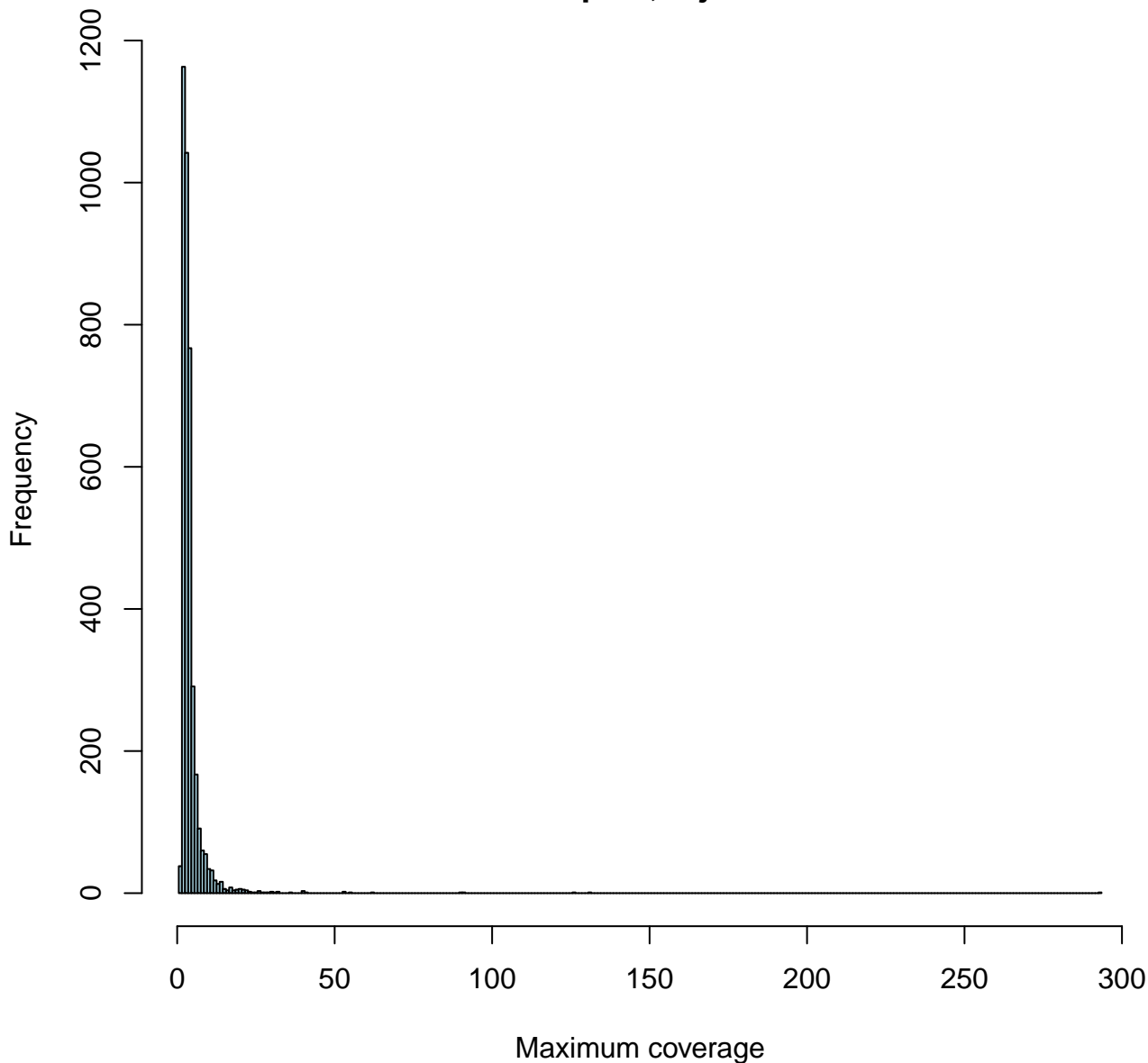
**jx in Intropolis but not "annotated" by knownGene:
21 samples; n jx: 4277**



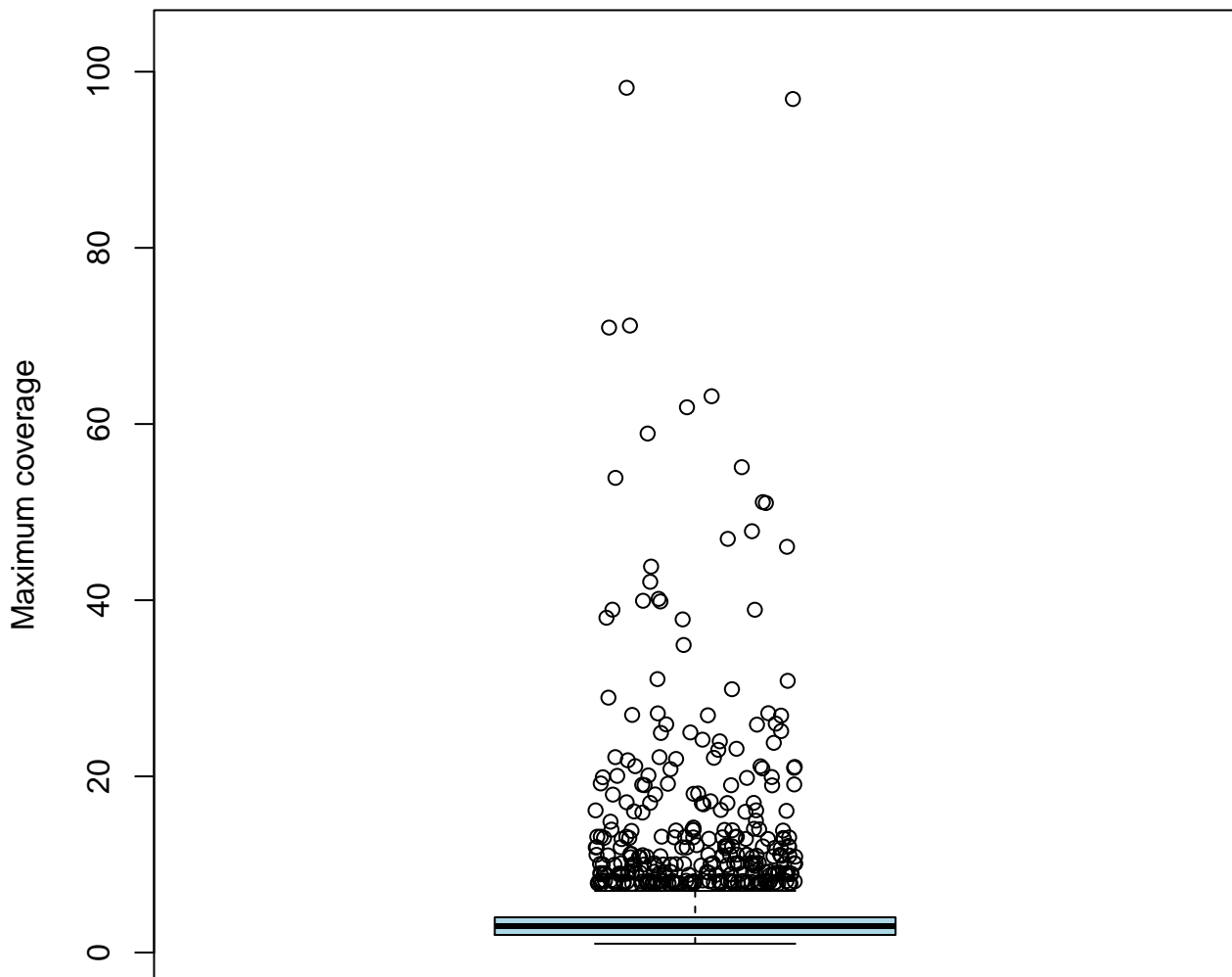
**jx in Intropolis but not "annotated" by knownGene:
22 samples; n jx: 3858**



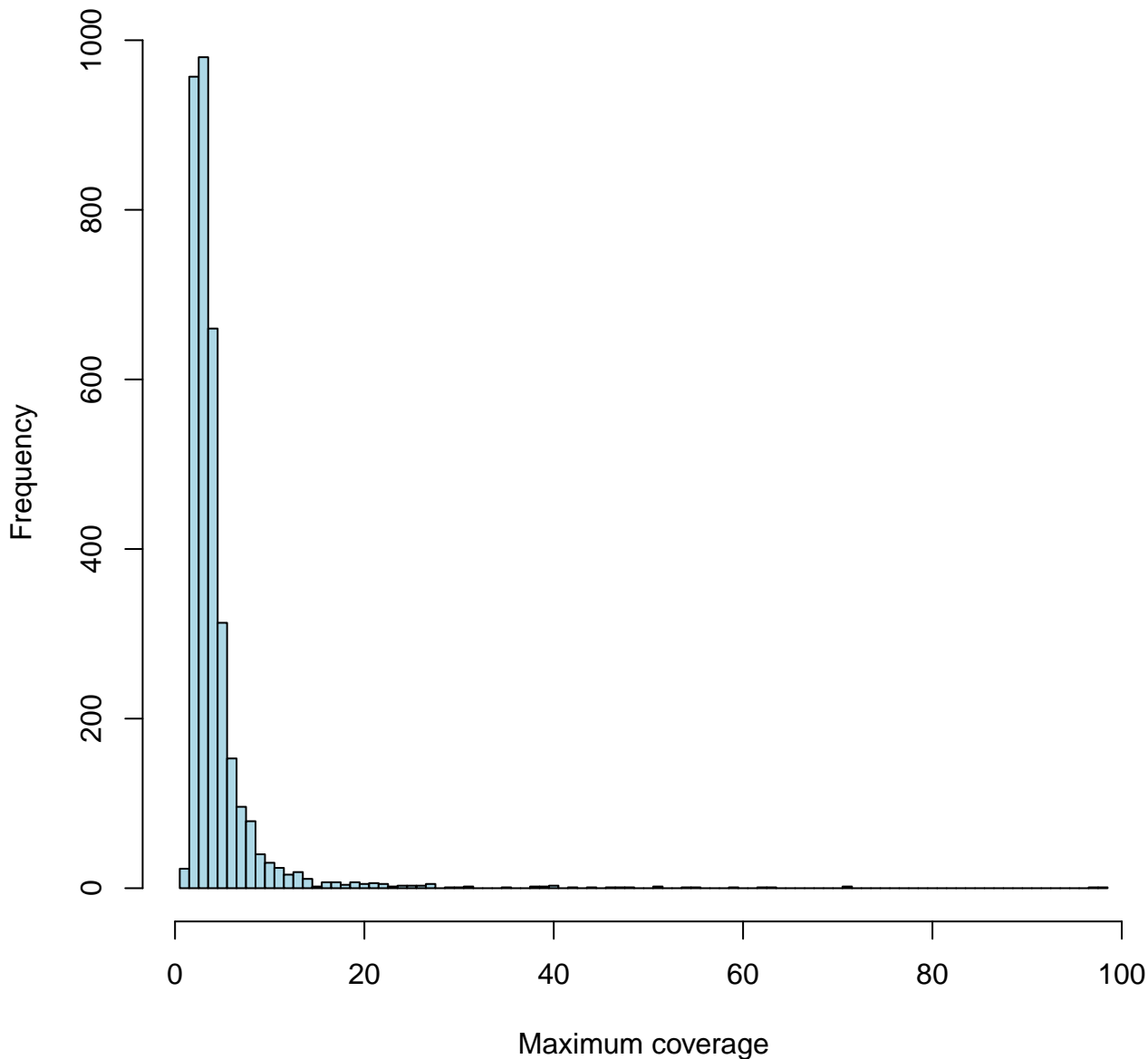
**jx in Intropolis but not "annotated" by knownGene:
22 samples; n jx: 3858**



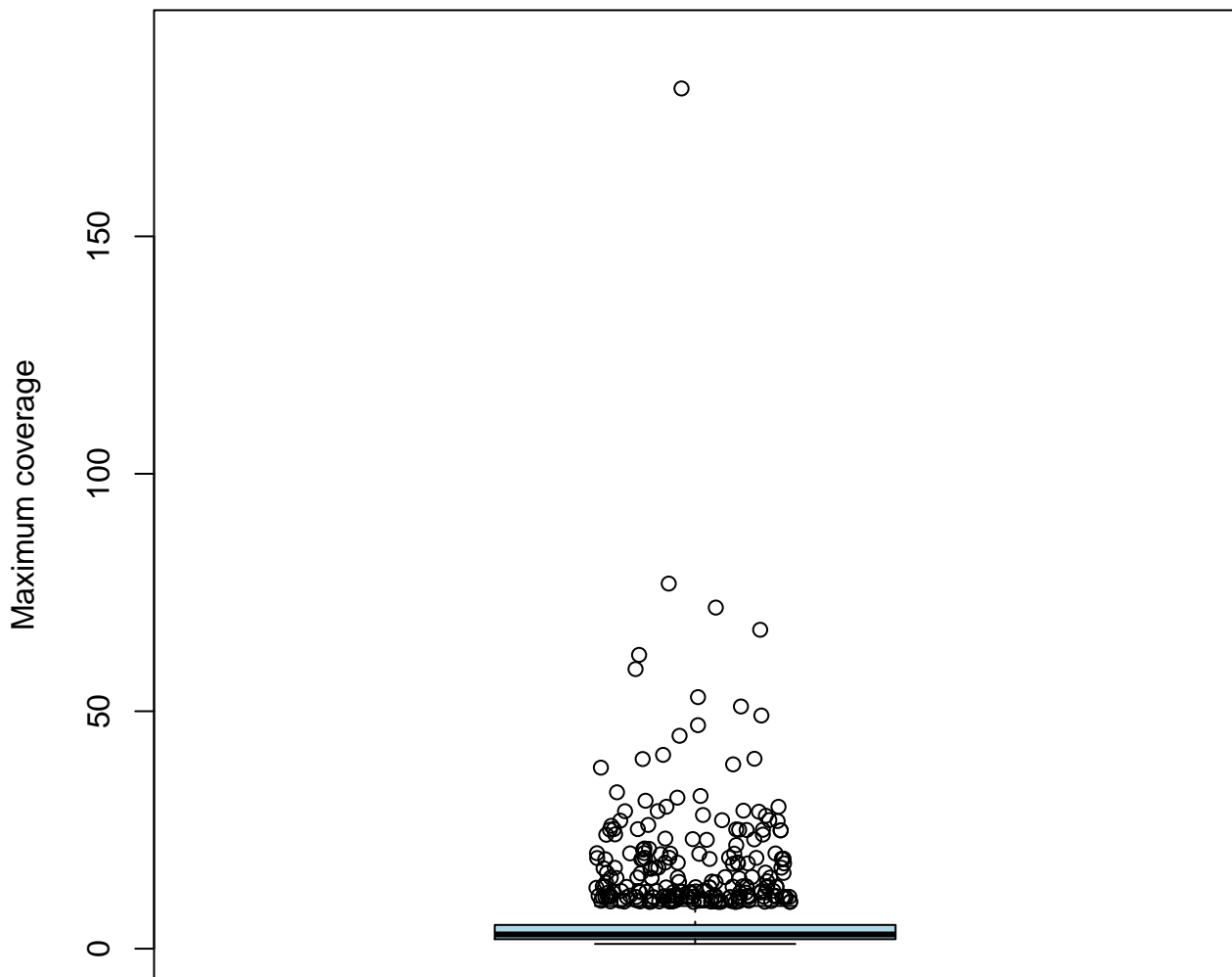
**jx in Intropolis but not "annotated" by knownGene:
23 samples; n jx: 3488**



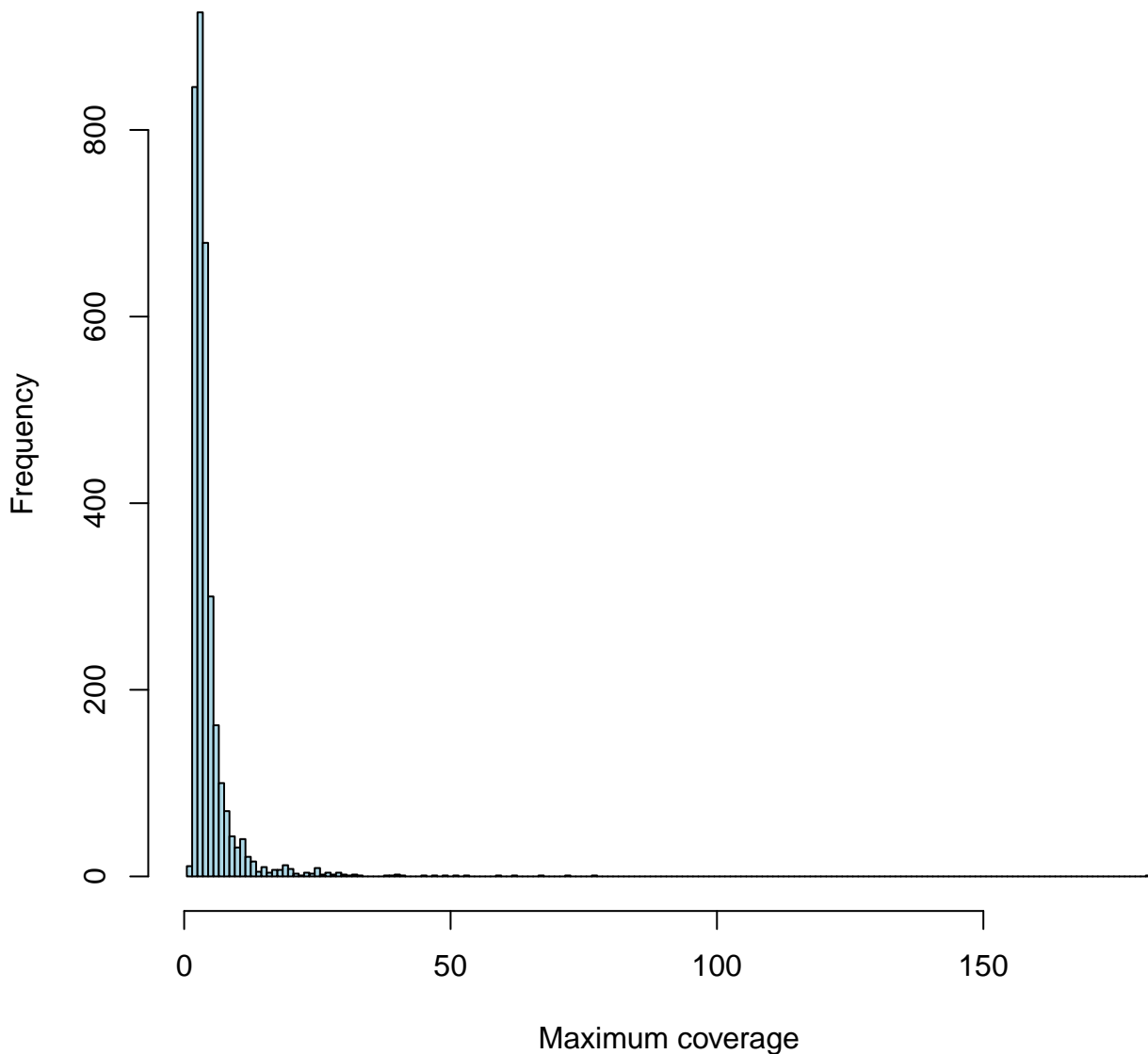
**jx in Intropolis but not "annotated" by knownGene:
23 samples; n jx: 3488**



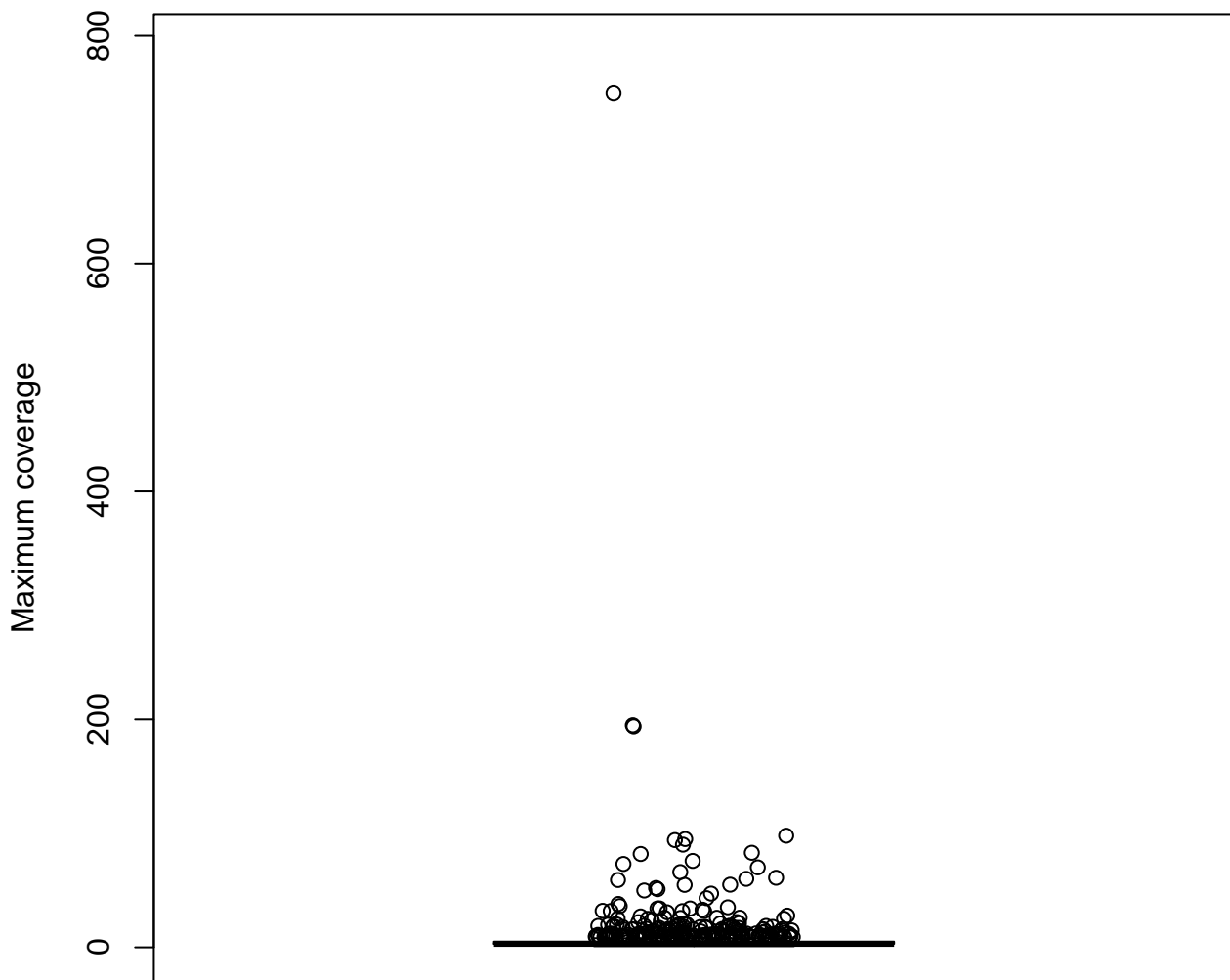
**jx in Intropolis but not "annotated" by knownGene:
24 samples; n jx: 3352**



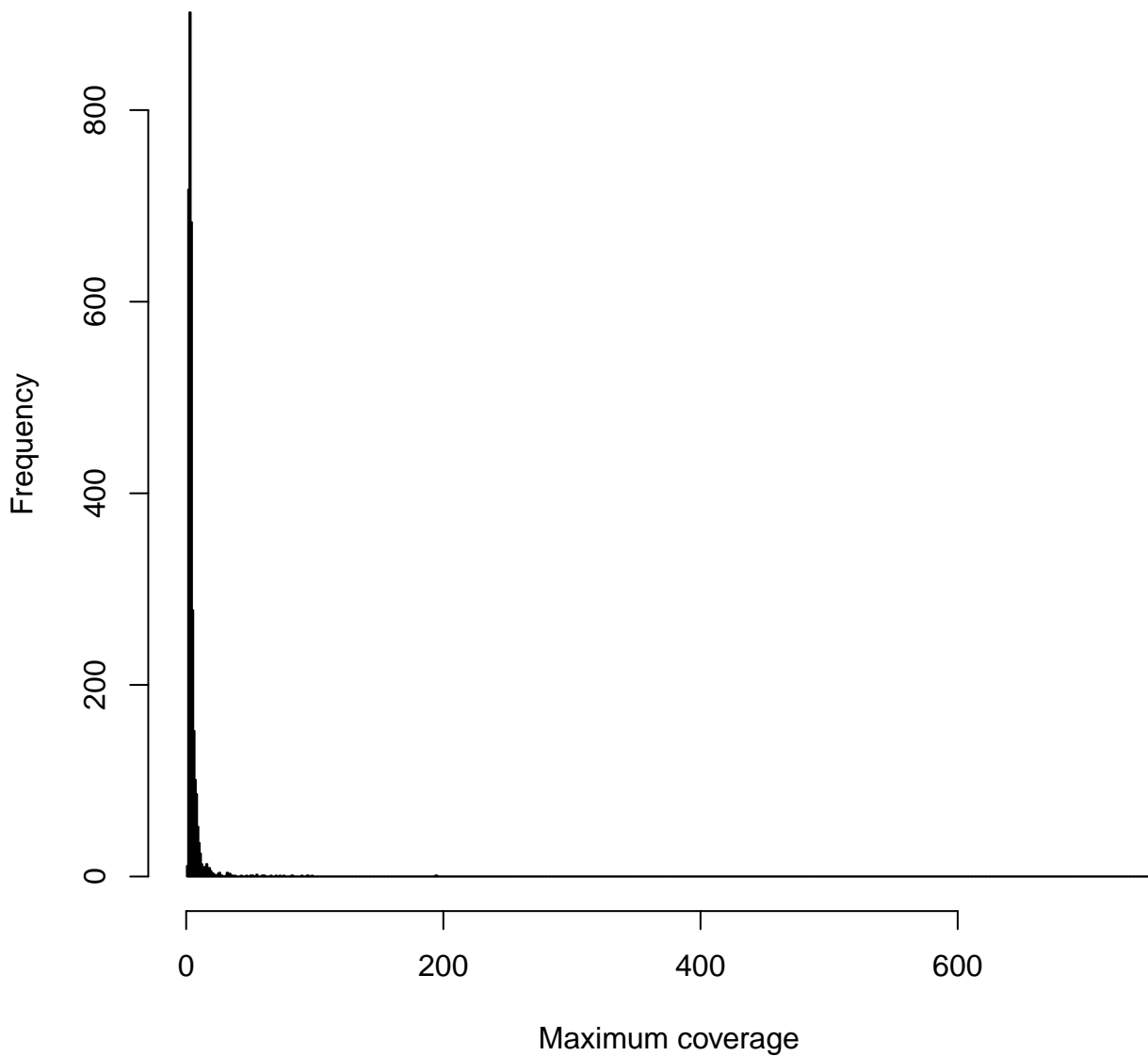
**jx in Intropolis but not "annotated" by knownGene:
24 samples; n jx: 3352**



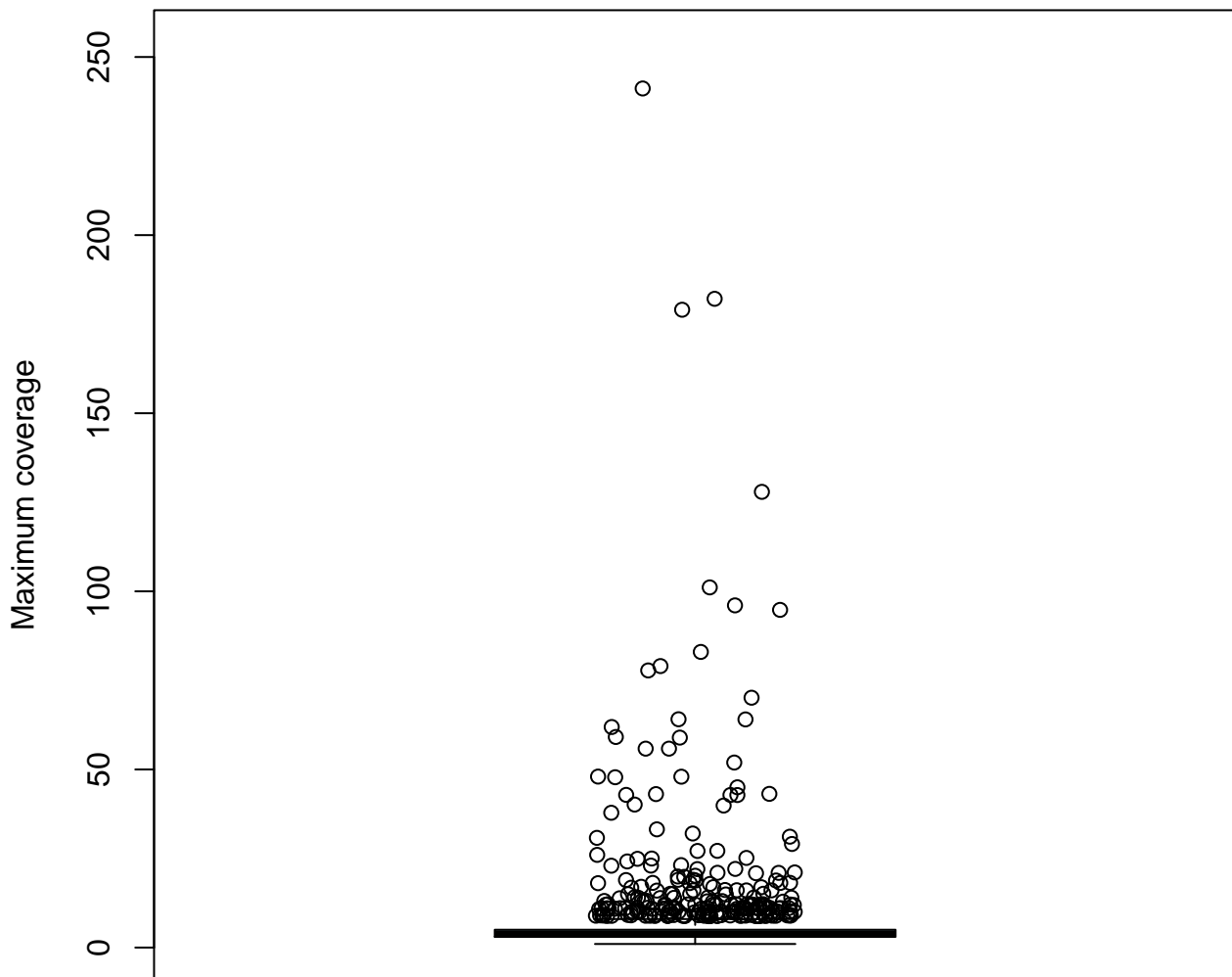
**jx in Intropolis but not "annotated" by knownGene:
25 samples; n jx: 3158**



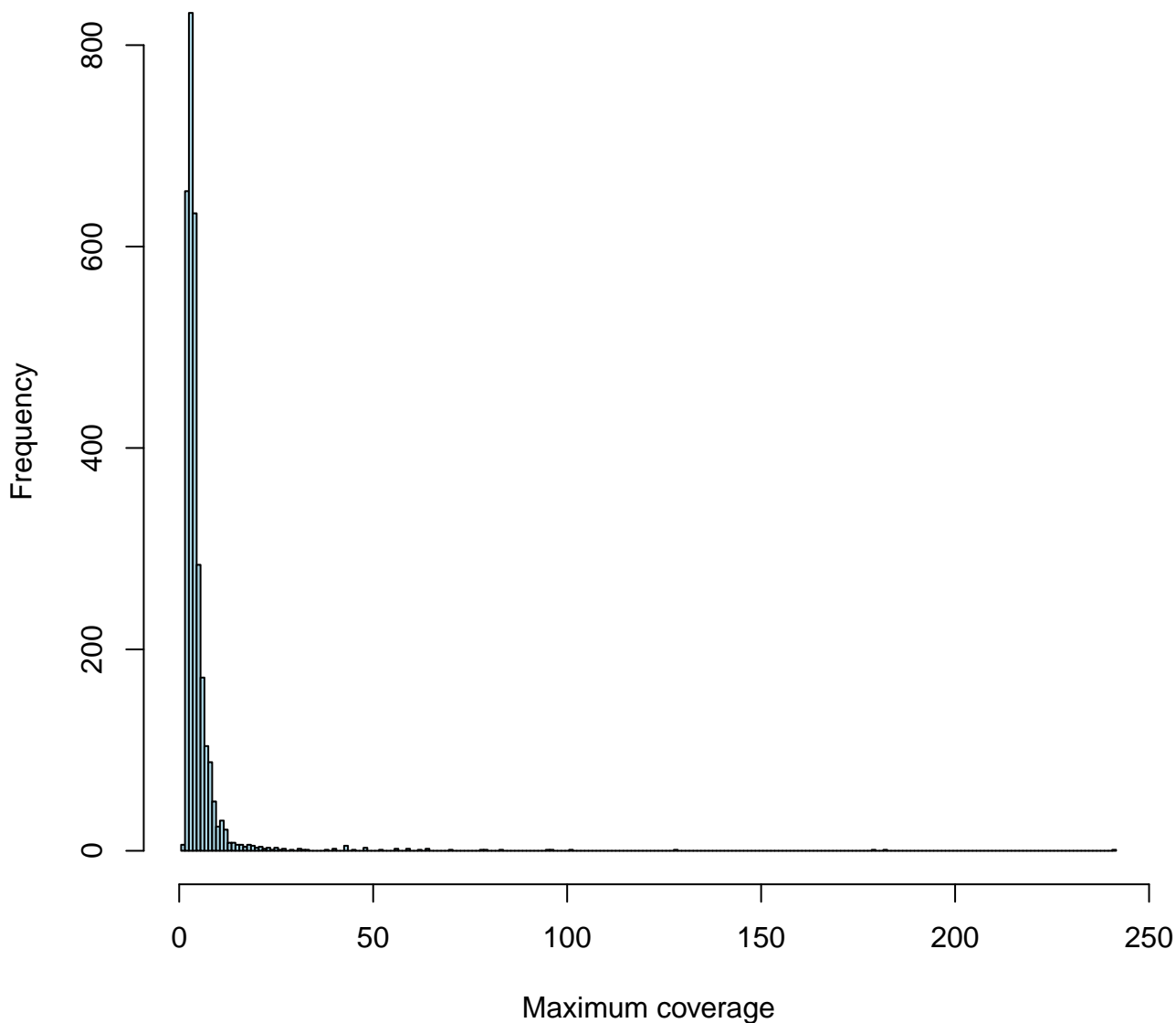
**jx in Intropolis but not "annotated" by knownGene:
25 samples; n jx: 3158**



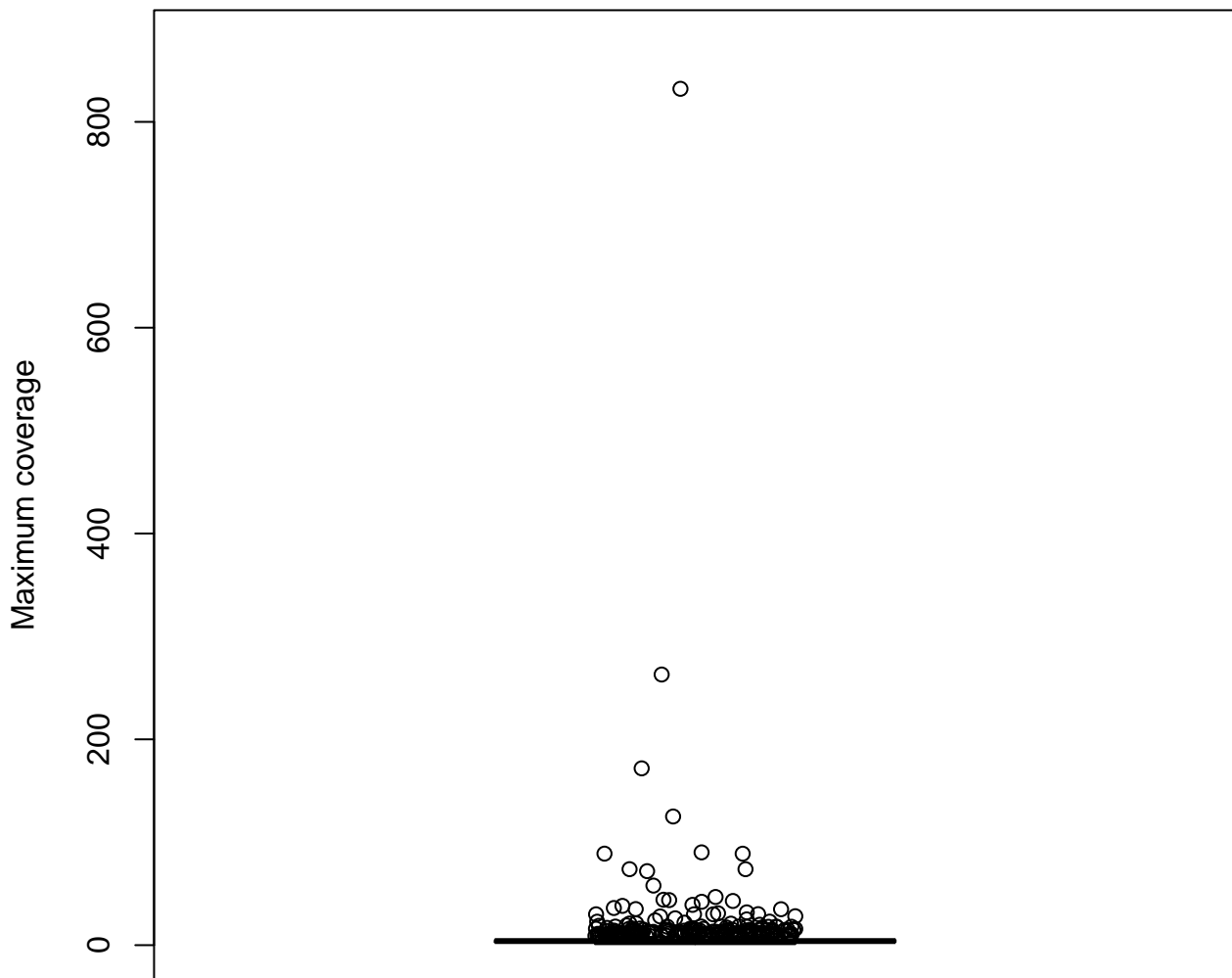
**jx in Intropolis but not "annotated" by knownGene:
26 samples; n jx: 2996**



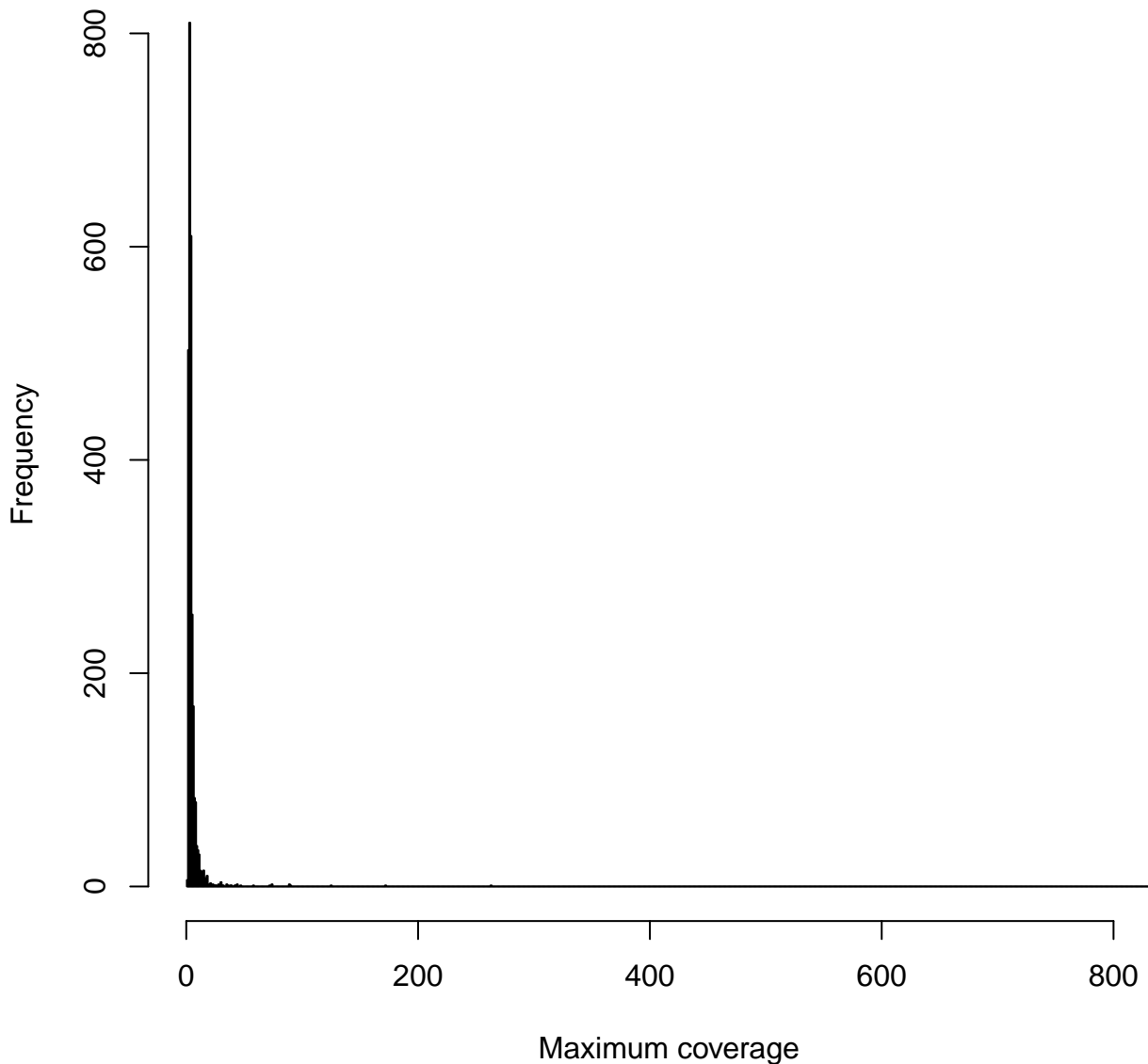
**jx in Intropolis but not "annotated" by knownGene:
26 samples; n jx: 2996**



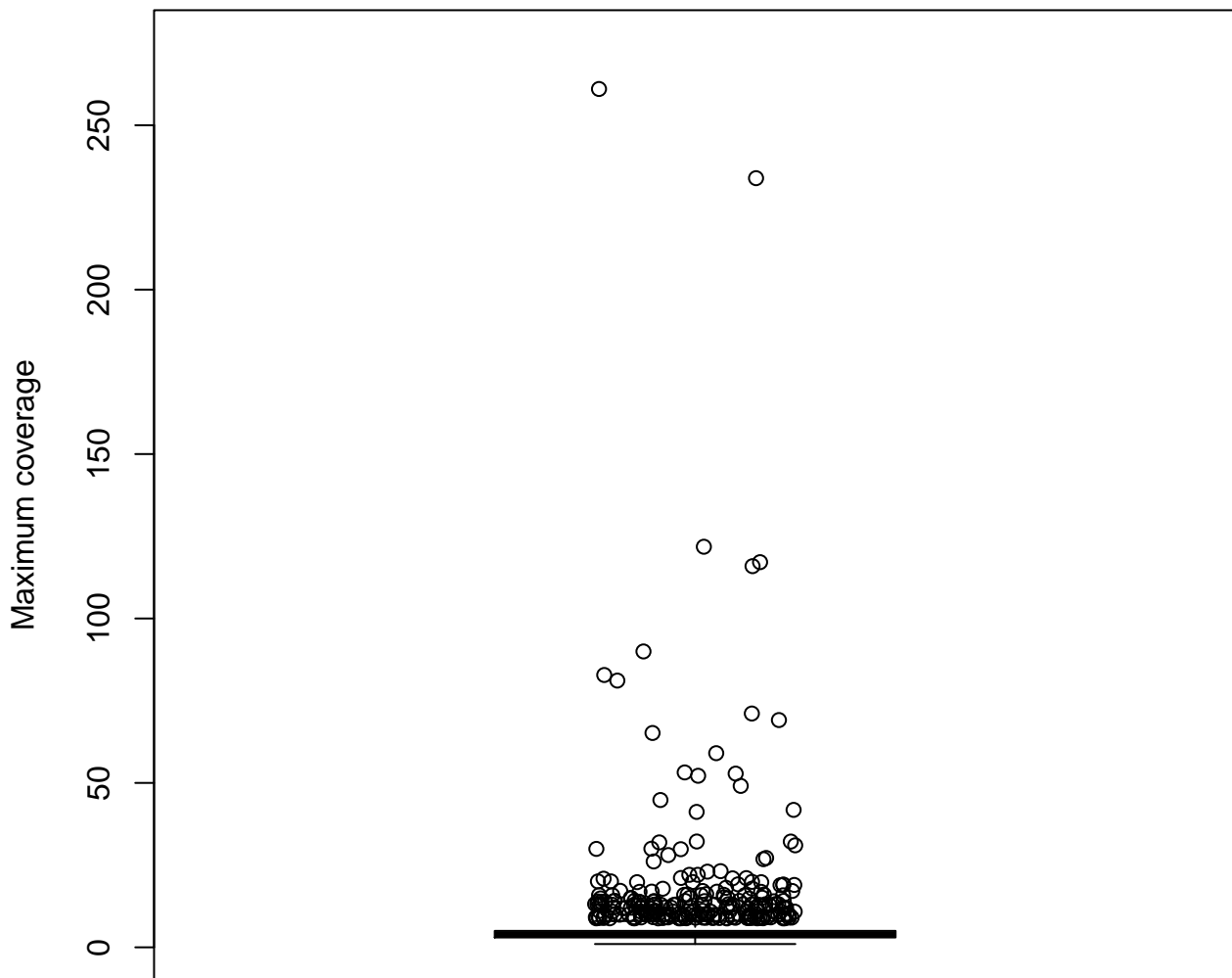
**jx in Intropolis but not "annotated" by knownGene:
27 samples; n jx: 2734**



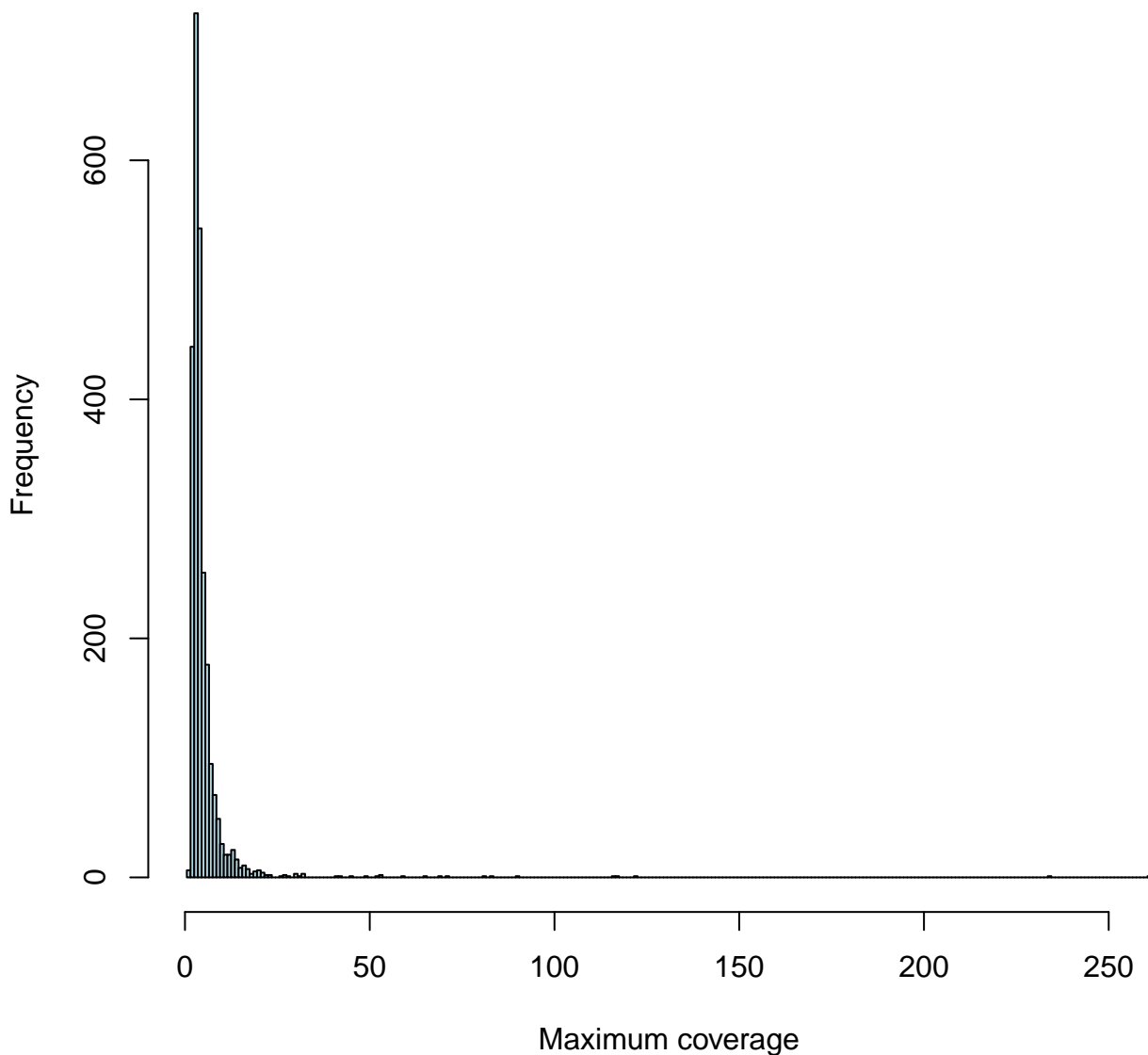
**jx in Intropolis but not "annotated" by knownGene:
27 samples; n jx: 2734**



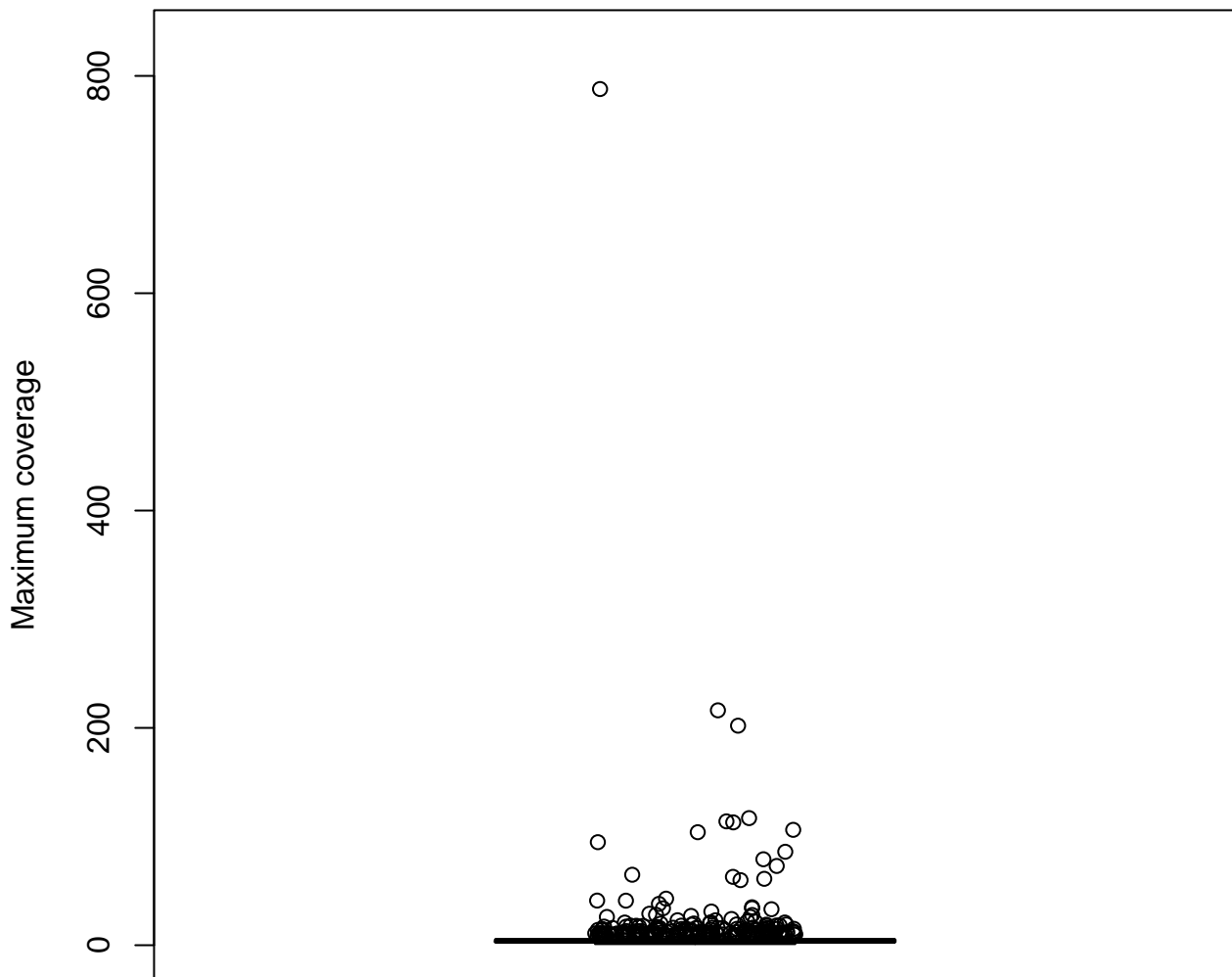
**jx in Intropolis but not "annotated" by knownGene:
28 samples; n jx: 2543**



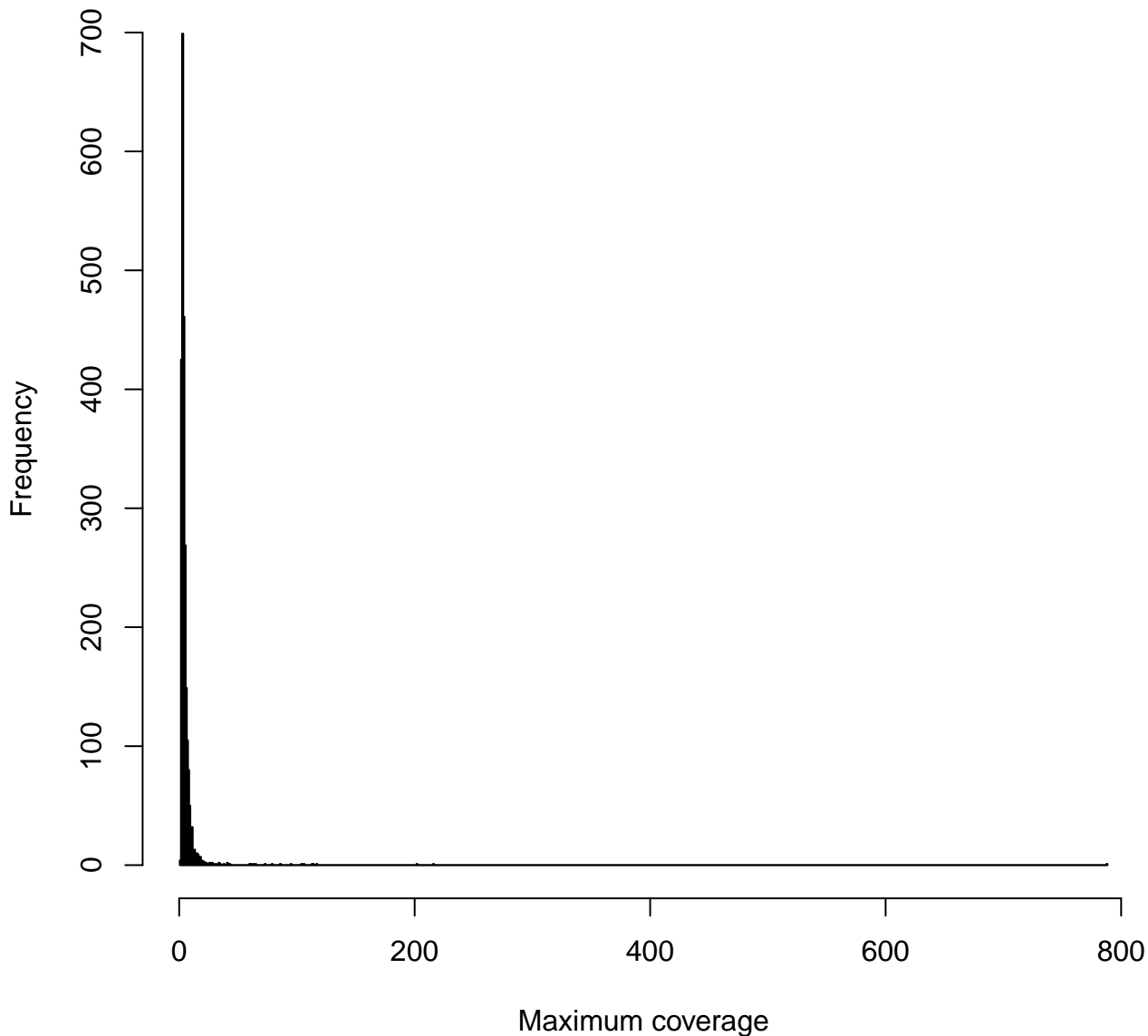
**jx in Intropolis but not "annotated" by knownGene:
28 samples; n jx: 2543**



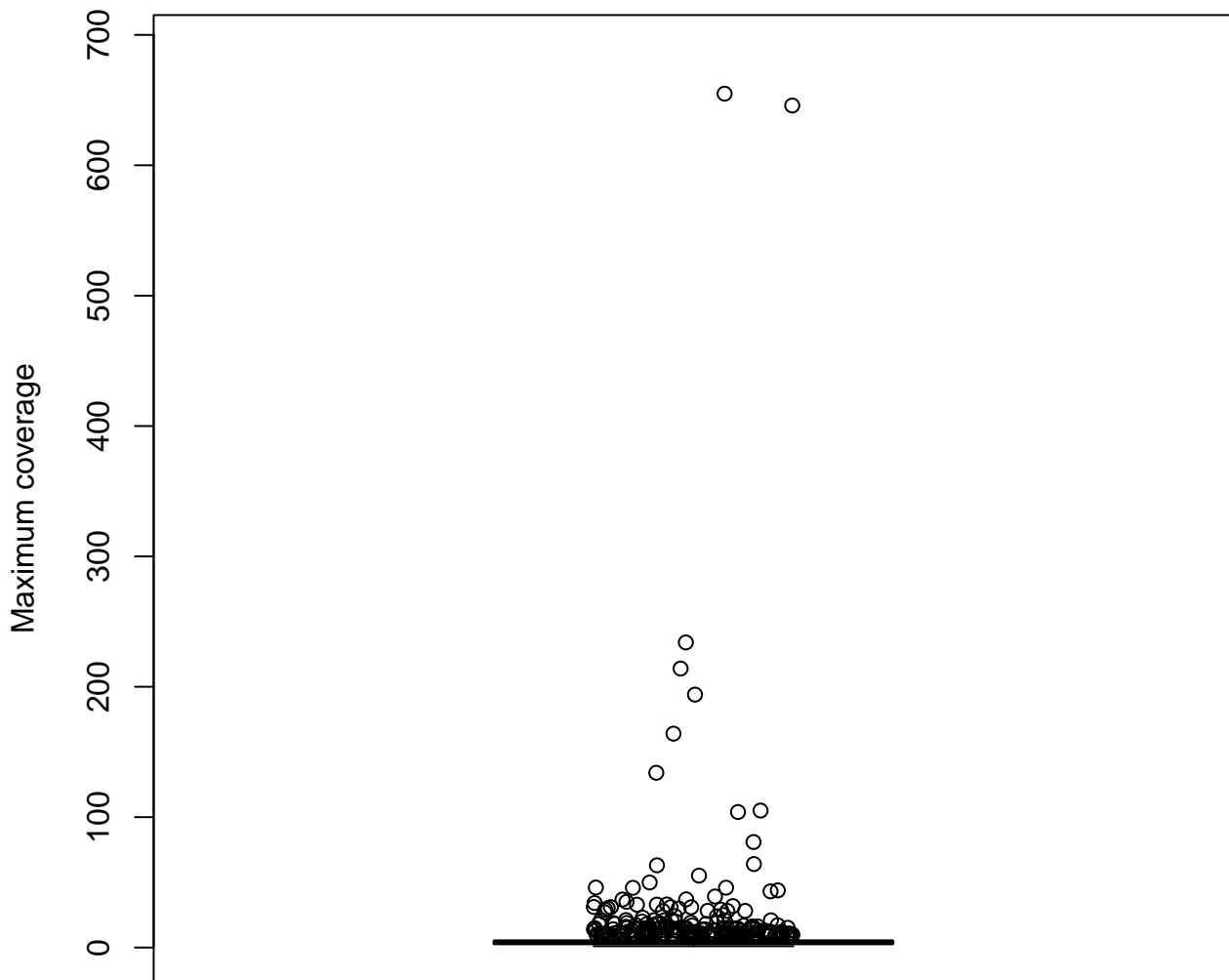
**jx in Intropolis but not "annotated" by knownGene:
29 samples; n jx: 2413**



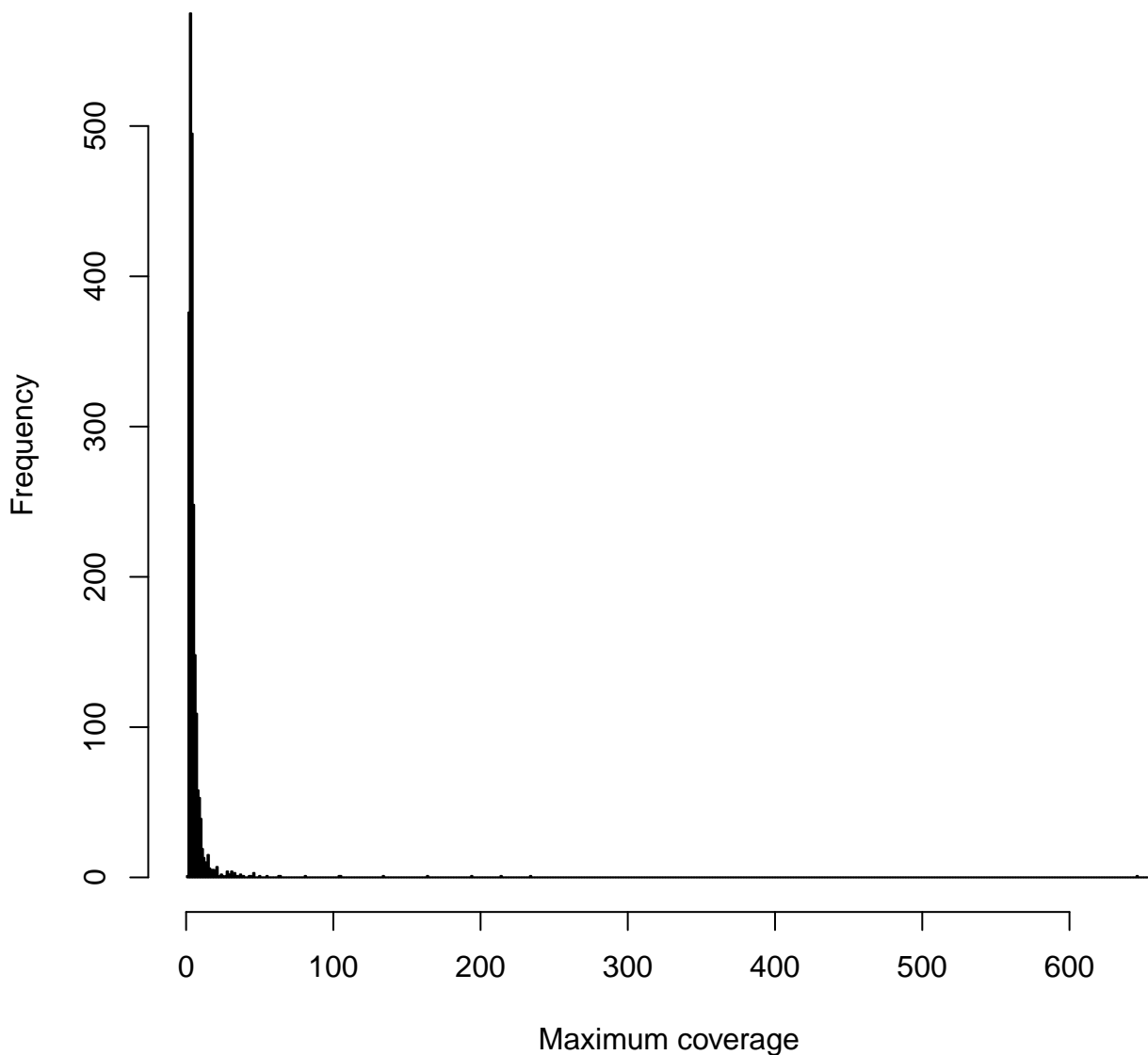
**jx in Intropolis but not "annotated" by knownGene:
29 samples; n jx: 2413**



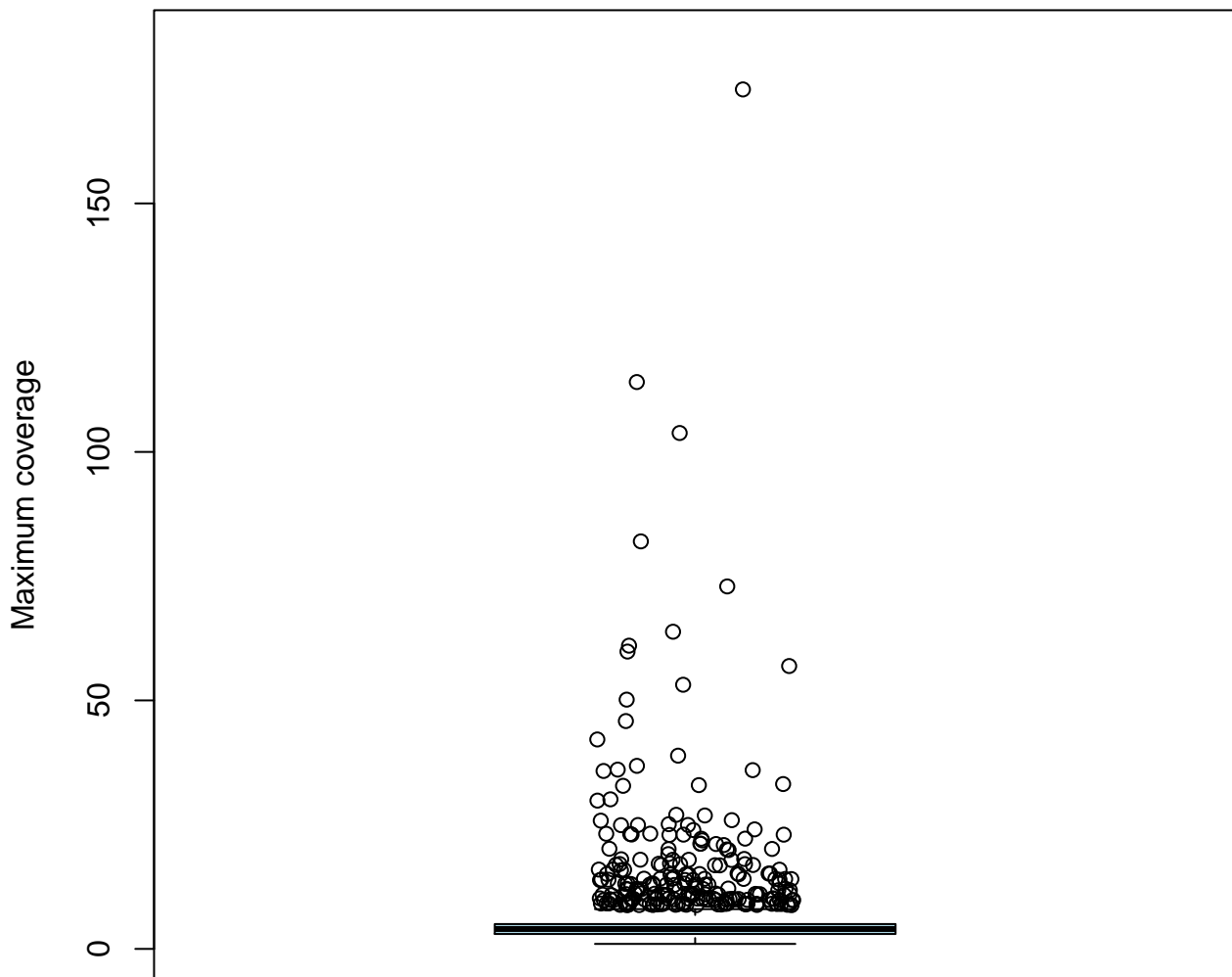
**jx in Intropolis but not "annotated" by knownGene:
30 samples; n jx: 2244**



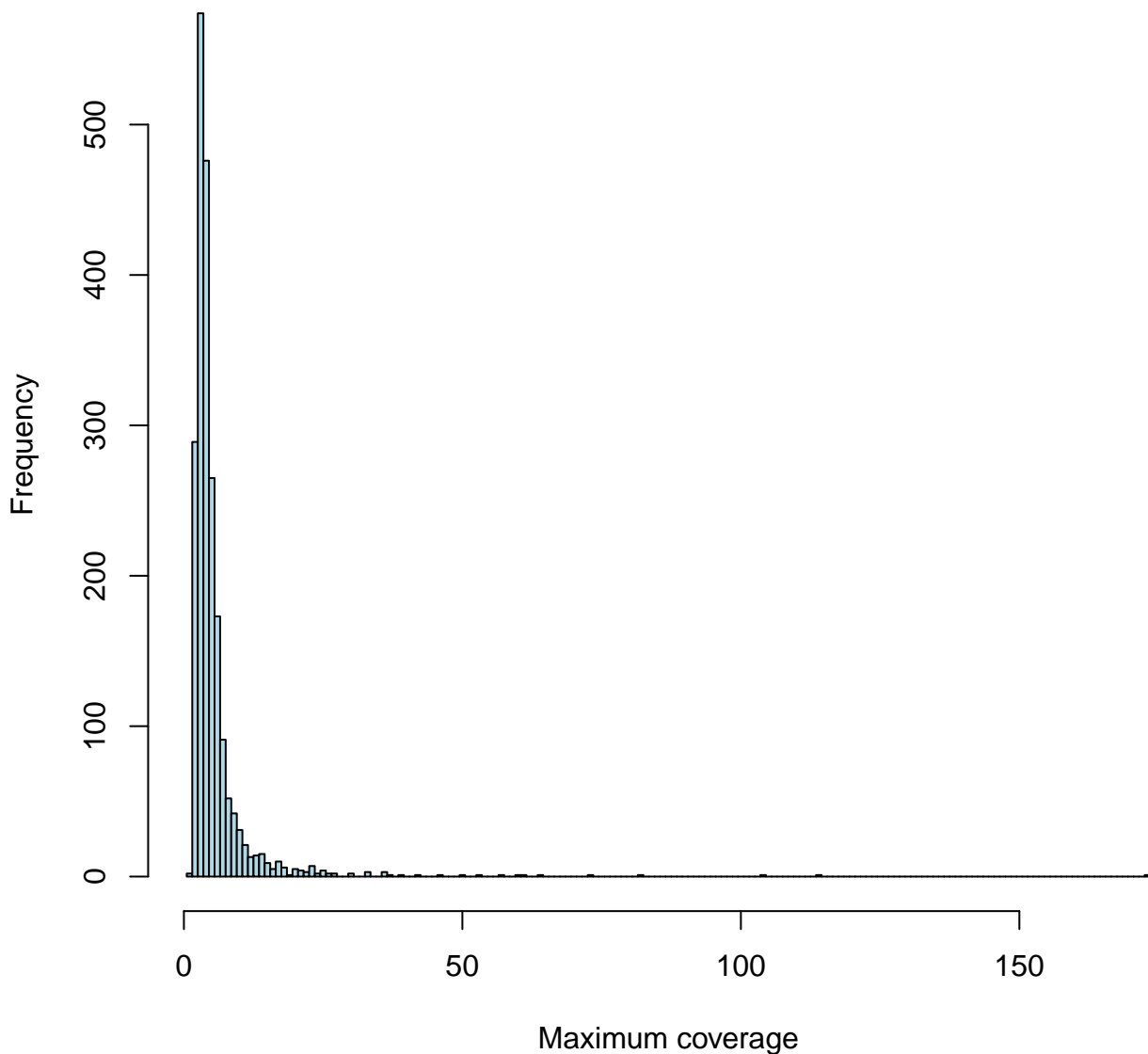
**jx in Intropolis but not "annotated" by knownGene:
30 samples; n jx: 2244**



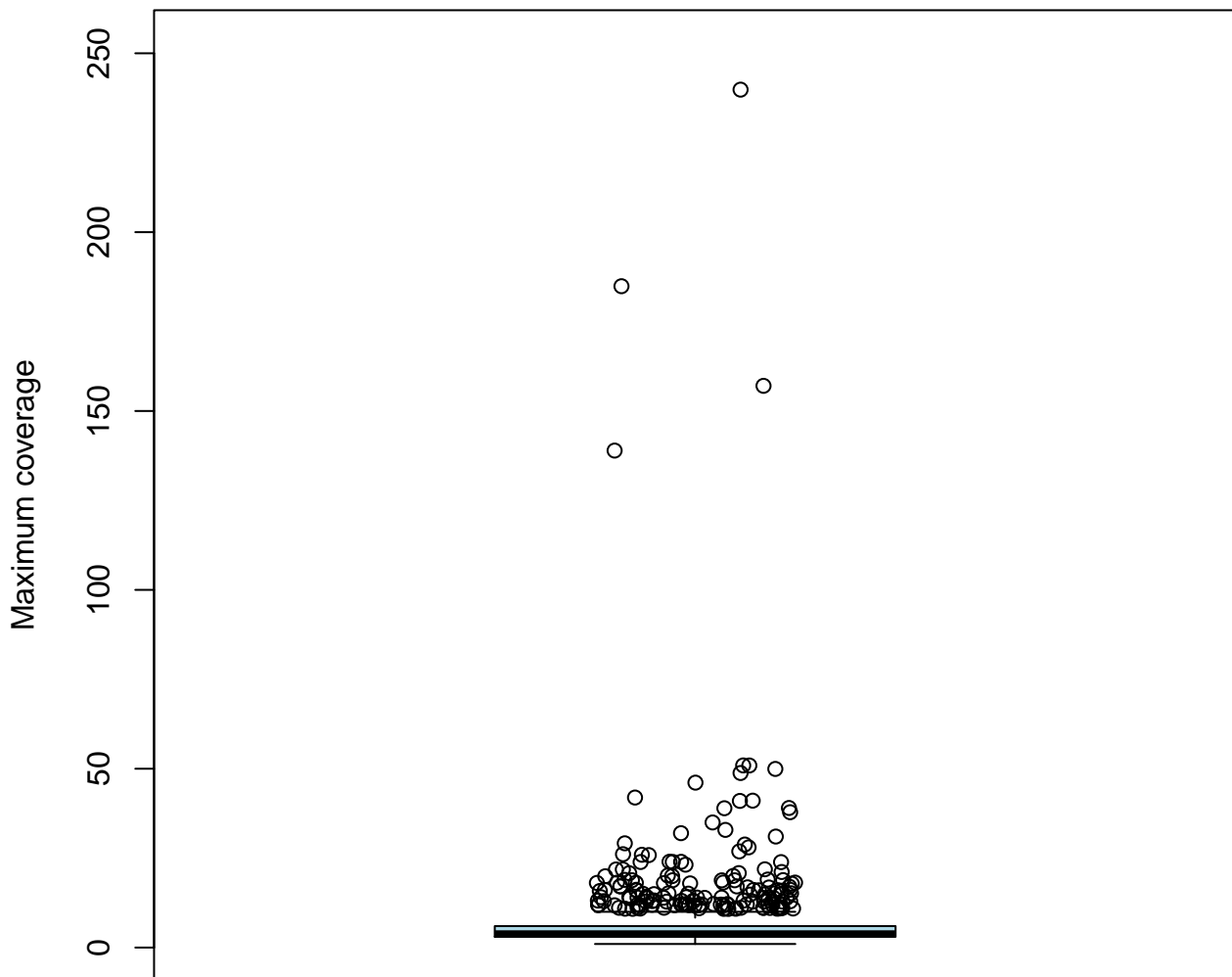
**jx in Intropolis but not "annotated" by knownGene:
31 samples; n jx: 2141**



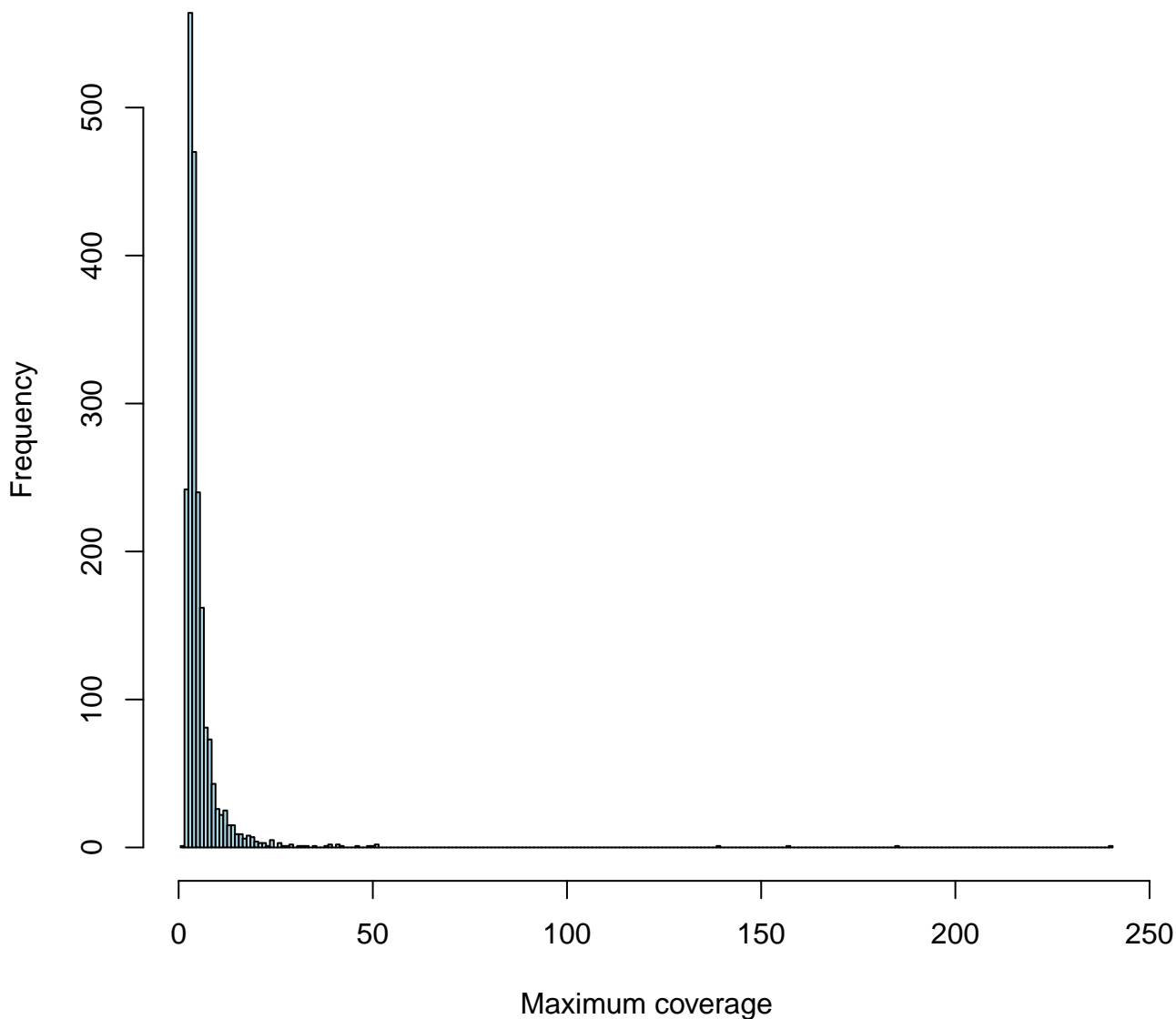
**jx in Intropolis but not "annotated" by knownGene:
31 samples; n jx: 2141**



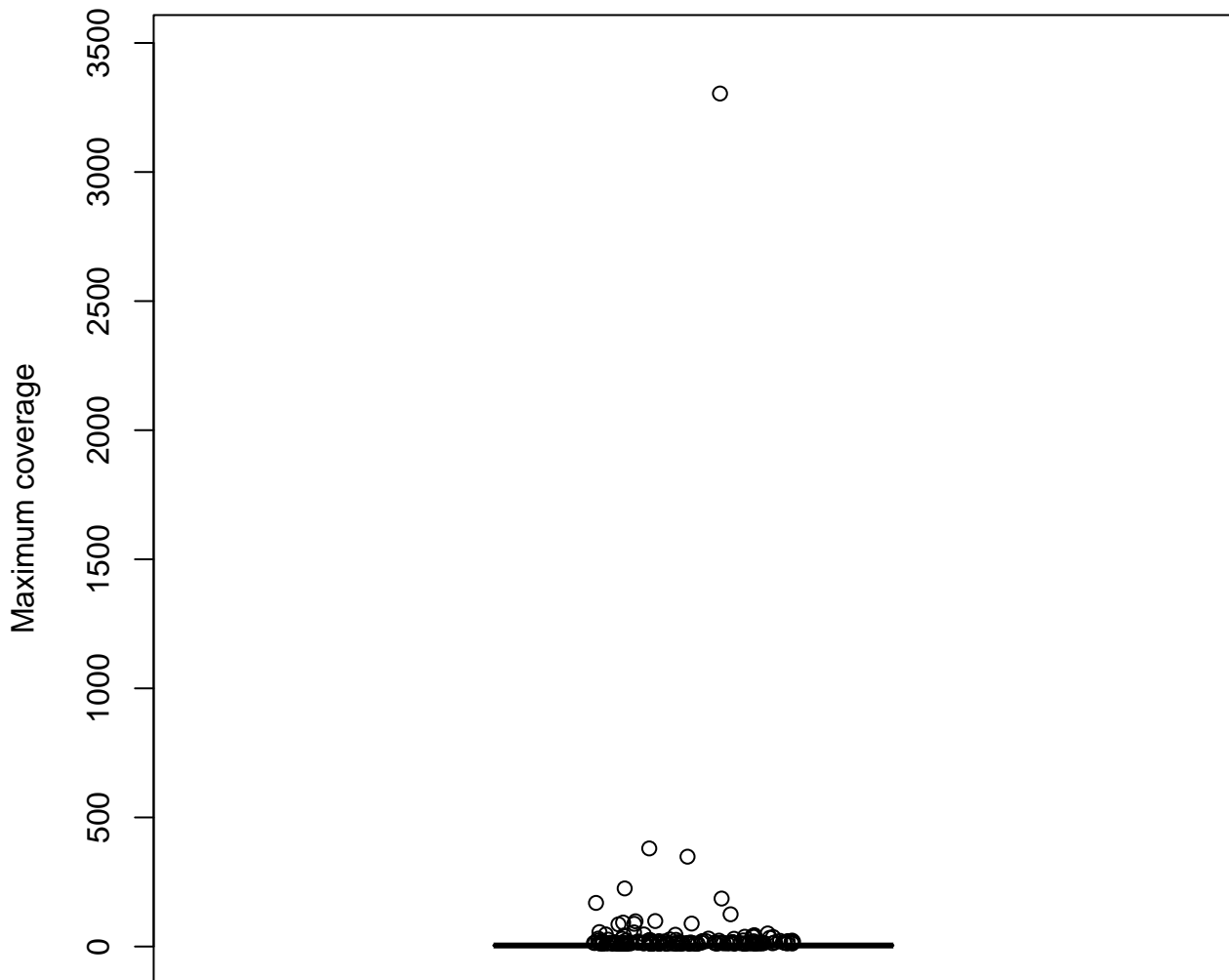
**jx in Intropolis but not "annotated" by knownGene:
32 samples; n jx: 2060**



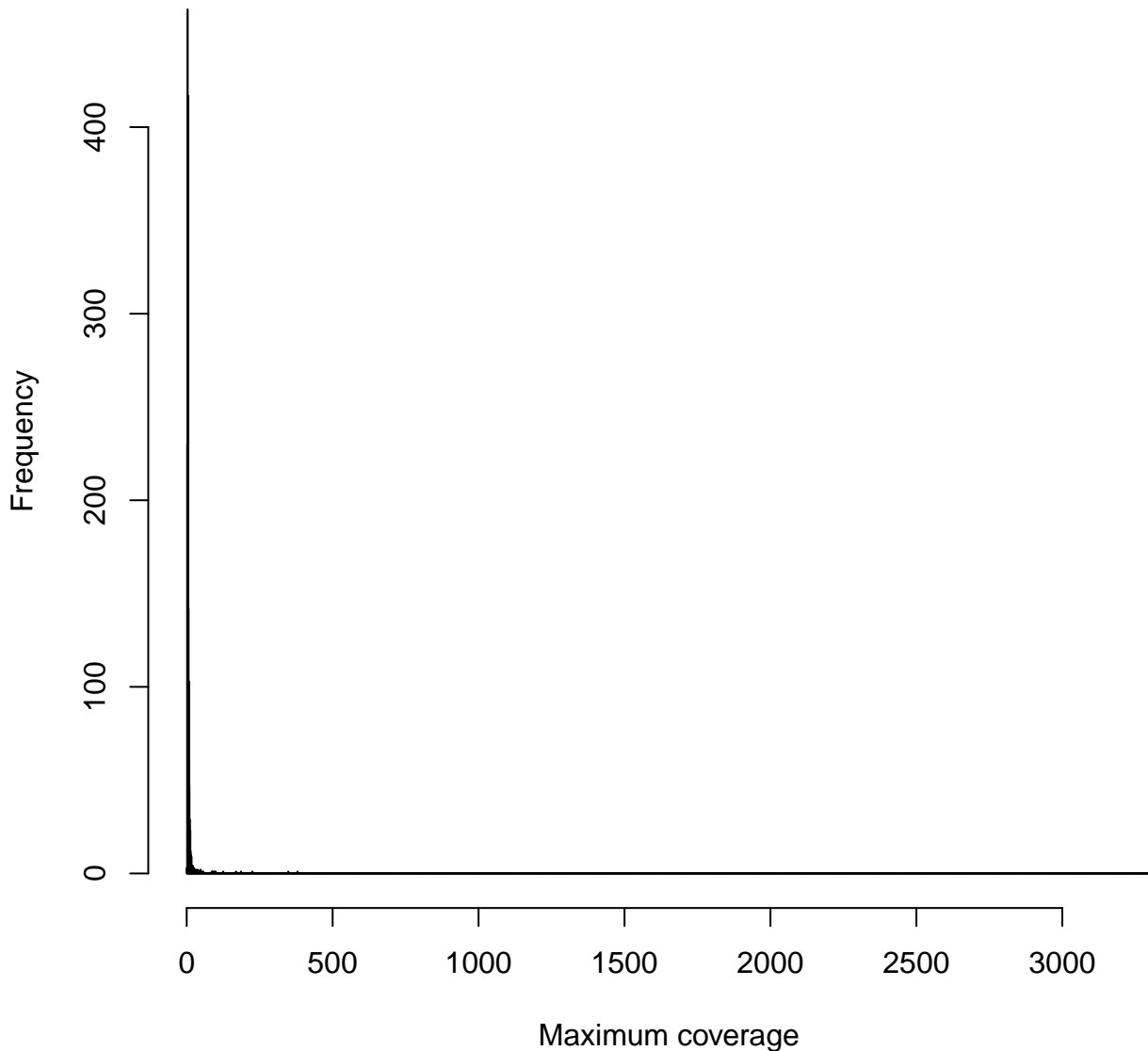
**jx in Intropolis but not "annotated" by knownGene:
32 samples; n jx: 2060**



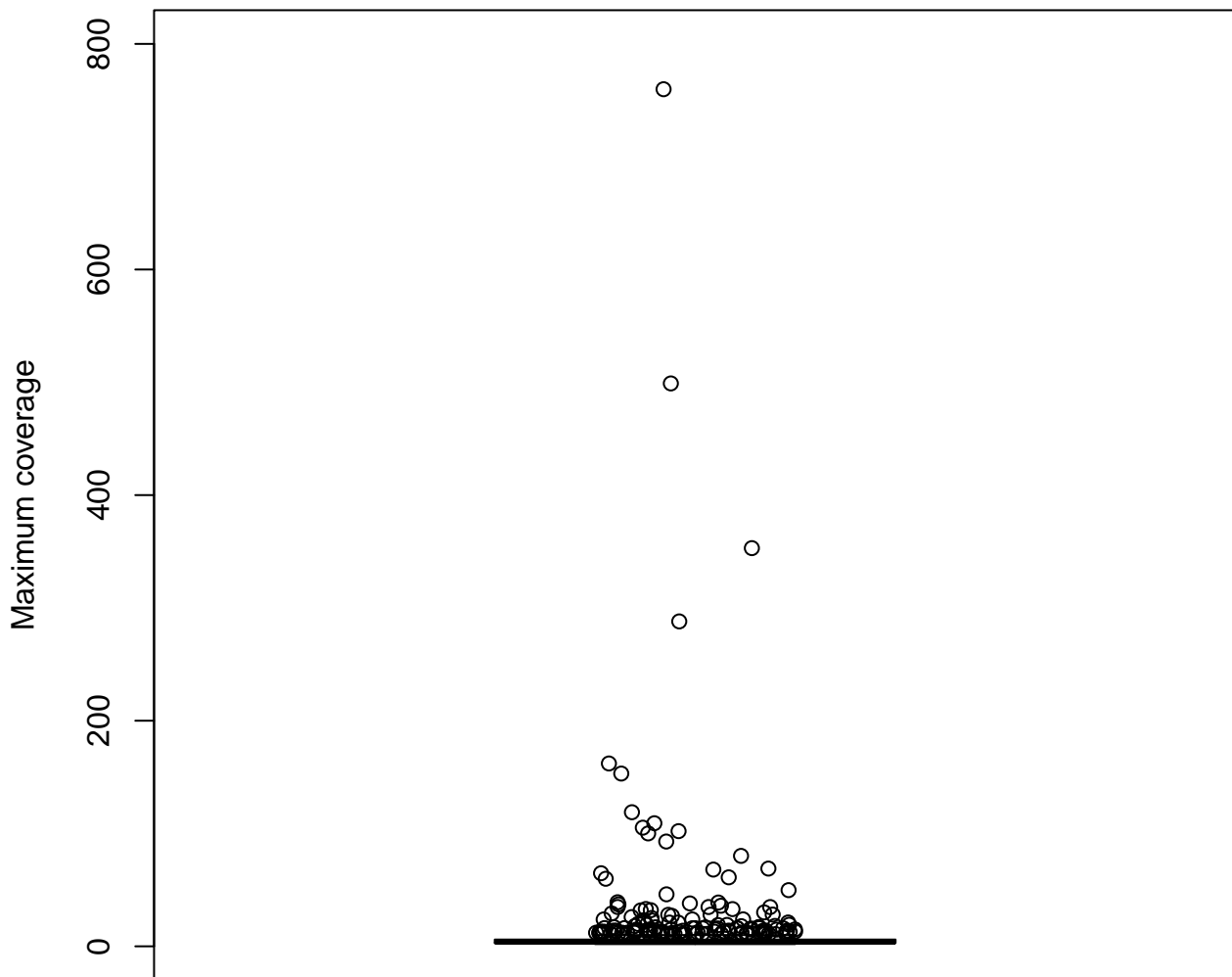
**jx in Intropolis but not "annotated" by knownGene:
33 samples; n jx: 1839**



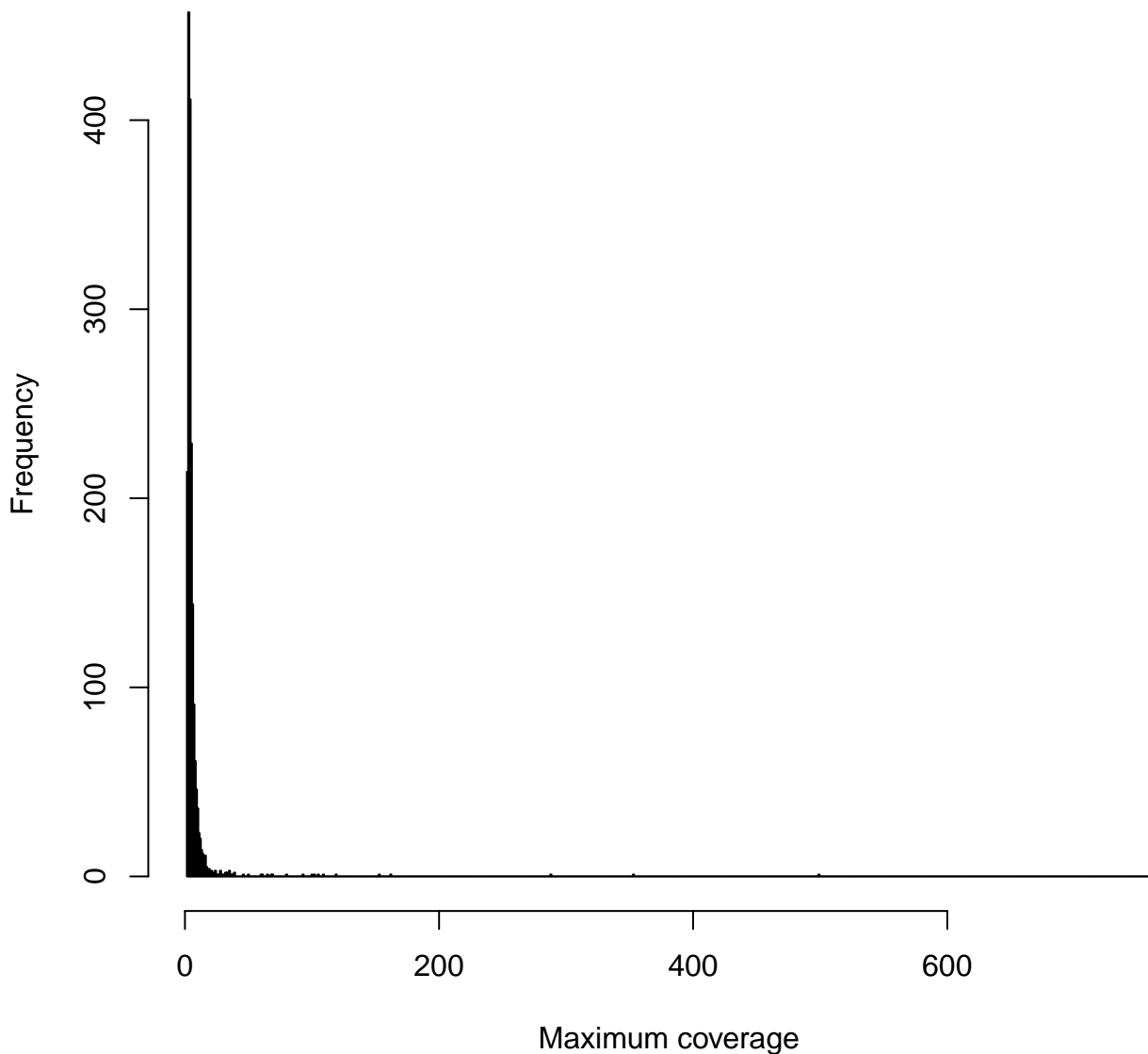
**jx in Intropolis but not "annotated" by knownGene:
33 samples; n jx: 1839**



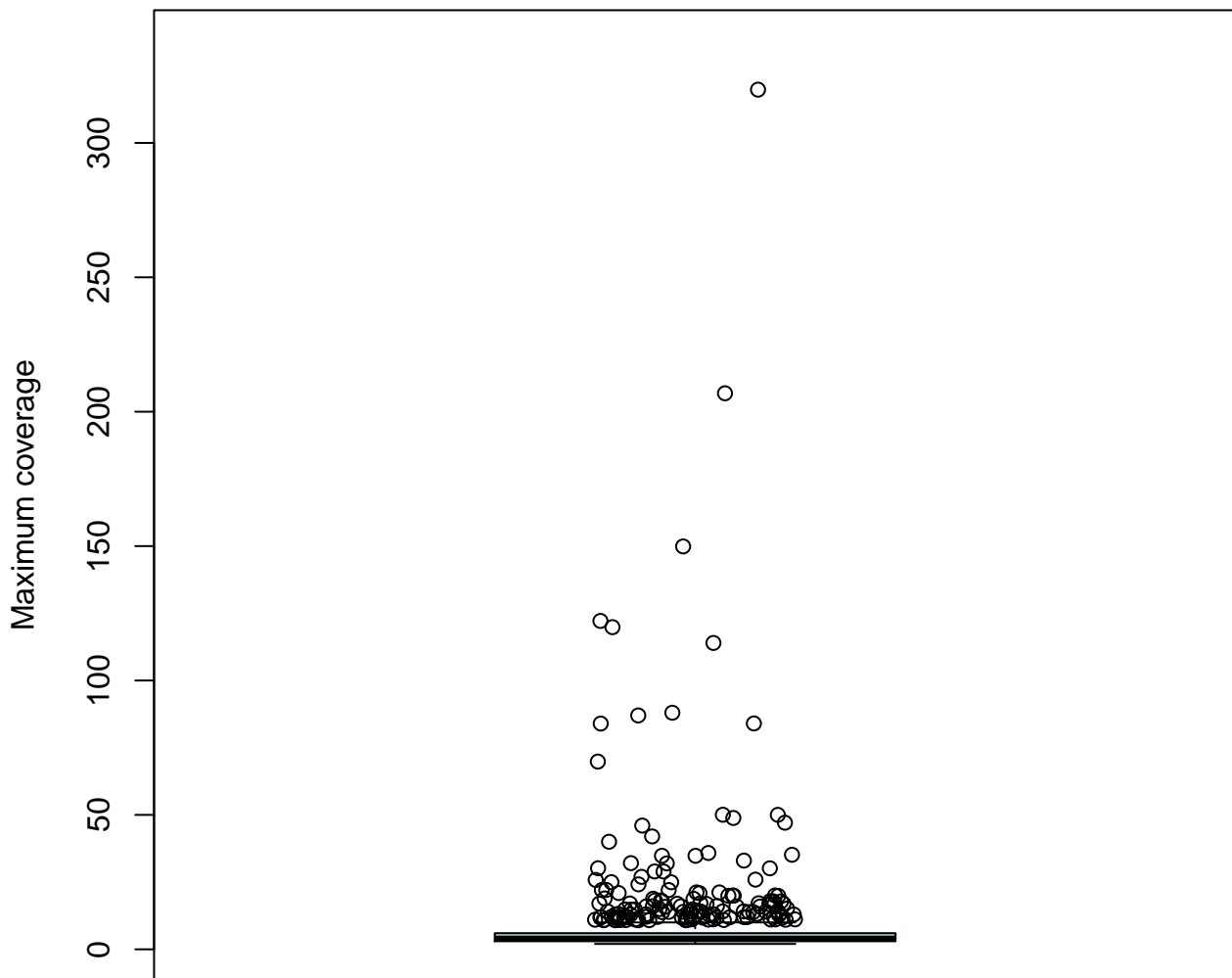
**jx in Intropolis but not "annotated" by knownGene:
34 samples; n jx: 1838**



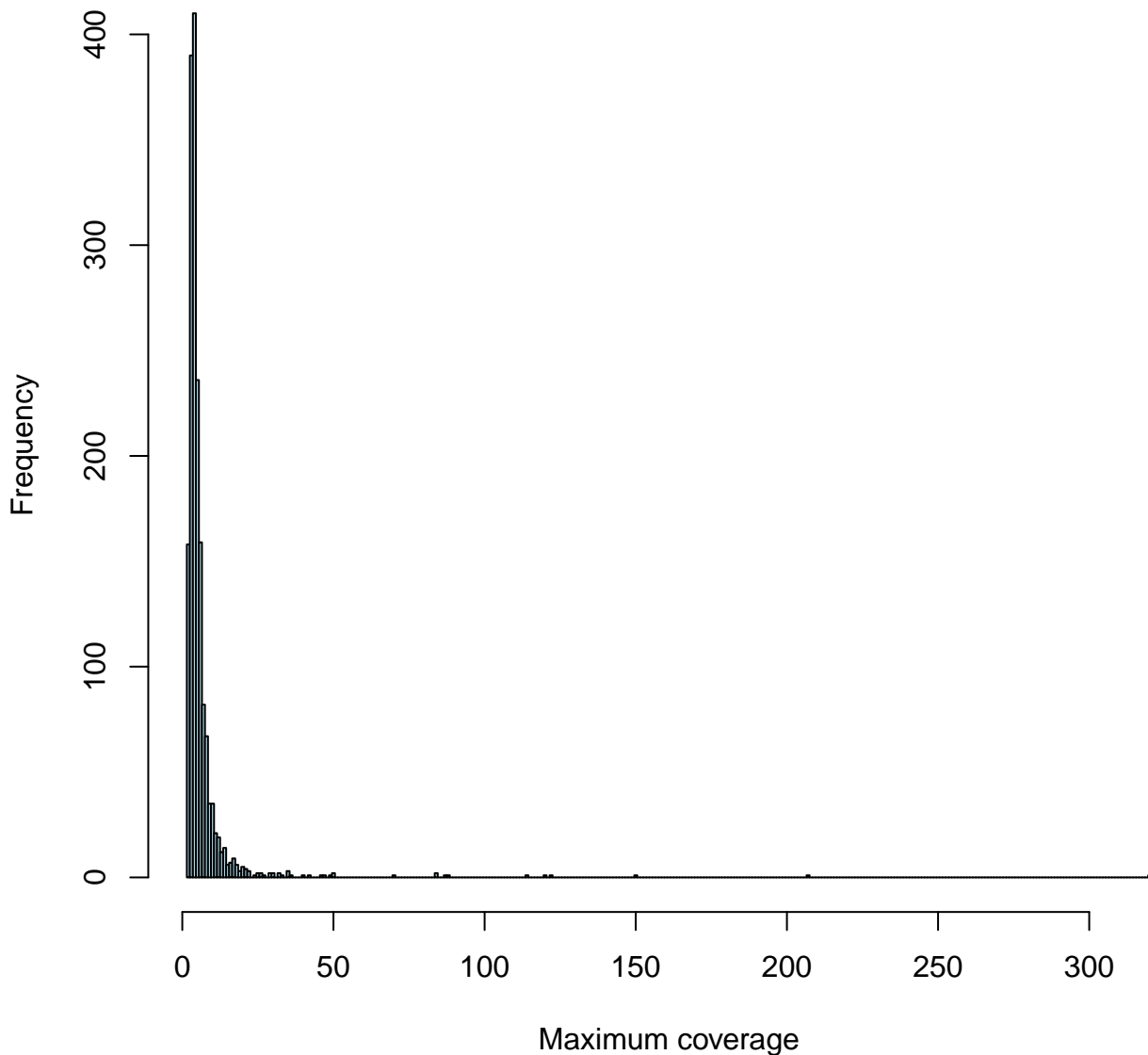
**jx in Intropolis but not "annotated" by knownGene:
34 samples; n jx: 1838**



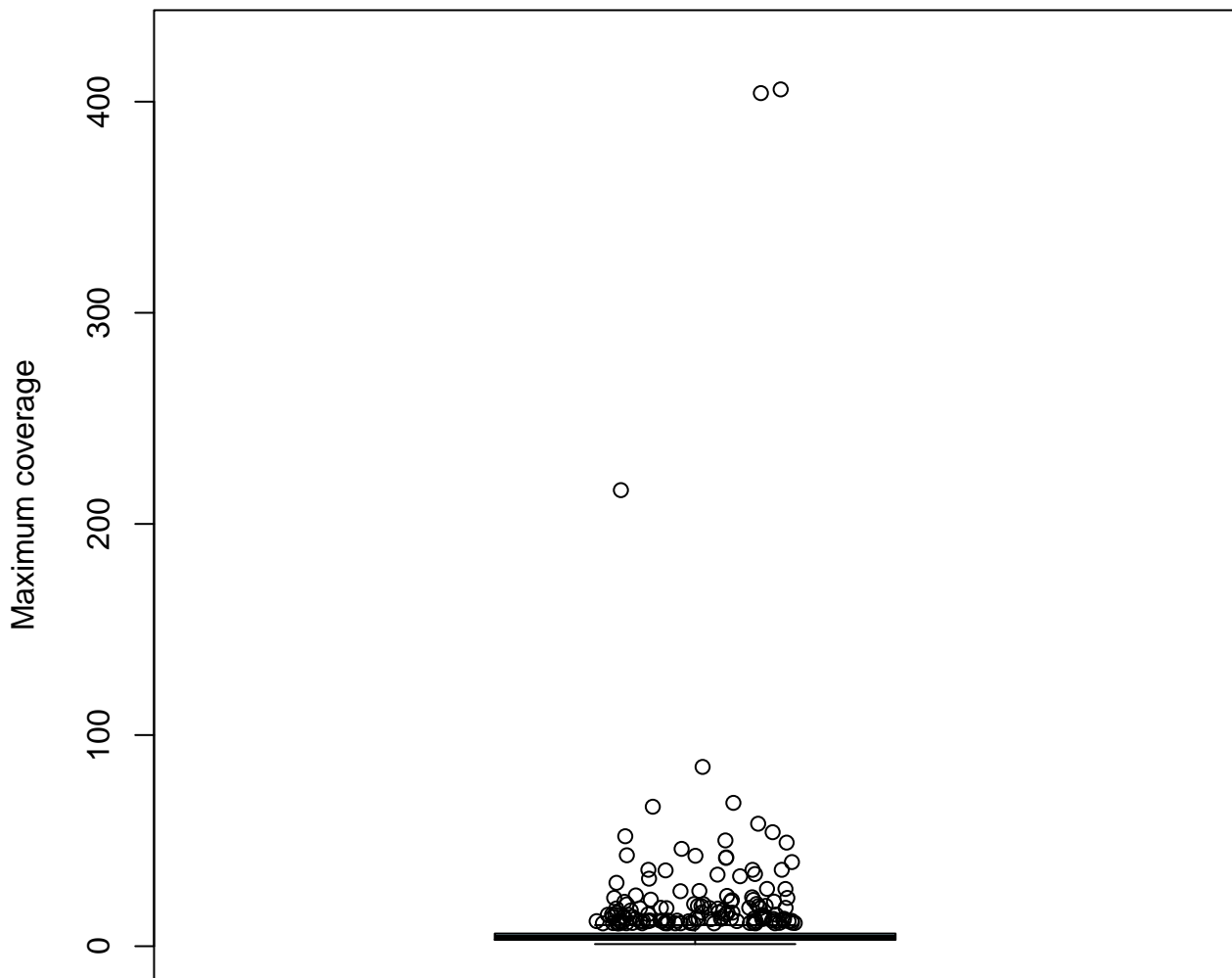
**jx in Intropolis but not "annotated" by knownGene:
35 samples; n jx: 1716**



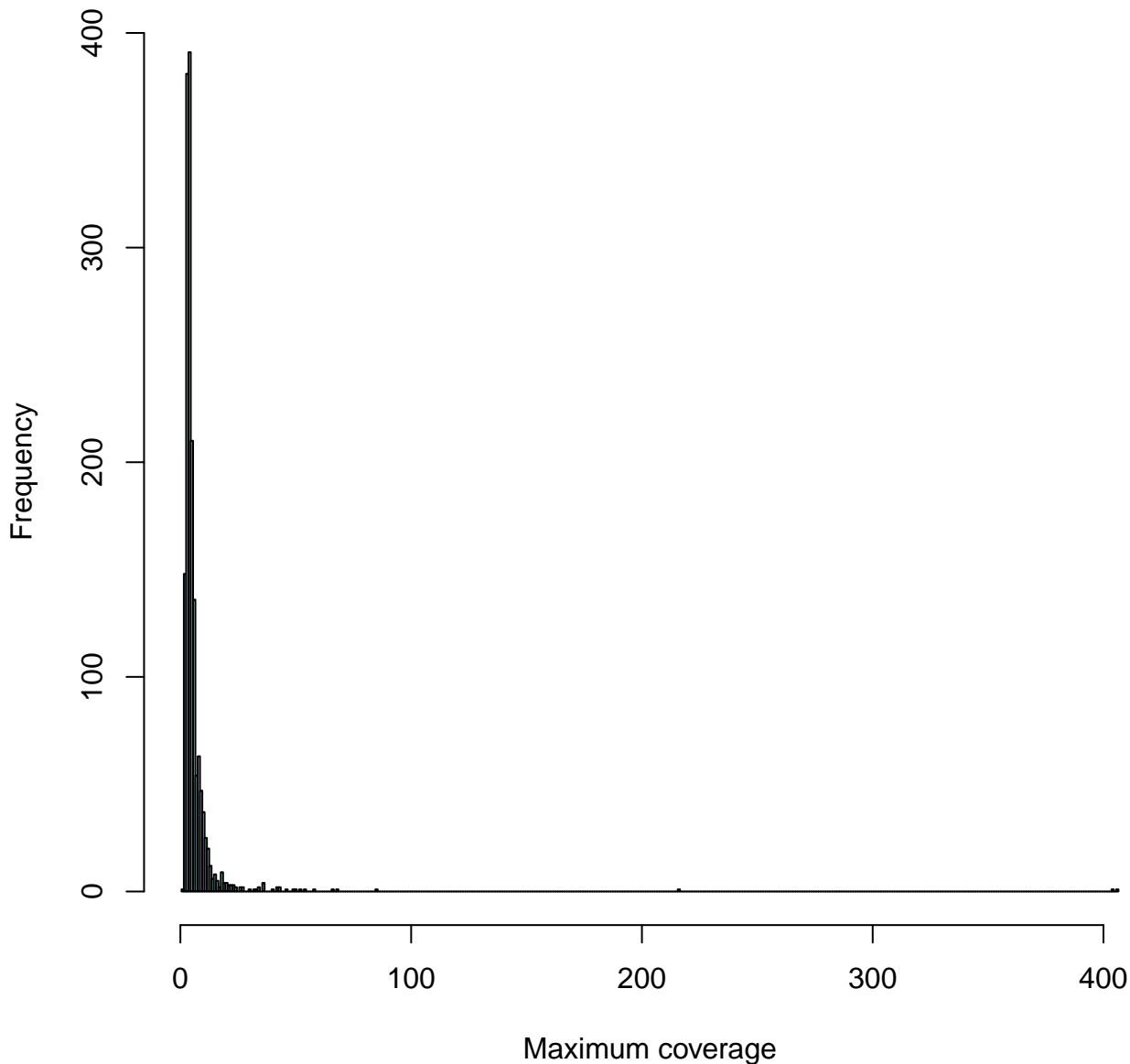
**jx in Intropolis but not "annotated" by knownGene:
35 samples; n jx: 1716**



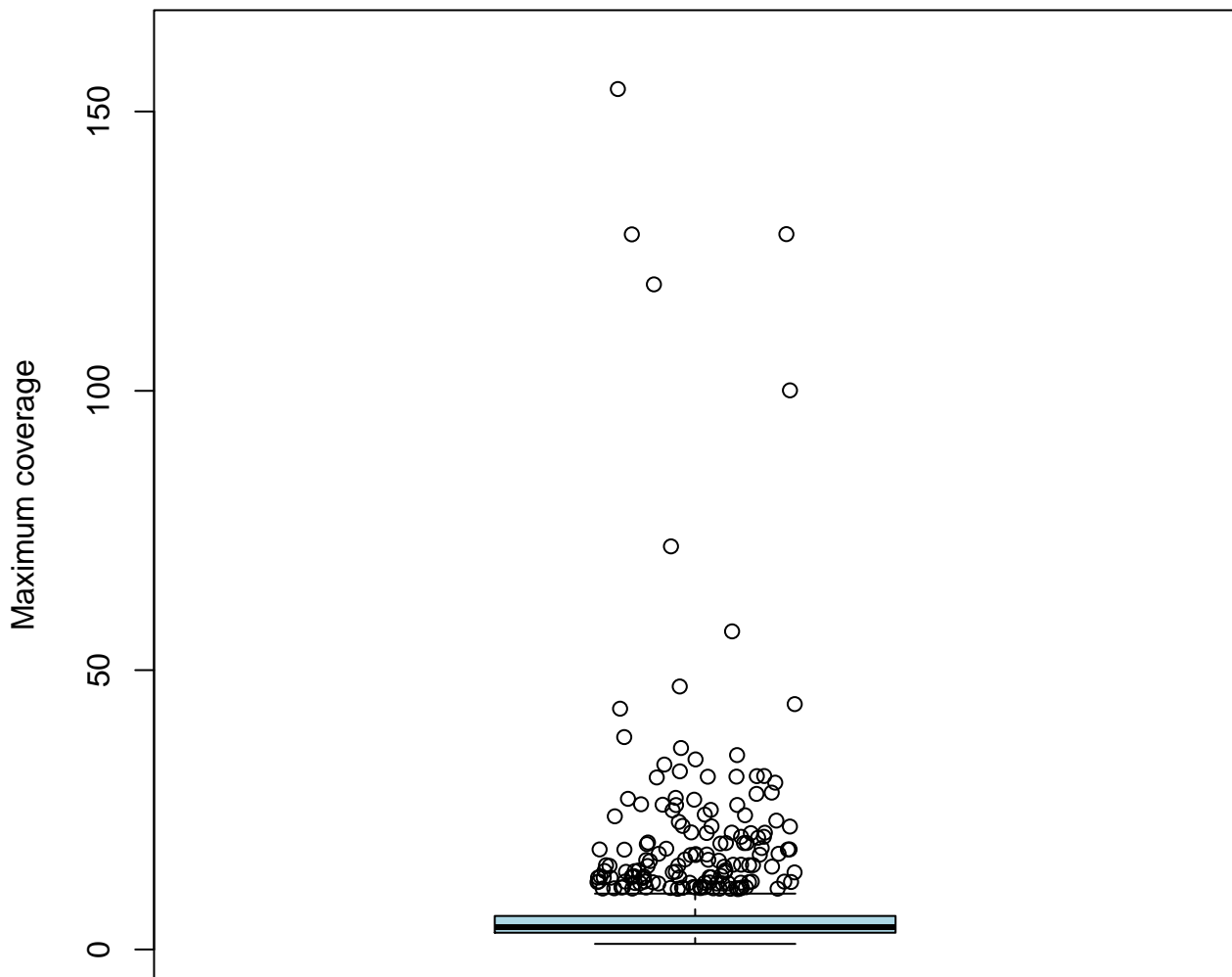
**jx in Intropolis but not "annotated" by knownGene:
36 samples; n jx: 1604**



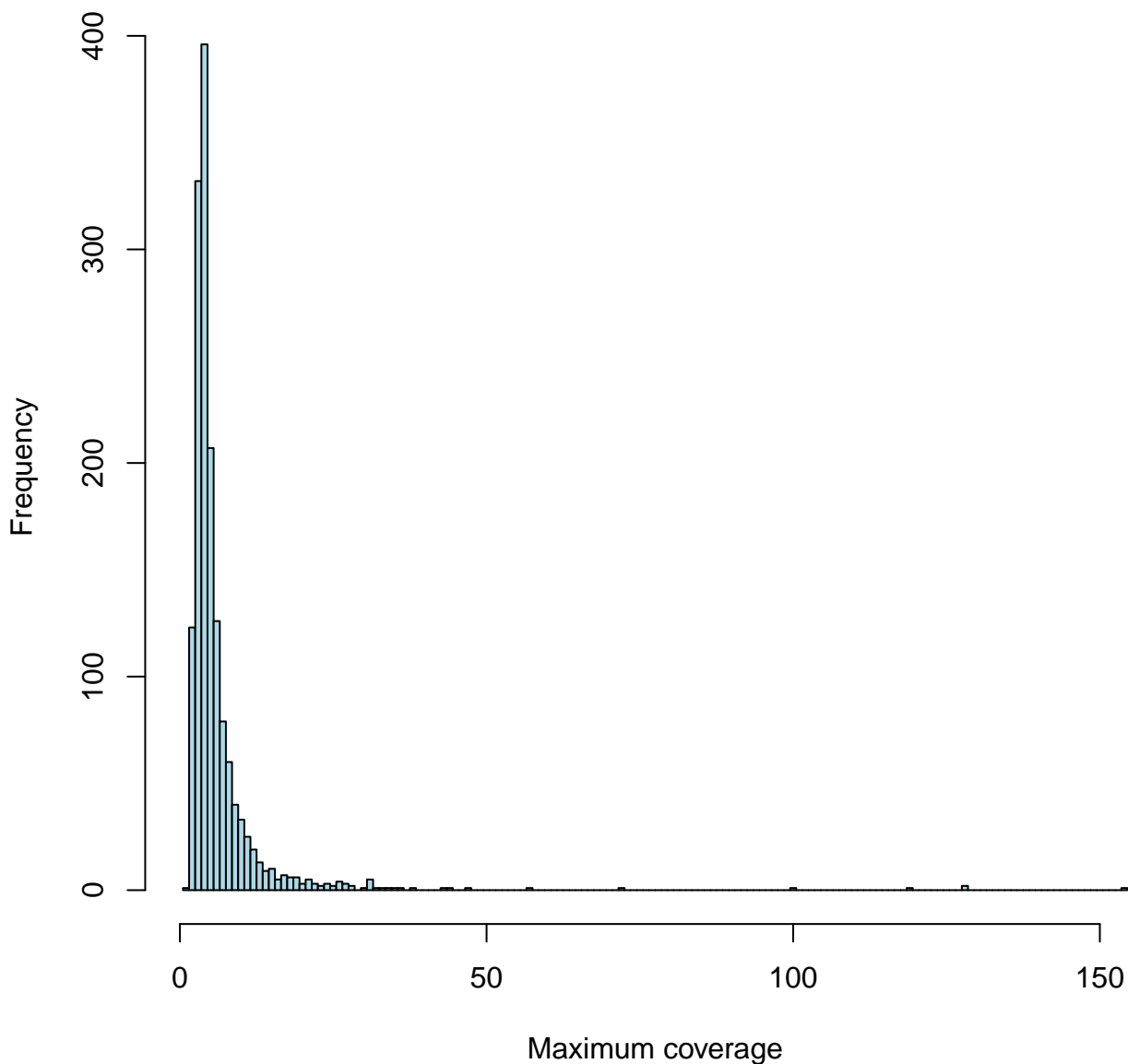
**jx in Intropolis but not "annotated" by knownGene:
36 samples; n jx: 1604**



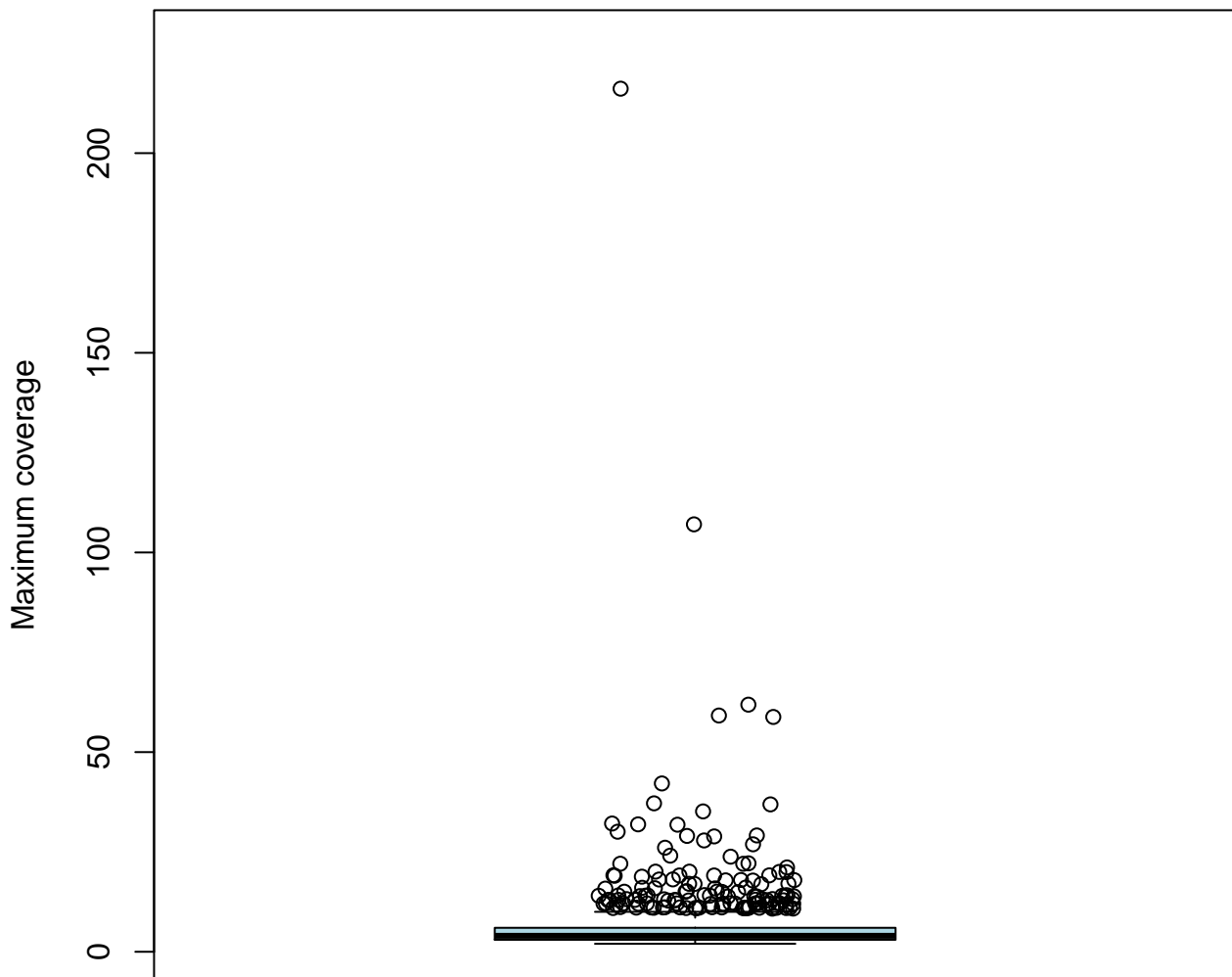
**jx in Intropolis but not "annotated" by knownGene:
37 samples; n jx: 1546**



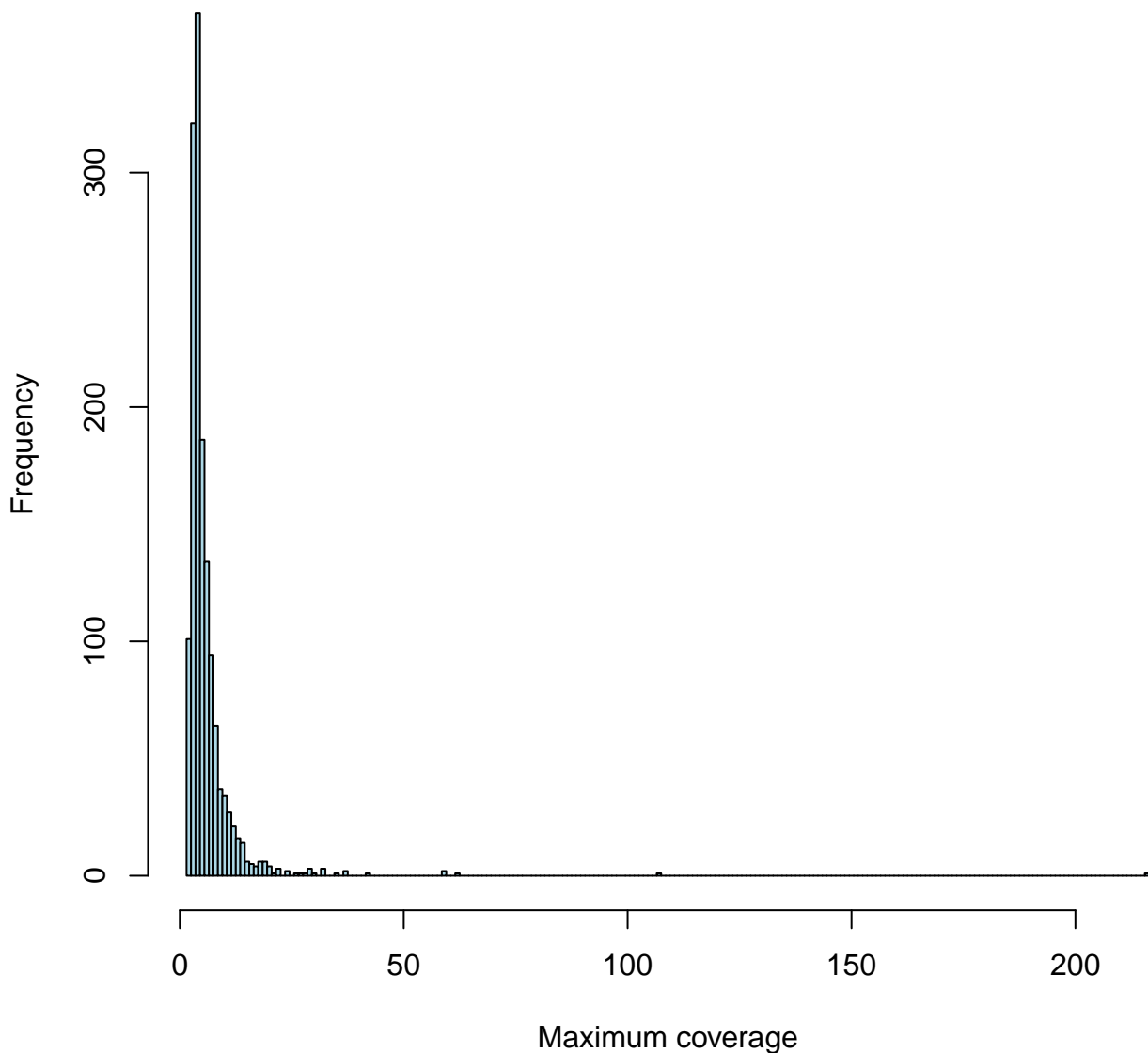
**jx in Intropolis but not "annotated" by knownGene:
37 samples; n jx: 1546**



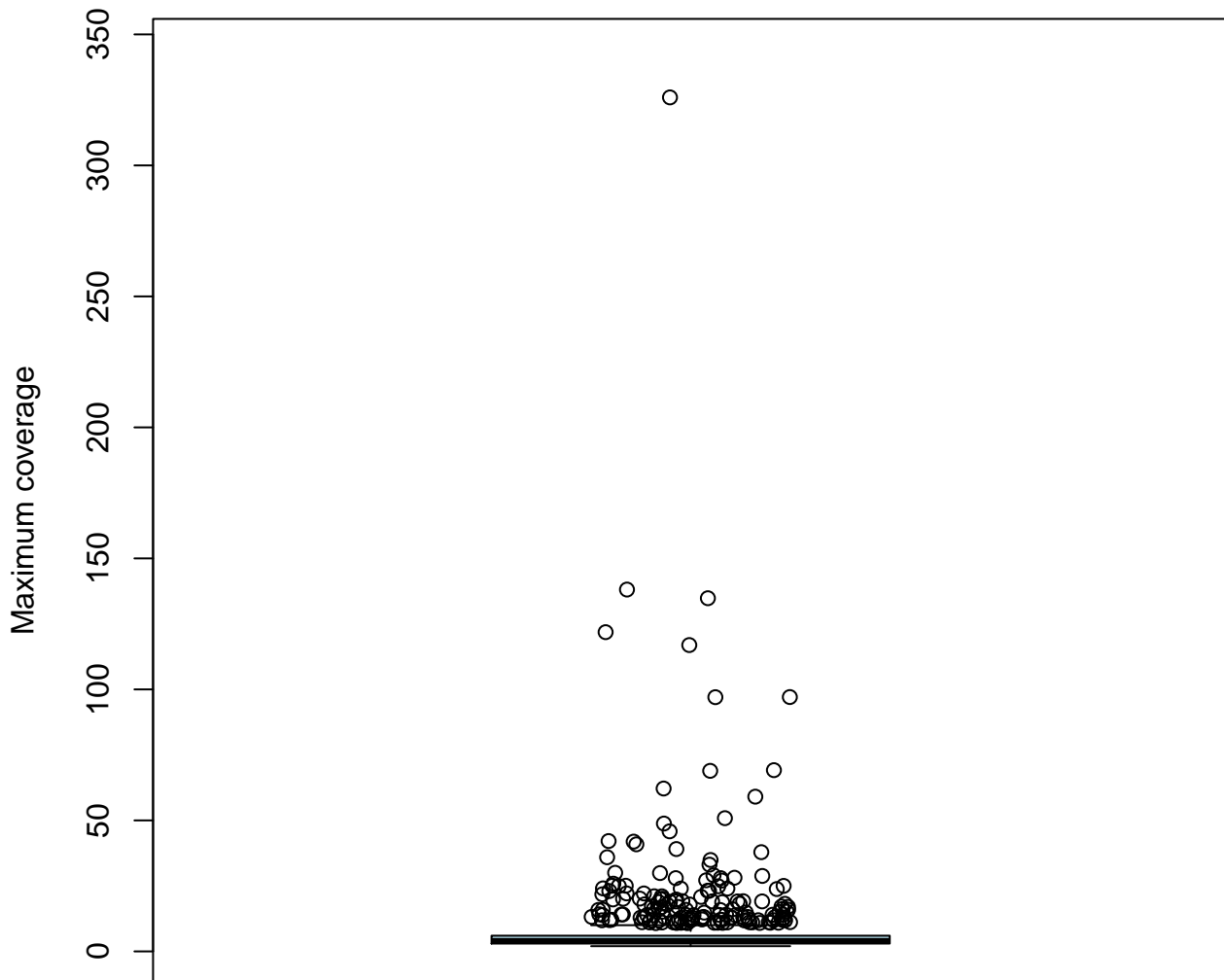
**jx in Intropolis but not "annotated" by knownGene:
38 samples; n jx: 1473**



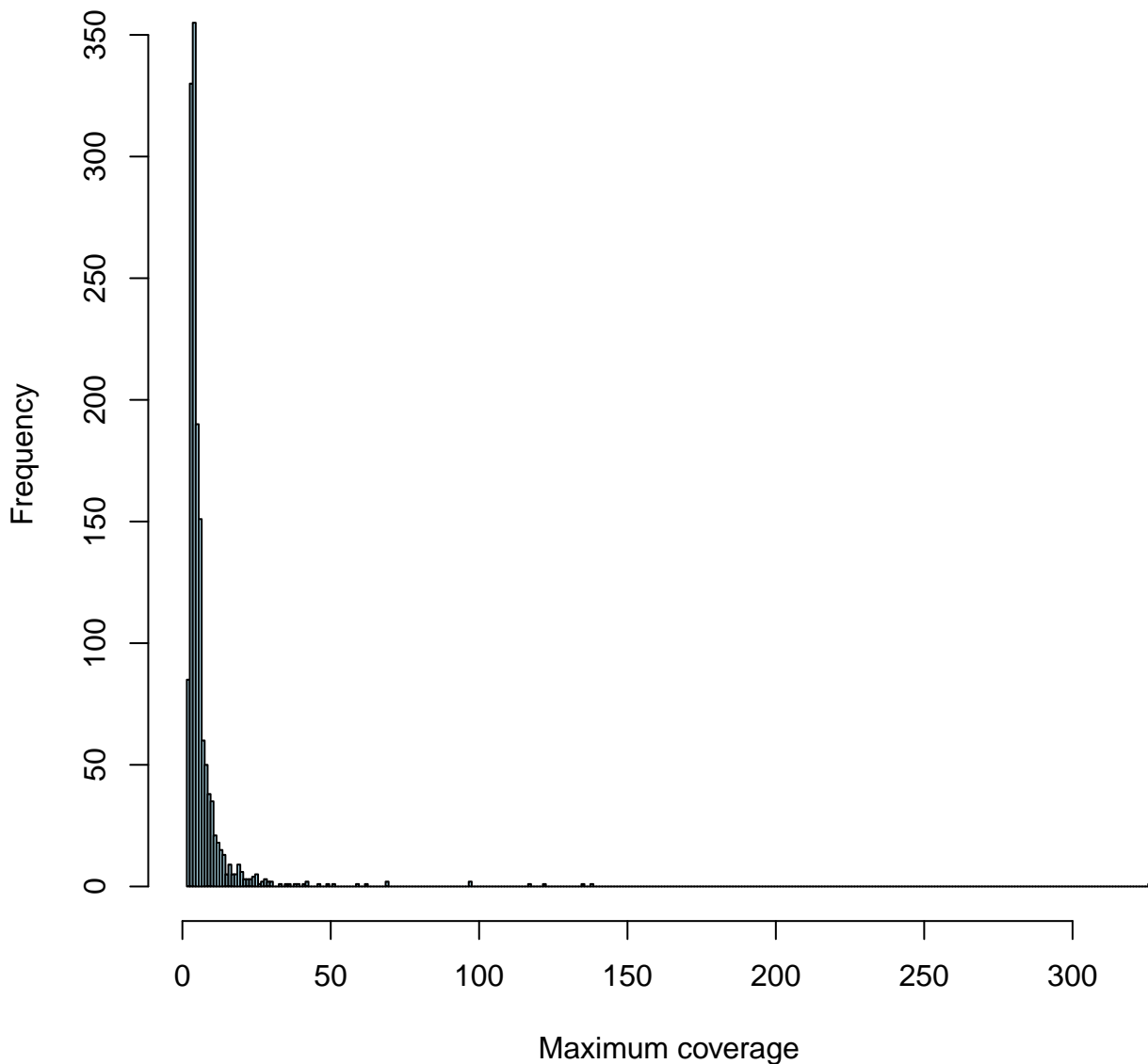
**jx in Intropolis but not "annotated" by knownGene:
38 samples; n jx: 1473**



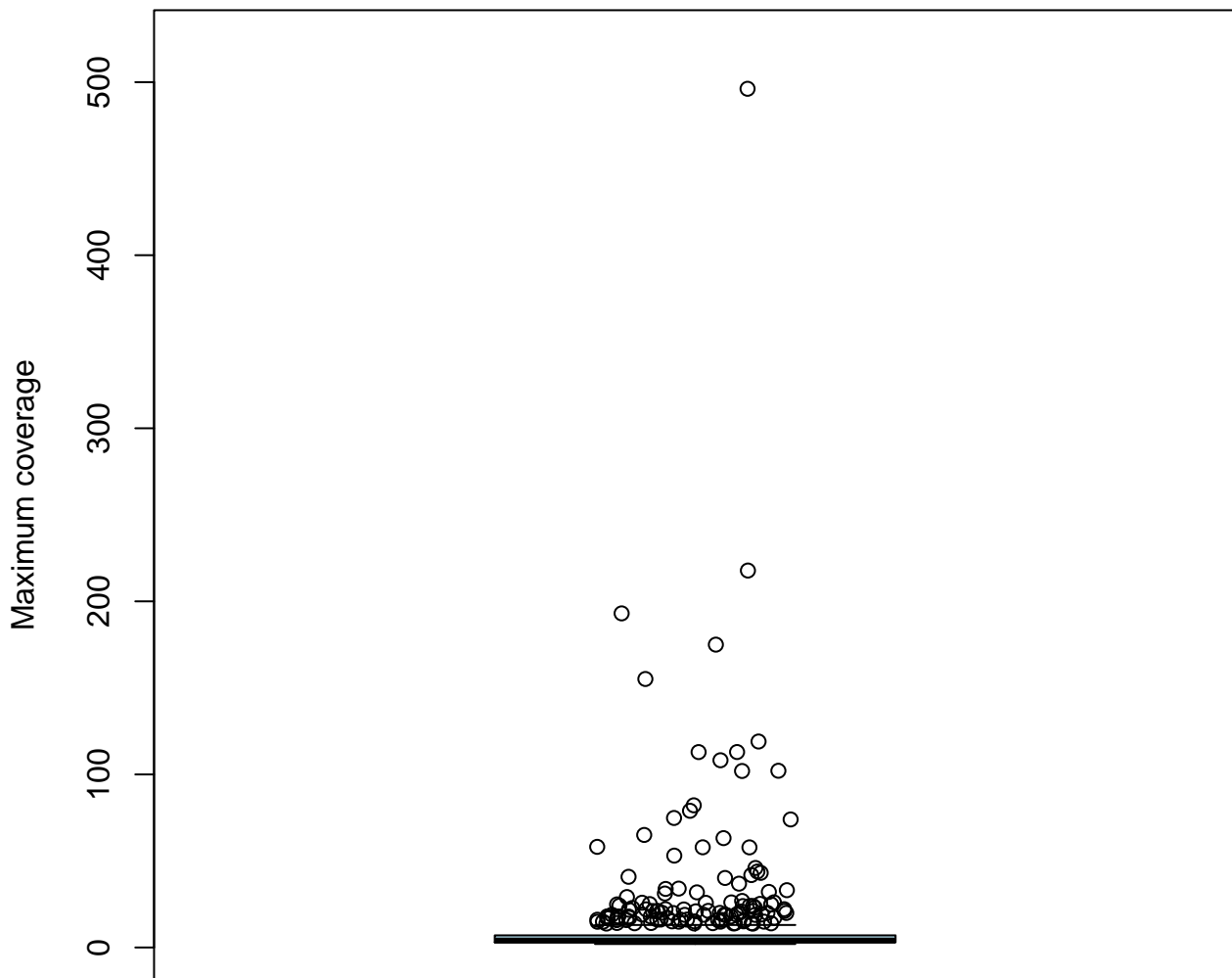
**jx in Intropolis but not "annotated" by knownGene:
39 samples; n jx: 1450**



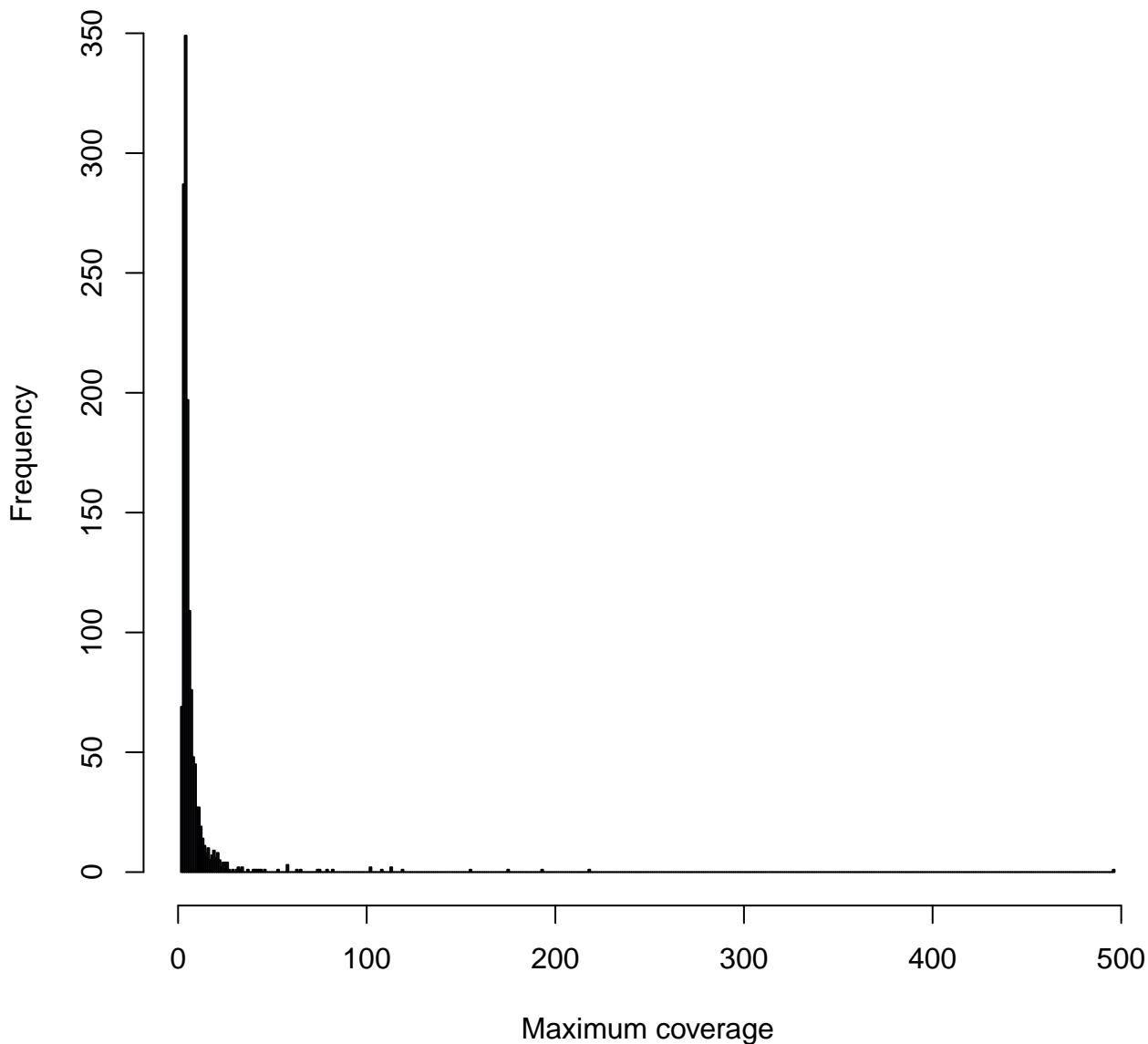
**jx in Intropolis but not "annotated" by knownGene:
39 samples; n jx: 1450**



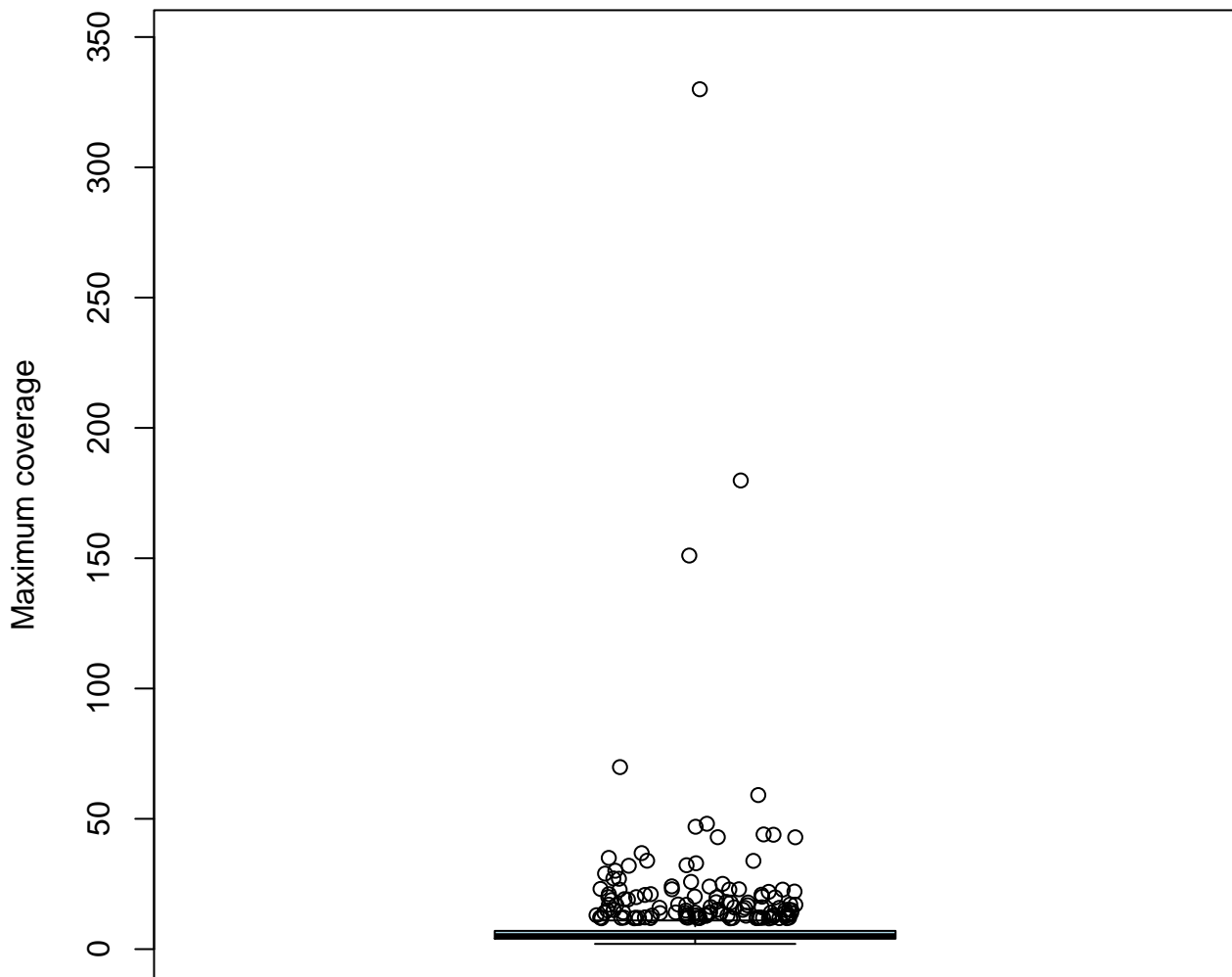
**jx in Intropolis but not "annotated" by knownGene:
40 samples; n jx: 1386**



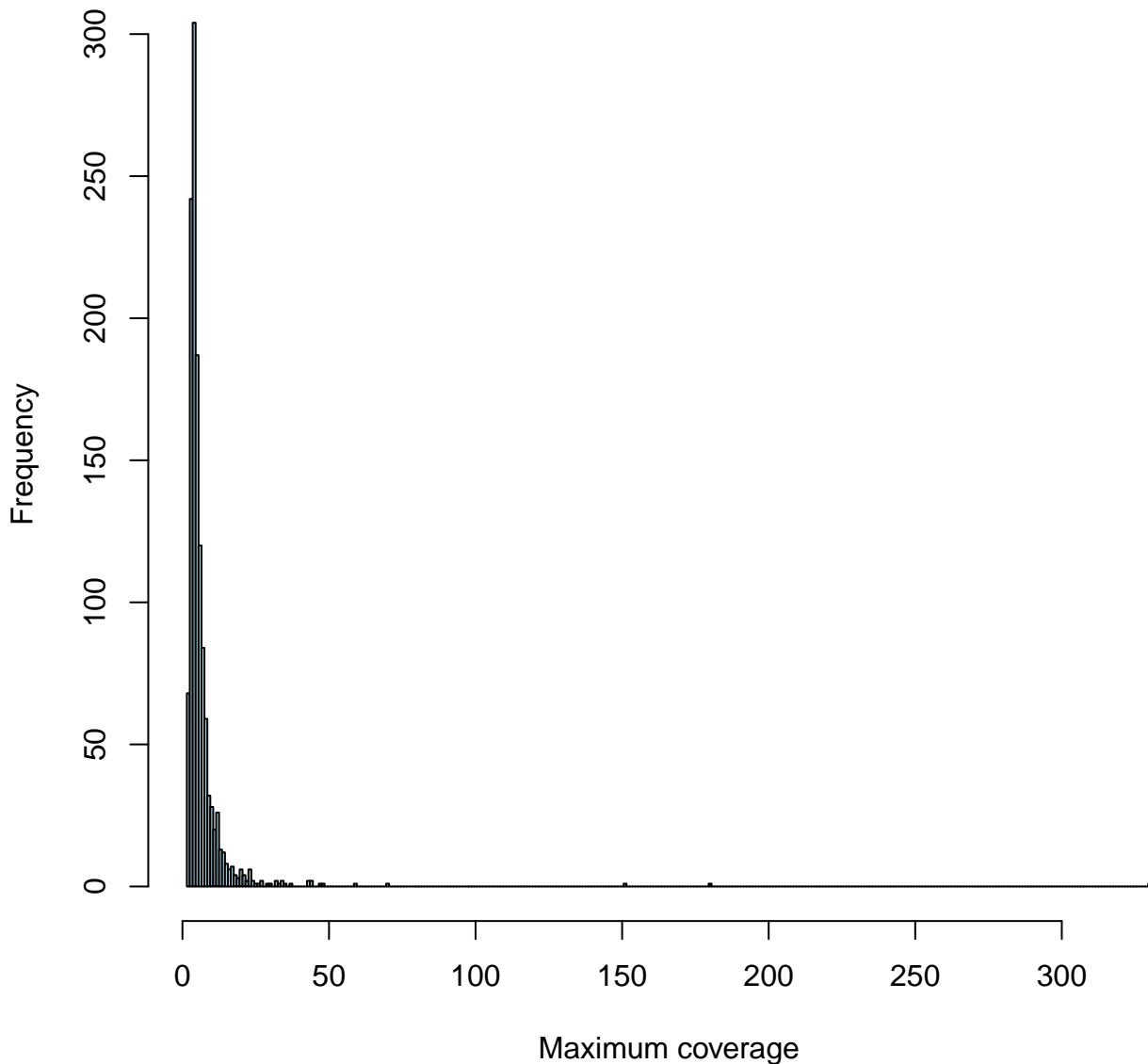
**jx in Intropolis but not "annotated" by knownGene:
40 samples; n jx: 1386**



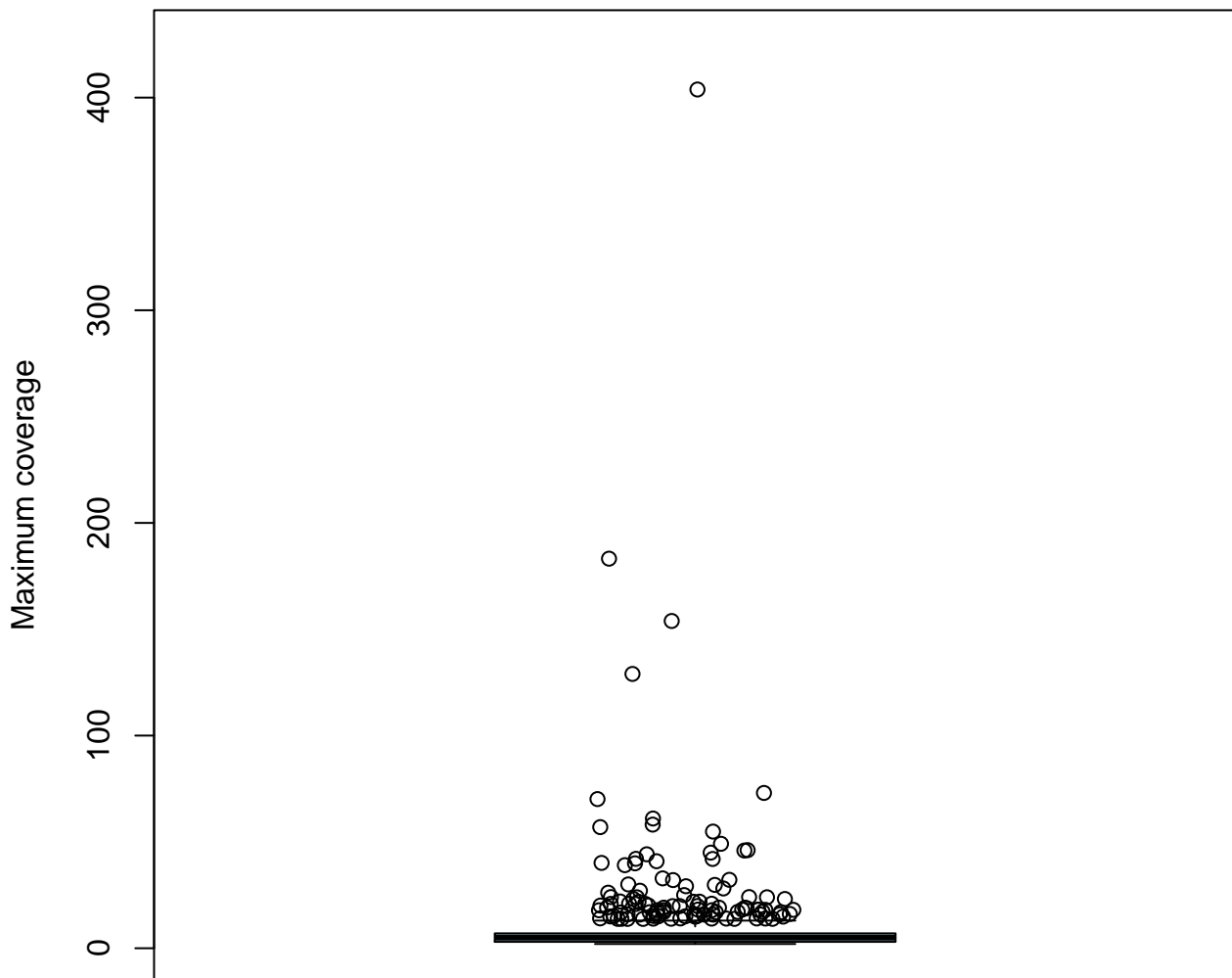
**jx in Intropolis but not "annotated" by knownGene:
41 samples; n jx: 1267**



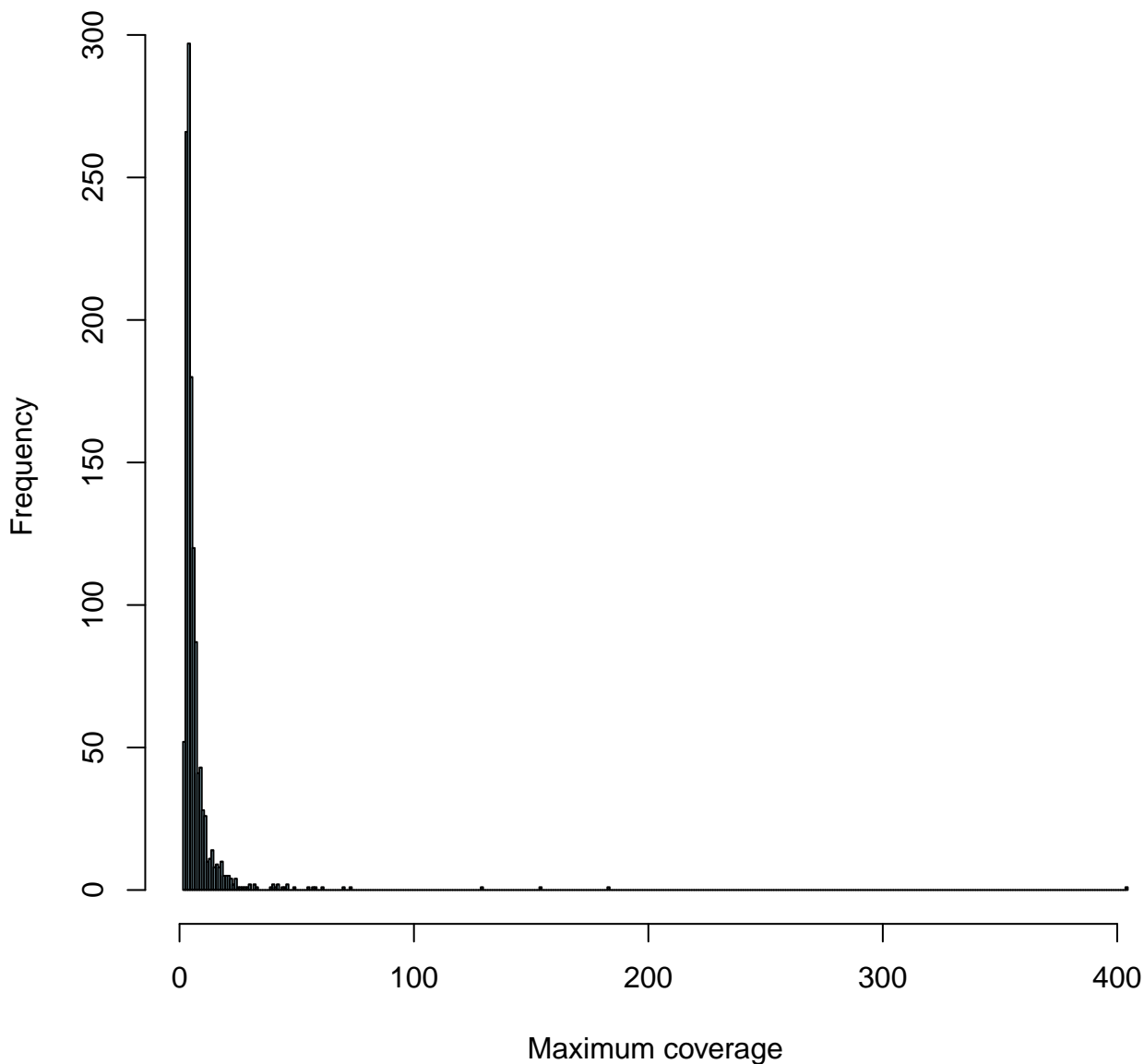
**jx in Intropolis but not "annotated" by knownGene:
41 samples; n jx: 1267**



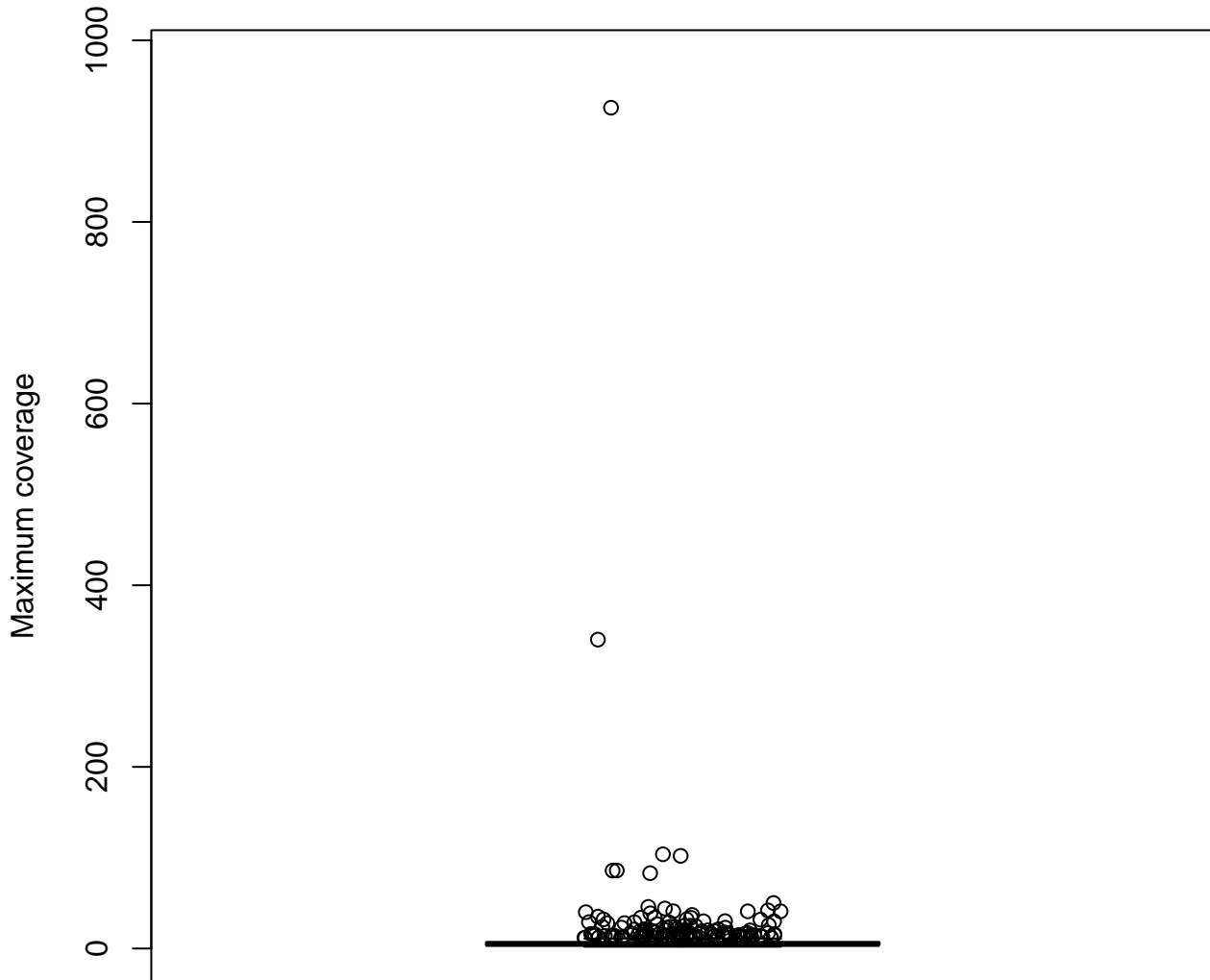
**jx in Intropolis but not "annotated" by knownGene:
42 samples; n jx: 1266**



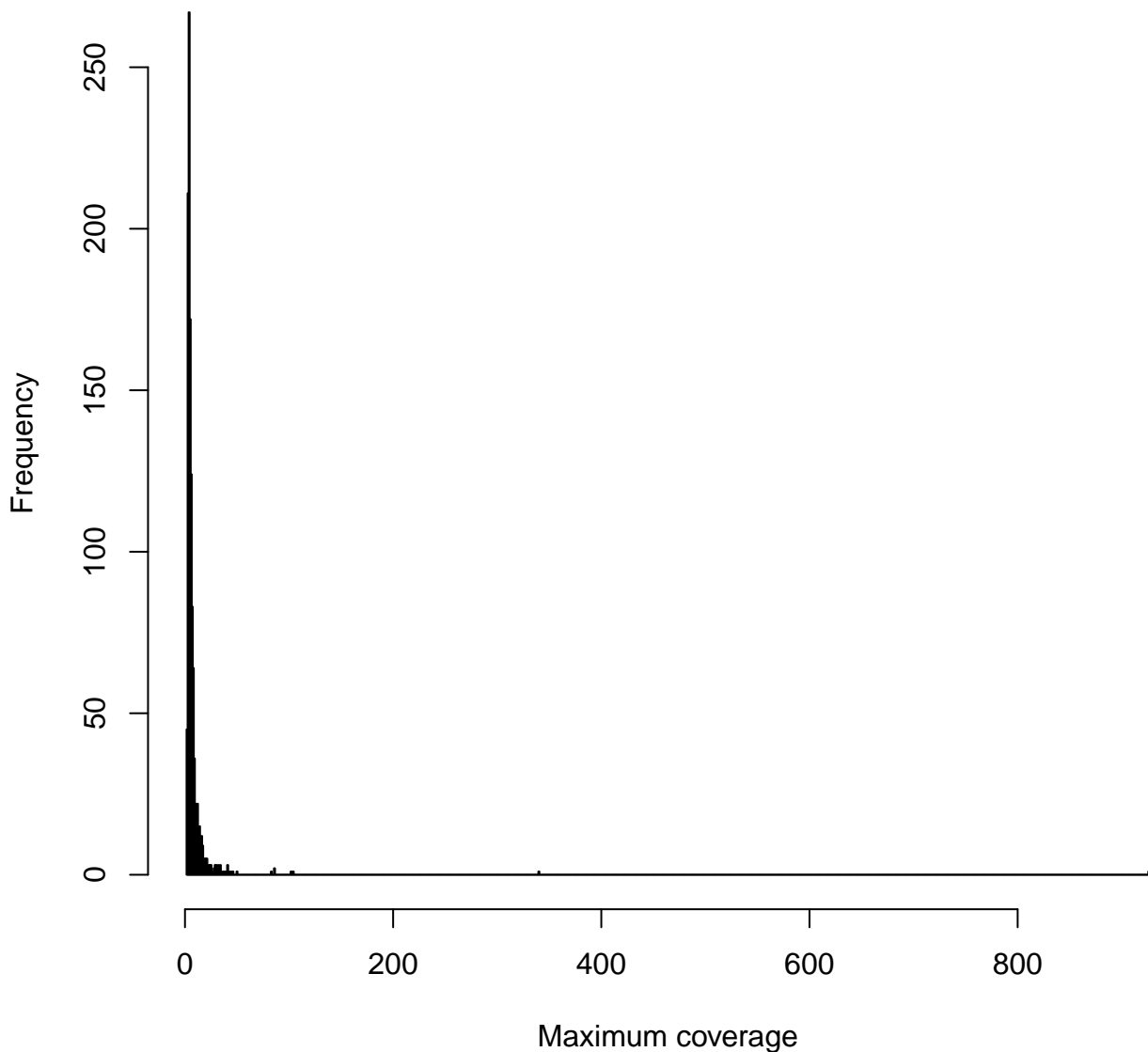
**jx in Intropolis but not "annotated" by knownGene:
42 samples; n jx: 1266**



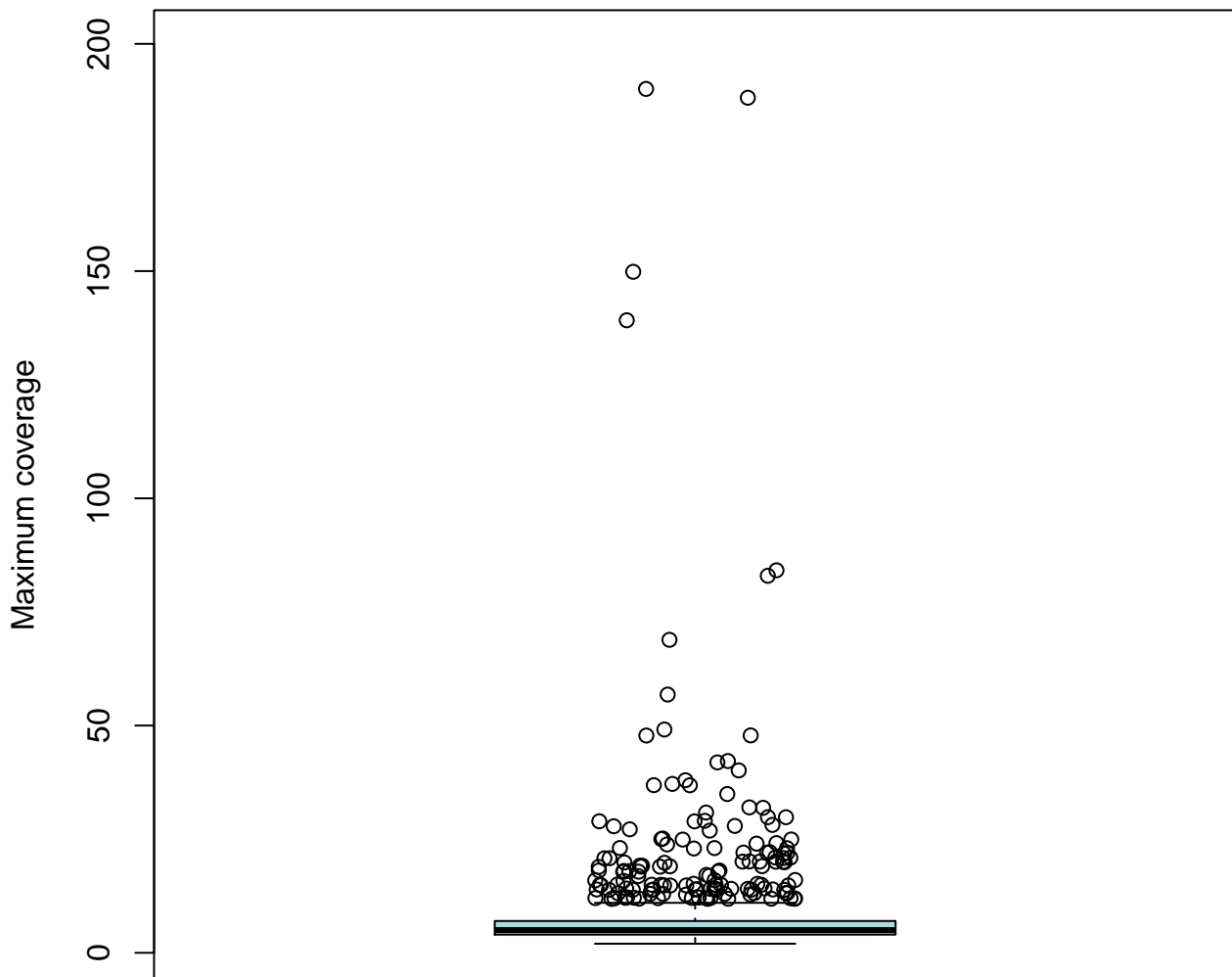
**jx in Intropolis but not "annotated" by knownGene:
43 samples; n jx: 1182**



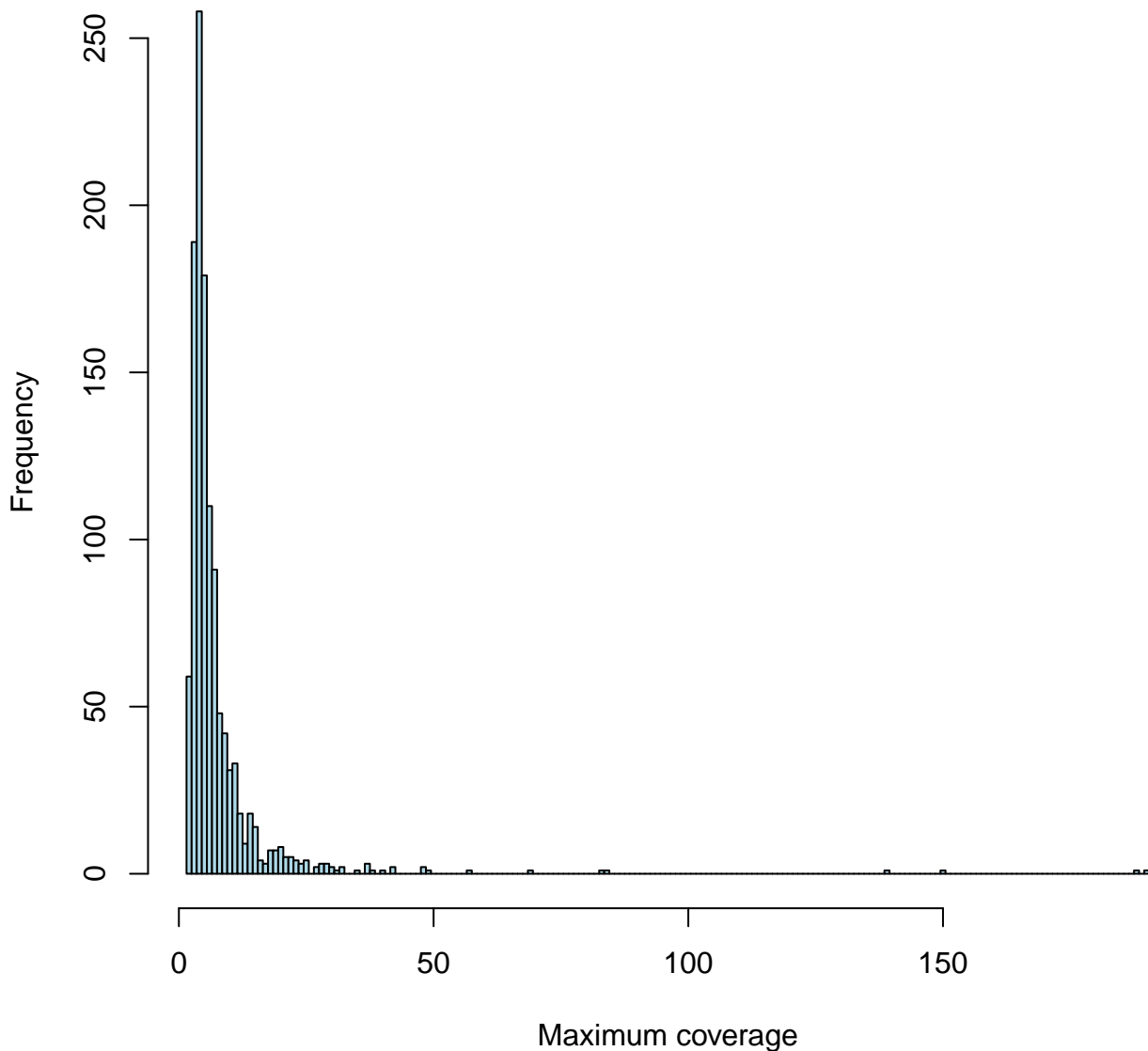
**jx in Intropolis but not "annotated" by knownGene:
43 samples; n jx: 1182**



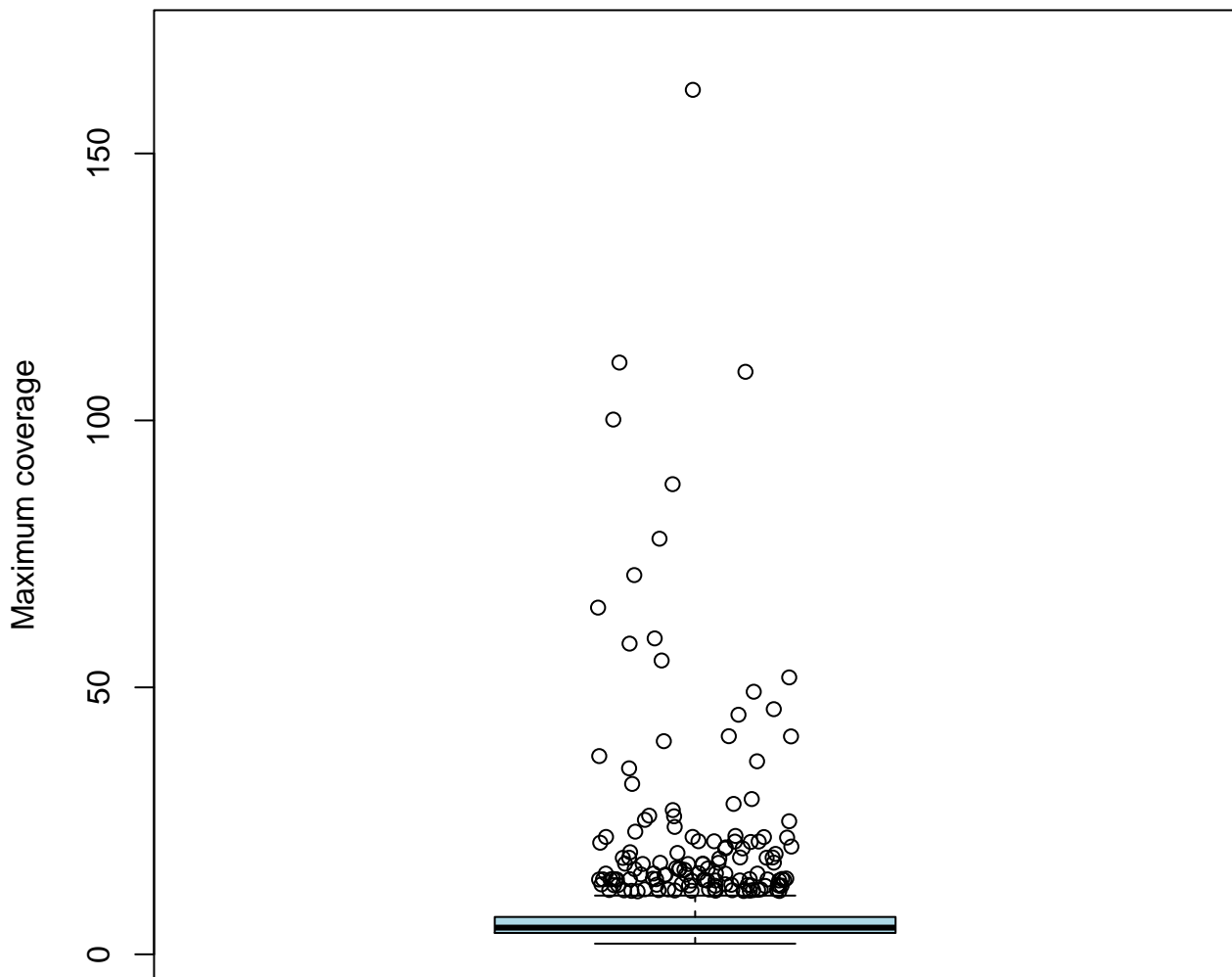
**jx in Intropolis but not "annotated" by knownGene:
44 samples; n jx: 1181**



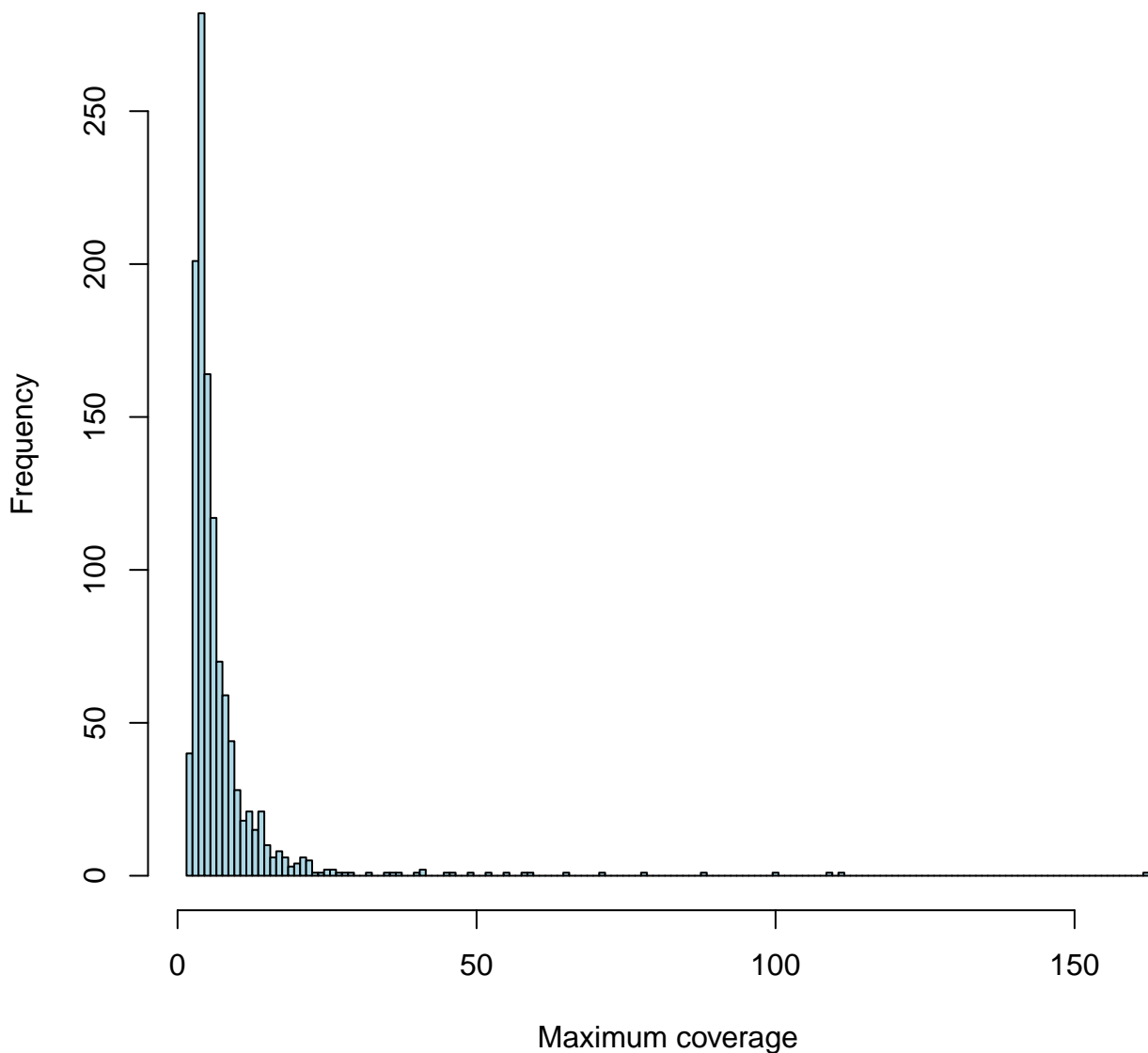
**jx in Intropolis but not "annotated" by knownGene:
44 samples; n jx: 1181**



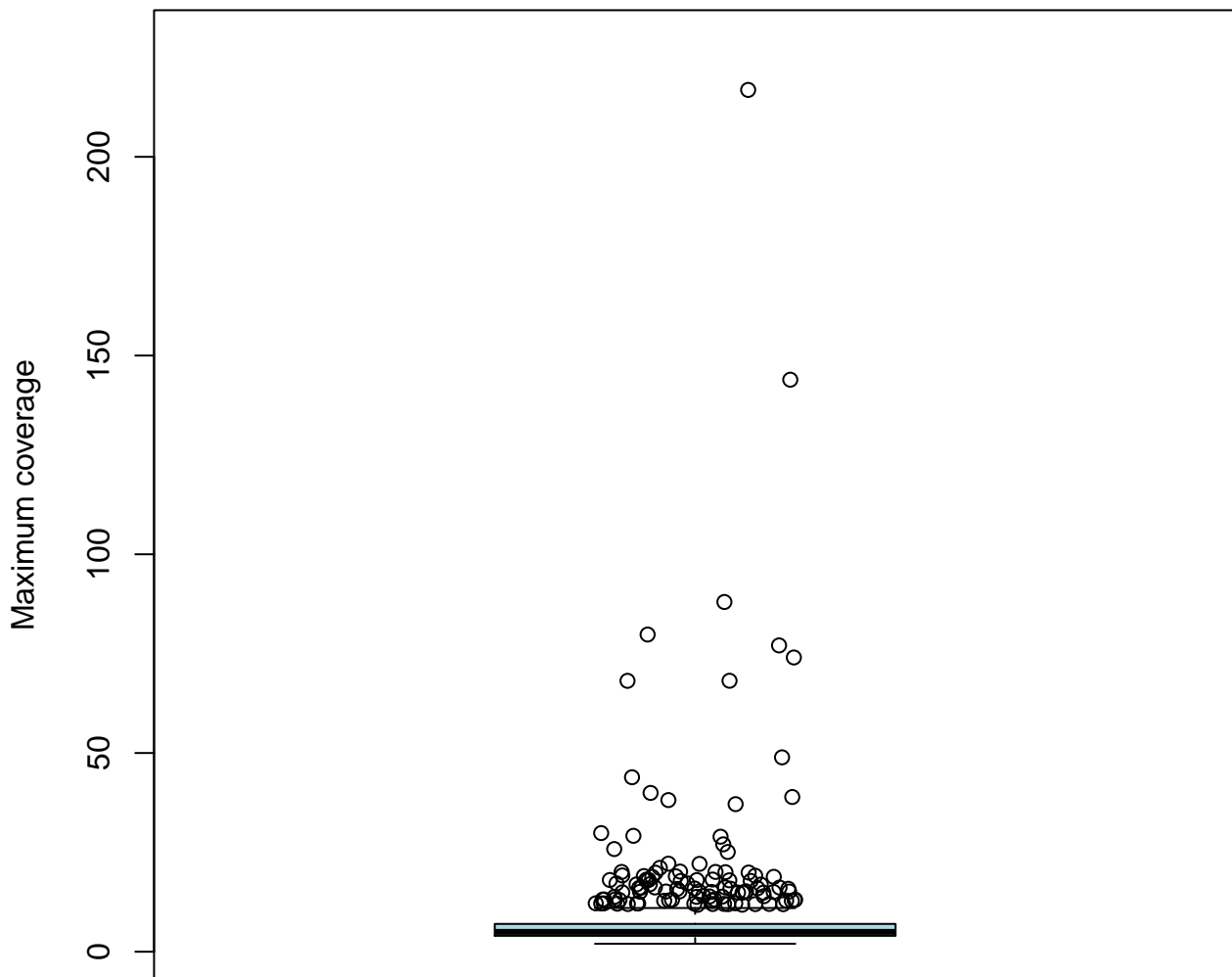
**jx in Intropolis but not "annotated" by knownGene:
45 samples; n jx: 1159**



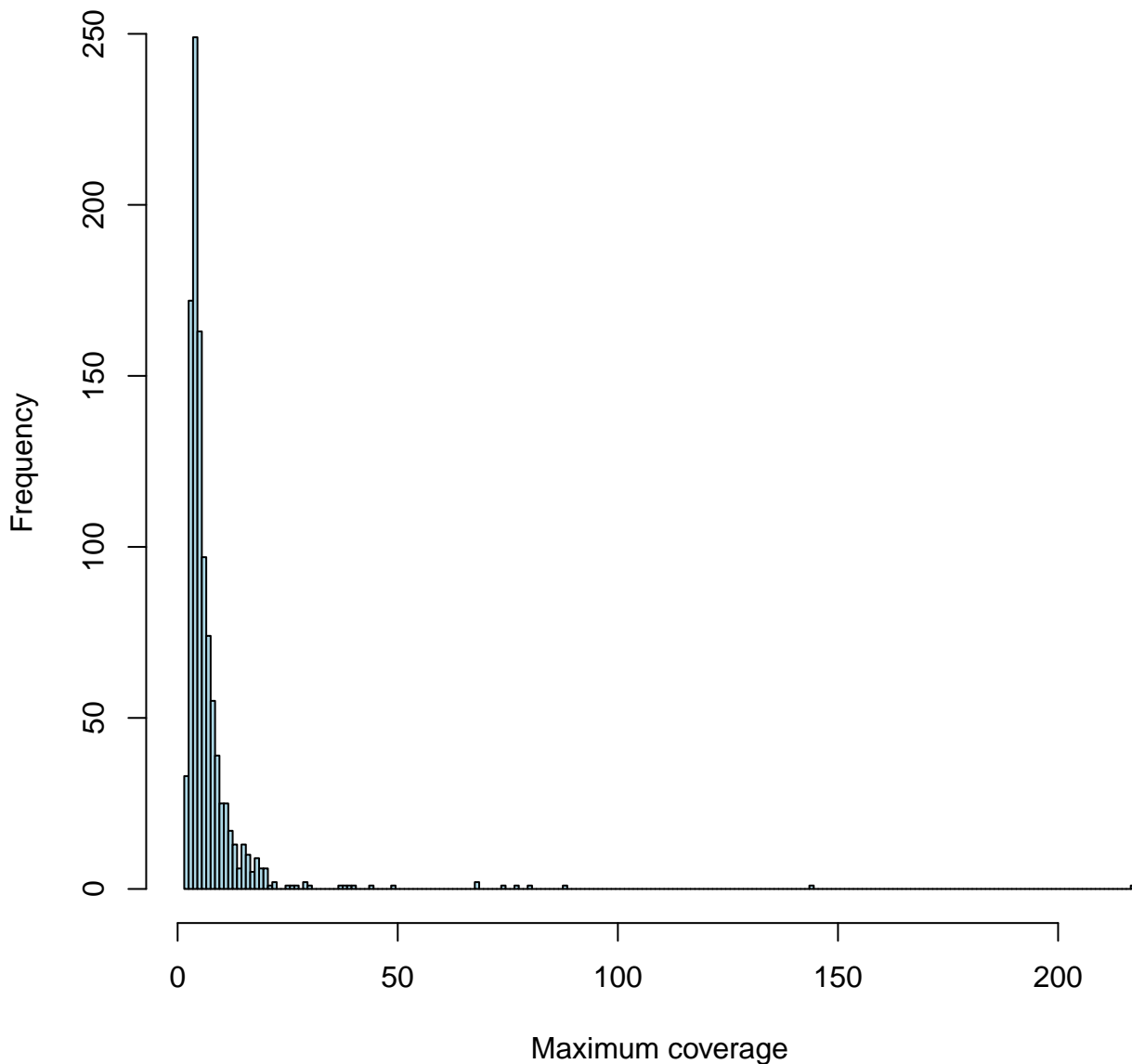
**jx in Intropolis but not "annotated" by knownGene:
45 samples; n jx: 1159**



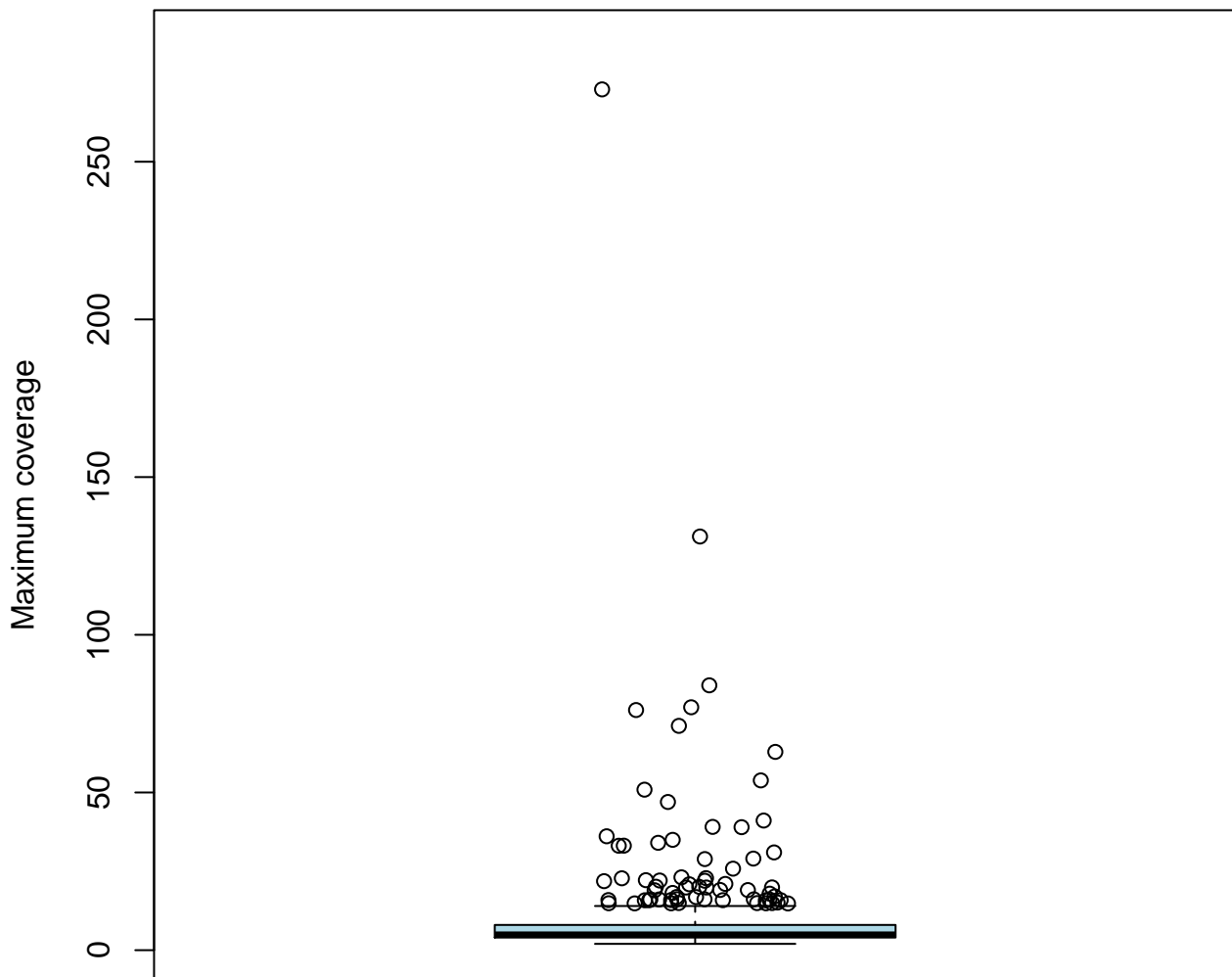
**jx in Intropolis but not "annotated" by knownGene:
46 samples; n jx: 1040**



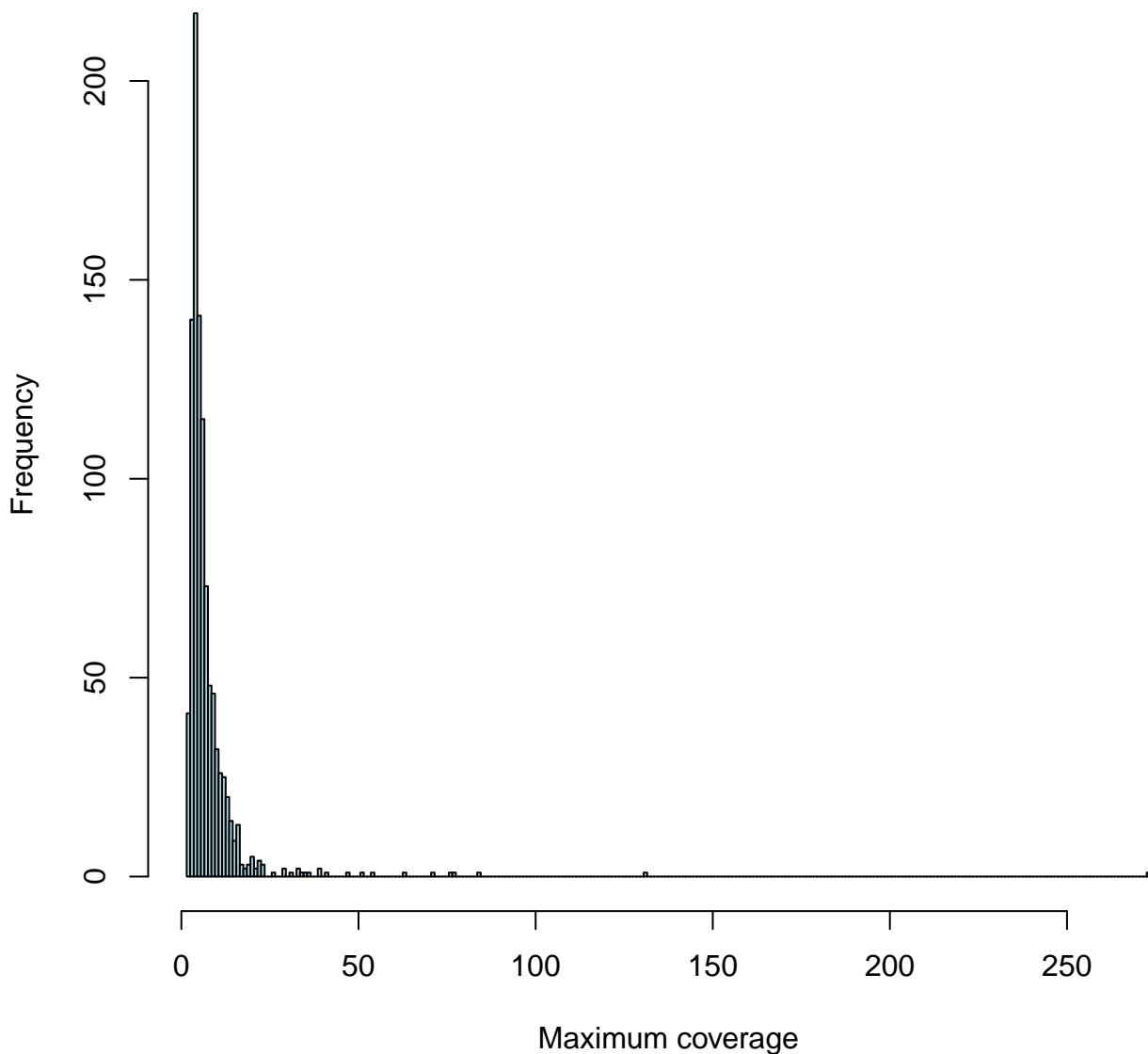
**jx in Intropolis but not "annotated" by knownGene:
46 samples; n jx: 1040**



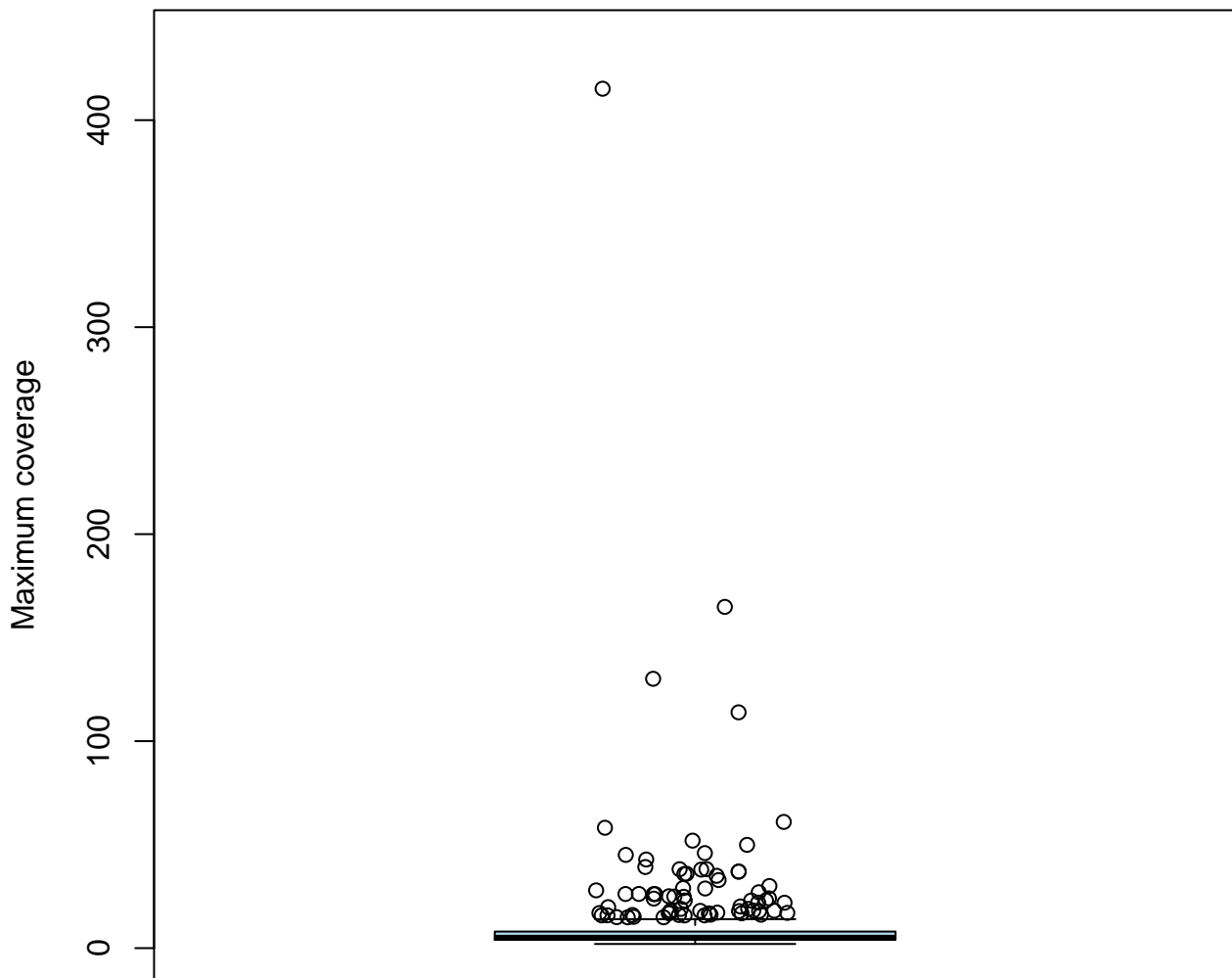
**jx in Intropolis but not "annotated" by knownGene:
47 samples; n jx: 1004**



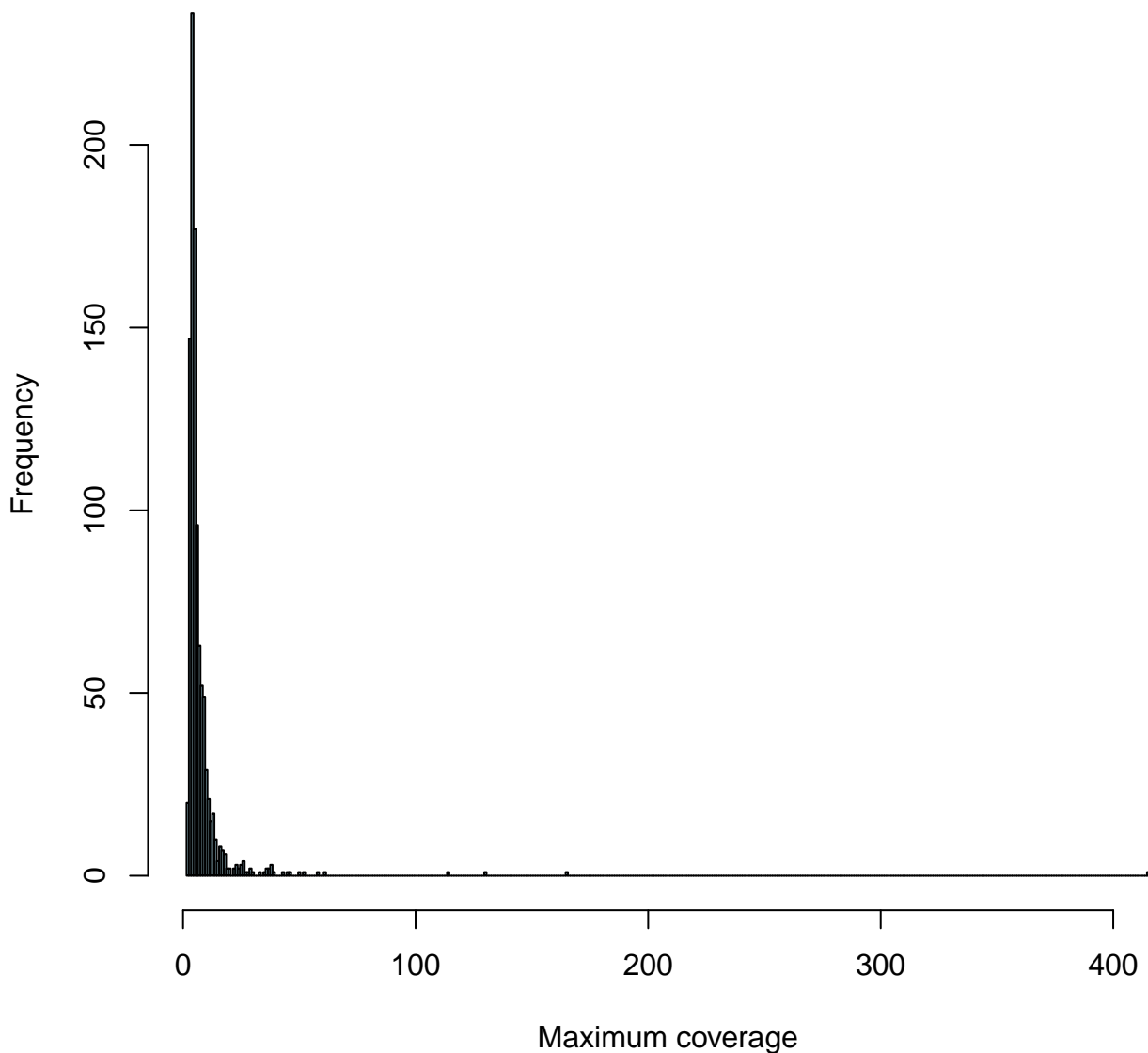
**jx in Intropolis but not "annotated" by knownGene:
47 samples; n jx: 1004**



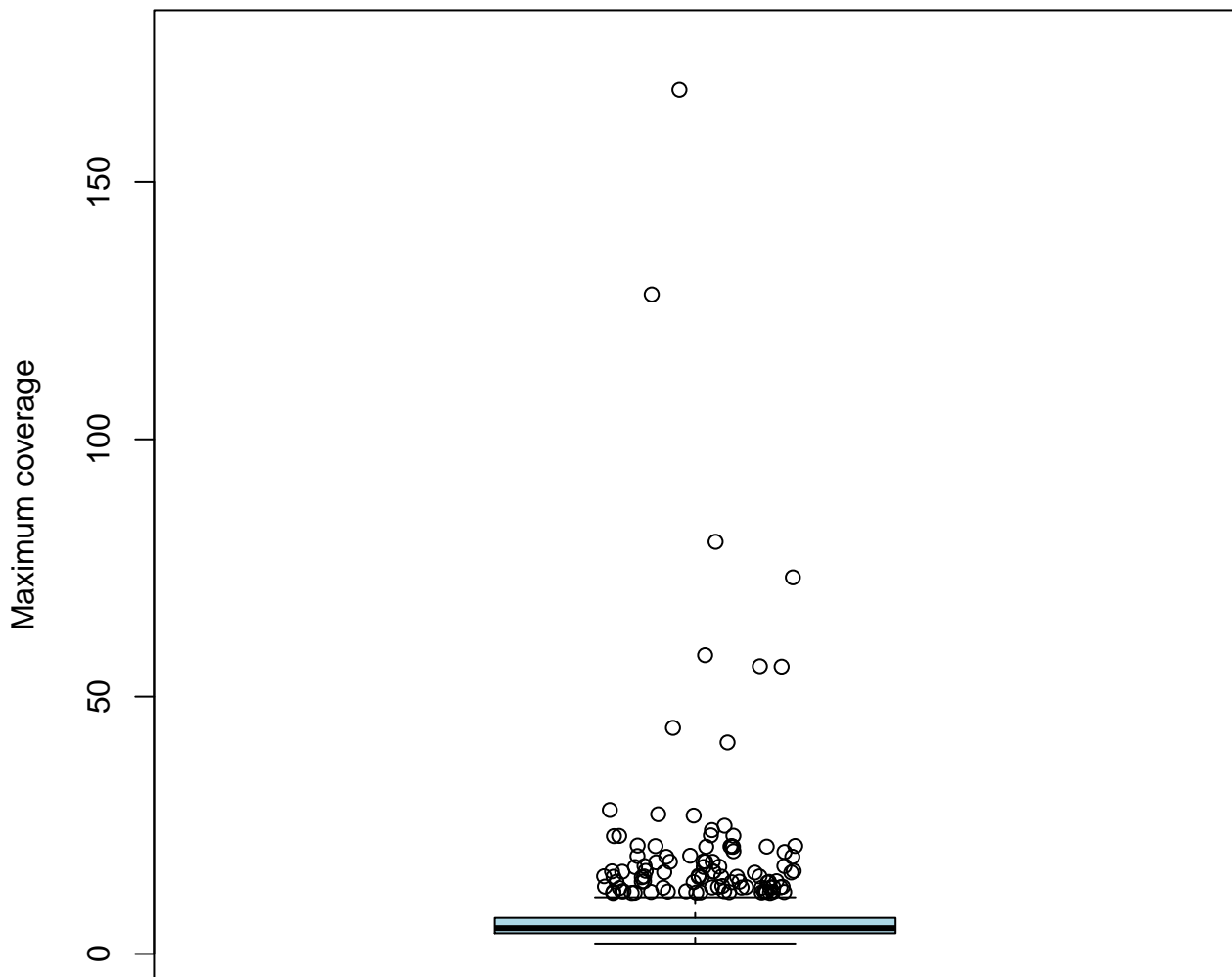
**jx in Intropolis but not "annotated" by knownGene:
48 samples; n jx: 1001**



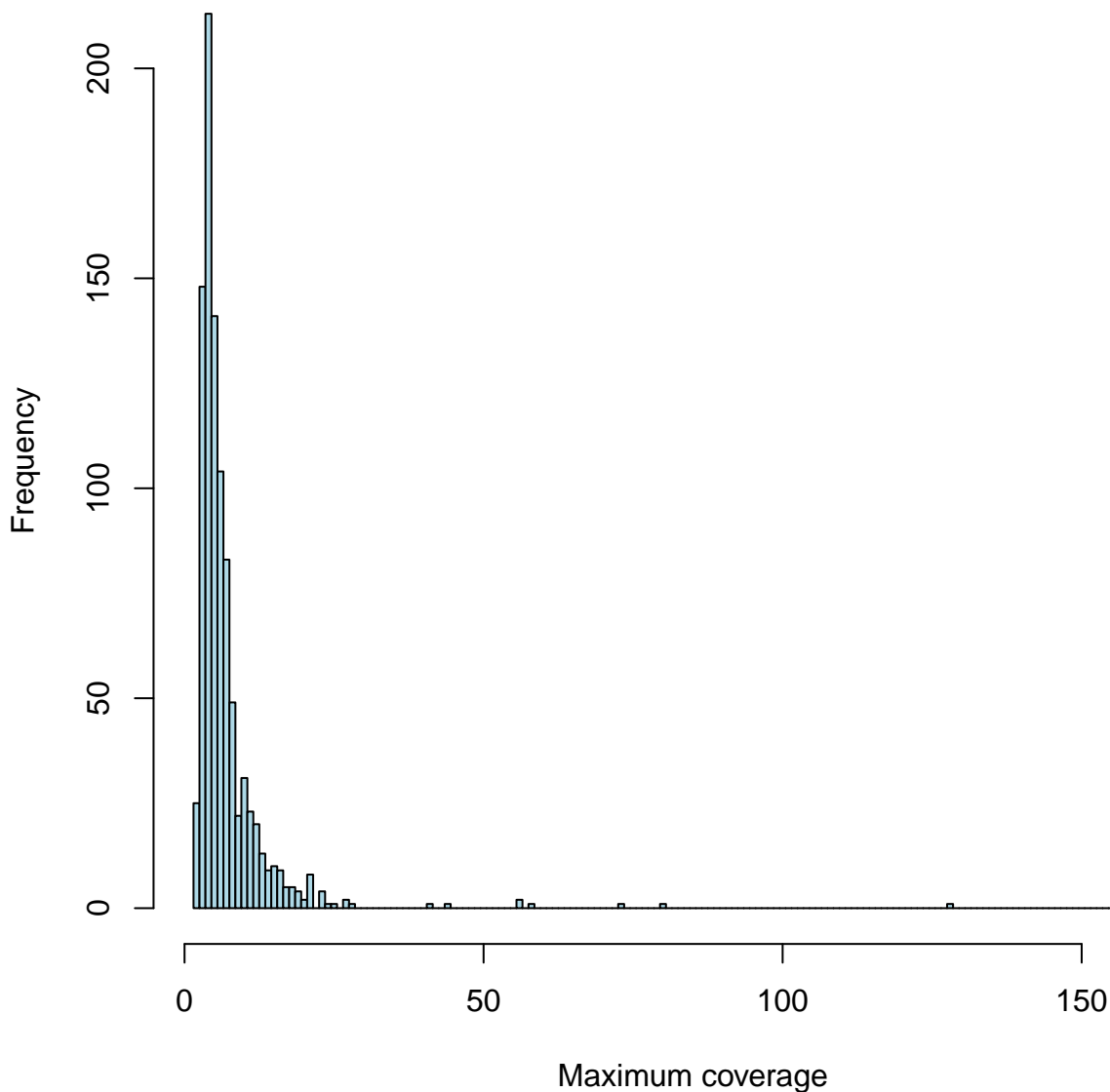
**jx in Intropolis but not "annotated" by knownGene:
48 samples; n jx: 1001**



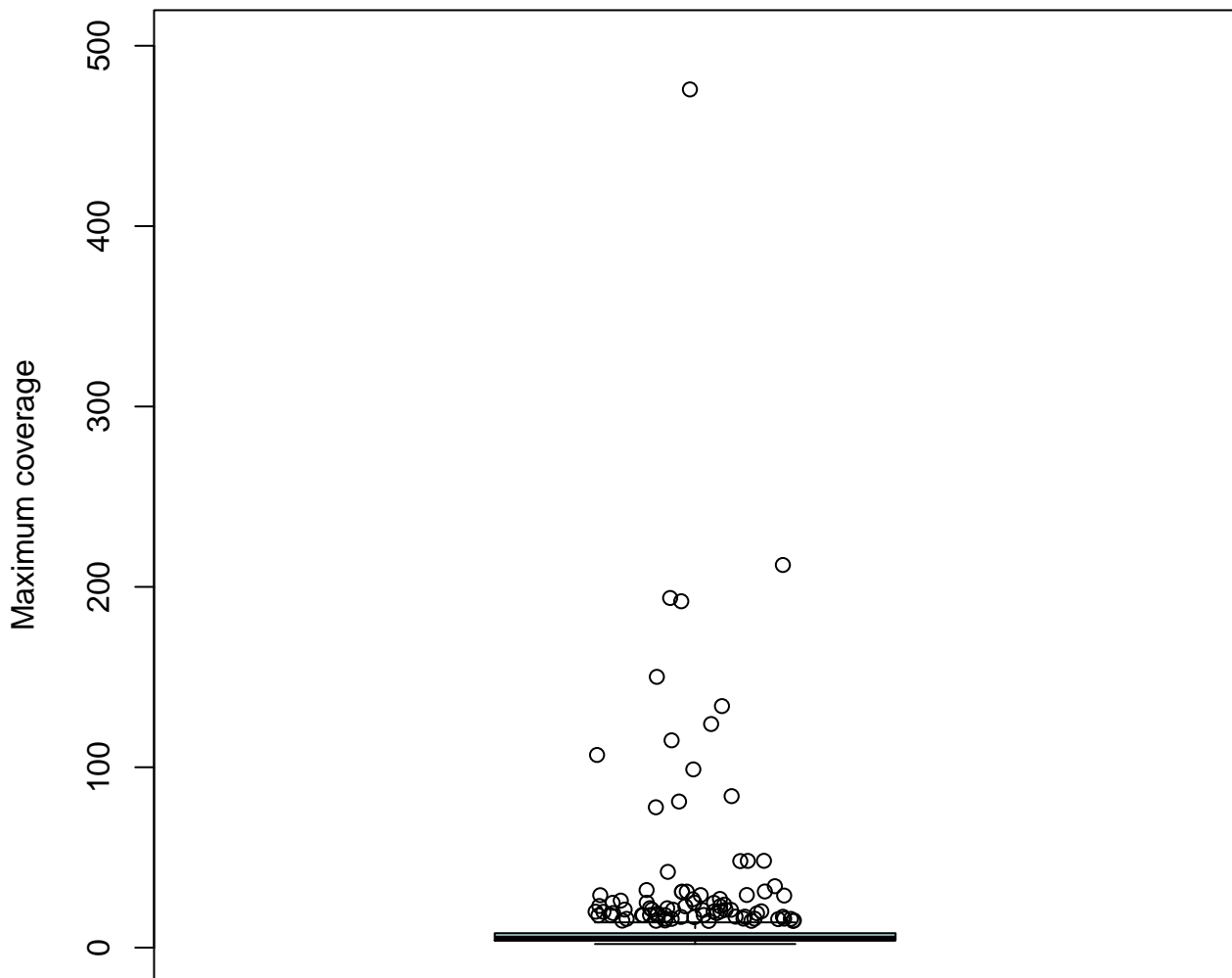
**jx in Intropolis but not "annotated" by knownGene:
49 samples; n jx: 942**



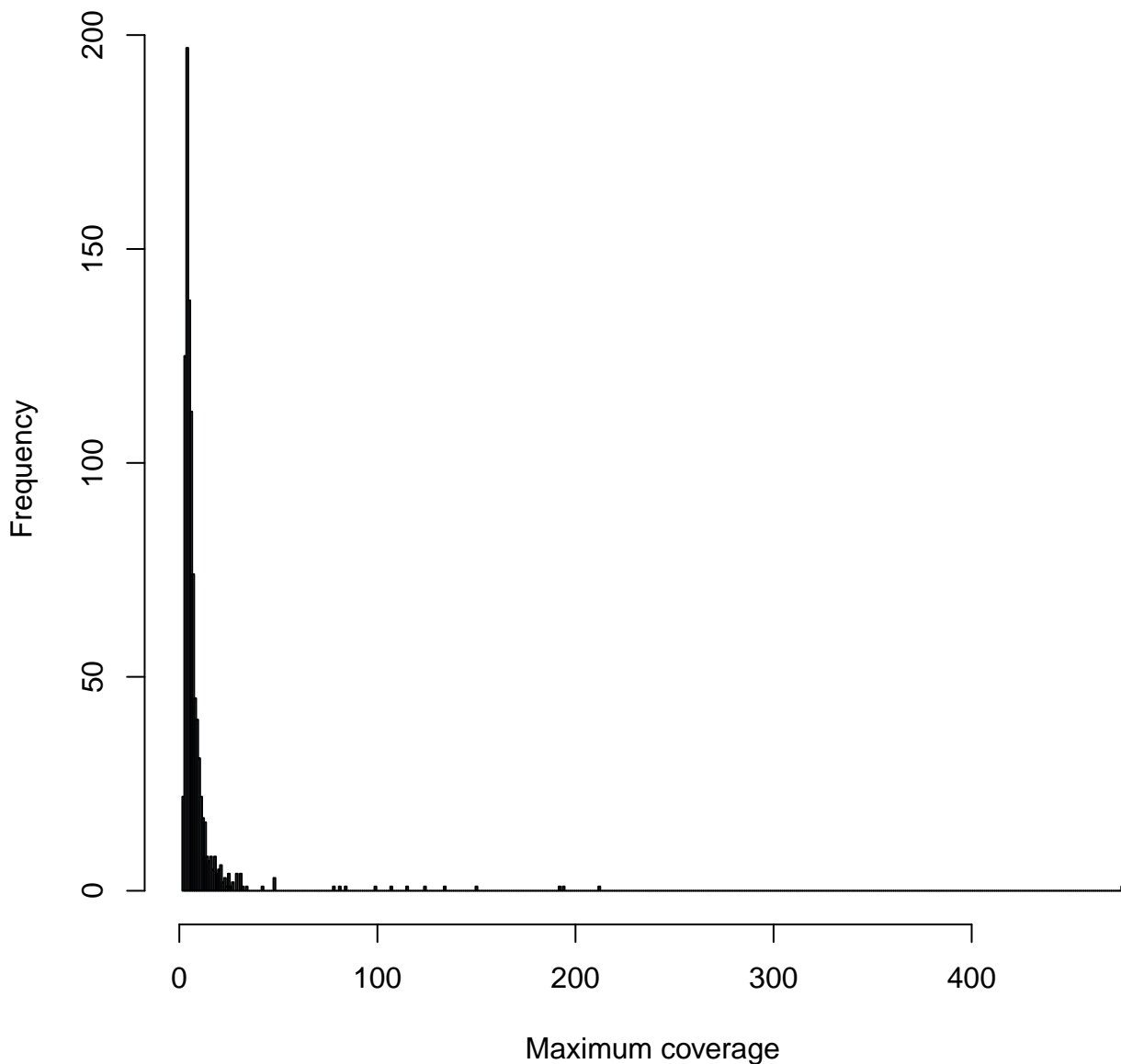
**jx in Intropolis but not "annotated" by knownGene:
49 samples; n jx: 942**



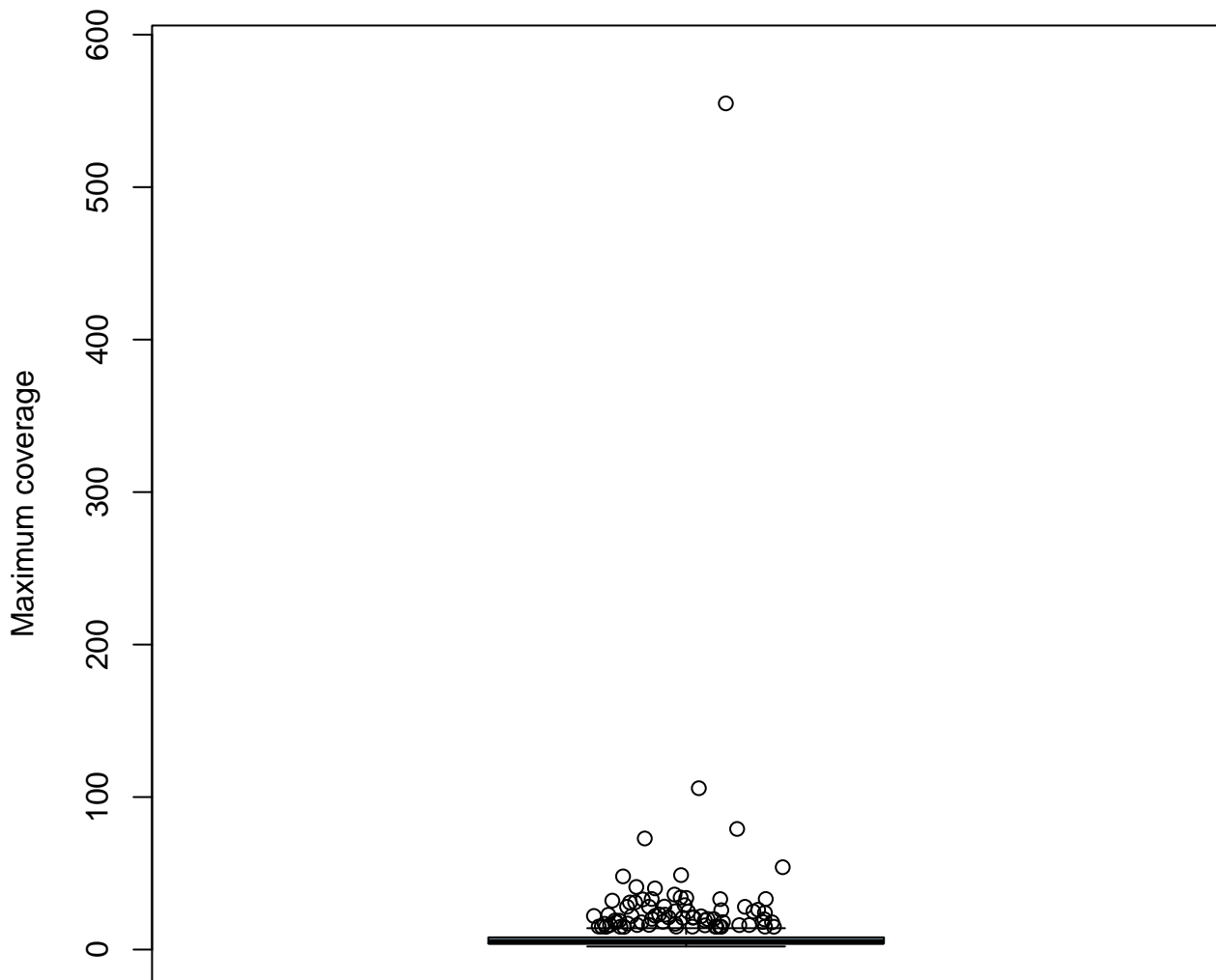
**jx in Intropolis but not "annotated" by knownGene:
50 samples; n jx: 930**



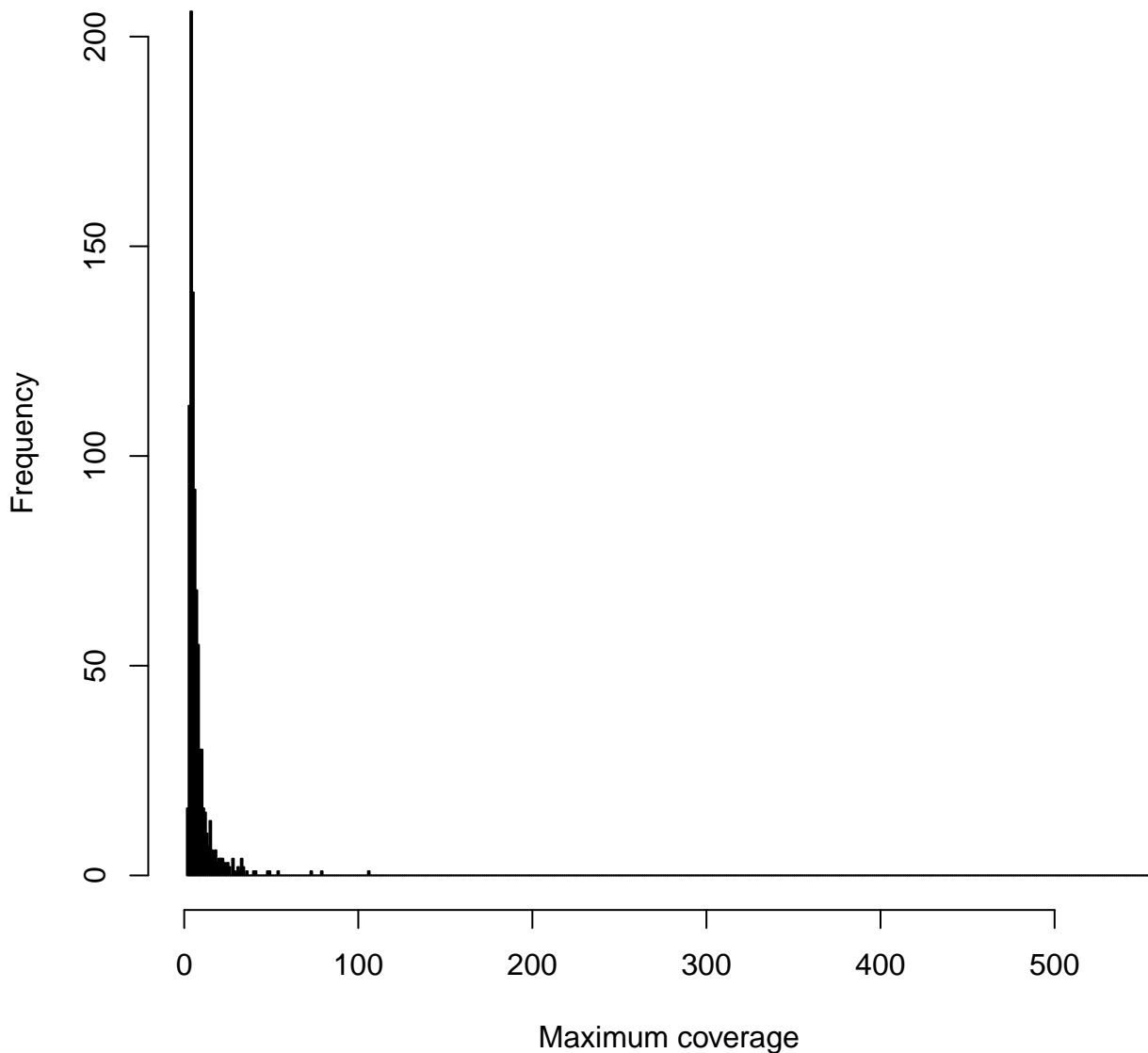
**jx in Intropolis but not "annotated" by knownGene:
50 samples; n jx: 930**



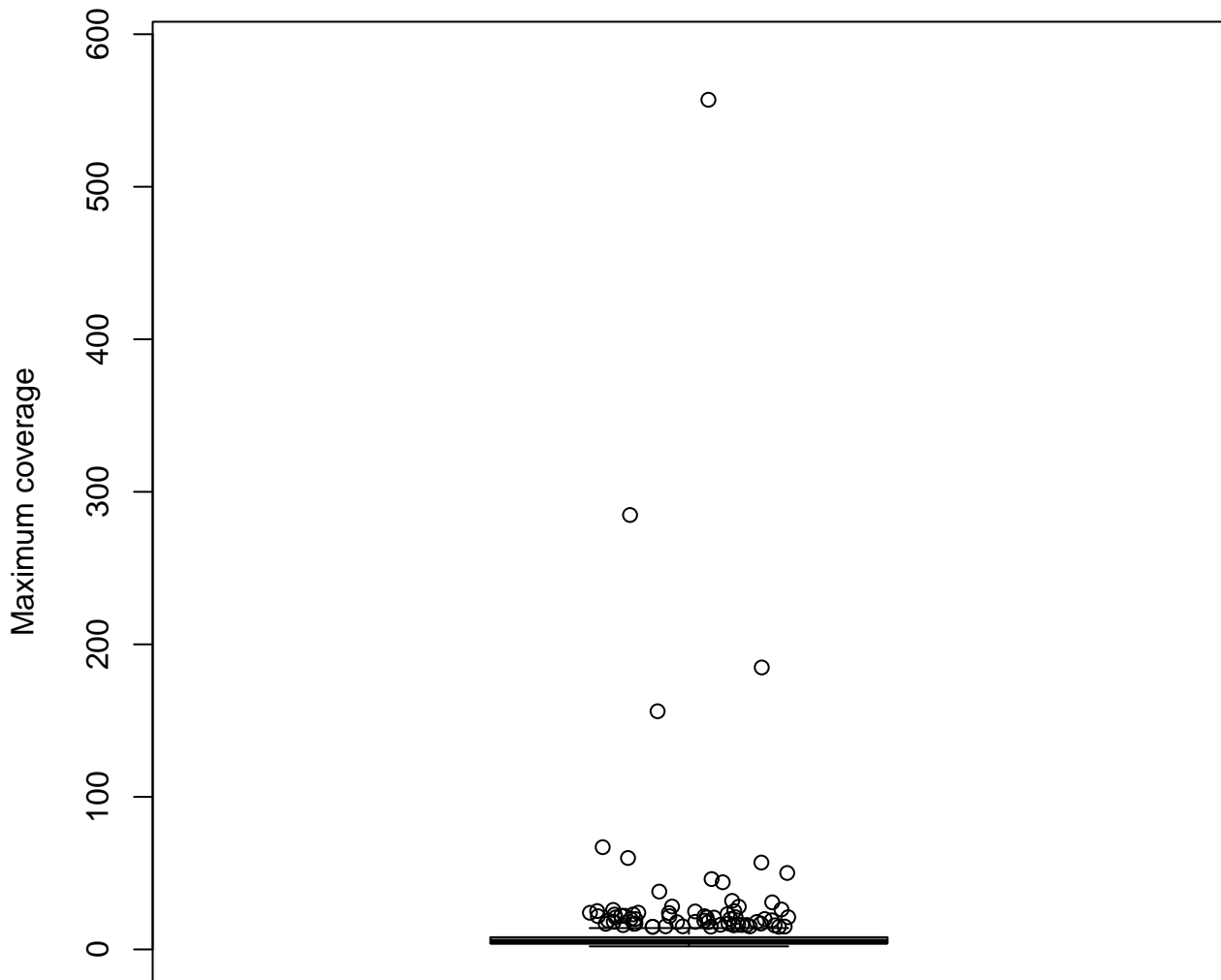
**jx in Intropolis but not "annotated" by knownGene:
51 samples; n jx: 872**



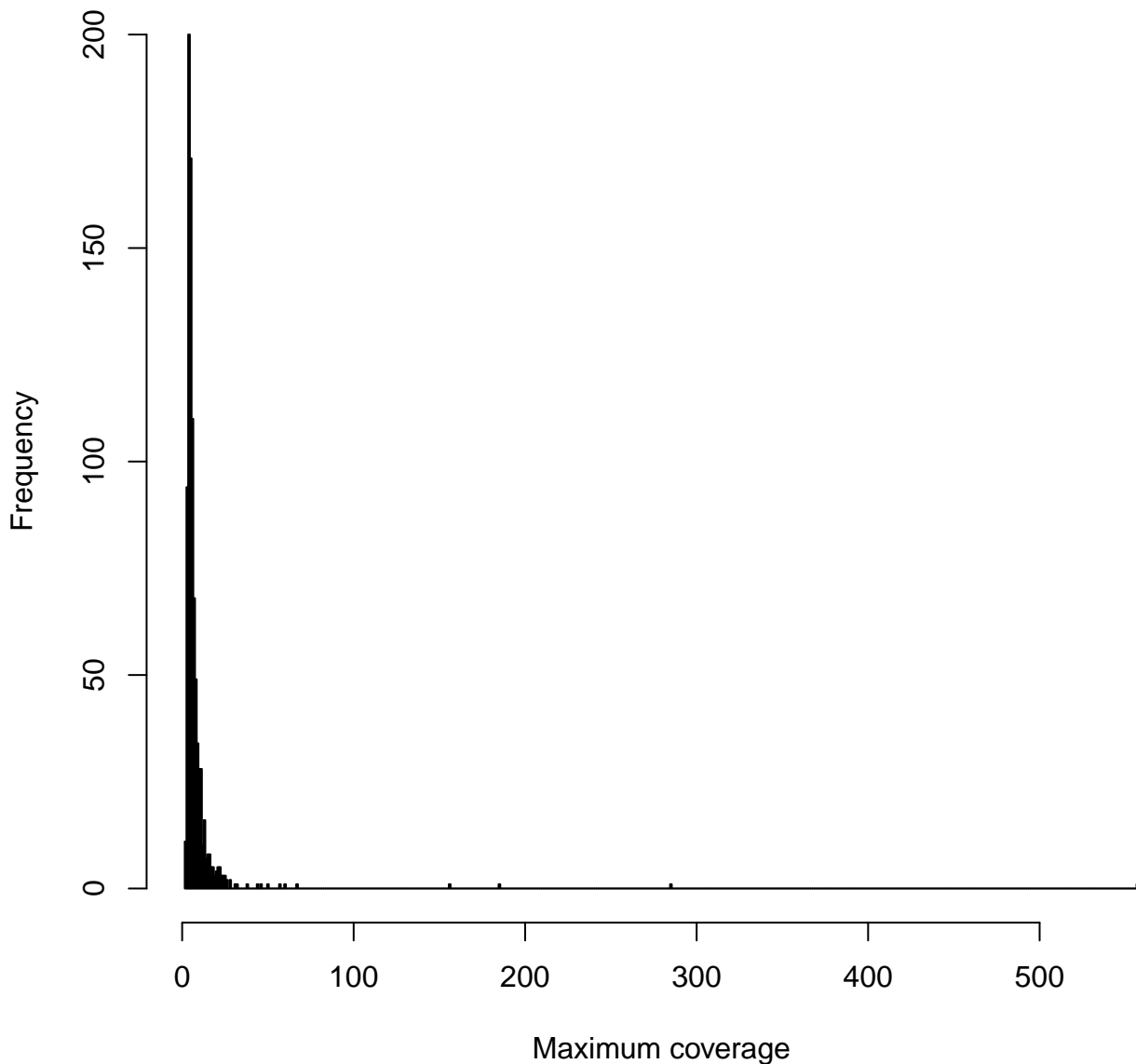
**jx in Intropolis but not "annotated" by knownGene:
51 samples; n jx: 872**



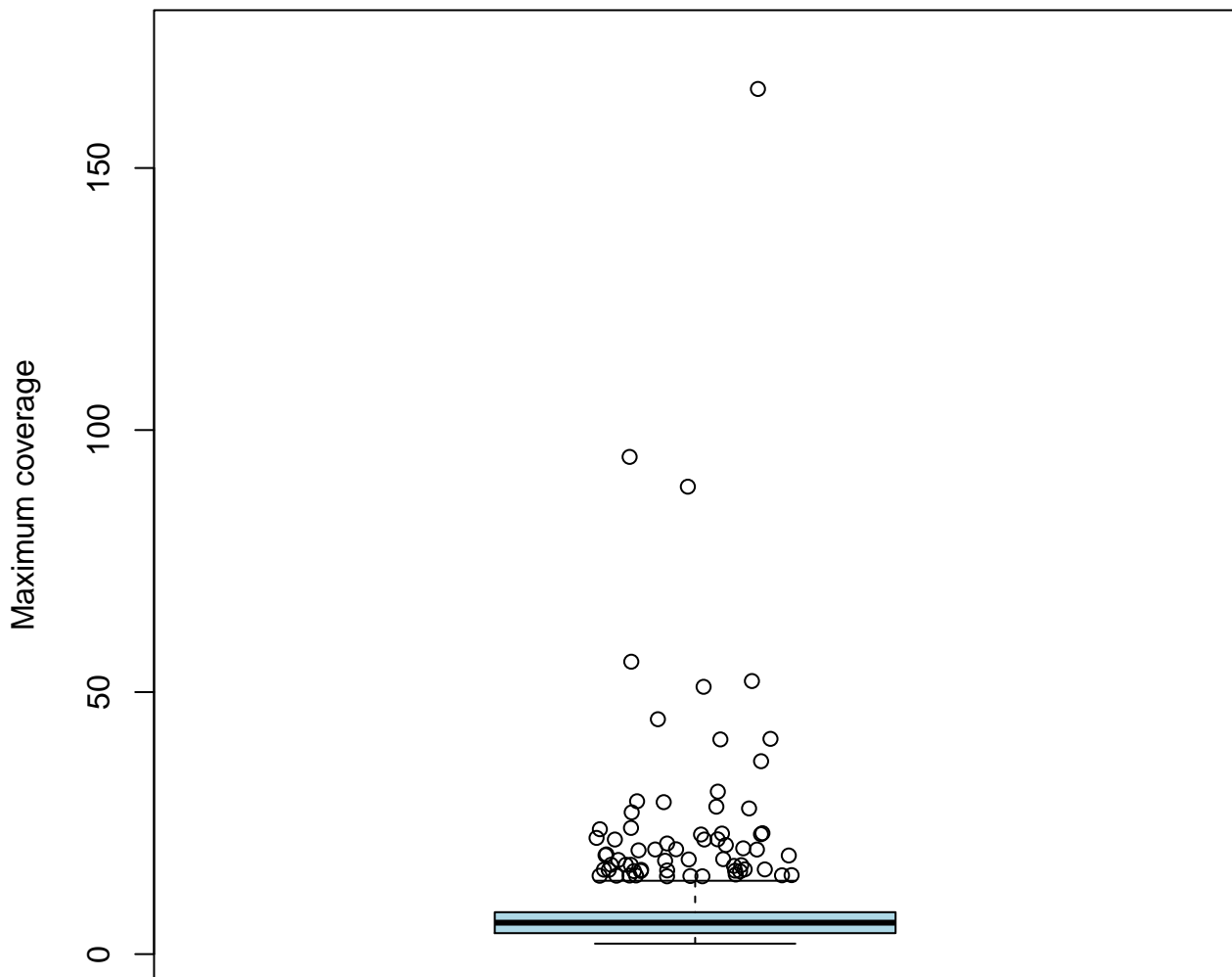
**jx in Intropolis but not "annotated" by knownGene:
52 samples; n jx: 895**



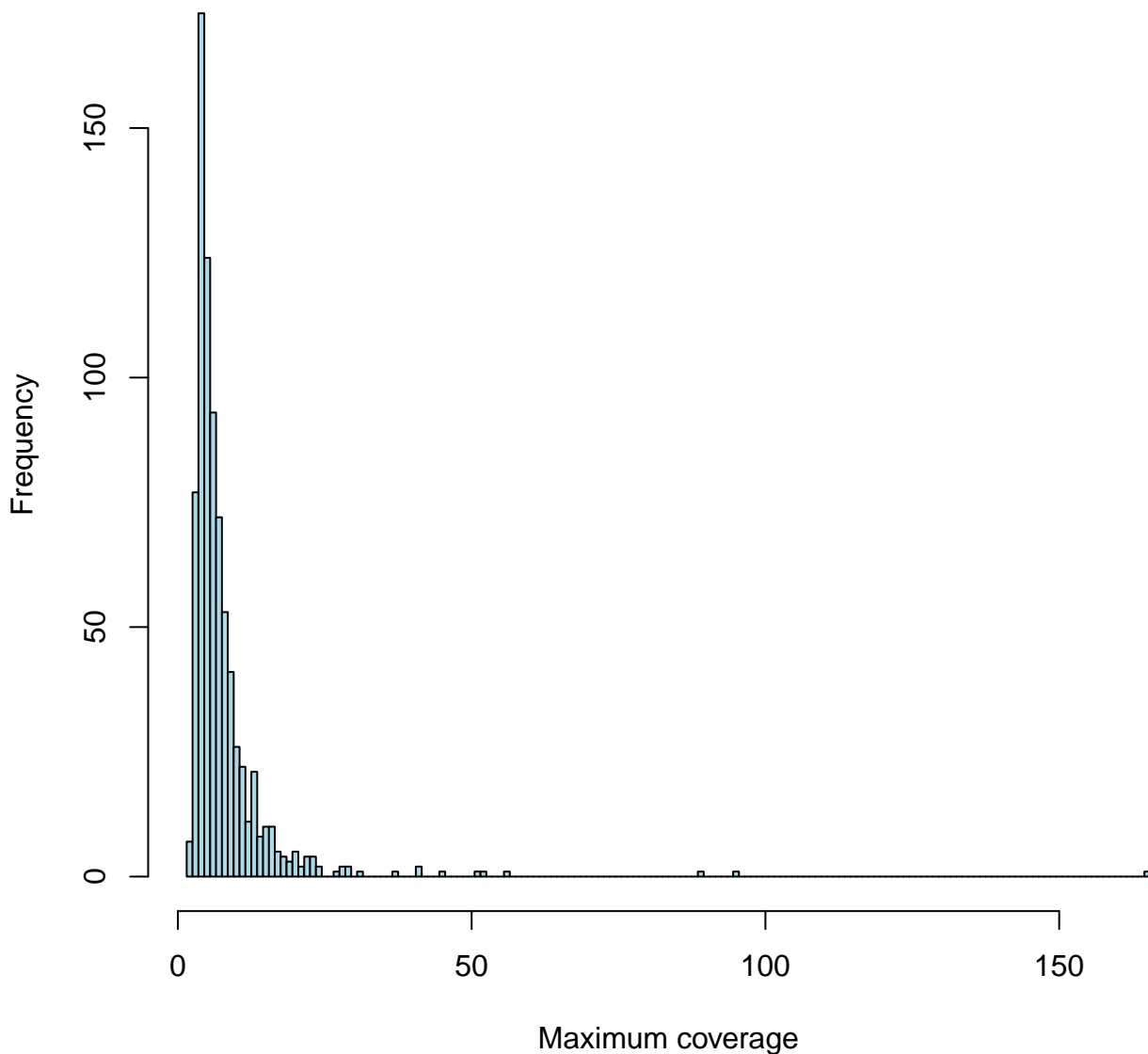
**jx in Intropolis but not "annotated" by knownGene:
52 samples; n jx: 895**



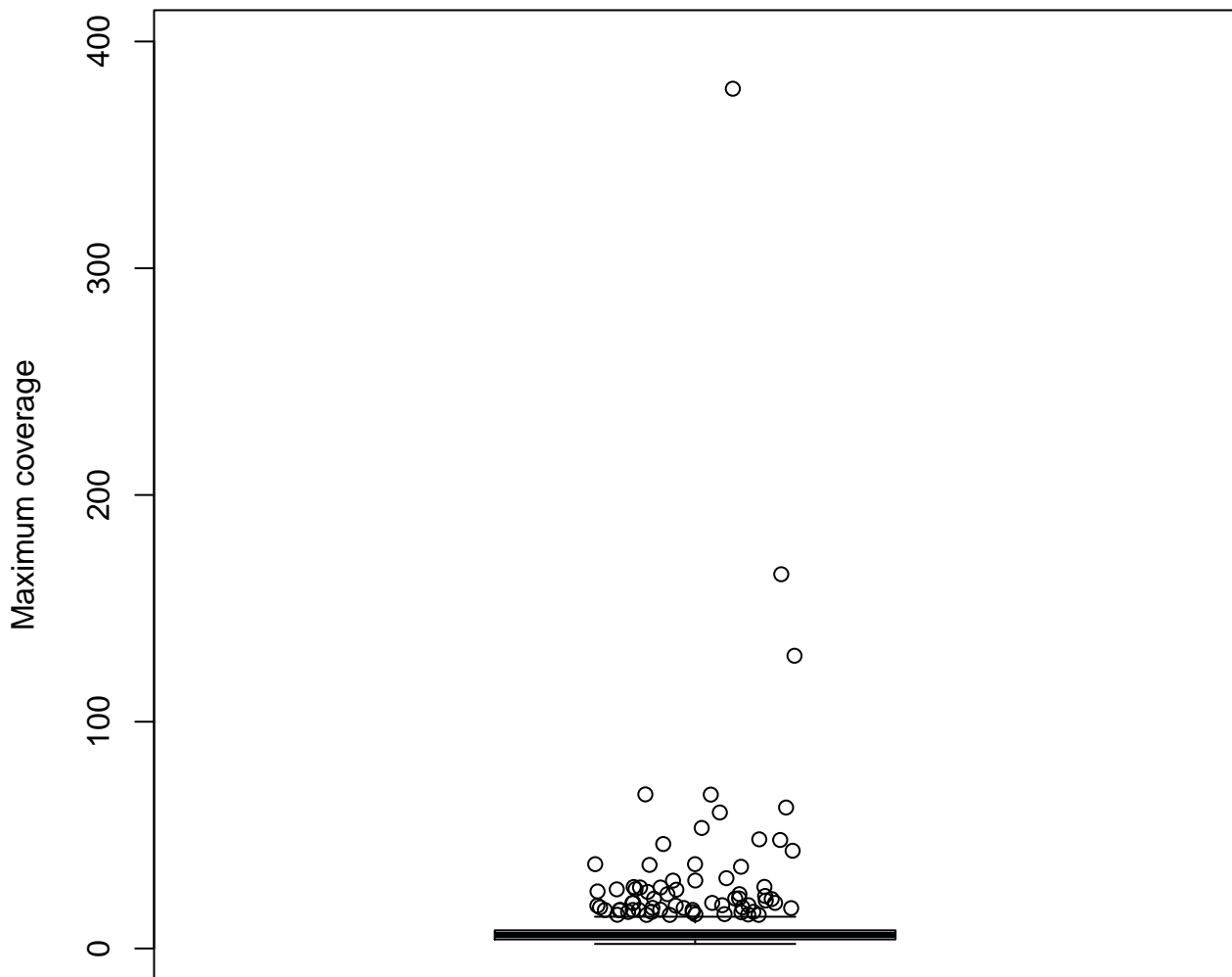
**jx in Intropolis but not "annotated" by knownGene:
53 samples; n jx: 793**



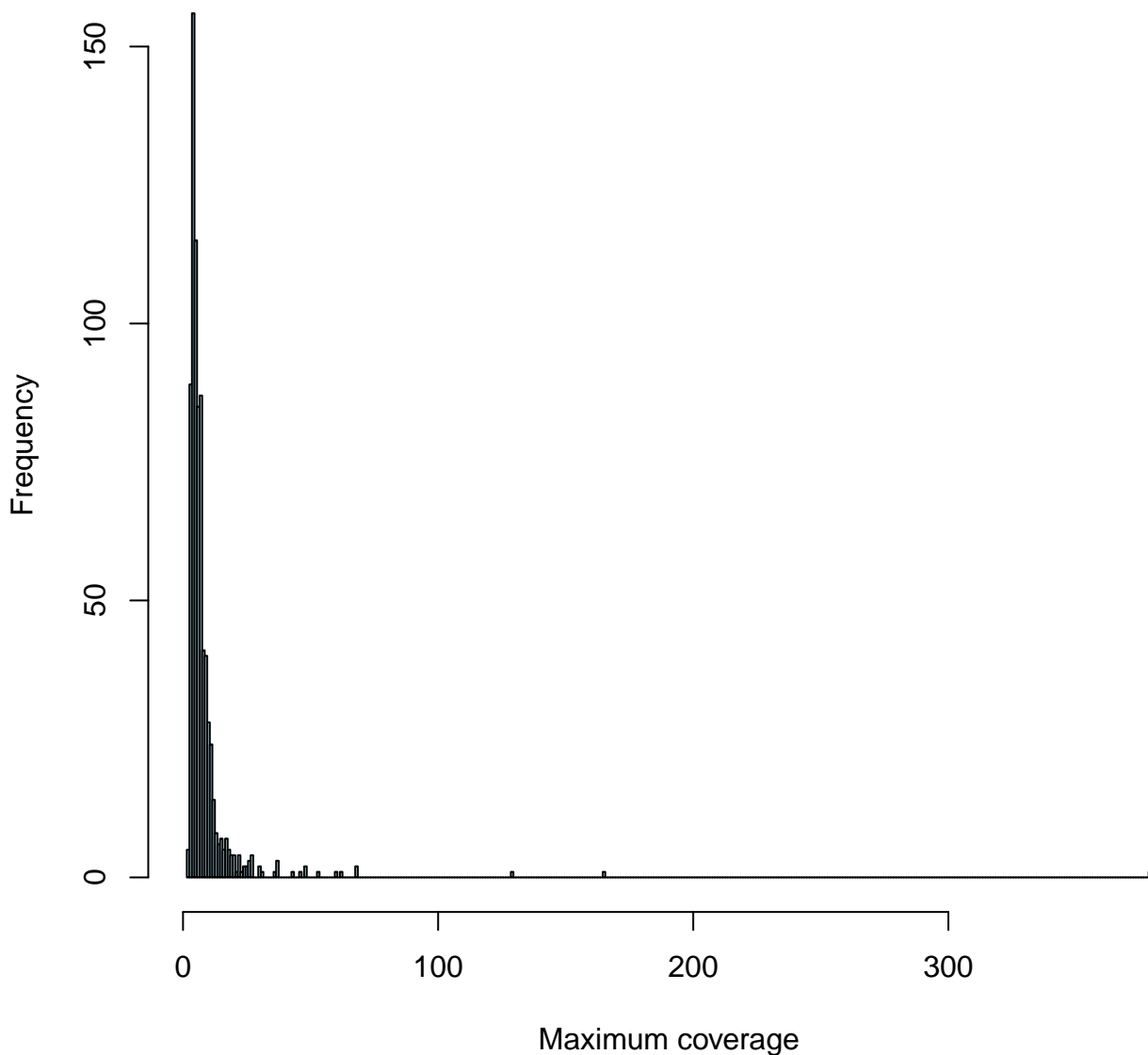
**jx in Intropolis but not "annotated" by knownGene:
53 samples; n jx: 793**



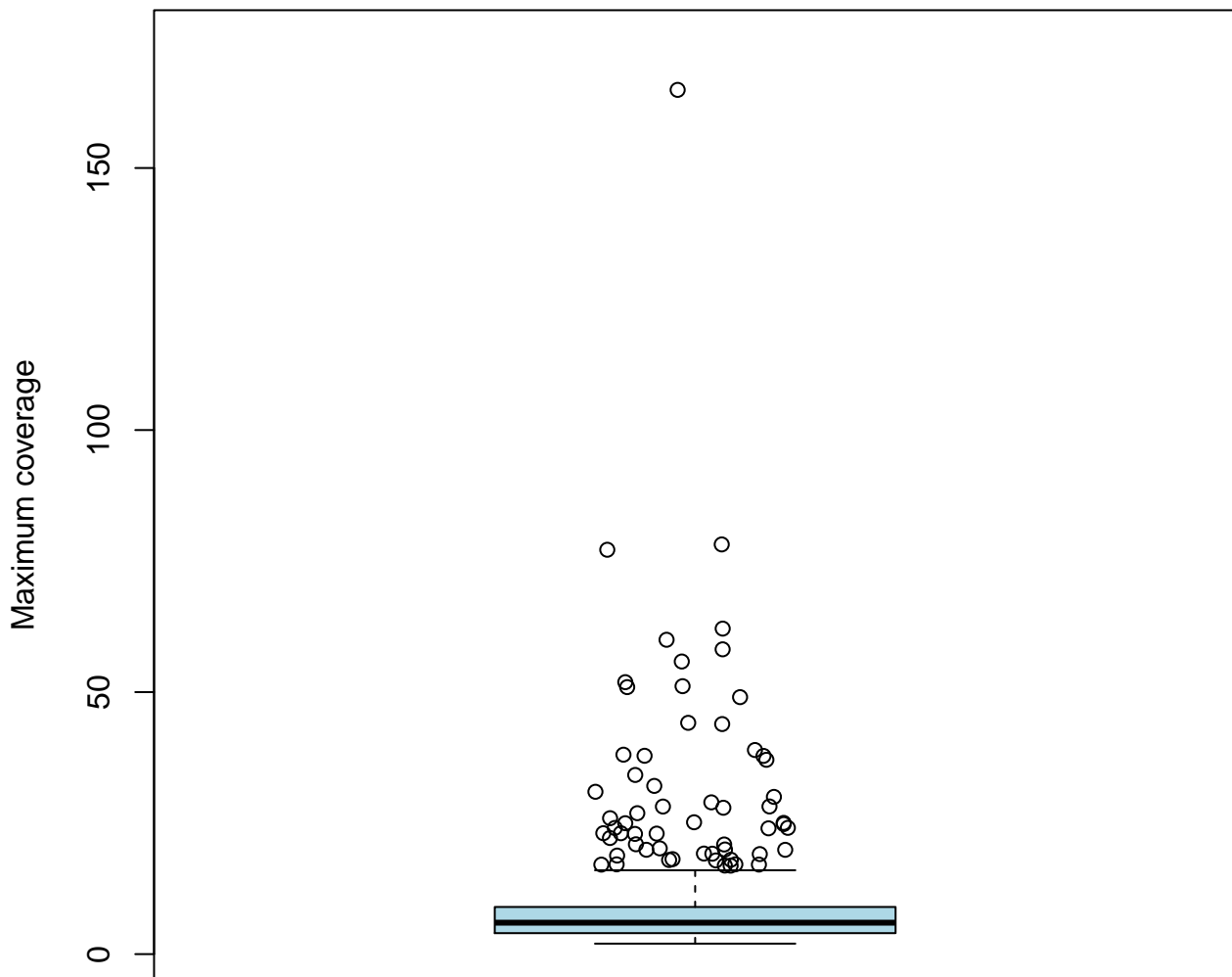
**jx in Intropolis but not "annotated" by knownGene:
54 samples; n jx: 766**



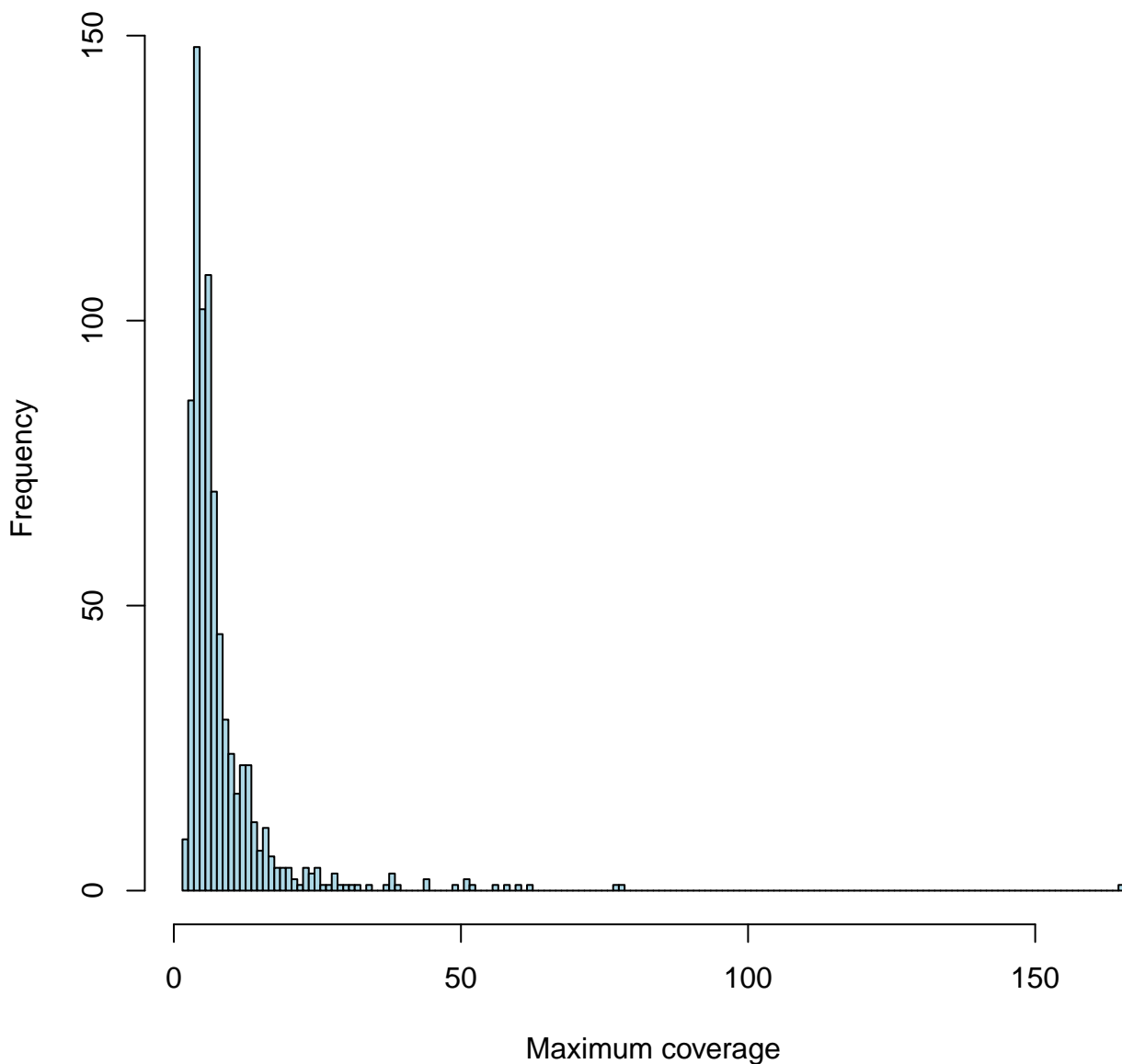
**jx in Intropolis but not "annotated" by knownGene:
54 samples; n jx: 766**



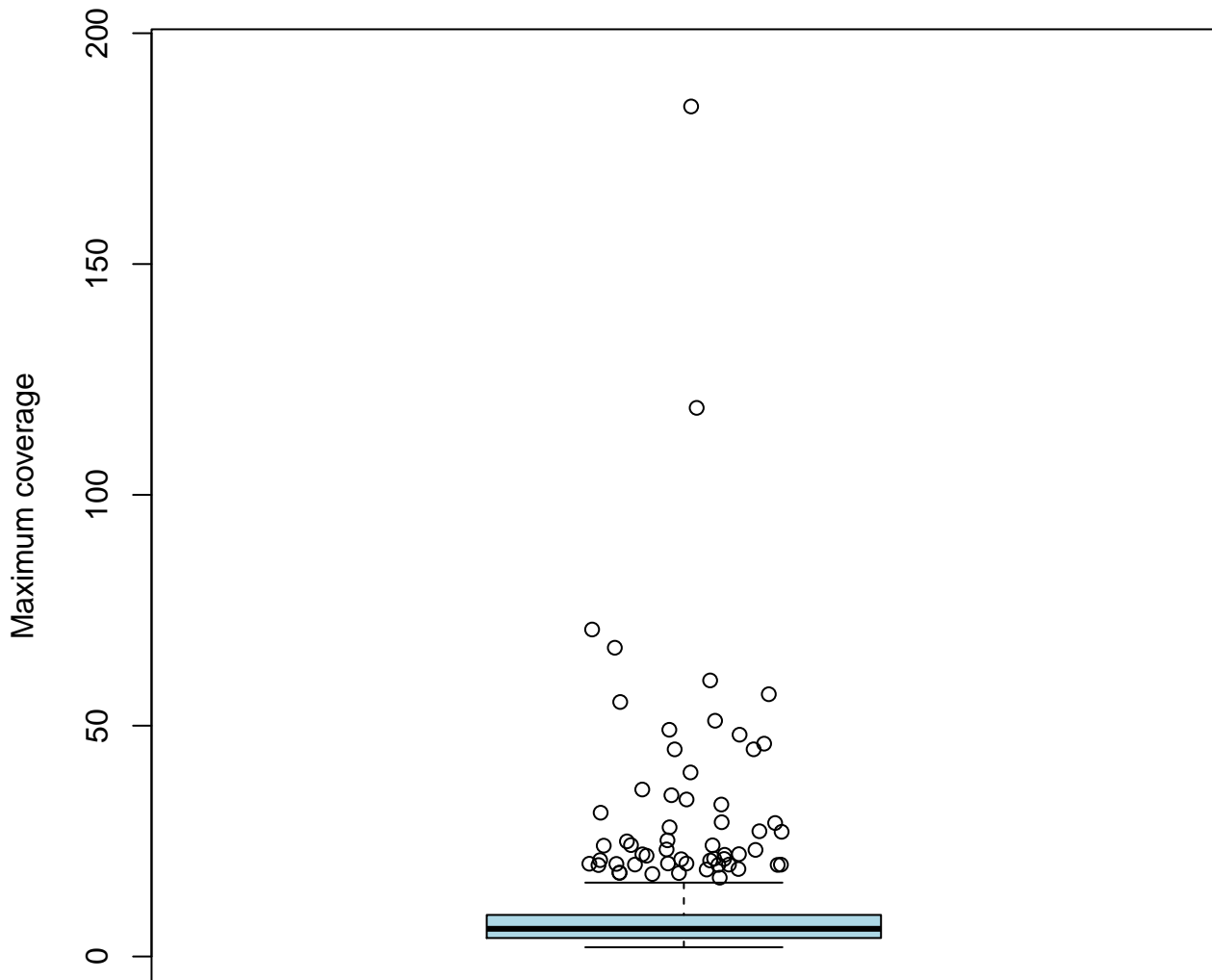
**jx in Intropolis but not "annotated" by knownGene:
55 samples; n jx: 773**



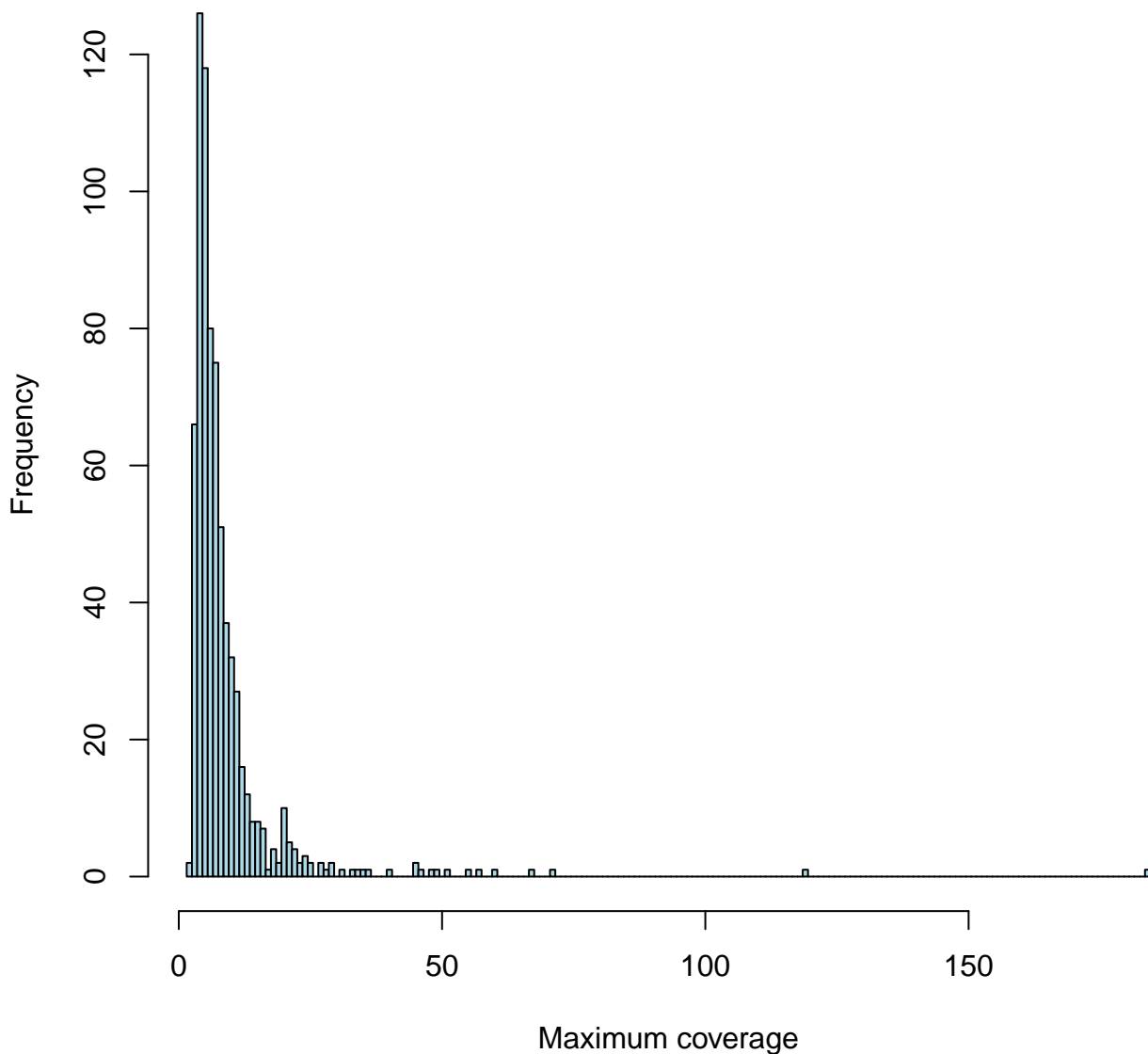
**jx in Intropolis but not "annotated" by knownGene:
55 samples; n jx: 773**



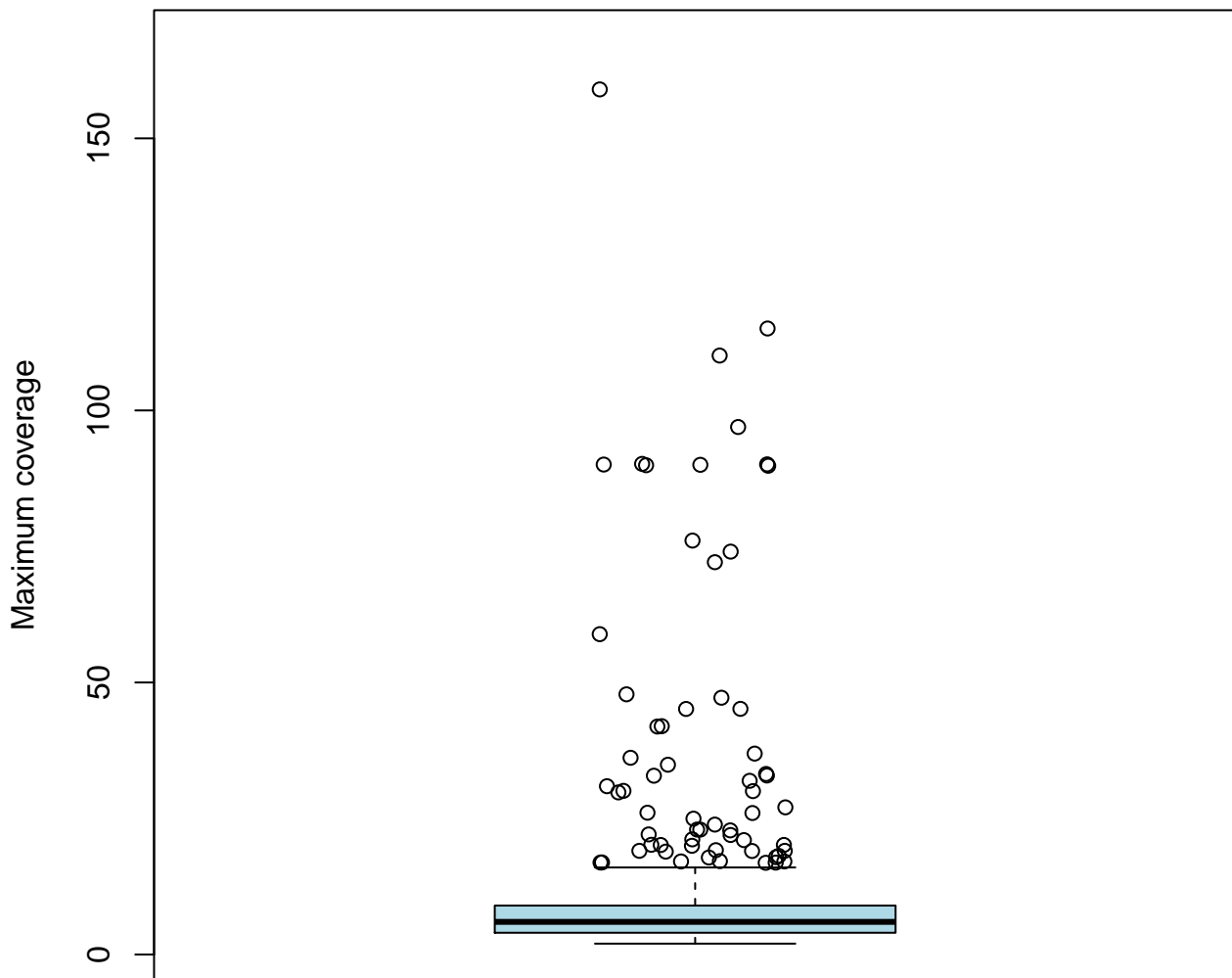
**jx in Intropolis but not "annotated" by knownGene:
56 samples; n jx: 722**



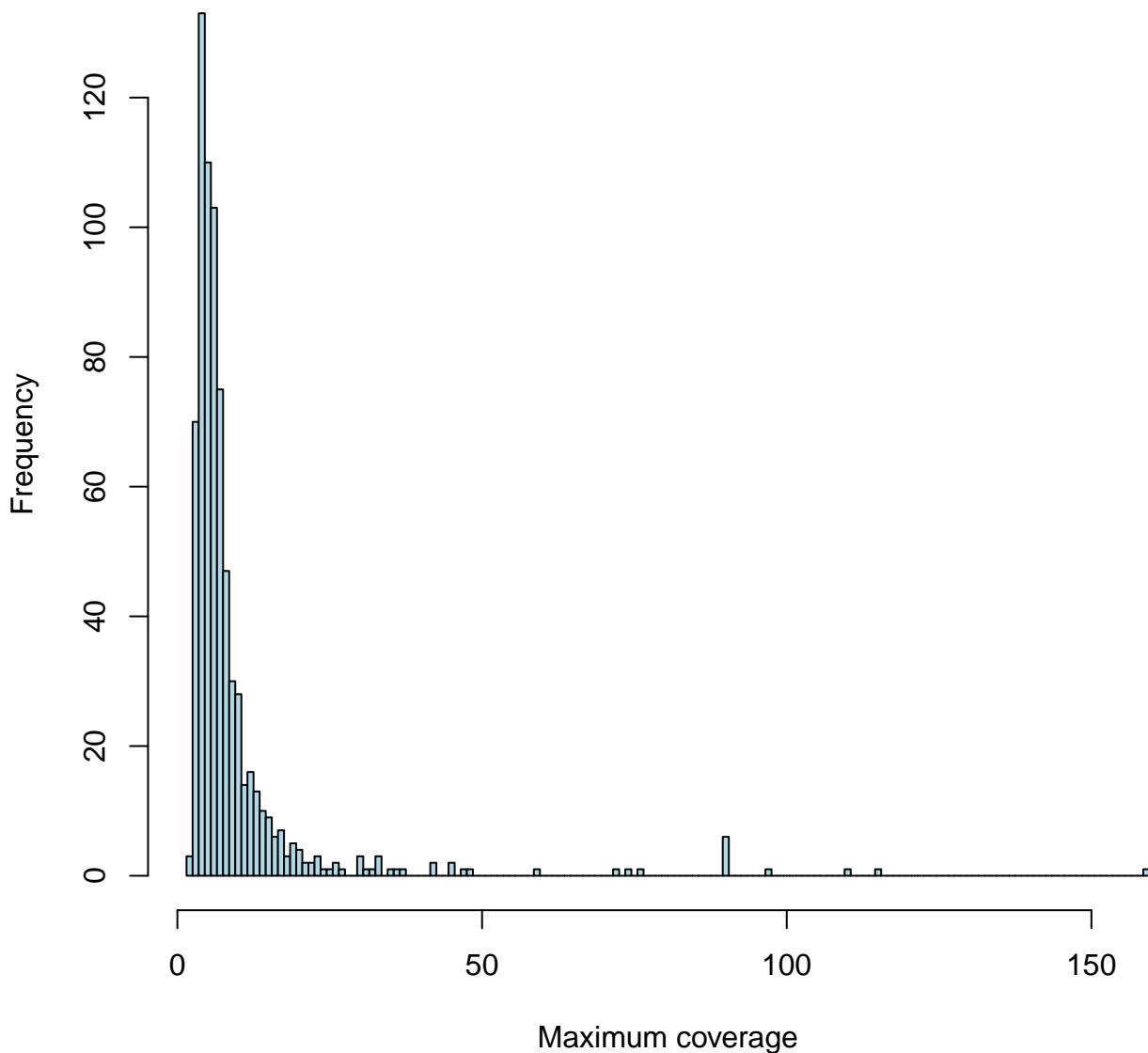
**jx in Intropolis but not "annotated" by knownGene:
56 samples; n jx: 722**



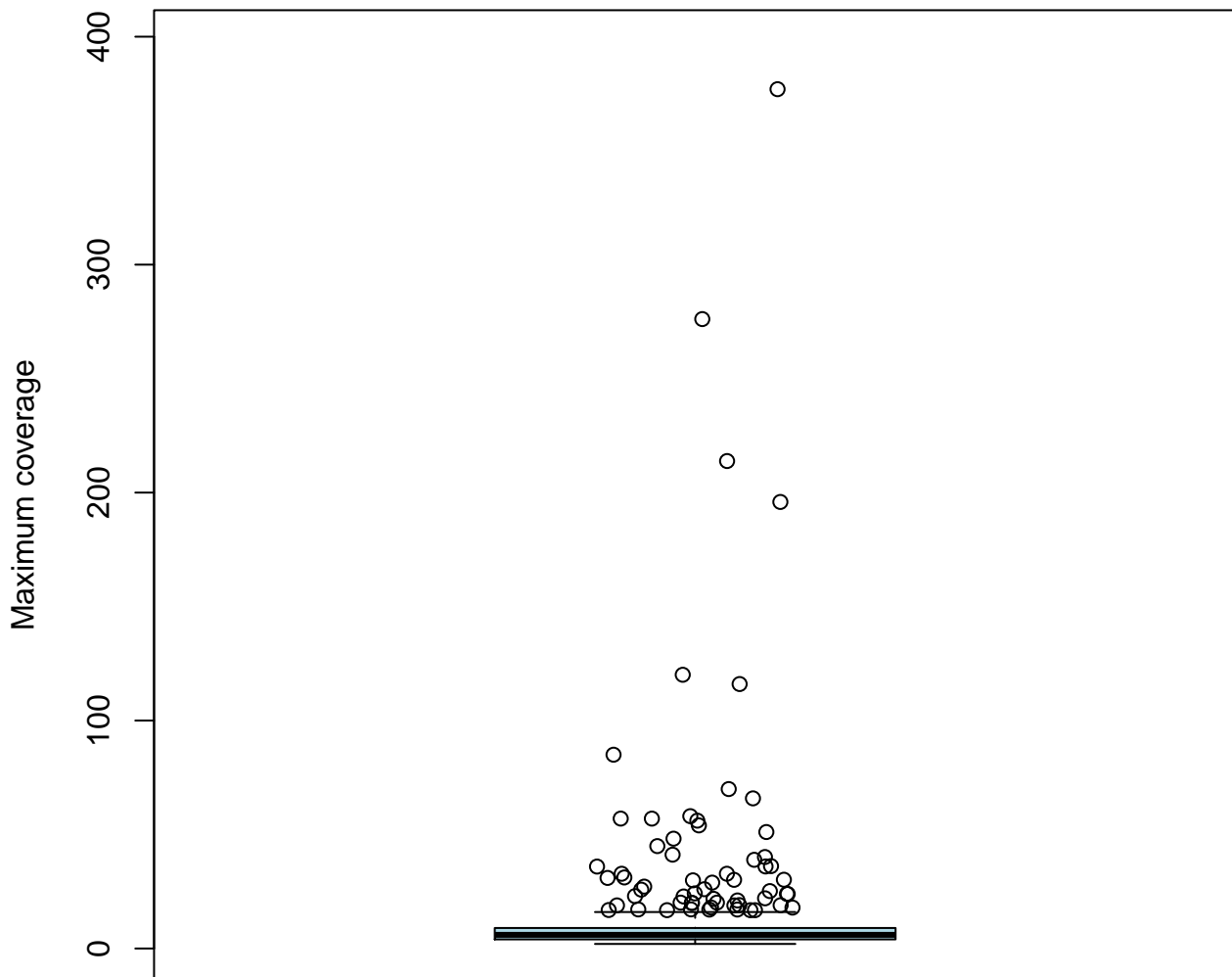
**jx in Intropolis but not "annotated" by knownGene:
57 samples; n jx: 729**



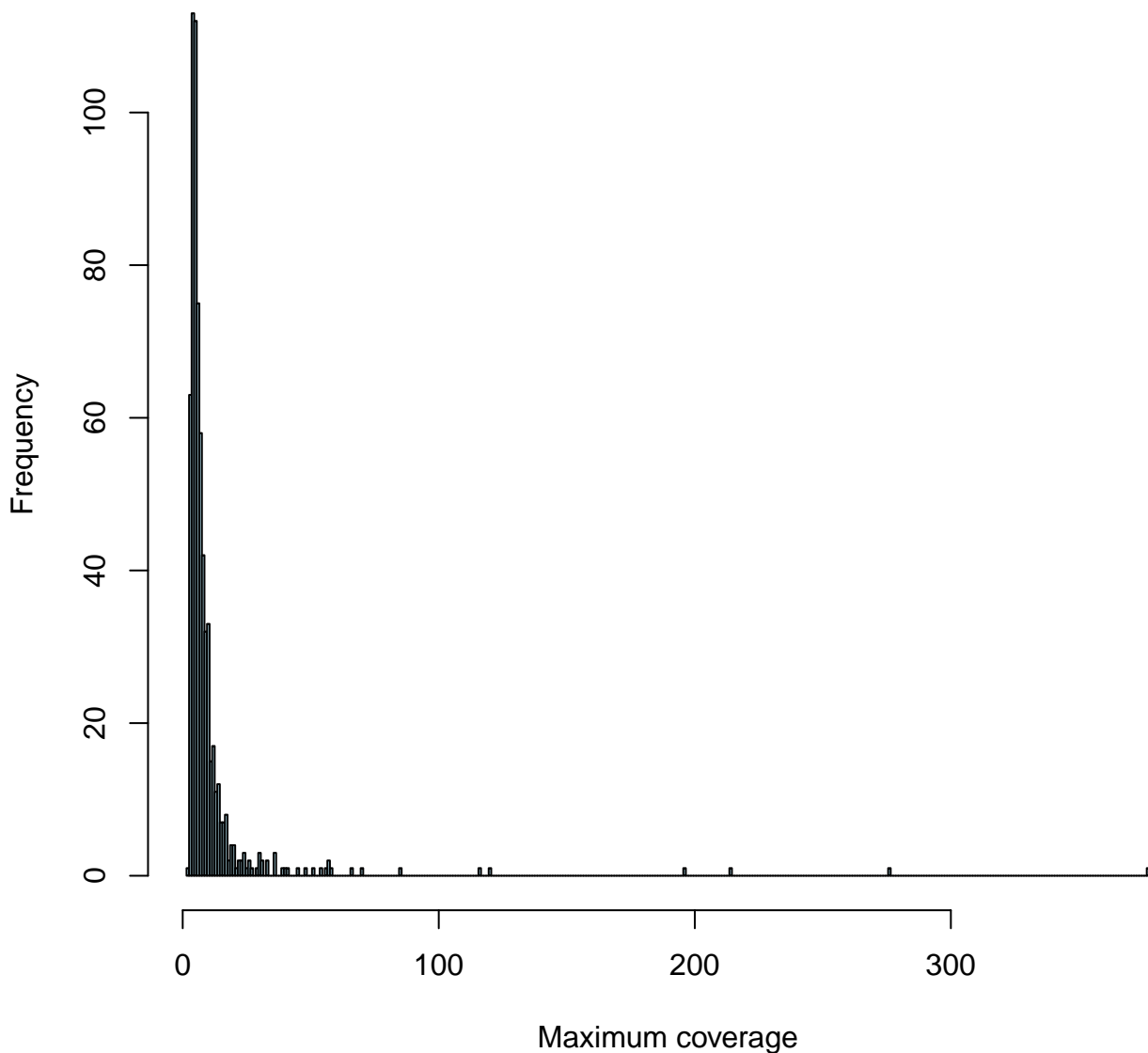
**jx in Intropolis but not "annotated" by knownGene:
57 samples; n jx: 729**



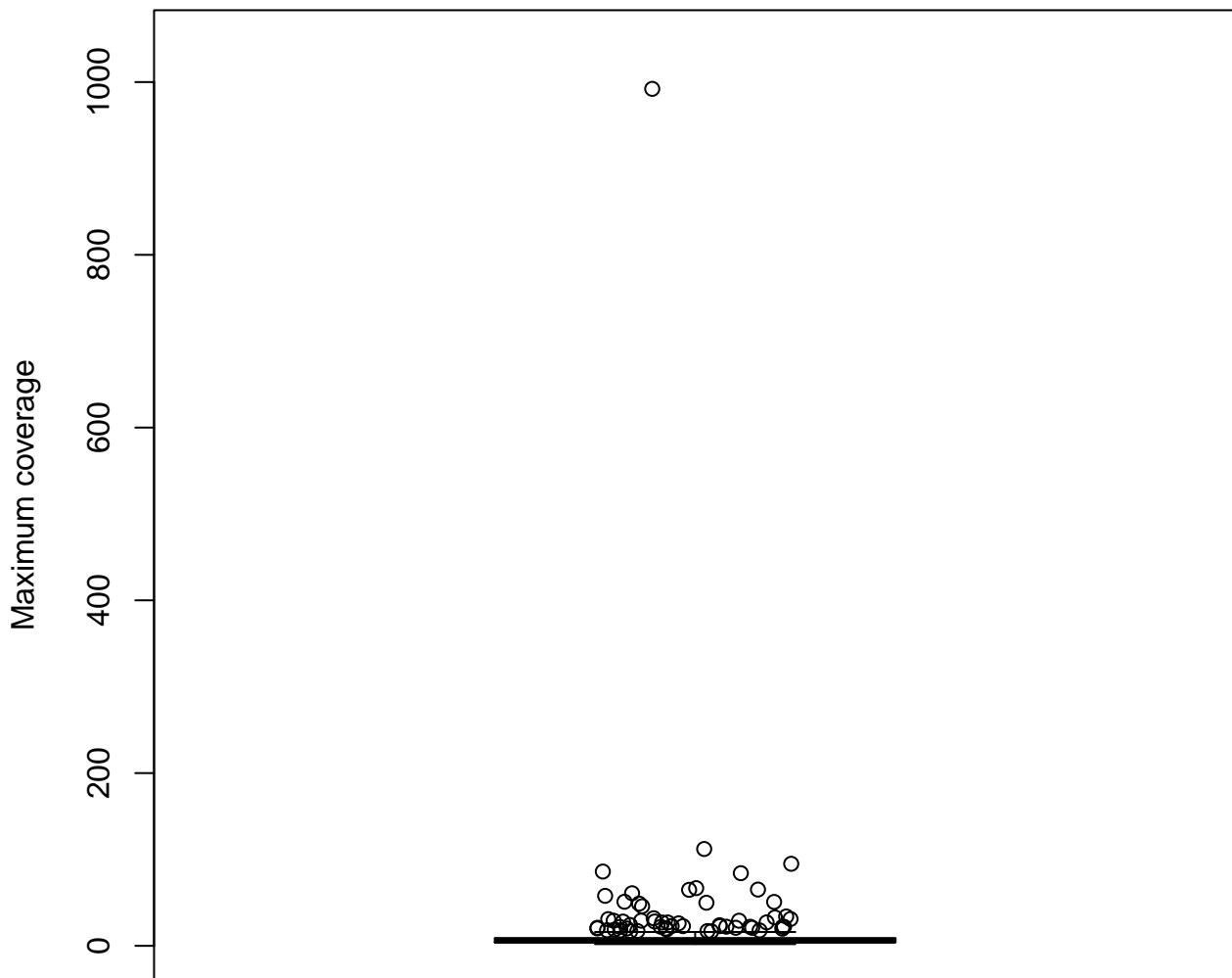
**jx in Intropolis but not "annotated" by knownGene:
58 samples; n jx: 659**



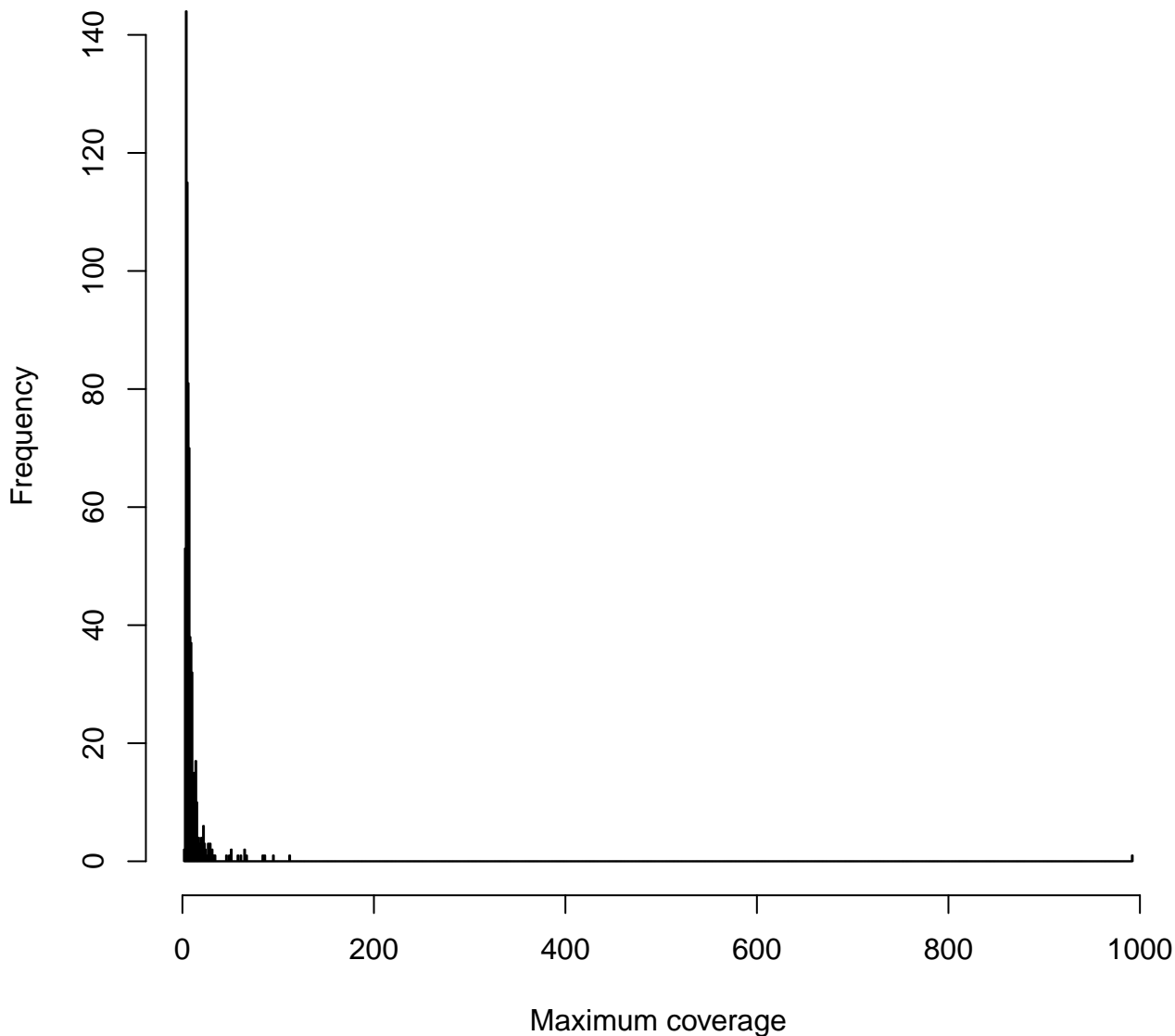
**jx in Intropolis but not "annotated" by knownGene:
58 samples; n jx: 659**



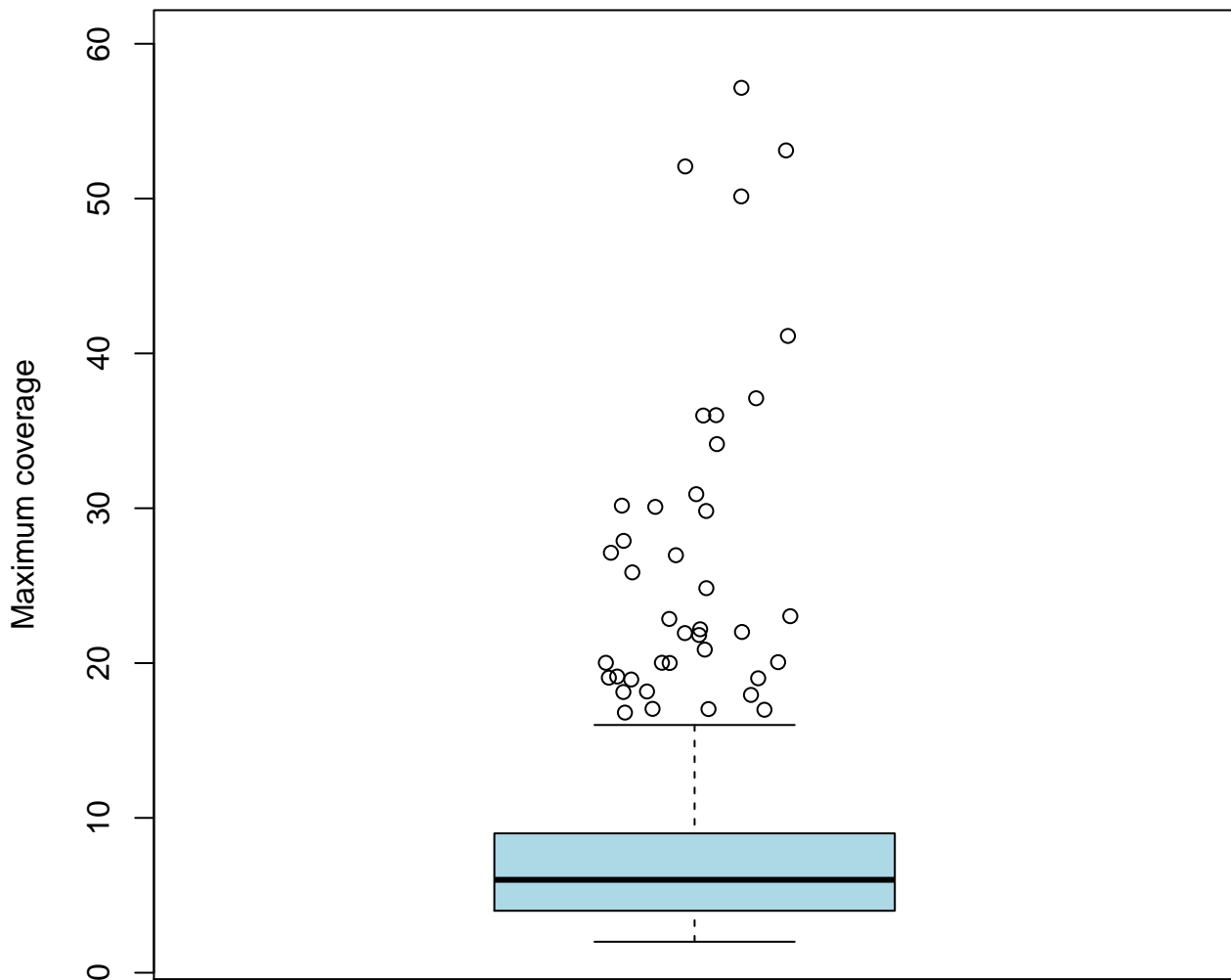
**jx in Intropolis but not "annotated" by knownGene:
59 samples; n jx: 702**



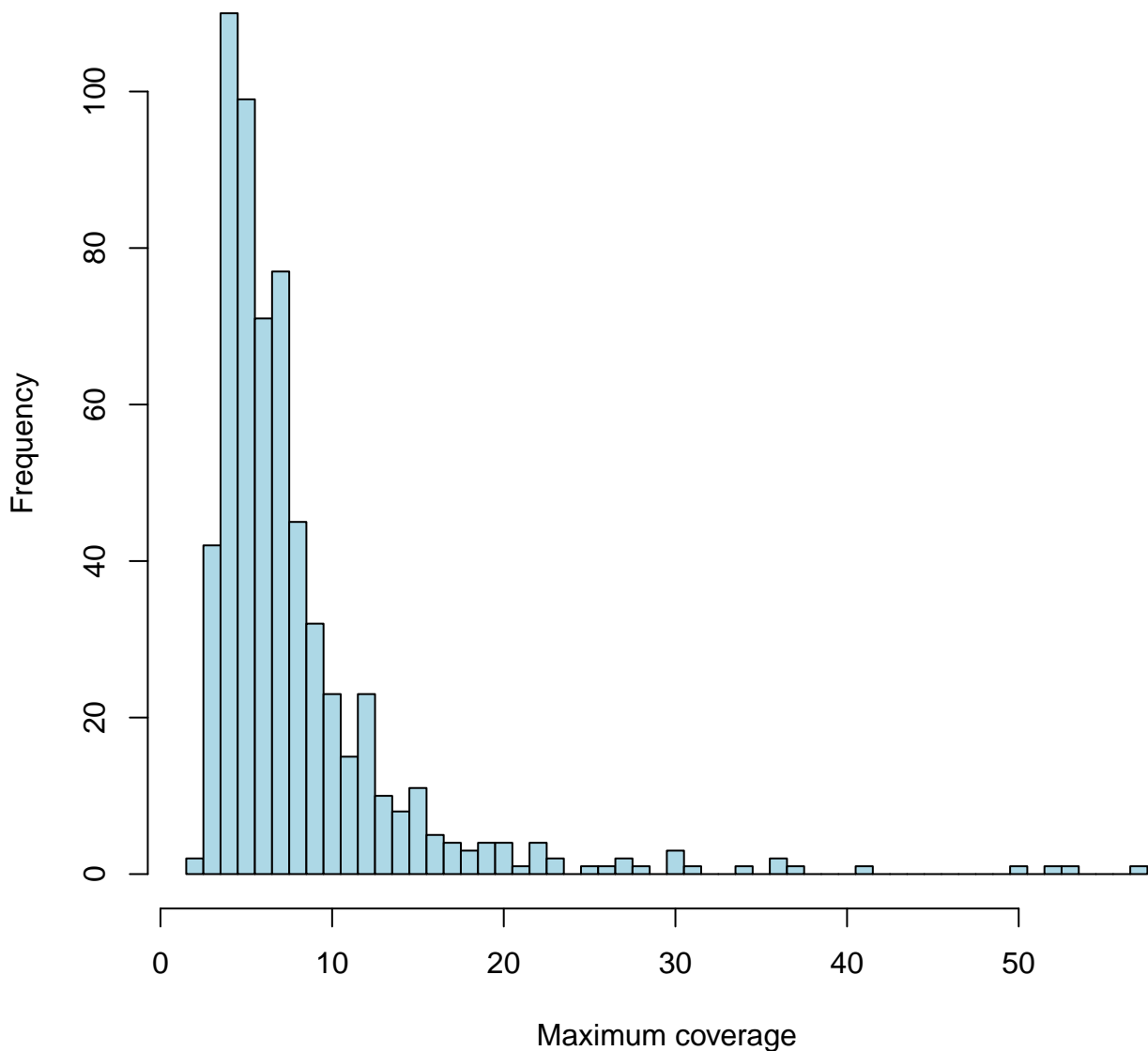
**jx in Intropolis but not "annotated" by knownGene:
59 samples; n jx: 702**



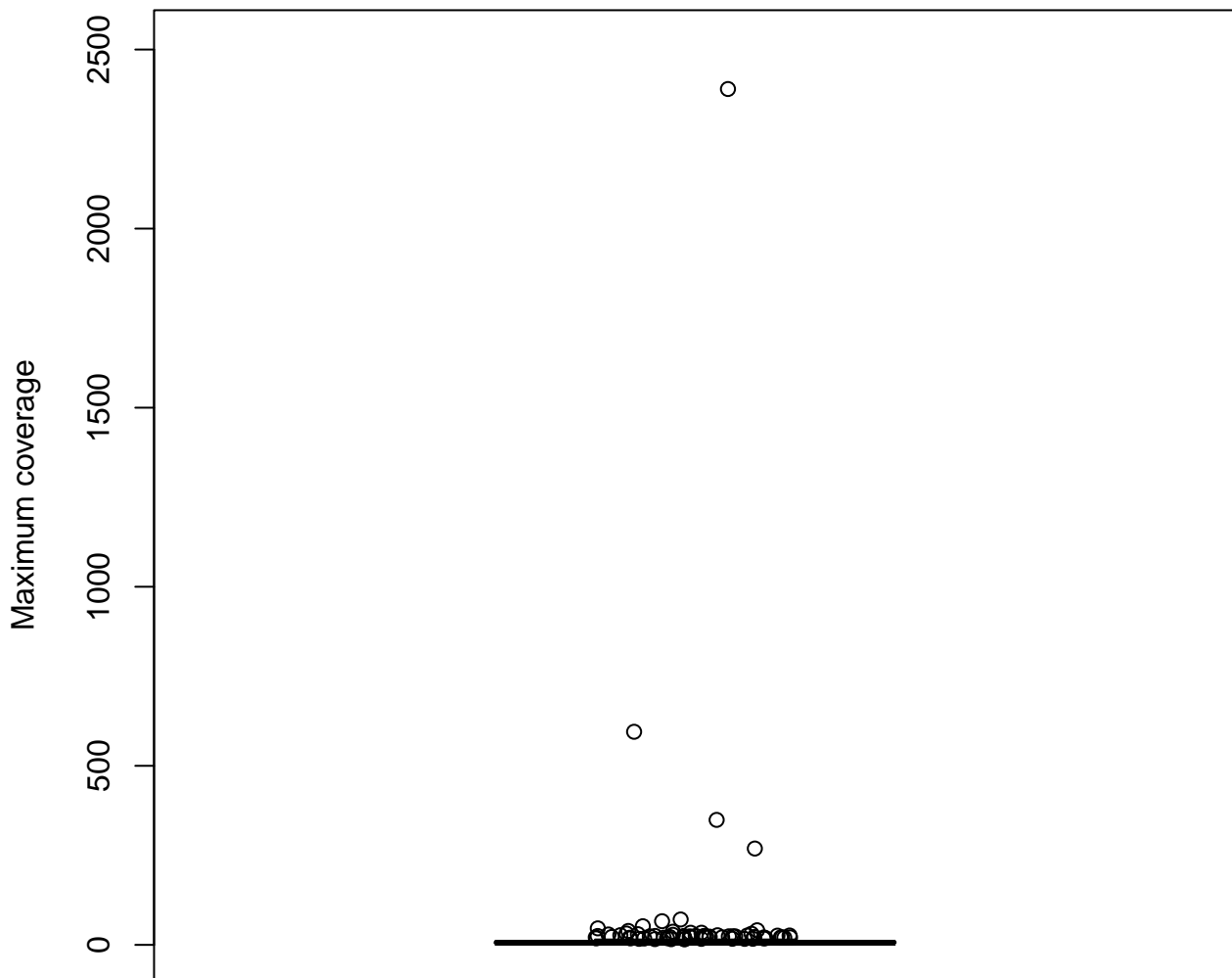
**jx in Intropolis but not "annotated" by knownGene:
60 samples; n jx: 613**



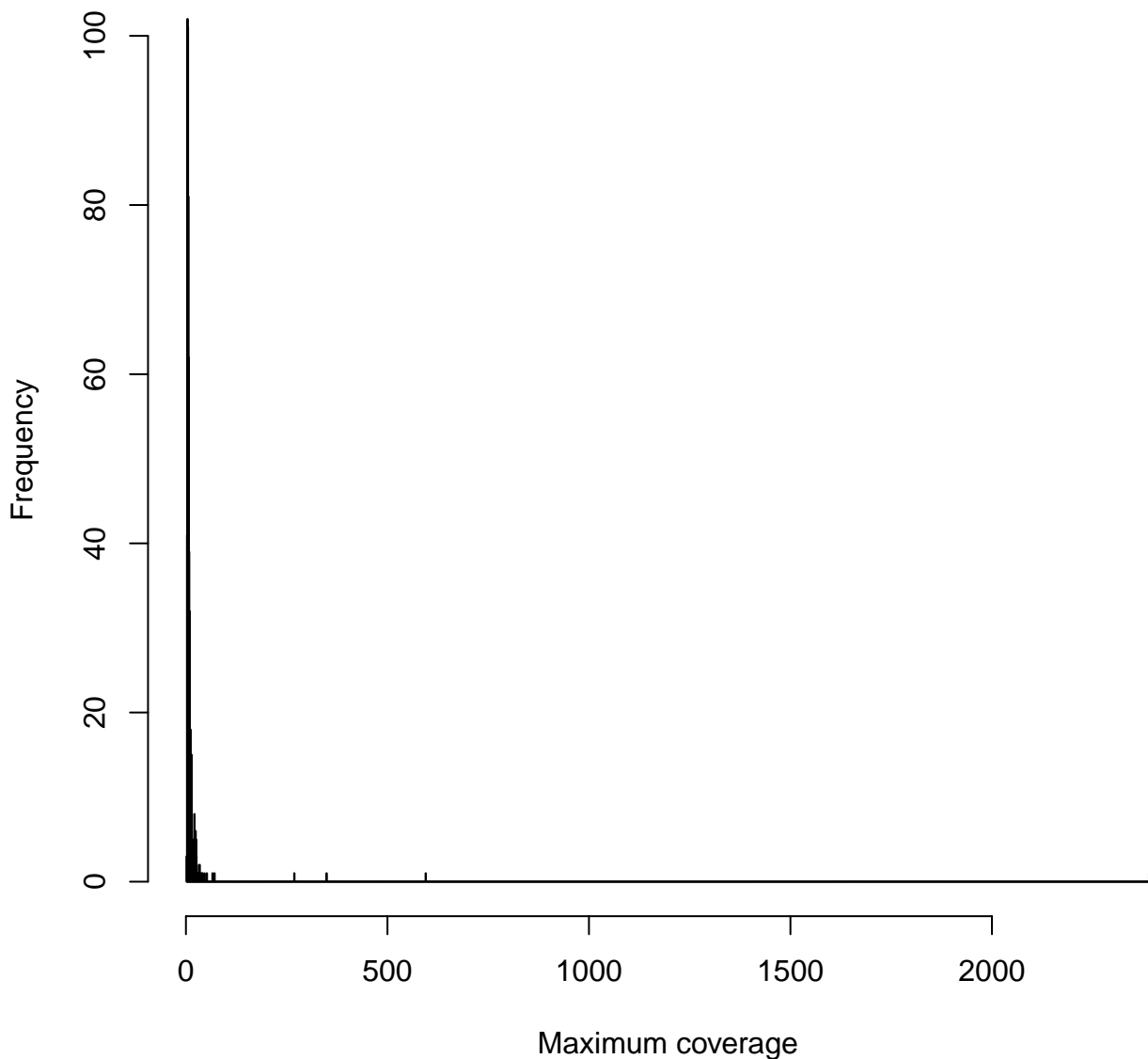
**jx in Intropolis but not "annotated" by knownGene:
60 samples; n jx: 613**



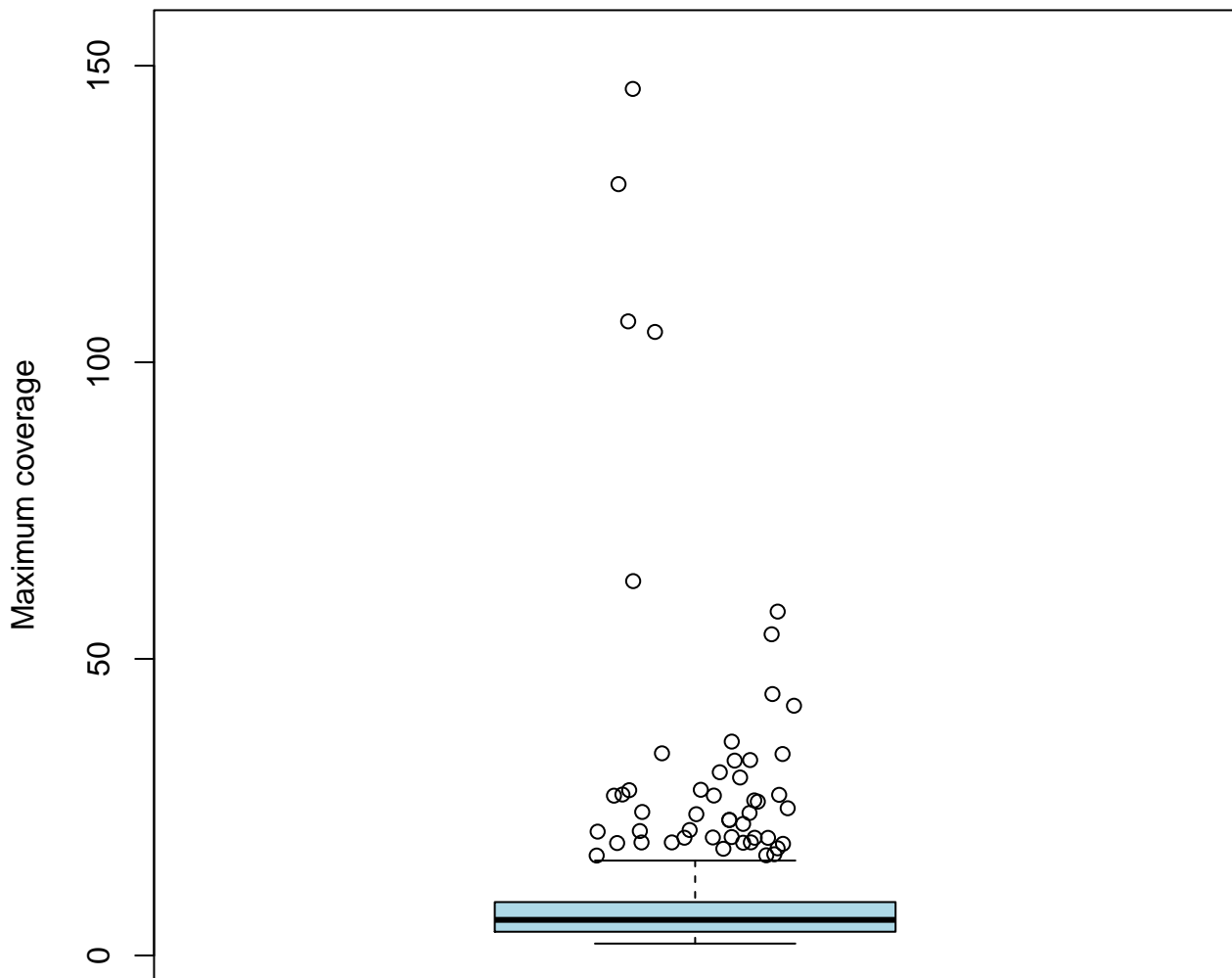
**jx in Intropolis but not "annotated" by knownGene:
61 samples; n jx: 611**



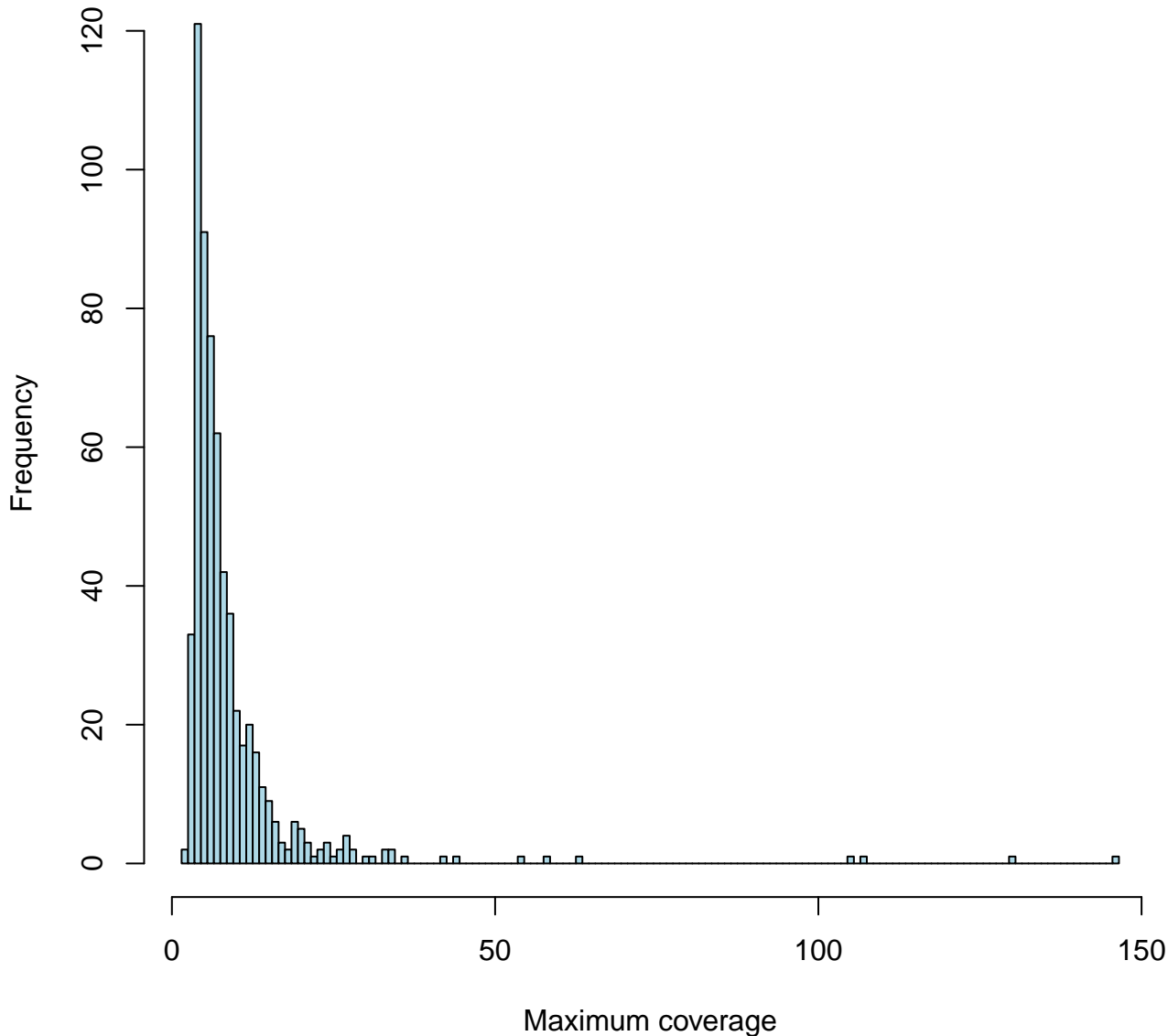
**jx in Intropolis but not "annotated" by knownGene:
61 samples; n jx: 611**



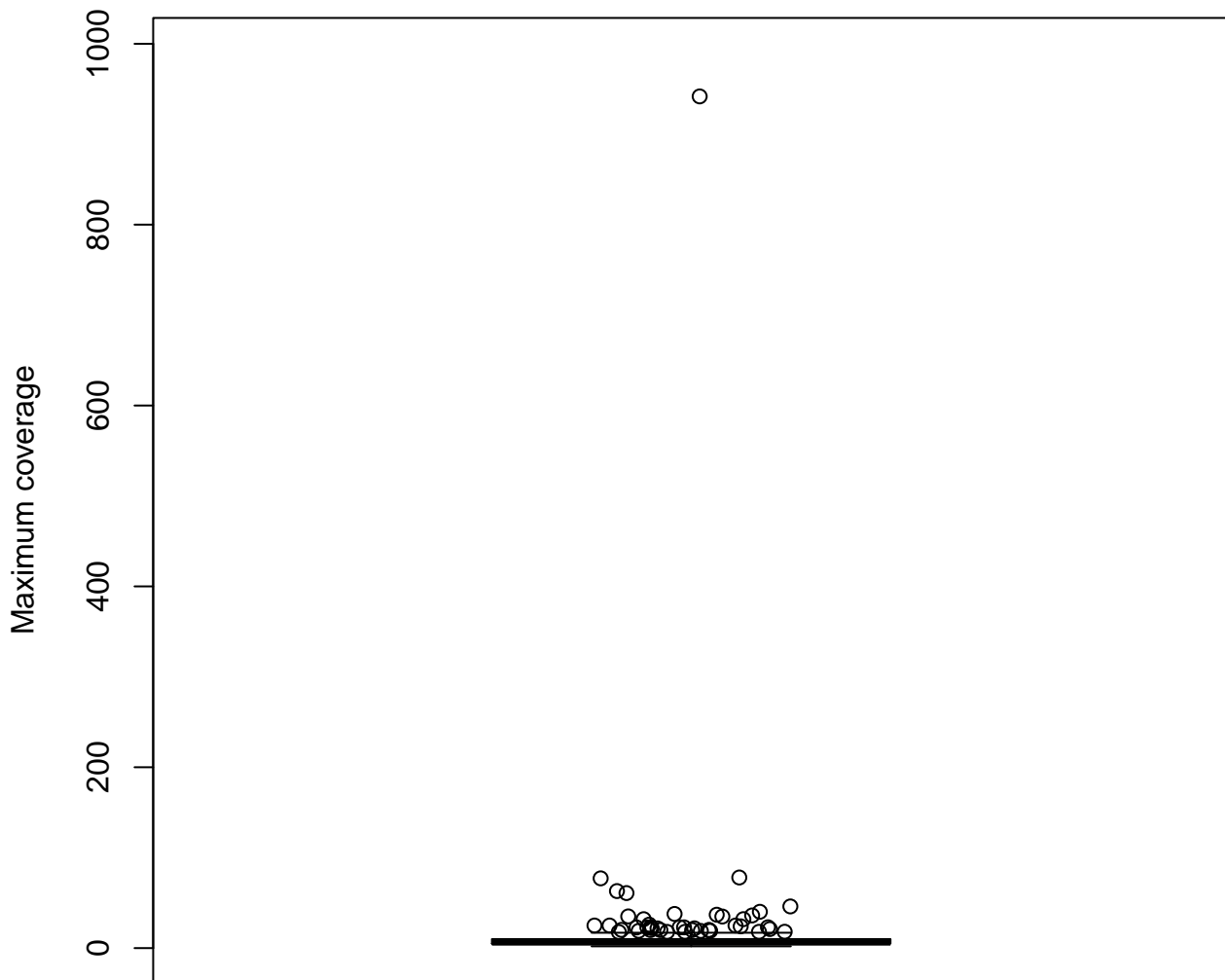
**jx in Intropolis but not "annotated" by knownGene:
62 samples; n jx: 614**



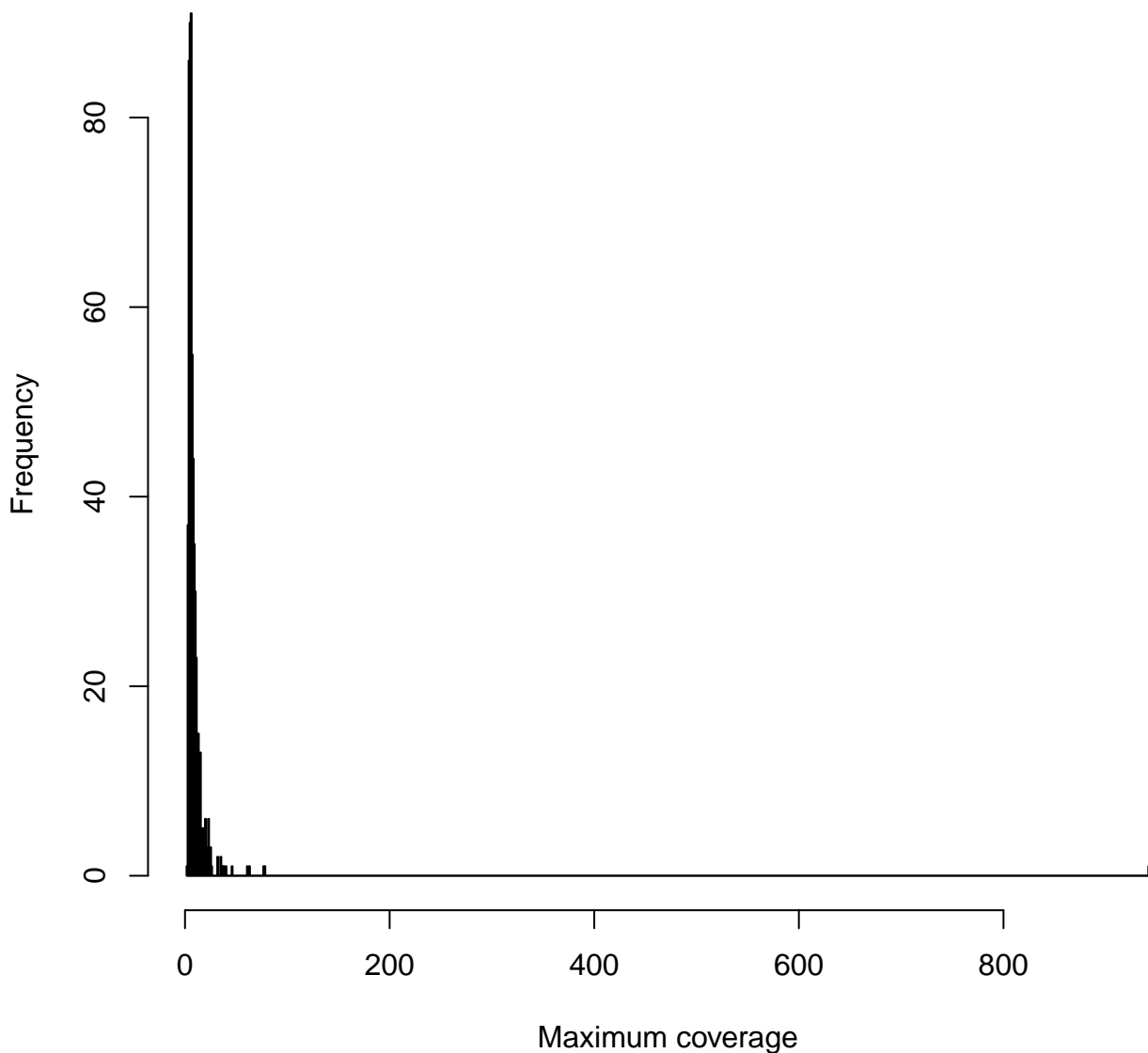
**jx in Intropolis but not "annotated" by knownGene:
62 samples; n jx: 614**



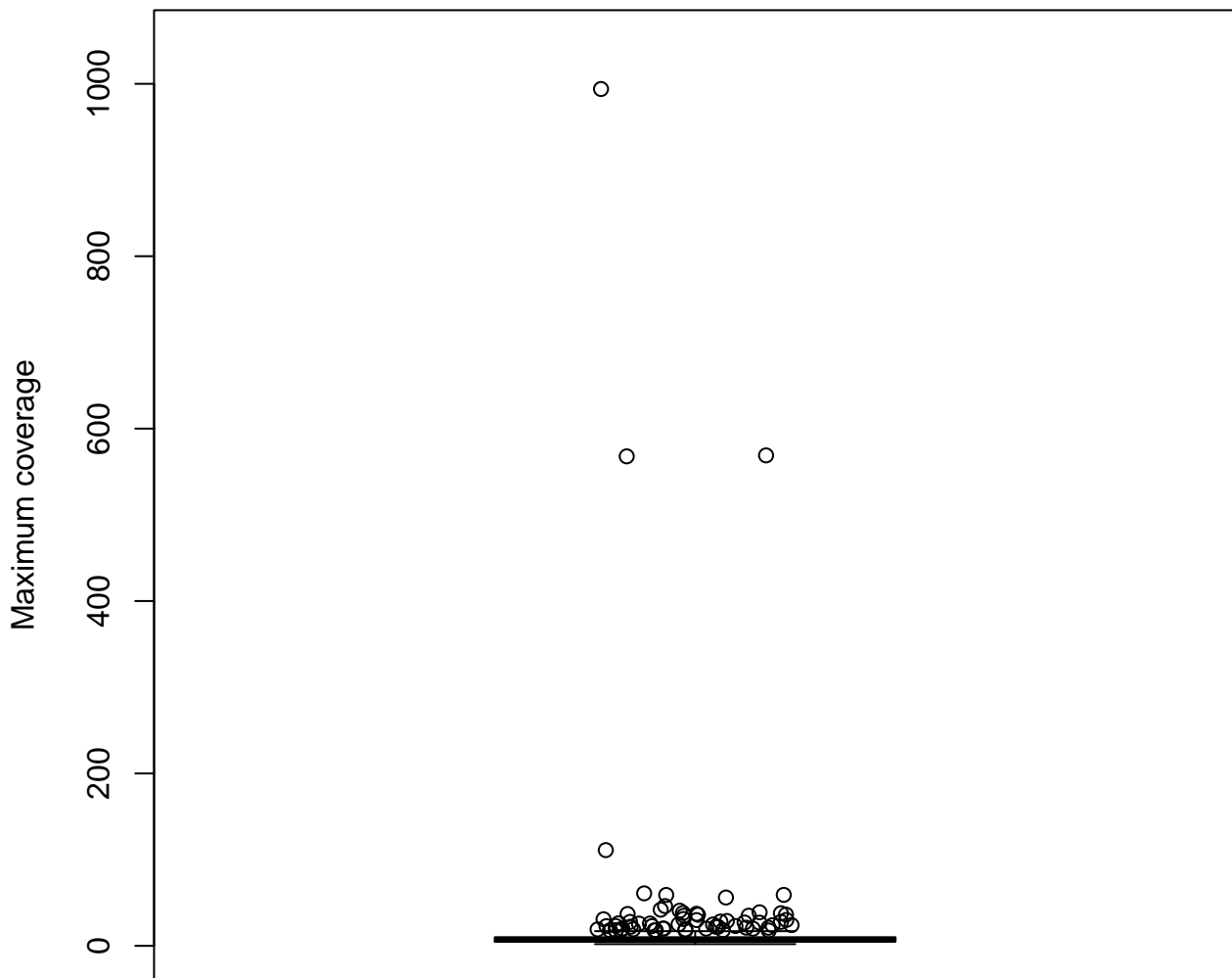
**jx in Intropolis but not "annotated" by knownGene:
63 samples; n jx: 593**



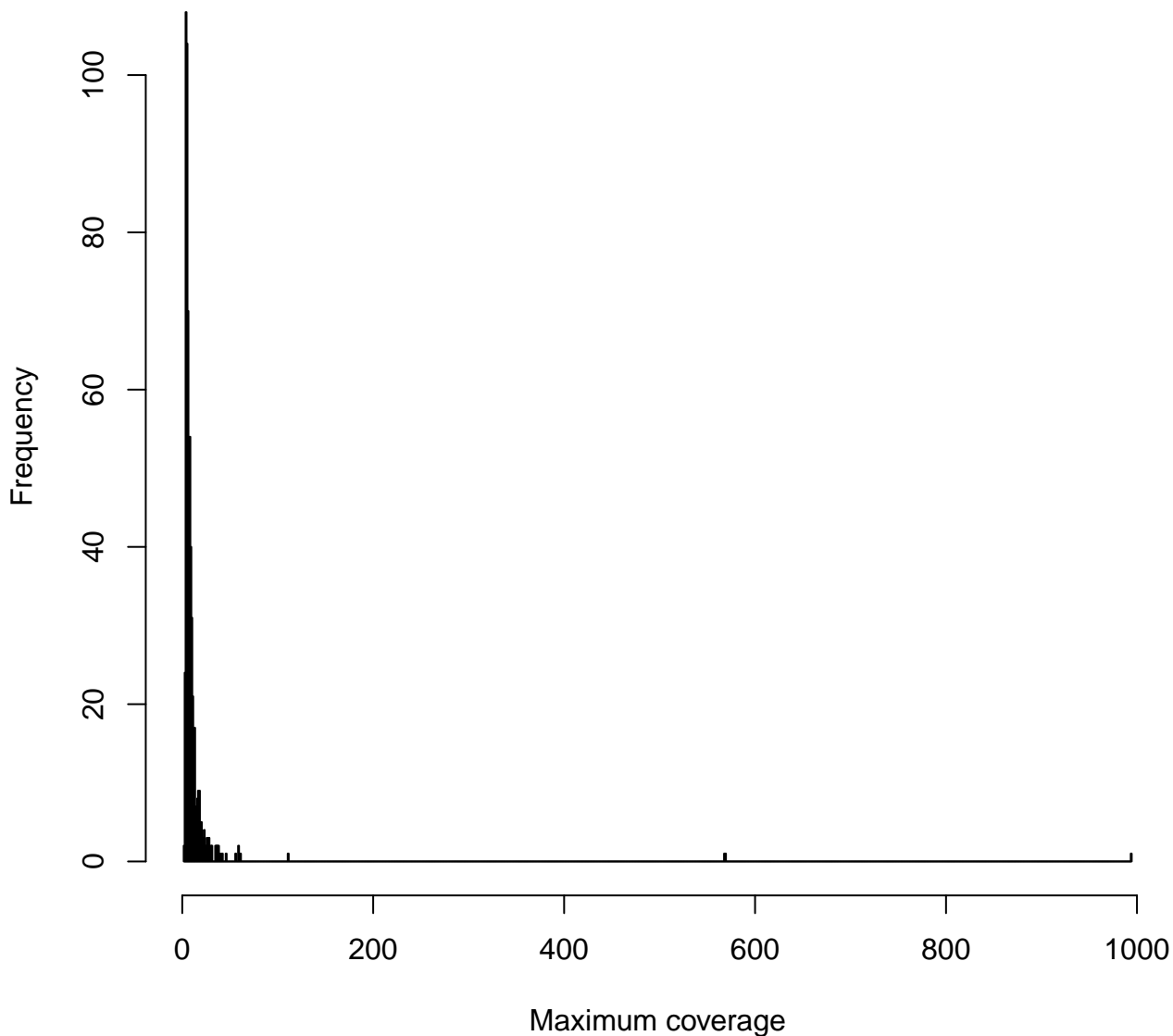
**jx in Intropolis but not "annotated" by knownGene:
63 samples; n jx: 593**



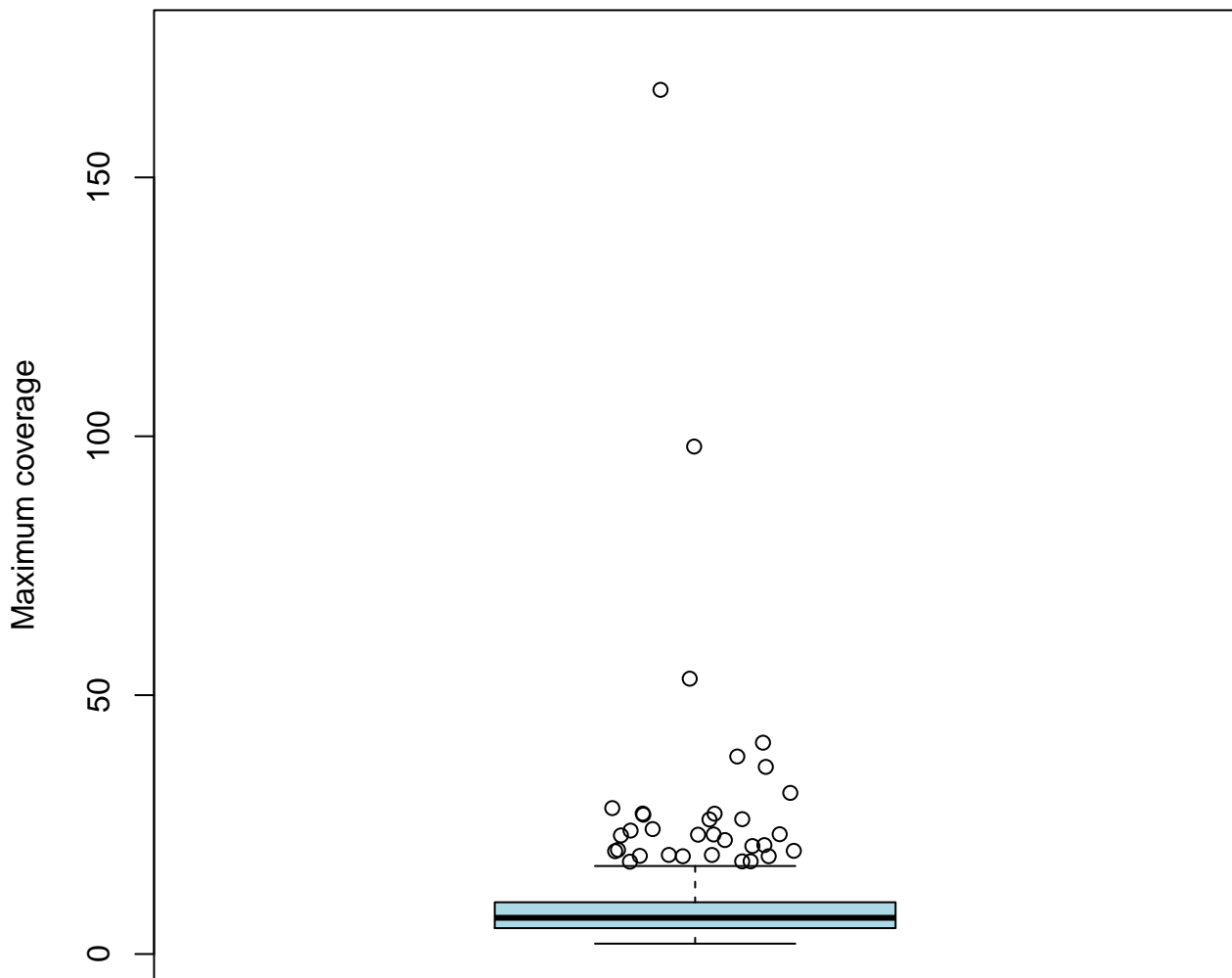
**jx in Intropolis but not "annotated" by knownGene:
64 samples; n jx: 620**



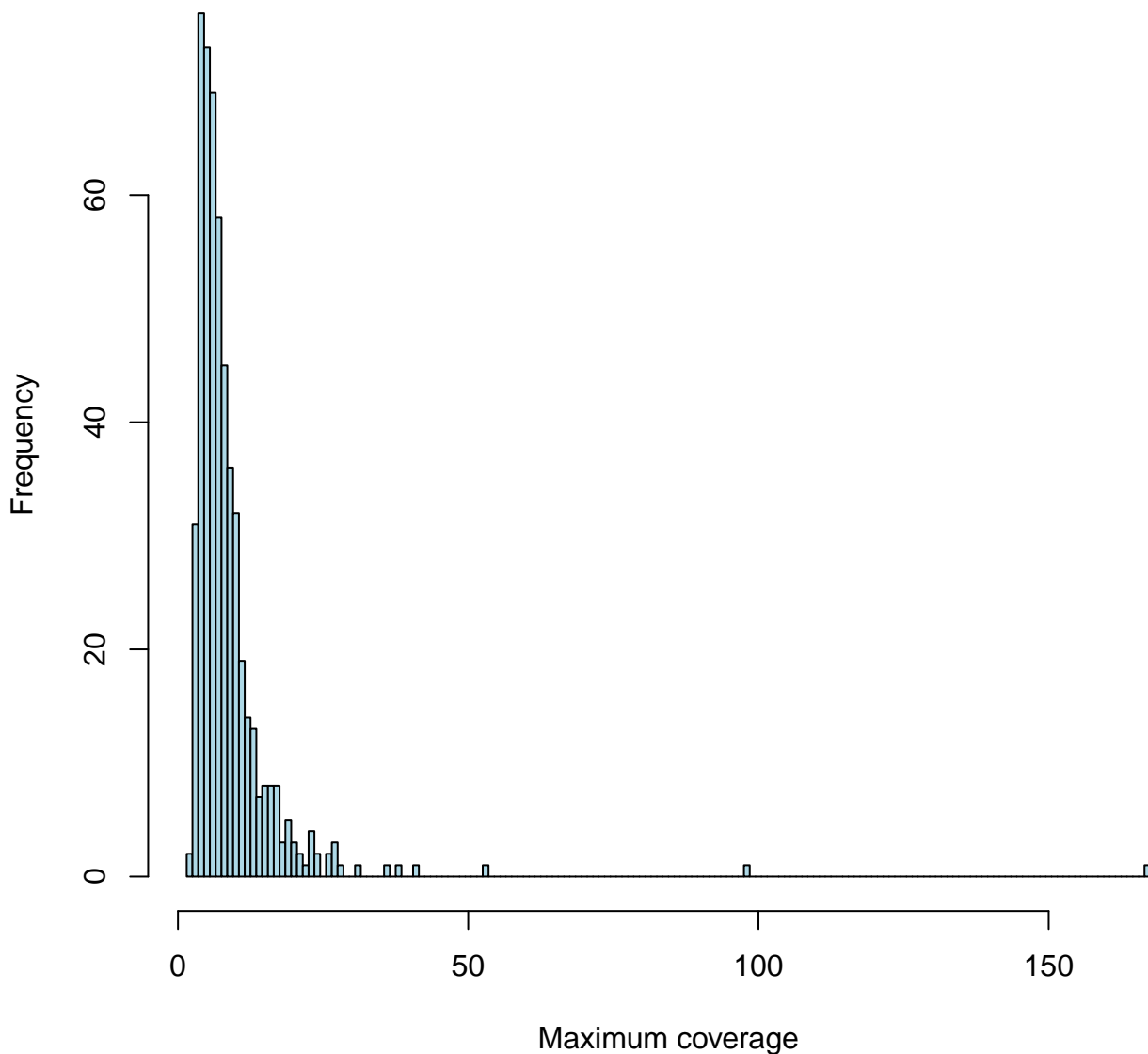
**jx in Intropolis but not "annotated" by knownGene:
64 samples; n jx: 620**



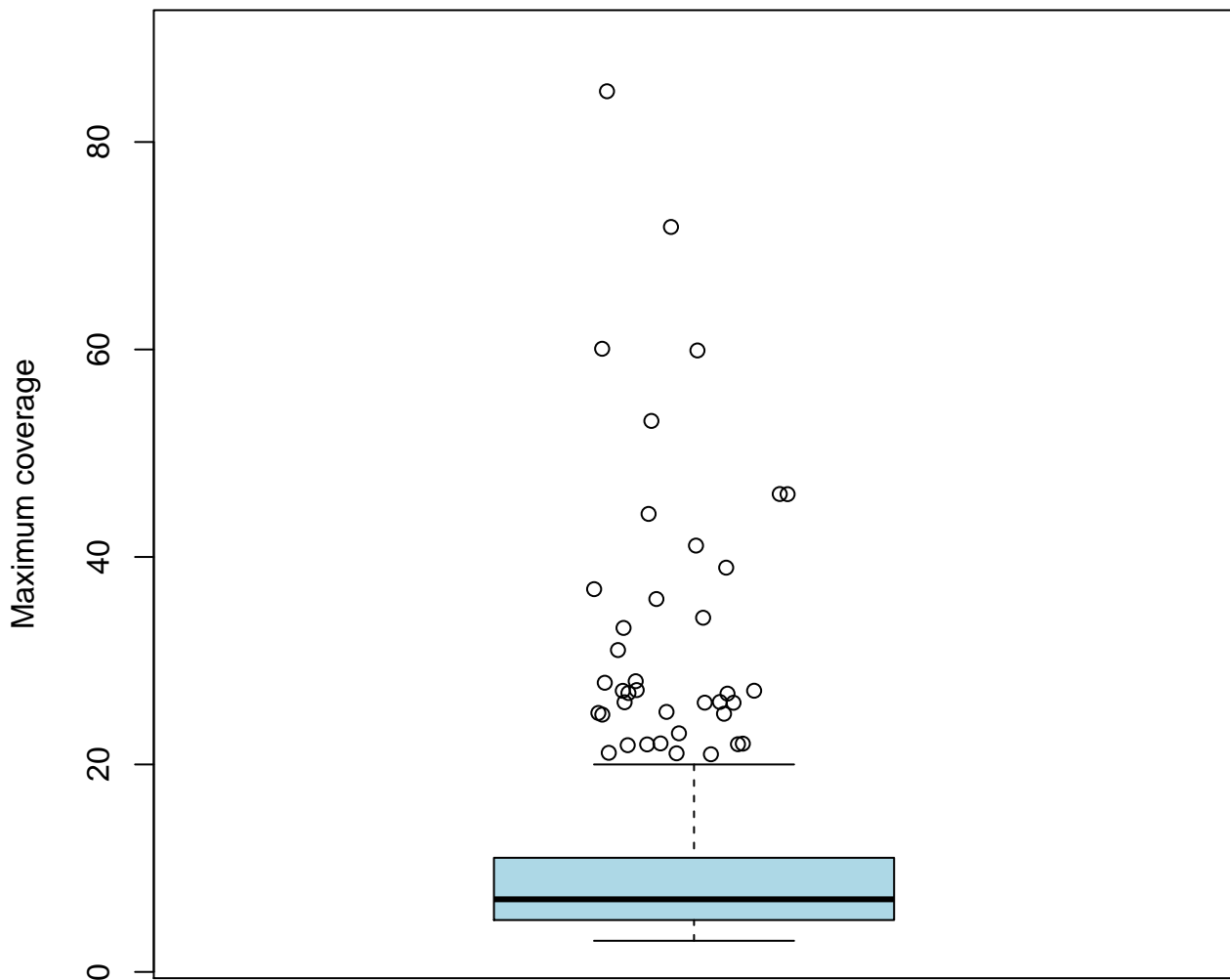
**jx in Intropolis but not "annotated" by knownGene:
65 samples; n jx: 532**



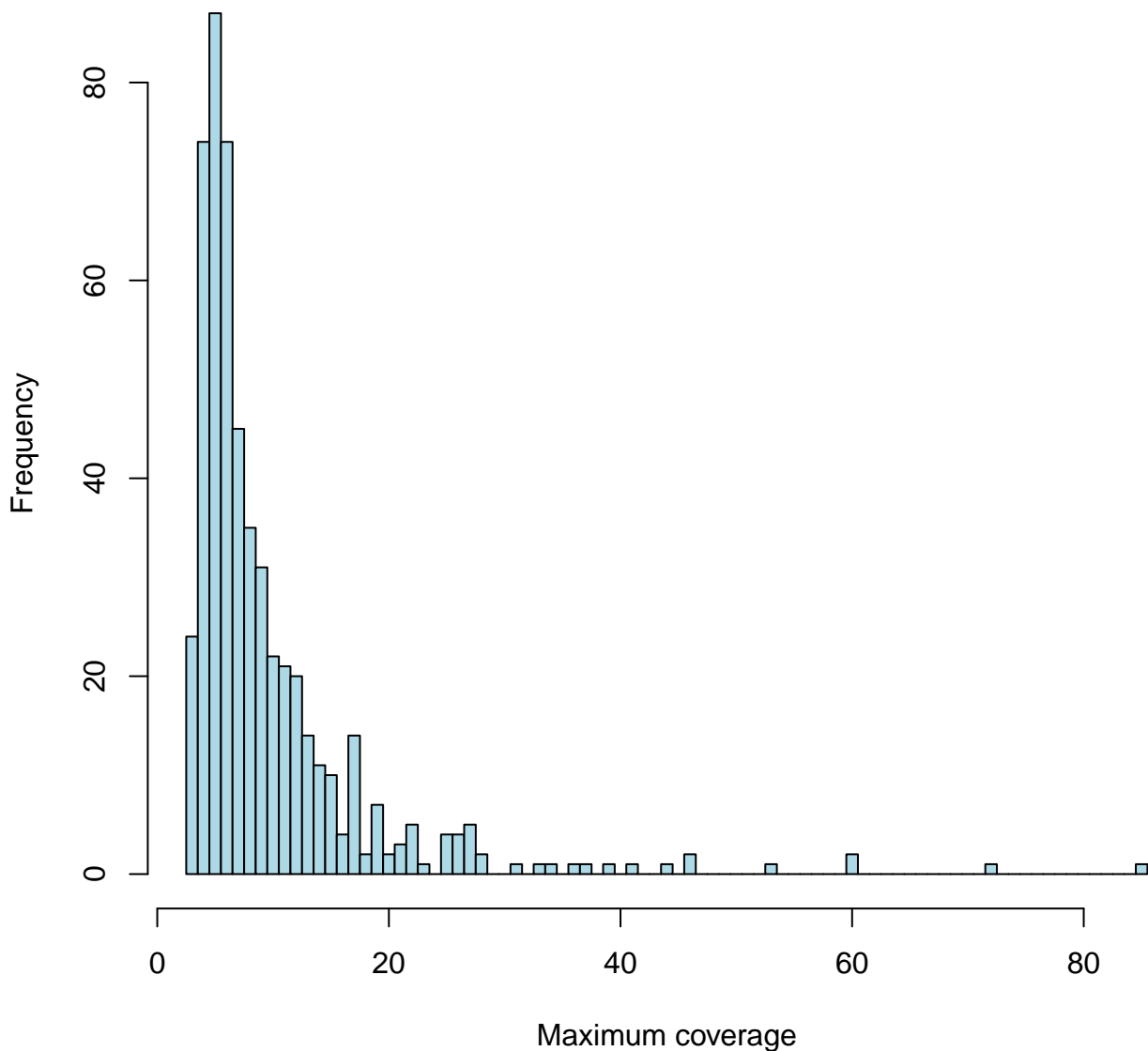
**jx in Intropolis but not "annotated" by knownGene:
65 samples; n jx: 532**



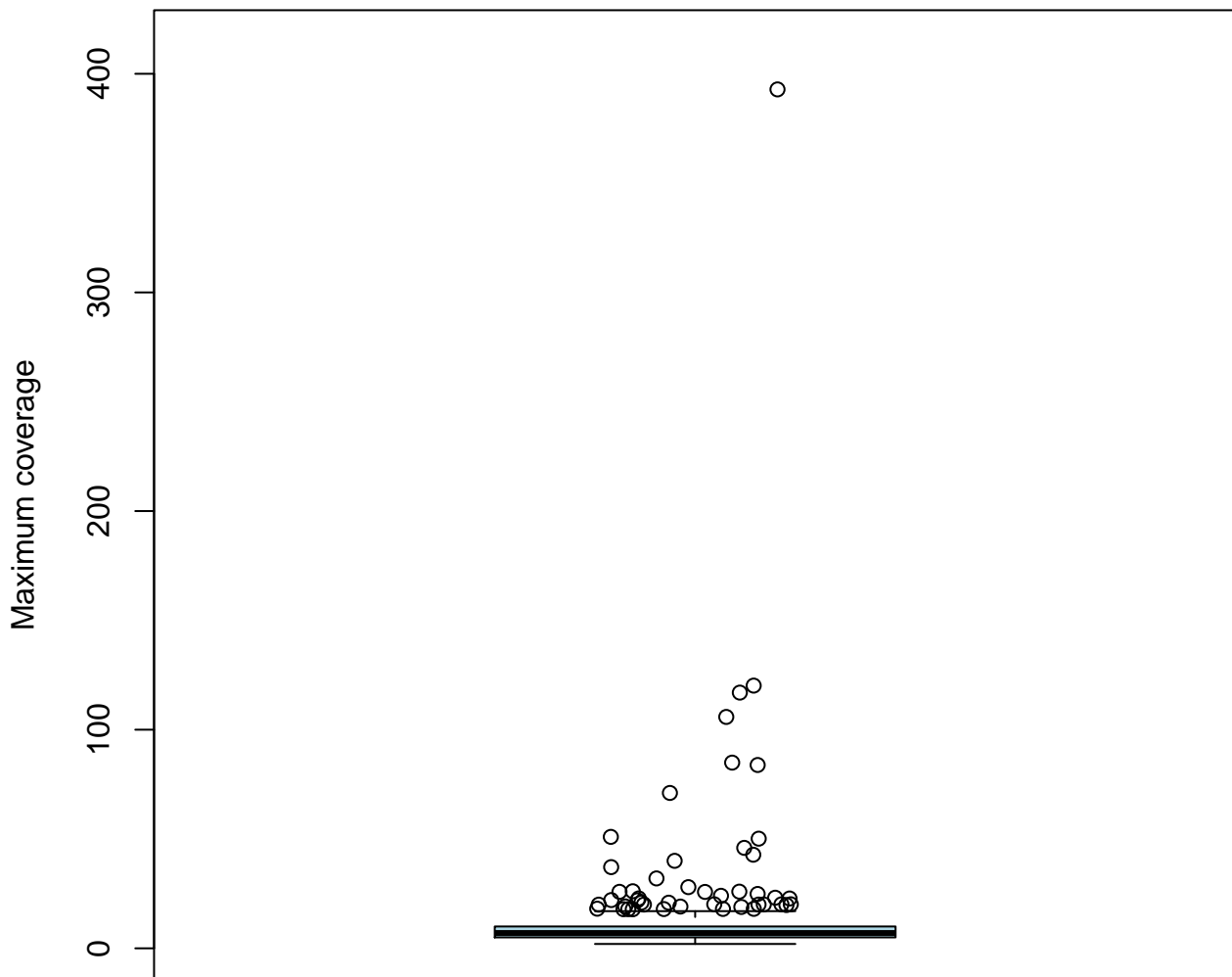
**jx in Intropolis but not "annotated" by knownGene:
66 samples; n jx: 536**



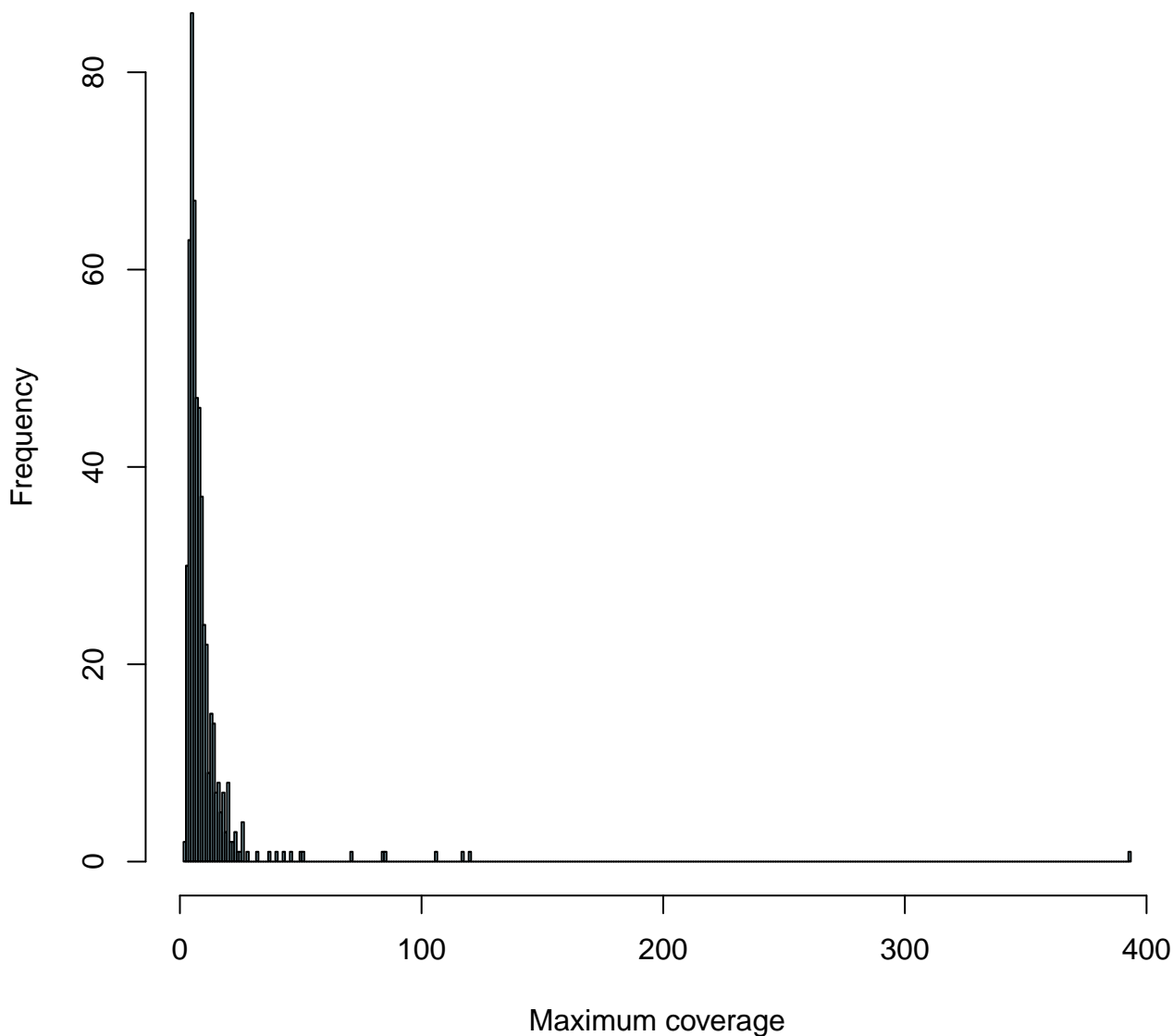
**jx in Intropolis but not "annotated" by knownGene:
66 samples; n jx: 536**



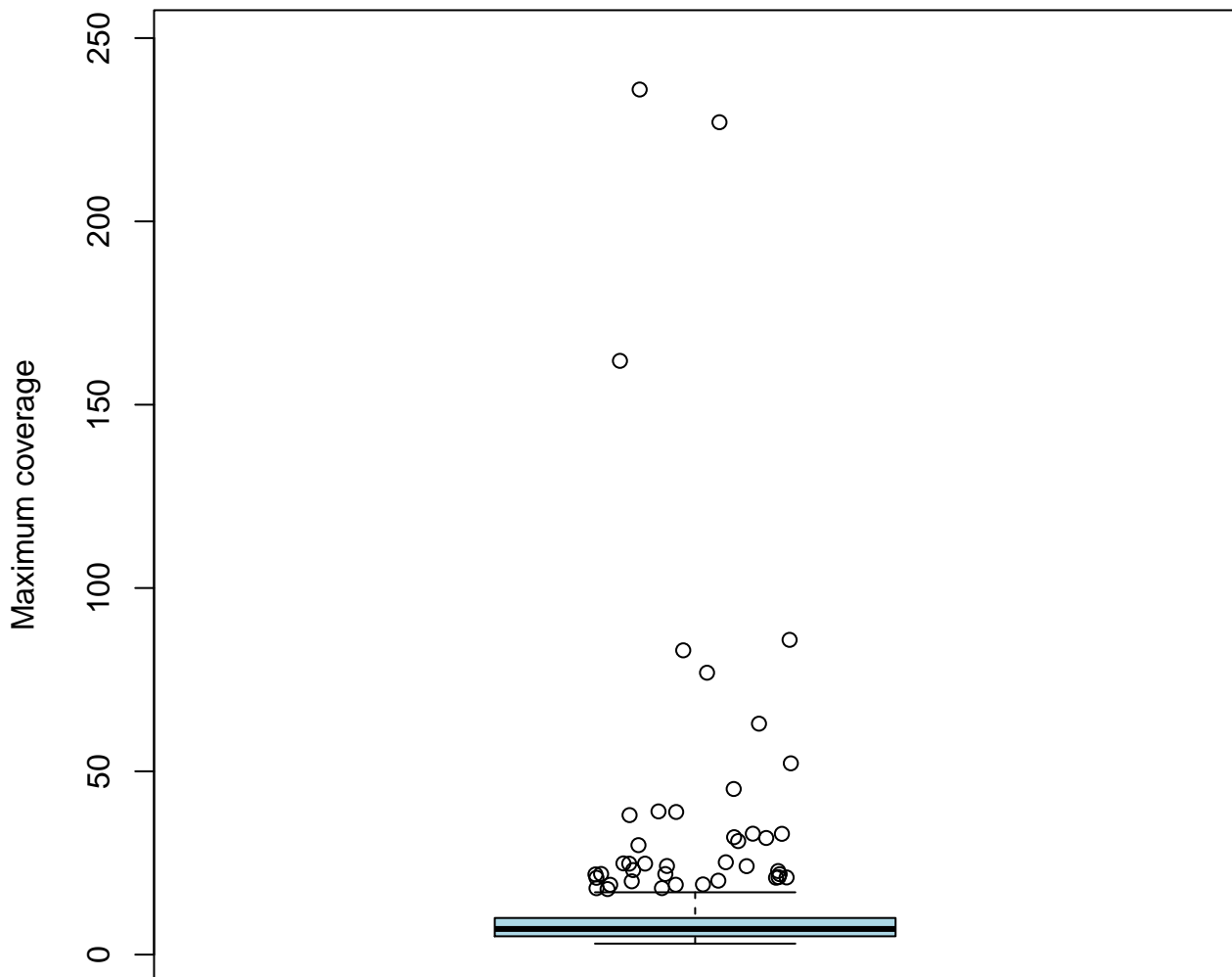
**jx in Intropolis but not "annotated" by knownGene:
67 samples; n jx: 528**



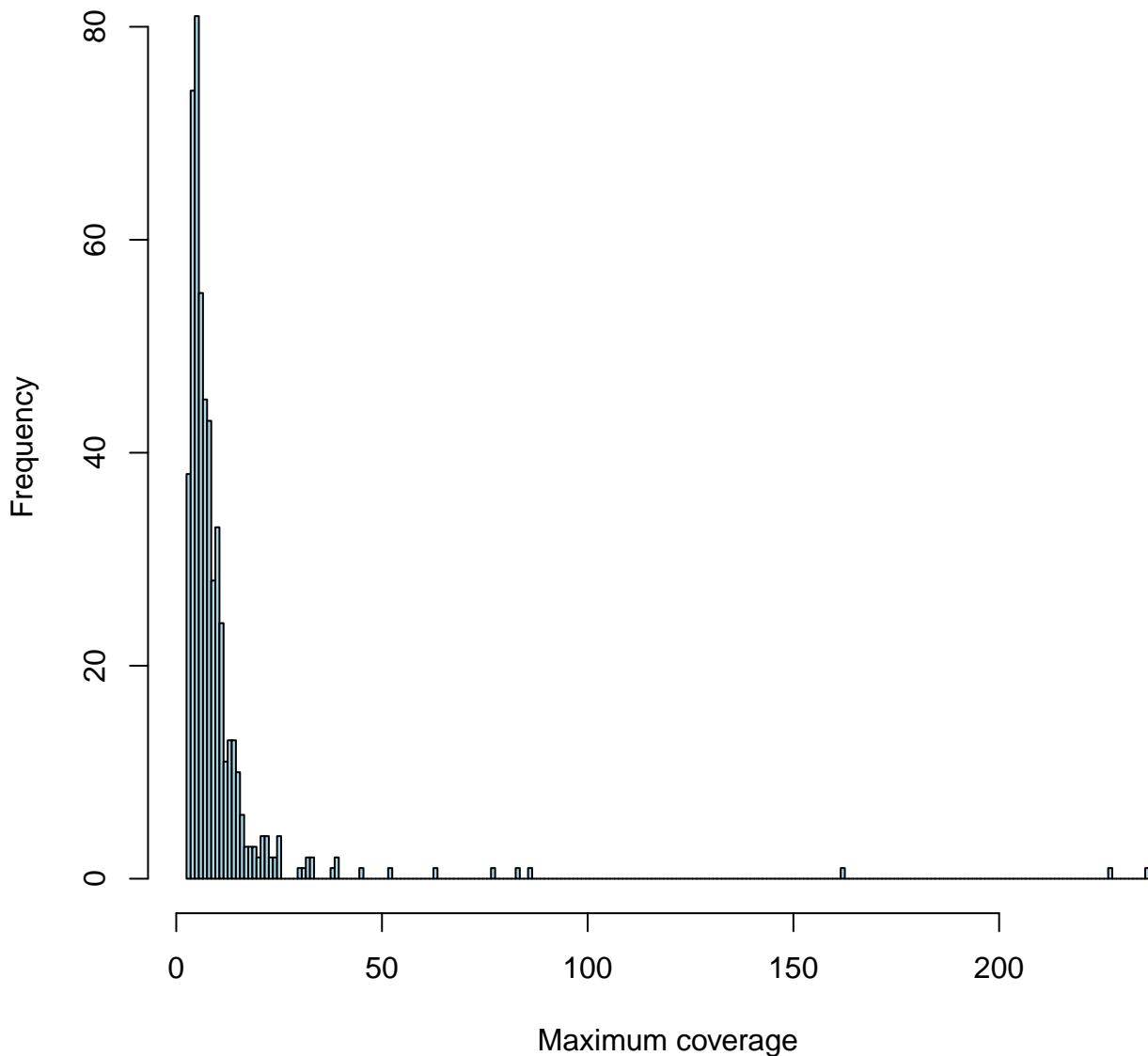
**jx in Intropolis but not "annotated" by knownGene:
67 samples; n jx: 528**



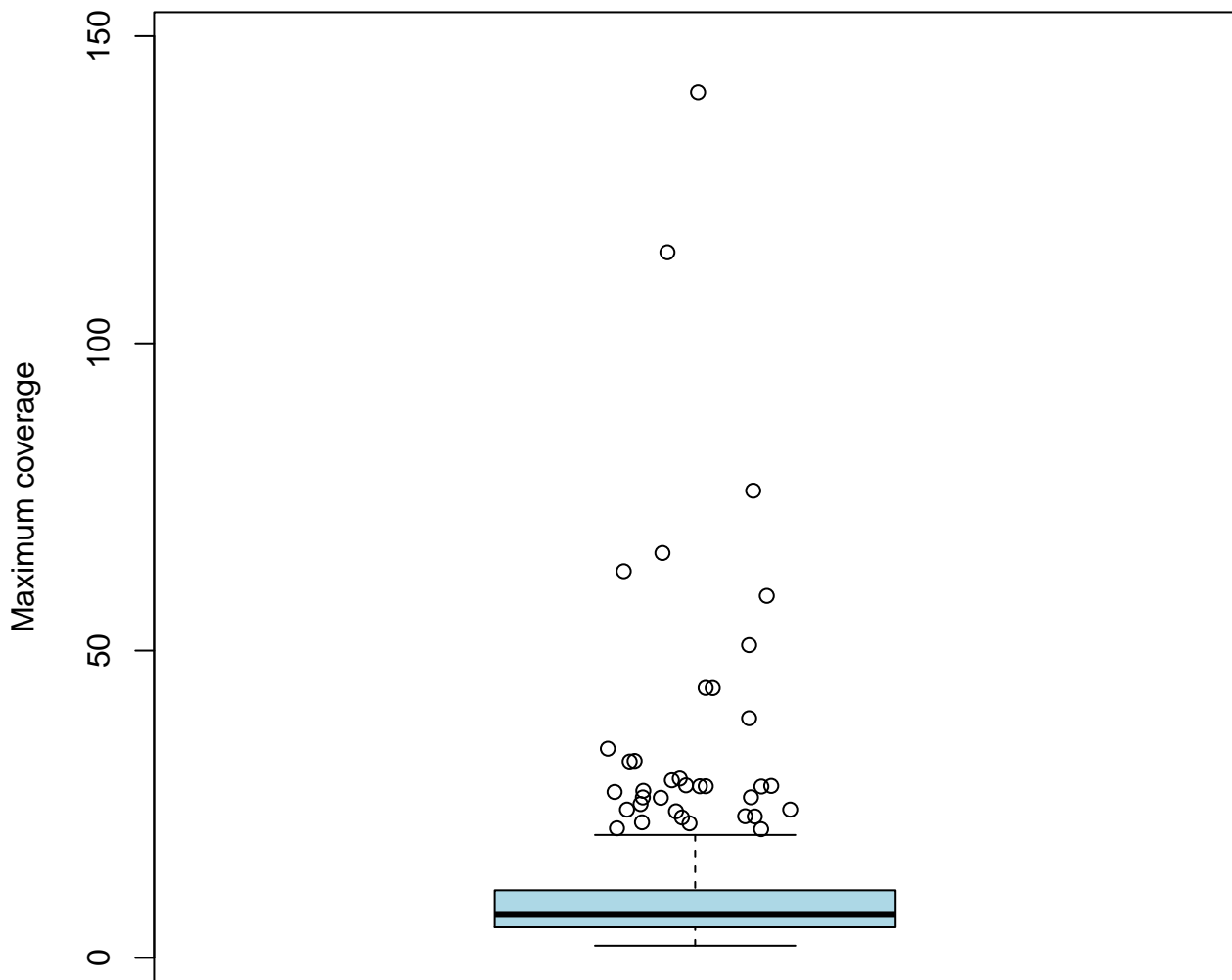
**jx in Intropolis but not "annotated" by knownGene:
68 samples; n jx: 519**



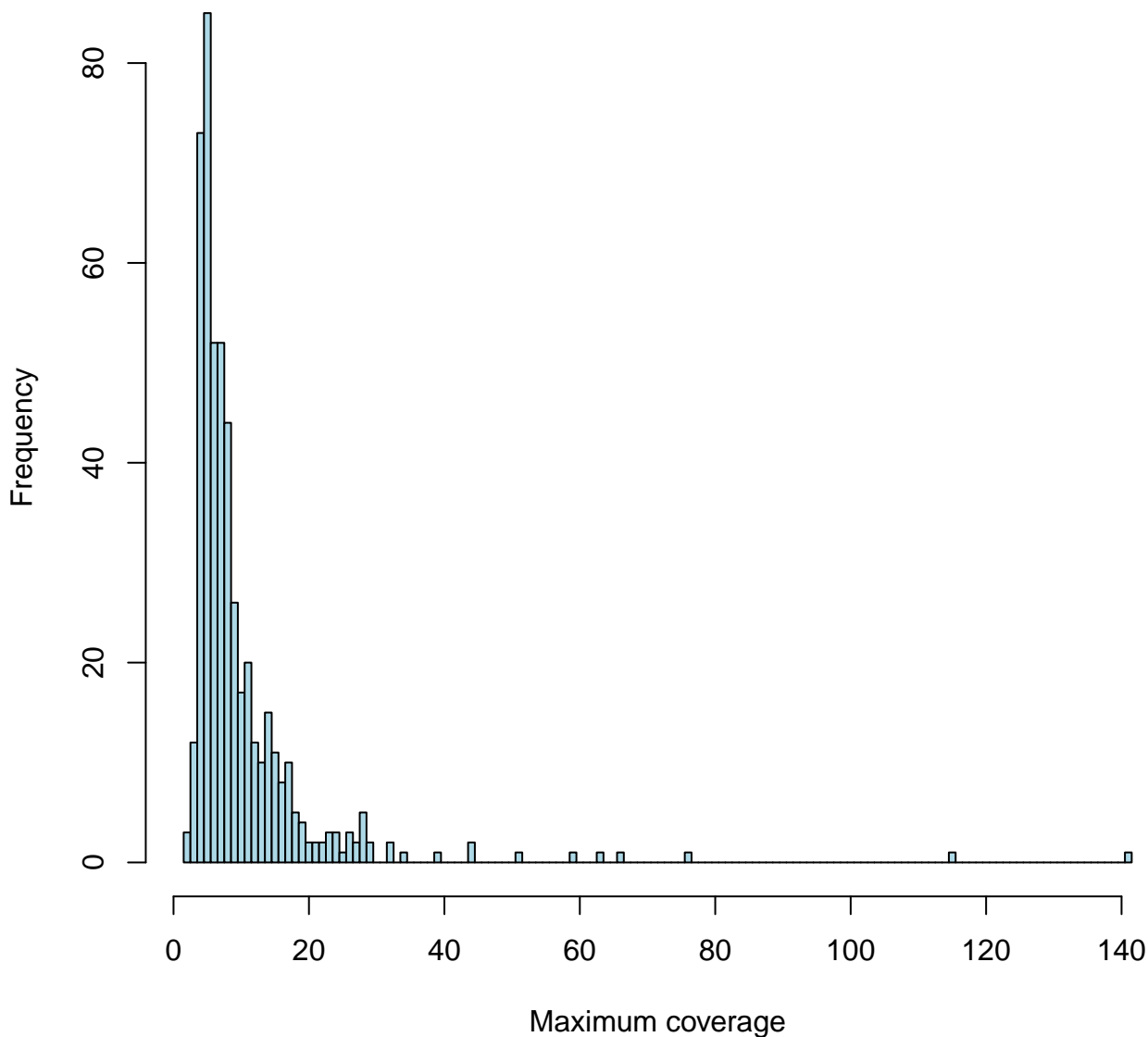
**jx in Intropolis but not "annotated" by knownGene:
68 samples; n jx: 519**



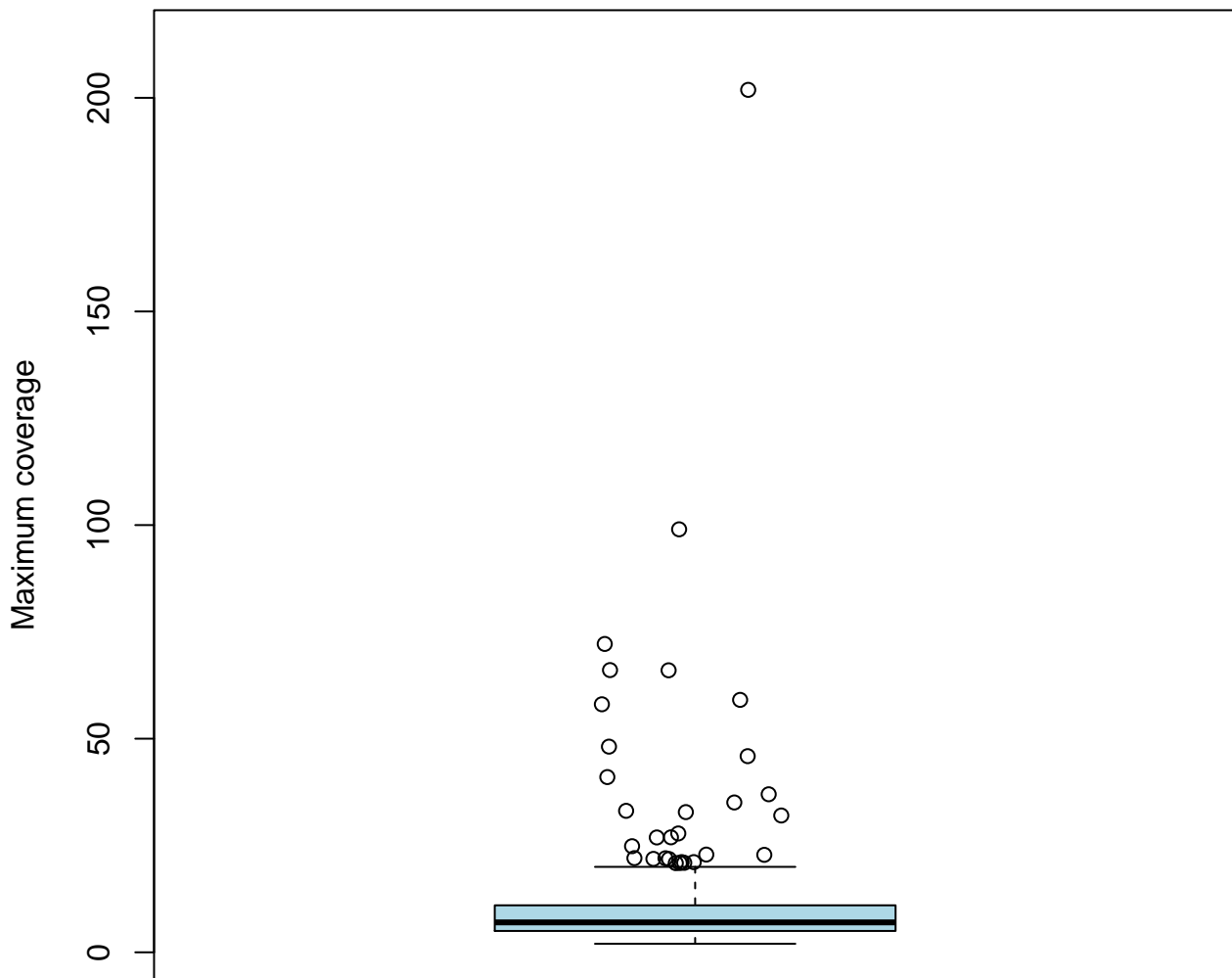
**jx in Intropolis but not "annotated" by knownGene:
69 samples; n jx: 497**



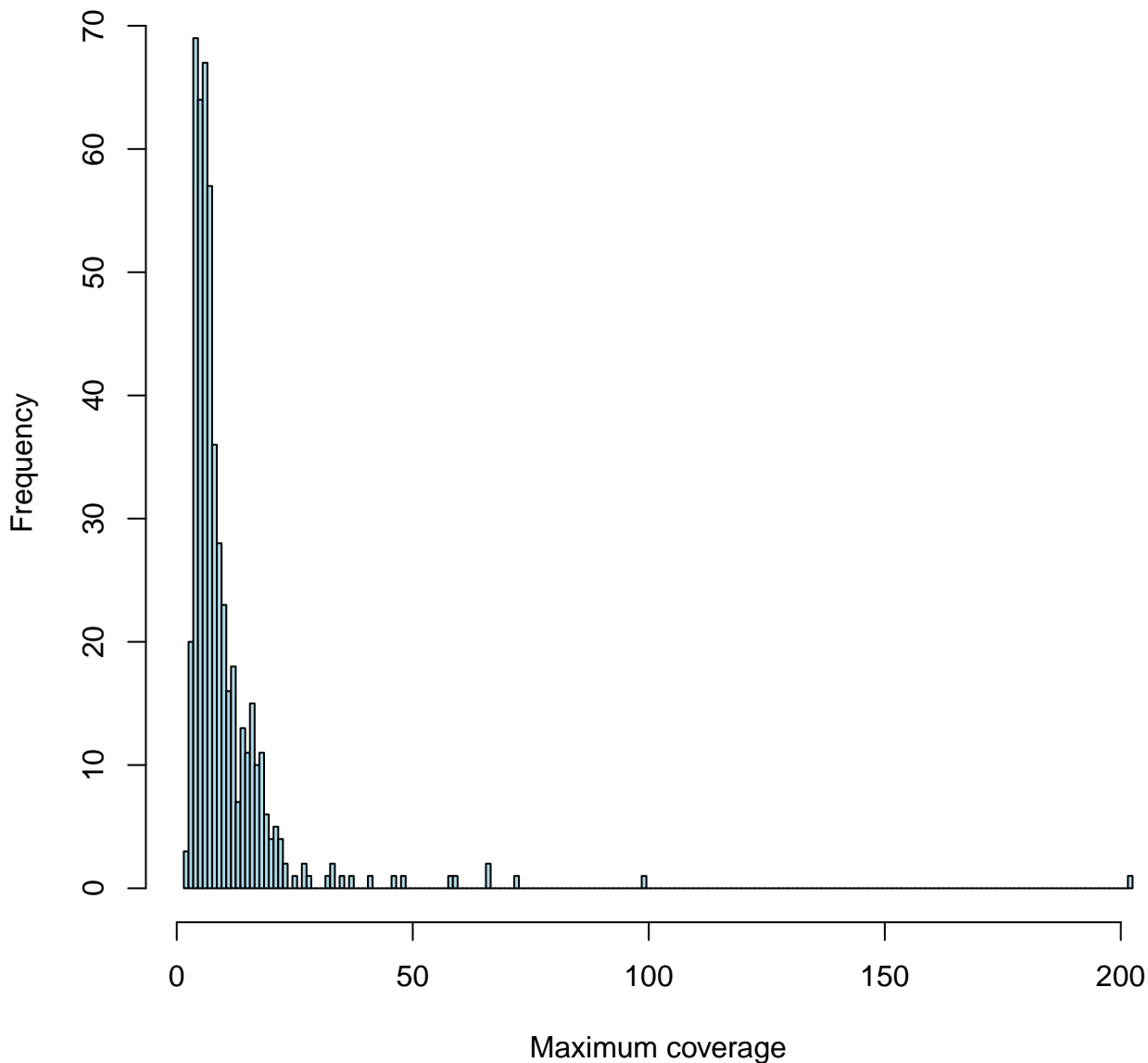
**jx in Intropolis but not "annotated" by knownGene:
69 samples; n jx: 497**



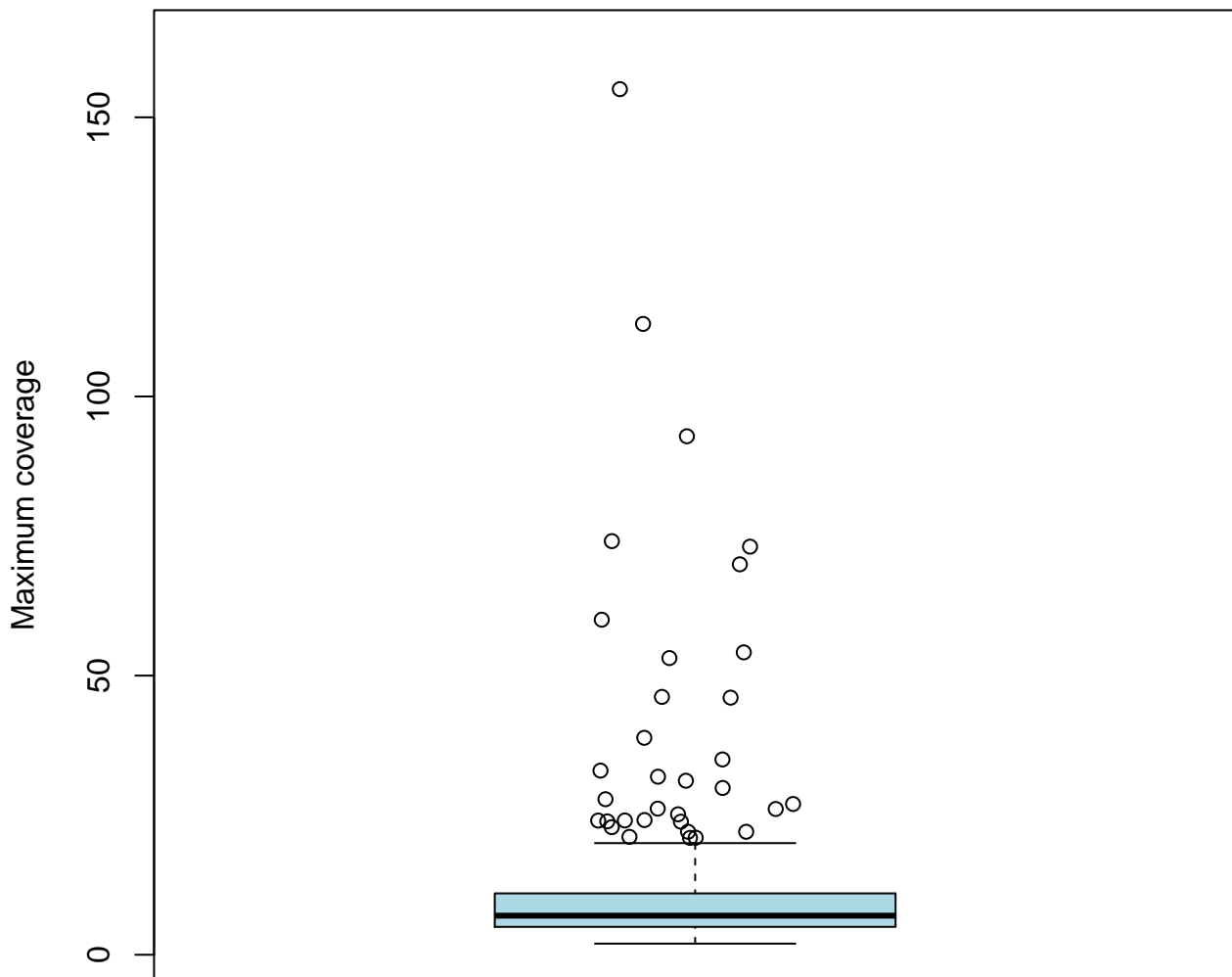
**jx in Intropolis but not "annotated" by knownGene:
70 samples; n jx: 508**



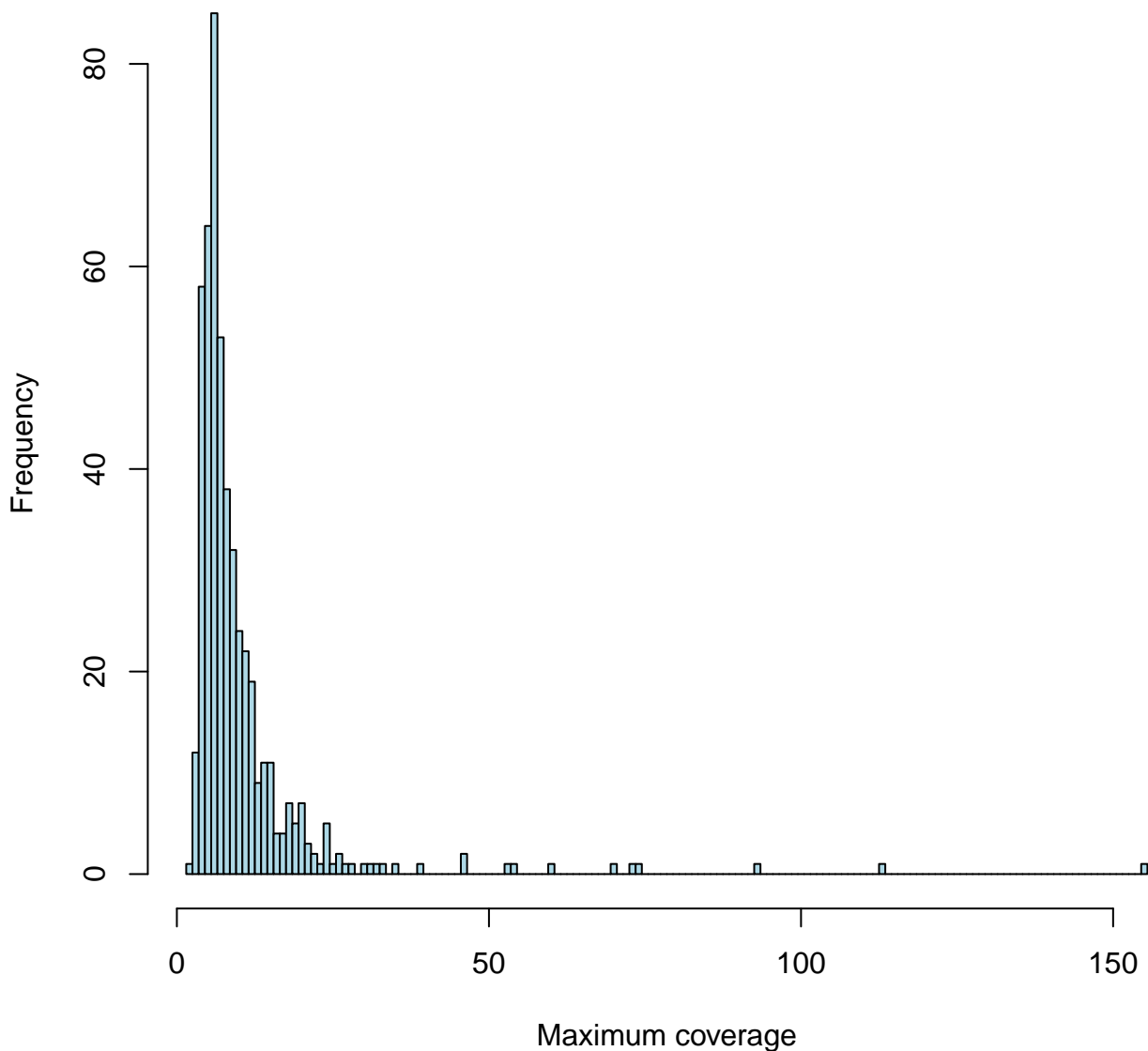
**jx in Intropolis but not "annotated" by knownGene:
70 samples; n jx: 508**



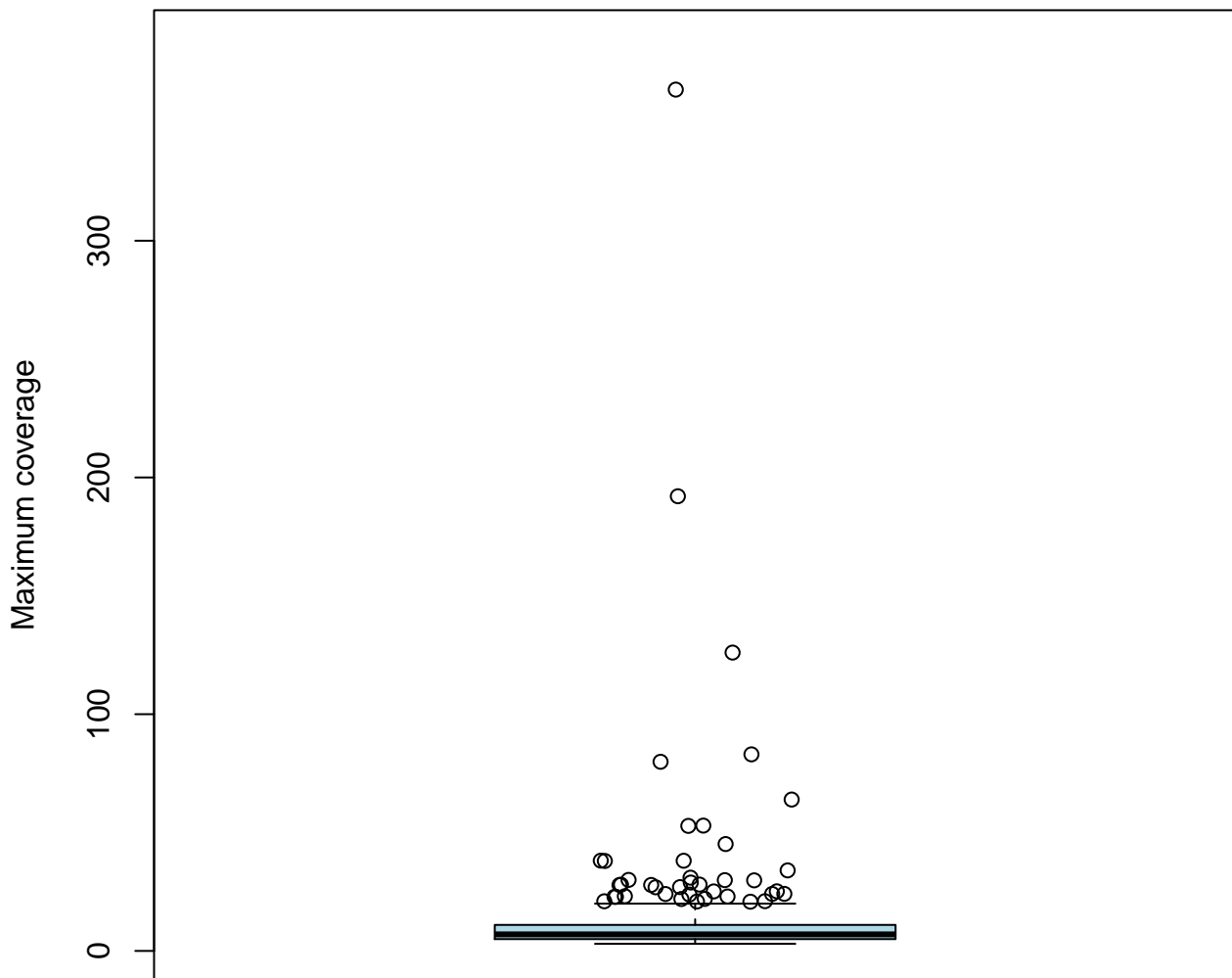
**jx in Intropolis but not "annotated" by knownGene:
71 samples; n jx: 499**



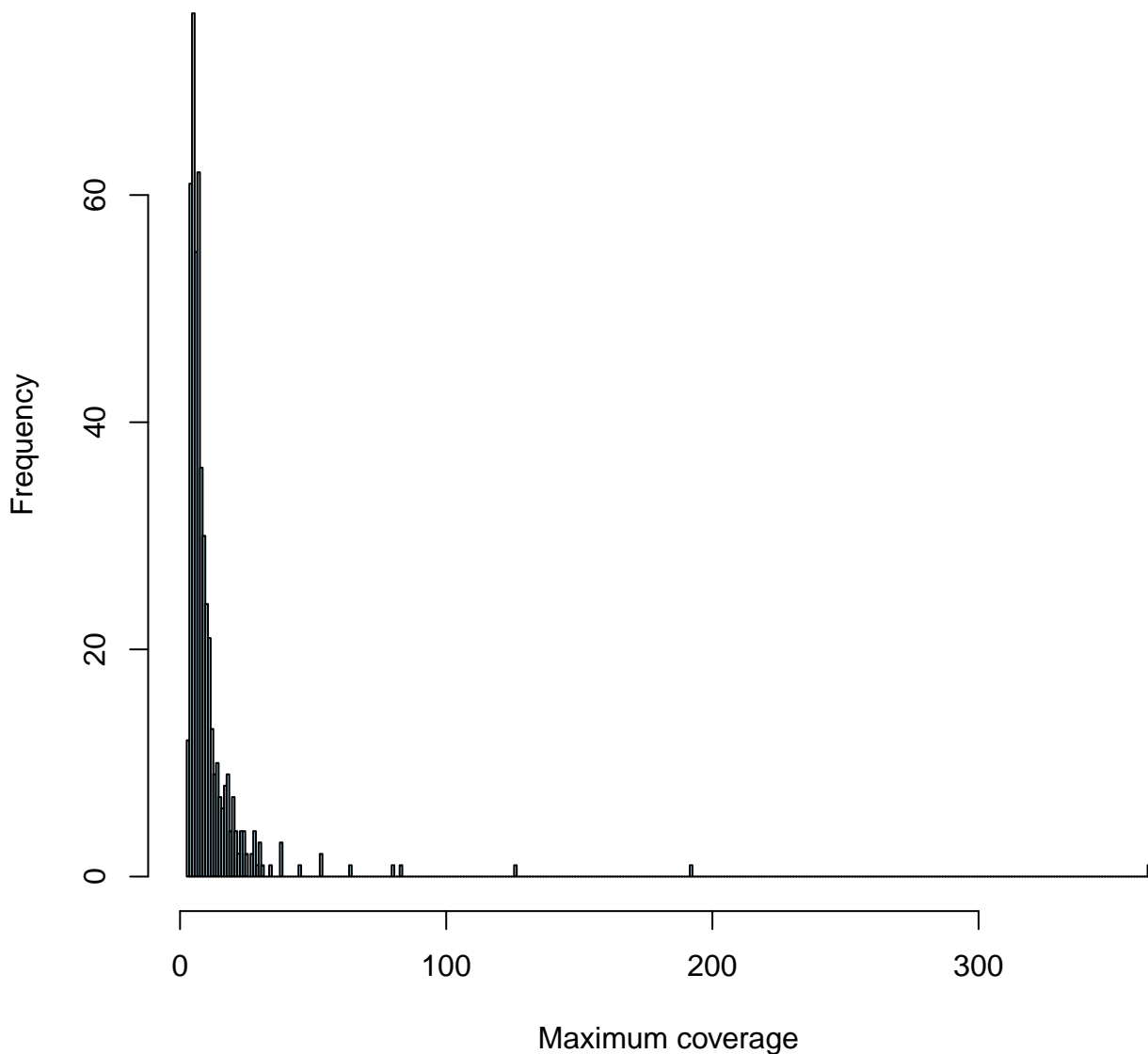
**jx in Intropolis but not "annotated" by knownGene:
71 samples; n jx: 499**



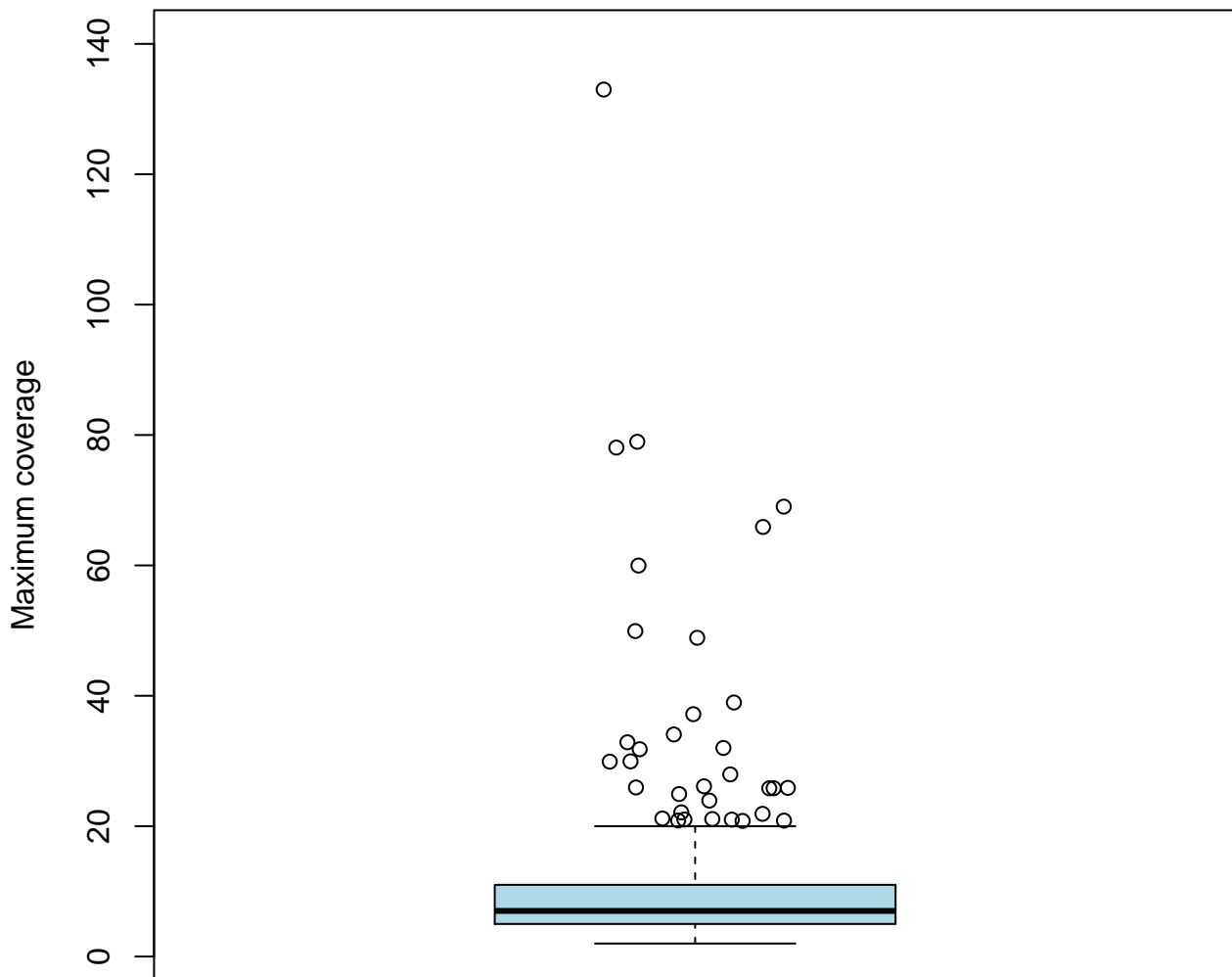
**jx in Intropolis but not "annotated" by knownGene:
72 samples; n jx: 490**



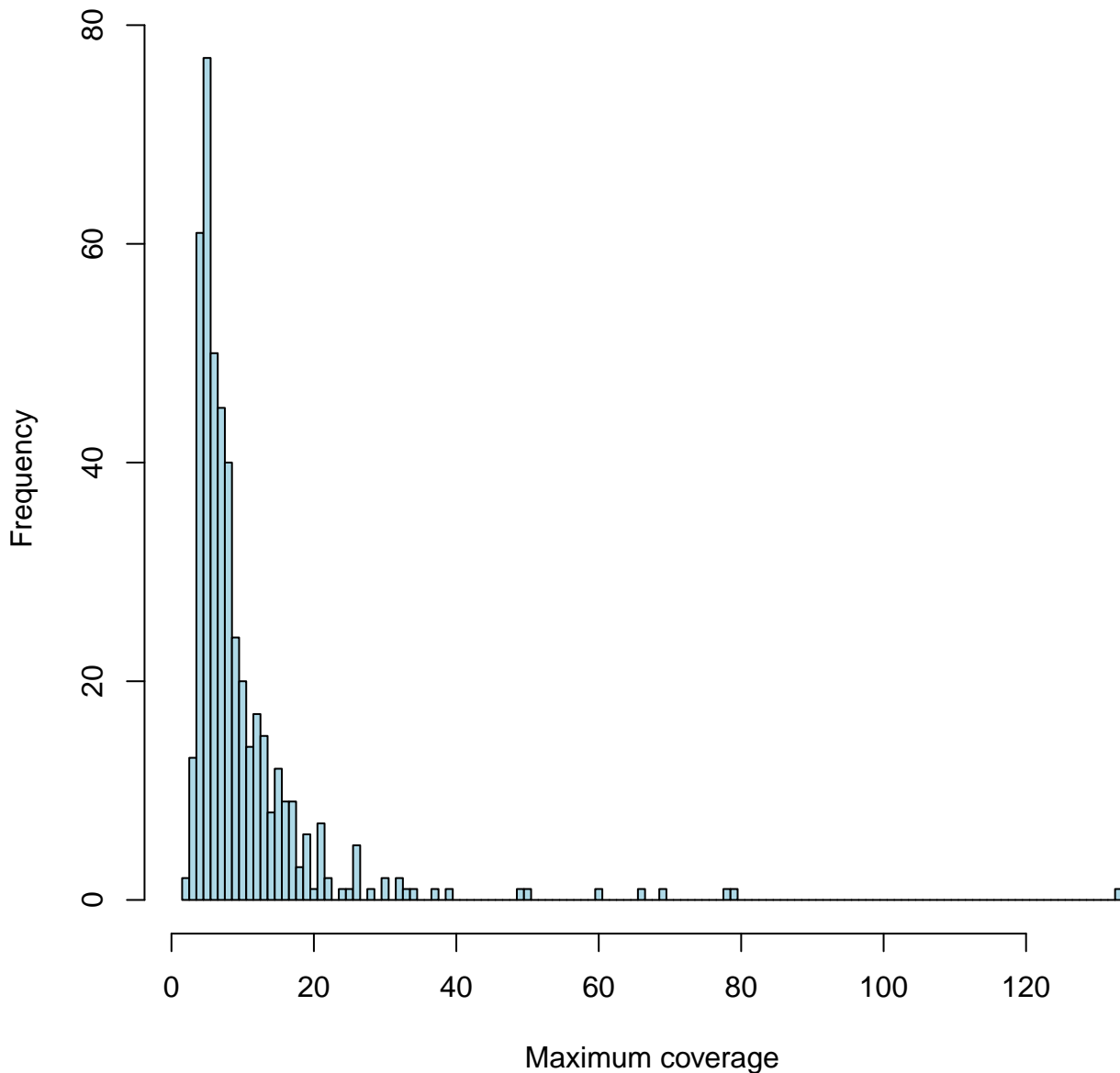
**jx in Intropolis but not "annotated" by knownGene:
72 samples; n jx: 490**



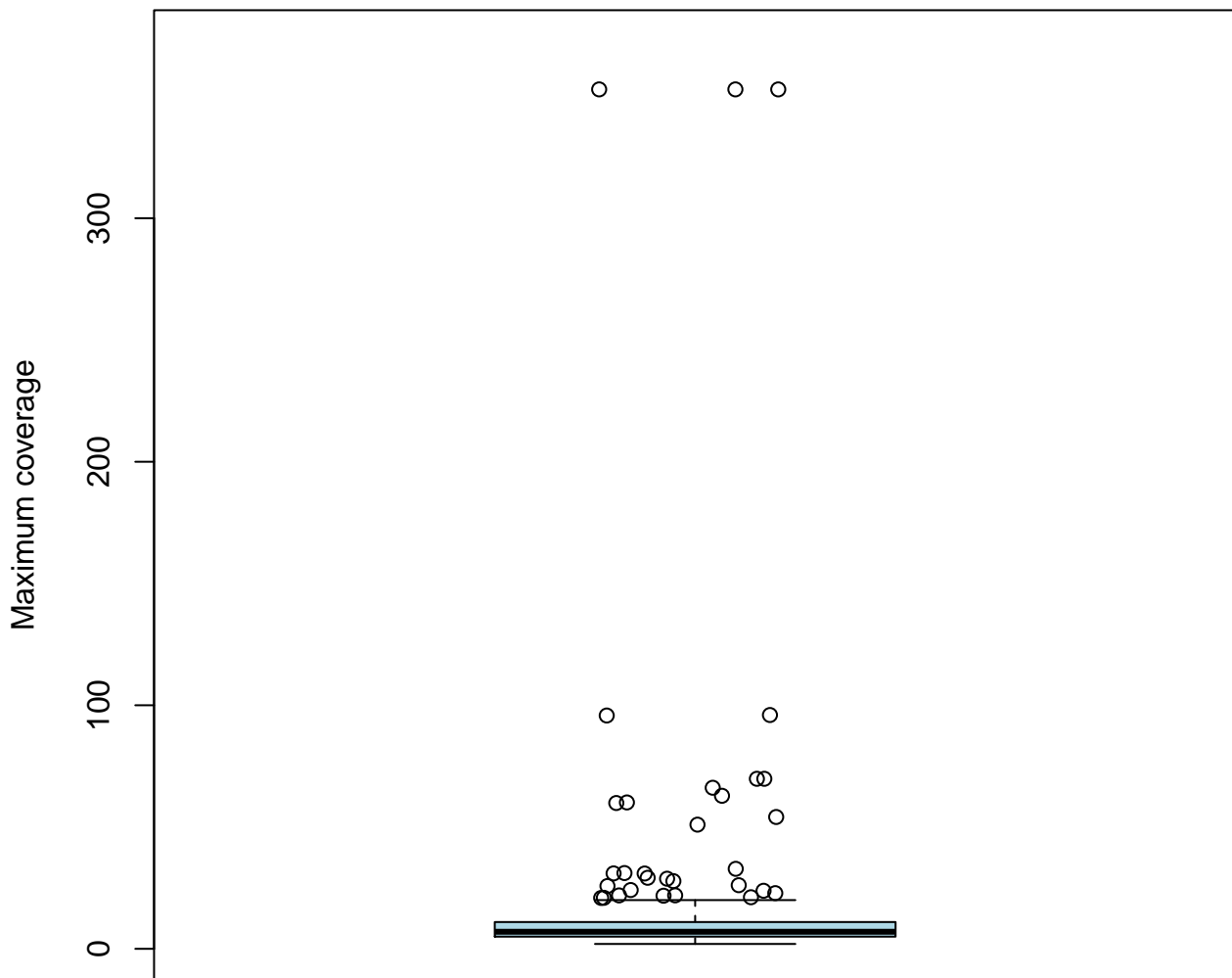
**jx in Intropolis but not "annotated" by knownGene:
73 samples; n jx: 459**



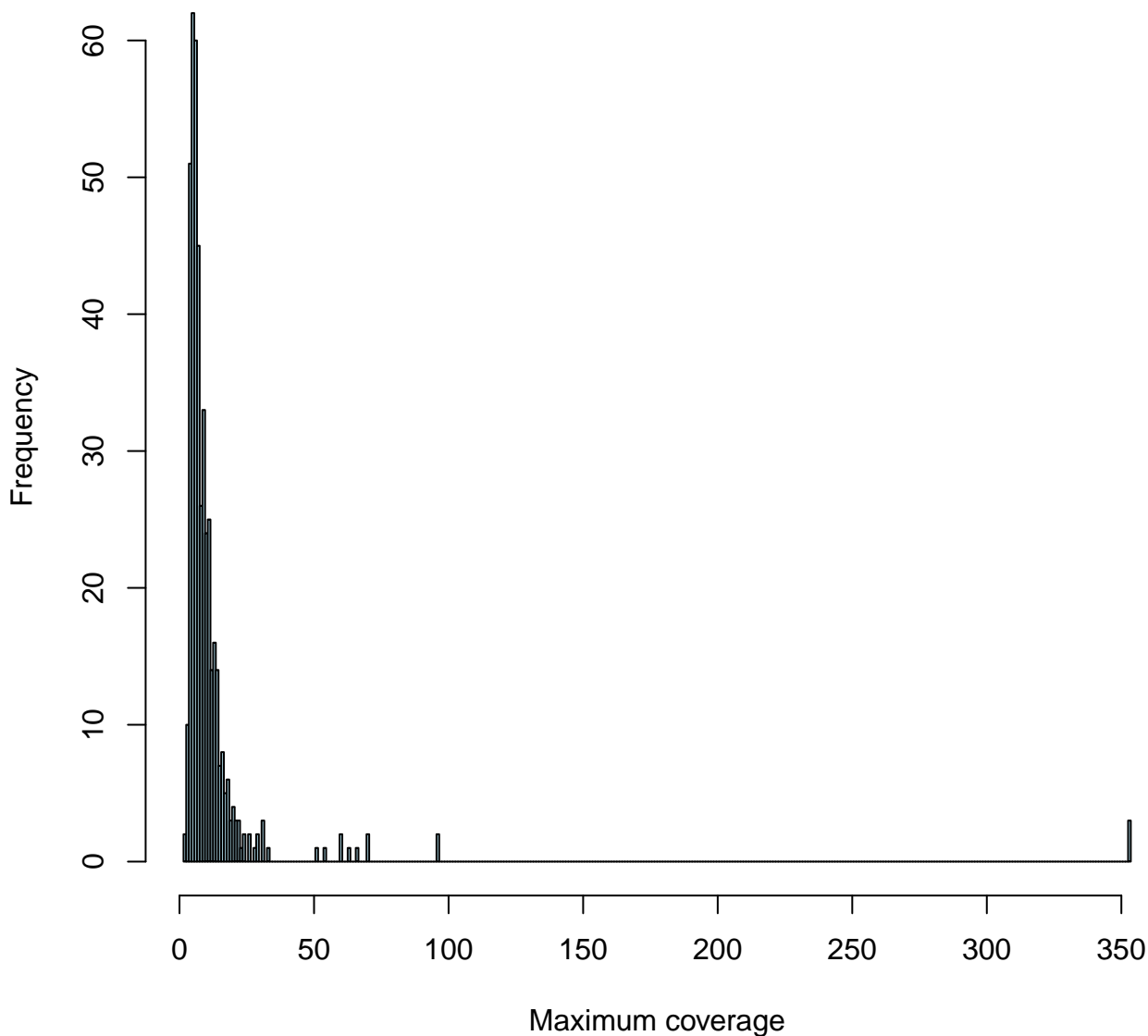
**jx in Intropolis but not "annotated" by knownGene:
73 samples; n jx: 459**



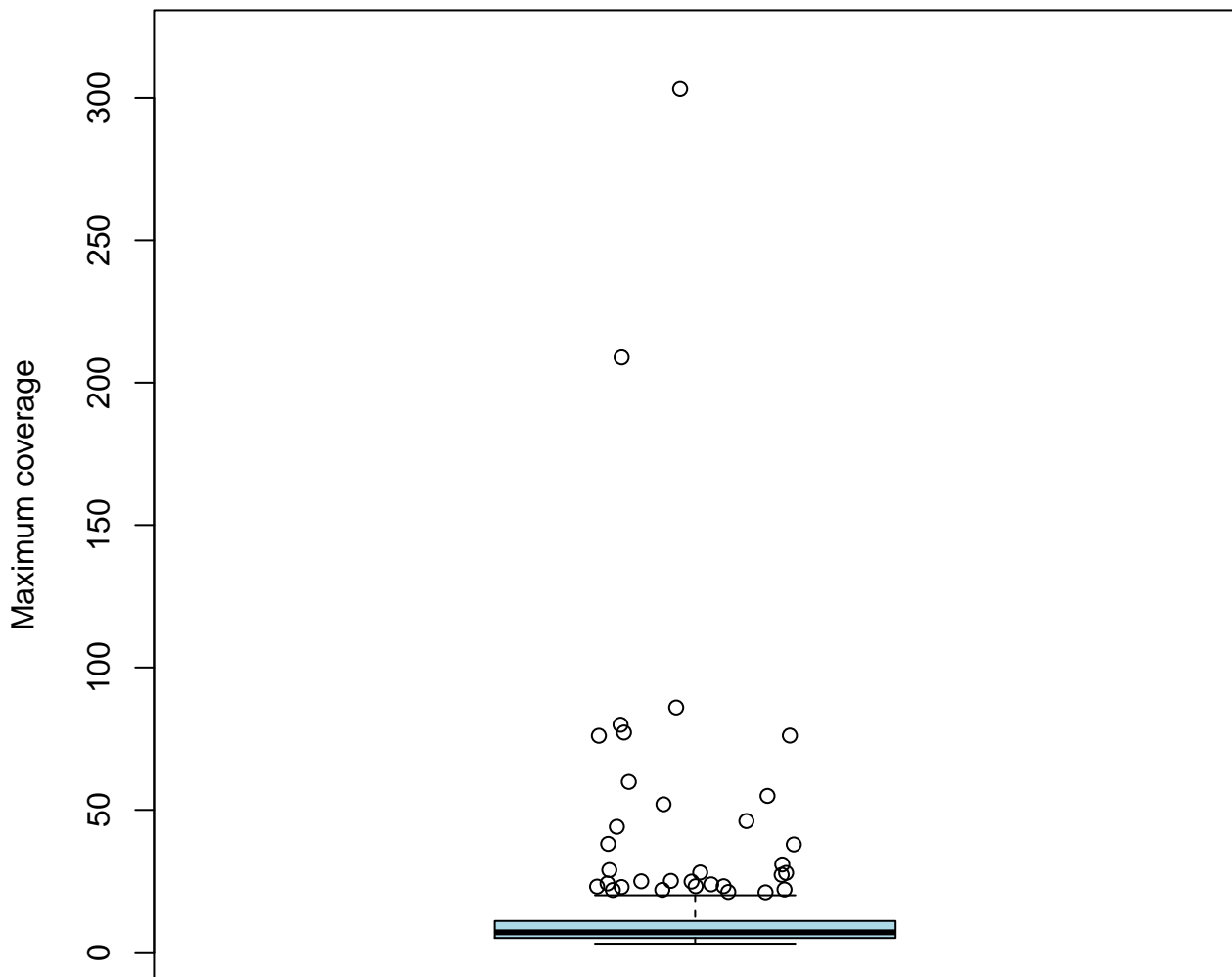
**jx in Intropolis but not "annotated" by knownGene:
74 samples; n jx: 446**



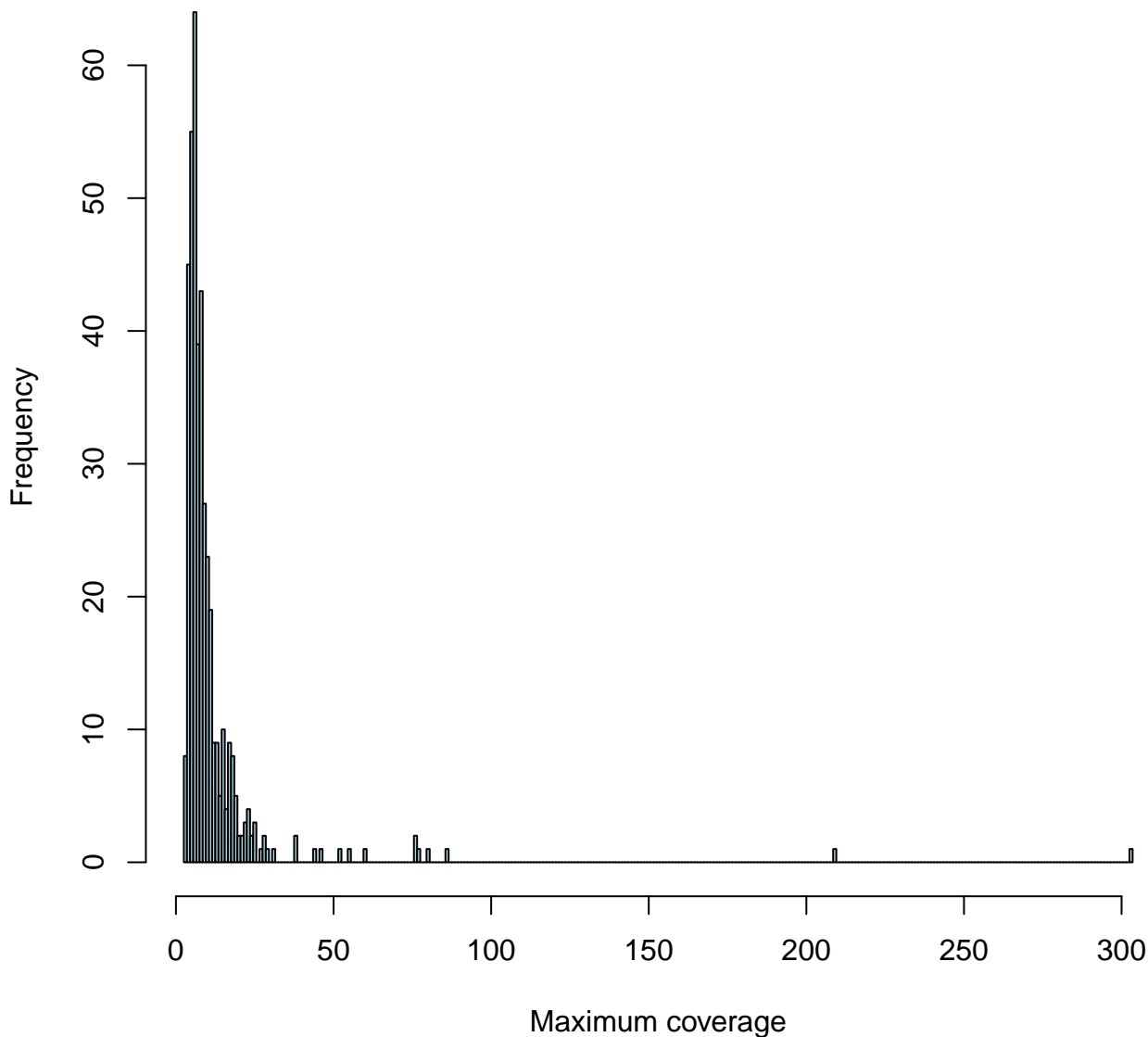
**jx in Intropolis but not "annotated" by knownGene:
74 samples; n jx: 446**



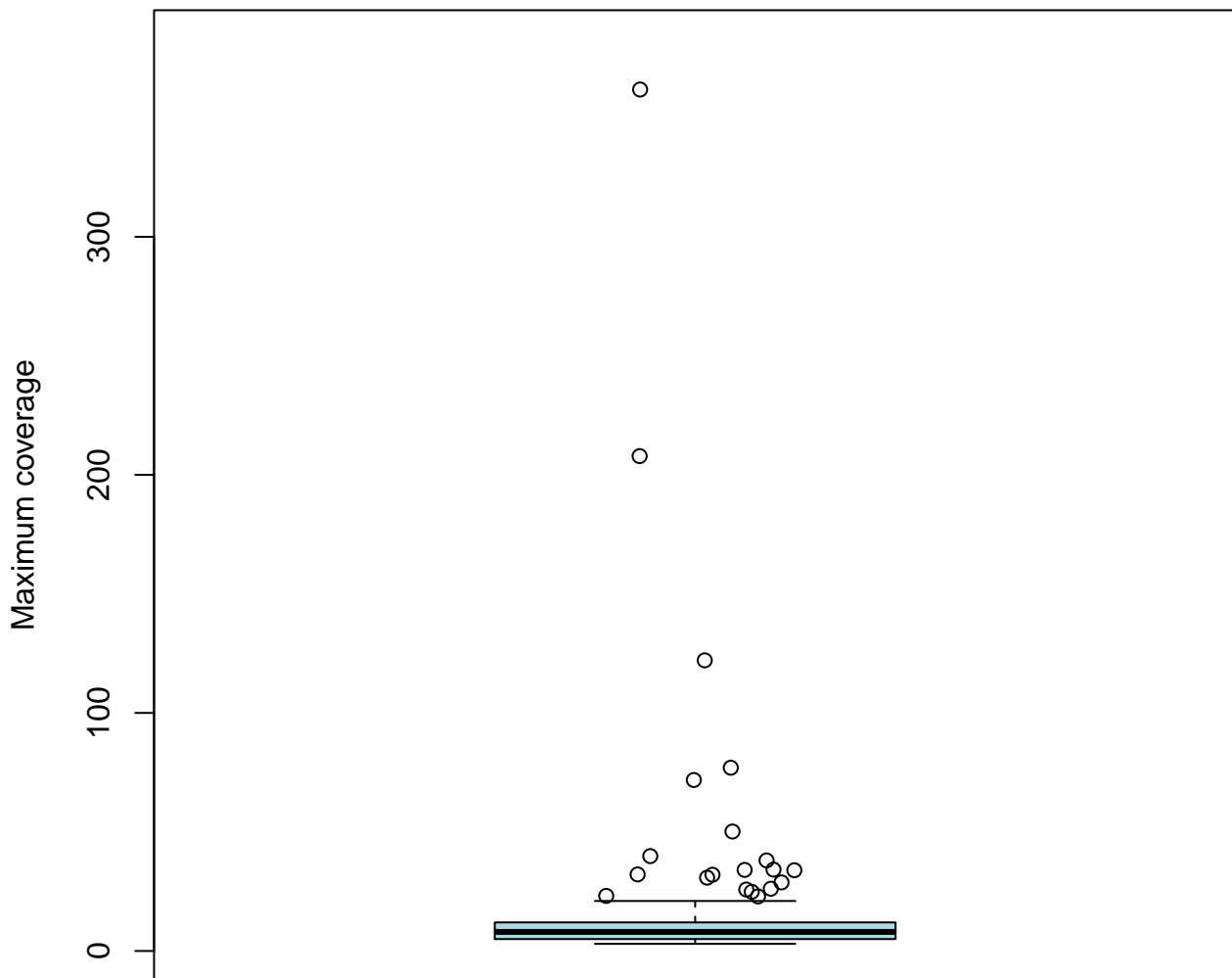
**jx in Intropolis but not "annotated" by knownGene:
75 samples; n jx: 417**



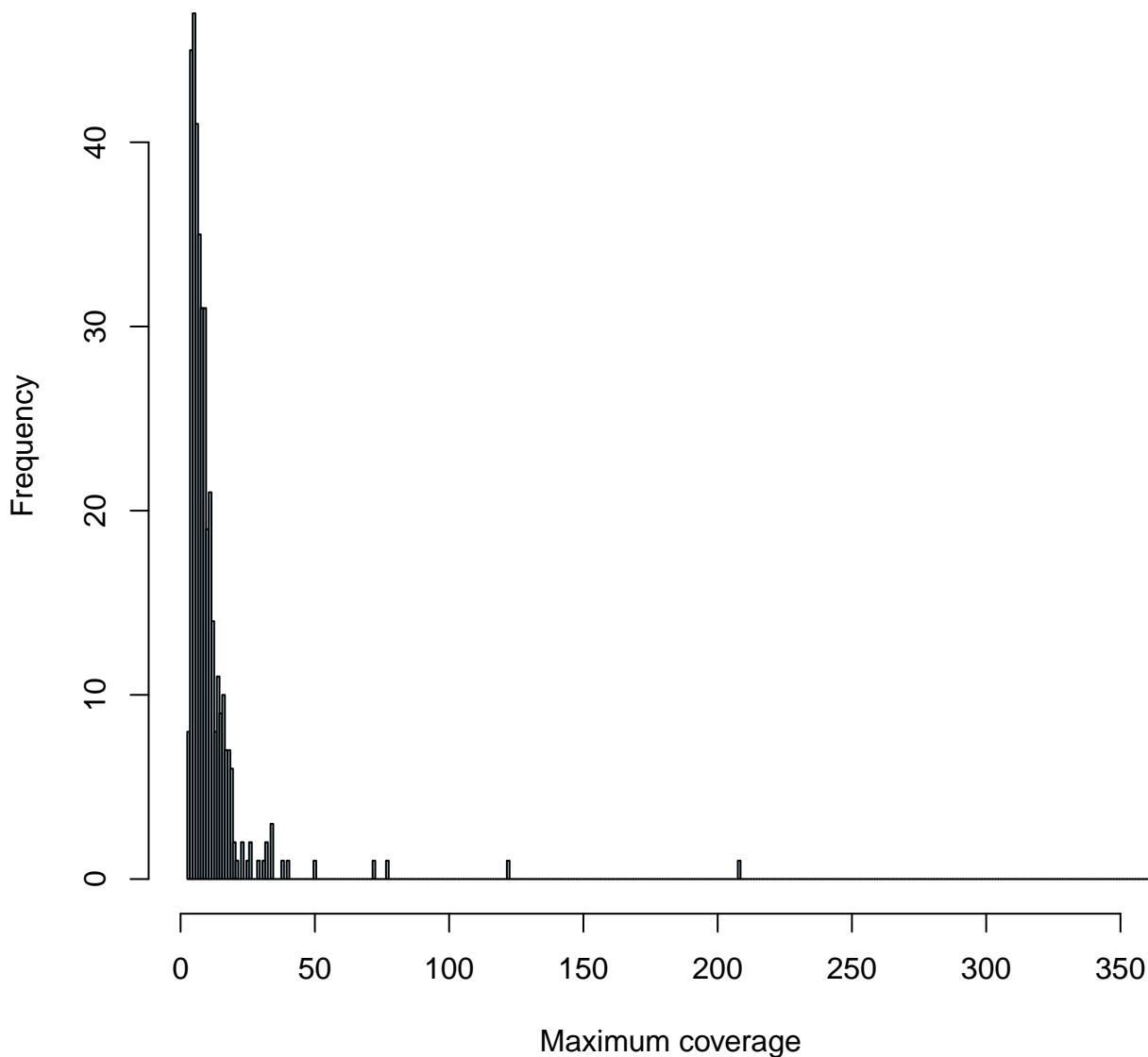
**jx in Intropolis but not "annotated" by knownGene:
75 samples; n jx: 417**



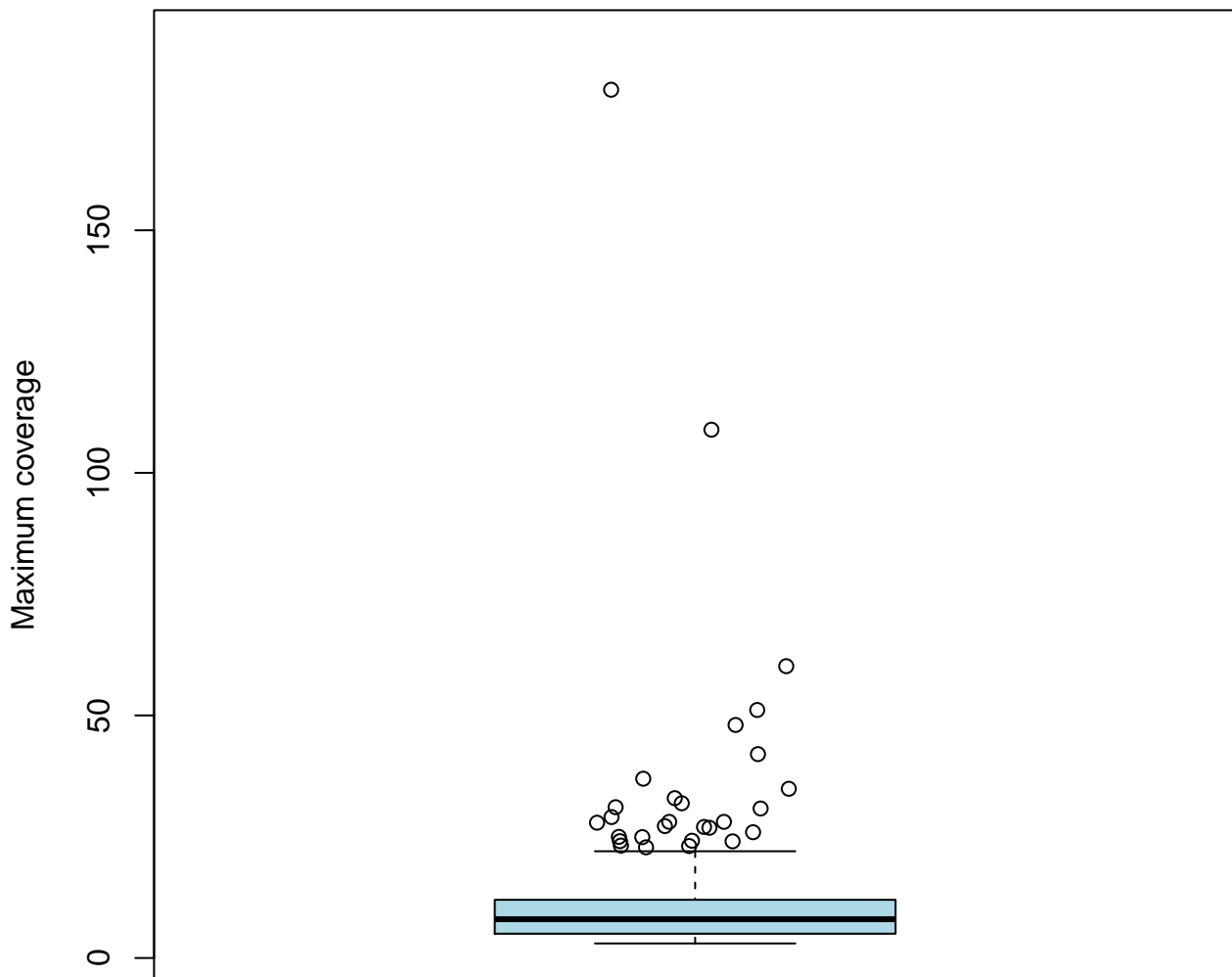
**jx in Intropolis but not "annotated" by knownGene:
76 samples; n jx: 373**



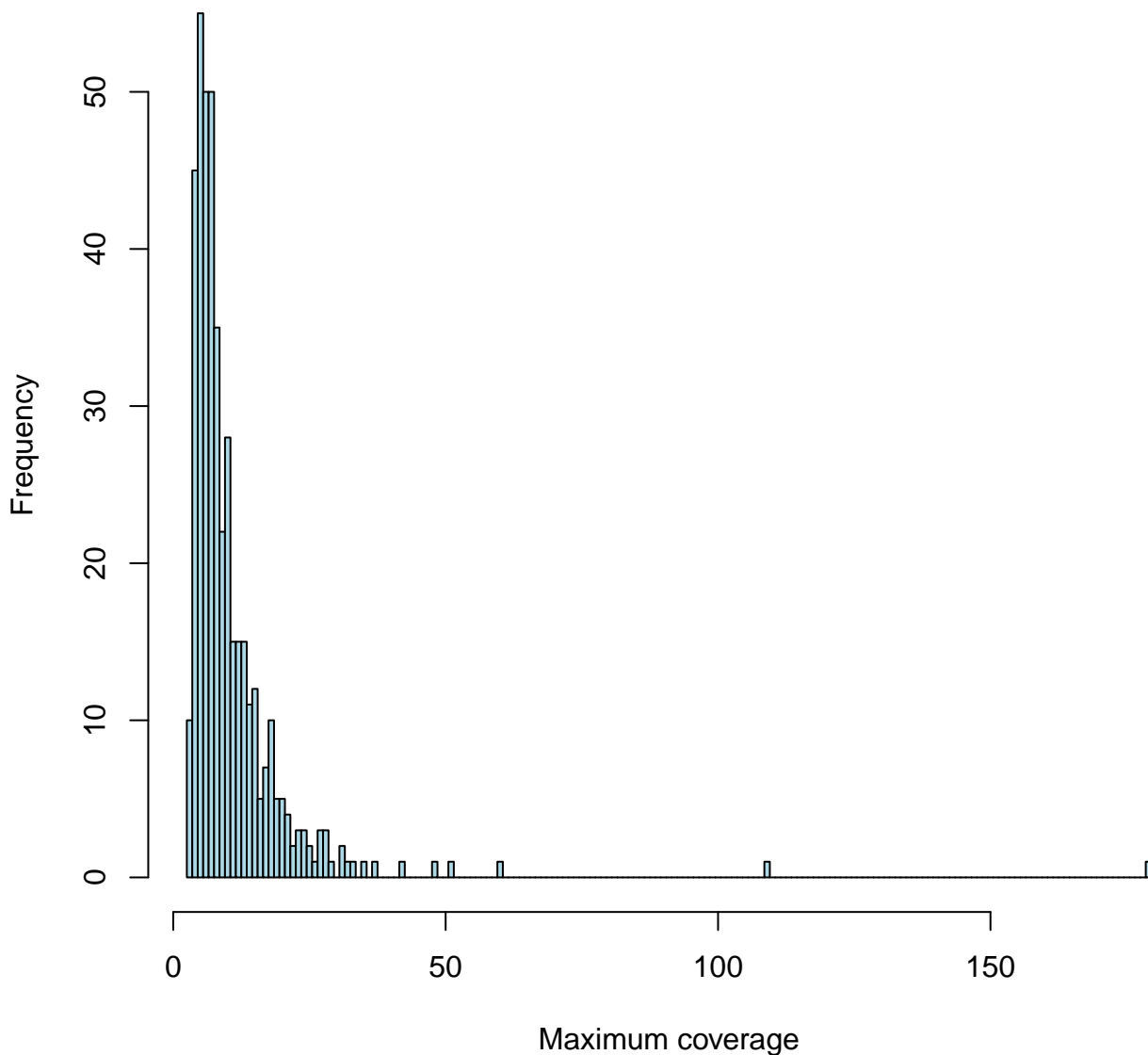
**jx in Intropolis but not "annotated" by knownGene:
76 samples; n jx: 373**



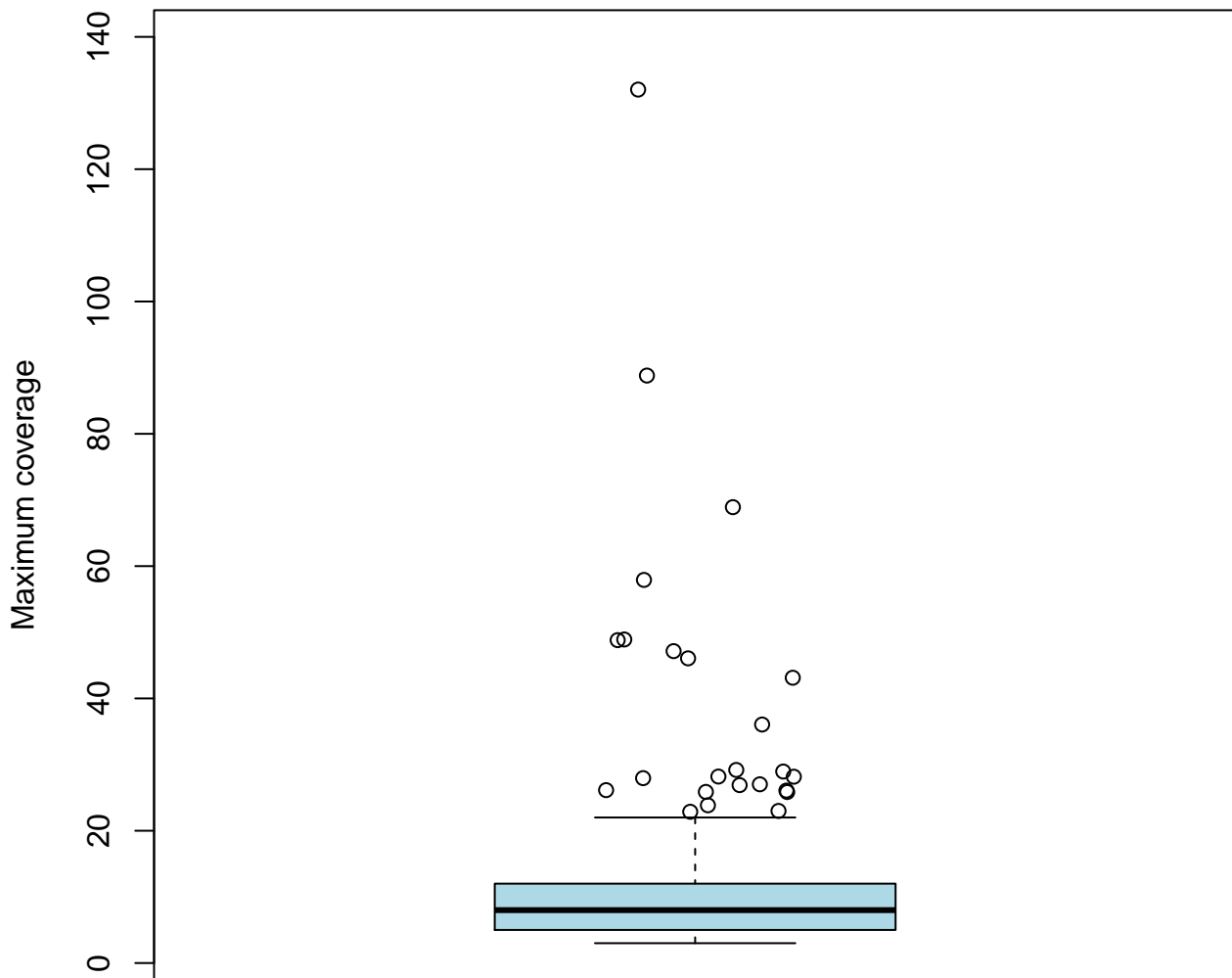
**jx in Intropolis but not "annotated" by knownGene:
77 samples; n jx: 429**



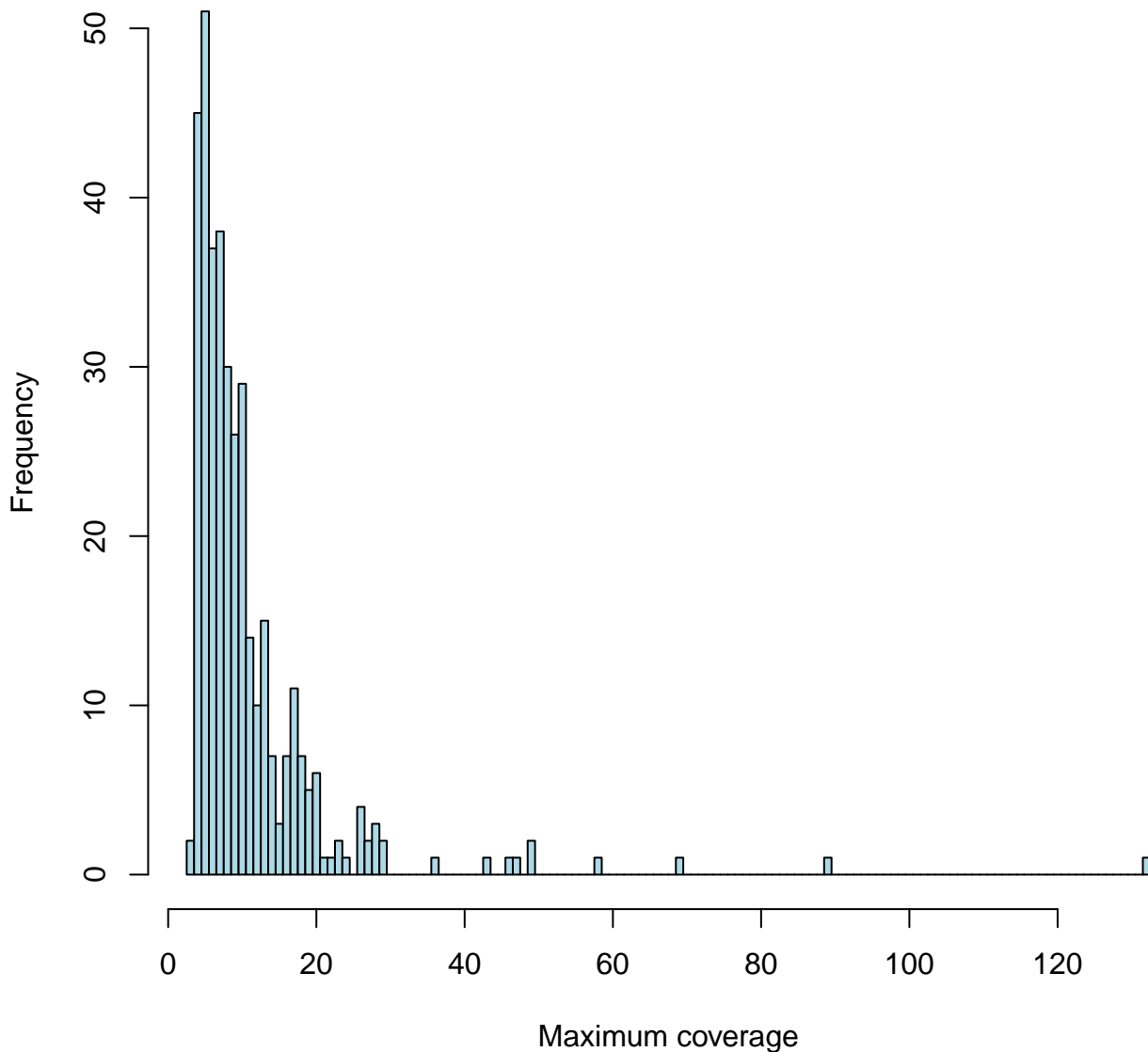
**jx in Intropolis but not "annotated" by knownGene:
77 samples; n jx: 429**



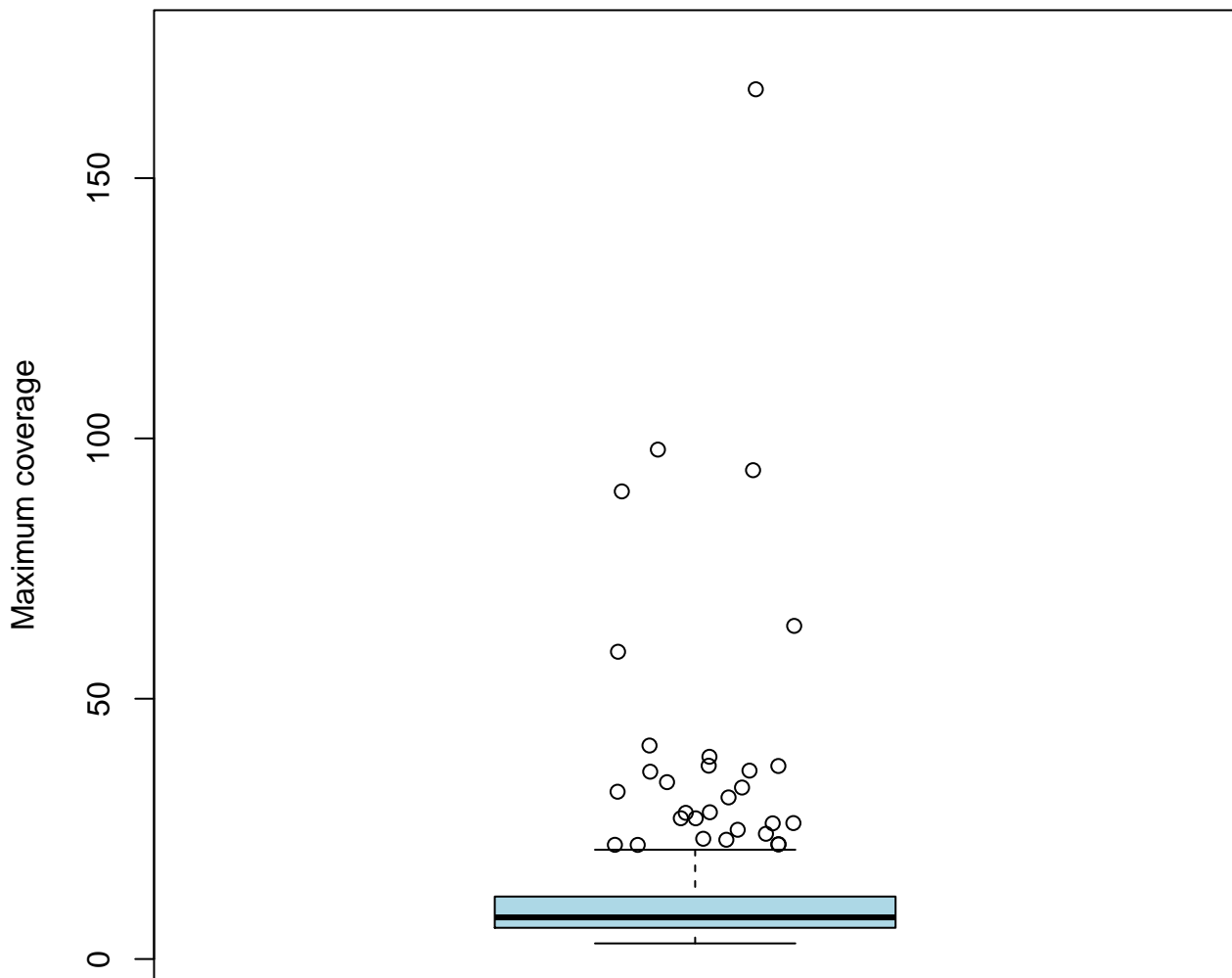
**jx in Intropolis but not "annotated" by knownGene:
78 samples; n jx: 369**



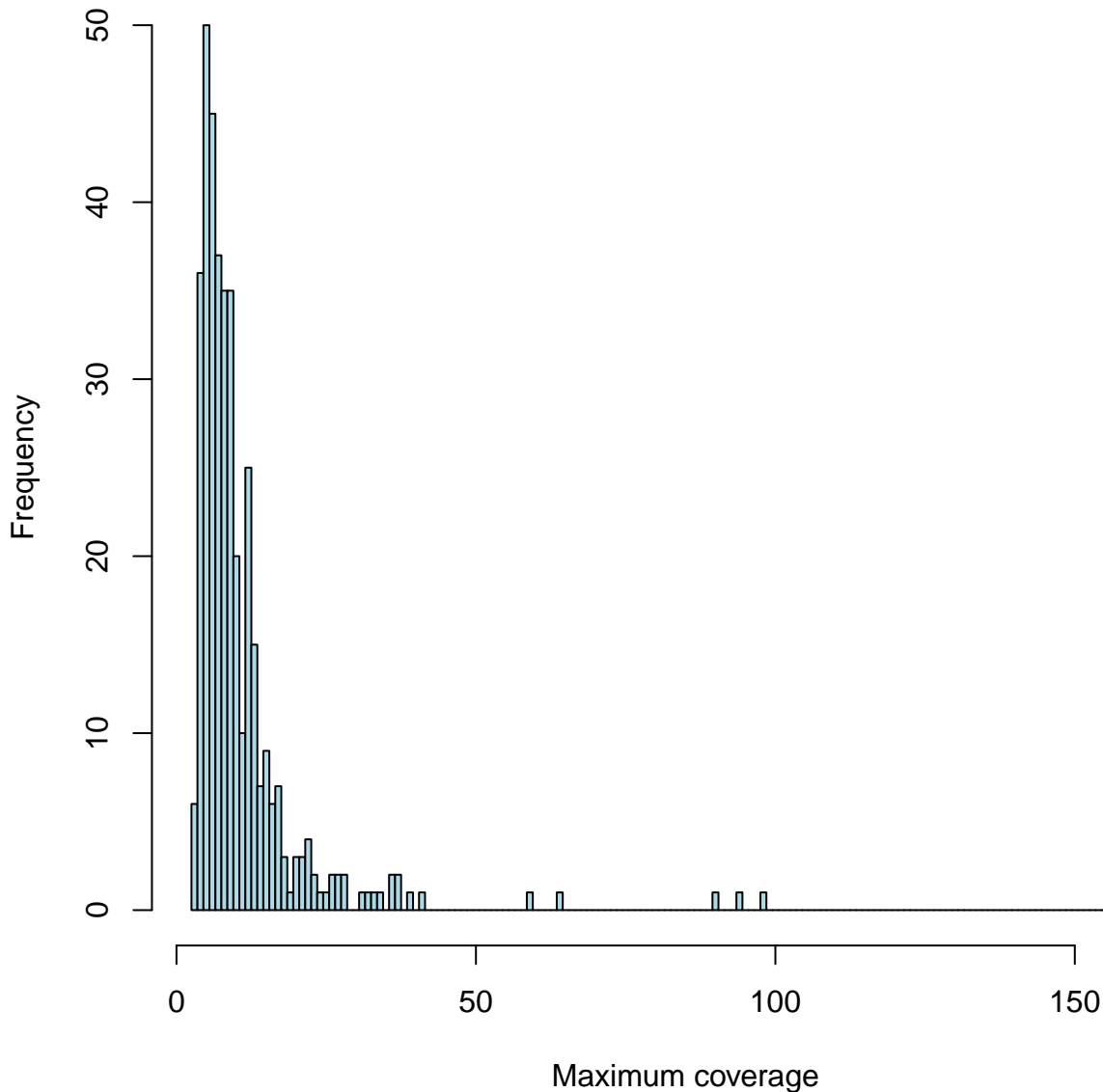
**jx in Intropolis but not "annotated" by knownGene:
78 samples; n jx: 369**



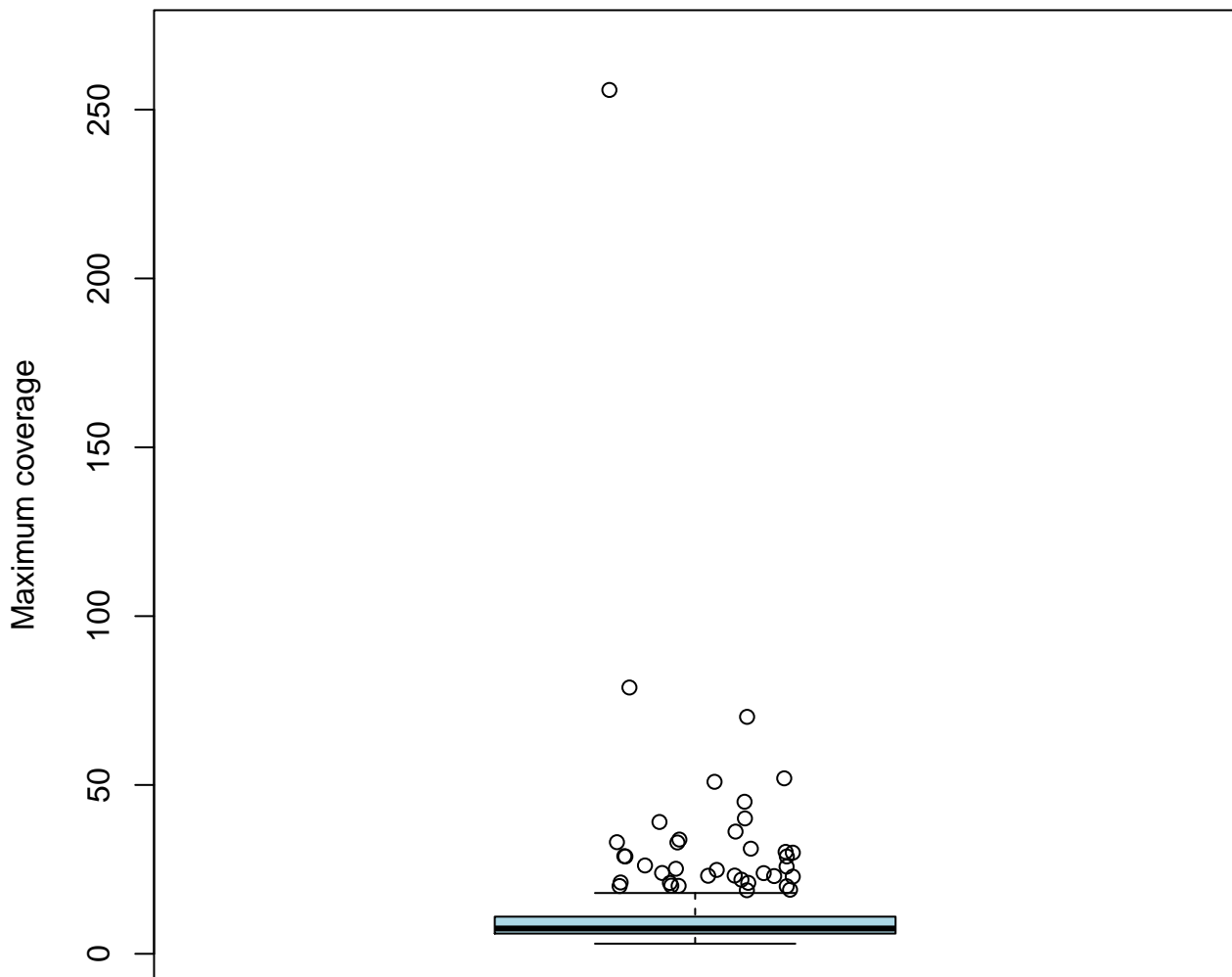
**jx in Intropolis but not "annotated" by knownGene:
79 samples; n jx: 383**



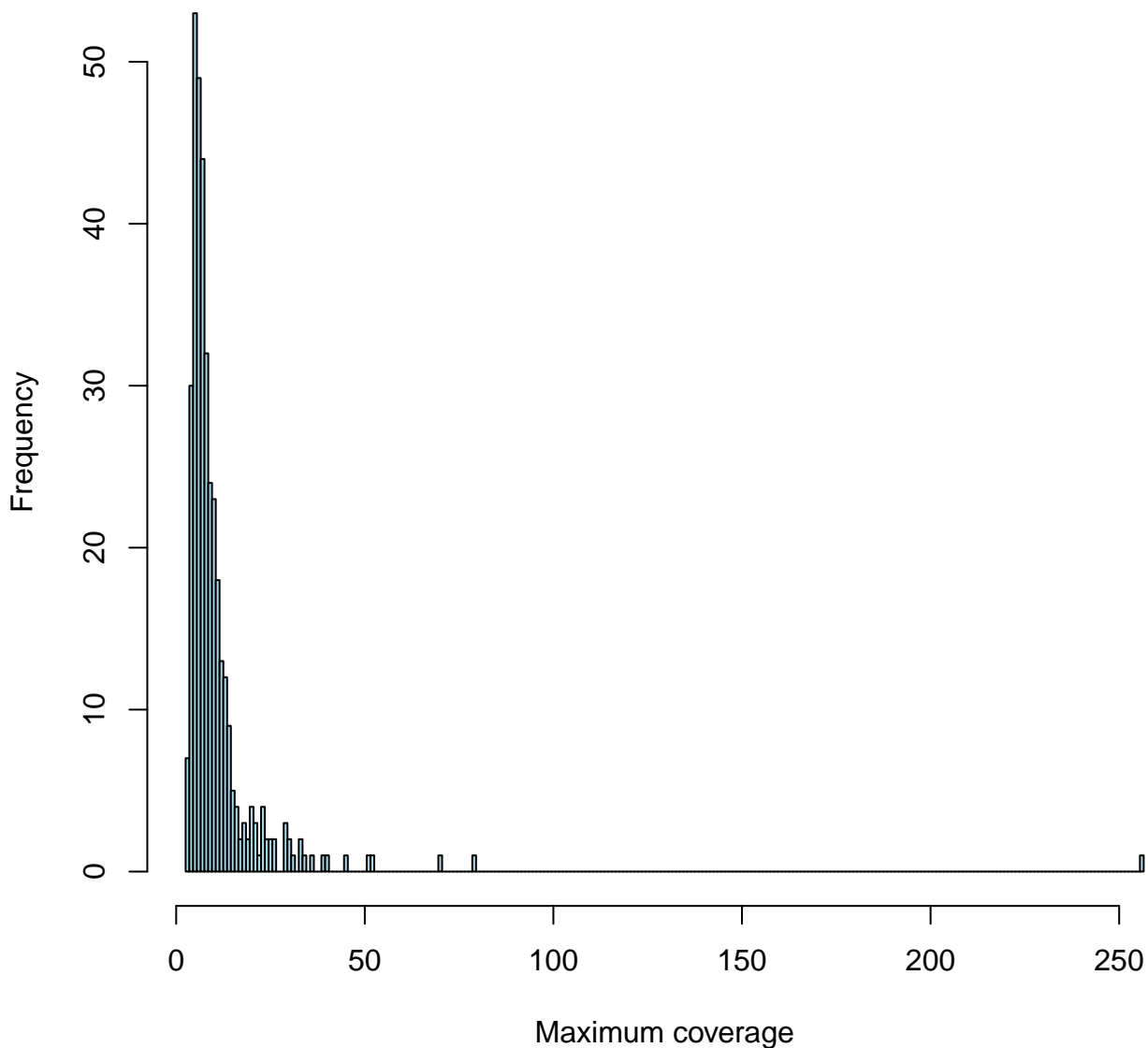
**jx in Intropolis but not "annotated" by knownGene:
79 samples; n jx: 383**



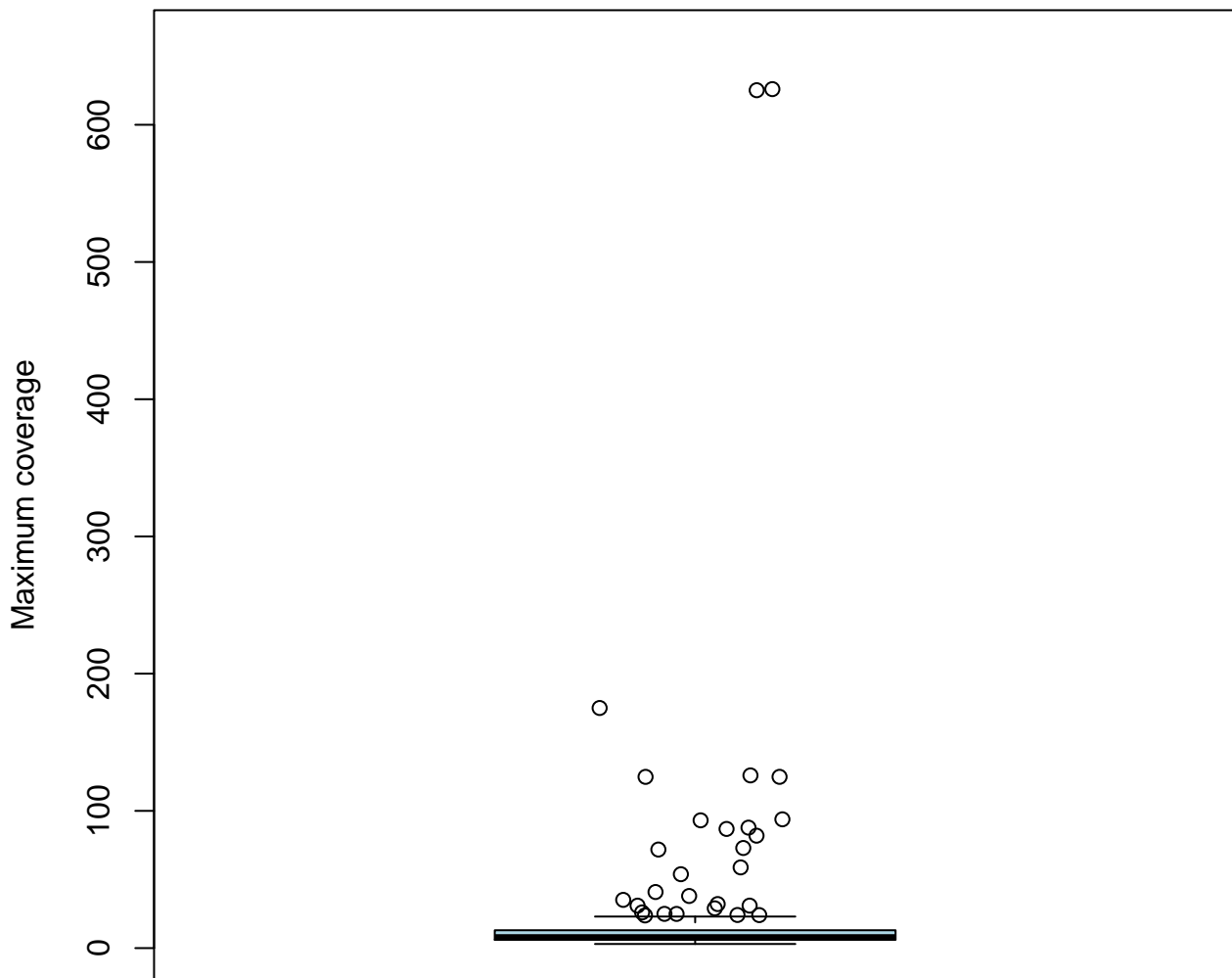
**jx in Intropolis but not "annotated" by knownGene:
80 samples; n jx: 366**



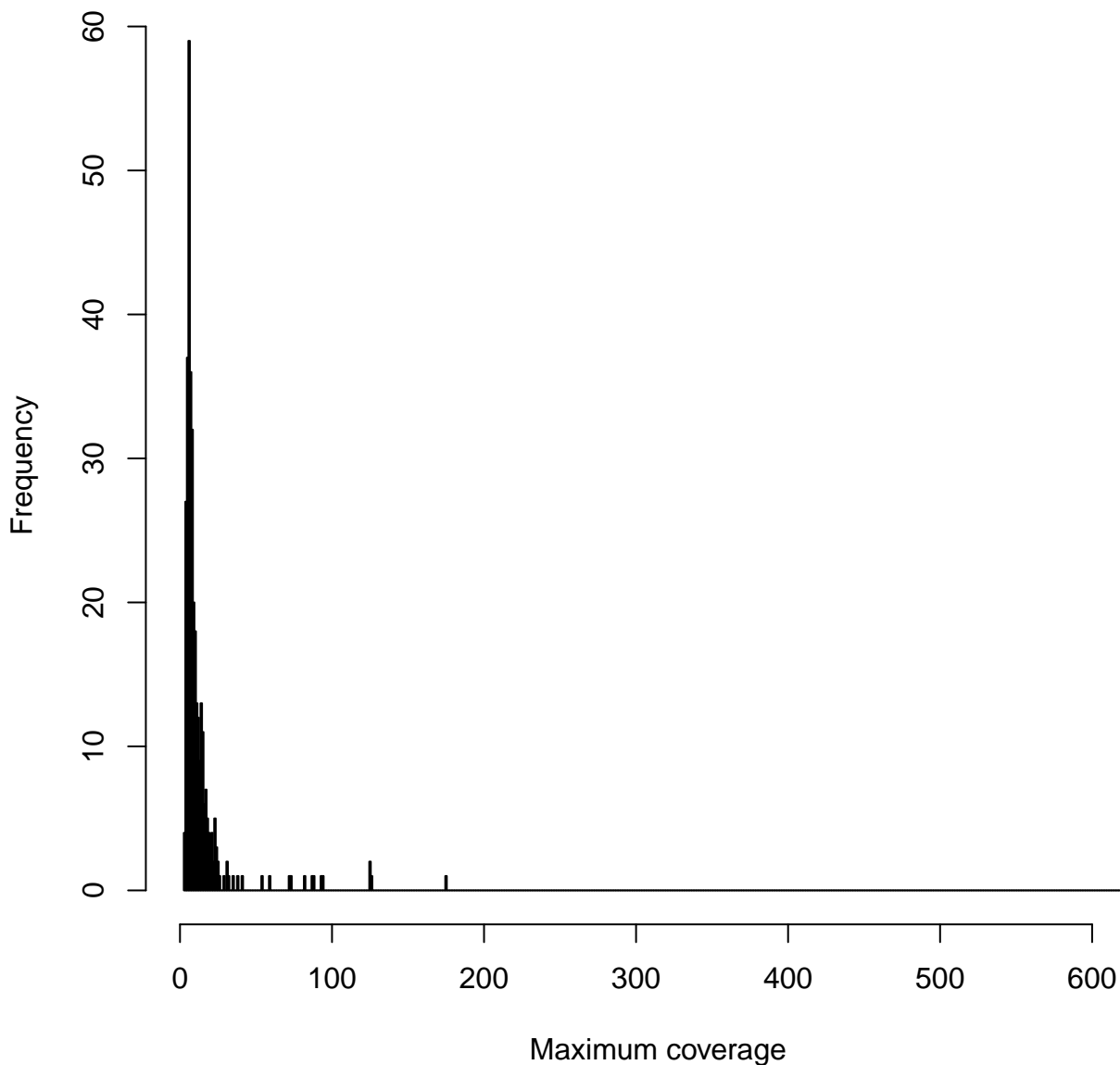
**jx in Intropolis but not "annotated" by knownGene:
80 samples; n jx: 366**



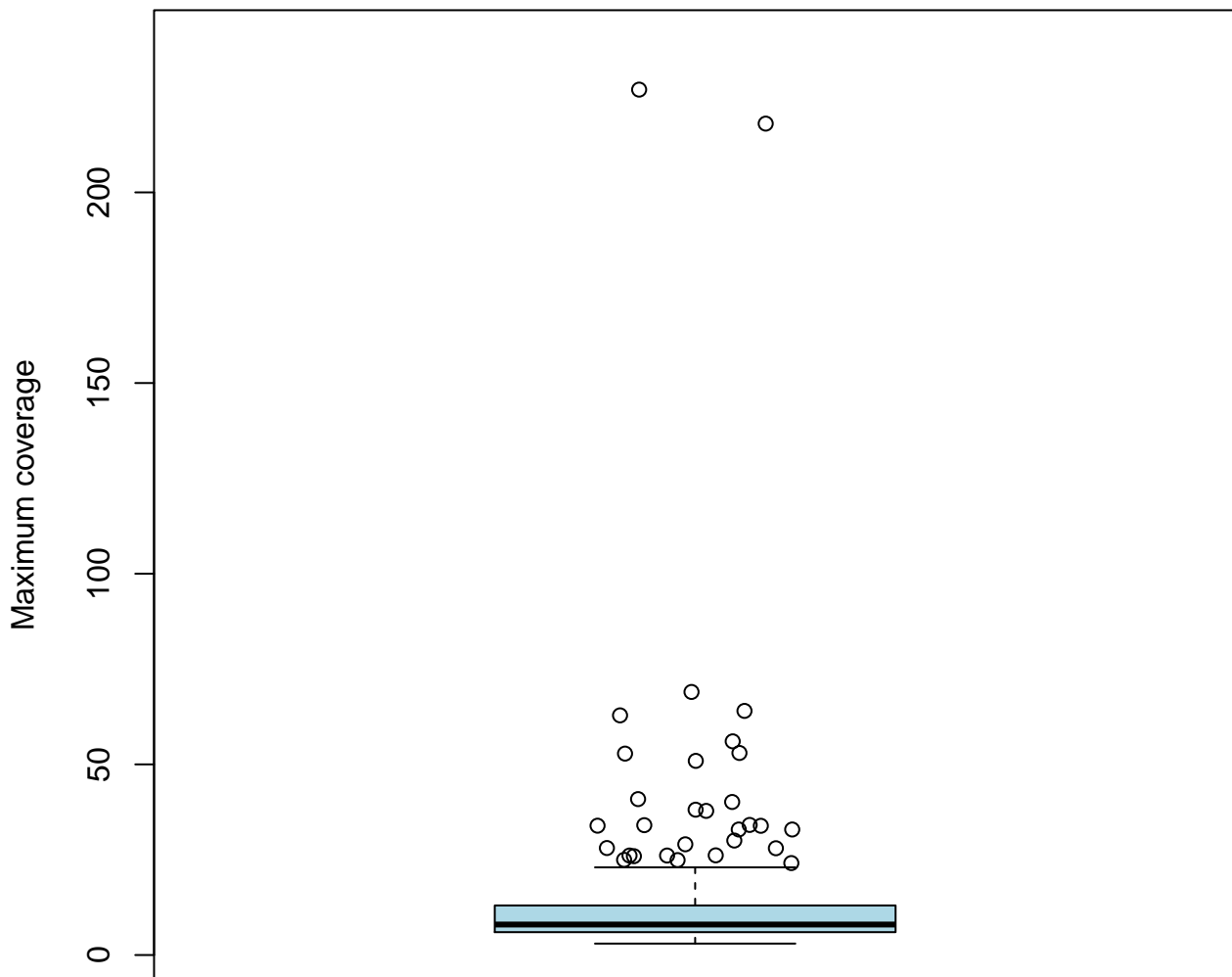
**jx in Intropolis but not "annotated" by knownGene:
81 samples; n jx: 354**



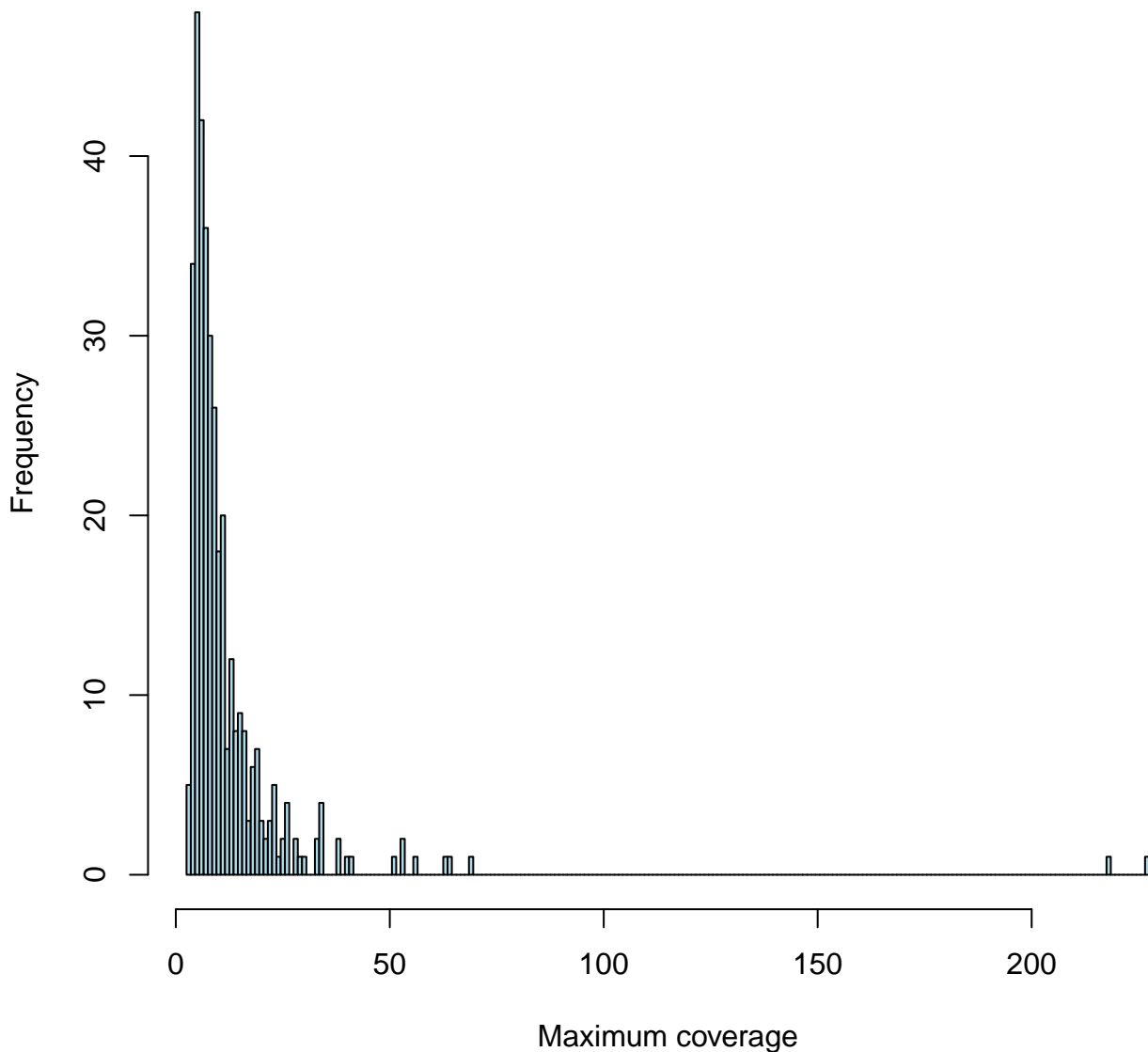
**jx in Intropolis but not "annotated" by knownGene:
81 samples; n jx: 354**



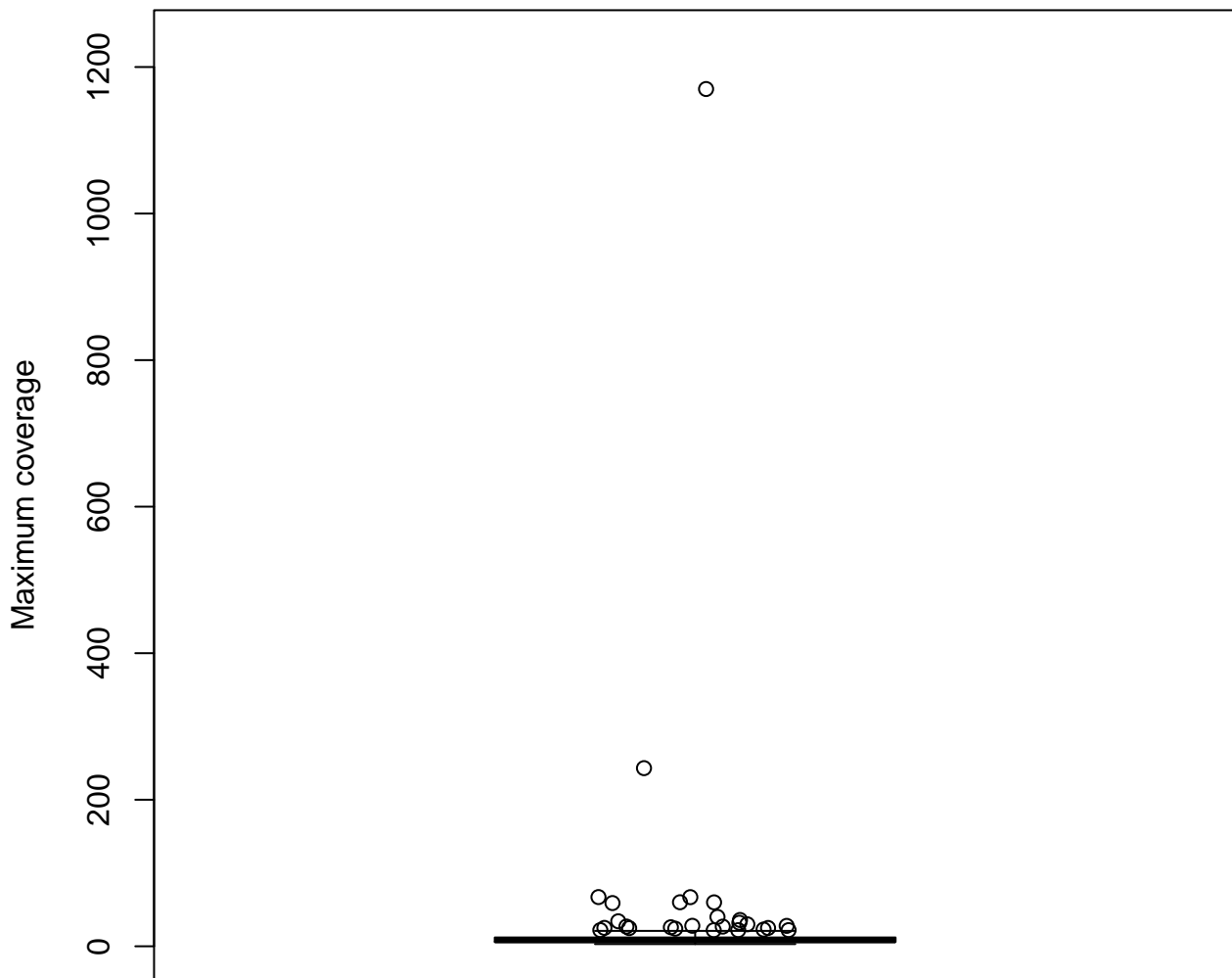
**jx in Intropolis but not "annotated" by knownGene:
82 samples; n jx: 362**



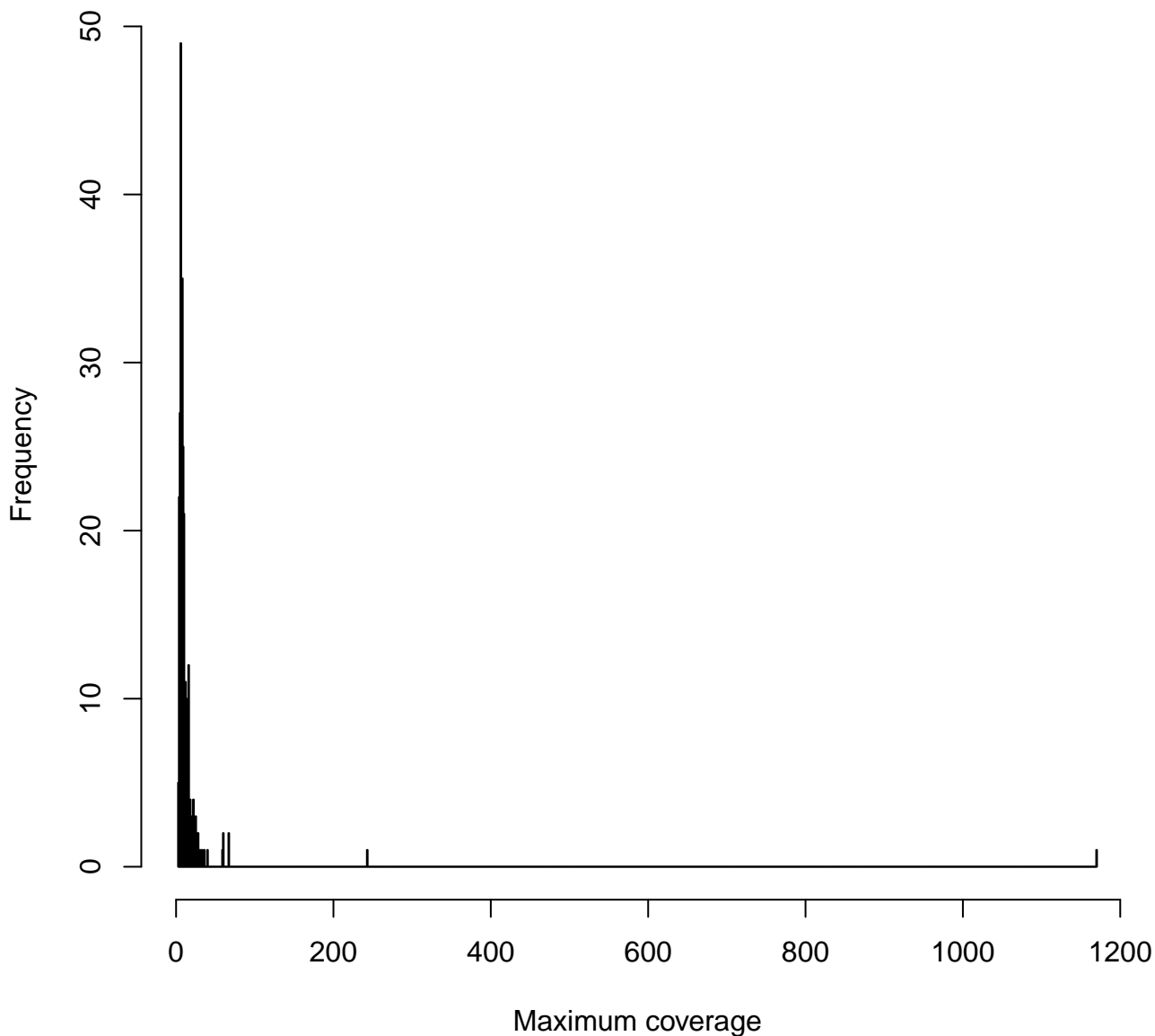
**jx in Intropolis but not "annotated" by knownGene:
82 samples; n jx: 362**



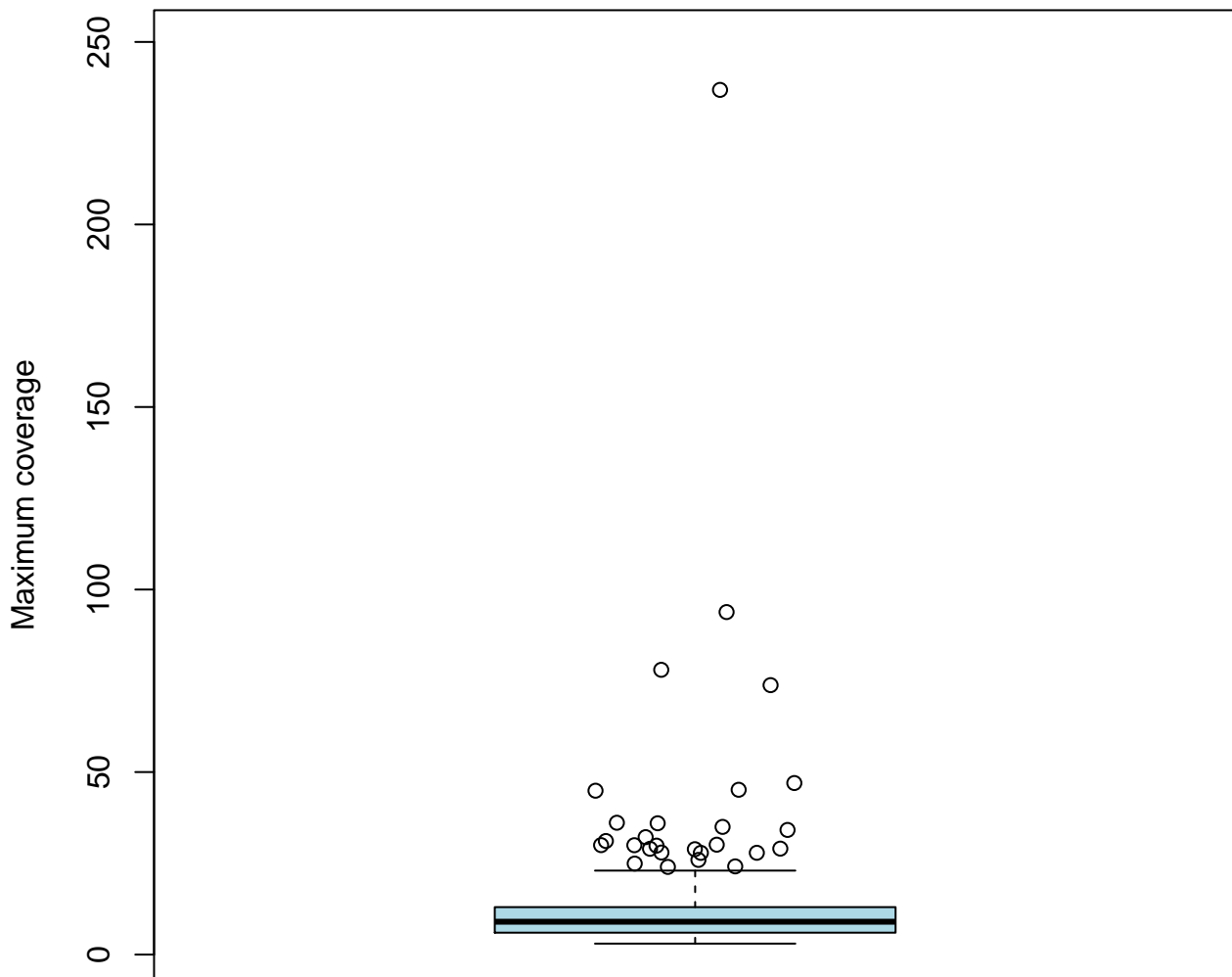
**jx in Intropolis but not "annotated" by knownGene:
83 samples; n jx: 314**



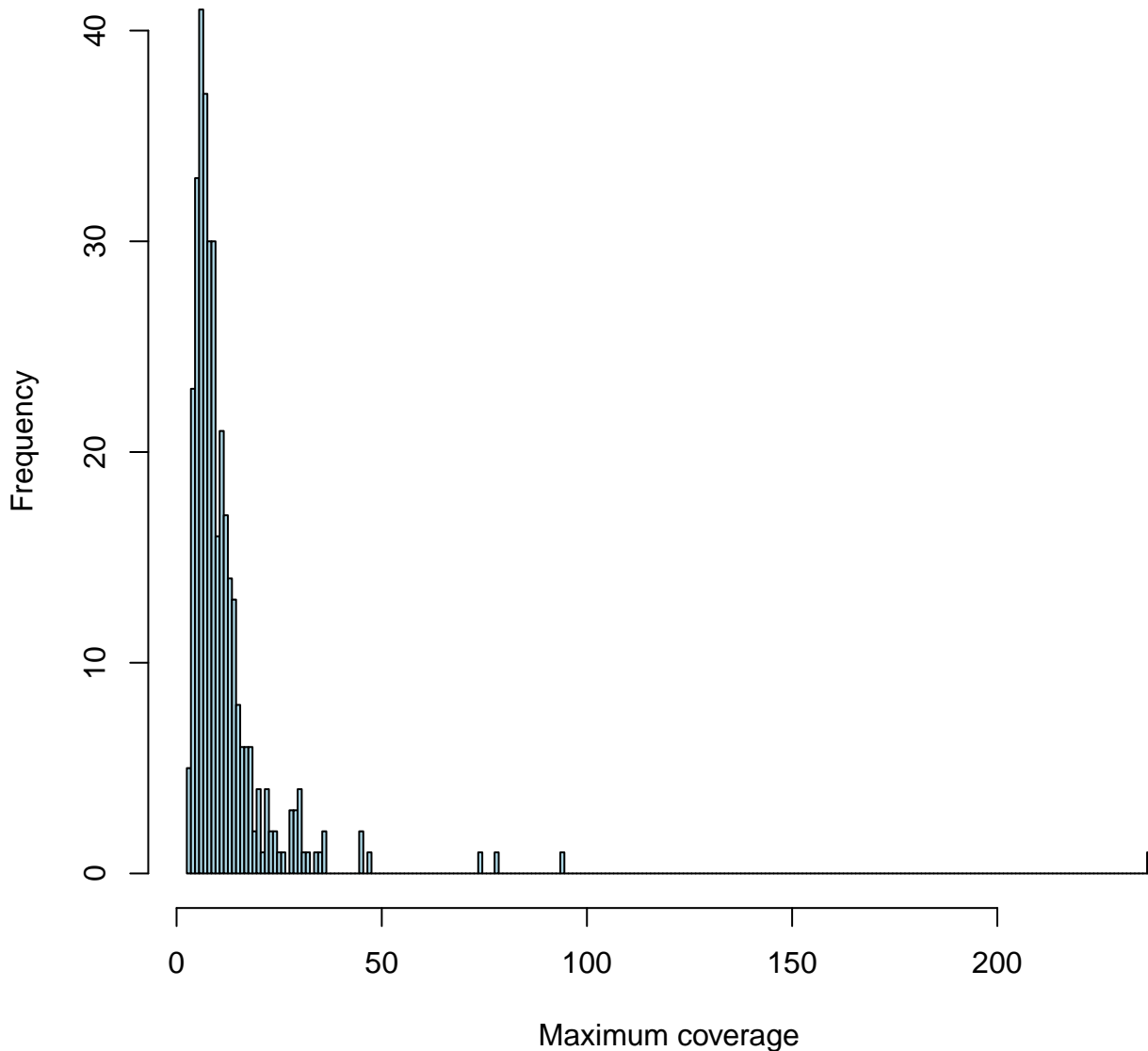
**jx in Intropolis but not "annotated" by knownGene:
83 samples; n jx: 314**



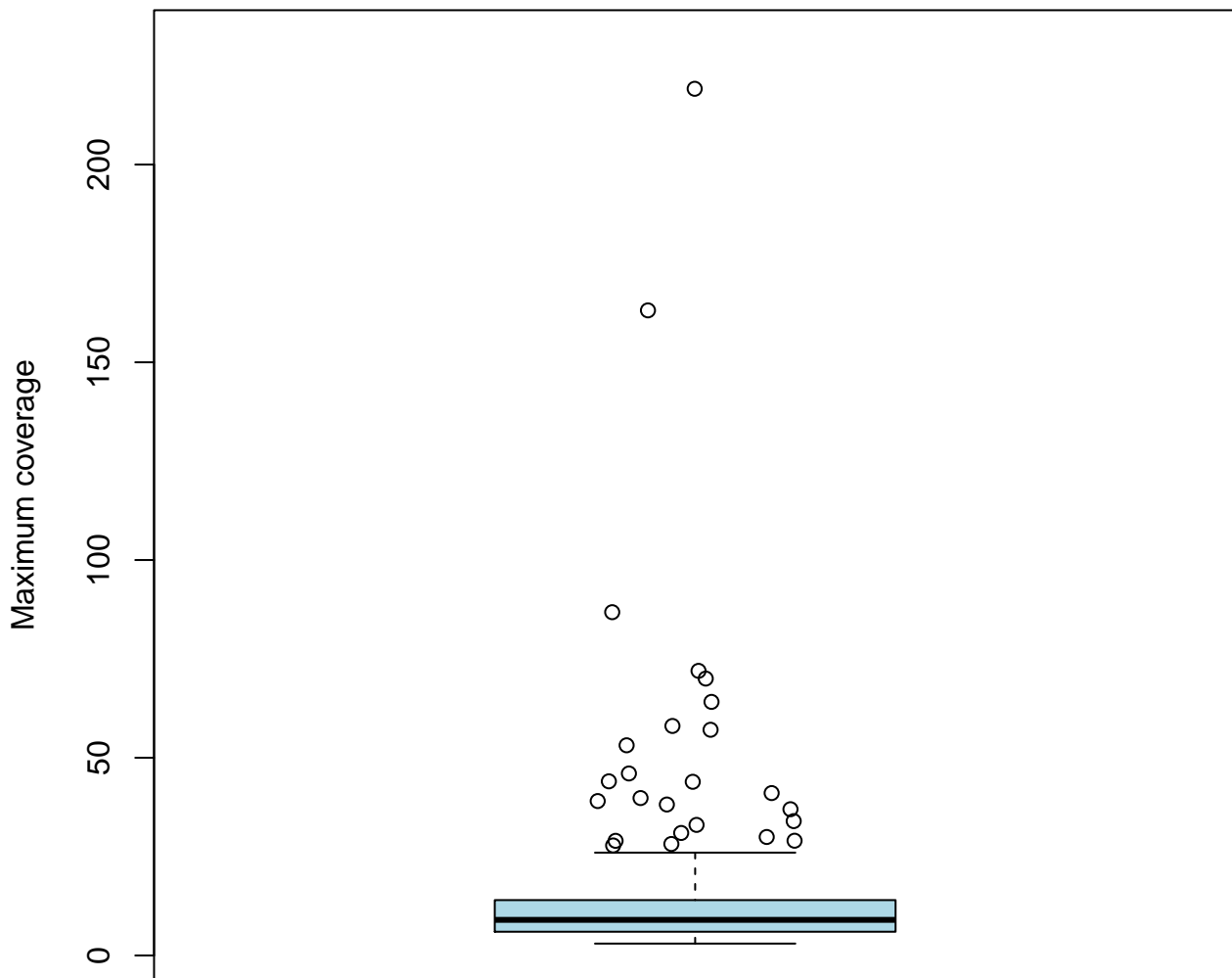
**jx in Intropolis but not "annotated" by knownGene:
84 samples; n jx: 346**



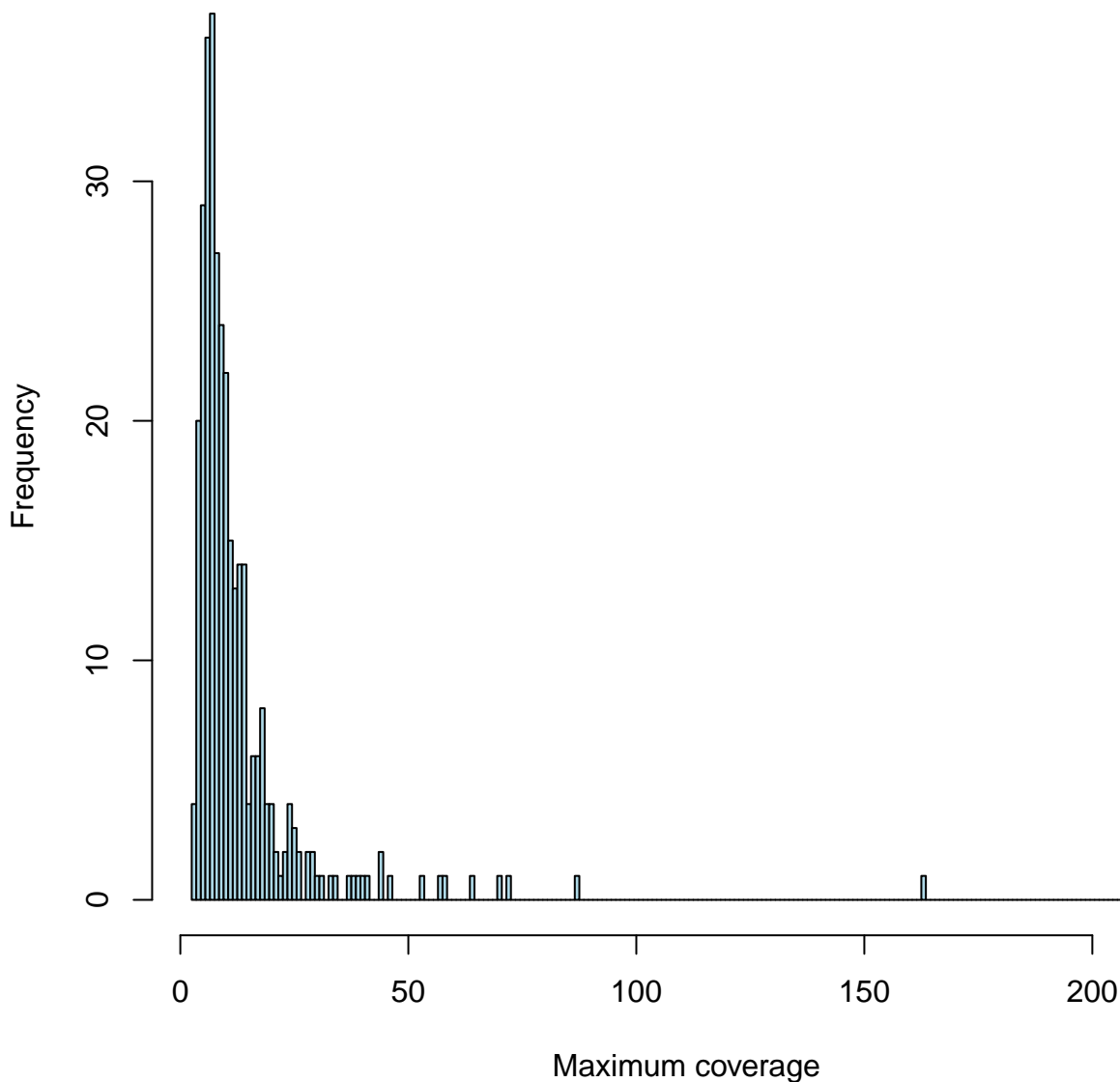
**jx in Intropolis but not "annotated" by knownGene:
84 samples; n jx: 346**



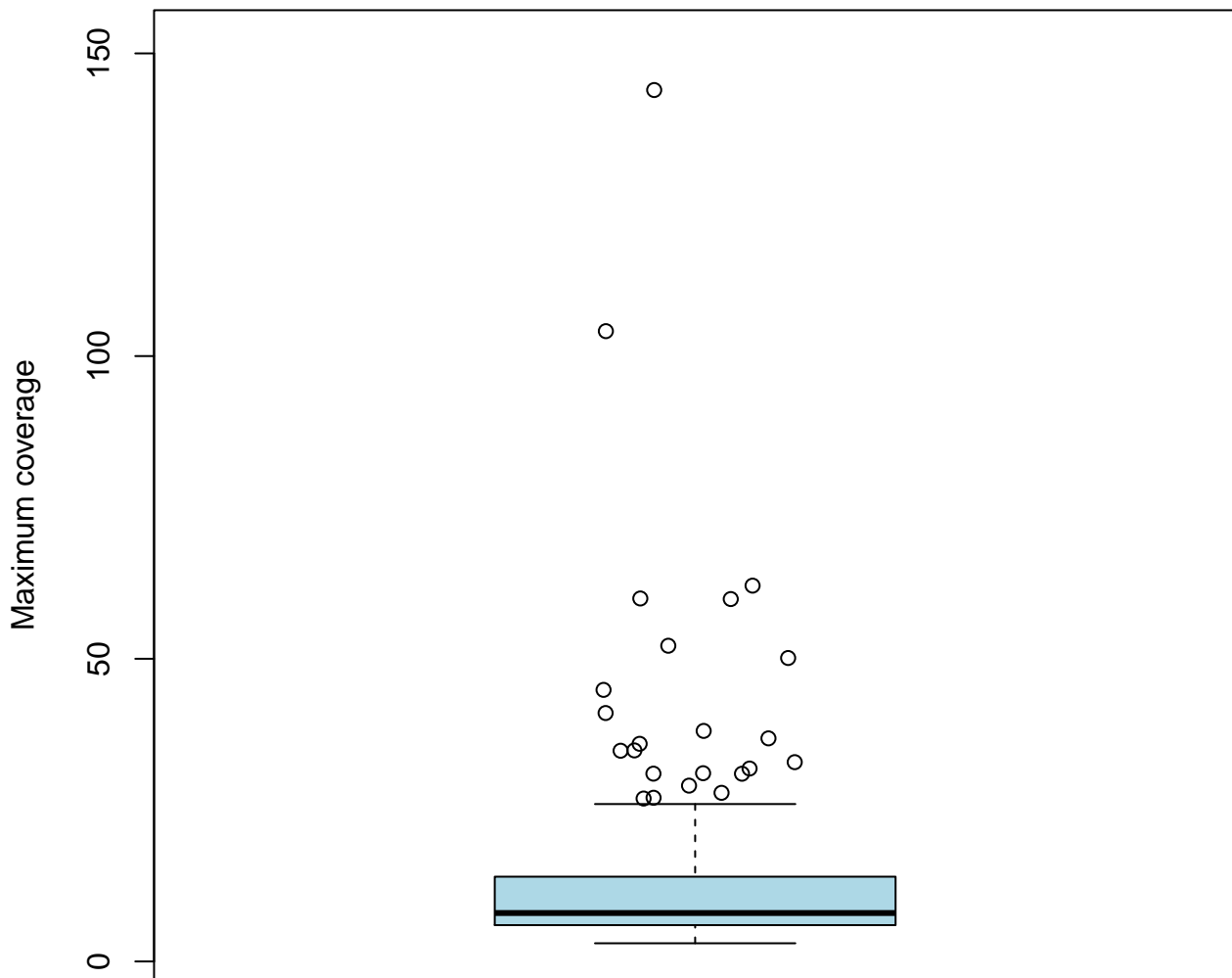
**jx in Intropolis but not "annotated" by knownGene:
85 samples; n jx: 326**



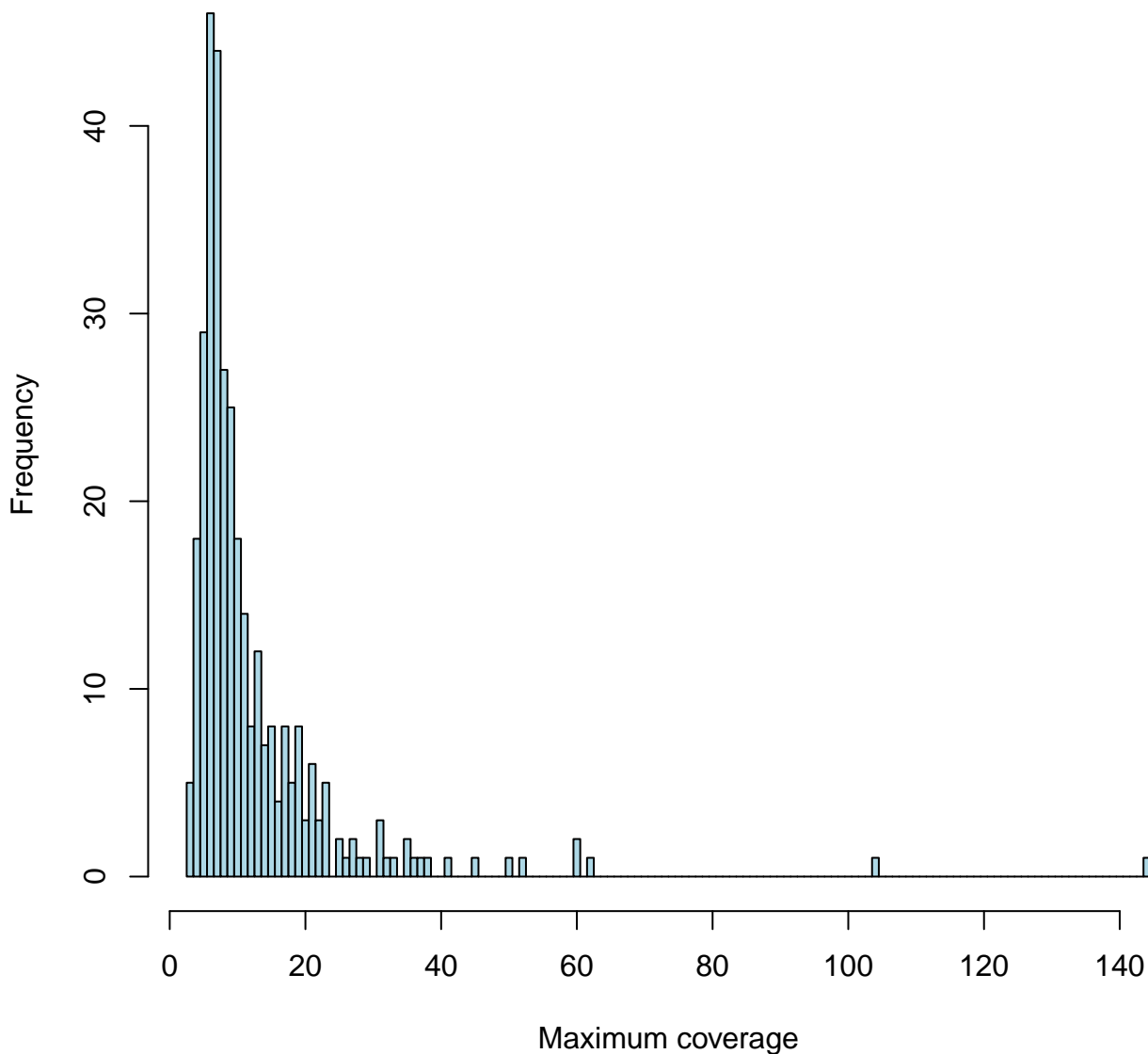
**jx in Intropolis but not "annotated" by knownGene:
85 samples; n jx: 326**



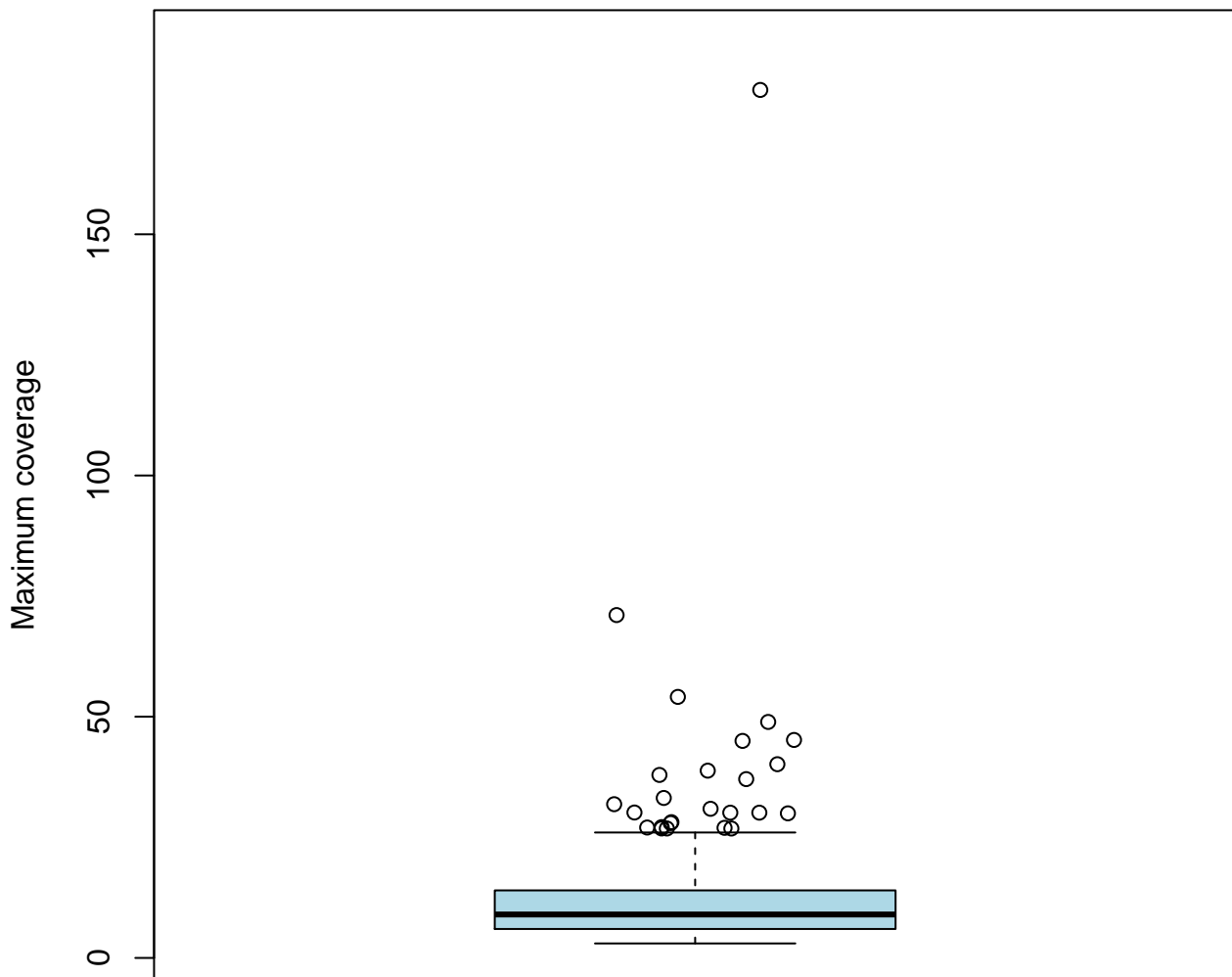
**jx in Intropolis but not "annotated" by knownGene:
86 samples; n jx: 329**



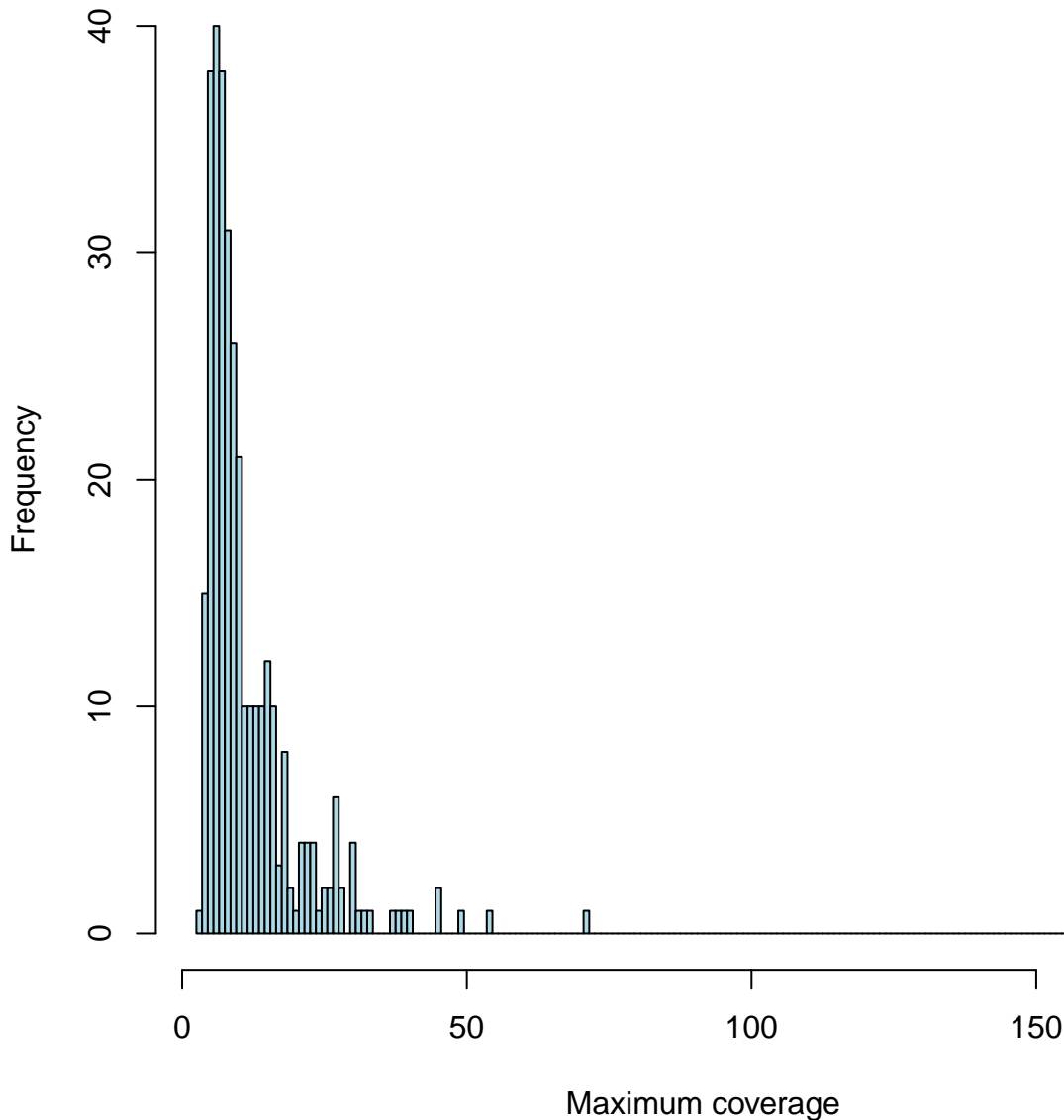
**jx in Intropolis but not "annotated" by knownGene:
86 samples; n jx: 329**



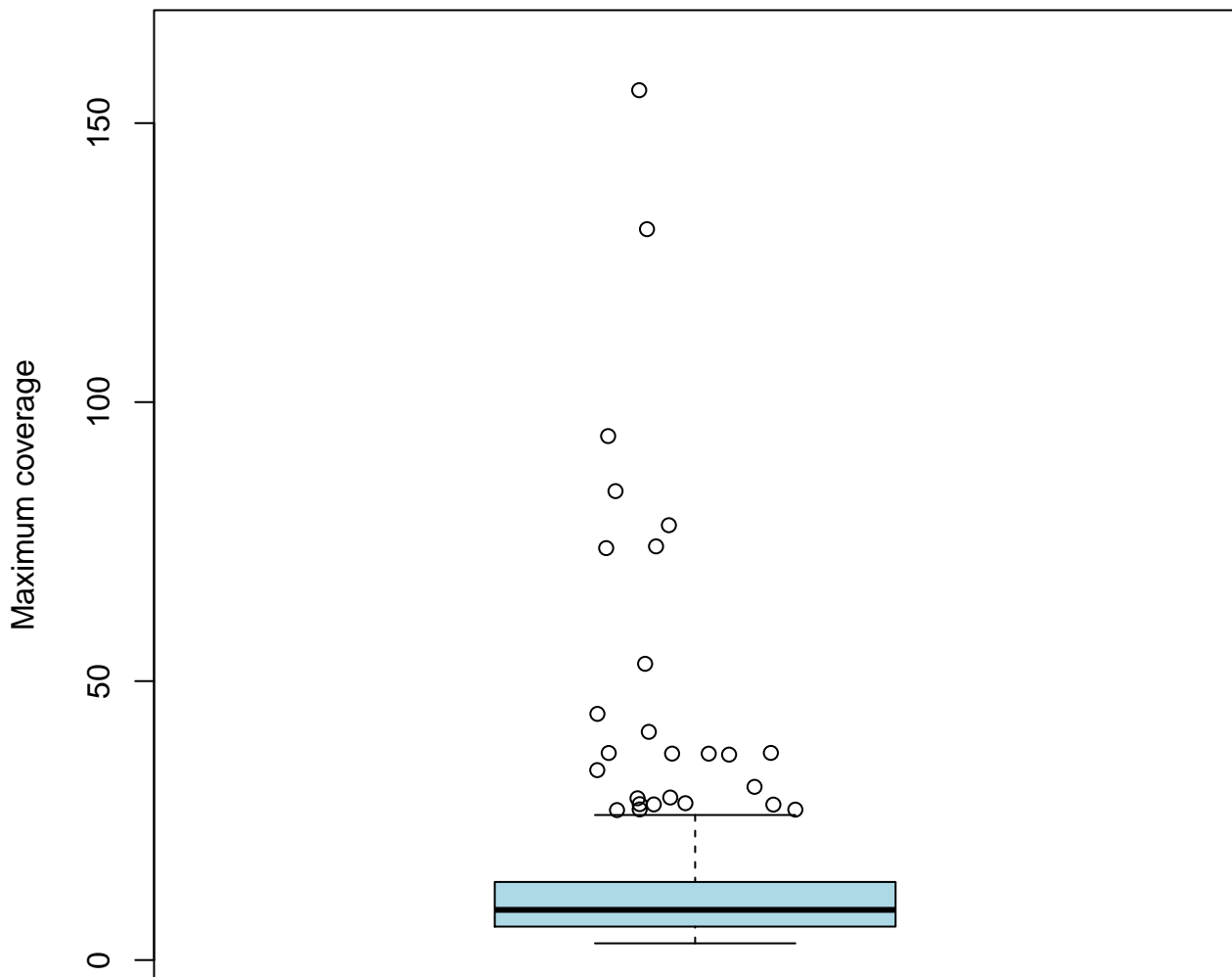
**jx in Intropolis but not "annotated" by knownGene:
87 samples; n jx: 328**



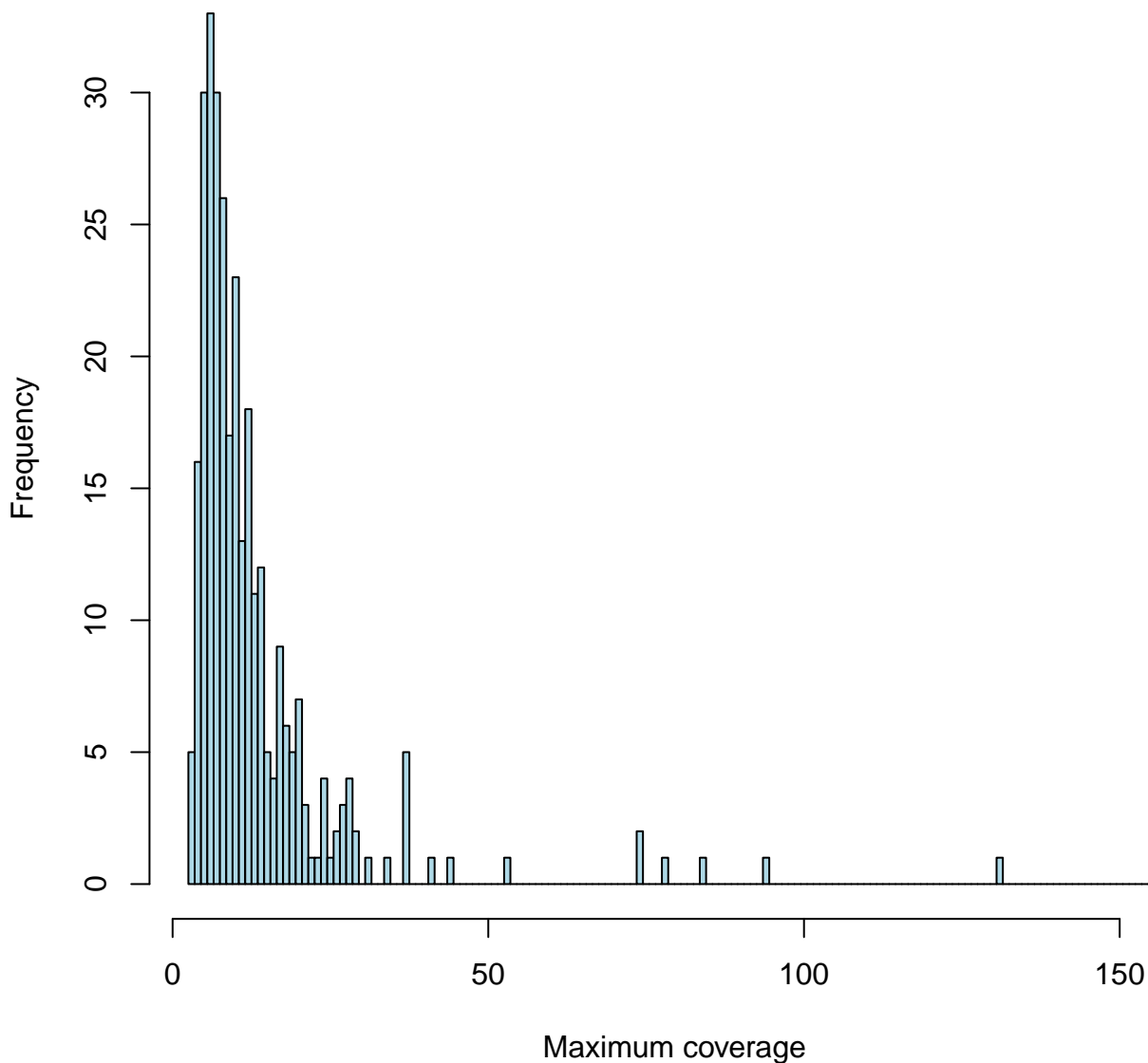
**jx in Intropolis but not "annotated" by knownGene:
87 samples; n jx: 328**



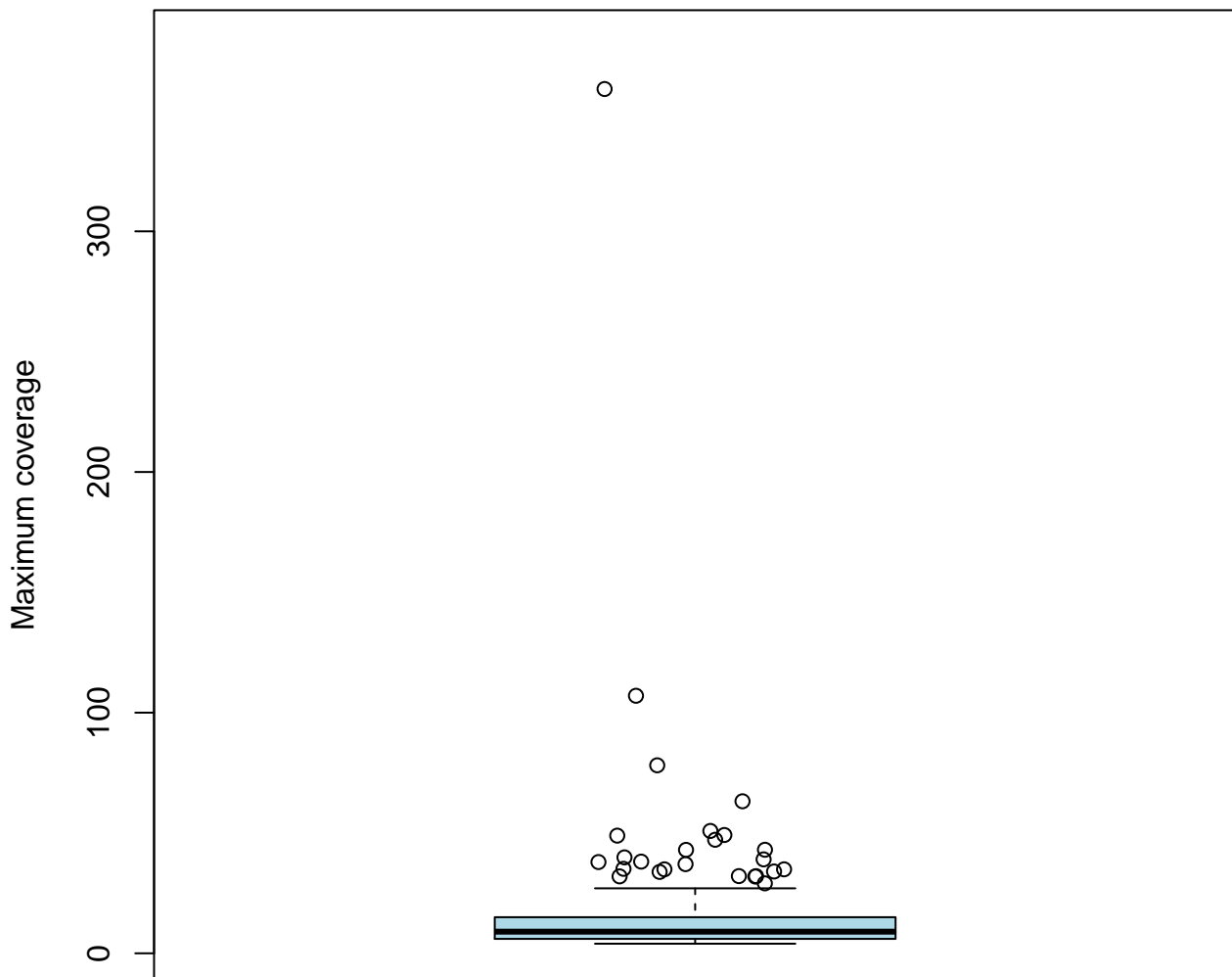
**jx in Intropolis but not "annotated" by knownGene:
88 samples; n jx: 308**



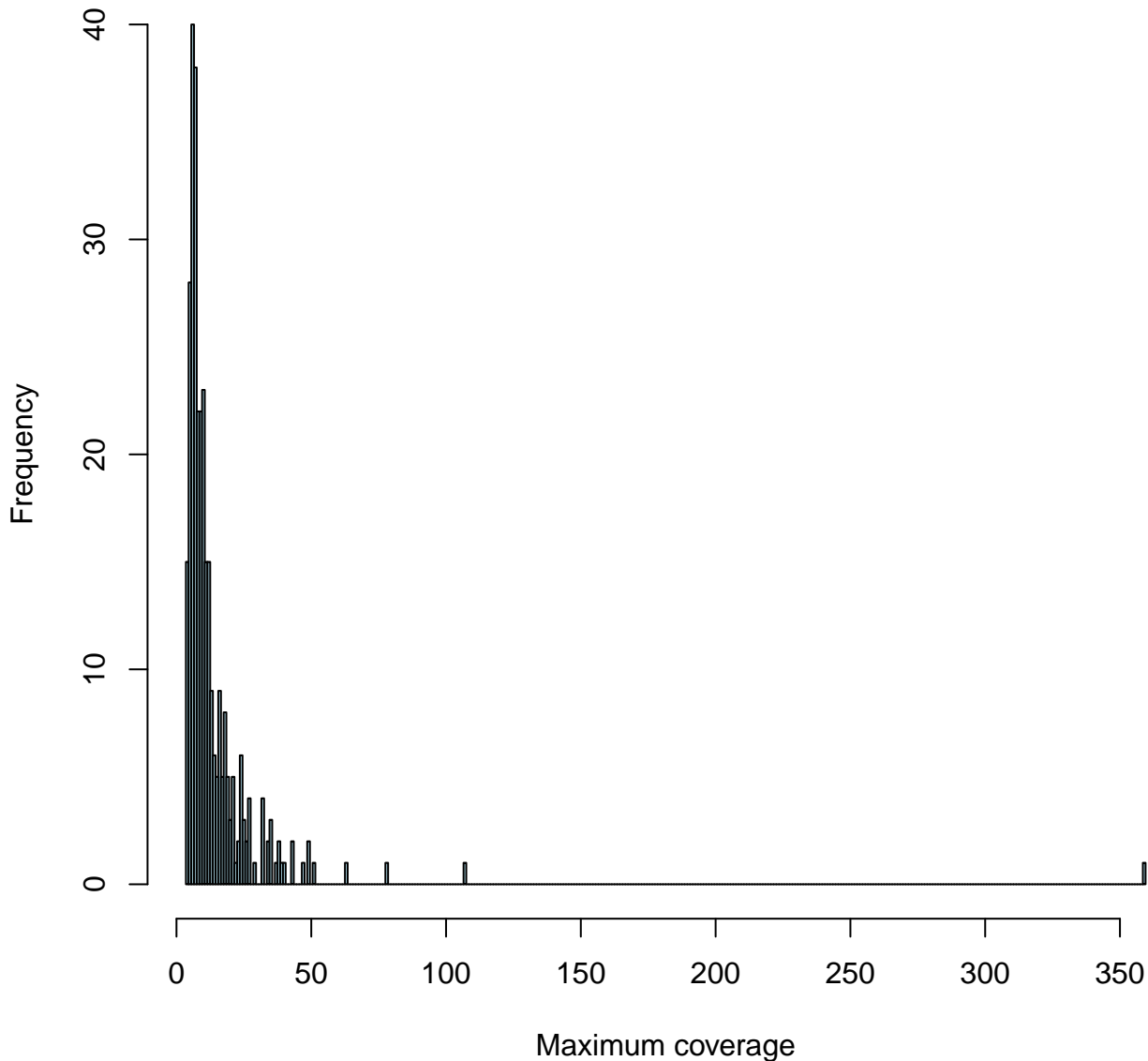
**jx in Intropolis but not "annotated" by knownGene:
88 samples; n jx: 308**



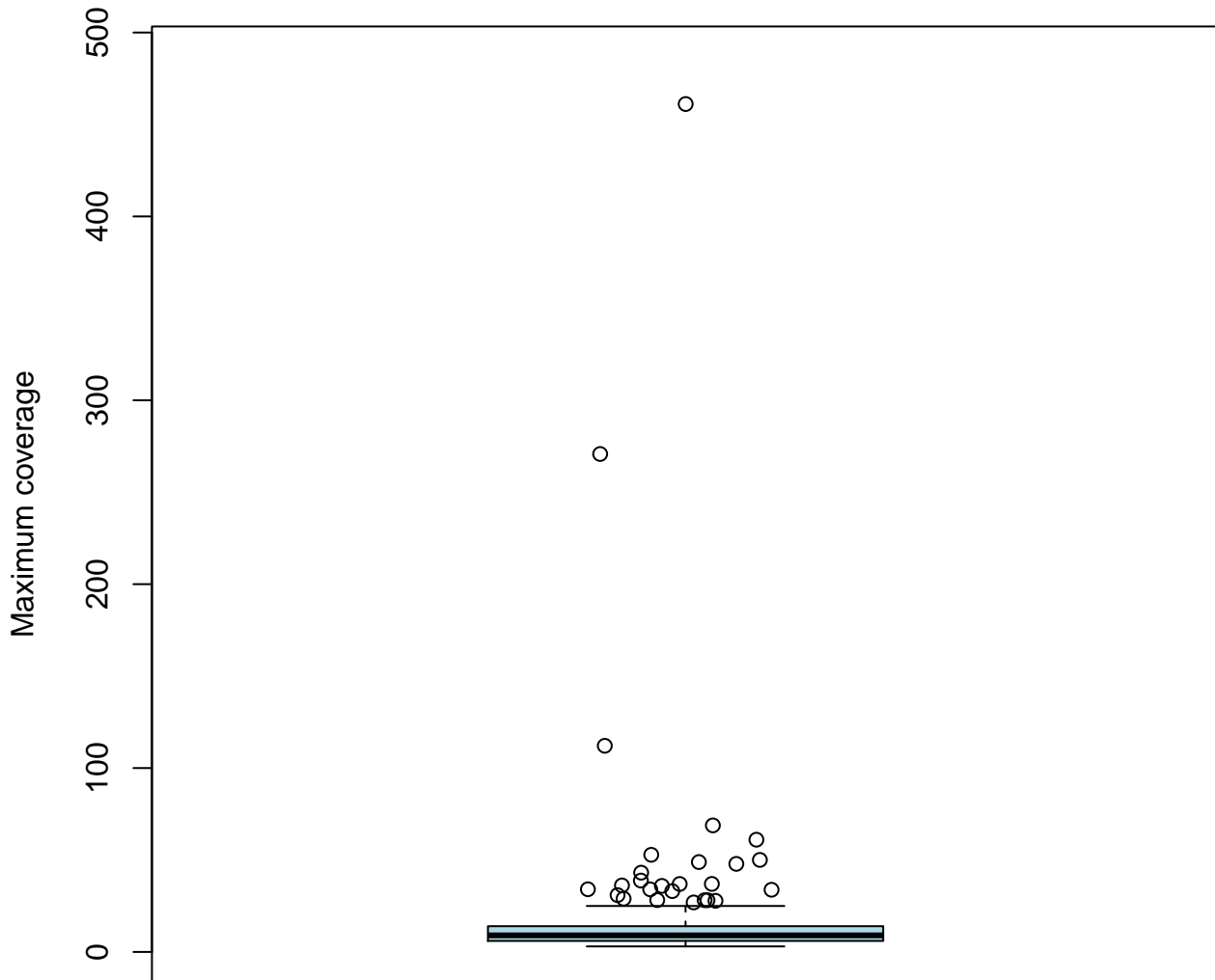
**jx in Intropolis but not "annotated" by knownGene:
89 samples; n jx: 316**



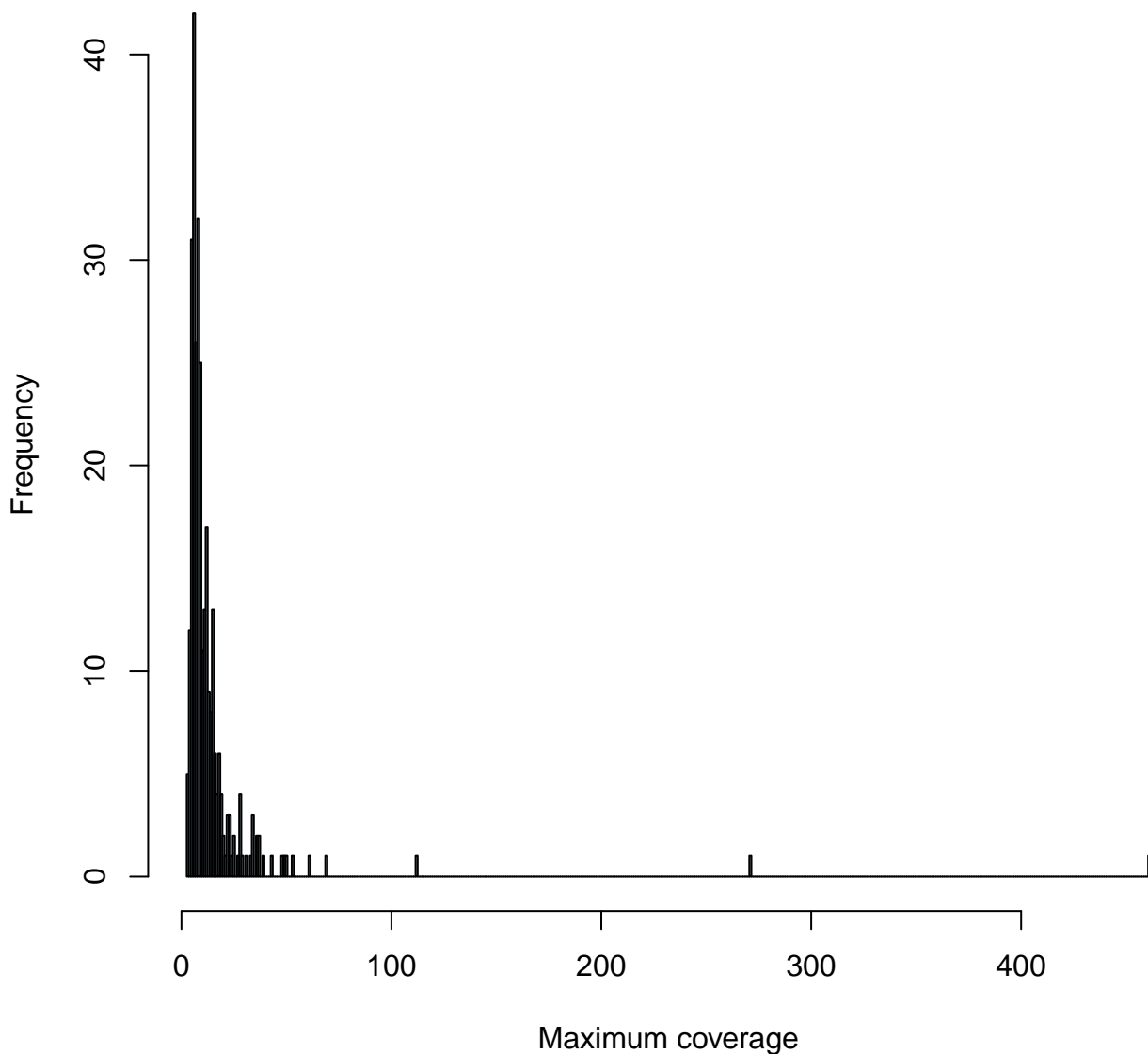
**jx in Intropolis but not "annotated" by knownGene:
89 samples; n jx: 316**



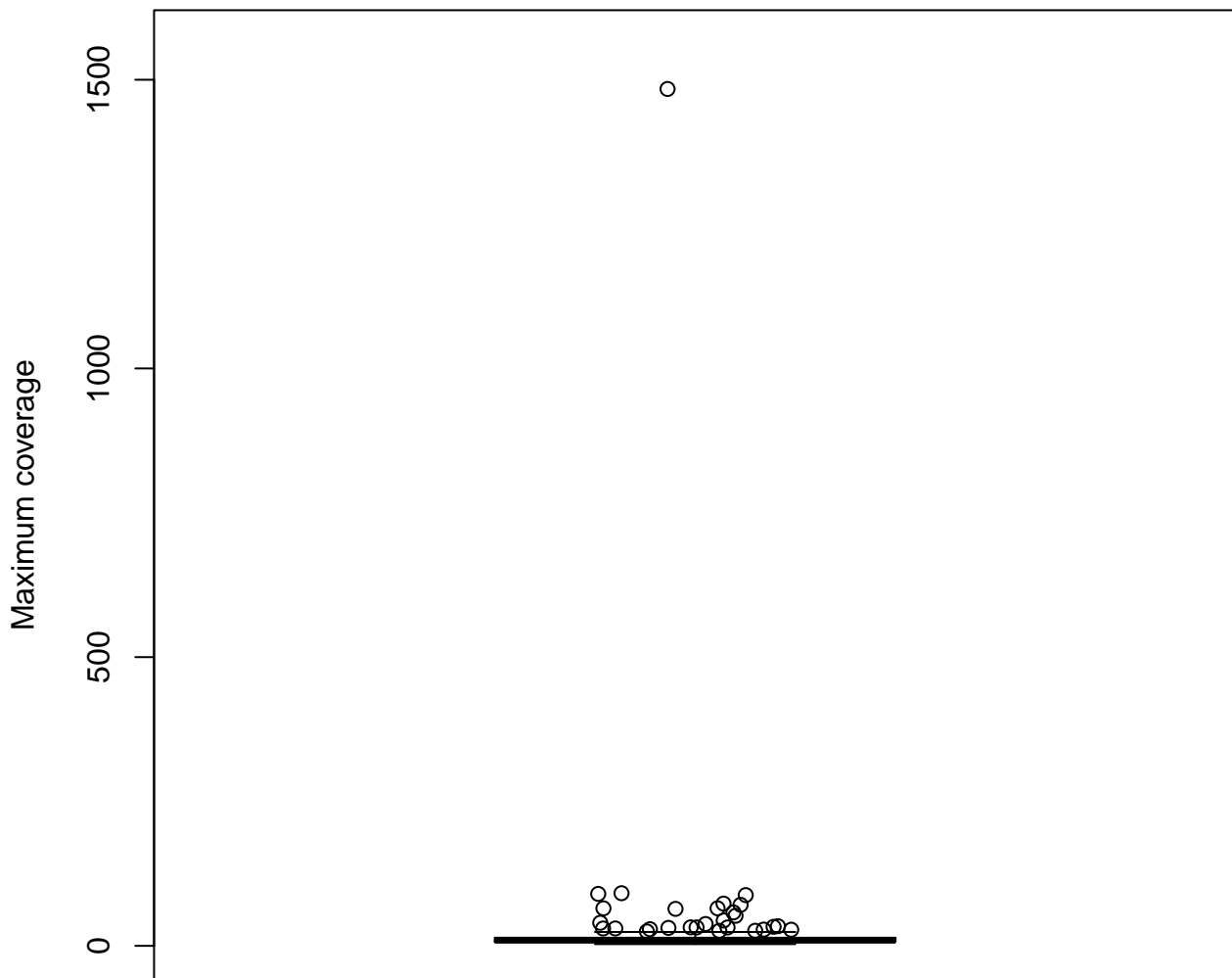
**jx in Intropolis but not "annotated" by knownGene:
90 samples; n jx: 302**



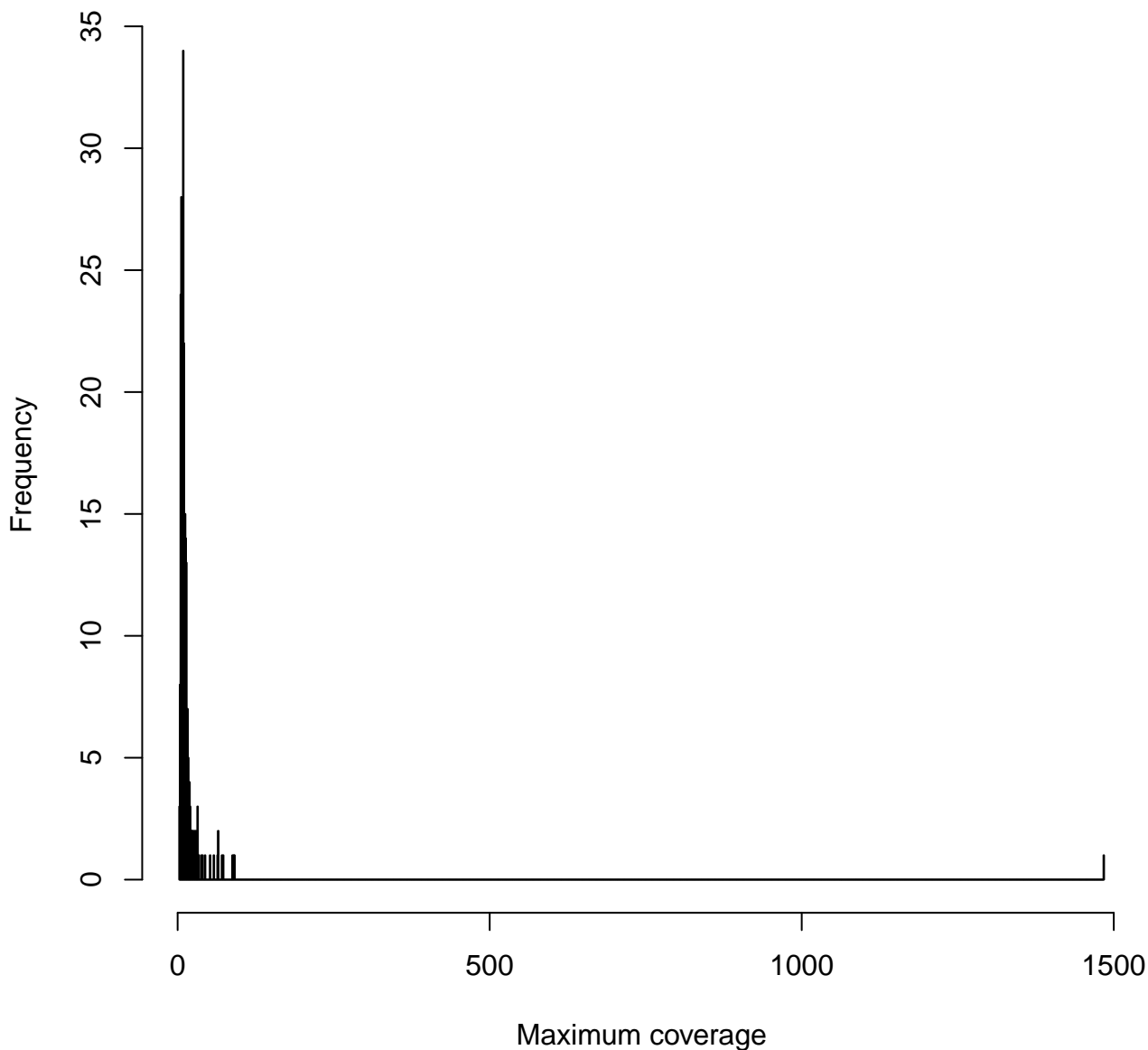
**jx in Intropolis but not "annotated" by knownGene:
90 samples; n jx: 302**



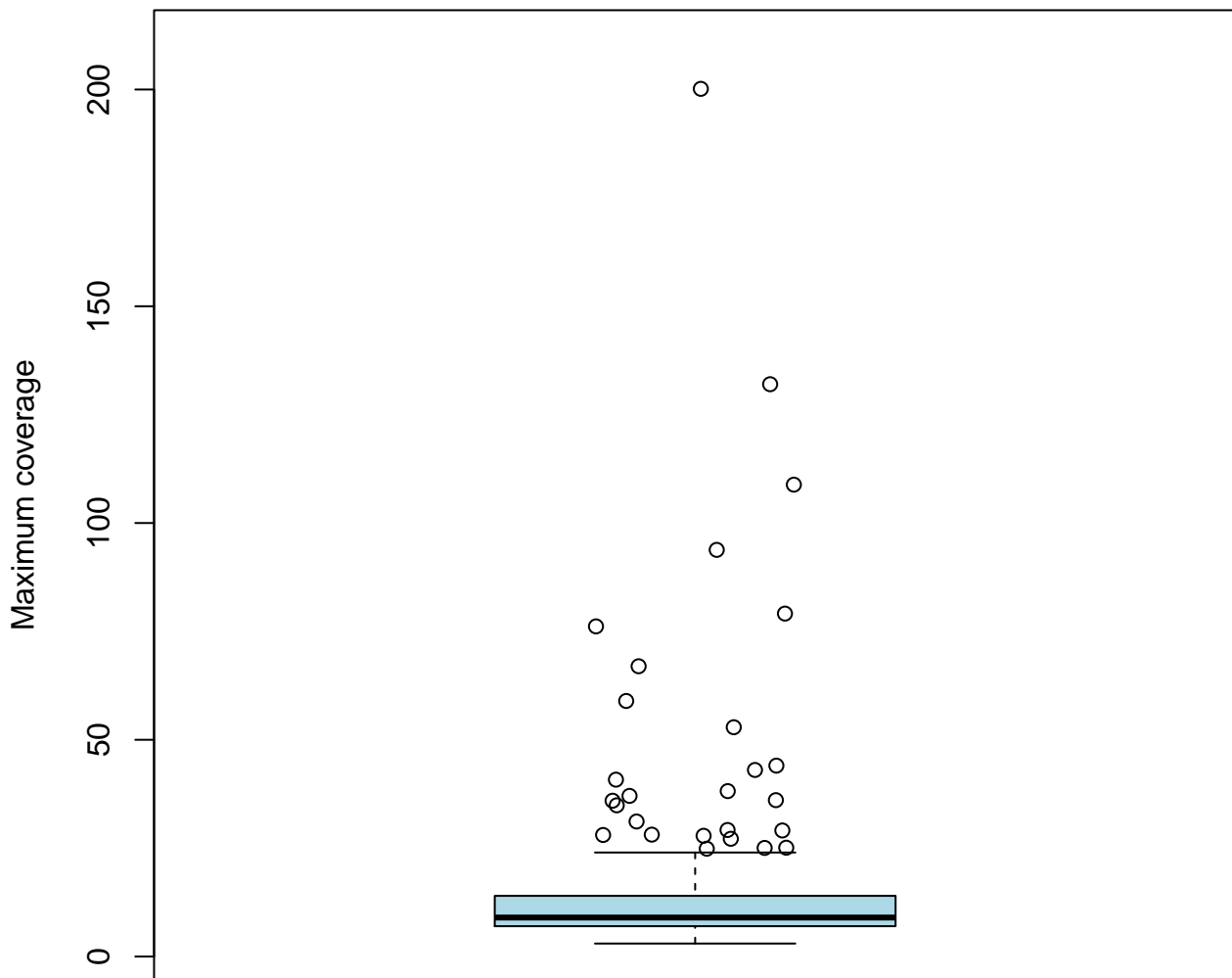
**jx in Intropolis but not "annotated" by knownGene:
91 samples; n jx: 285**



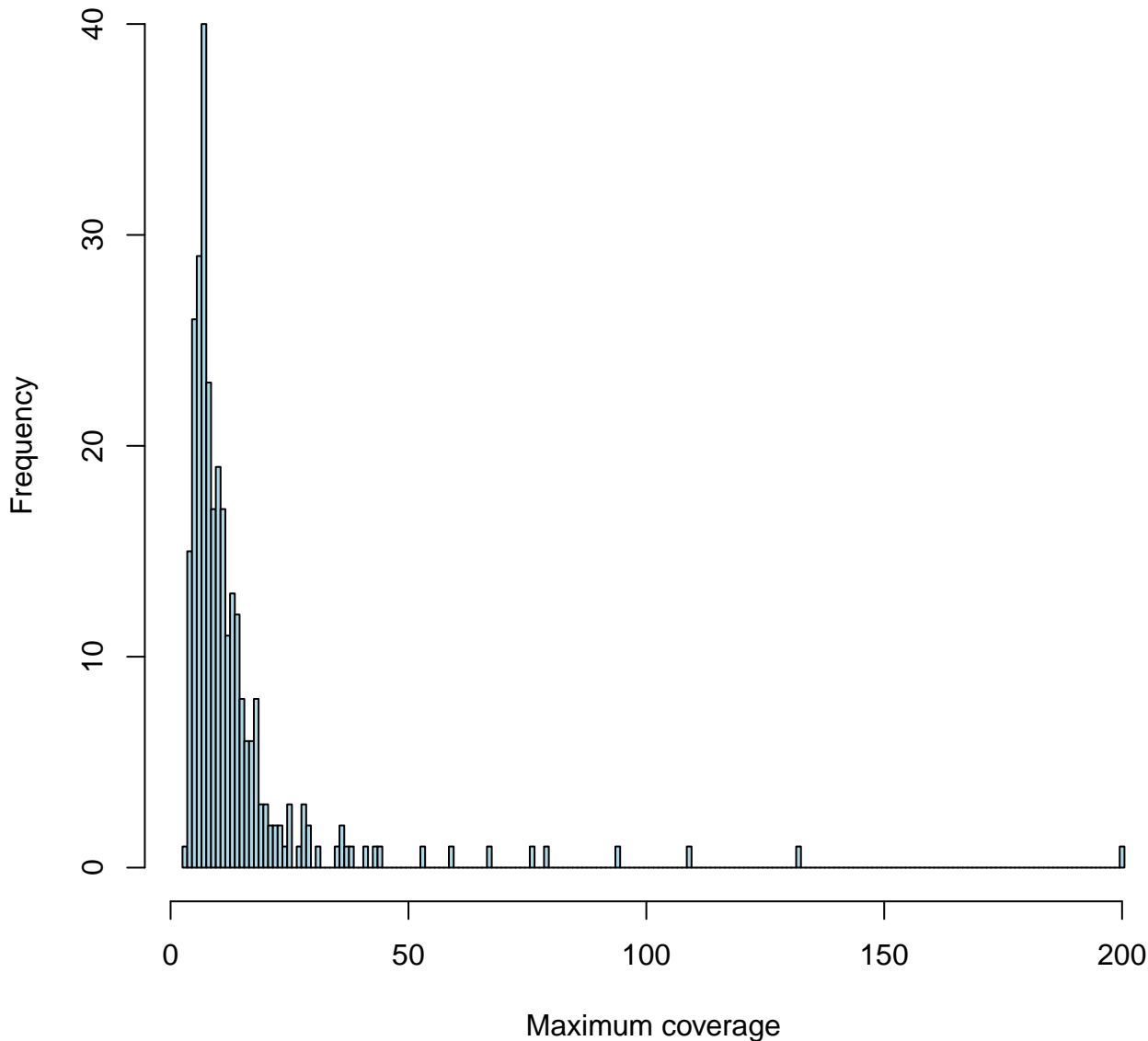
**jx in Intropolis but not "annotated" by knownGene:
91 samples; n jx: 285**



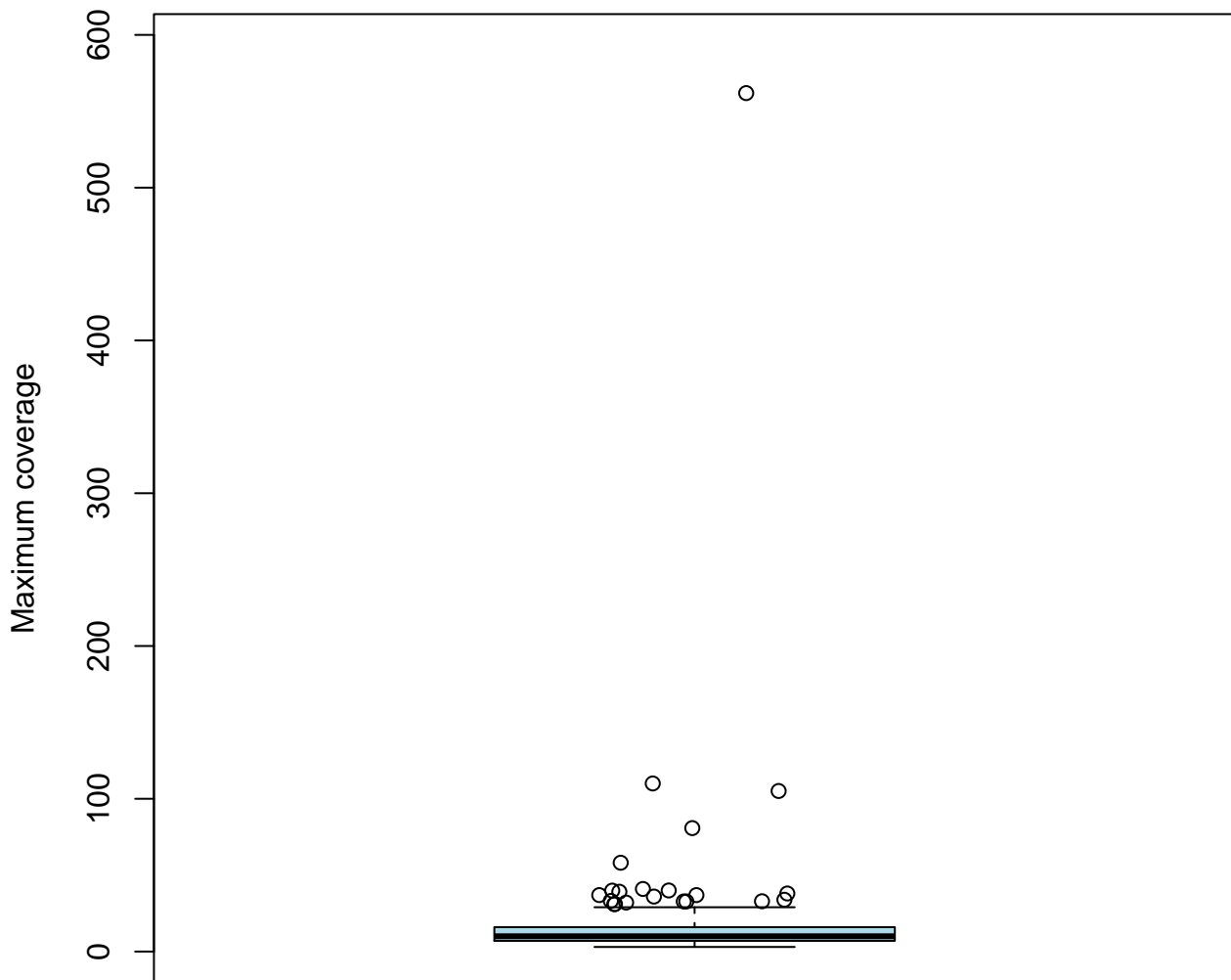
**jx in Intropolis but not "annotated" by knownGene:
92 samples; n jx: 291**



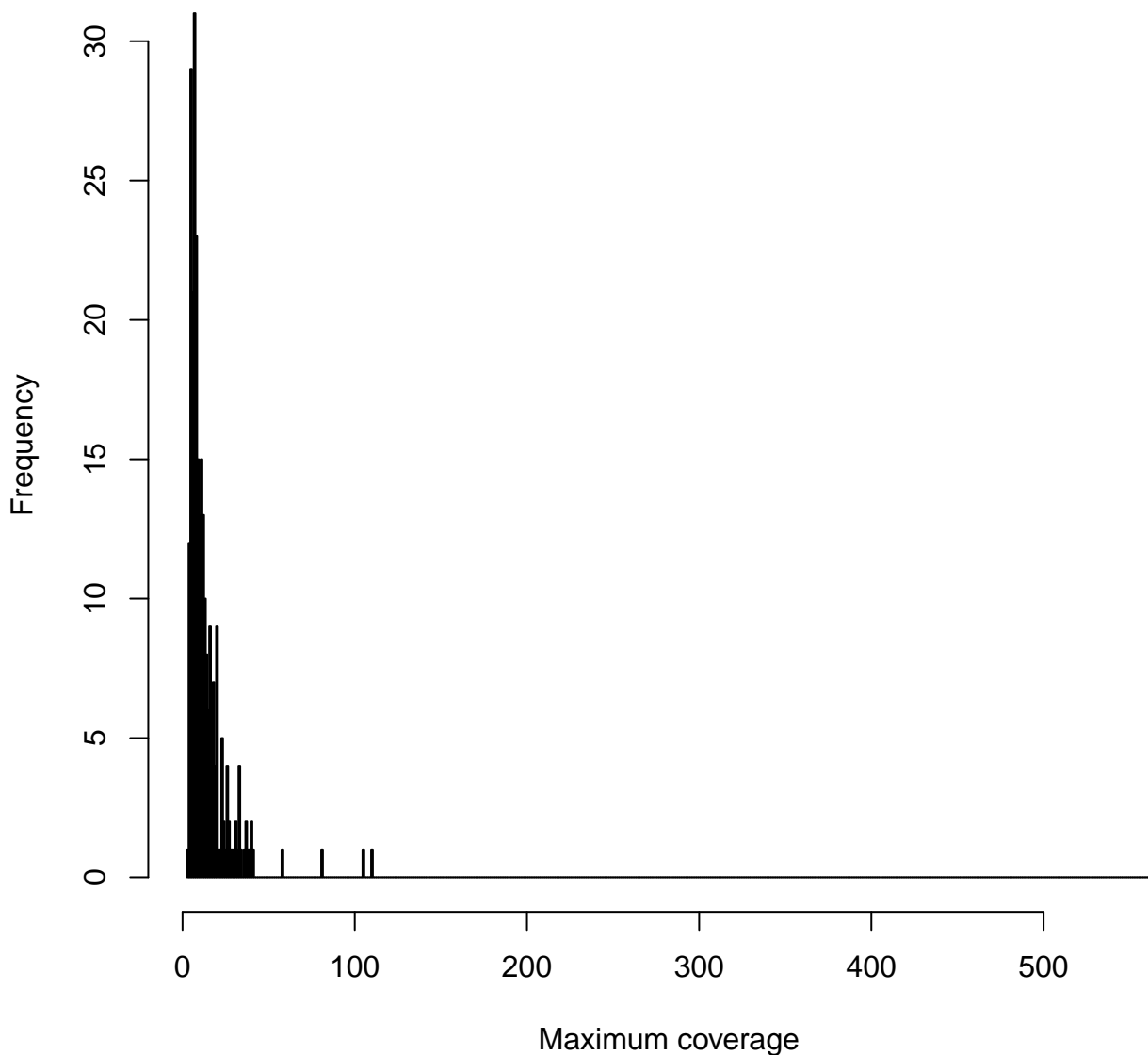
**jx in Intropolis but not "annotated" by knownGene:
92 samples; n jx: 291**



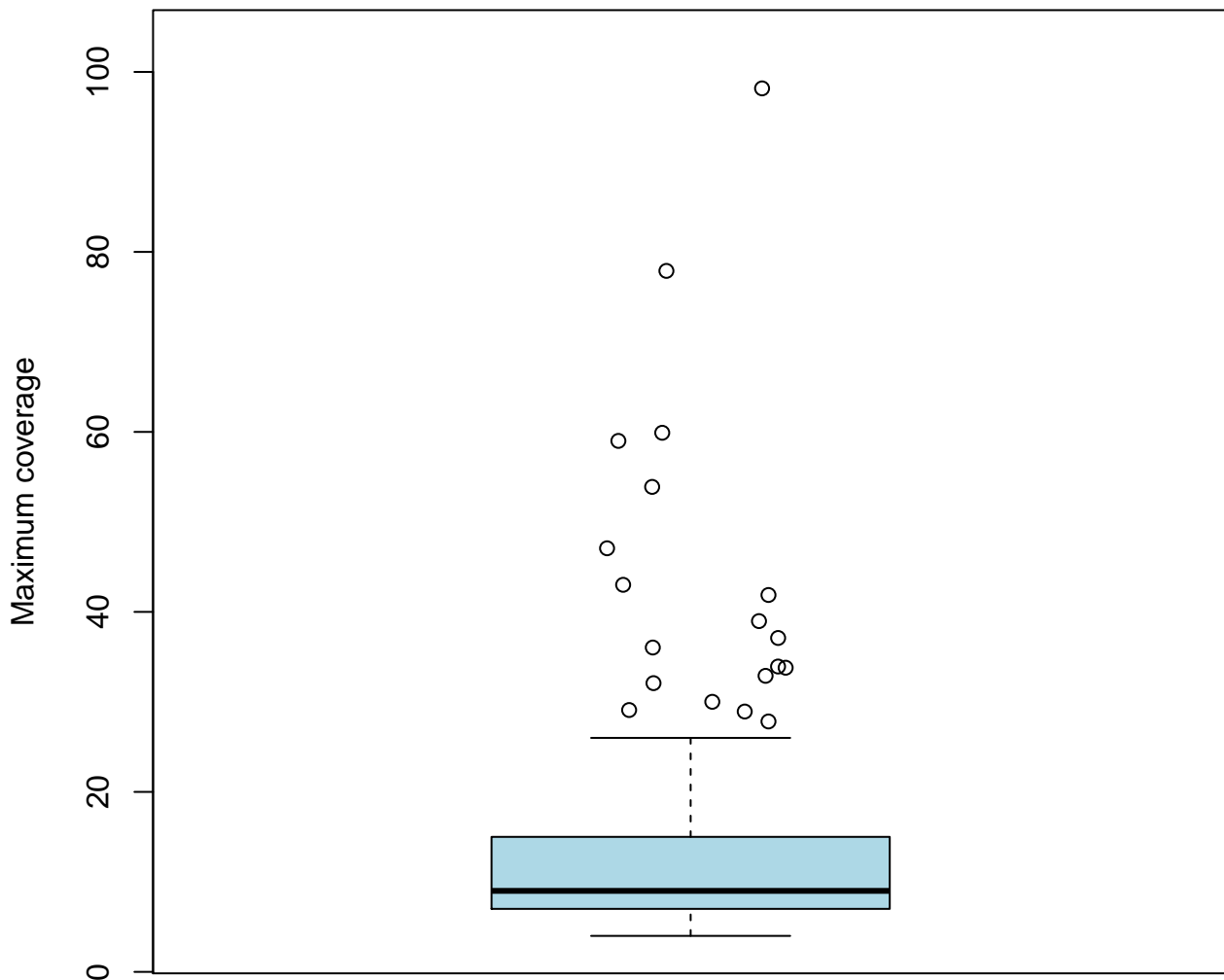
**jx in Intropolis but not "annotated" by knownGene:
93 samples; n jx: 271**



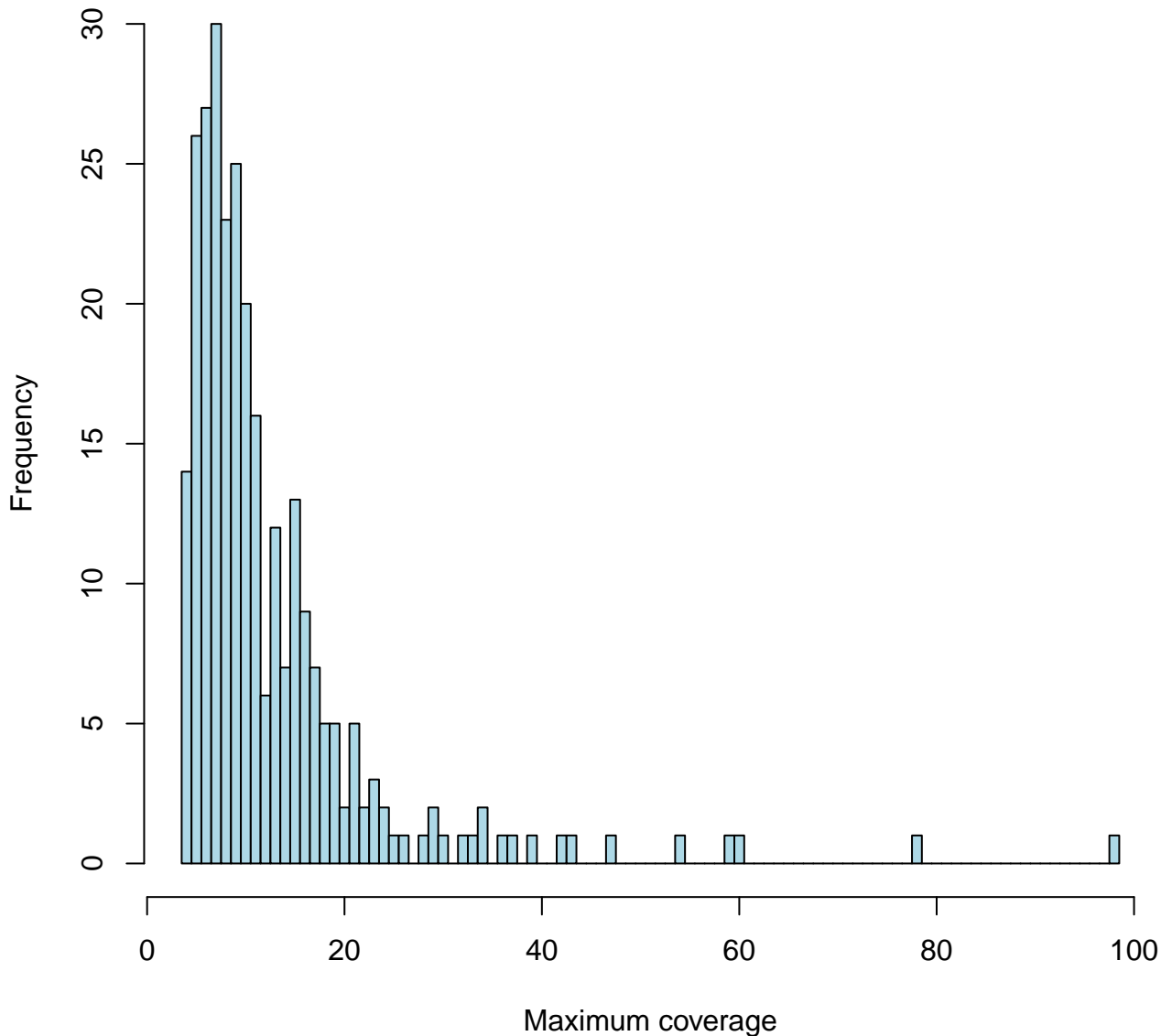
**jx in Intropolis but not "annotated" by knownGene:
93 samples; n jx: 271**



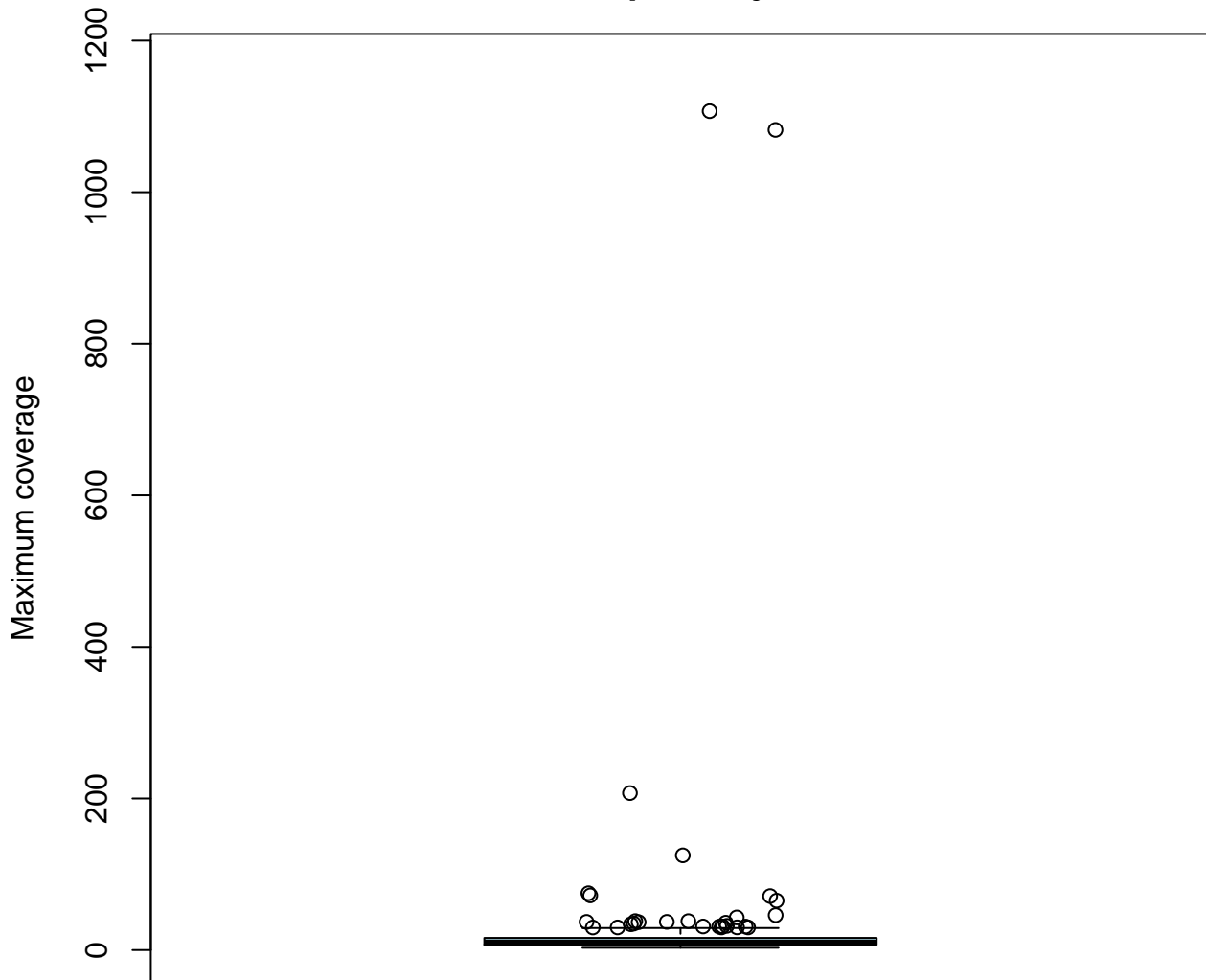
**jx in Intropolis but not "annotated" by knownGene:
94 samples; n jx: 280**



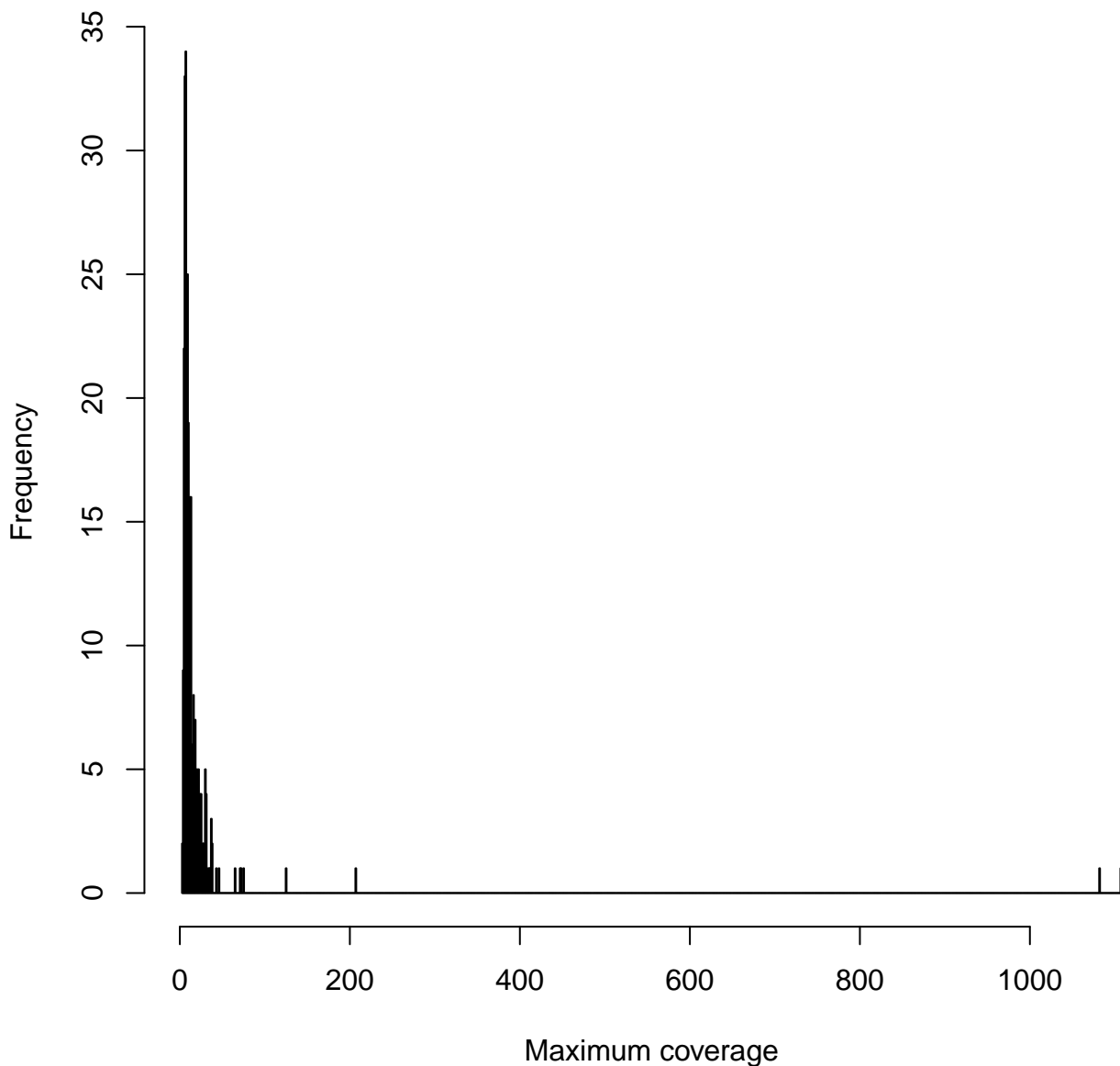
**jx in Intropolis but not "annotated" by knownGene:
94 samples; n jx: 280**



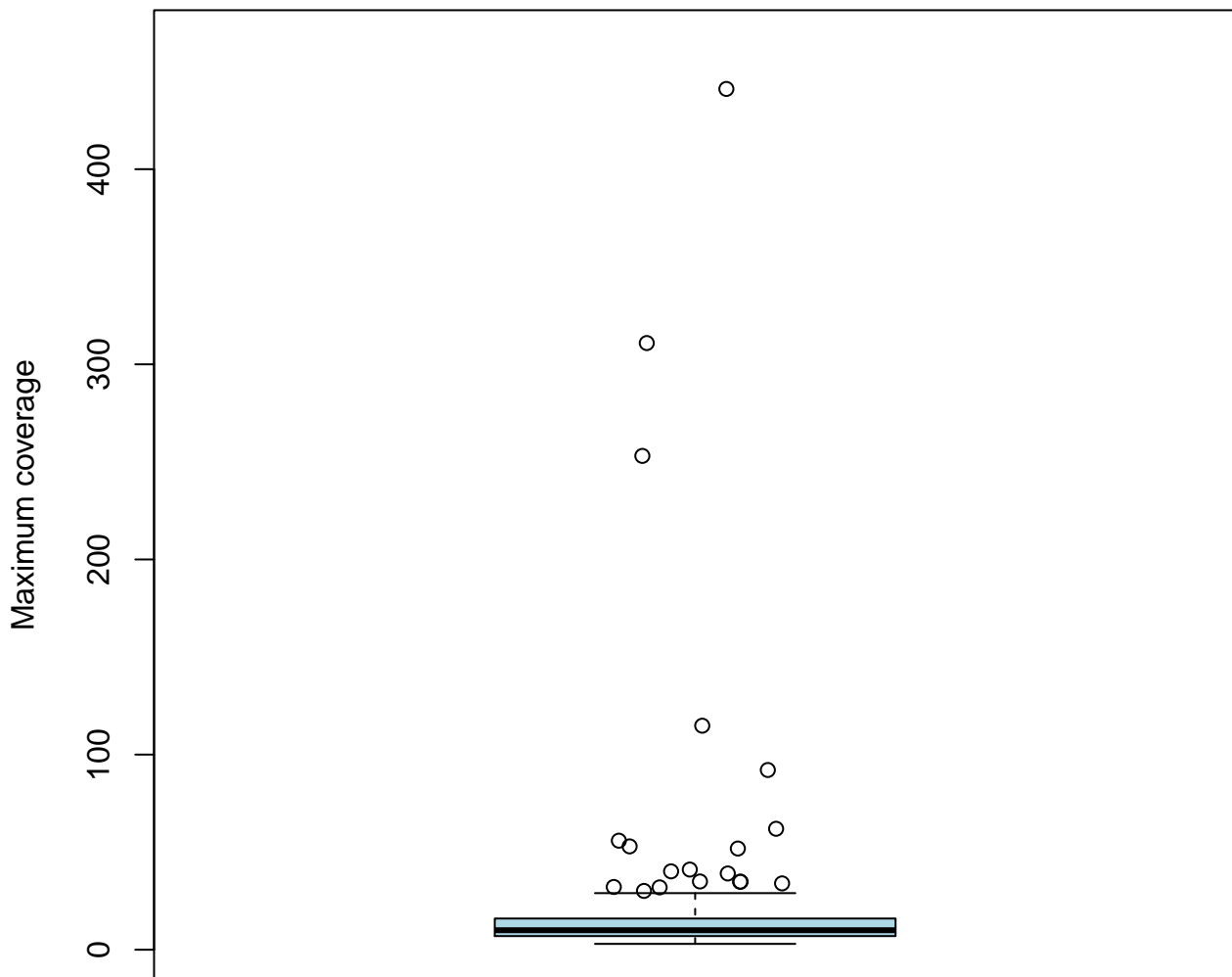
**jx in Intropolis but not "annotated" by knownGene:
95 samples; n jx: 293**



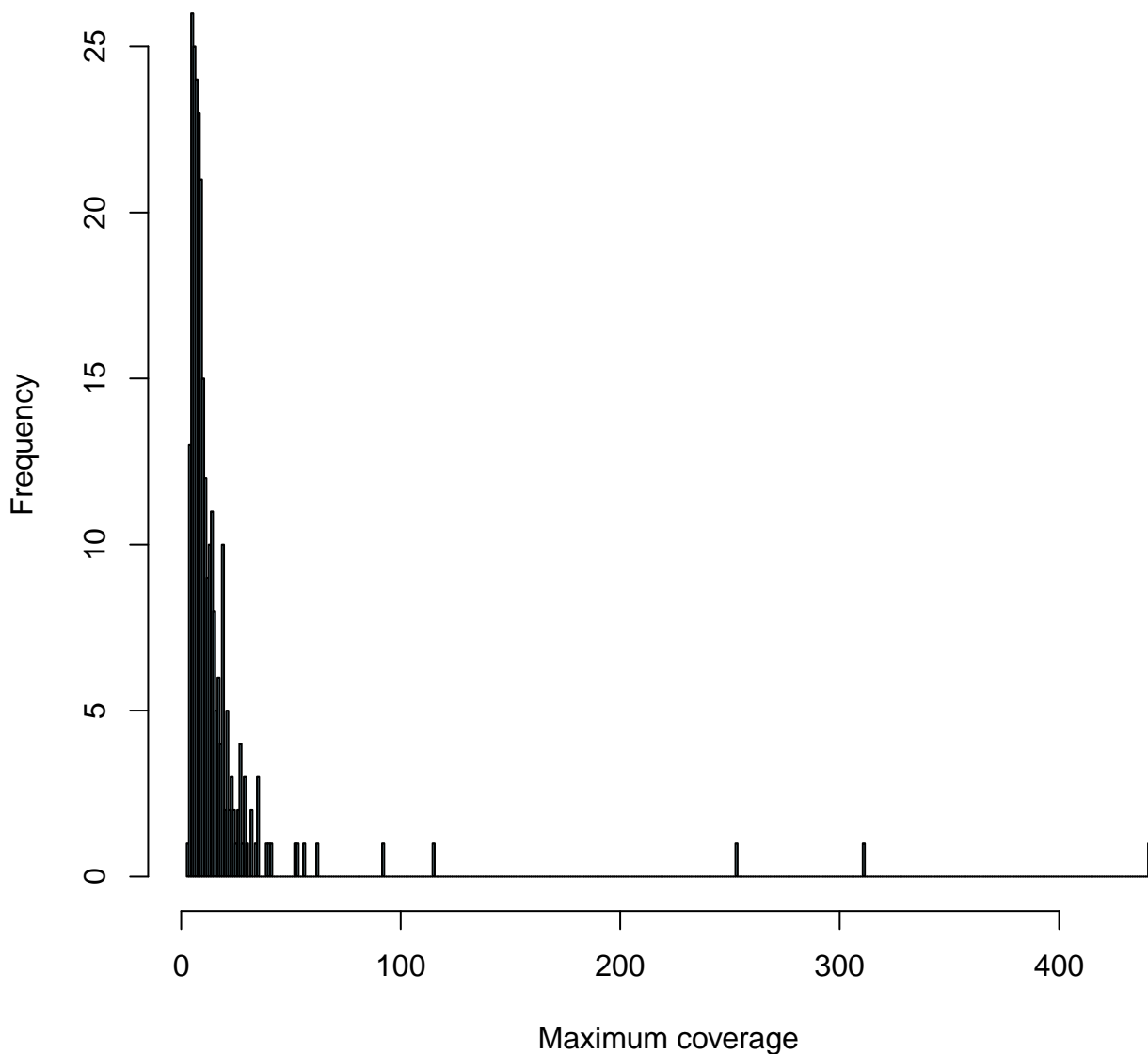
**jx in Intropolis but not "annotated" by knownGene:
95 samples; n jx: 293**



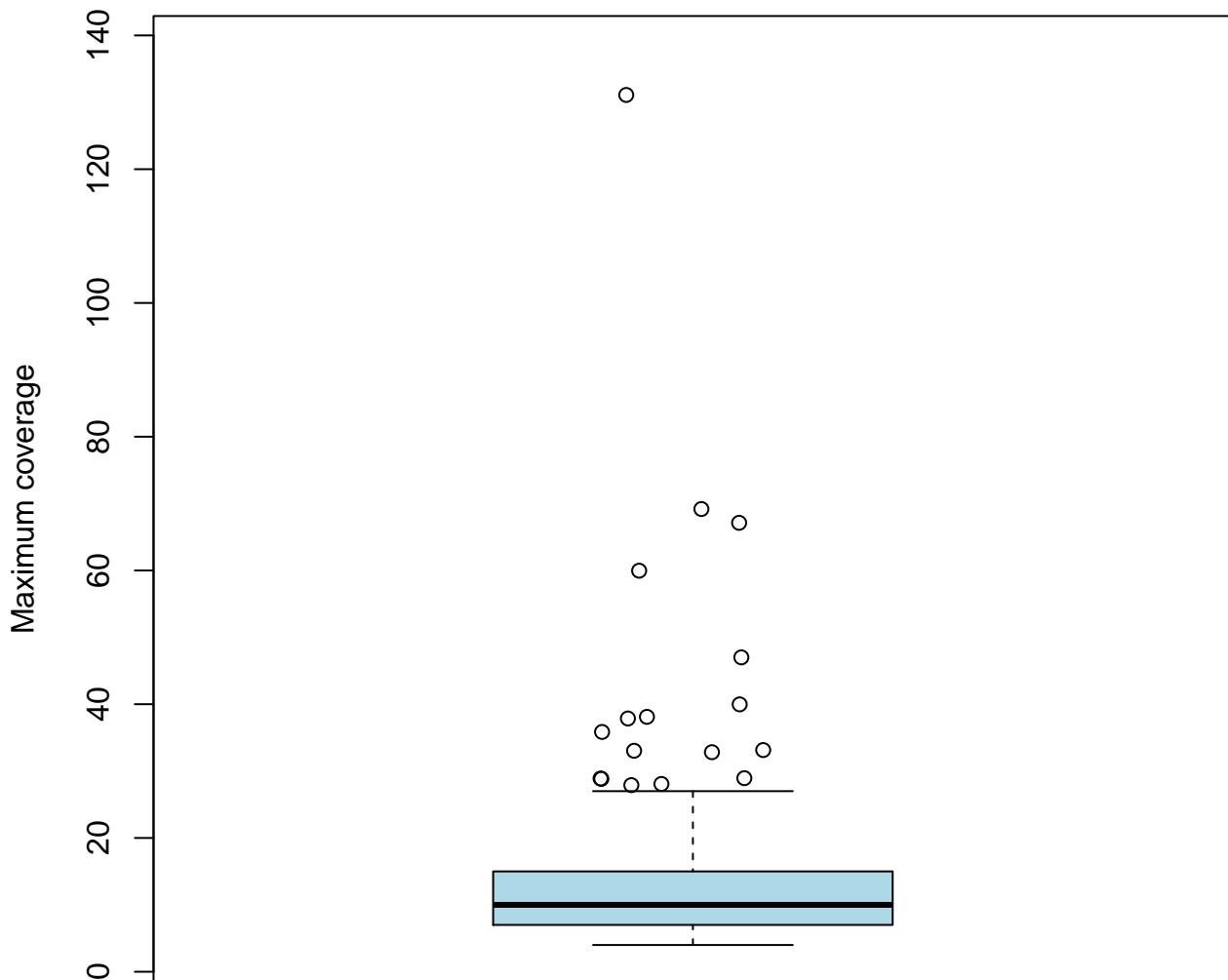
**jx in Intropolis but not "annotated" by knownGene:
96 samples; n jx: 267**



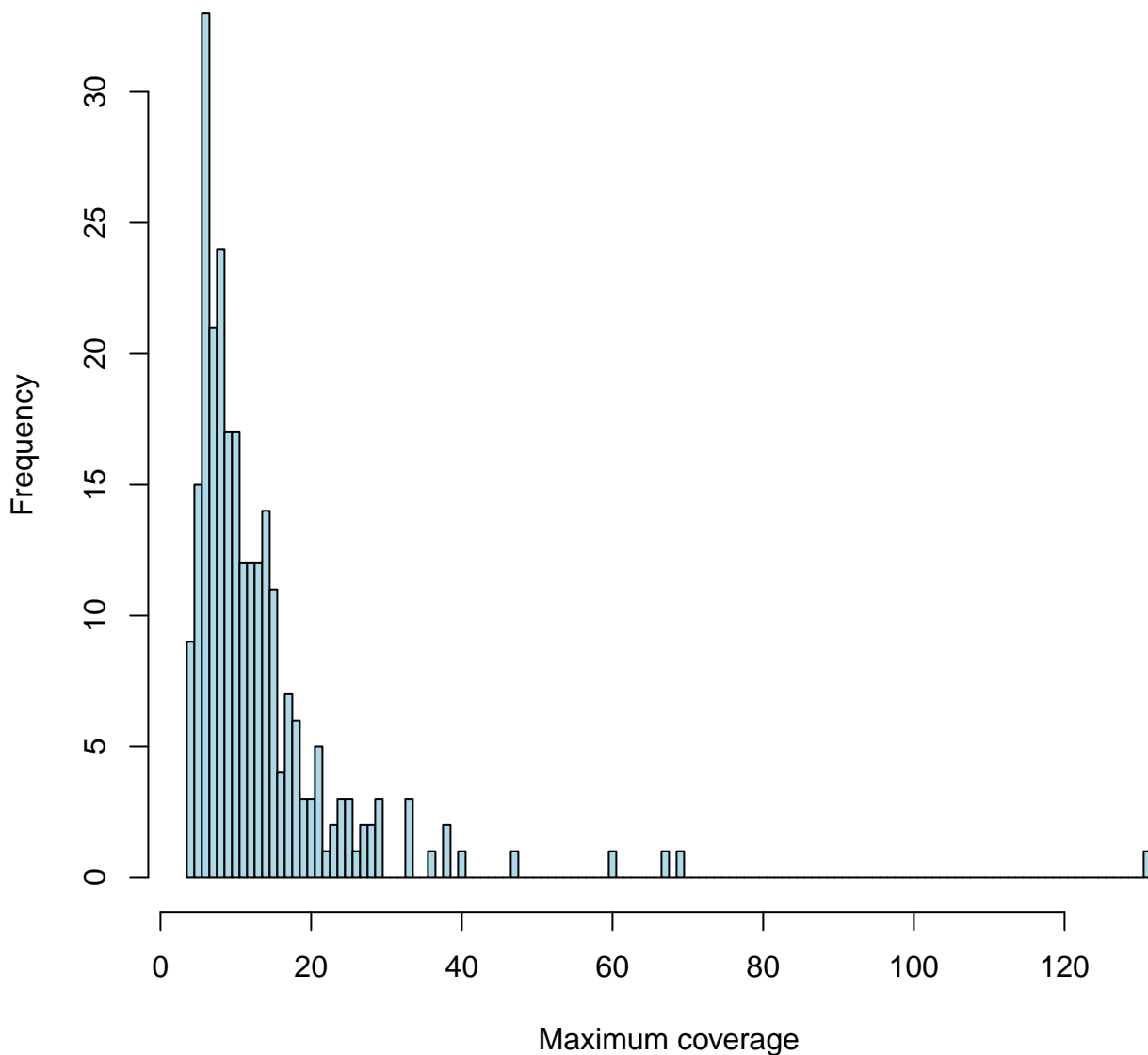
**jx in Intropolis but not "annotated" by knownGene:
96 samples; n jx: 267**



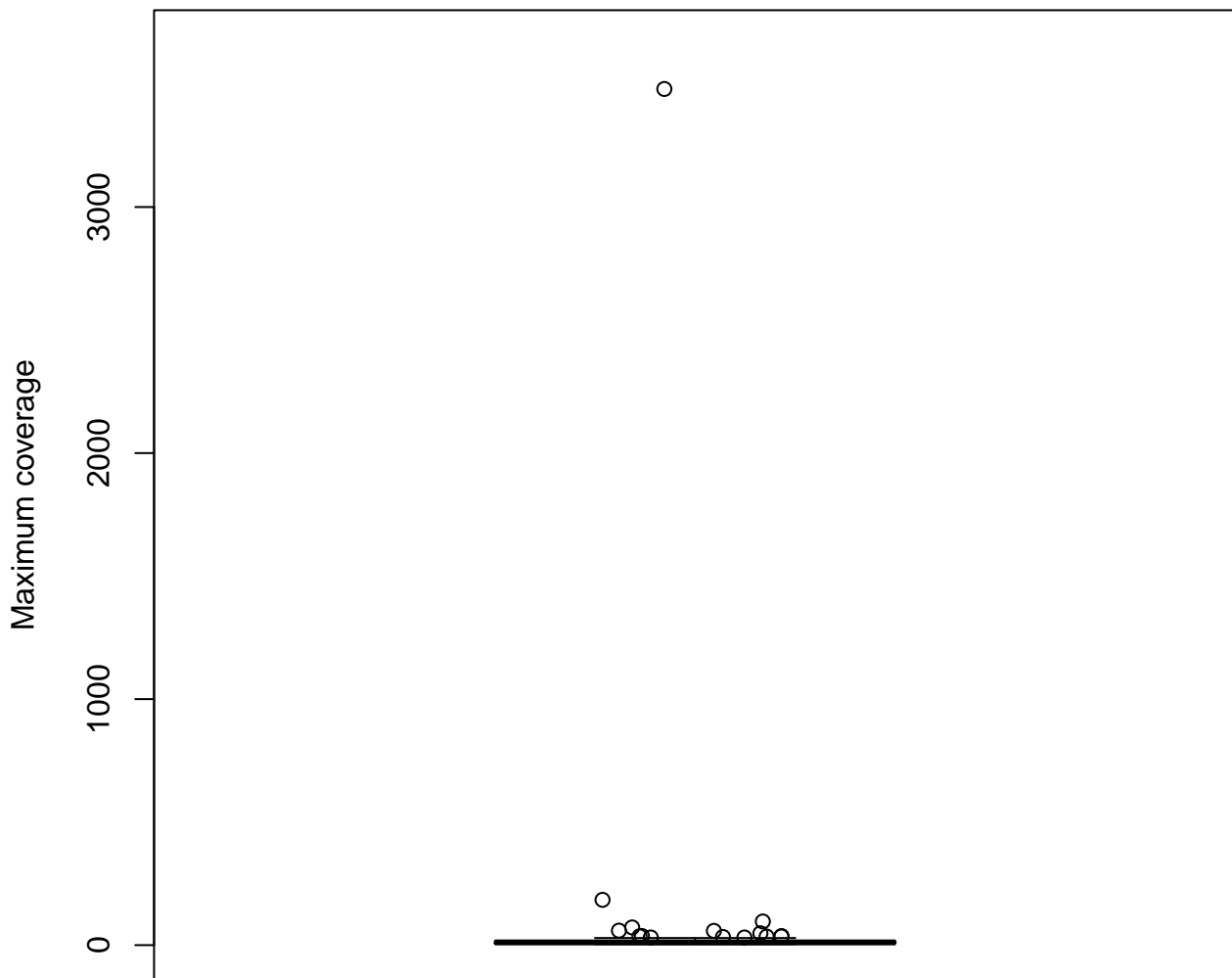
**jx in Intropolis but not "annotated" by knownGene:
97 samples; n jx: 254**



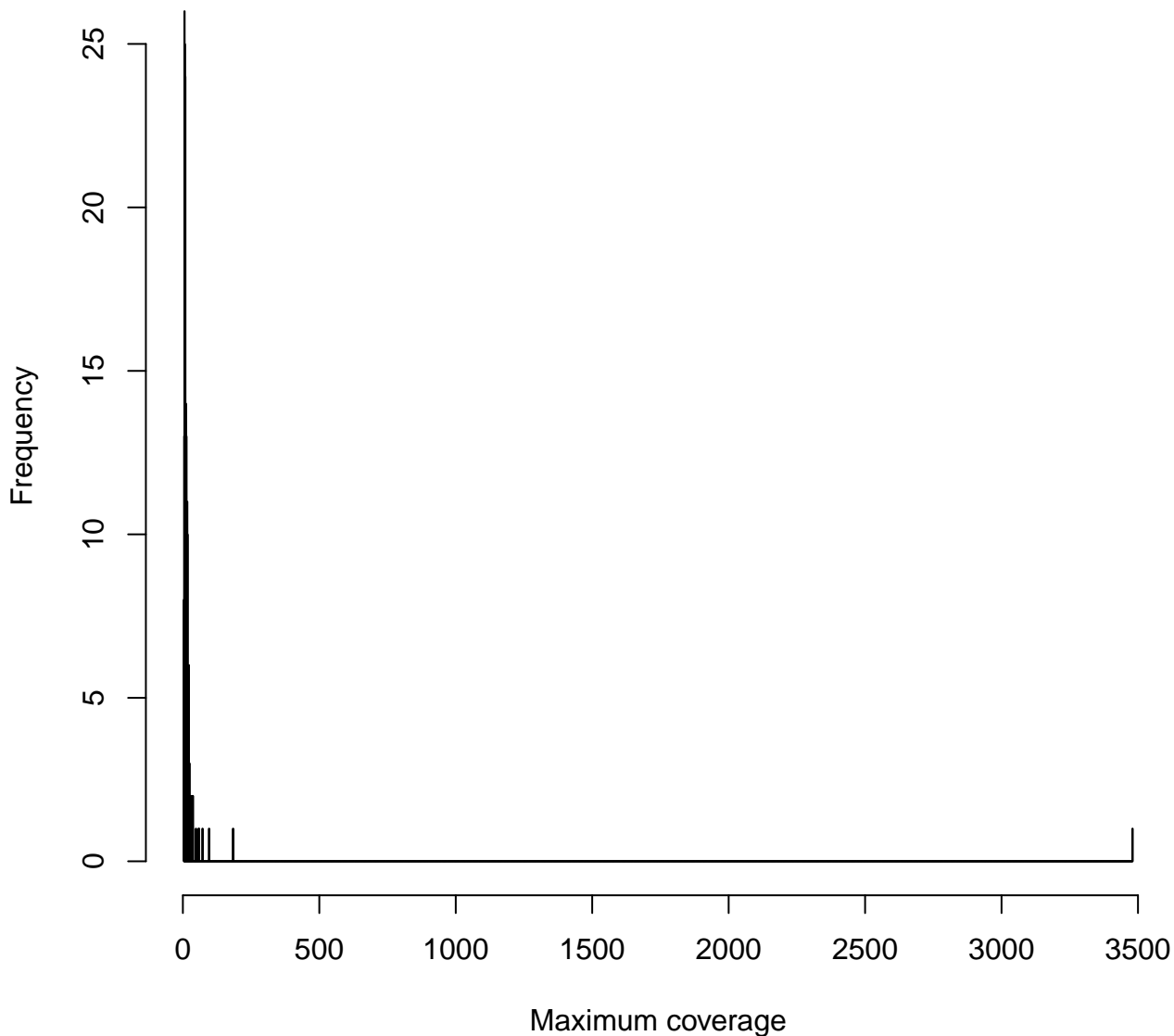
**jx in Intropolis but not "annotated" by knownGene:
97 samples; n jx: 254**



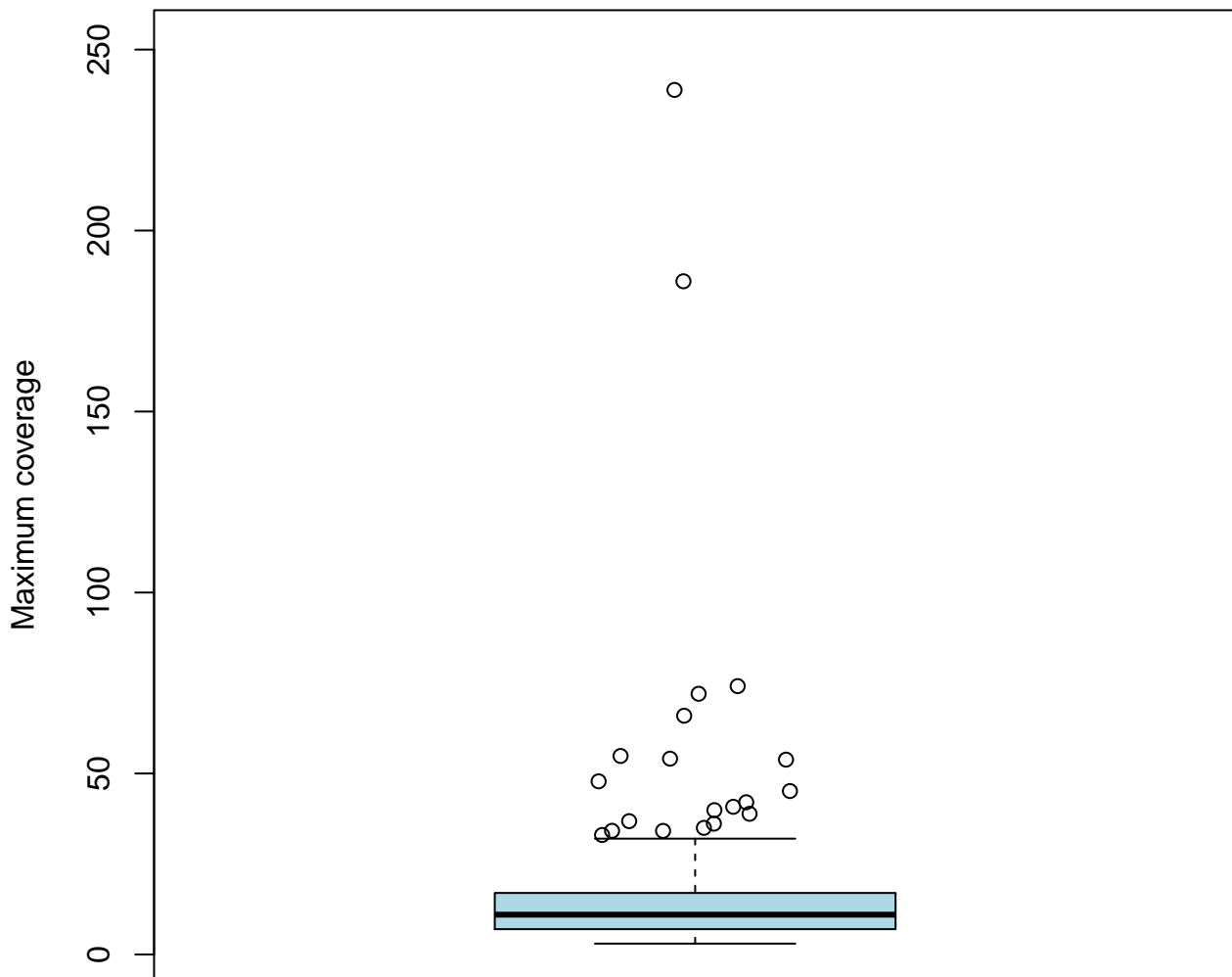
**jx in Intropolis but not "annotated" by knownGene:
98 samples; n jx: 247**



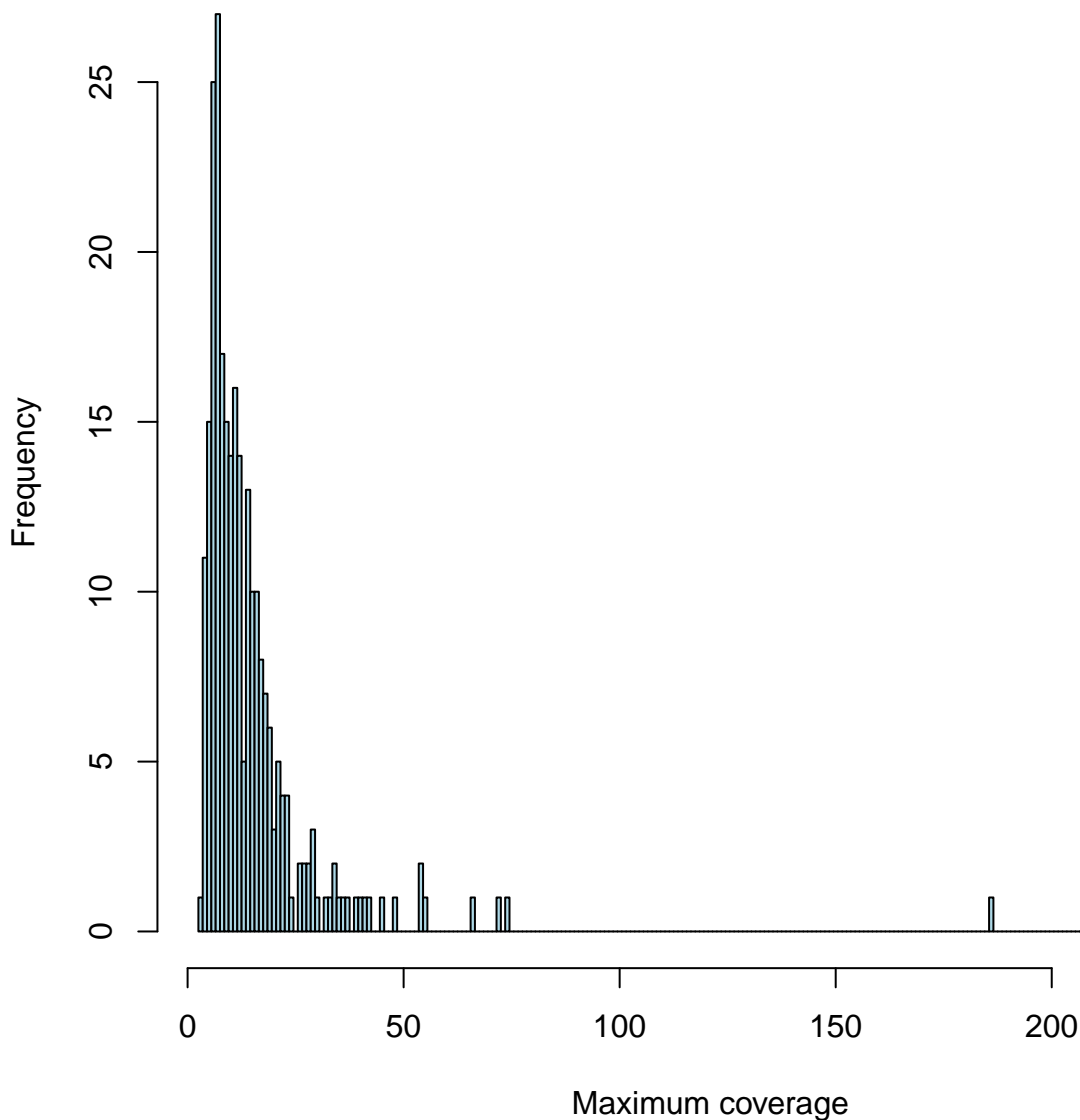
**jx in Intropolis but not "annotated" by knownGene:
98 samples; n jx: 247**



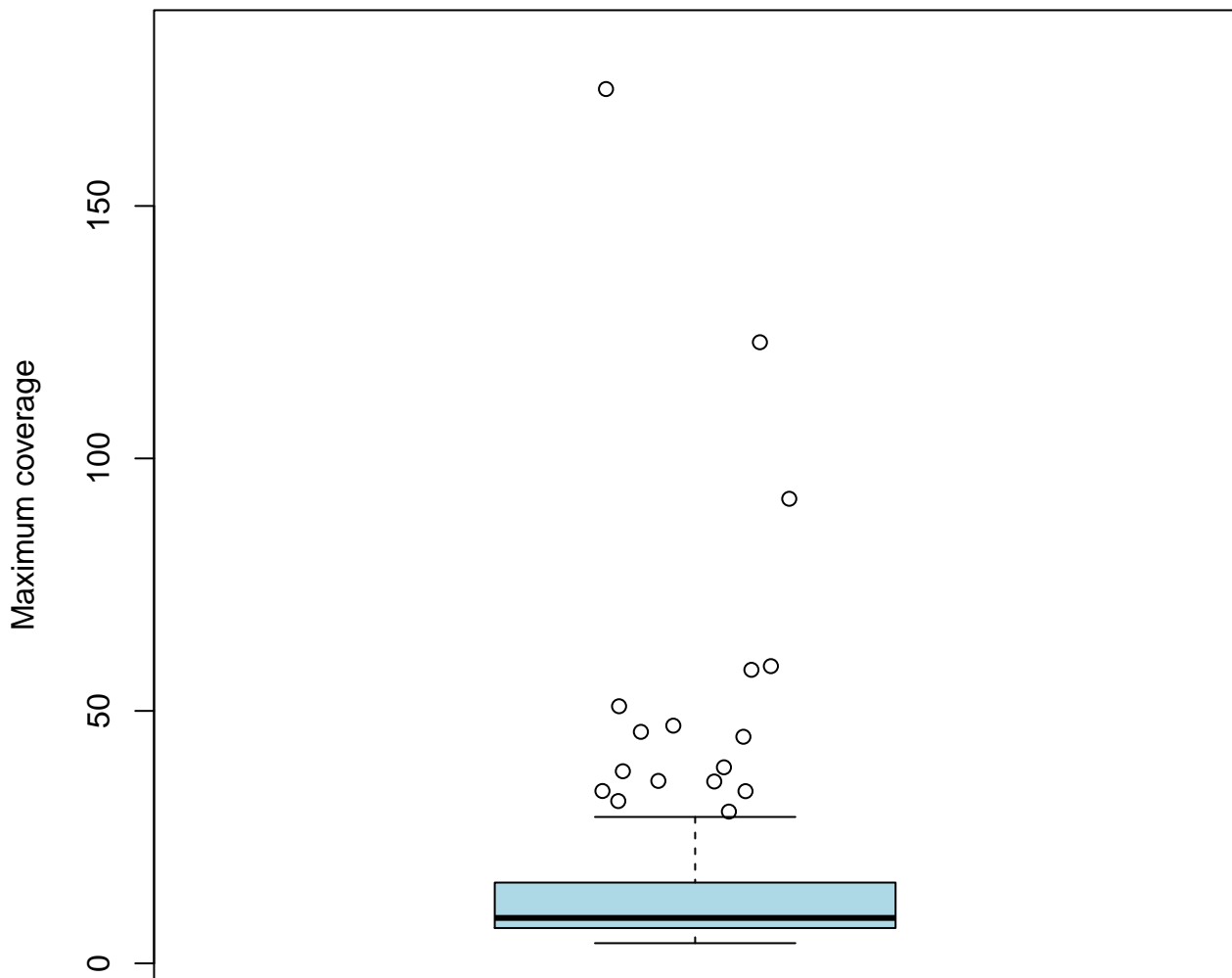
**jx in Intropolis but not "annotated" by knownGene:
99 samples; n jx: 262**



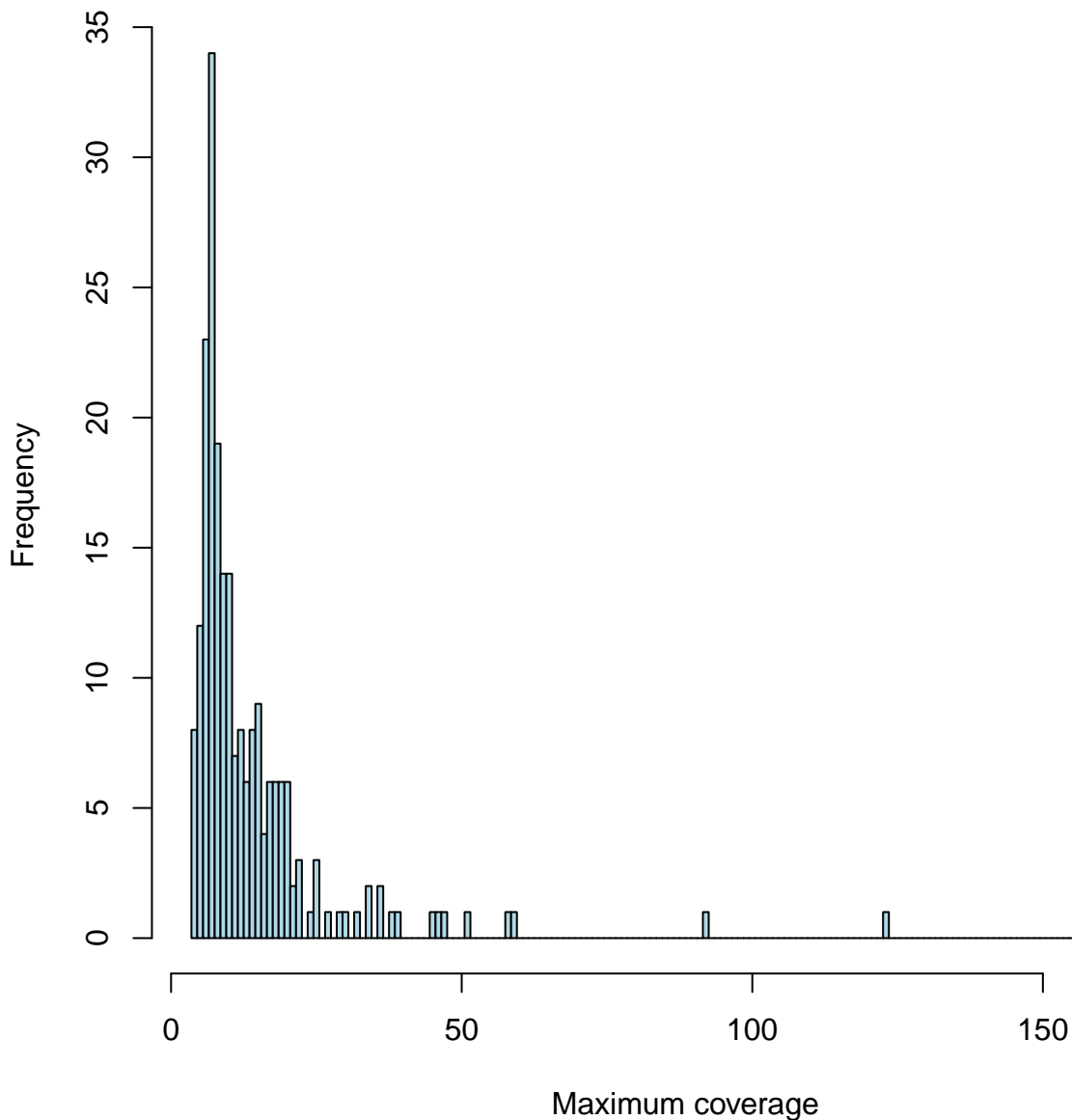
**jx in Intropolis but not "annotated" by knownGene:
99 samples; n jx: 262**



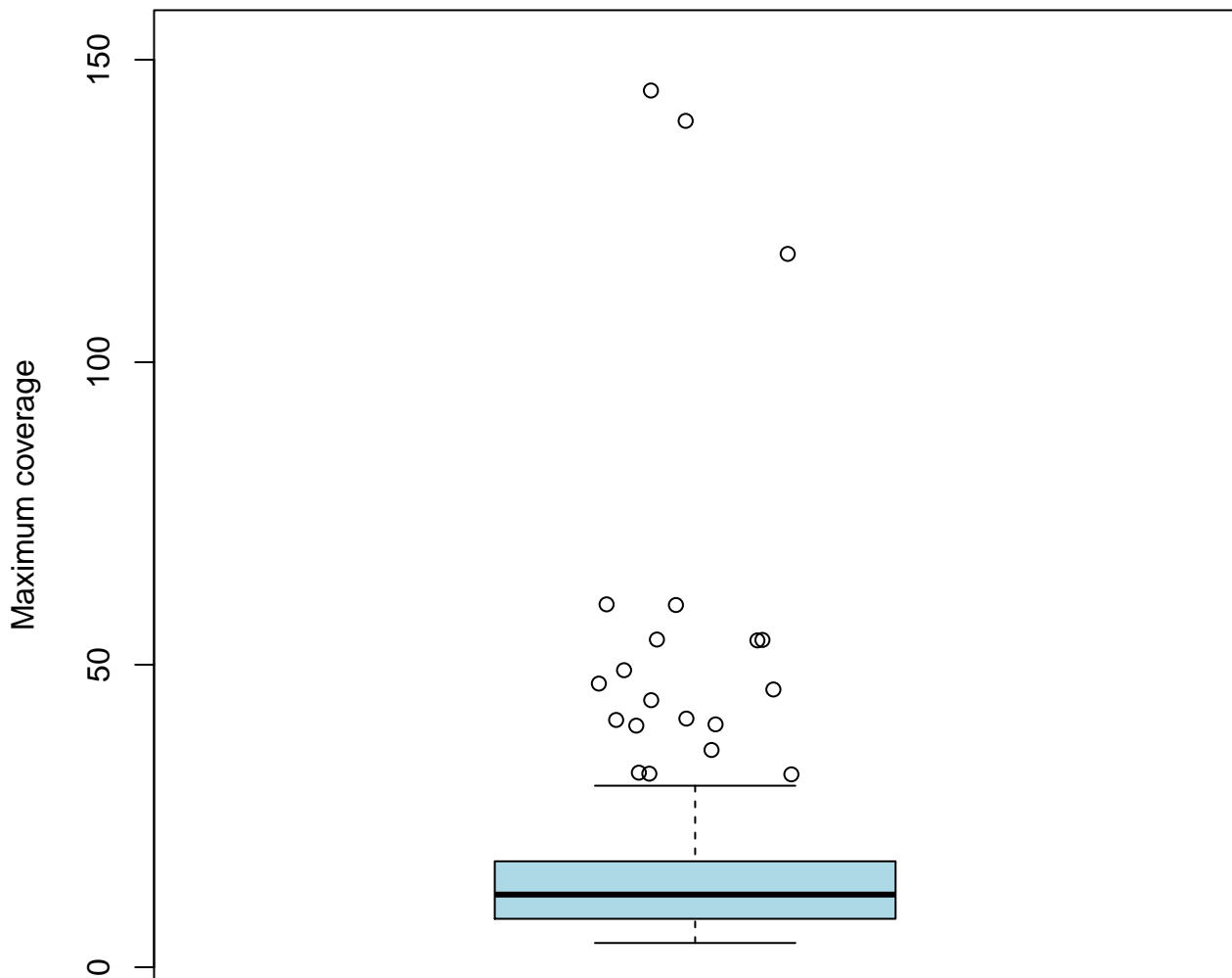
**jx in Intropolis but not "annotated" by knownGene:
100 samples; n jx: 218**



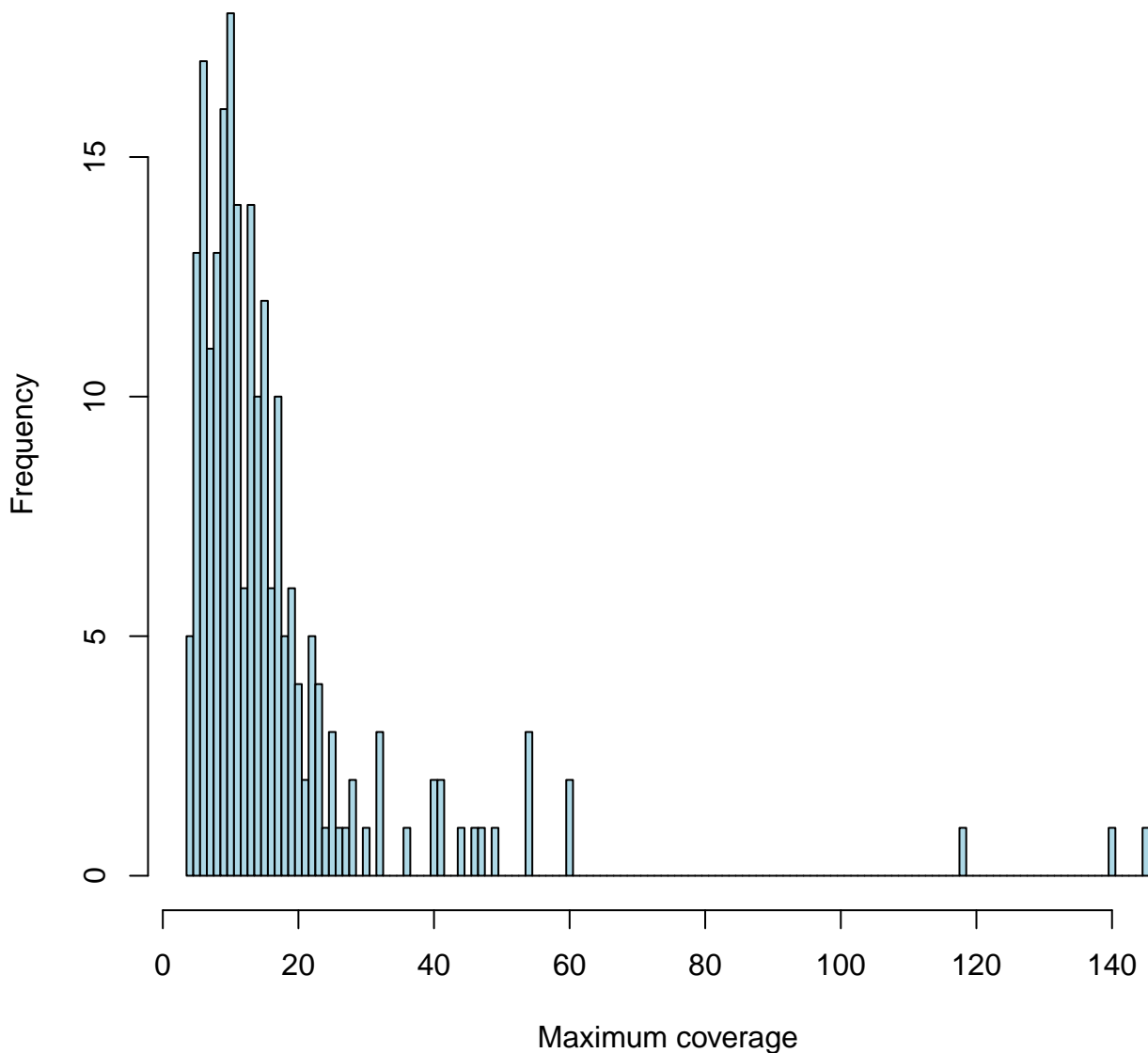
**jx in Intropolis but not "annotated" by knownGene:
100 samples; n jx: 218**



**jx in Intropolis but not "annotated" by knownGene:
101 samples; n jx: 220**



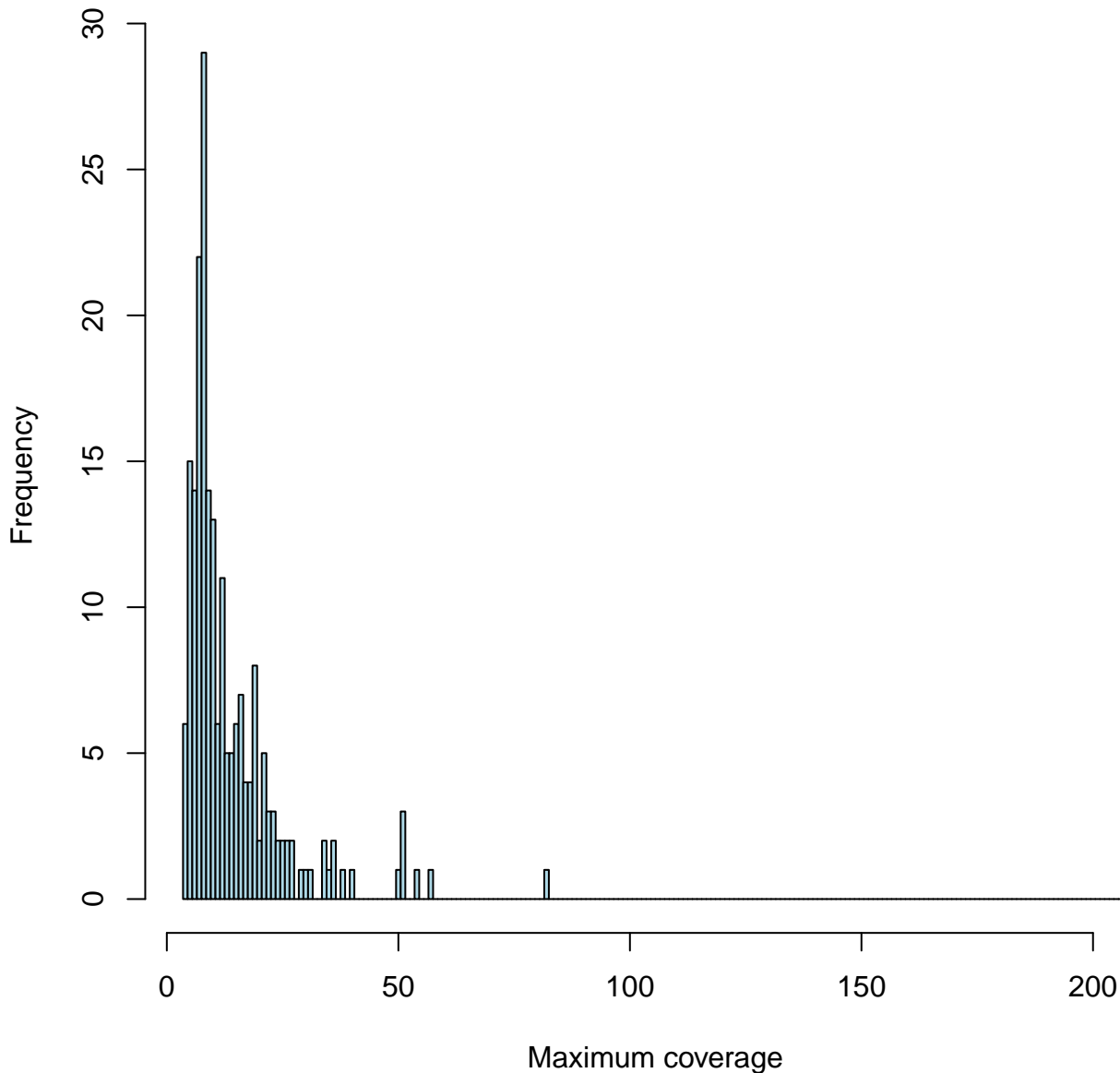
**jx in Intropolis but not "annotated" by knownGene:
101 samples; n jx: 220**



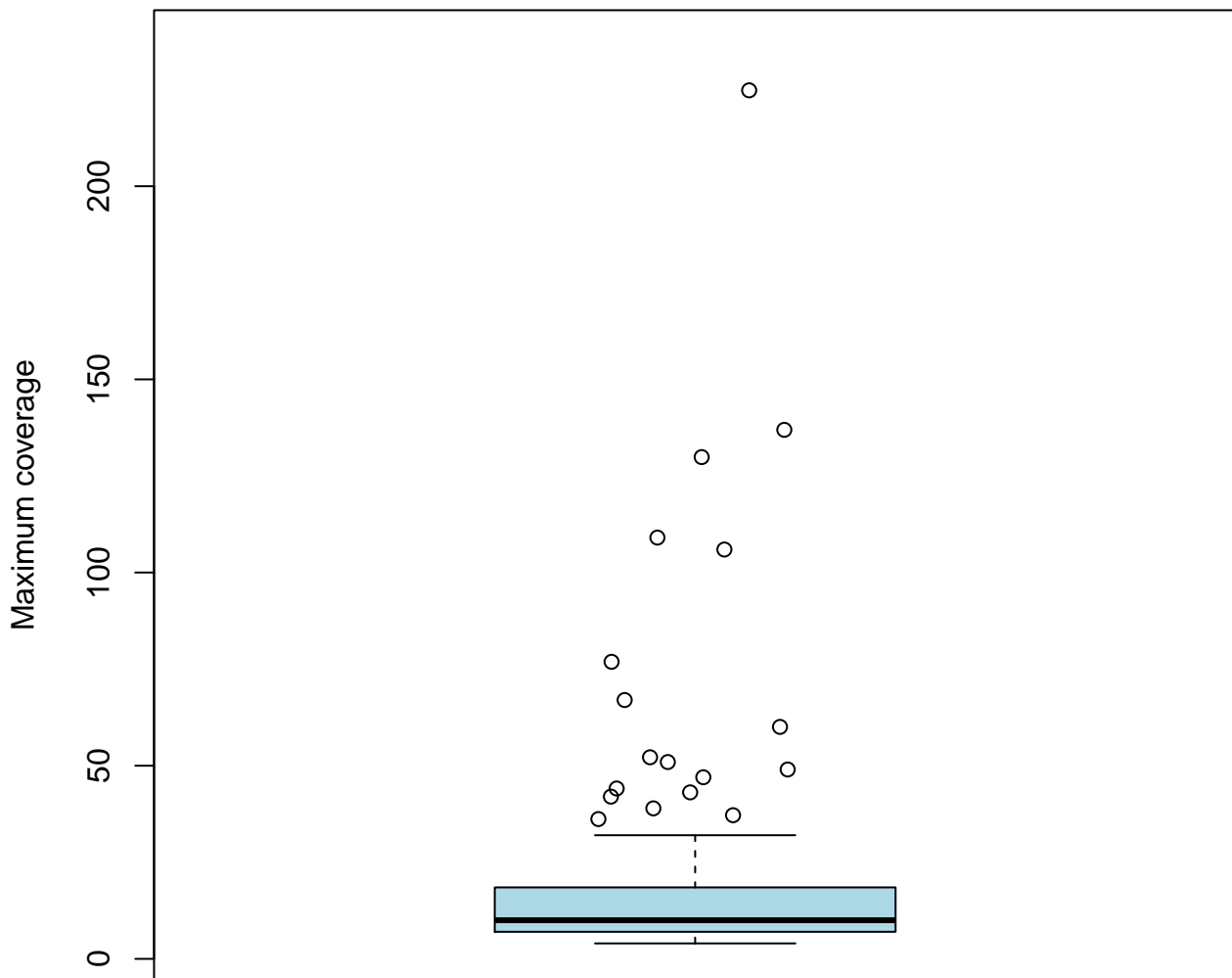
102 samples; n jx: 208



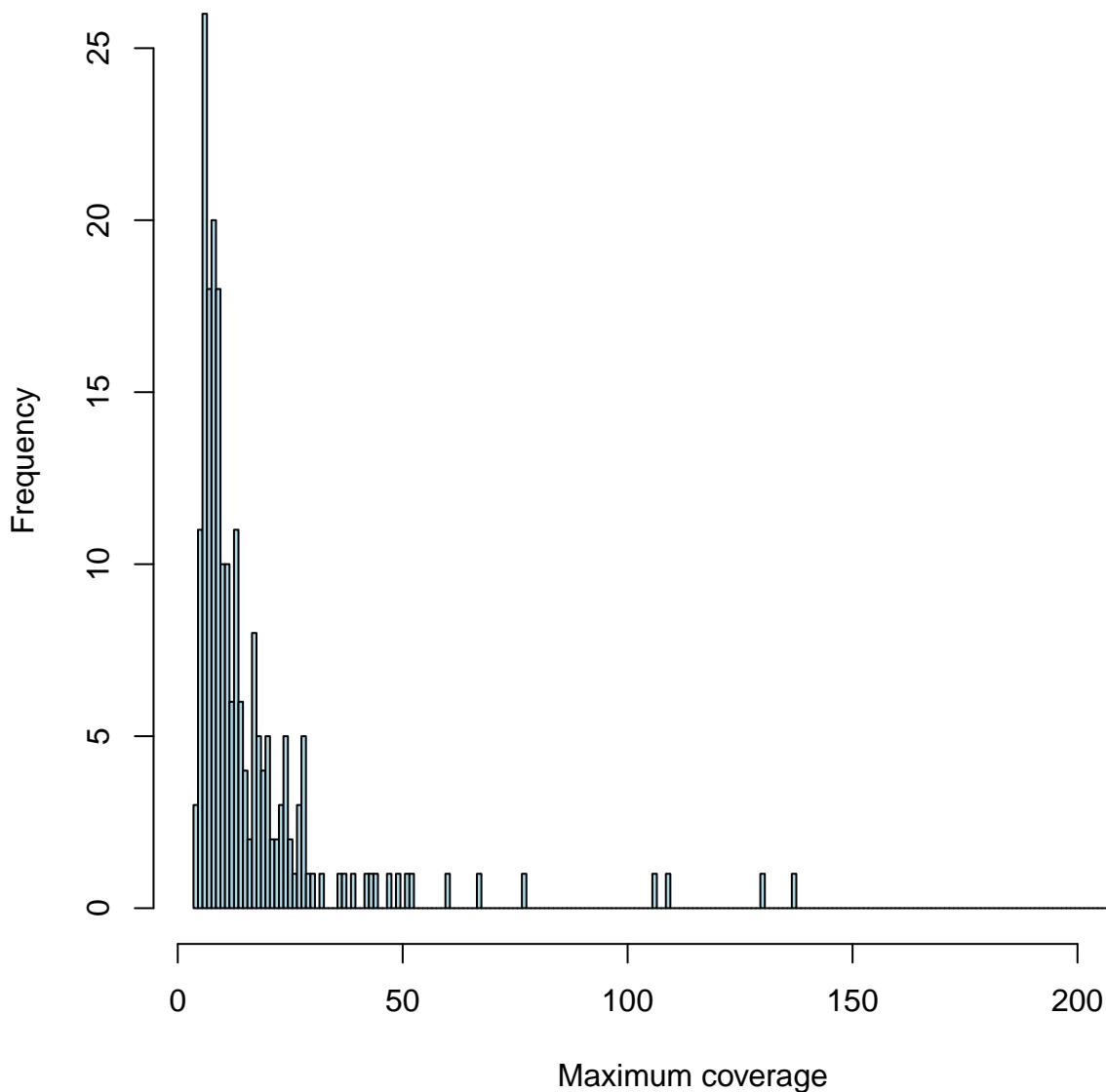
**jx in Intropolis but not "annotated" by knownGene:
102 samples; n jx: 208**



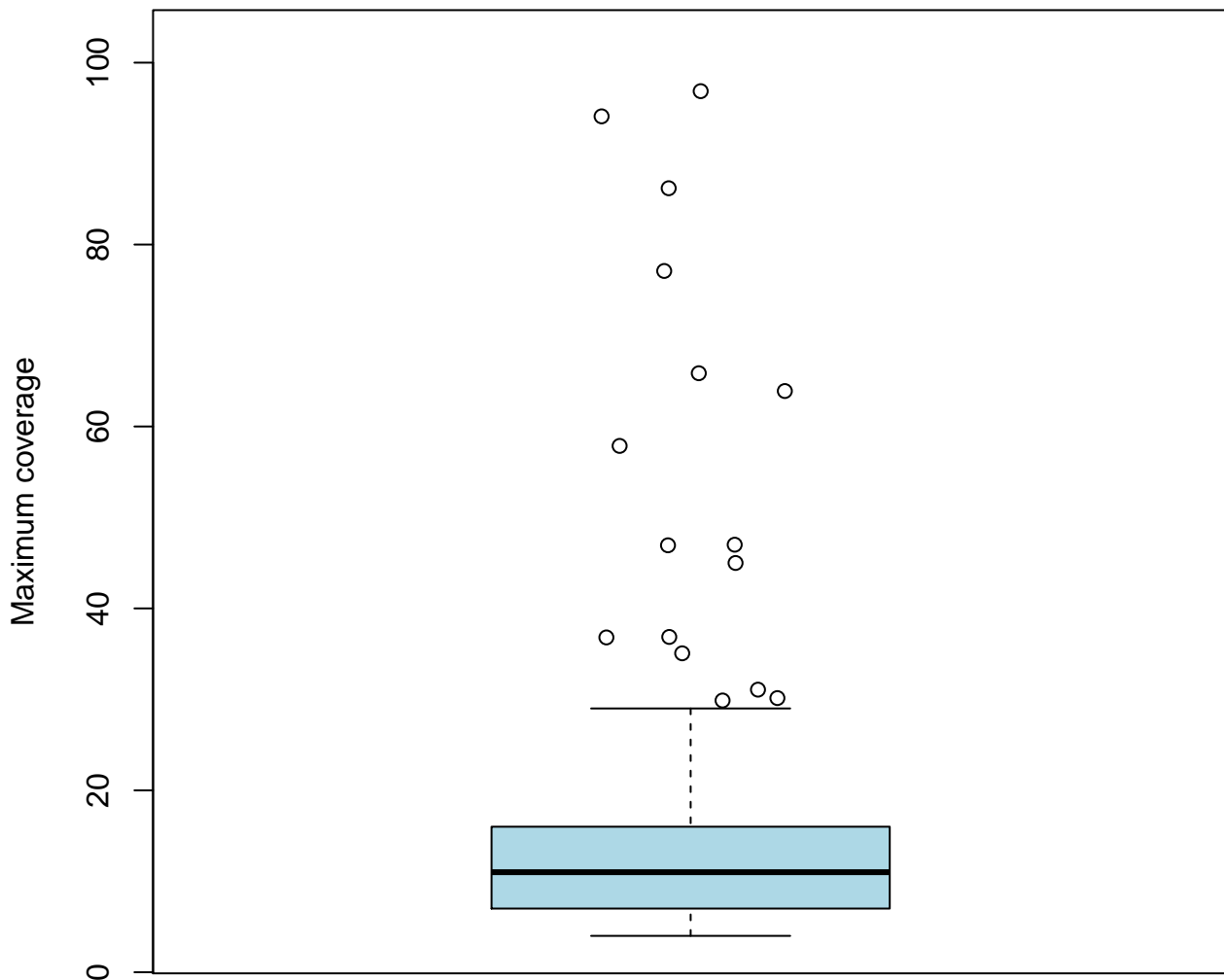
**jx in Intropolis but not "annotated" by knownGene:
103 samples; n jx: 211**



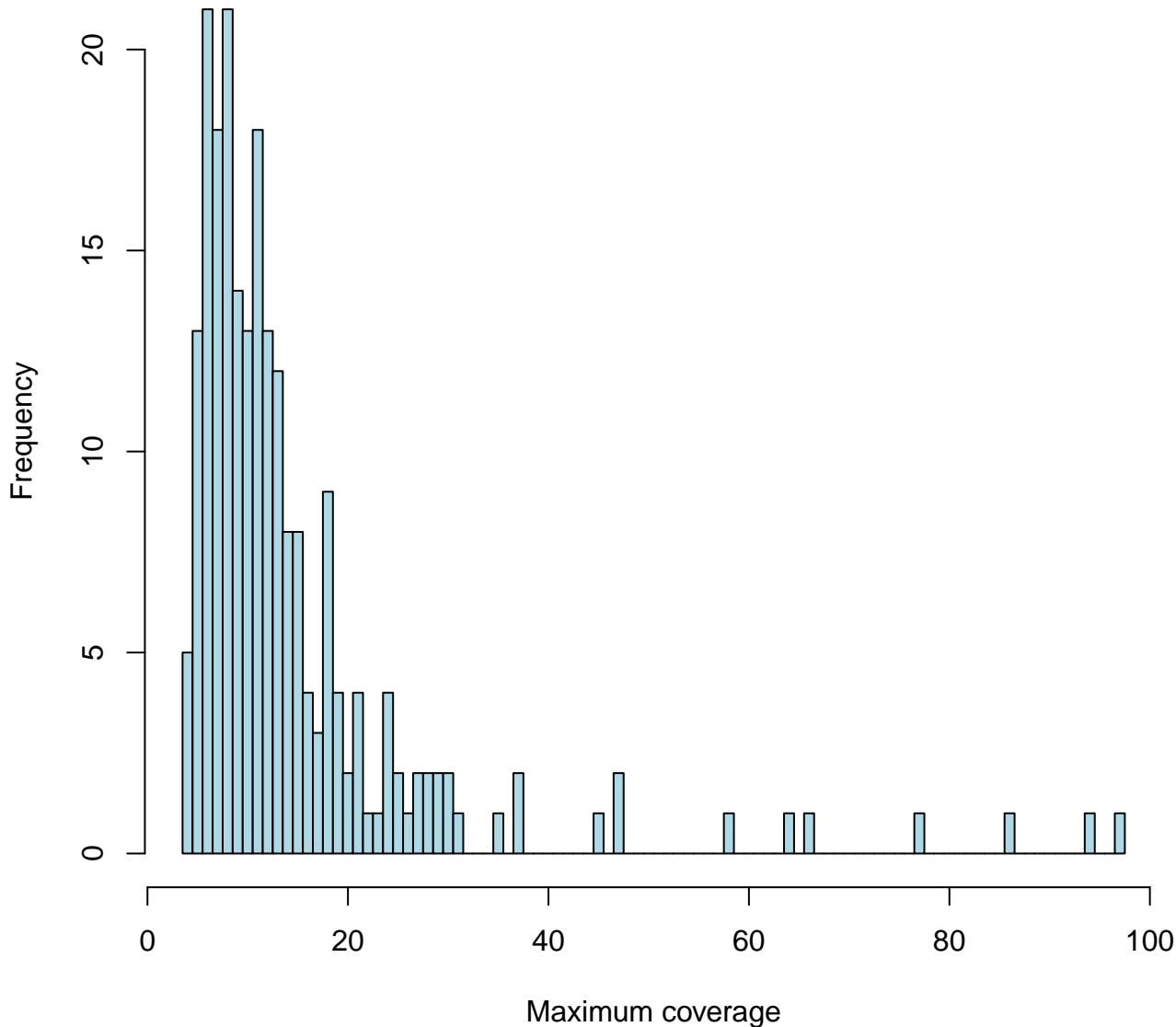
**jx in Intropolis but not "annotated" by knownGene:
103 samples; n jx: 211**



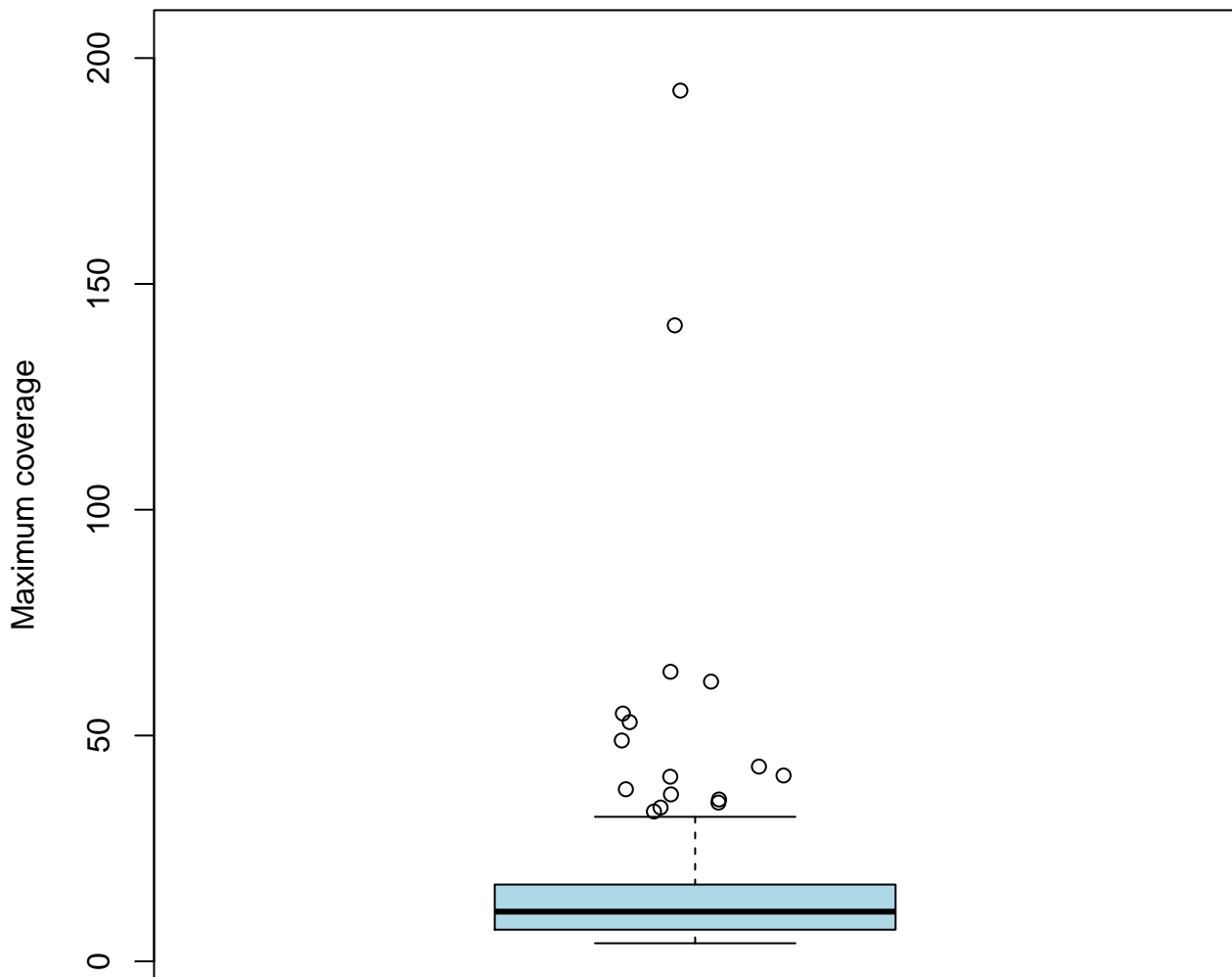
**jx in Intropolis but not "annotated" by knownGene:
104 samples; n jx: 221**



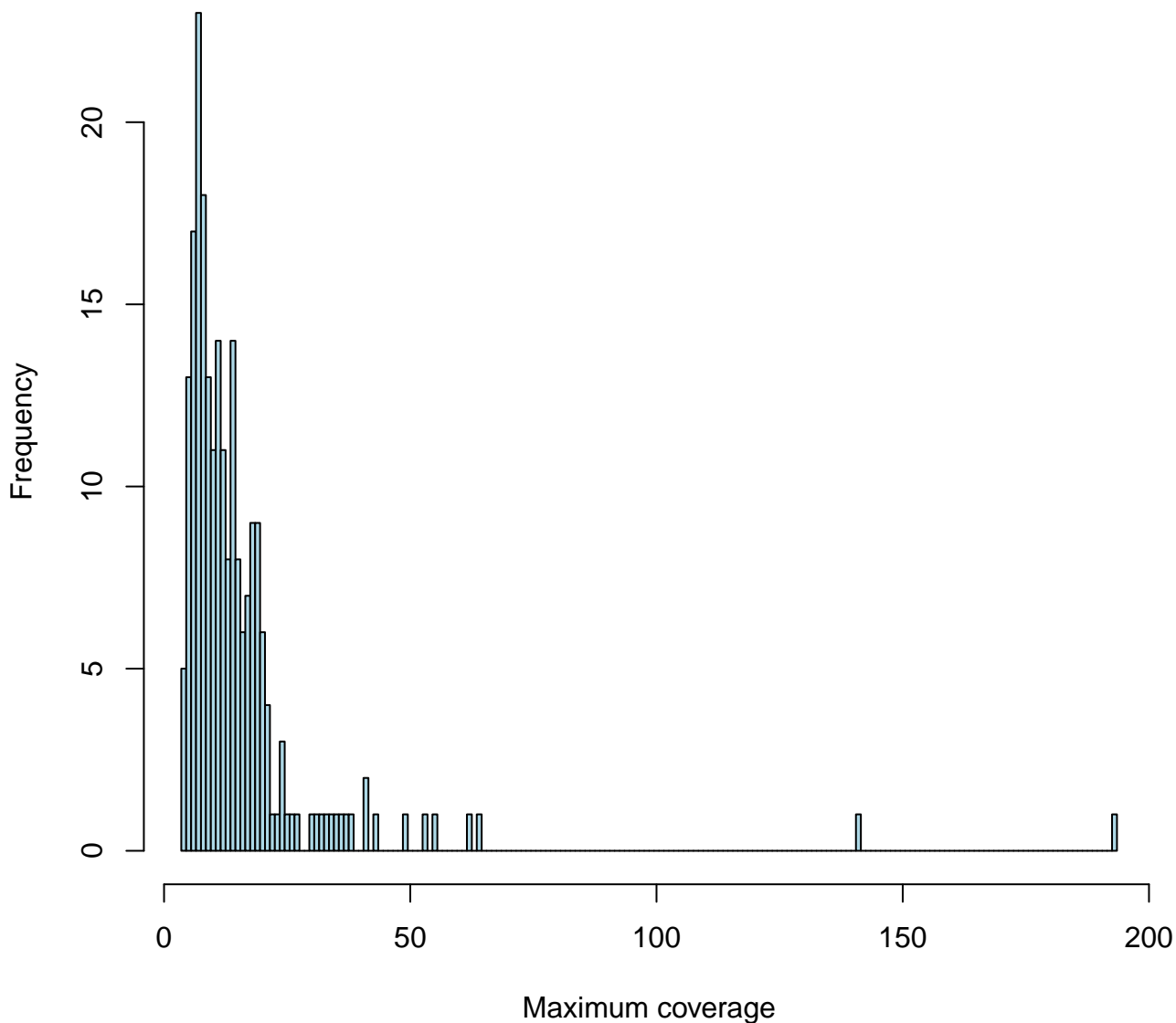
**jx in Intropolis but not "annotated" by knownGene:
104 samples; n jx: 221**



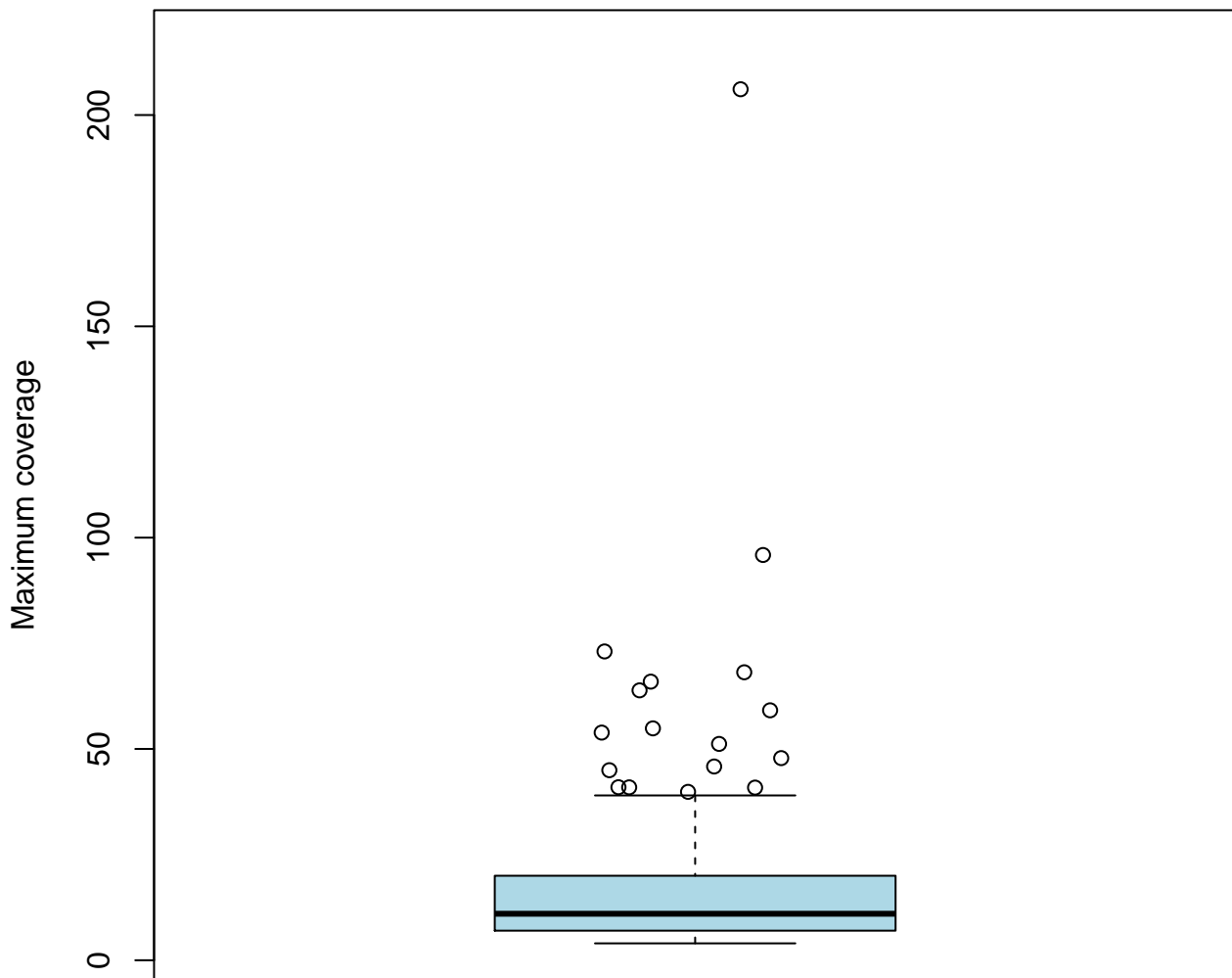
**jx in Intropolis but not "annotated" by knownGene:
105 samples; n jx: 223**



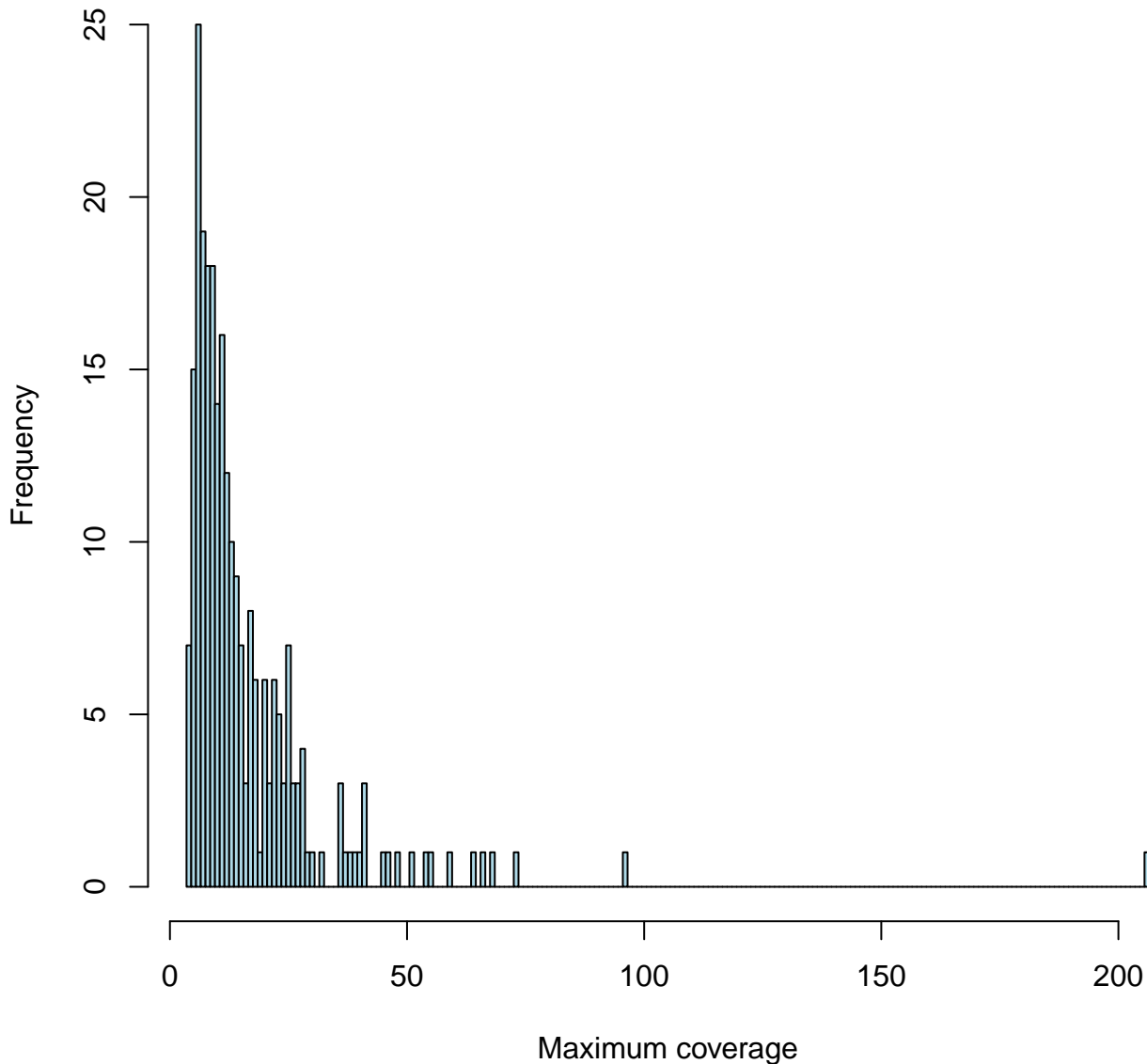
**jx in Intropolis but not "annotated" by knownGene:
105 samples; n jx: 223**



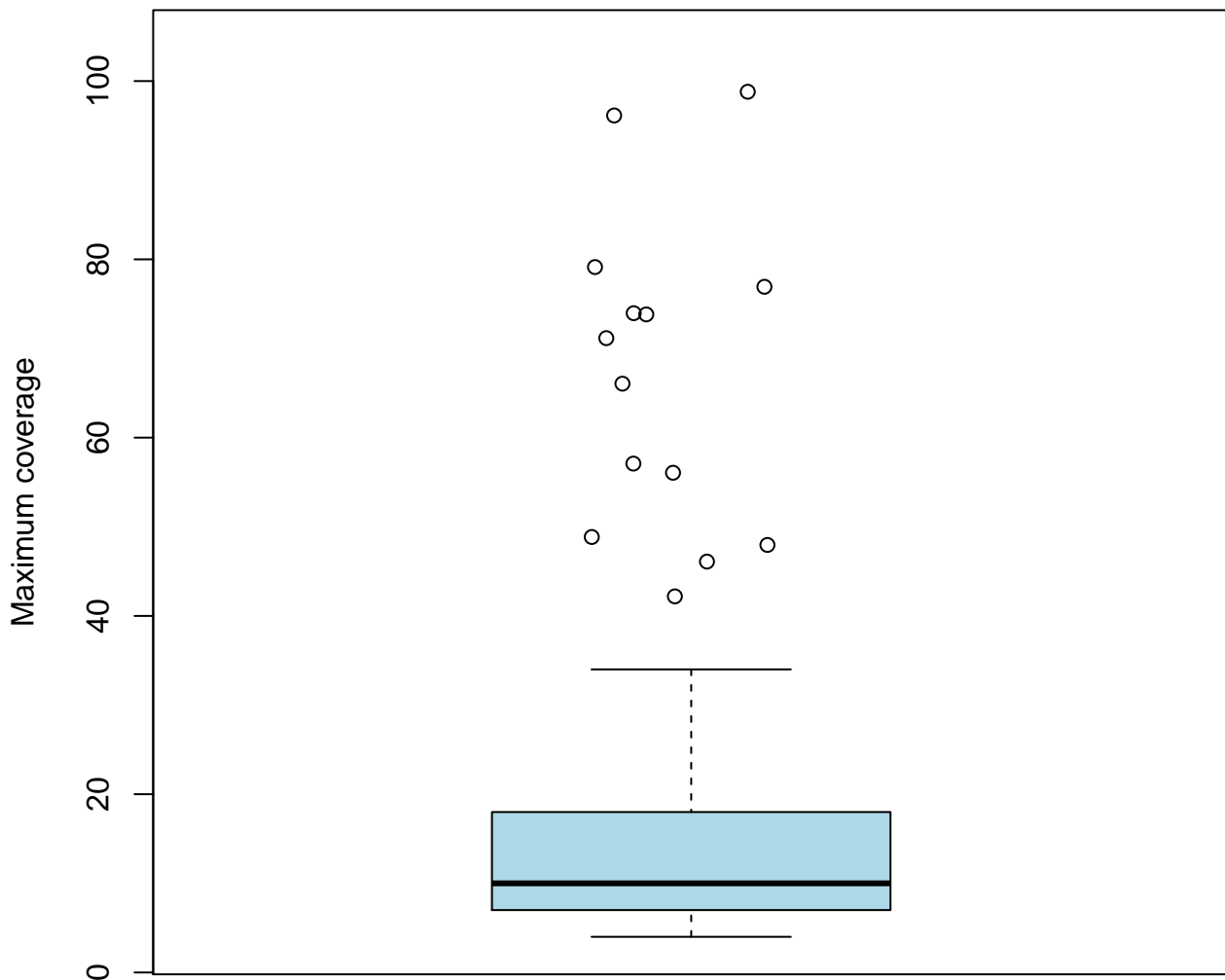
**jx in Intropolis but not "annotated" by knownGene:
106 samples; n jx: 254**



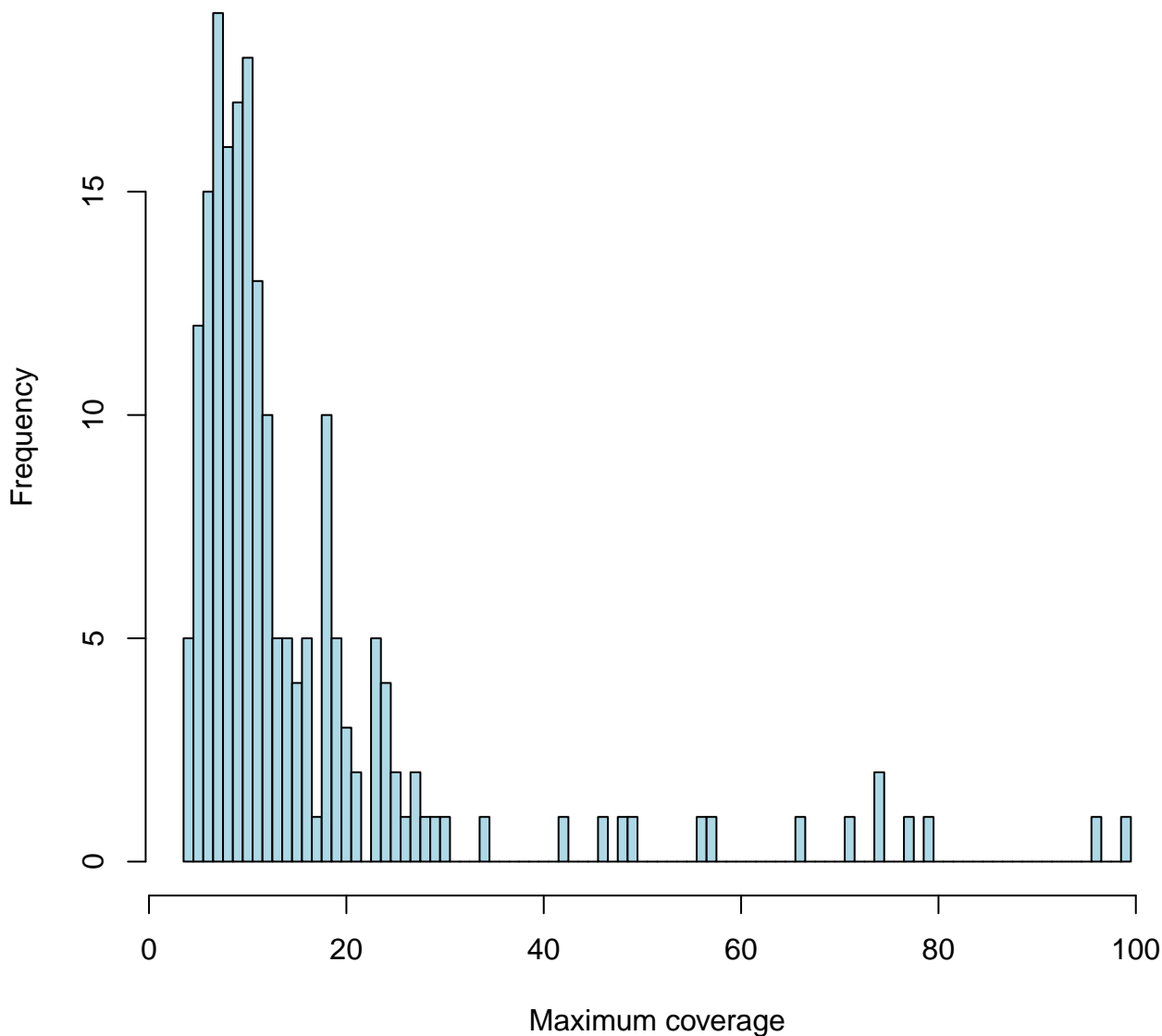
**jx in Intropolis but not "annotated" by knownGene:
106 samples; n jx: 254**



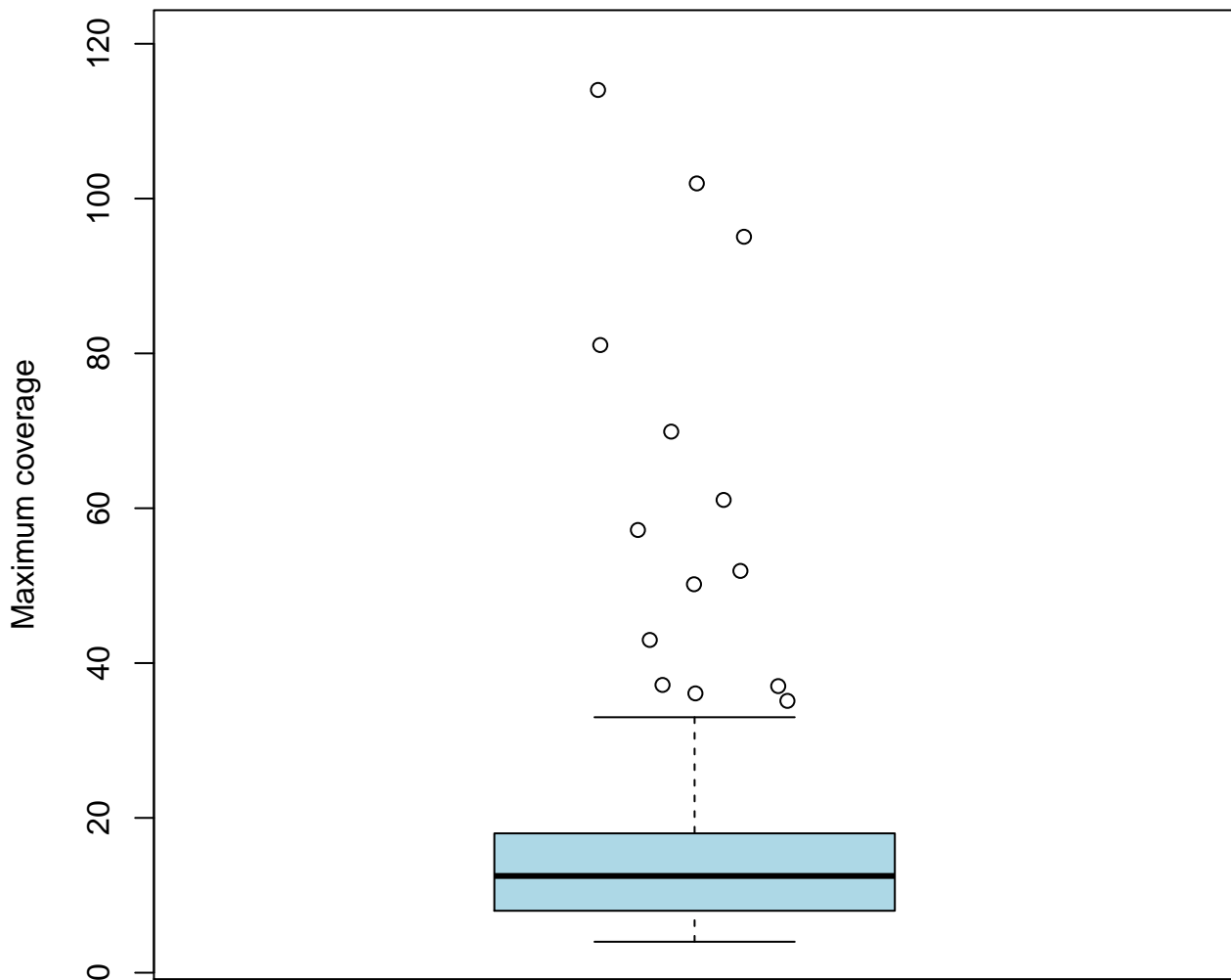
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107 samples; n jx: 197**



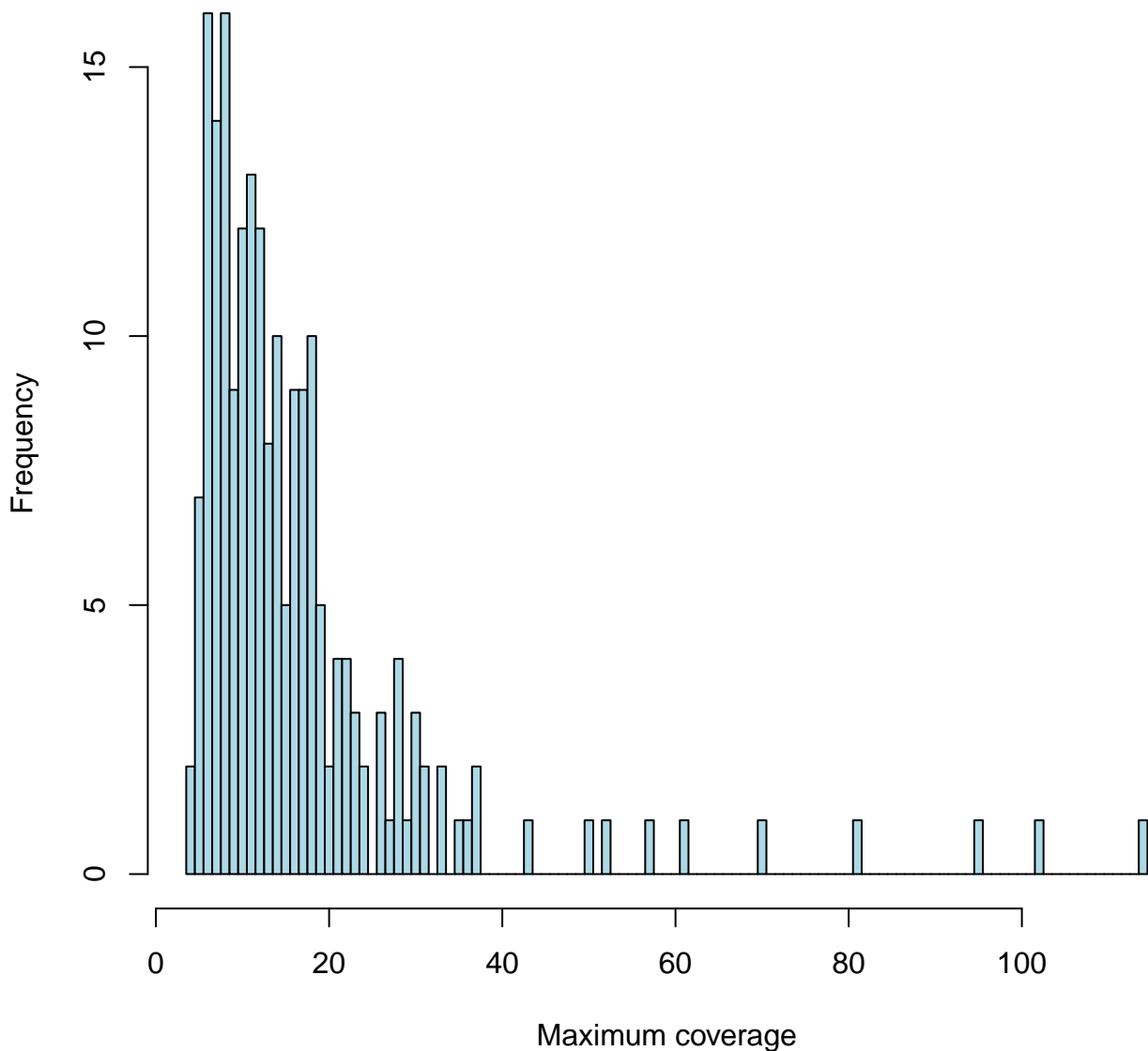
**jx in Intropolis but not "annotated" by knownGene:
107 samples; n jx: 197**



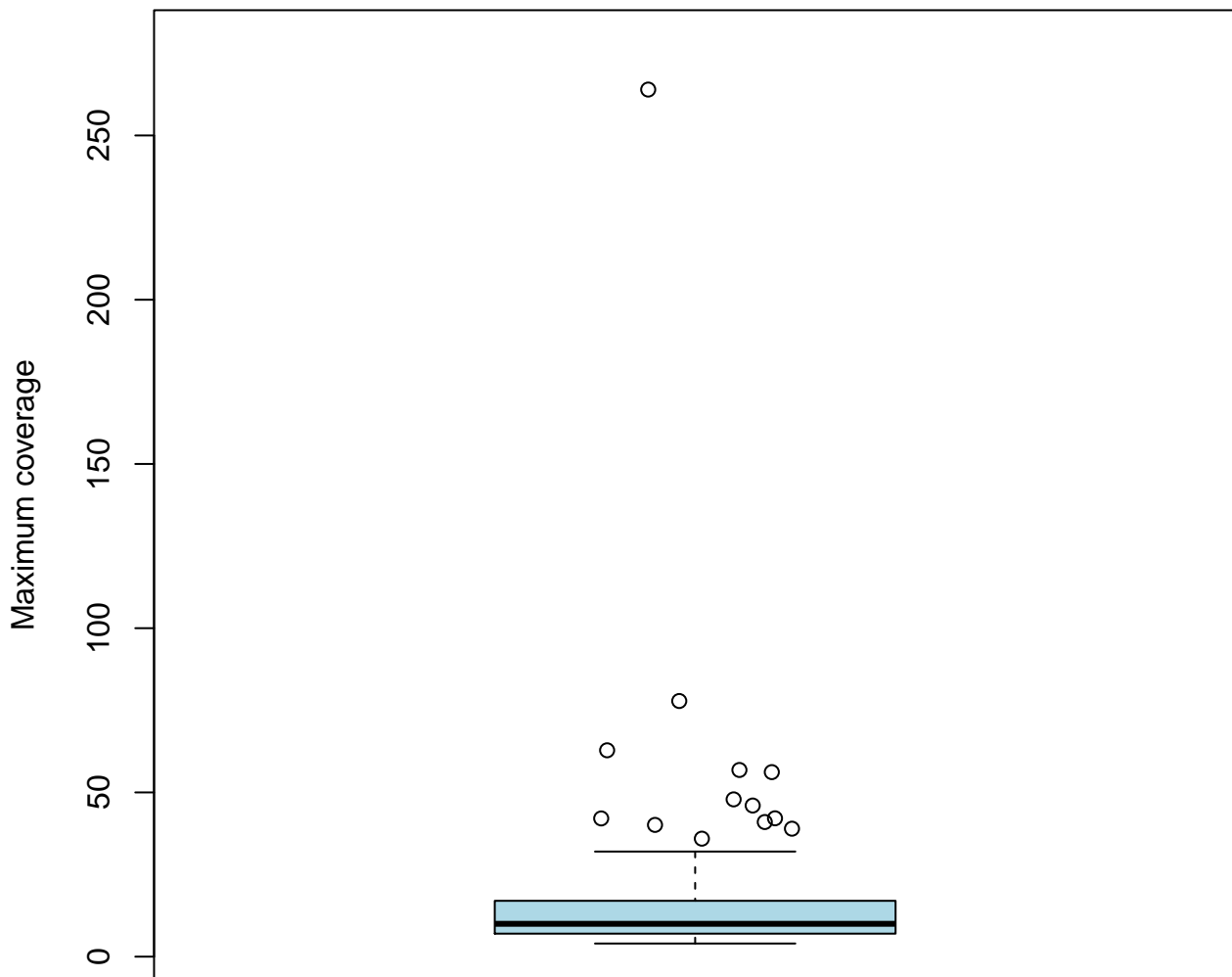
**jx in Intropolis but not "annotated" by knownGene:
108 samples; n jx: 202**



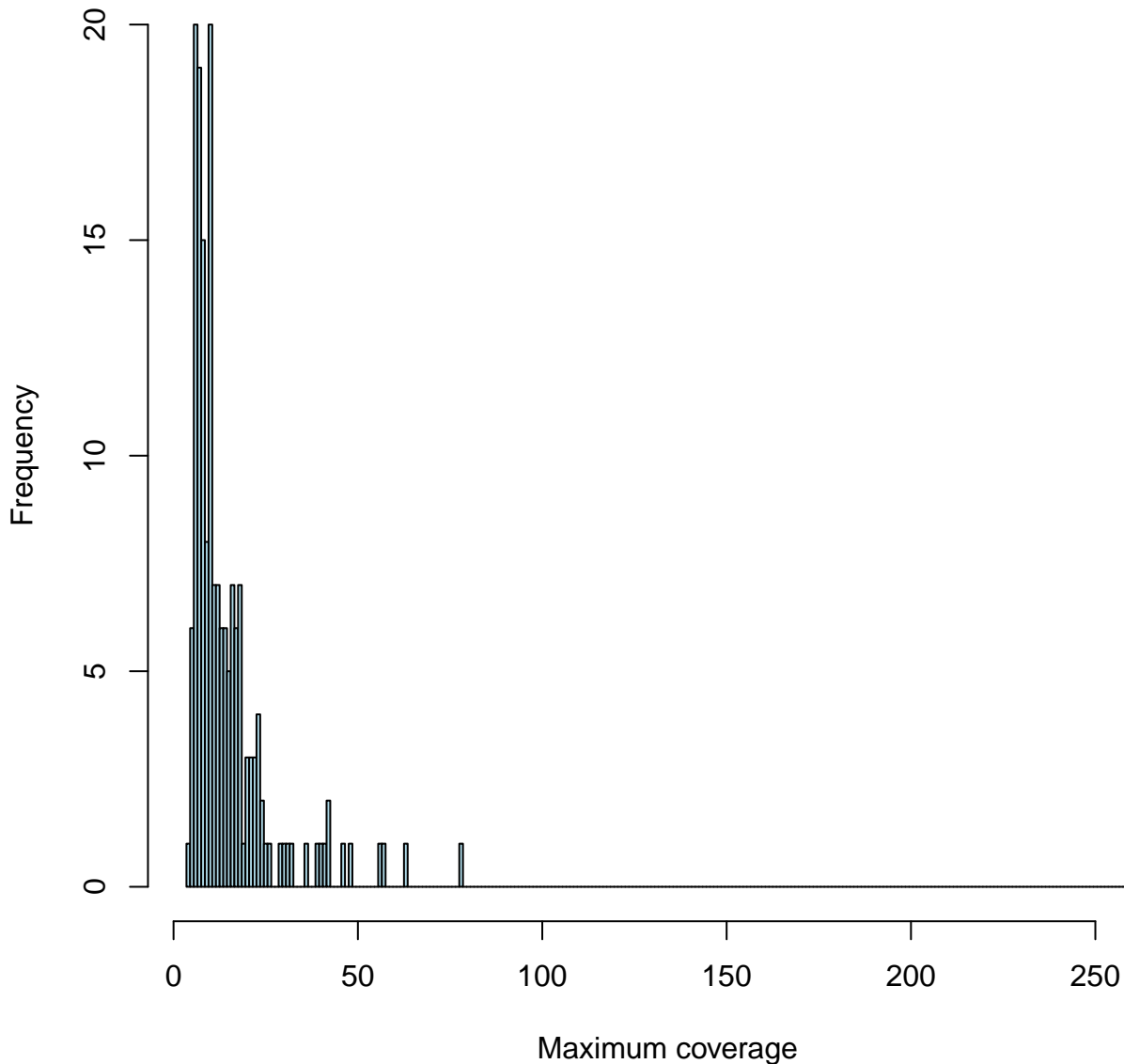
**jx in Intropolis but not "annotated" by knownGene:
108 samples; n jx: 202**



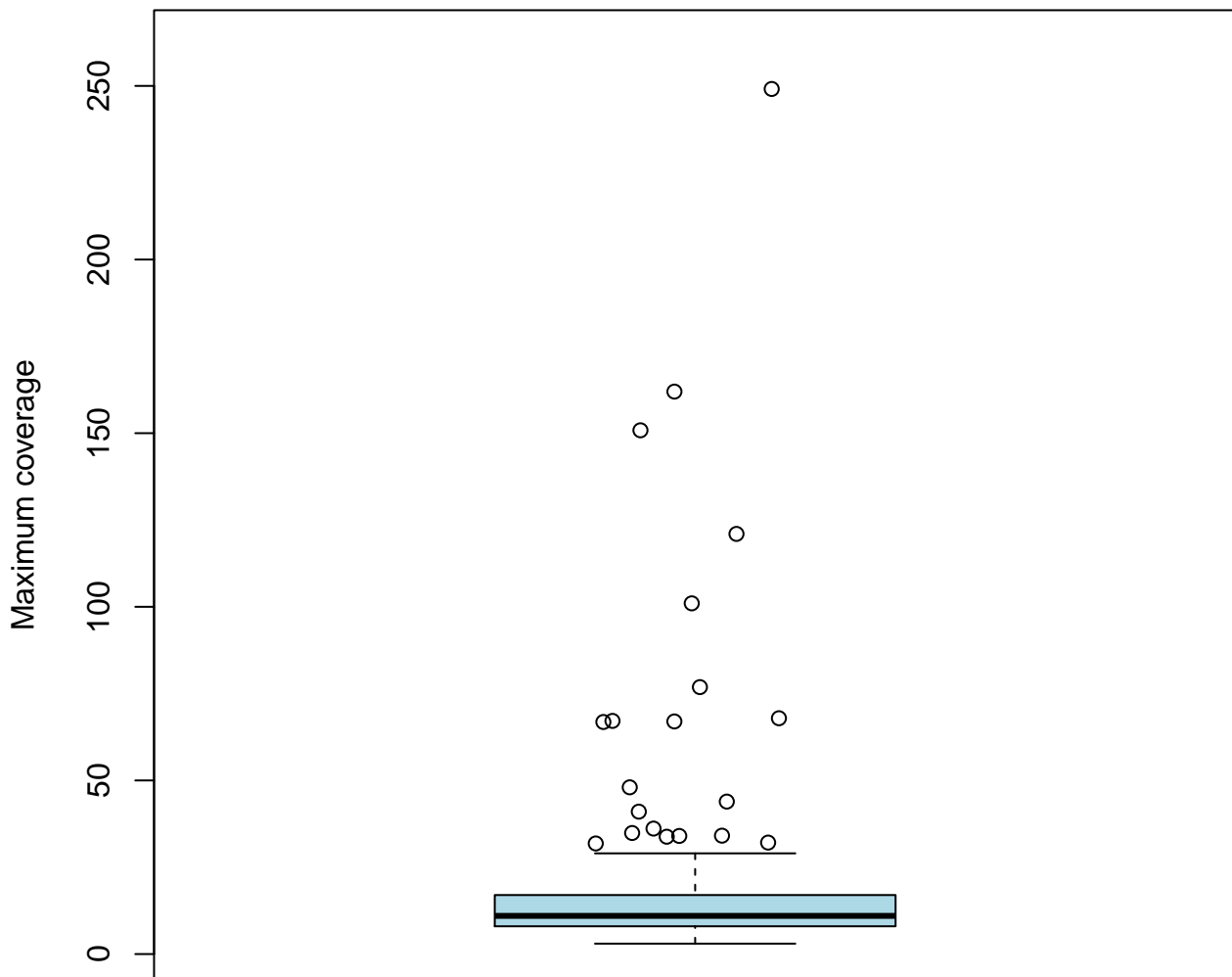
**jx in Intropolis but not "annotated" by knownGene:
109 samples; n jx: 175**



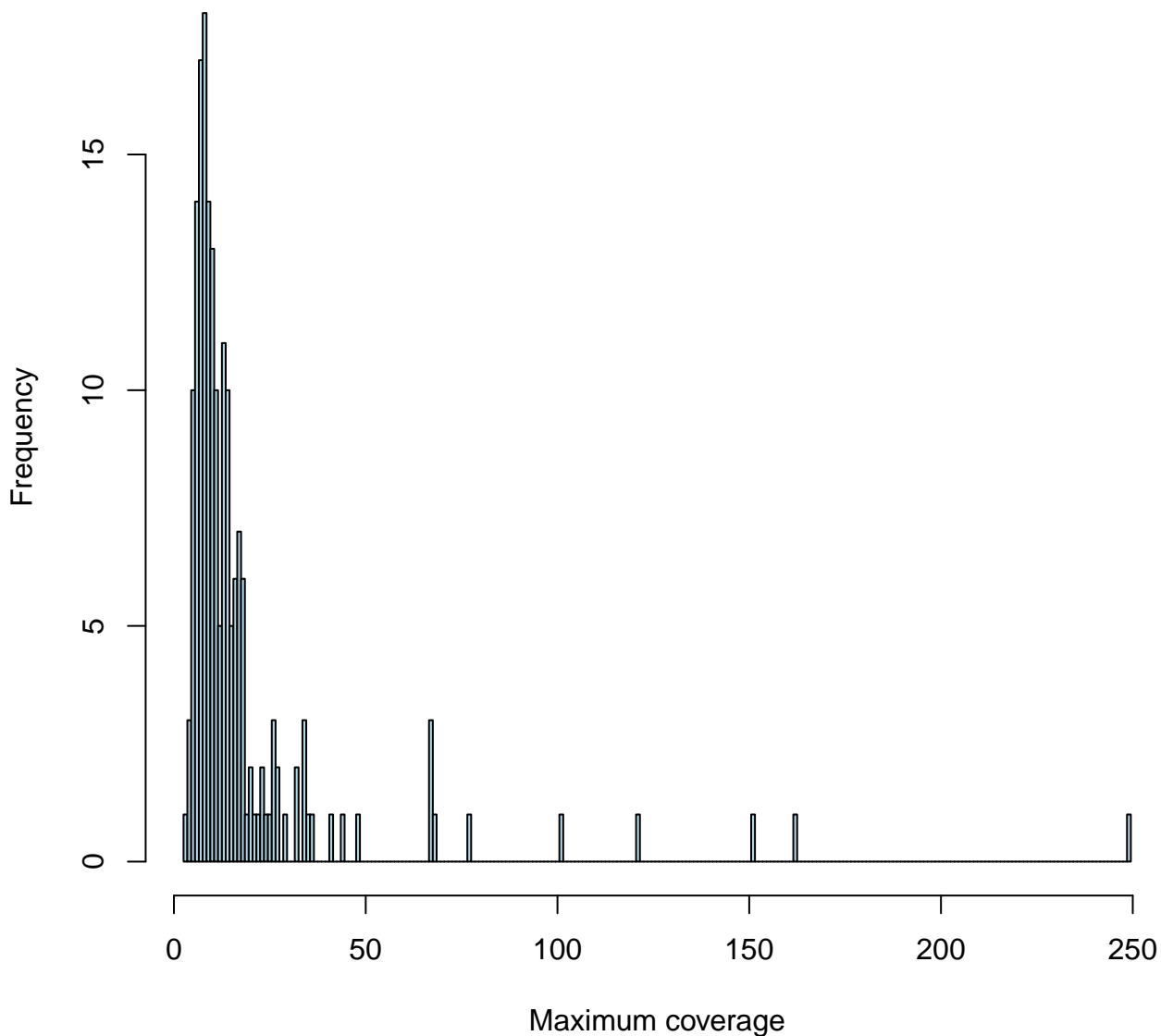
**jx in Intropolis but not "annotated" by knownGene:
109 samples; n jx: 175**



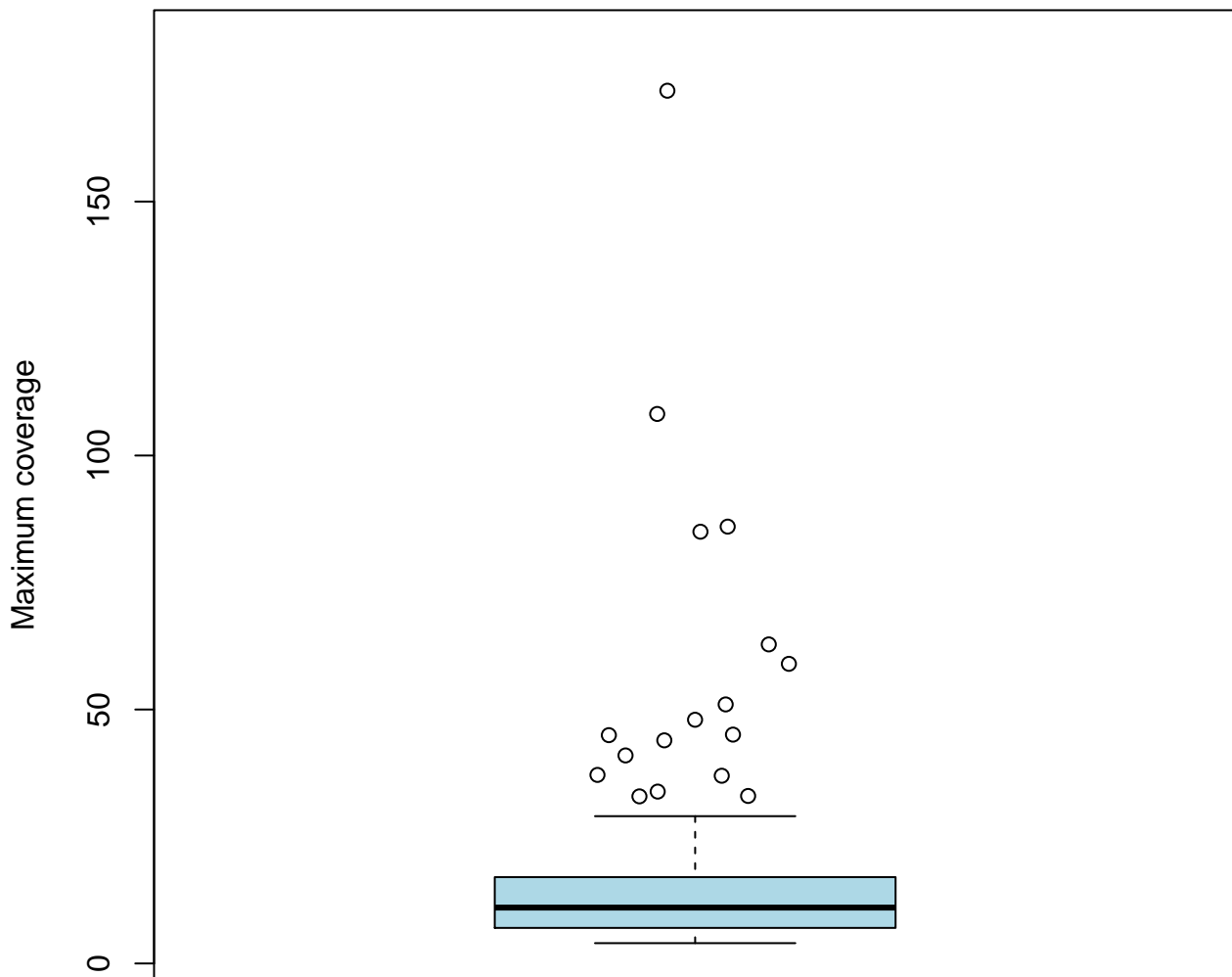
**jx in Intropolis but not "annotated" by knownGene:
110 samples; n jx: 185**



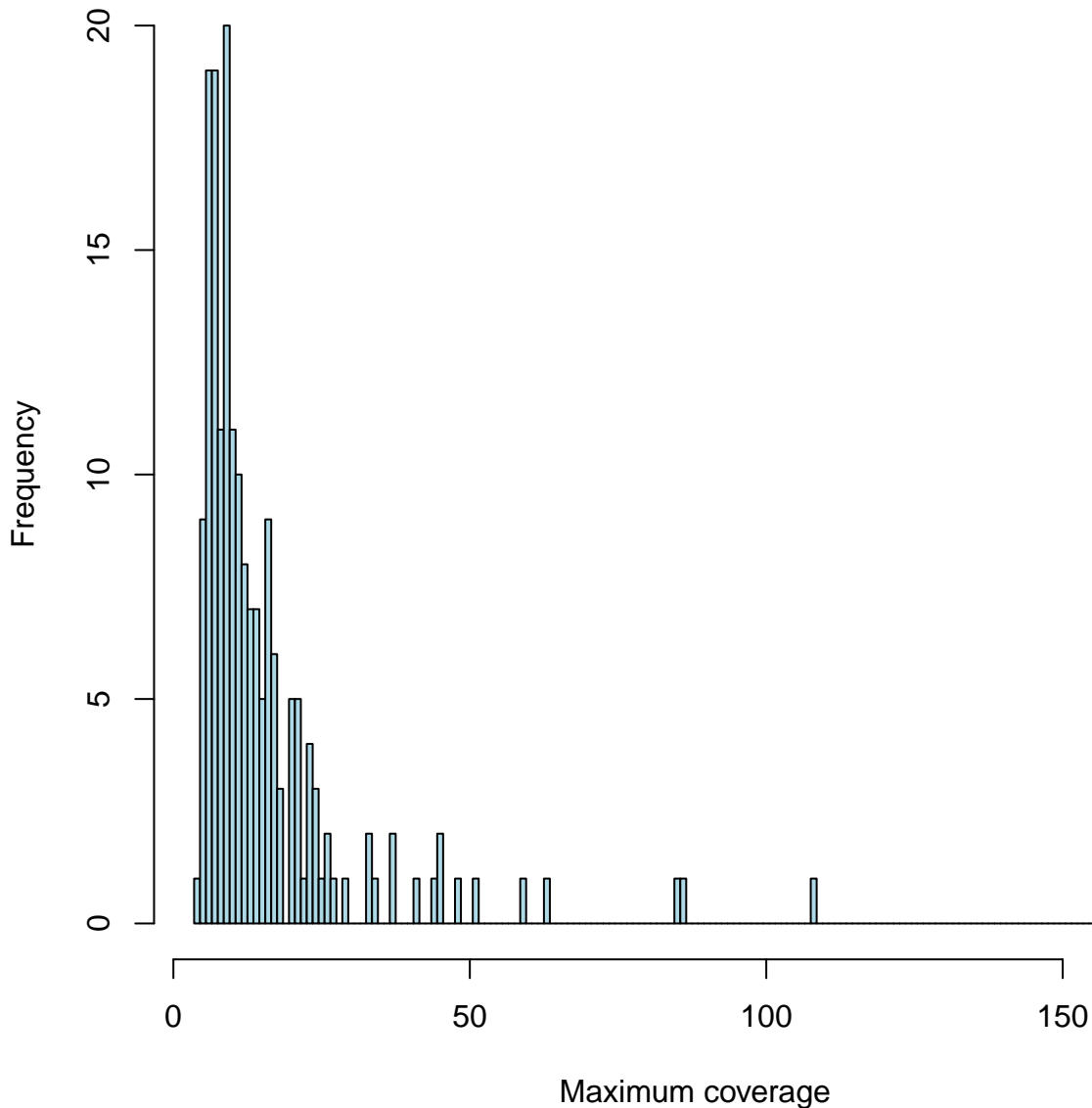
**jx in Intropolis but not "annotated" by knownGene:
110 samples; n jx: 185**



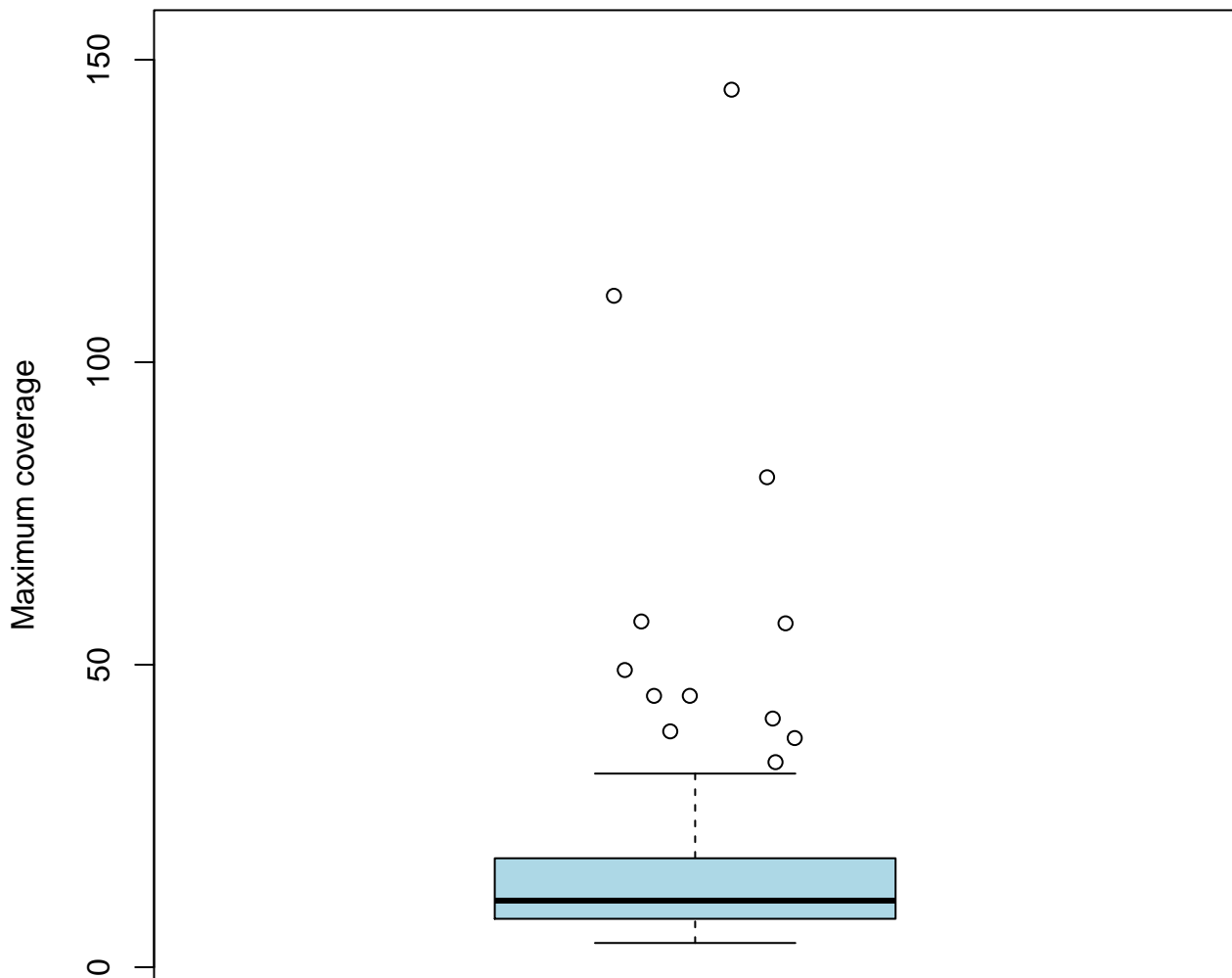
**jx in Intropolis but not "annotated" by knownGene:
111 samples; n jx: 185**



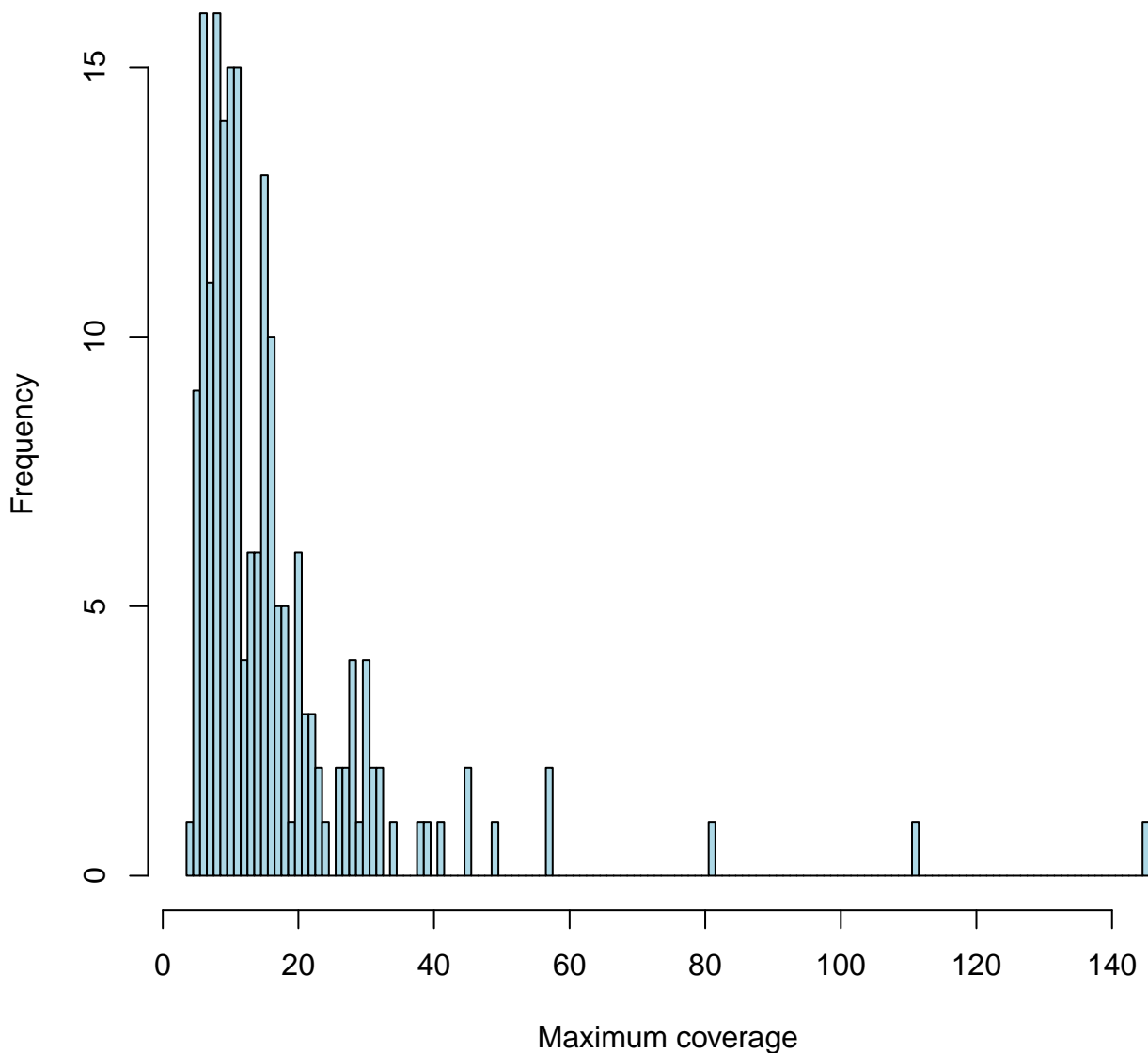
**jx in Intropolis but not "annotated" by knownGene:
111 samples; n jx: 185**



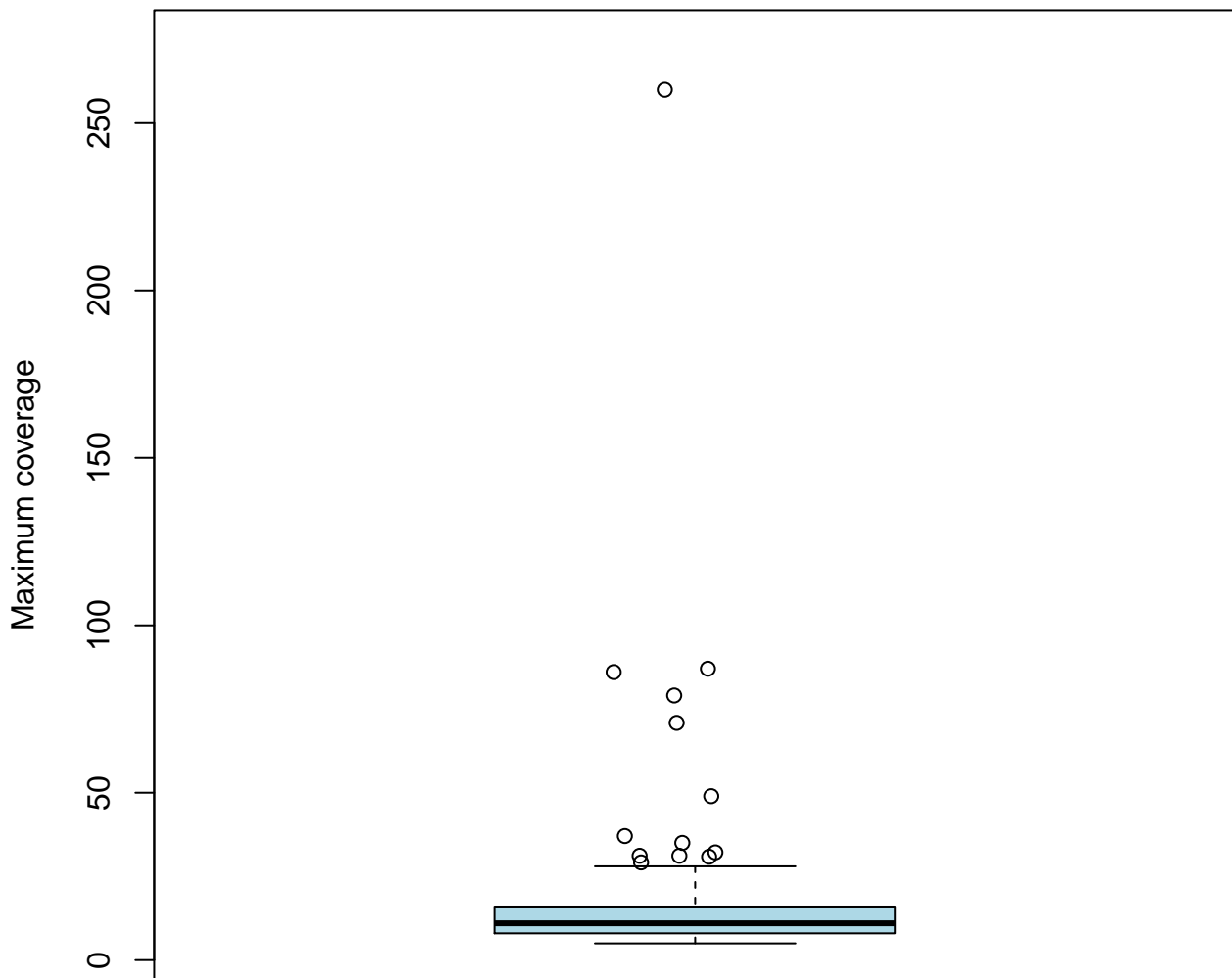
**jx in Intropolis but not "annotated" by knownGene:
112 samples; n jx: 191**



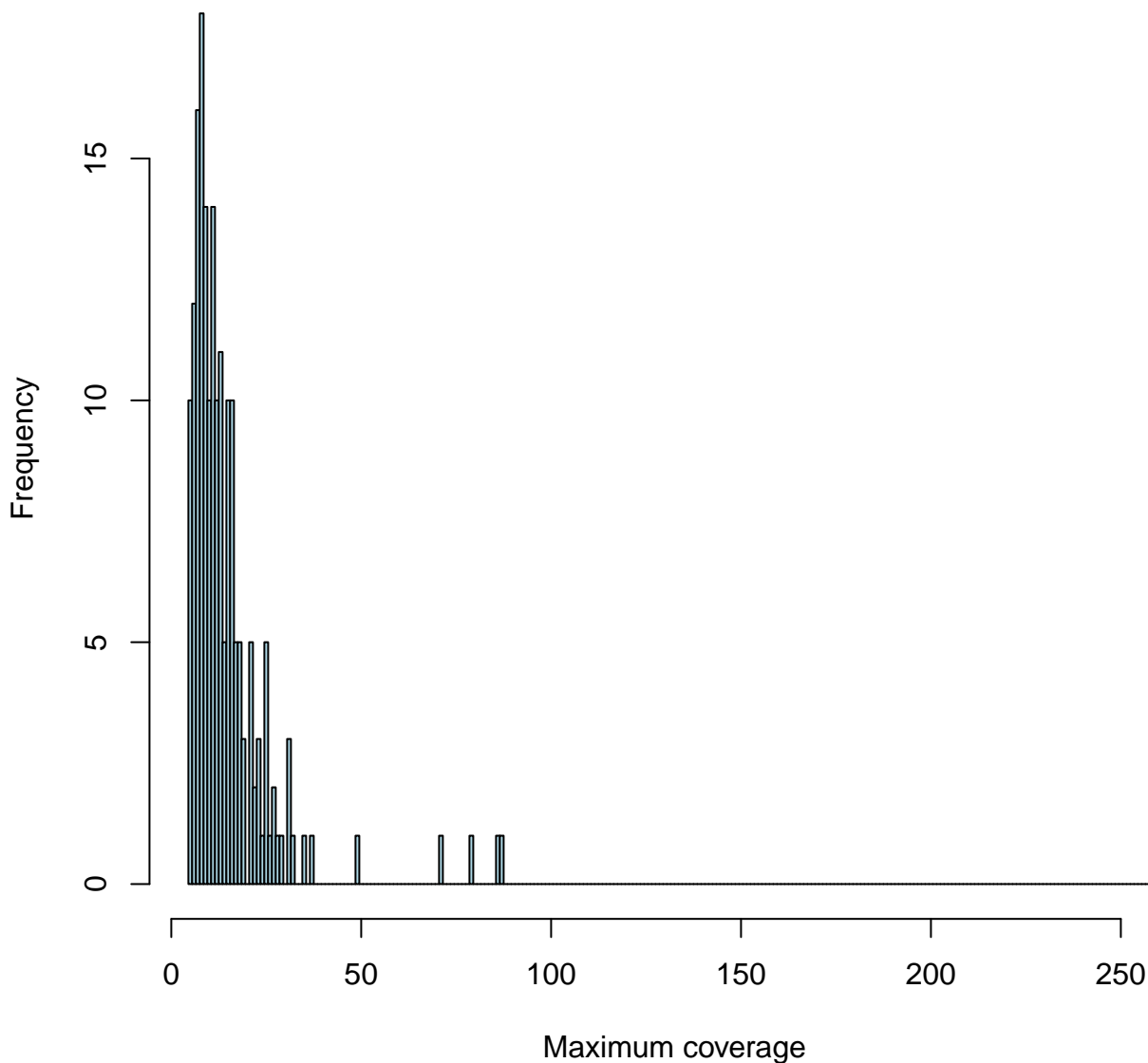
**jx in Intropolis but not "annotated" by knownGene:
112 samples; n jx: 191**



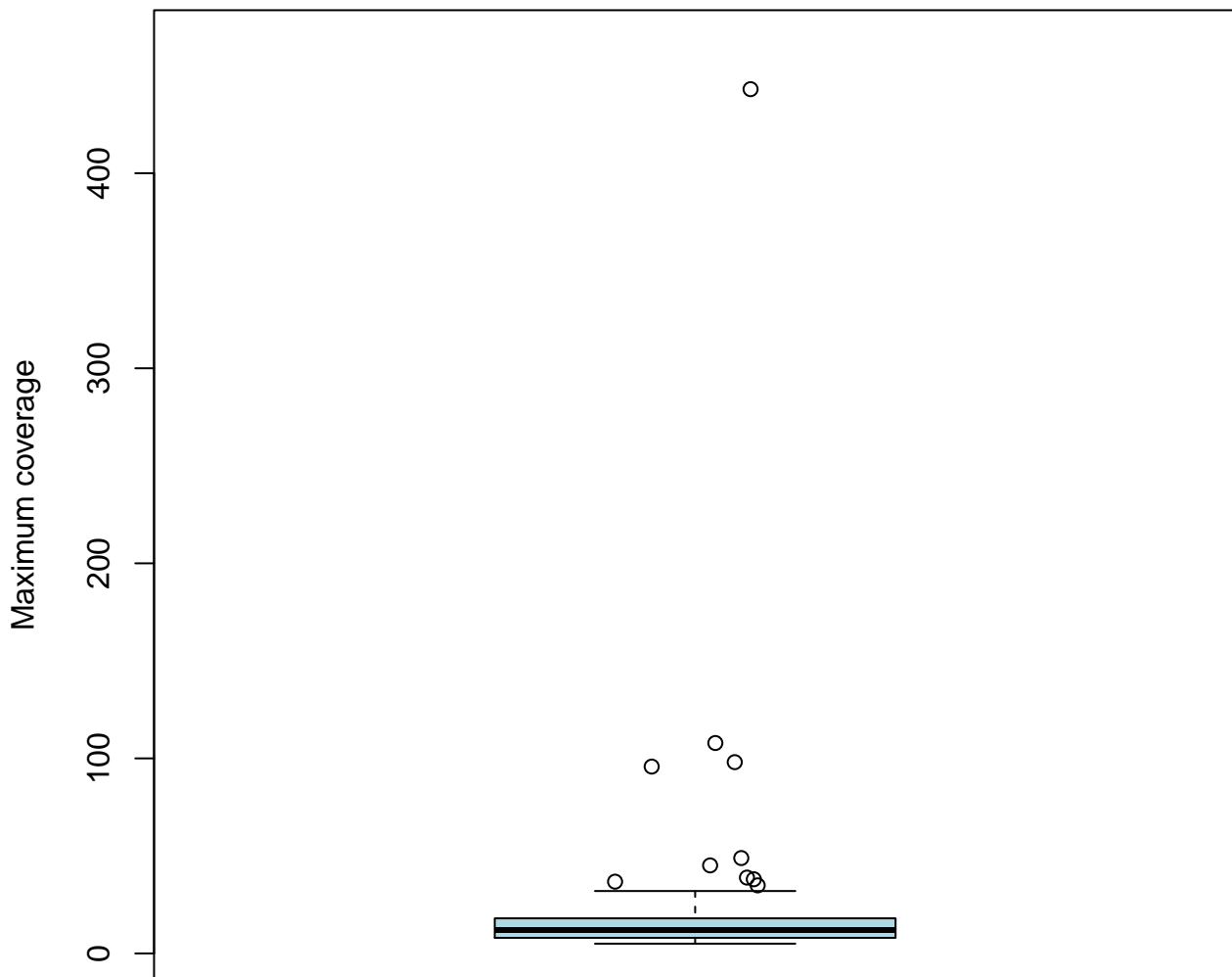
**jx in Intropolis but not "annotated" by knownGene:
113 samples; n jx: 186**



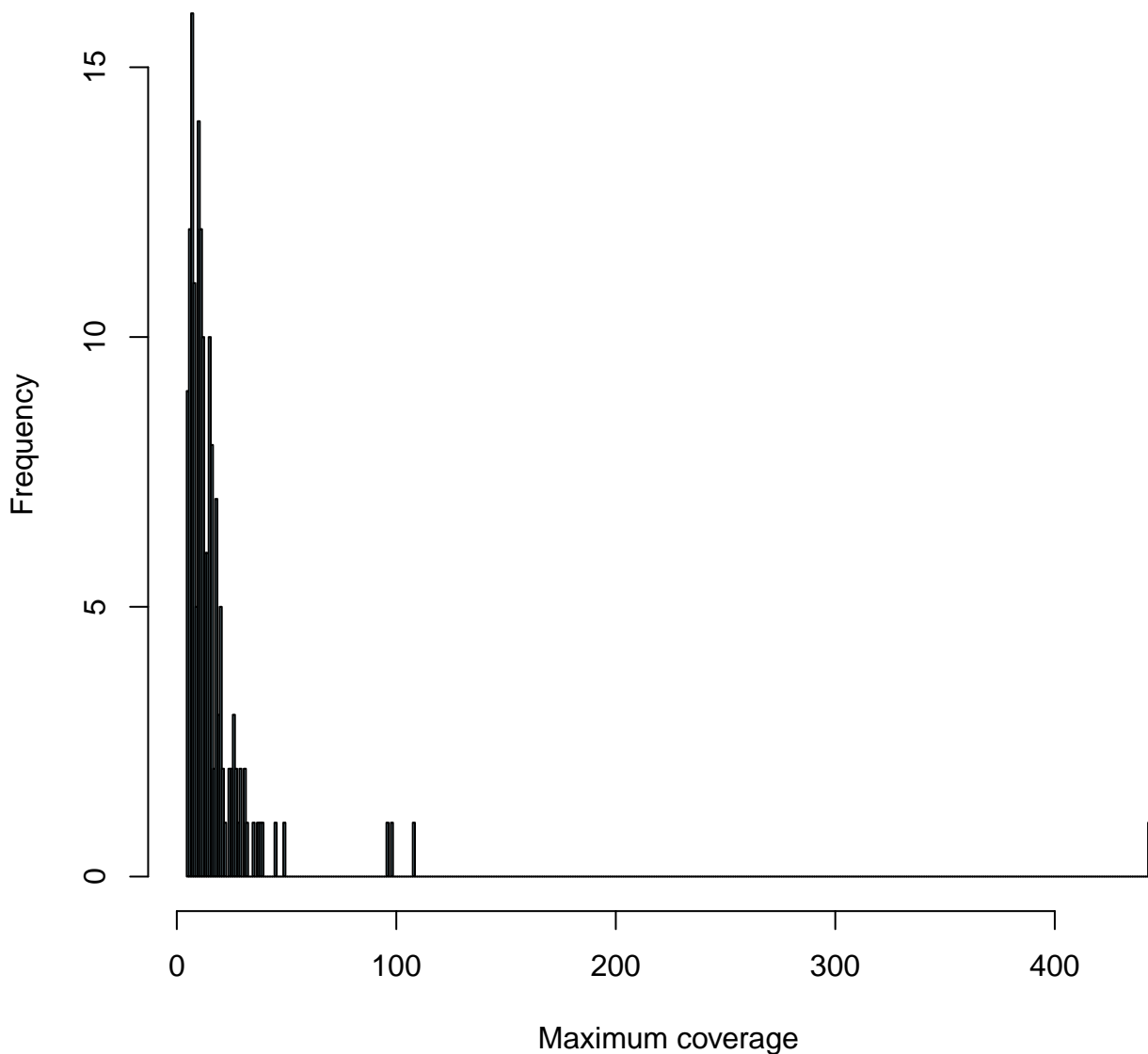
**jx in Intropolis but not "annotated" by knownGene:
113 samples; n jx: 186**



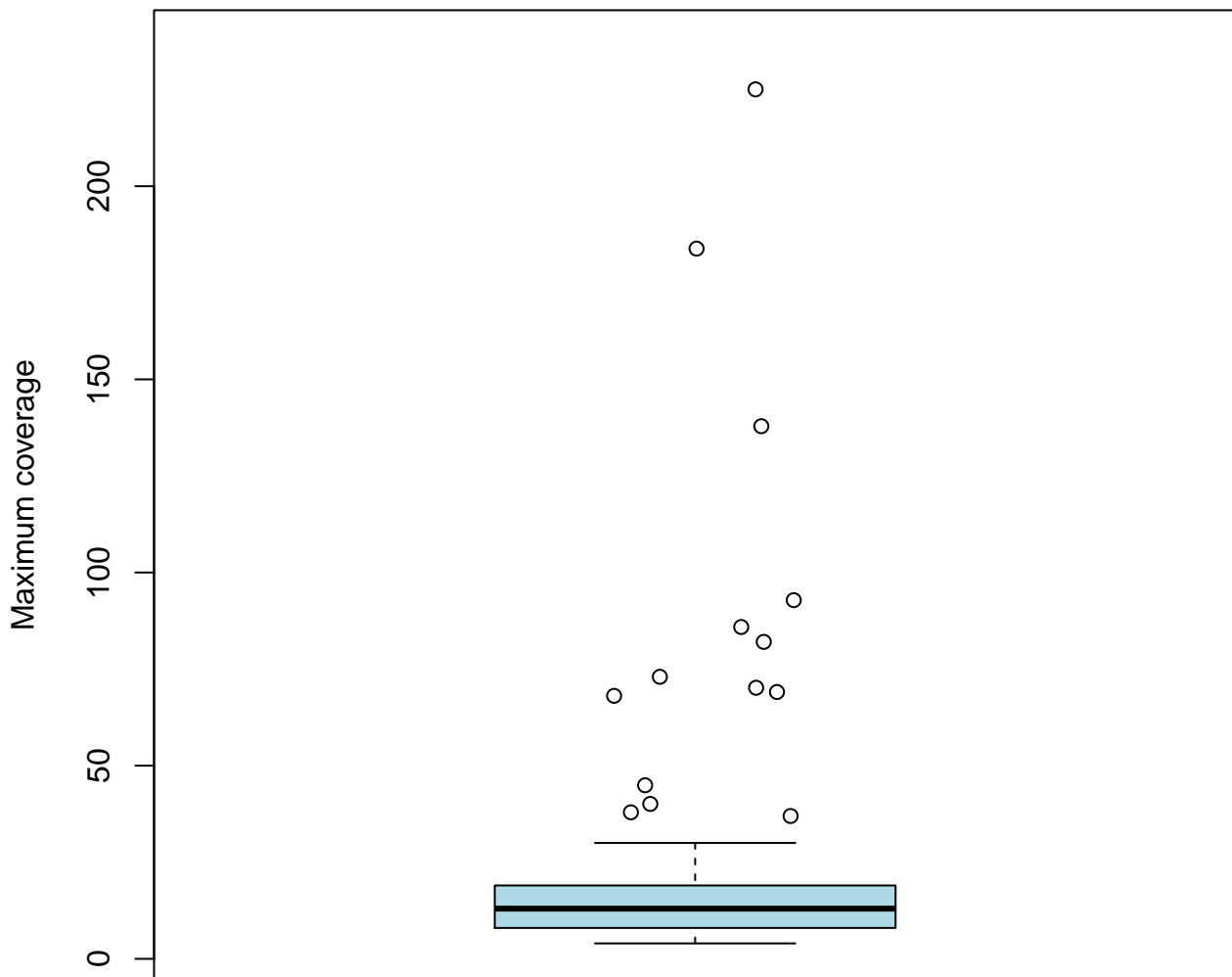
**jx in Intropolis but not "annotated" by knownGene:
114 samples; n jx: 164**



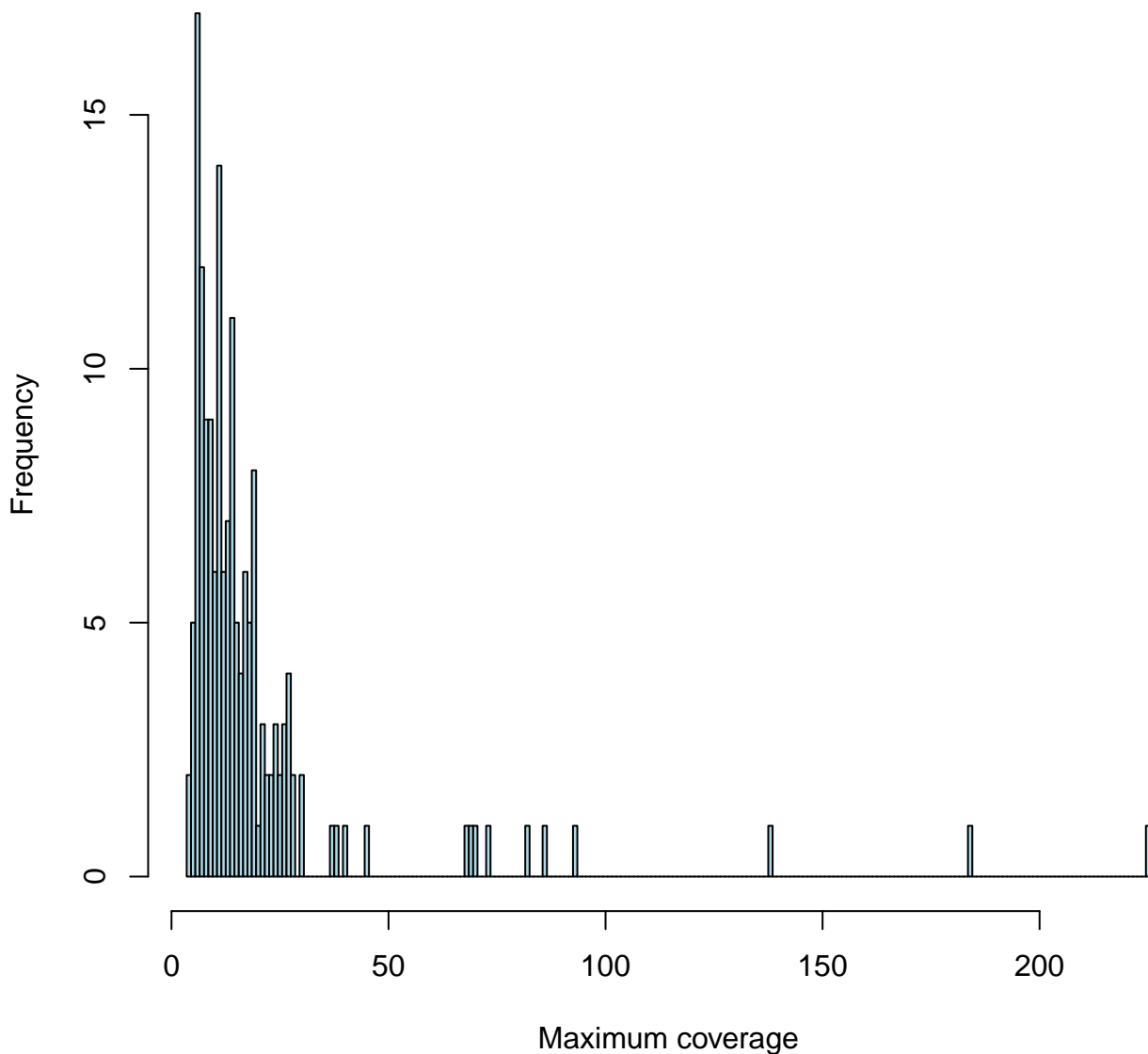
**jx in Intropolis but not "annotated" by knownGene:
114 samples; n jx: 164**



**jx in Intropolis but not "annotated" by knownGene:
115 samples; n jx: 164**



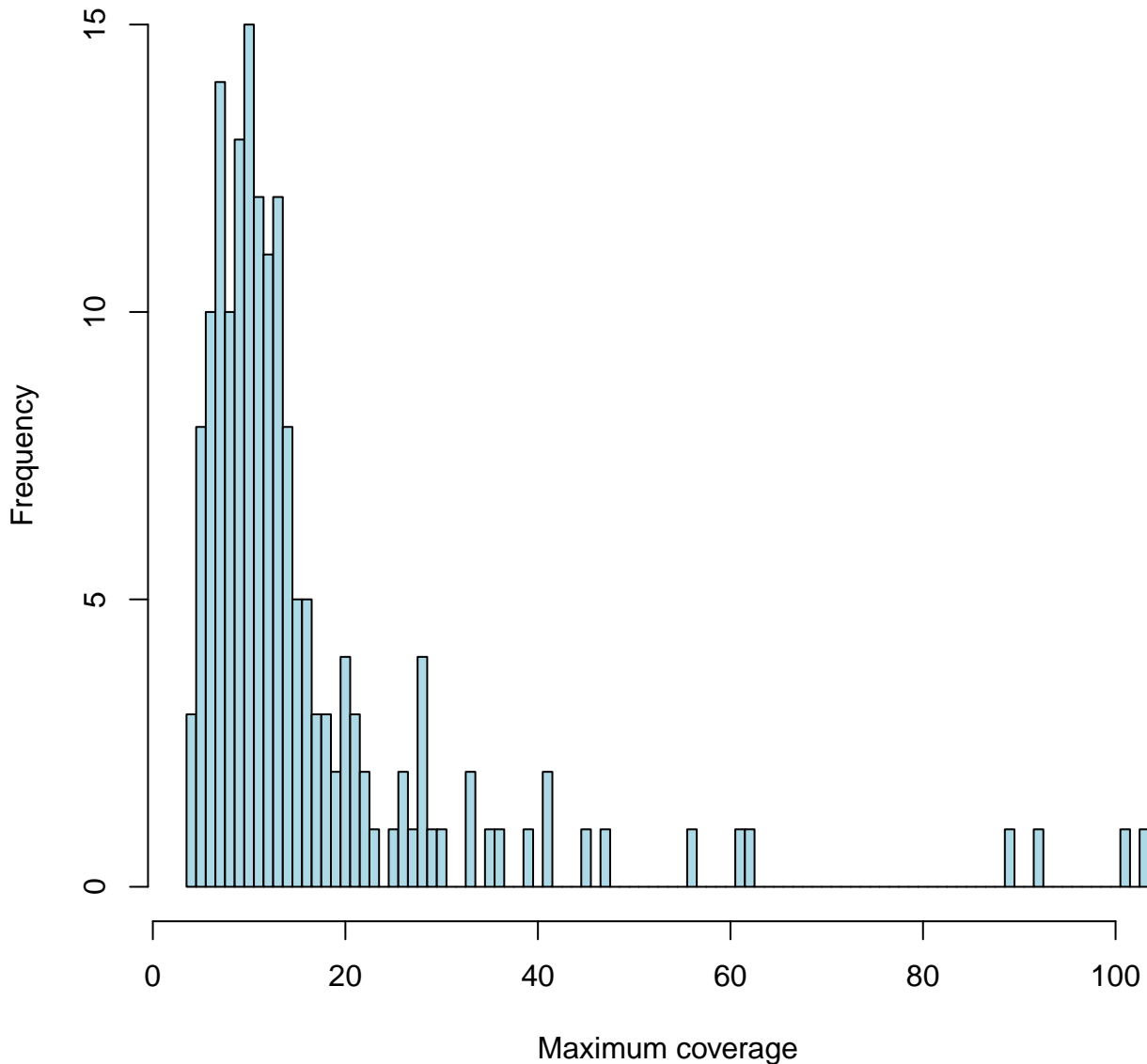
**jx in Intropolis but not "annotated" by knownGene:
115 samples; n jx: 164**



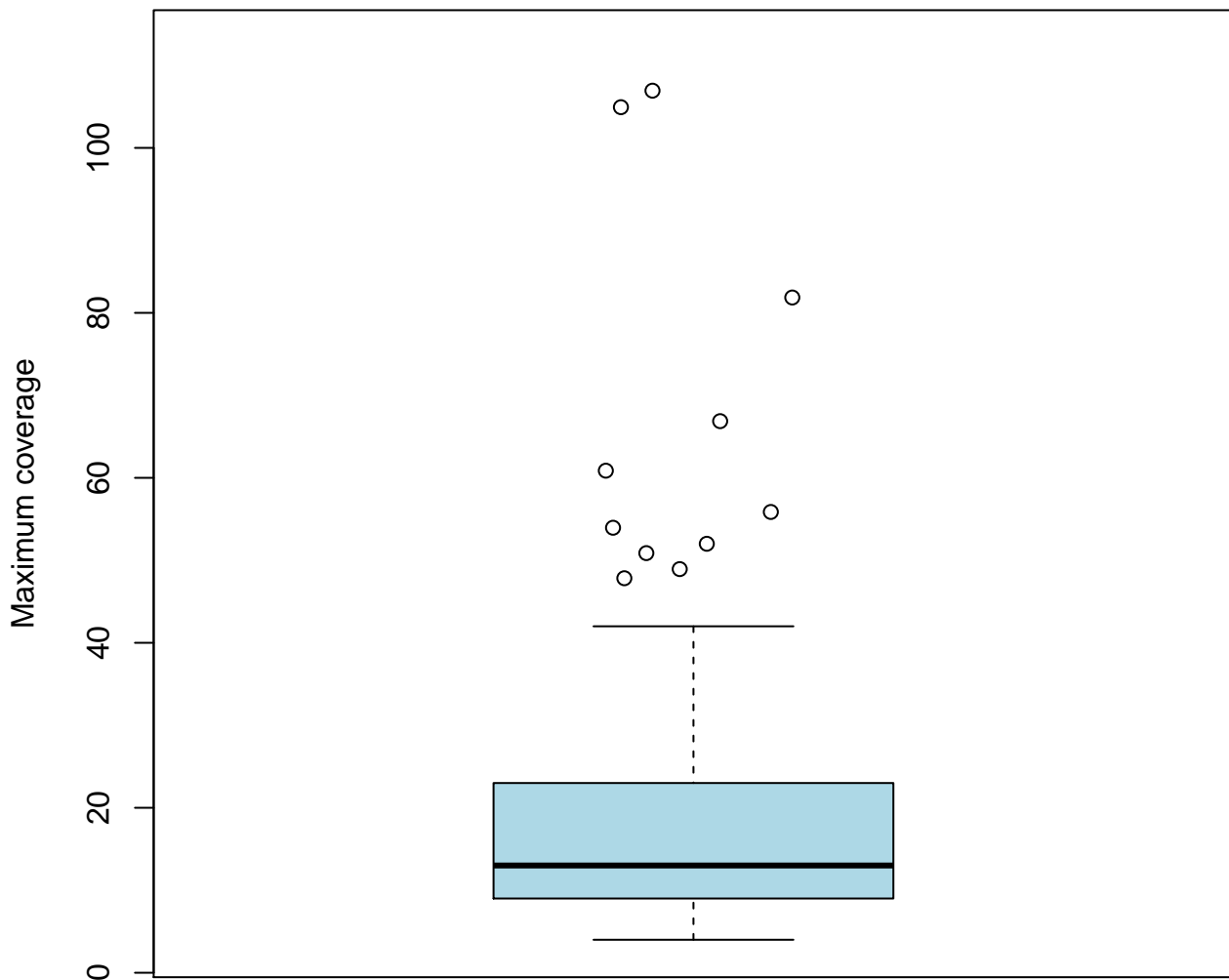
116 samples; n jx: 170



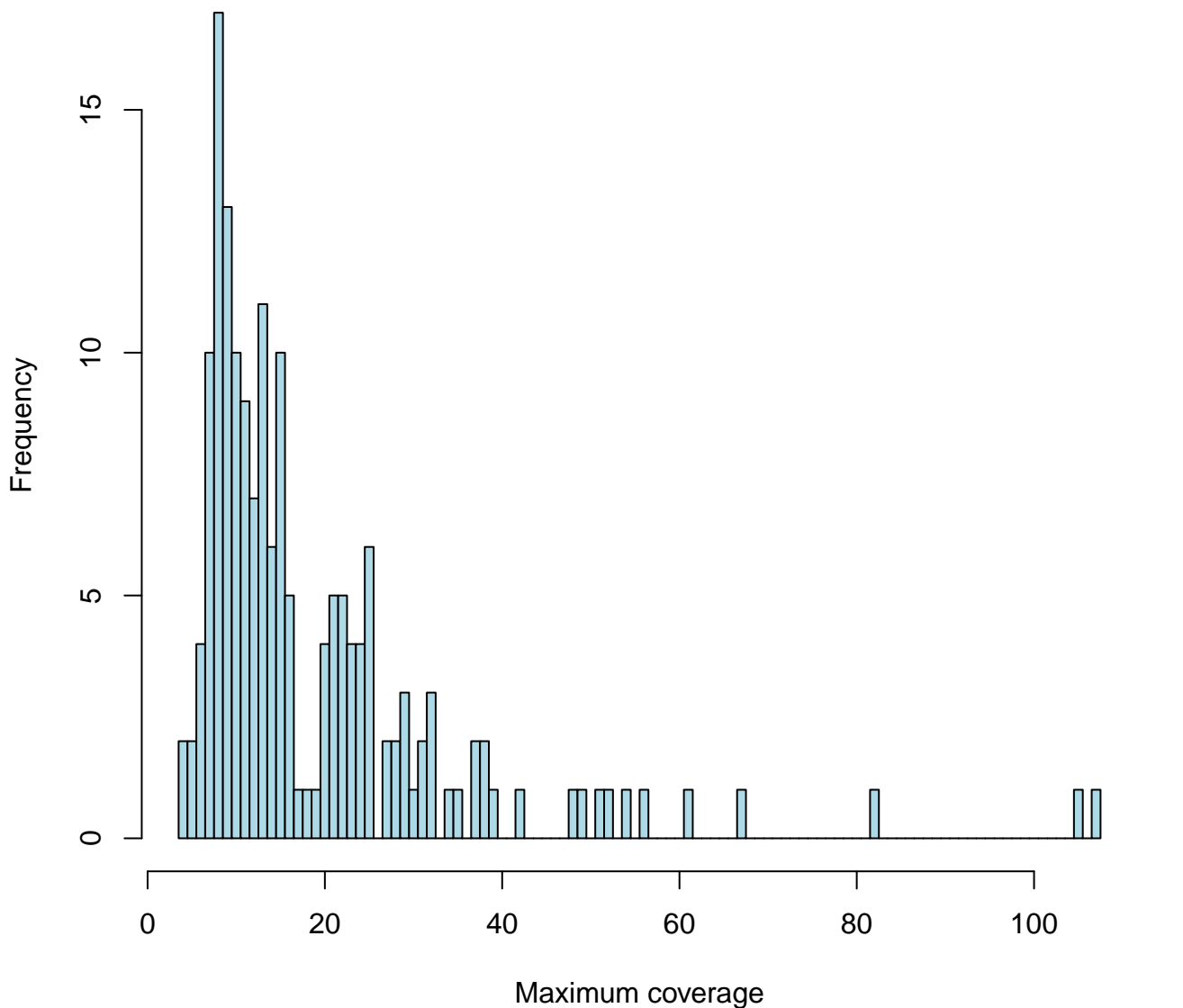
**jx in Intropolis but not "annotated" by knownGene:
116 samples; n jx: 170**



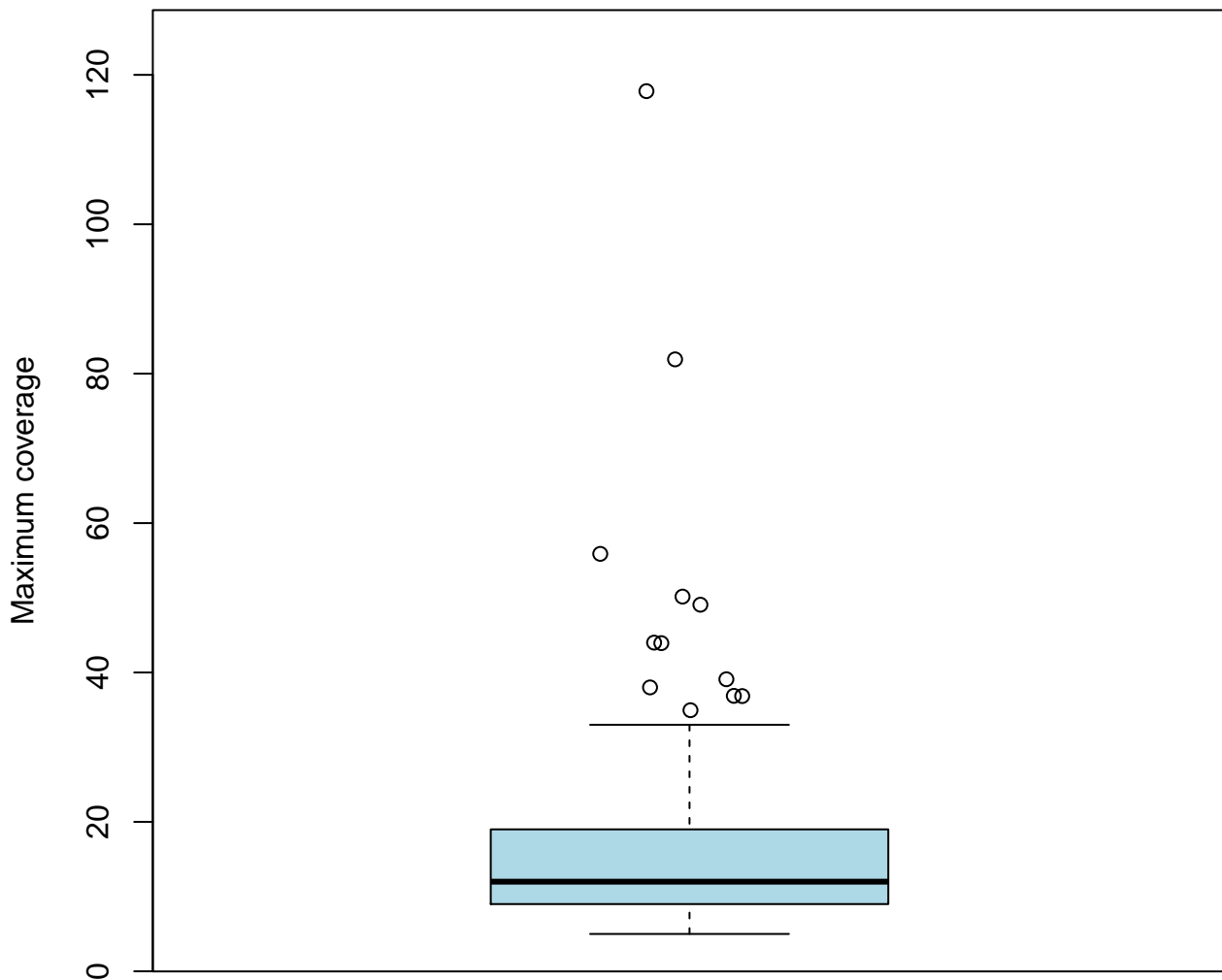
**jx in Intropolis but not "annotated" by knownGene:
117 samples; n jx: 169**



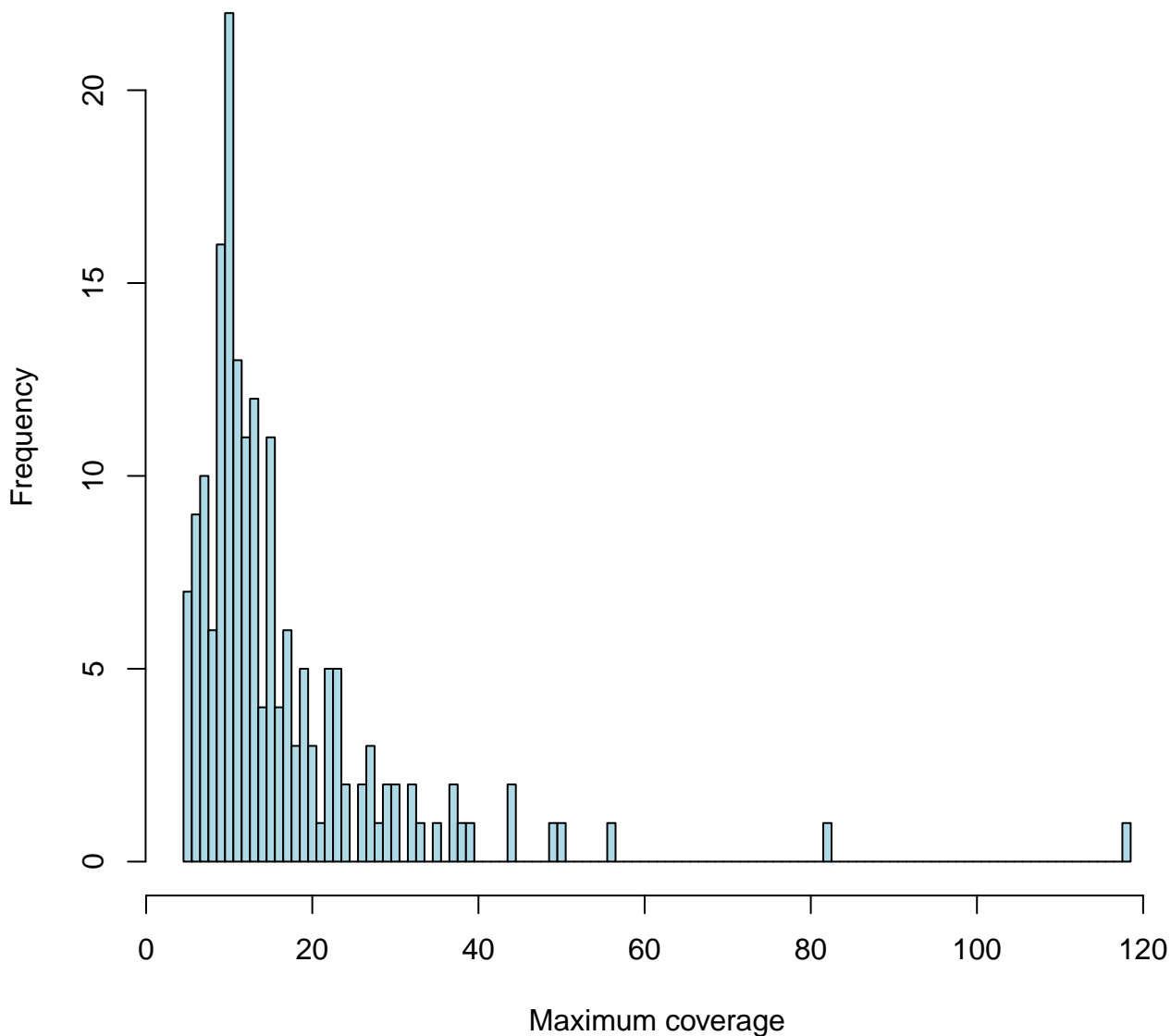
**jx in Intropolis but not "annotated" by knownGene:
117 samples; n jx: 169**



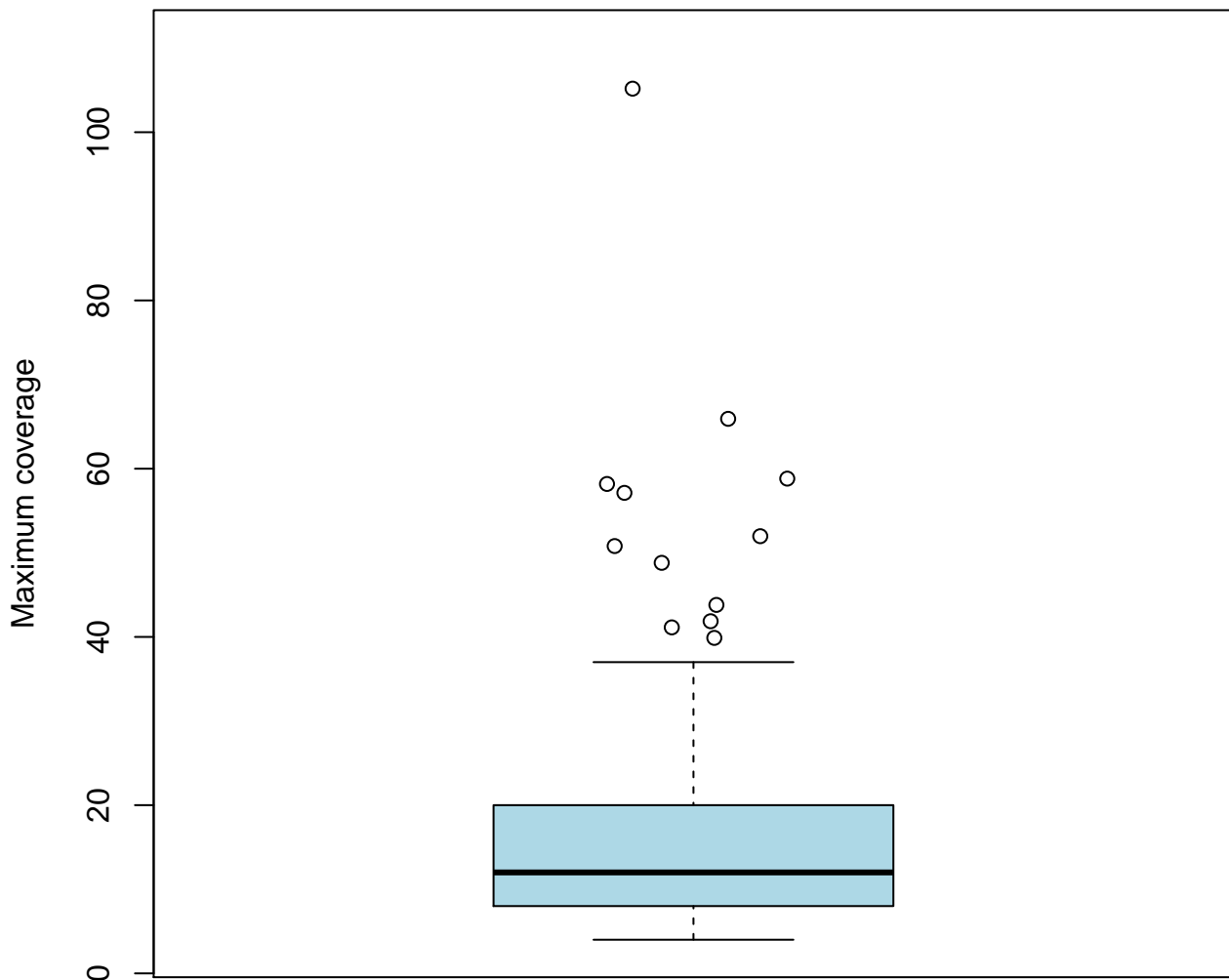
**jx in Intropolis but not "annotated" by knownGene:
118 samples; n jx: 180**



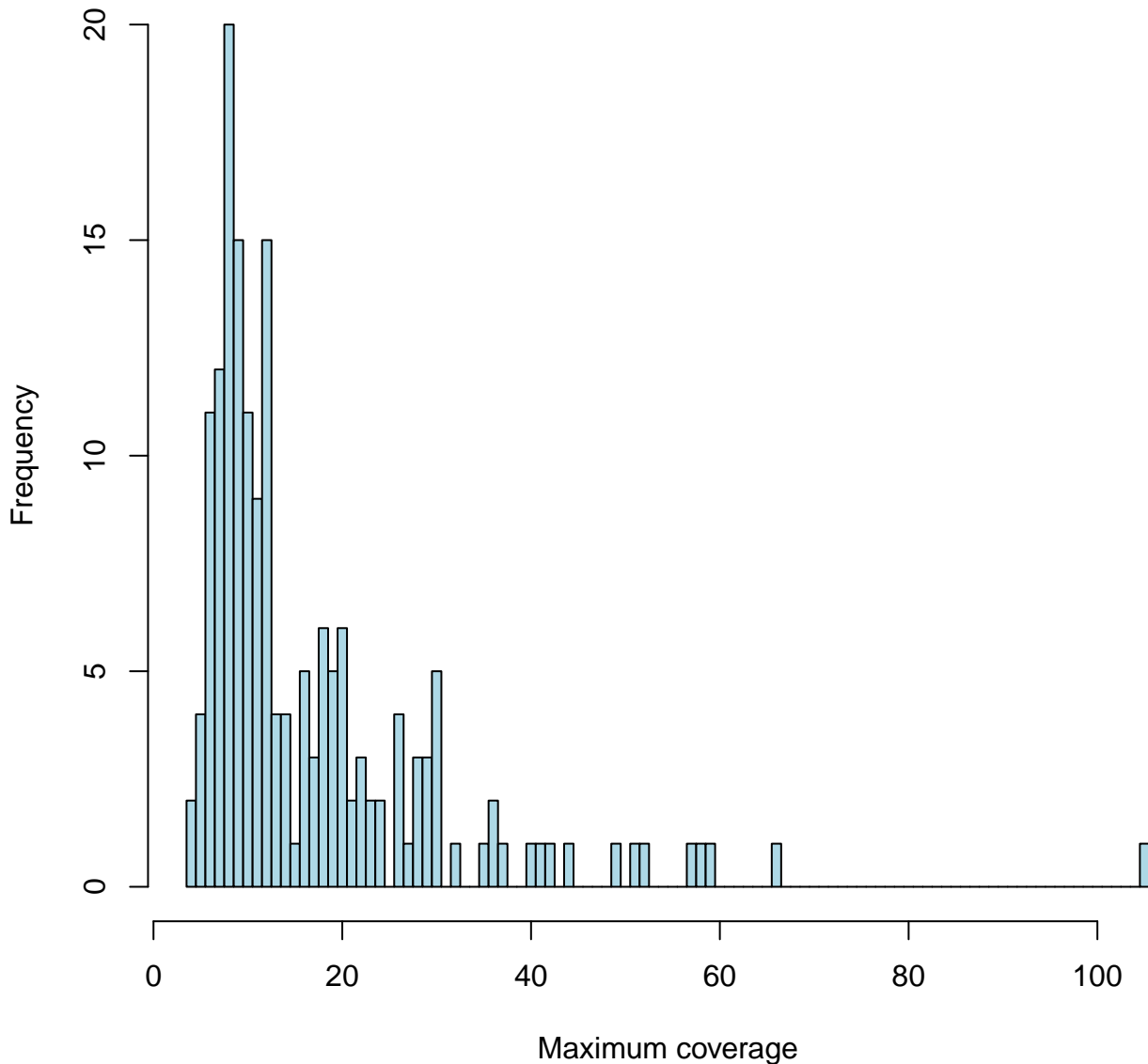
**jx in Intropolis but not "annotated" by knownGene:
118 samples; n jx: 180**



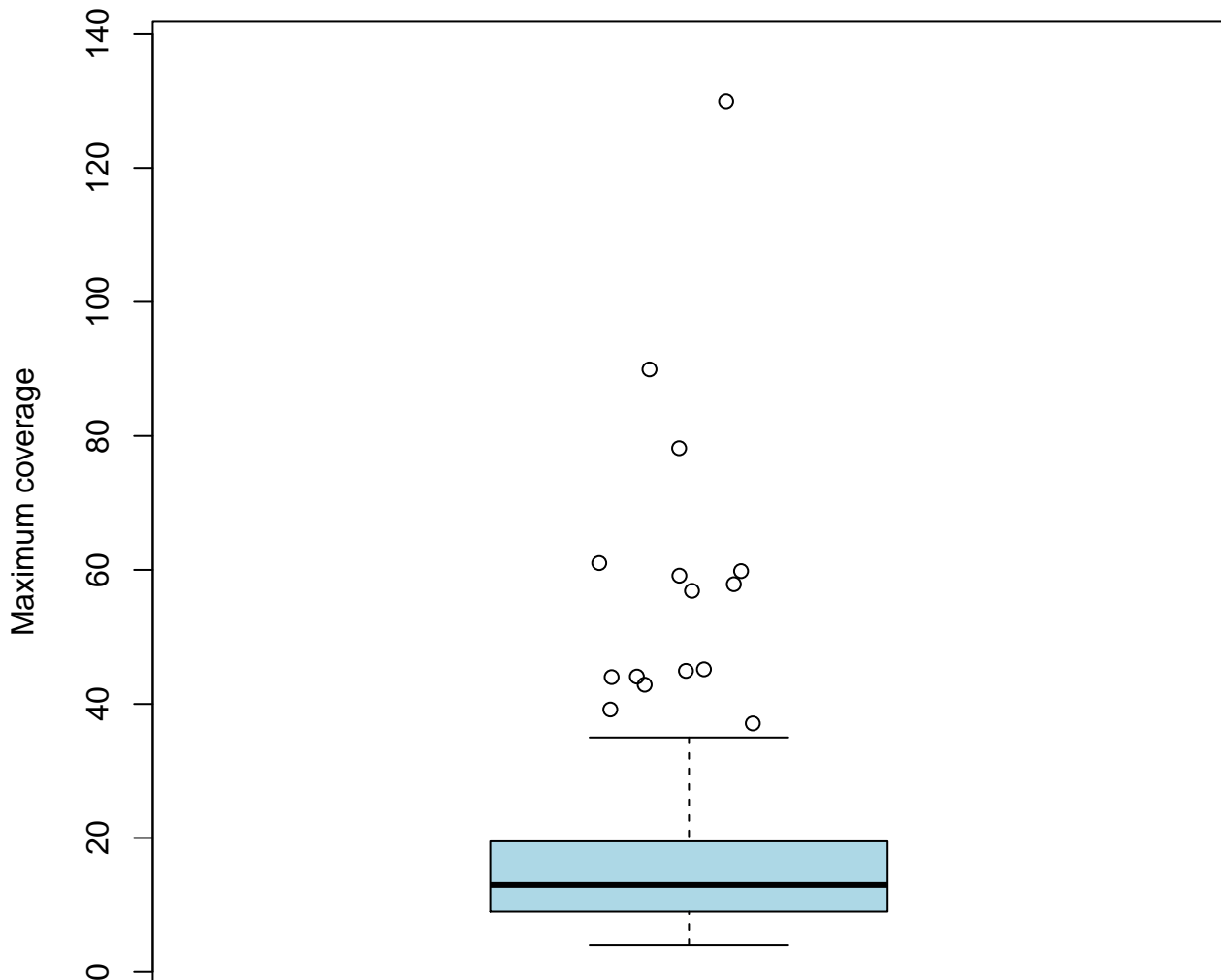
**jx in Intropolis but not "annotated" by knownGene:
119 samples; n jx: 175**



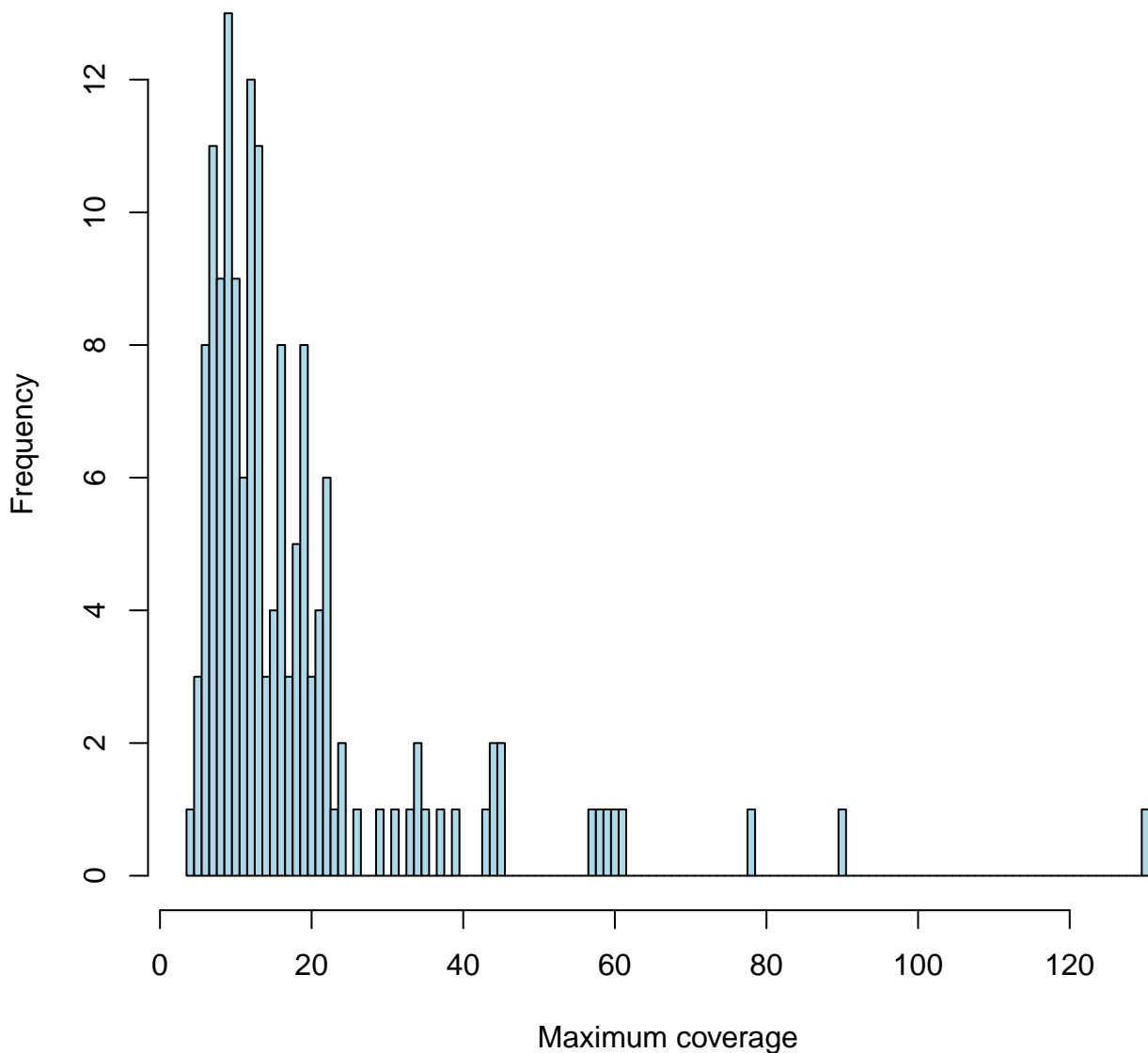
**jx in Intropolis but not "annotated" by knownGene:
119 samples; n jx: 175**



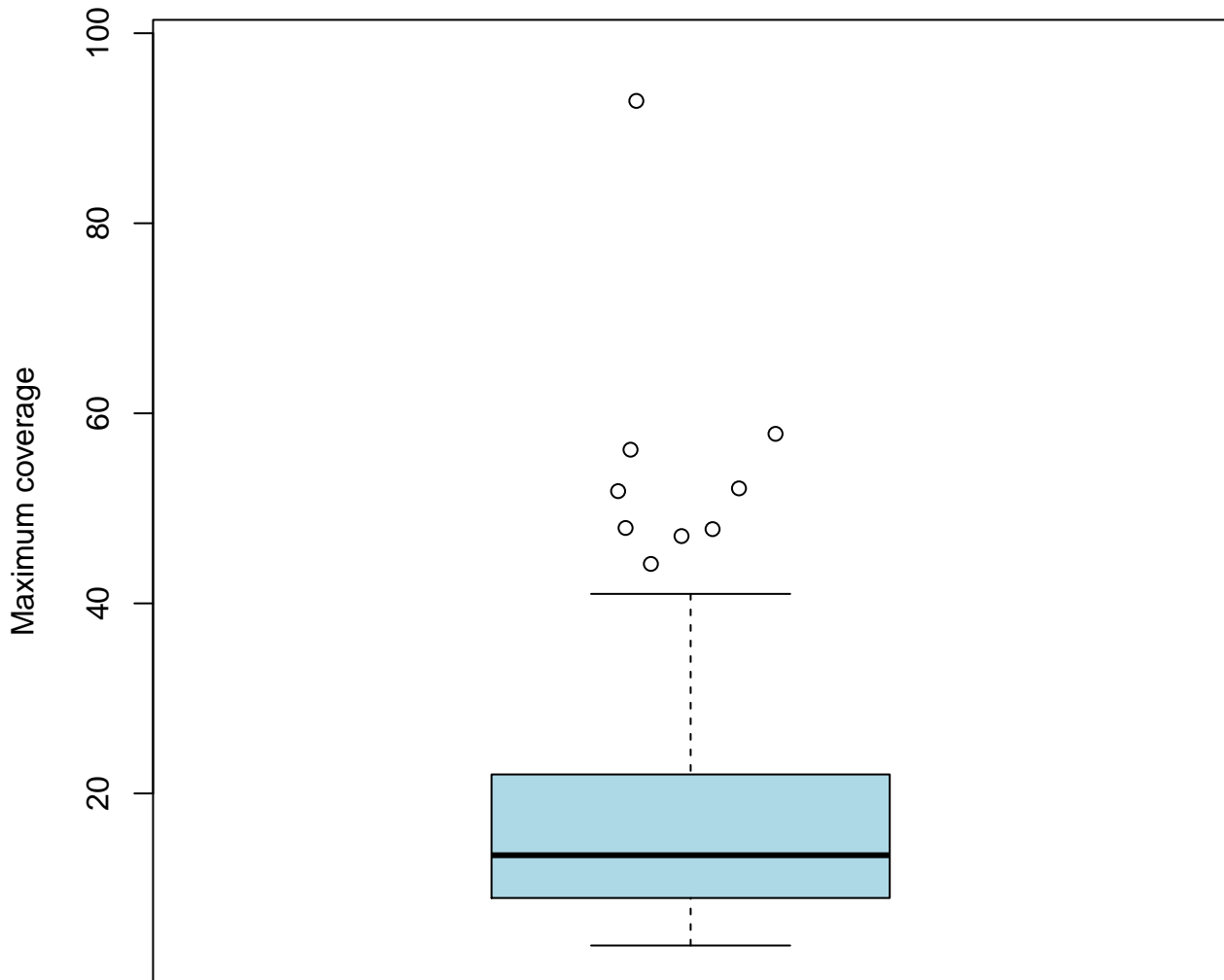
**jx in Intropolis but not "annotated" by knownGene:
120 samples; n jx: 152**



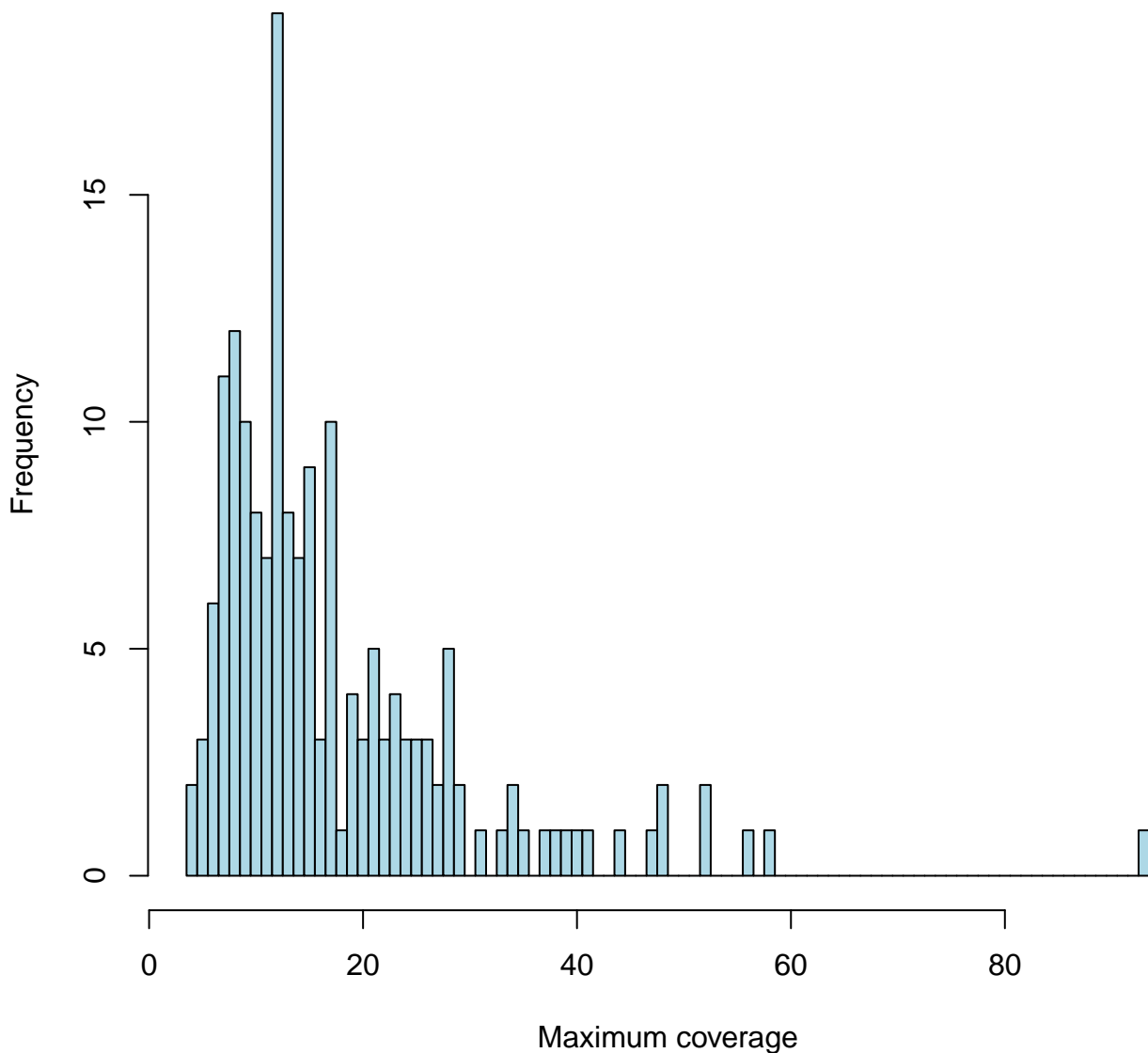
**jx in Intropolis but not "annotated" by knownGene:
120 samples; n jx: 152**



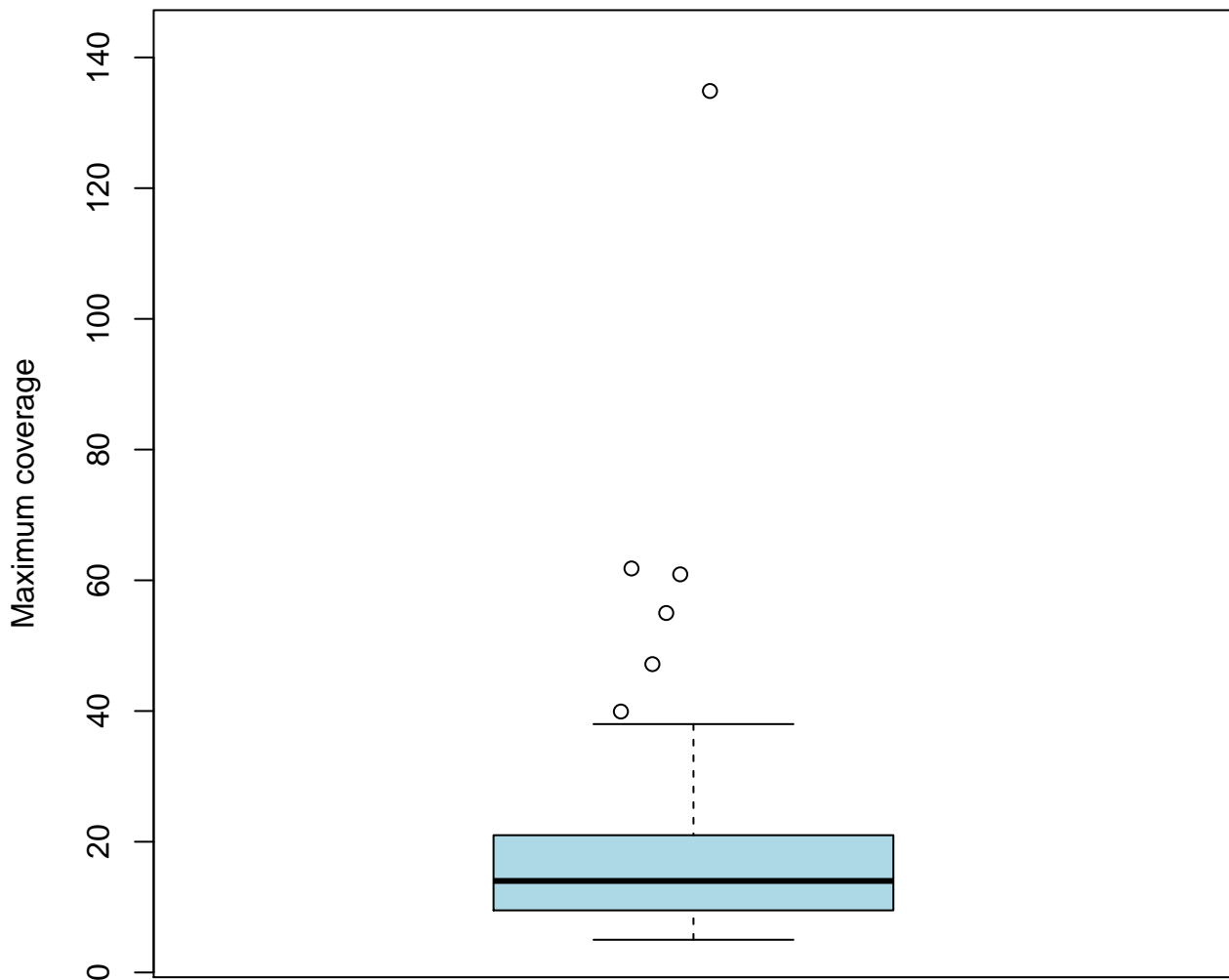
**jx in Intropolis but not "annotated" by knownGene:
121 samples; n jx: 172**



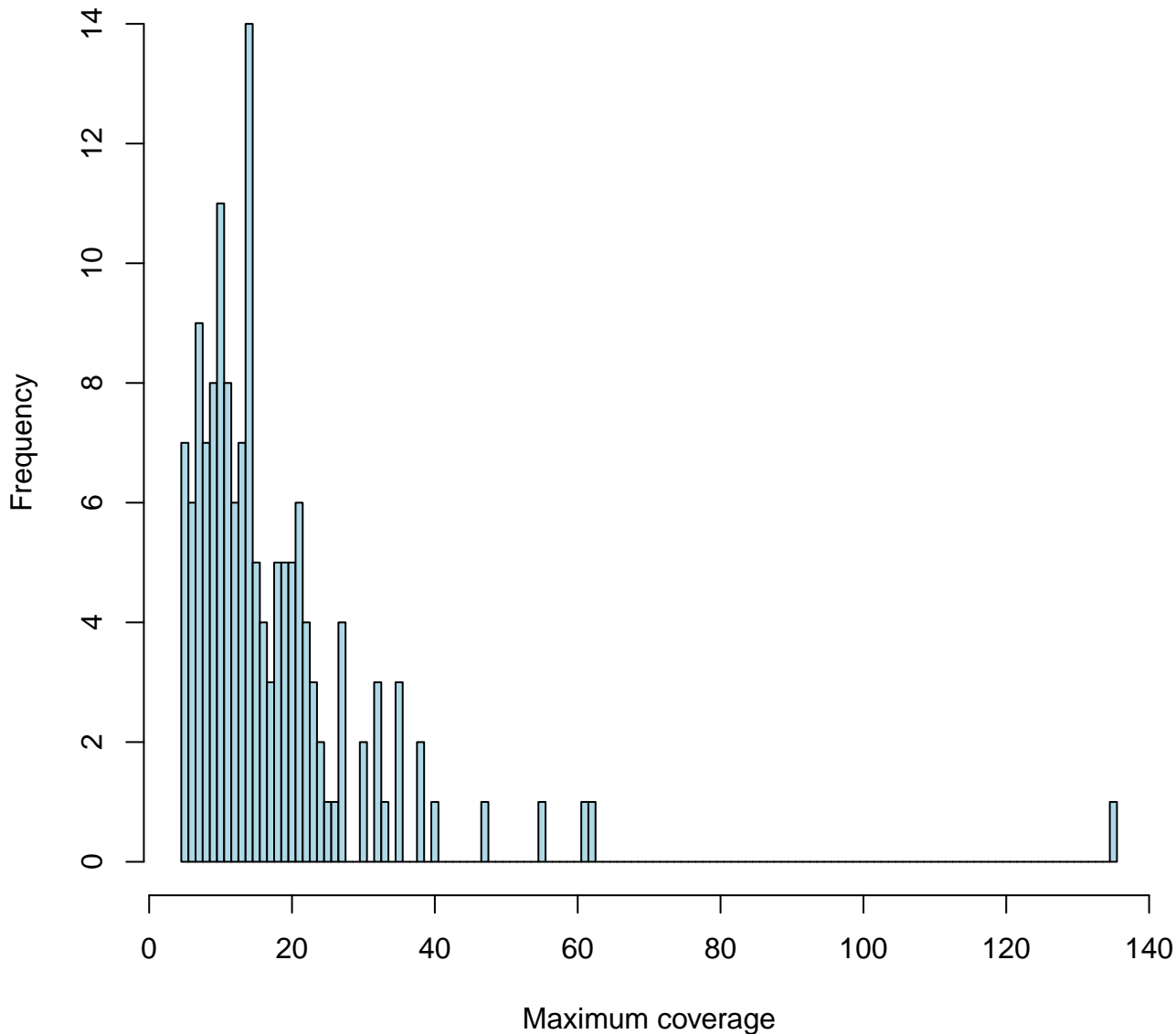
**jx in Intropolis but not "annotated" by knownGene:
121 samples; n jx: 172**



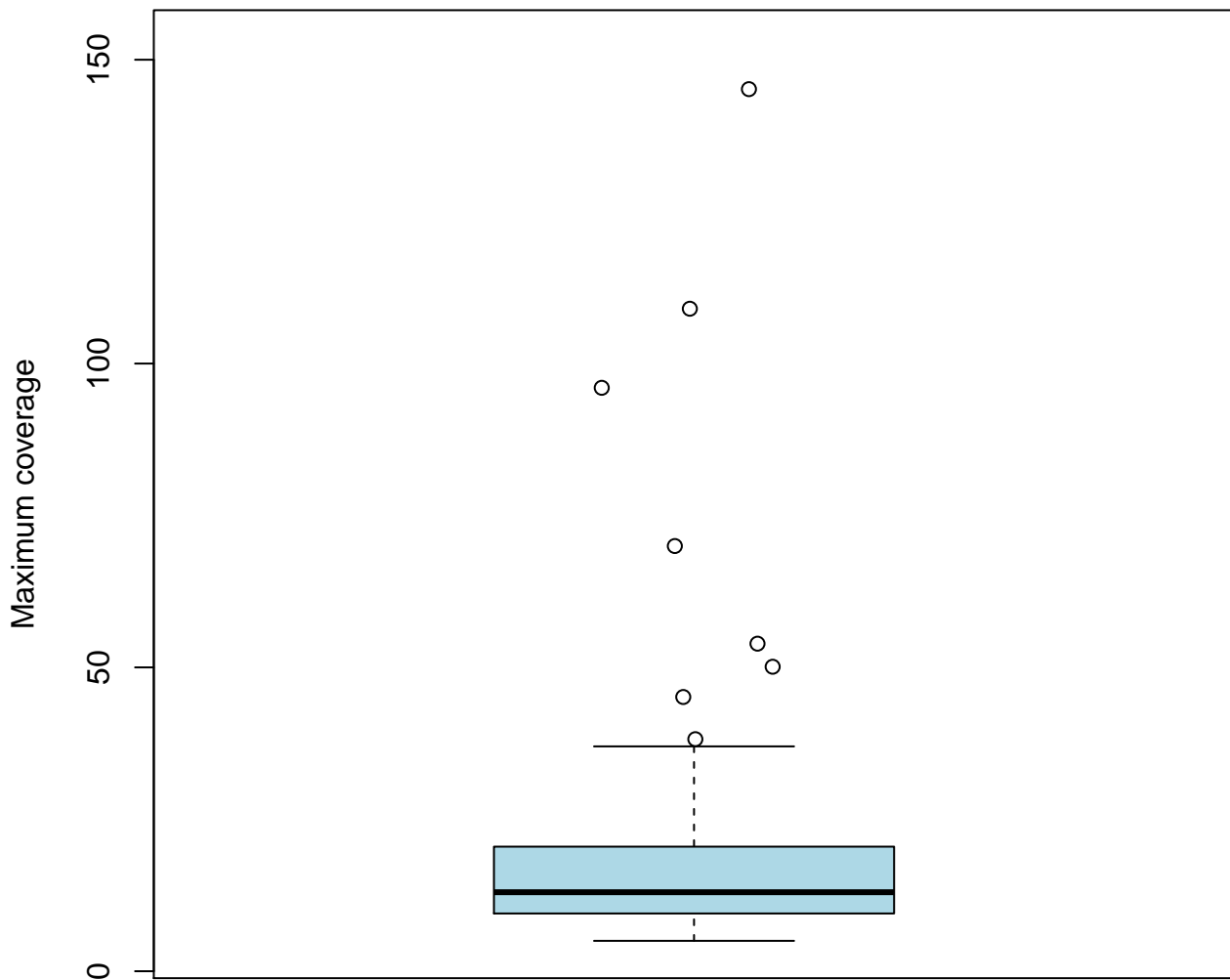
**jx in Intropolis but not "annotated" by knownGene:
122 samples; n jx: 148**



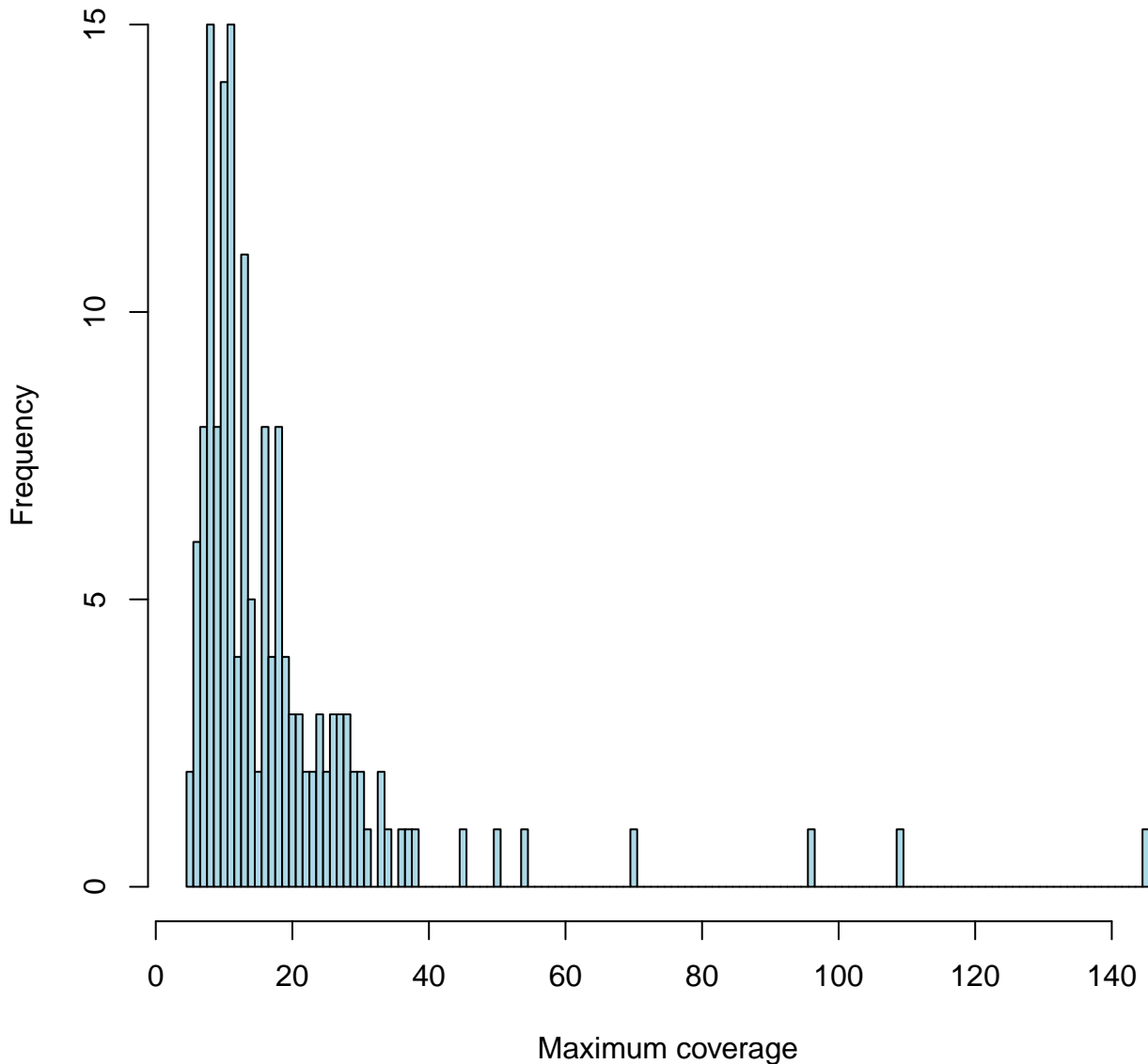
**jx in Intropolis but not "annotated" by knownGene:
122 samples; n jx: 148**



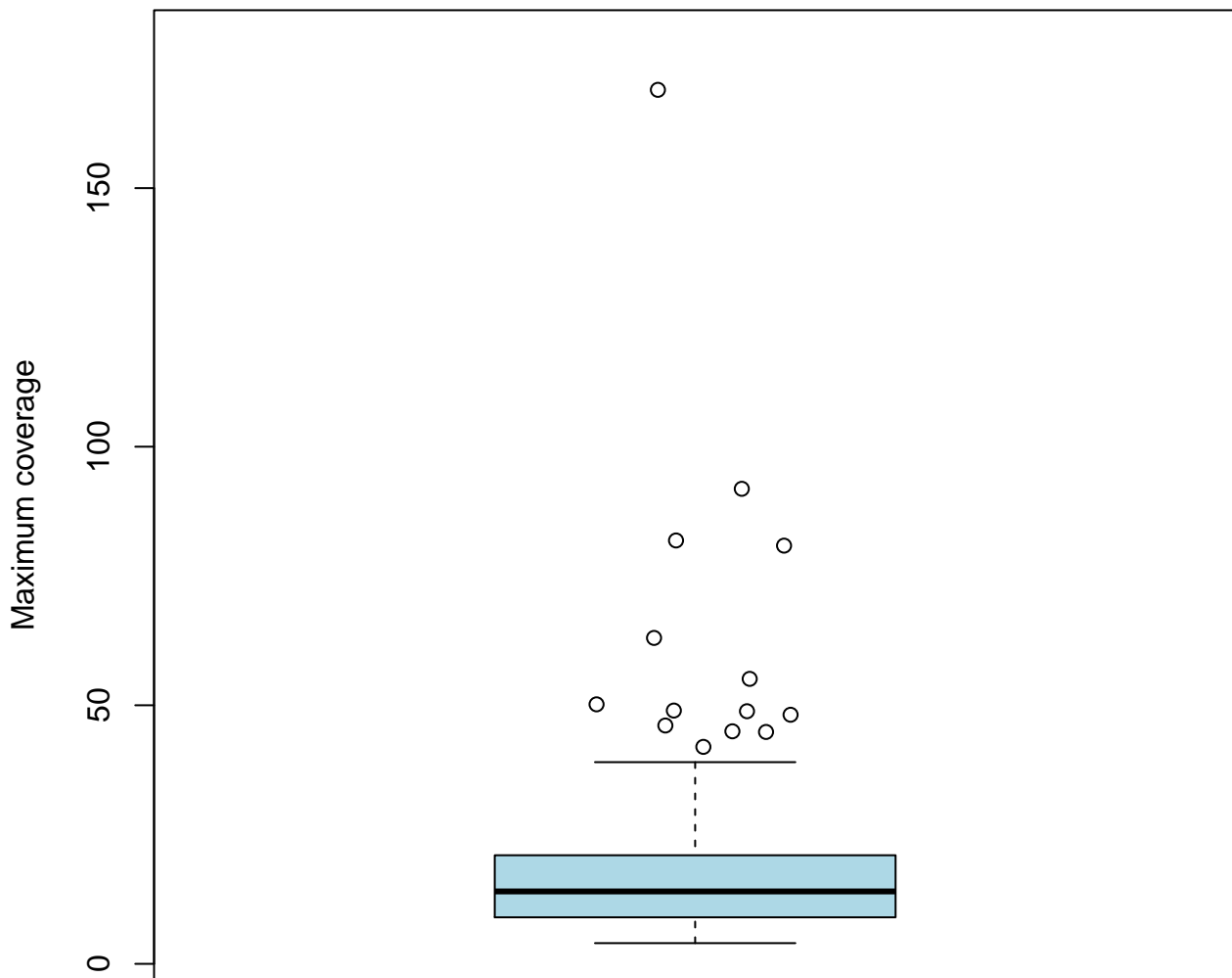
**jx in Intropolis but not "annotated" by knownGene:
123 samples; n jx: 156**



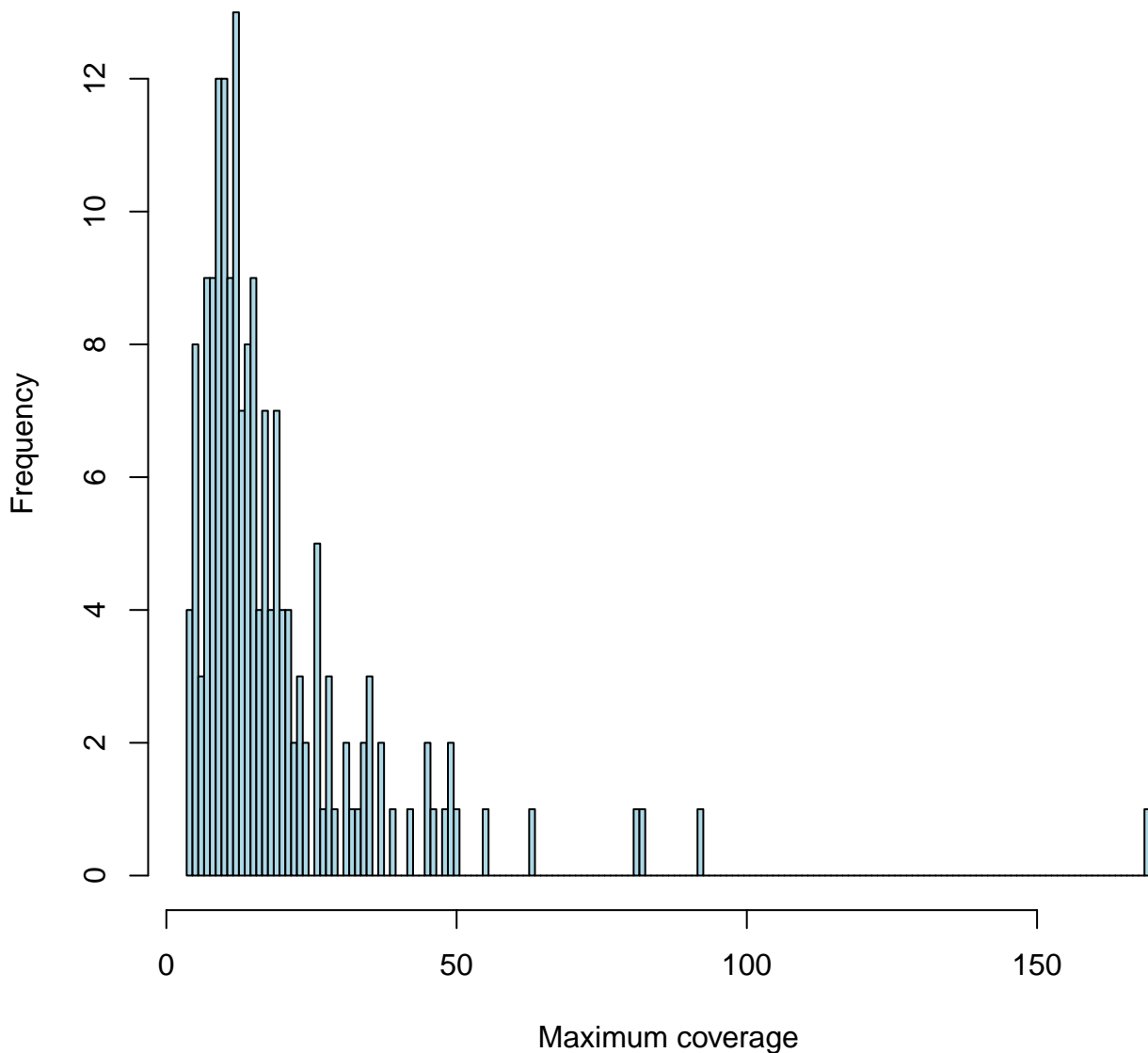
**jx in Intropolis but not "annotated" by knownGene:
123 samples; n jx: 156**



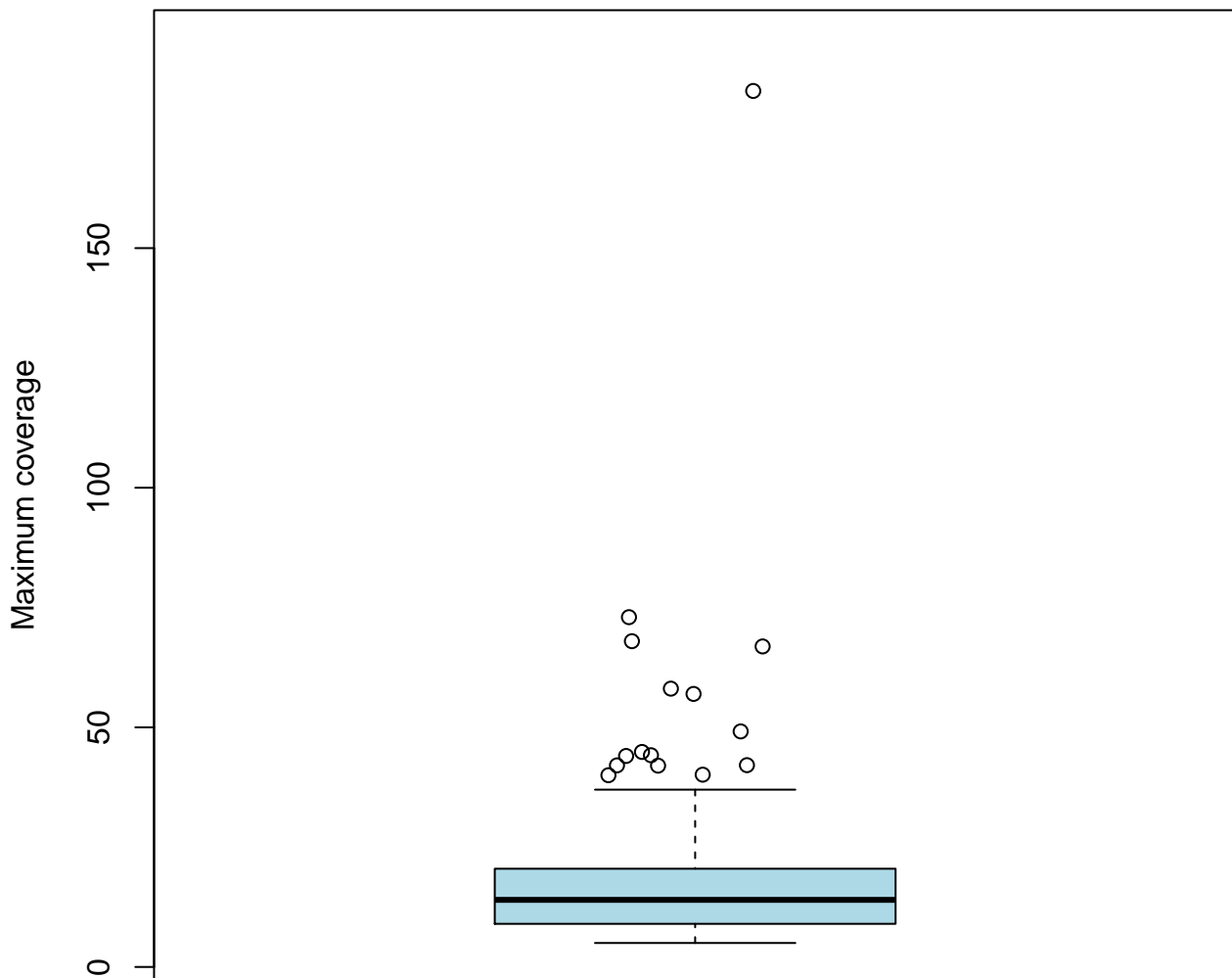
**jx in Intropolis but not "annotated" by knownGene:
124 samples; n jx: 176**



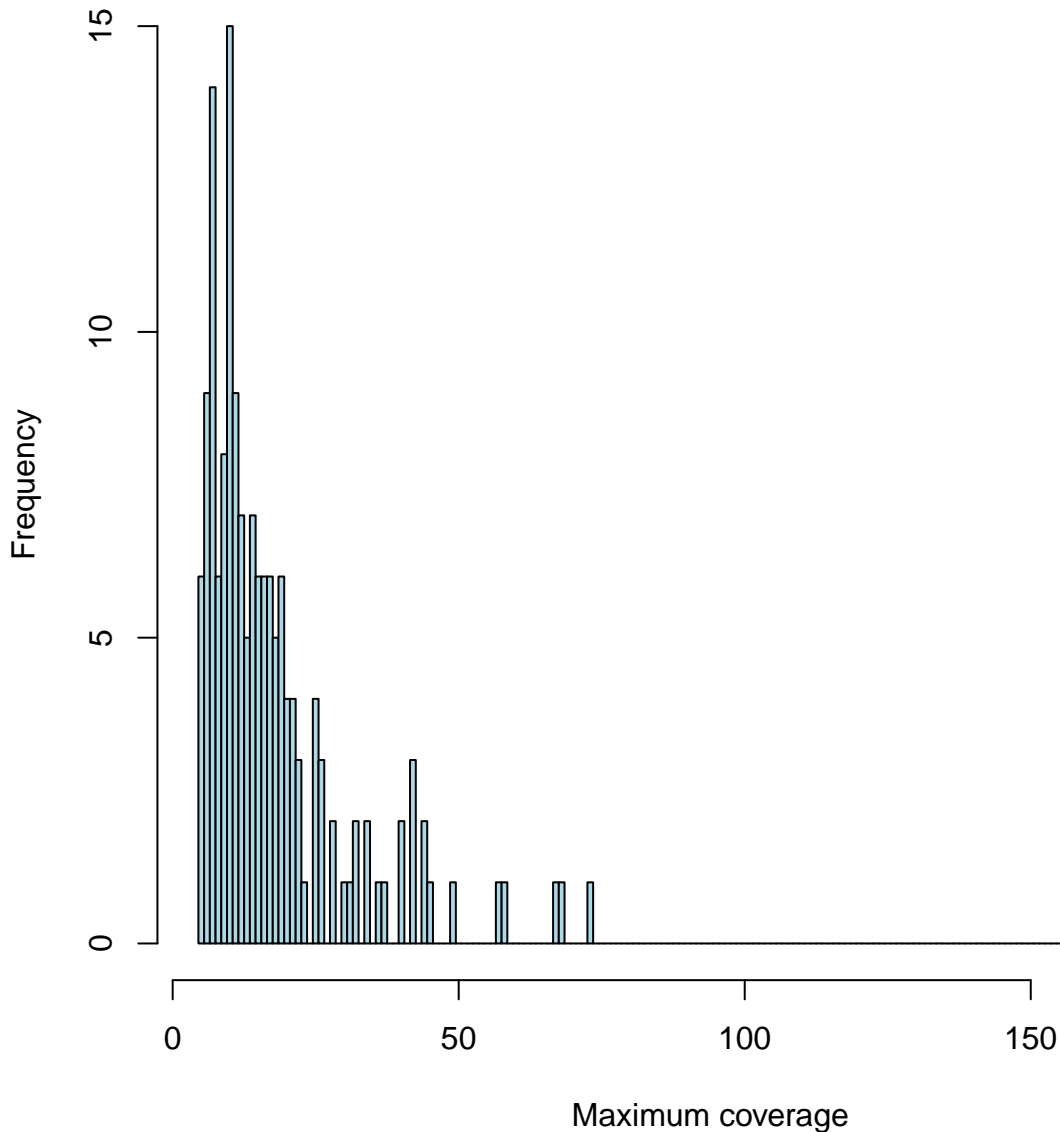
**jx in Intropolis but not "annotated" by knownGene:
124 samples; n jx: 176**



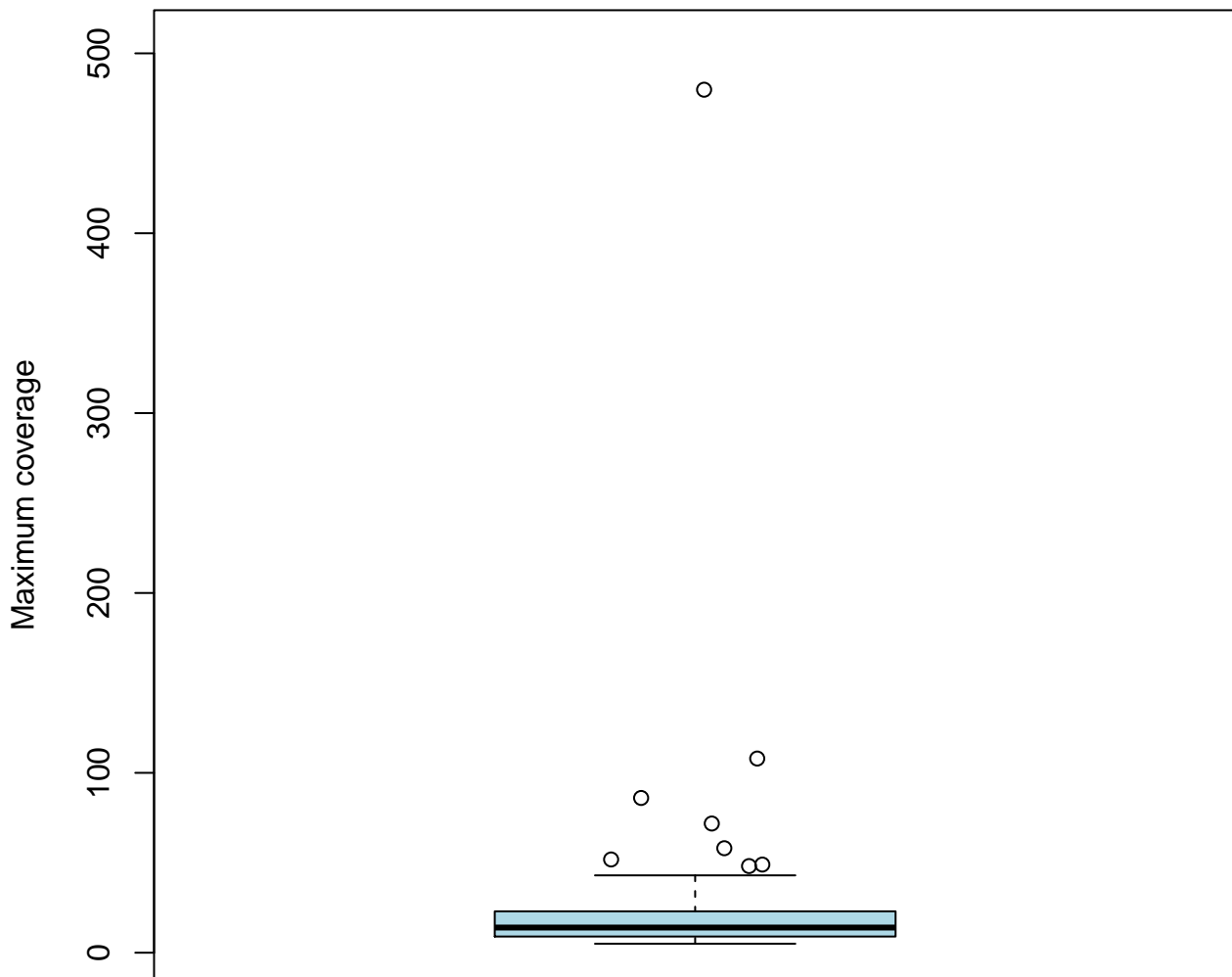
**jx in Intropolis but not "annotated" by knownGene:
125 samples; n jx: 159**



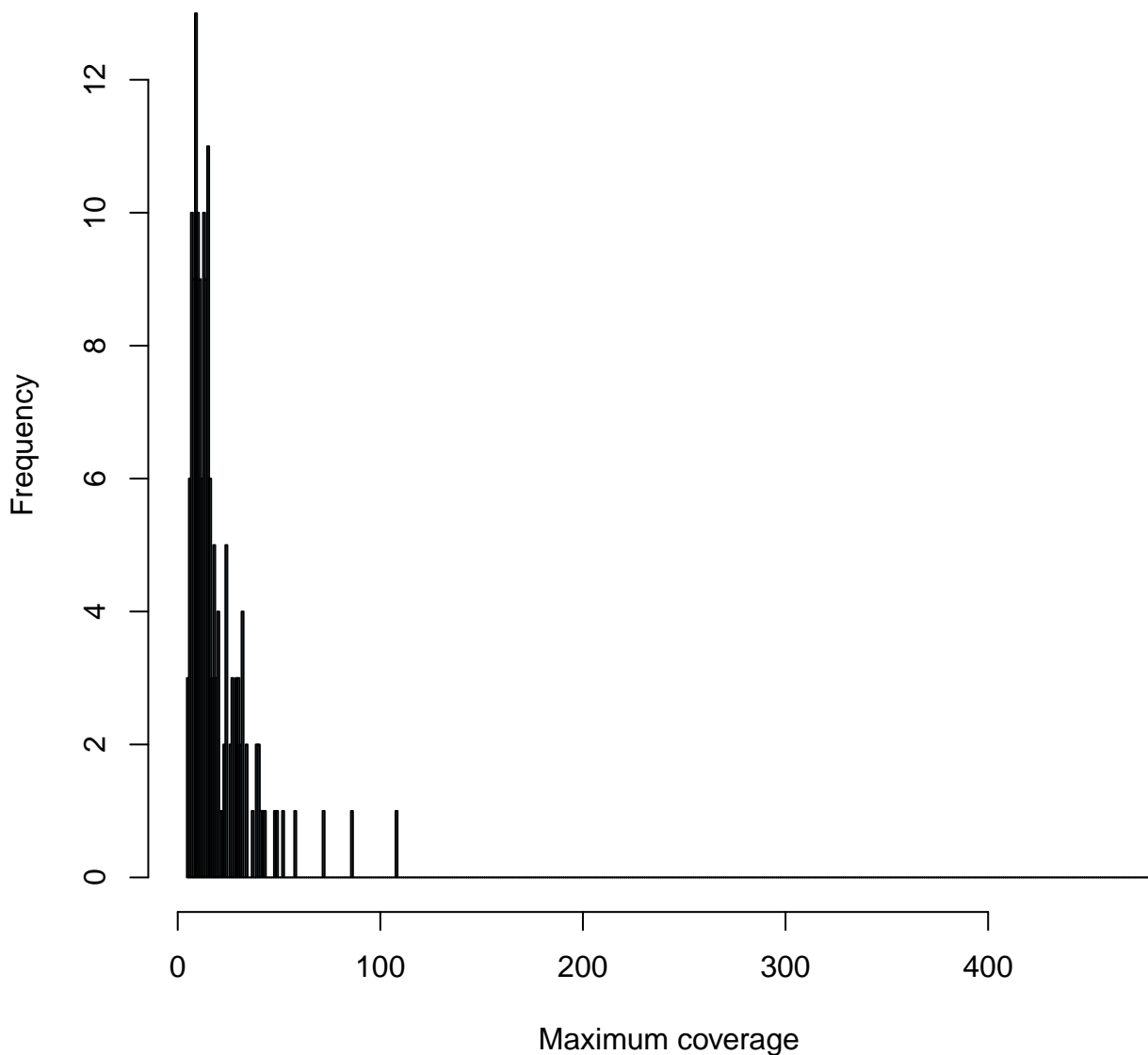
**jx in Intropolis but not "annotated" by knownGene:
125 samples; n jx: 159**



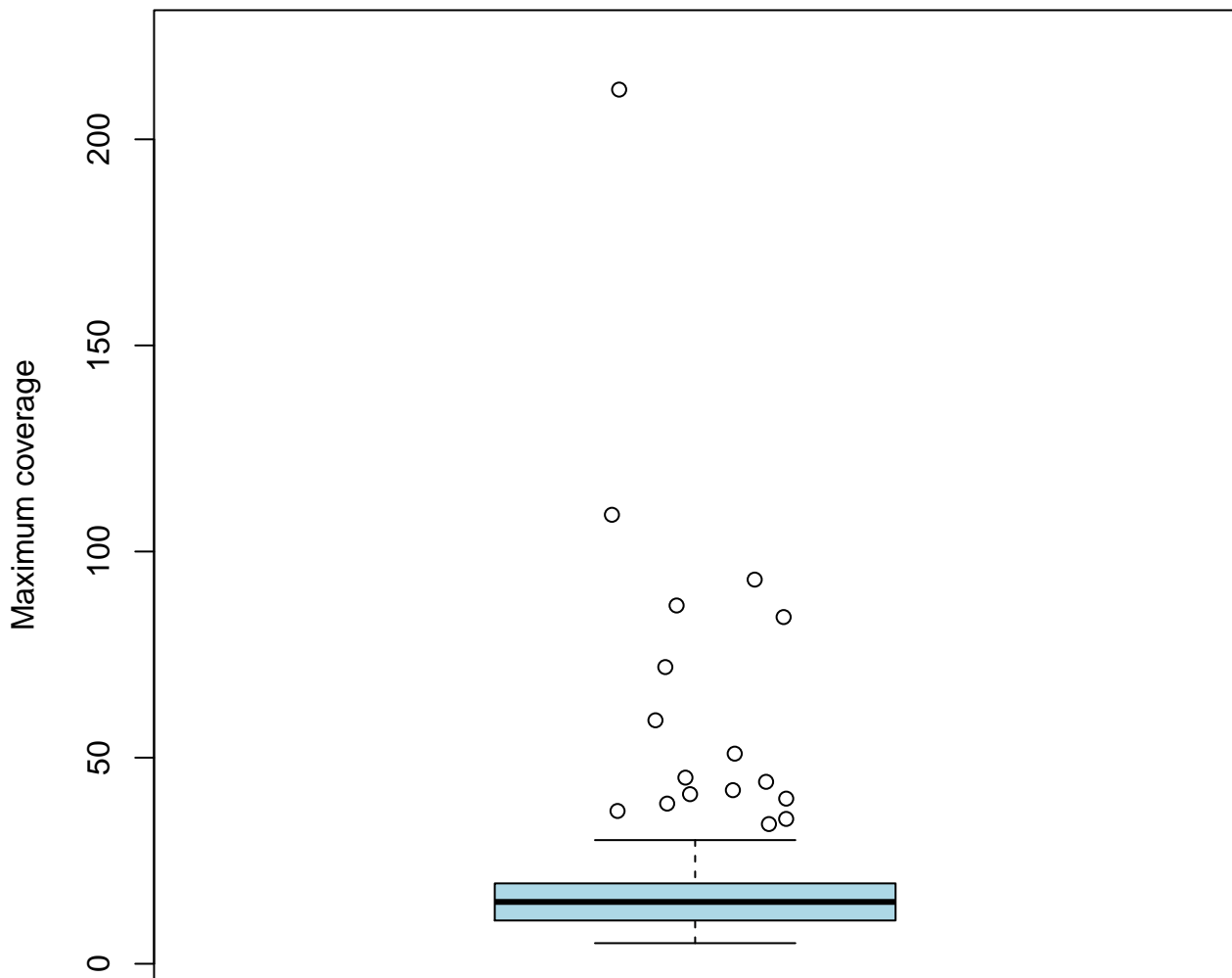
**jx in Intropolis but not "annotated" by knownGene:
126 samples; n jx: 161**



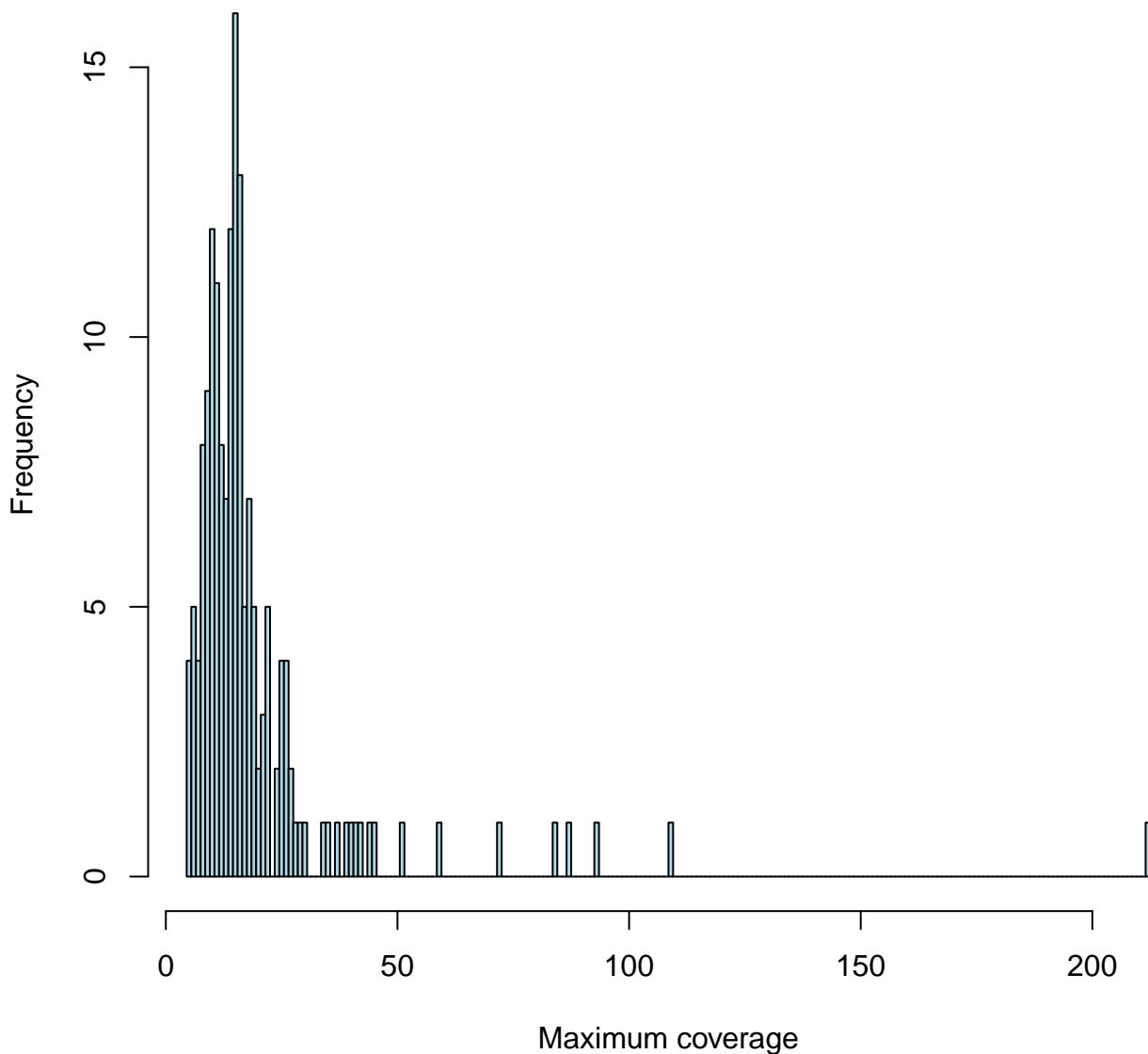
**jx in Intropolis but not "annotated" by knownGene:
126 samples; n jx: 161**



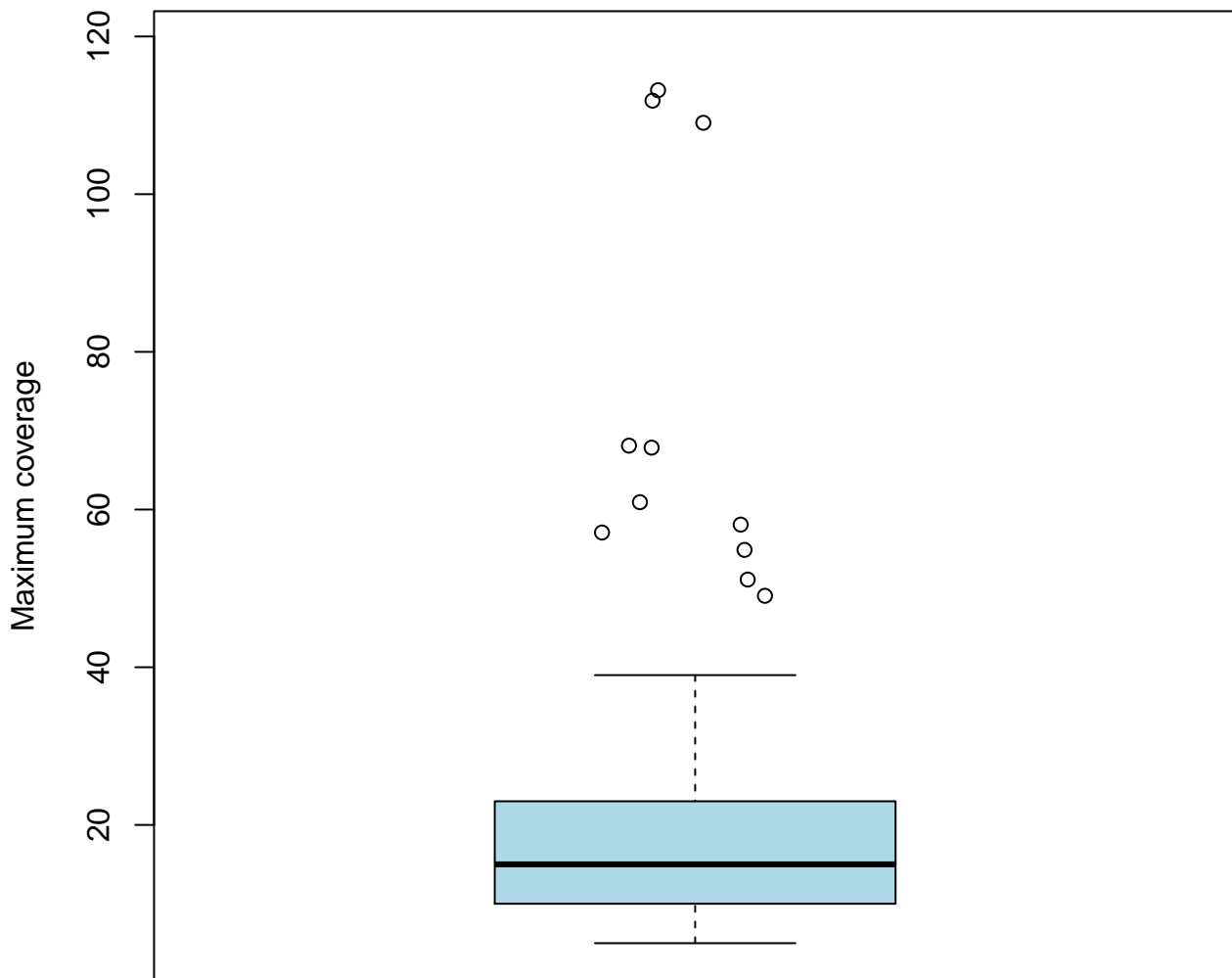
**jx in Intropolis but not "annotated" by knownGene:
127 samples; n jx: 168**



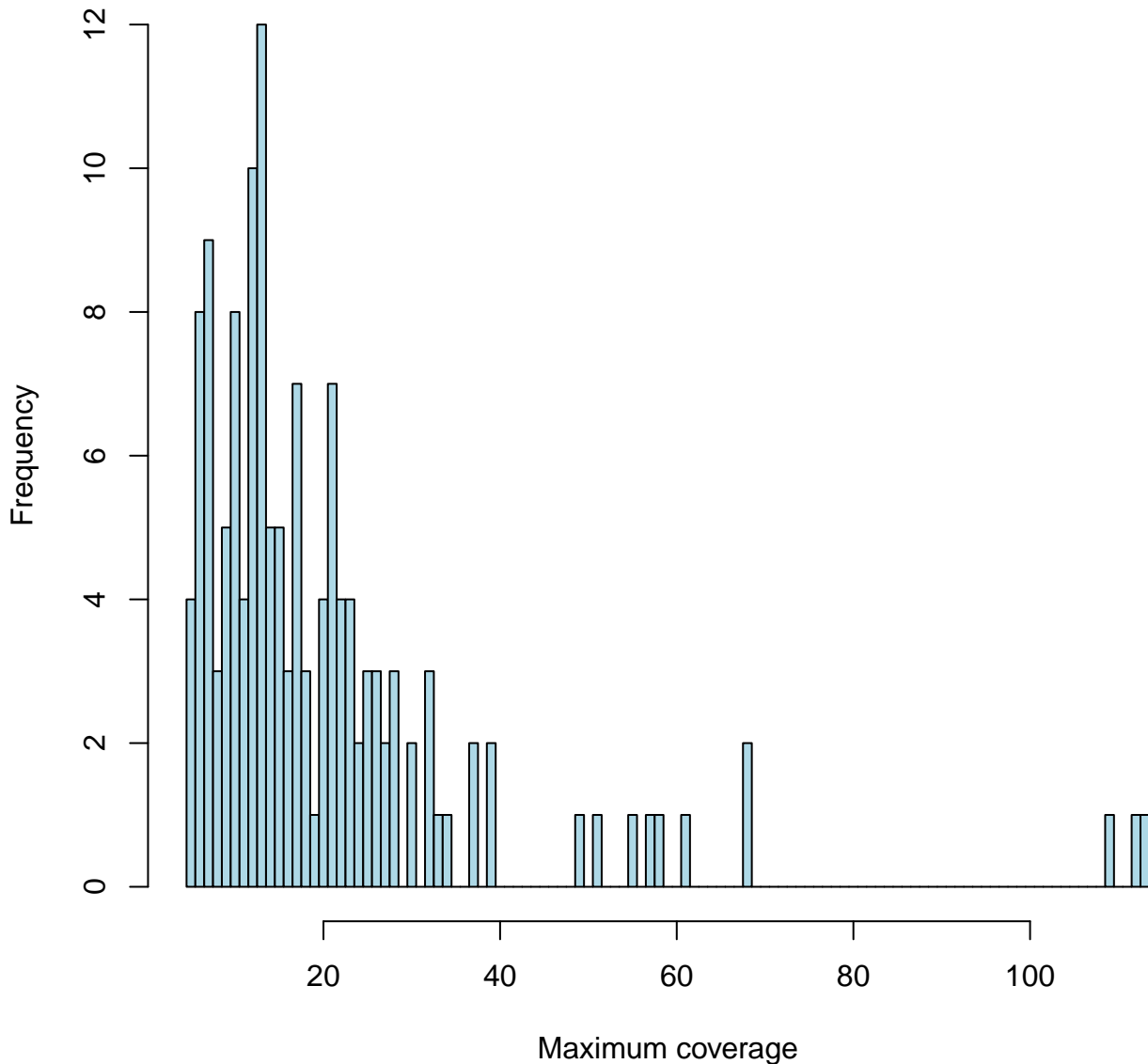
**jx in Intropolis but not "annotated" by knownGene:
127 samples; n jx: 168**



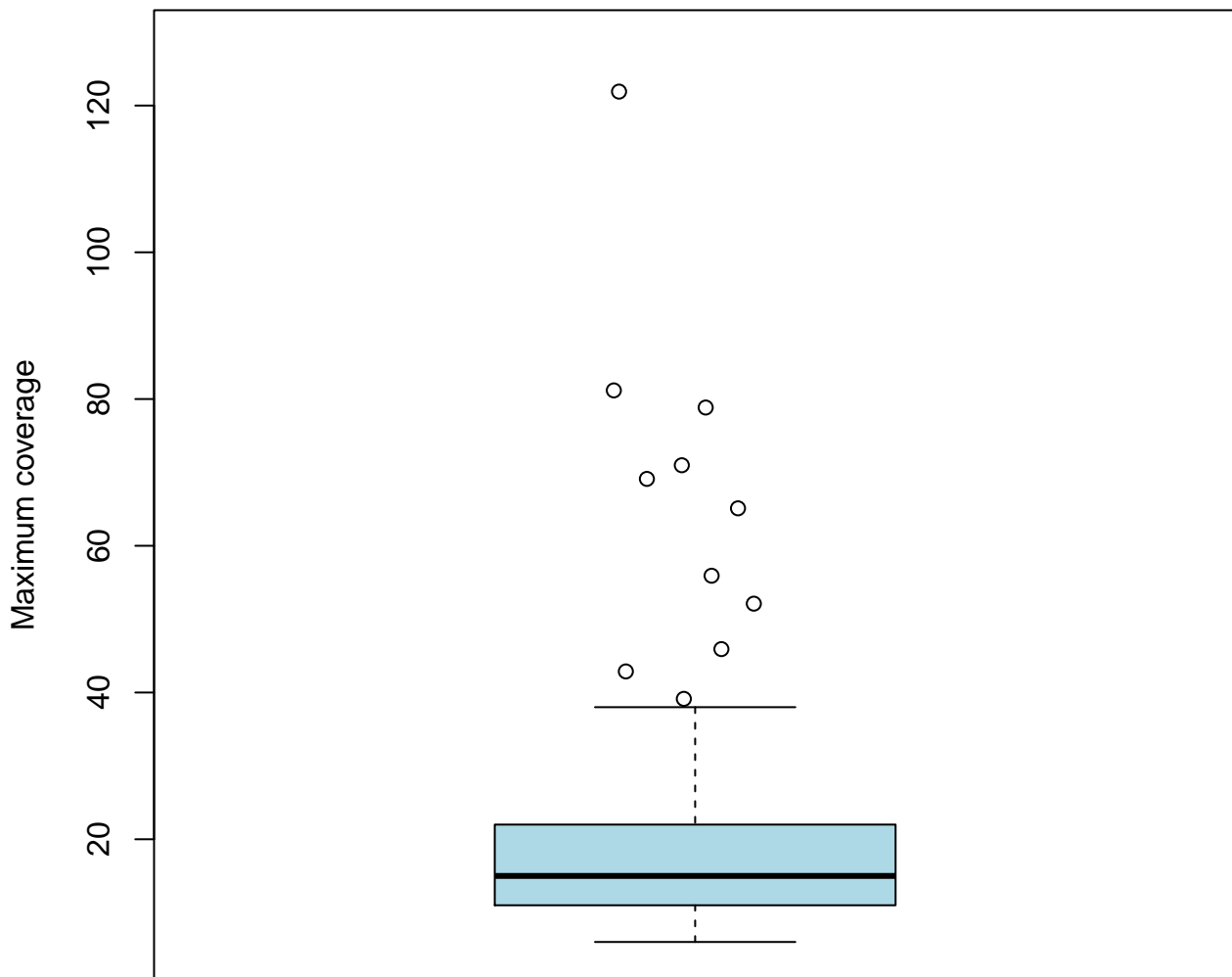
**jx in Intropolis but not "annotated" by knownGene:
128 samples; n jx: 141**



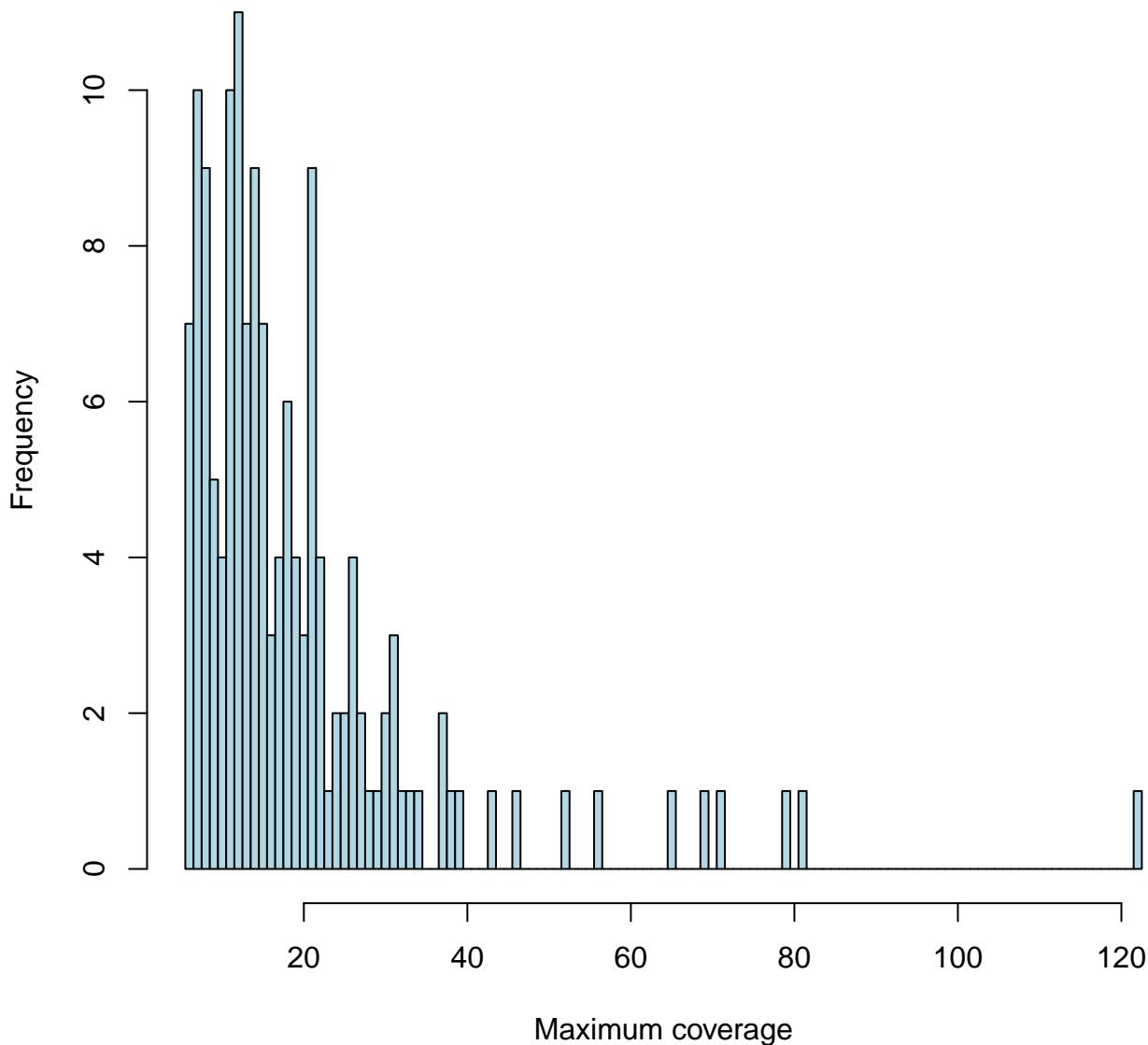
**jx in Intropolis but not "annotated" by knownGene:
128 samples; n jx: 141**



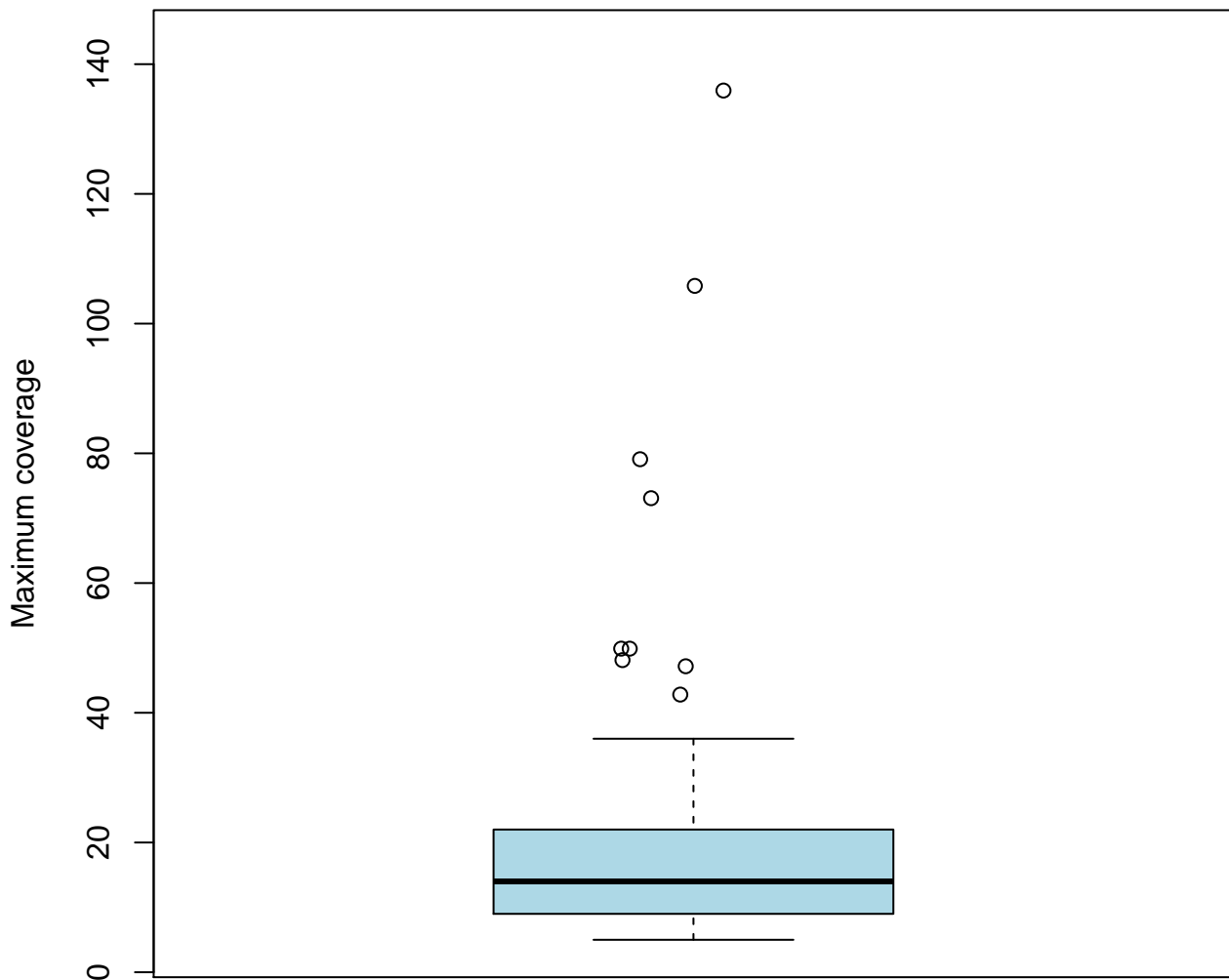
**jx in Intropolis but not "annotated" by knownGene:
129 samples; n jx: 147**



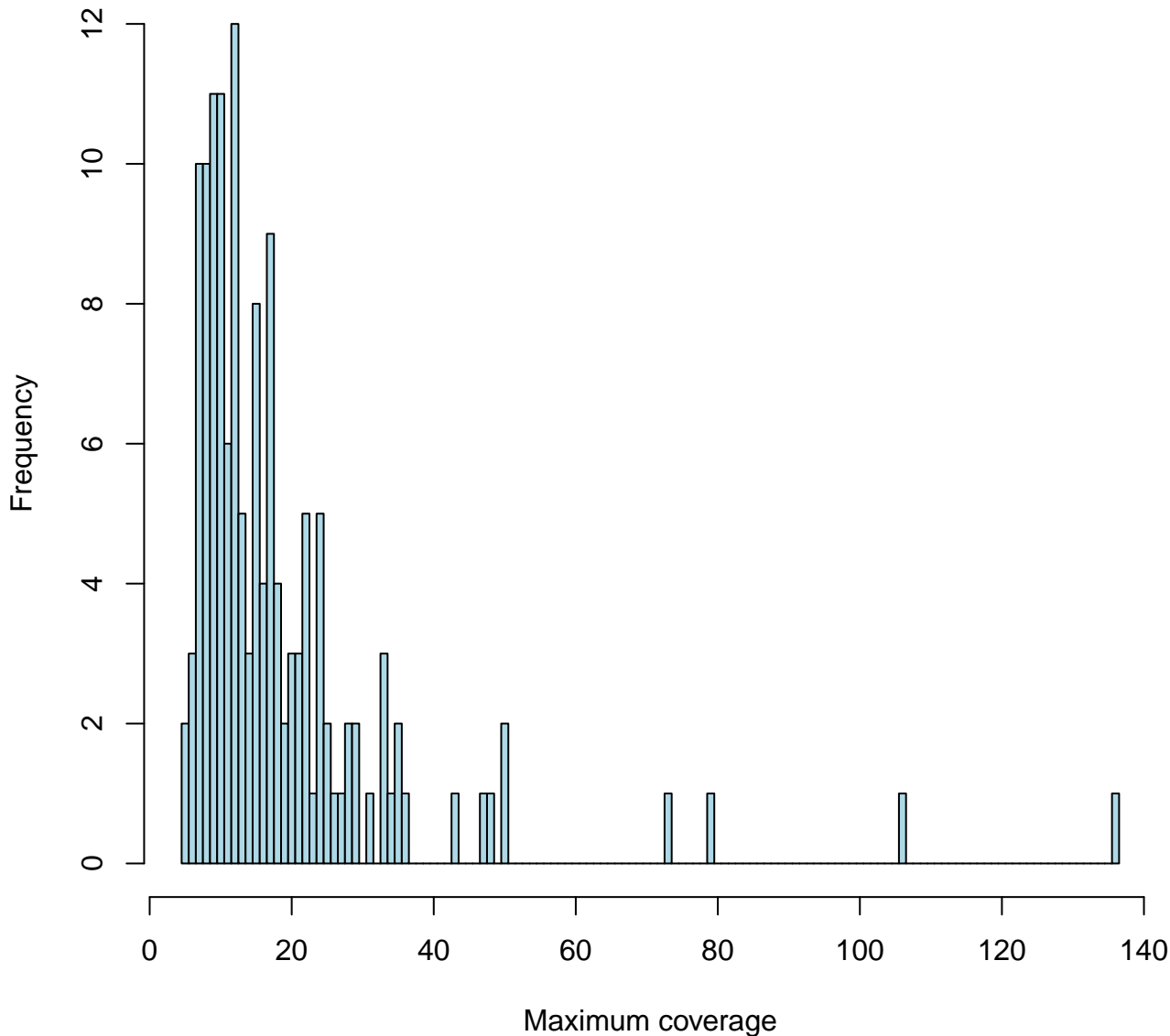
**jx in Intropolis but not "annotated" by knownGene:
129 samples; n jx: 147**



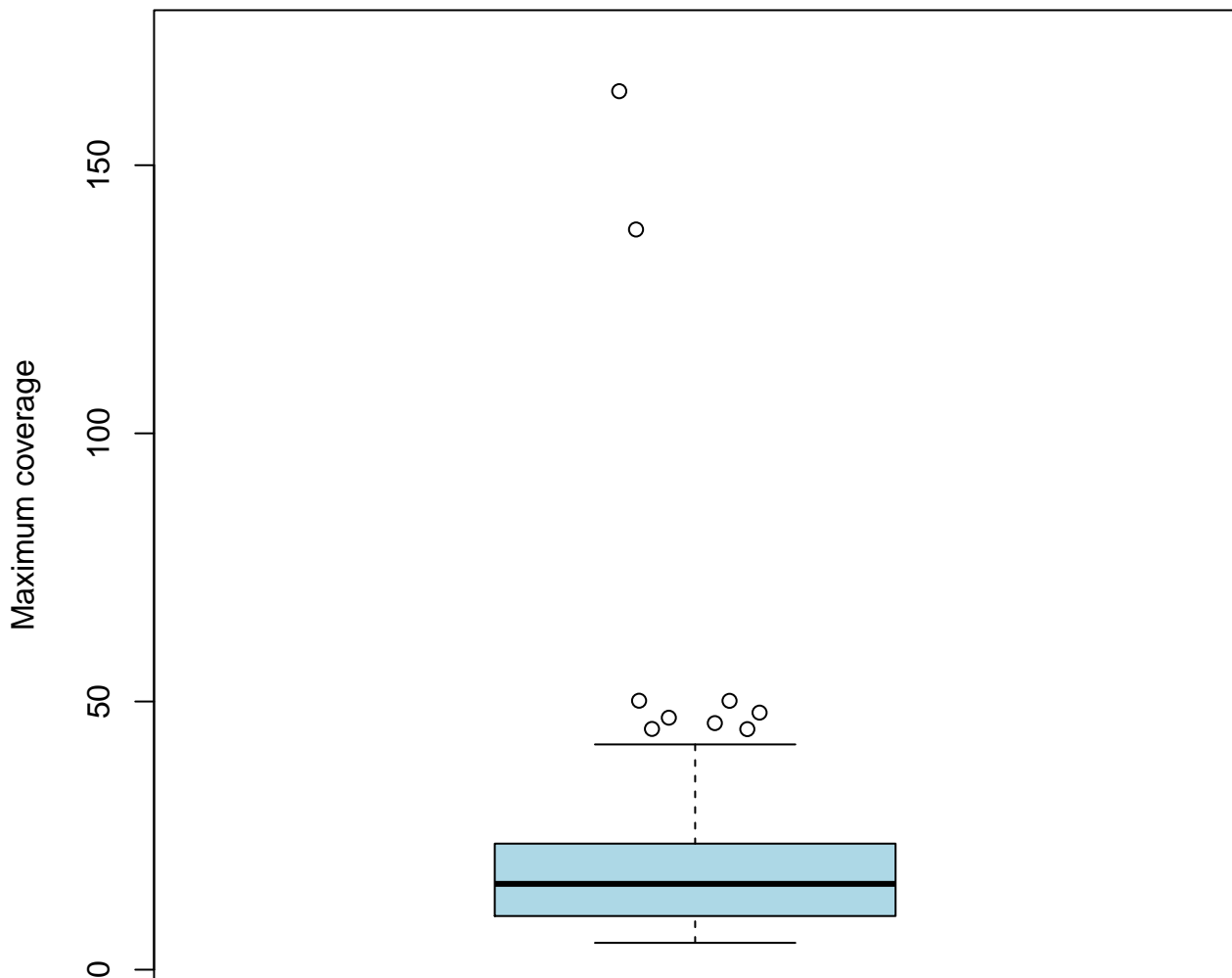
**jx in Intropolis but not "annotated" by knownGene:
130 samples; n jx: 142**



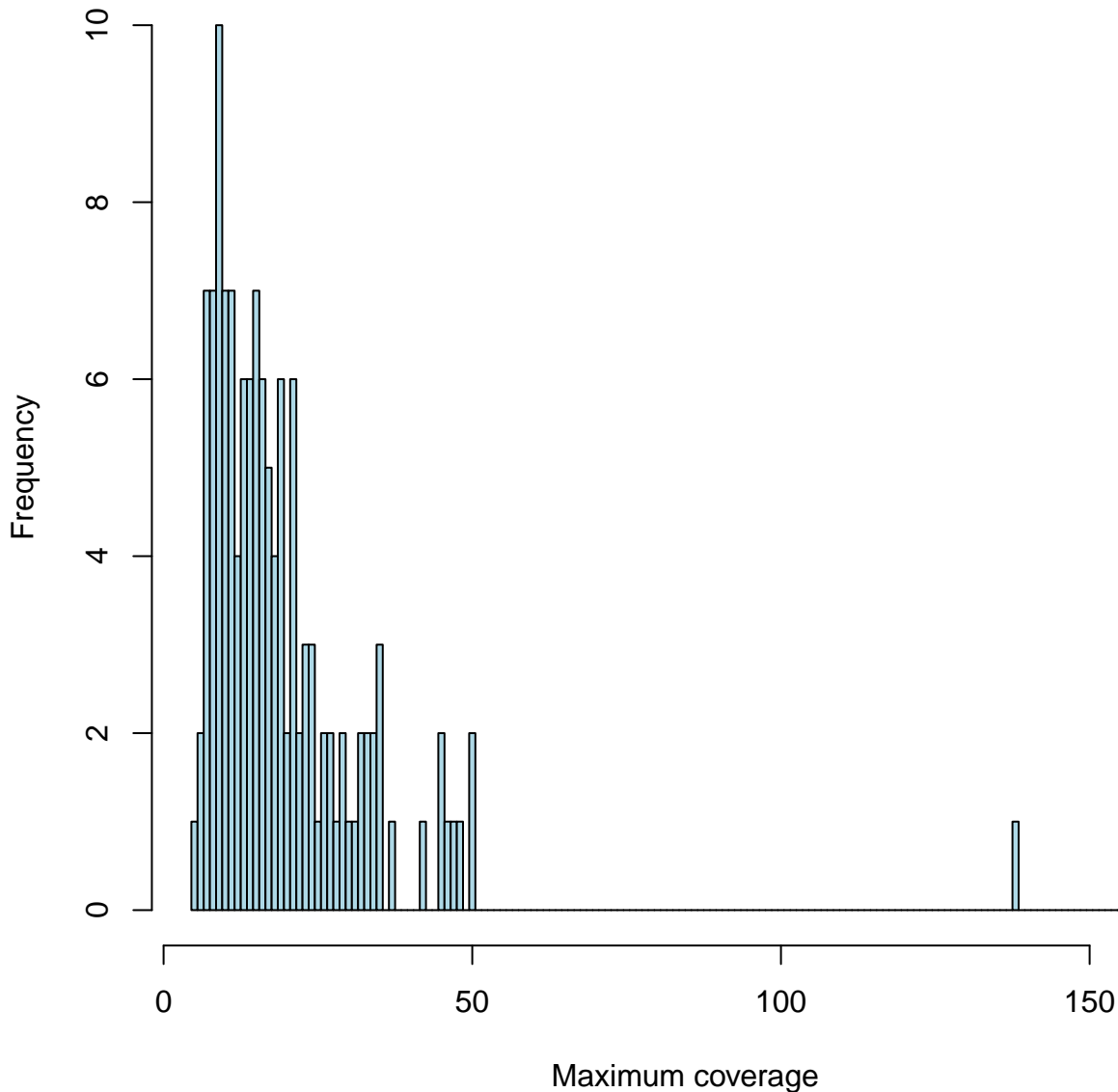
**jx in Intropolis but not "annotated" by knownGene:
130 samples; n jx: 142**



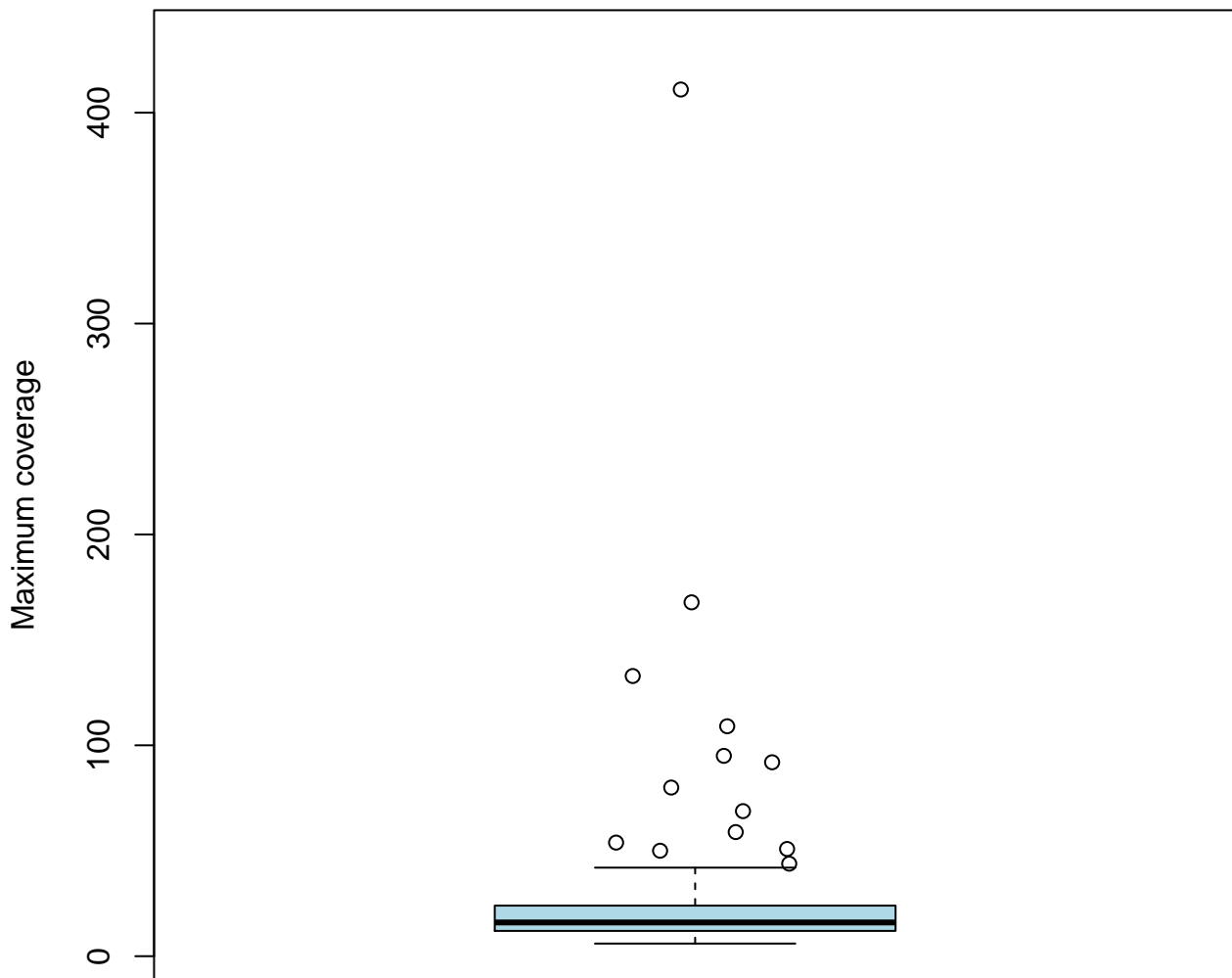
**jx in Intropolis but not "annotated" by knownGene:
131 samples; n jx: 131**



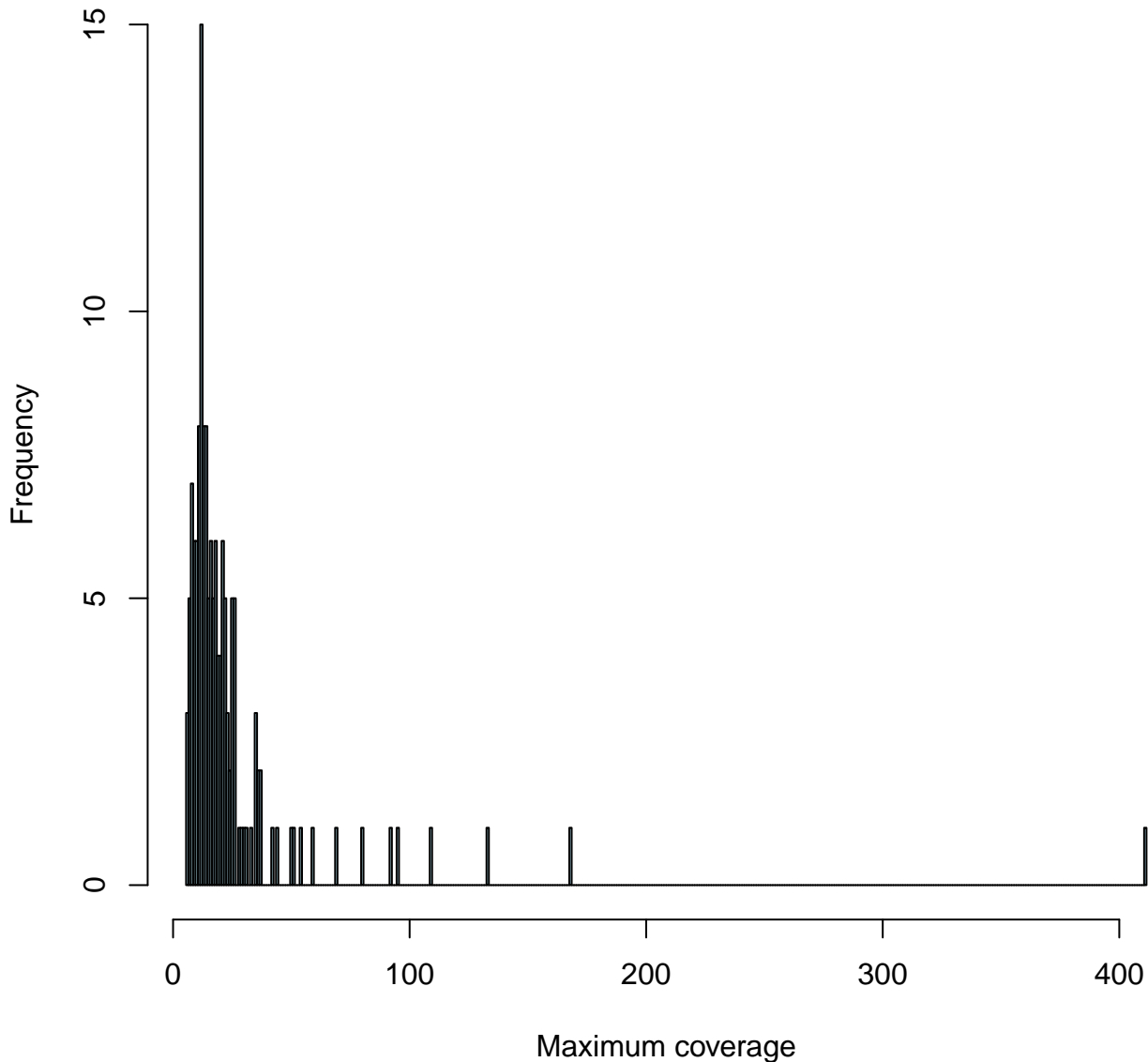
**jx in Intropolis but not "annotated" by knownGene:
131 samples; n jx: 131**



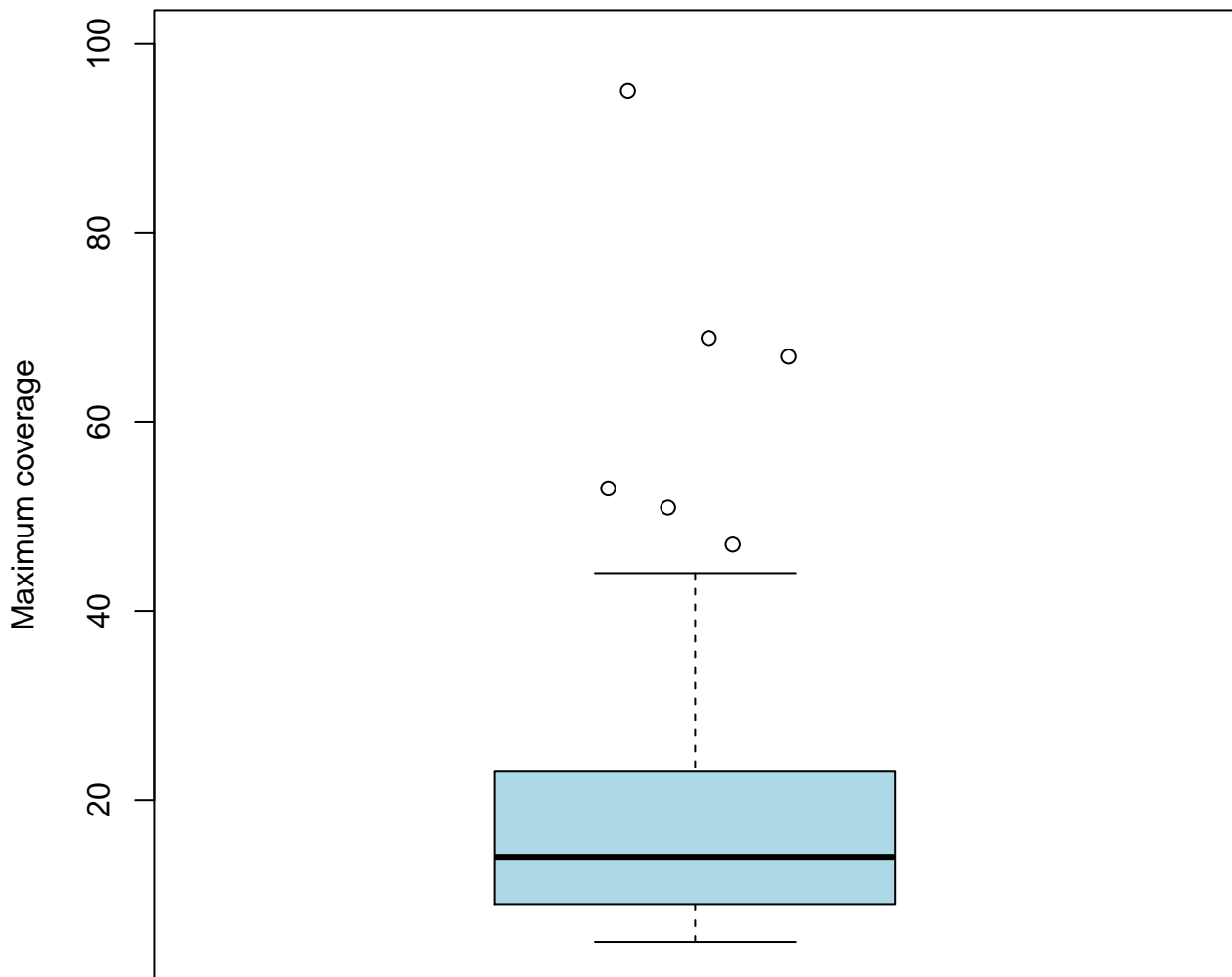
**jx in Intropolis but not "annotated" by knownGene:
132 samples; n jx: 148**



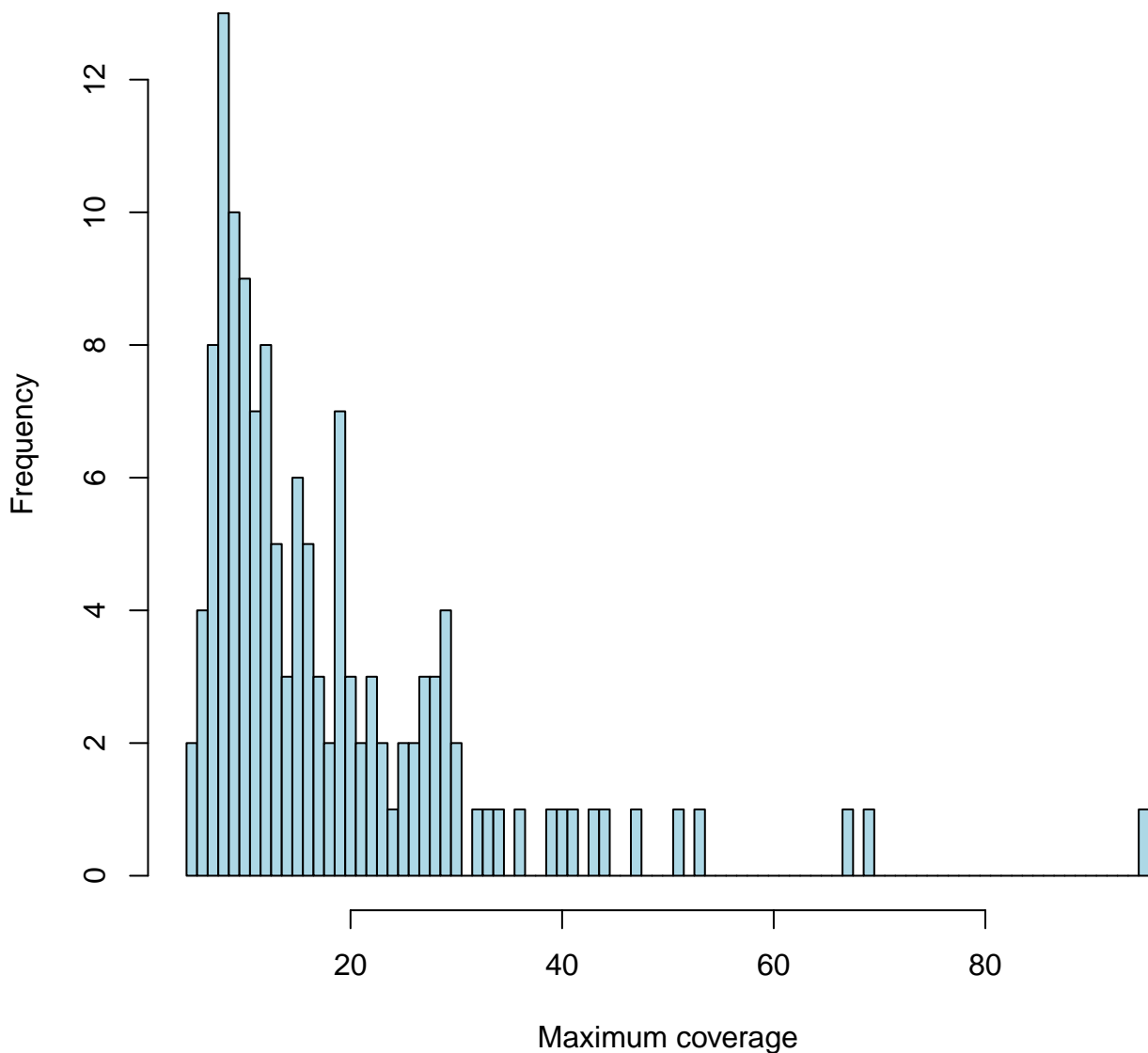
**jx in Intropolis but not "annotated" by knownGene:
132 samples; n jx: 148**



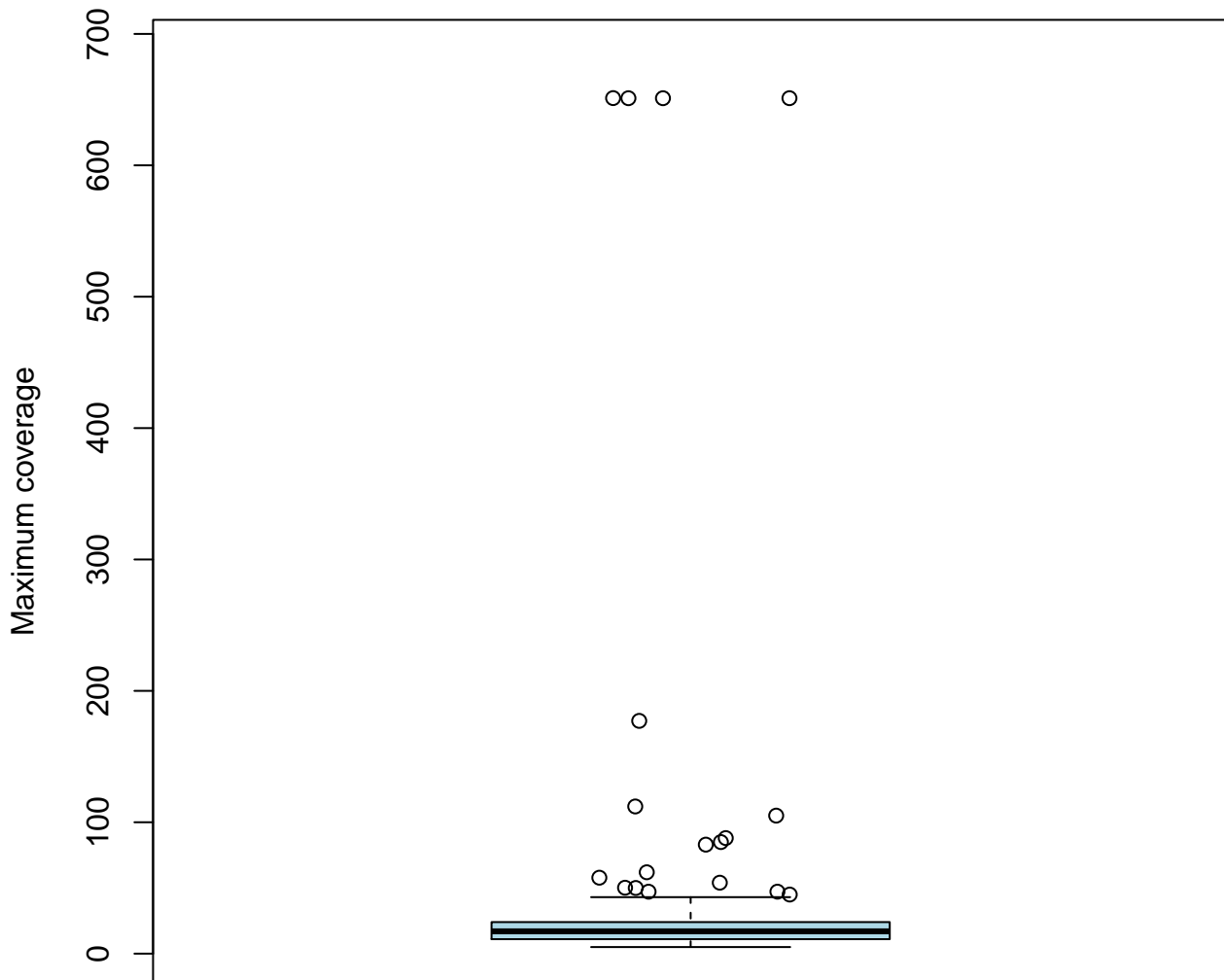
**jx in Intropolis but not "annotated" by knownGene:
133 samples; n jx: 134**



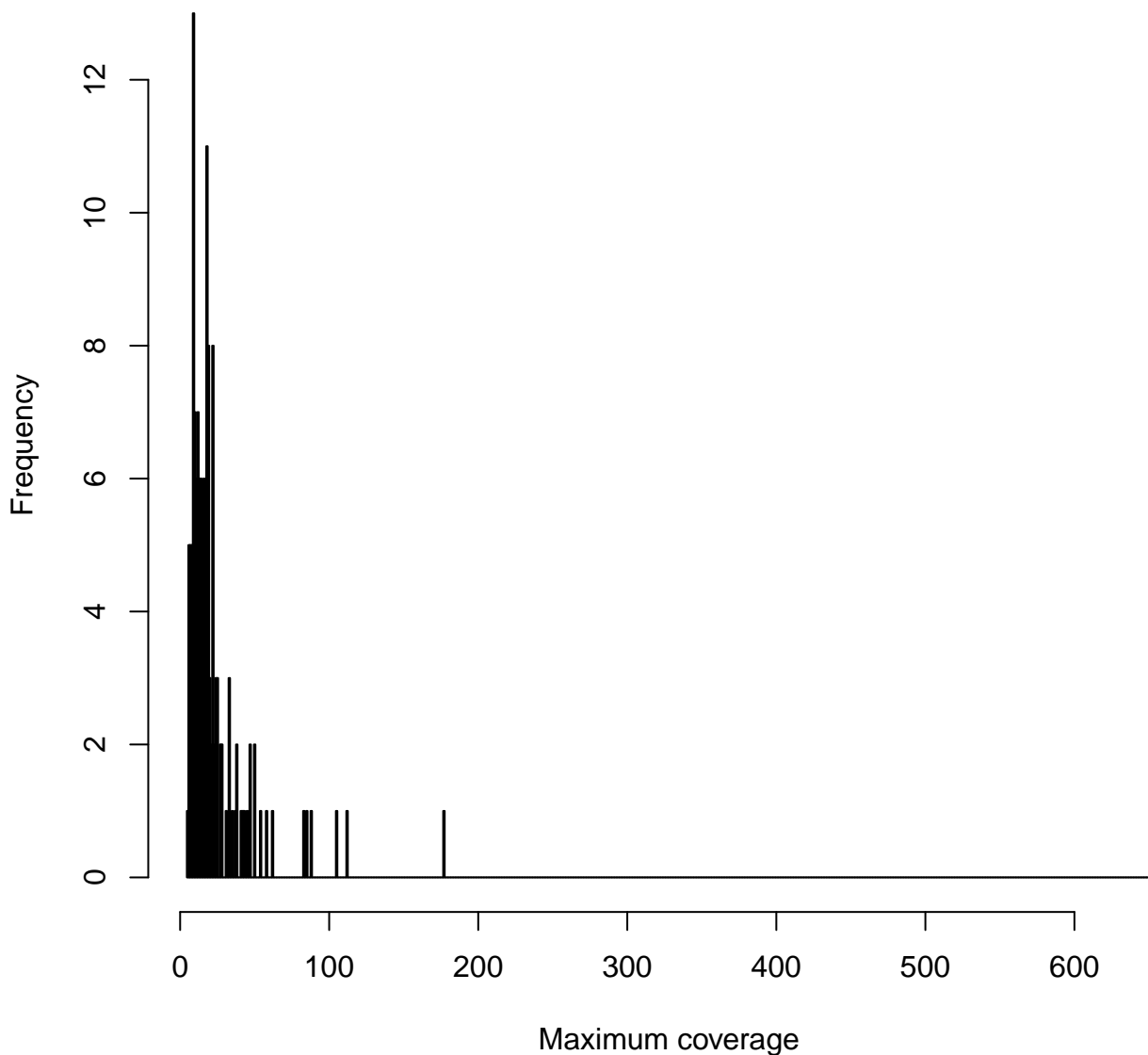
**jx in Intropolis but not "annotated" by knownGene:
133 samples; n jx: 134**



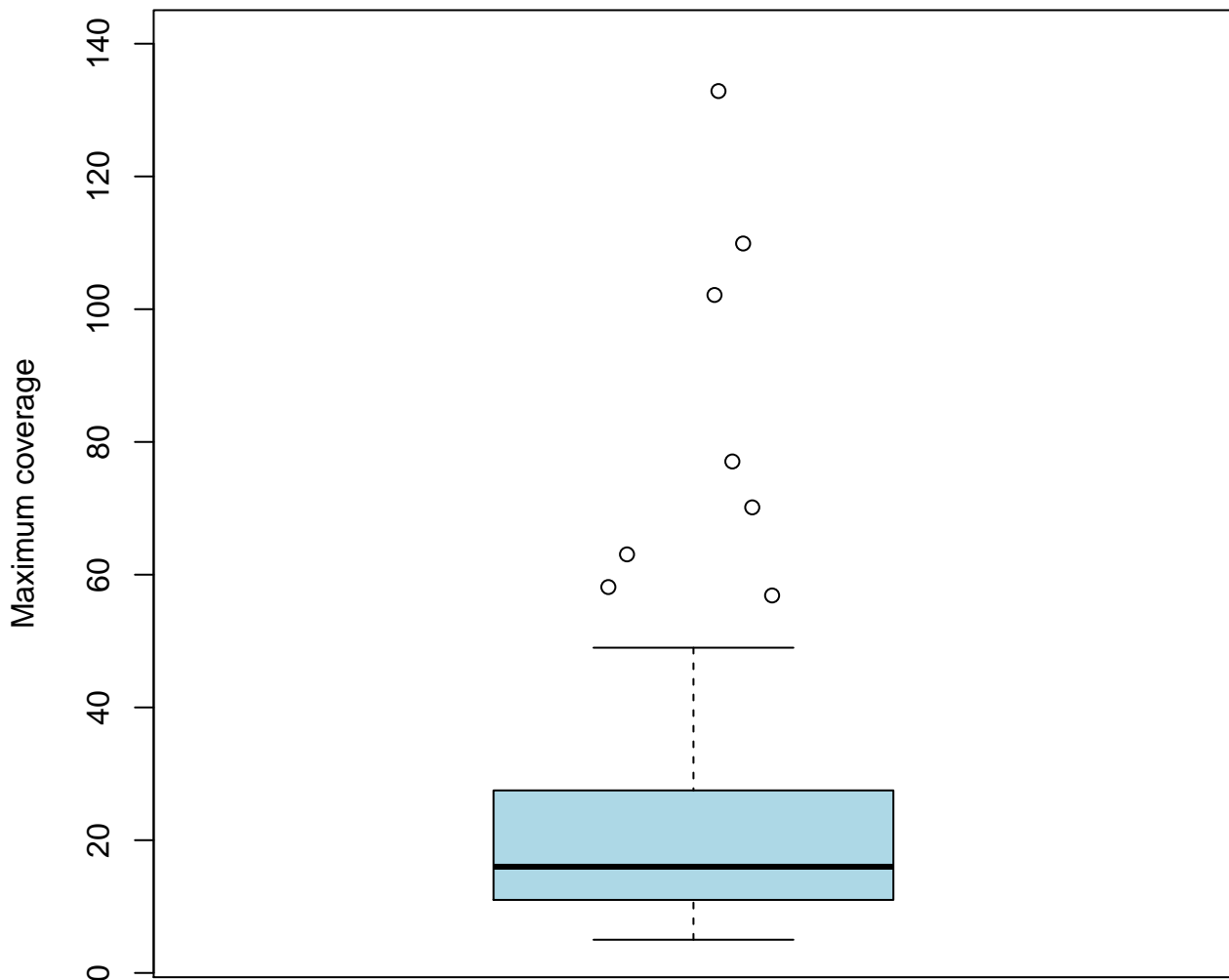
**jx in Intropolis but not "annotated" by knownGene:
134 samples; n jx: 148**



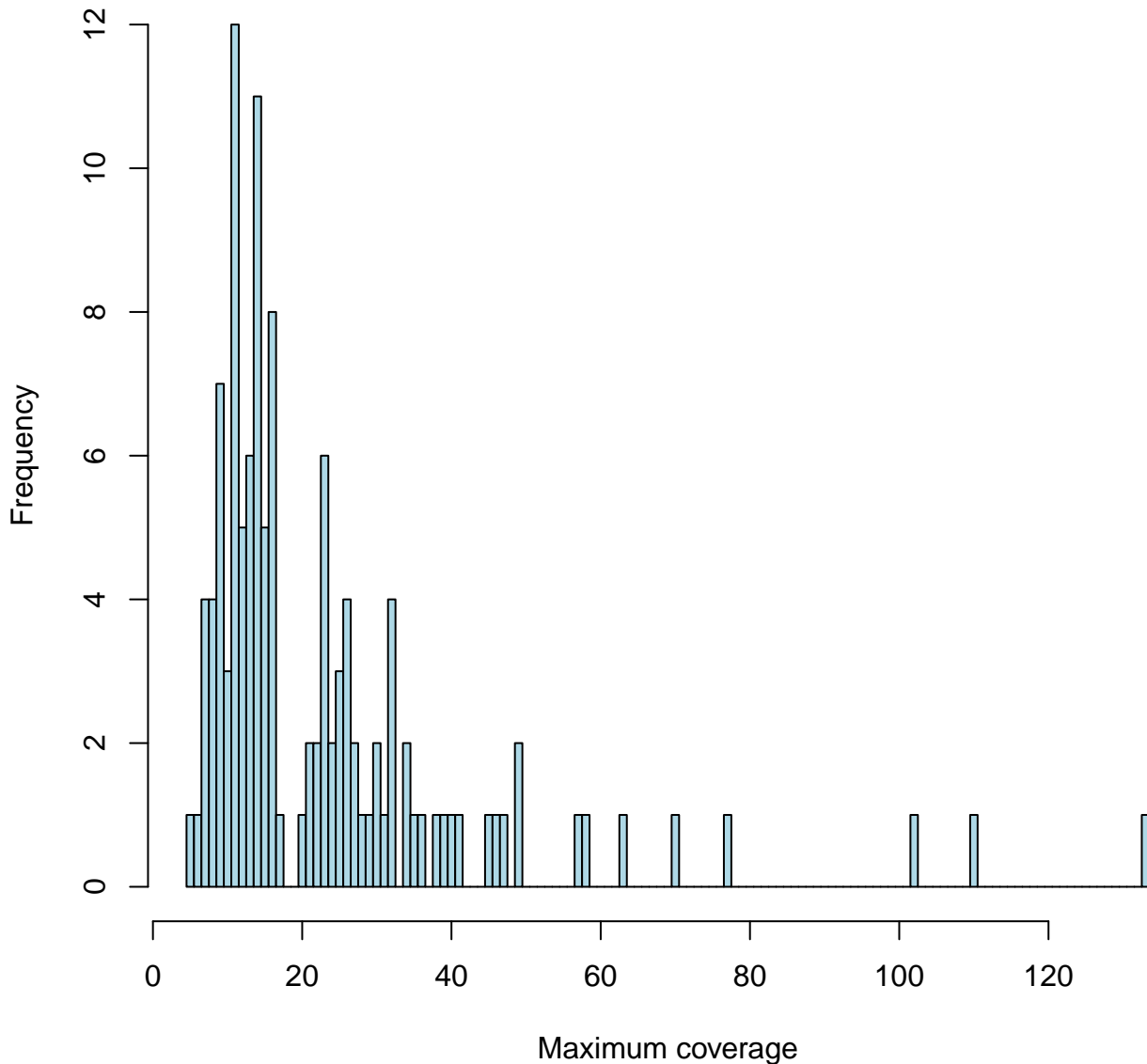
**jx in Intropolis but not "annotated" by knownGene:
134 samples; n jx: 148**



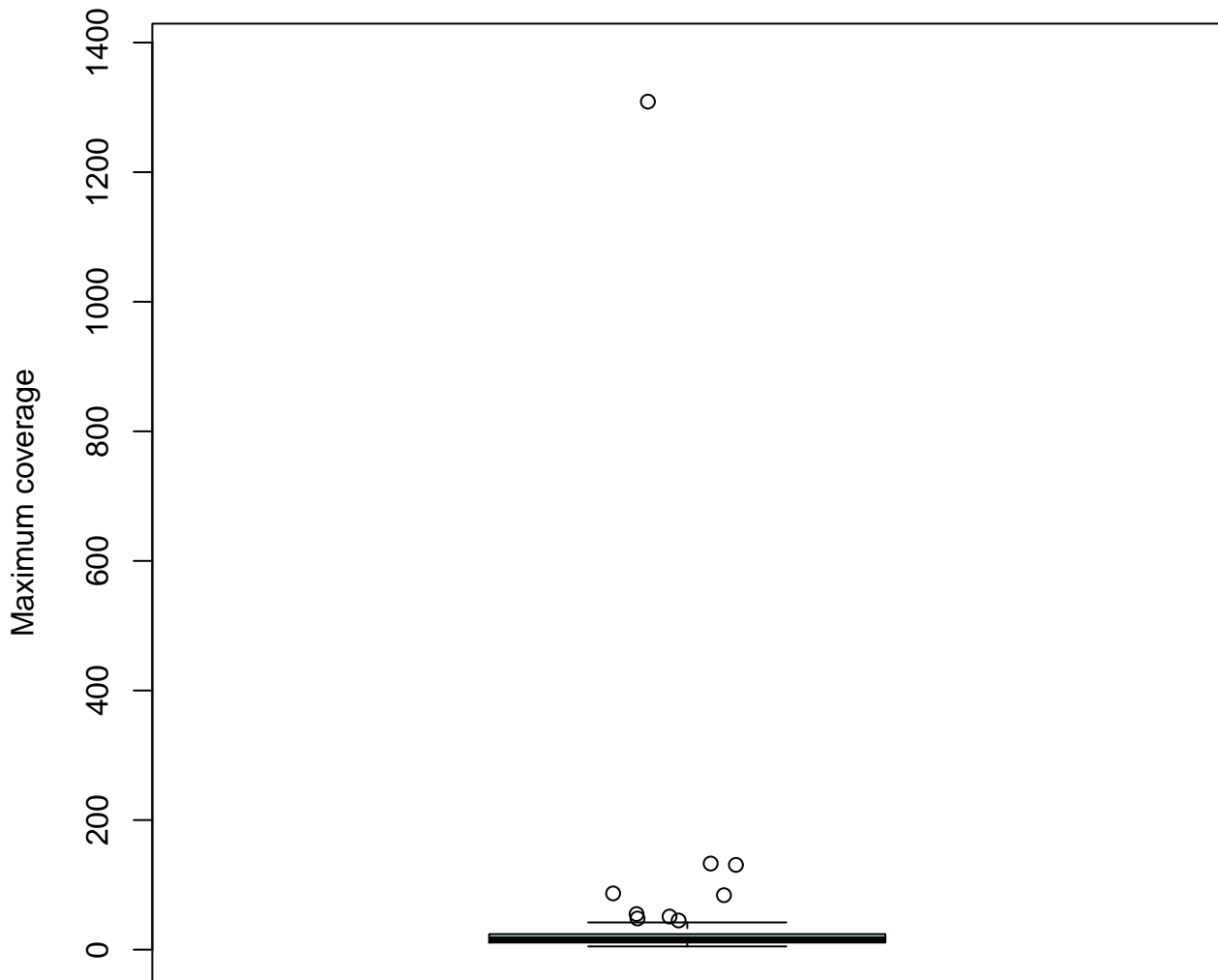
**jx in Intropolis but not "annotated" by knownGene:
135 samples; n jx: 120**



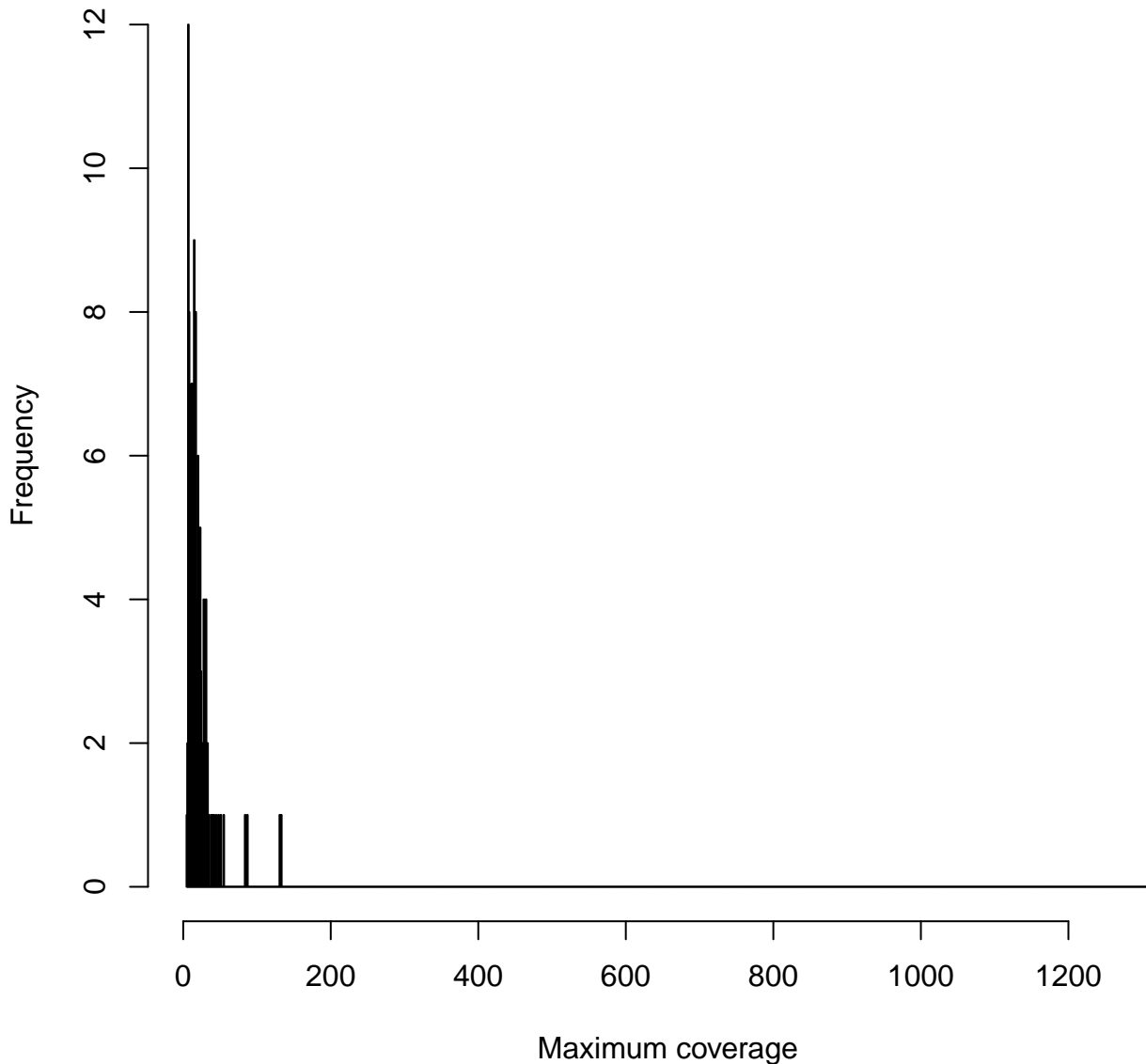
**jx in Intropolis but not "annotated" by knownGene:
135 samples; n jx: 120**



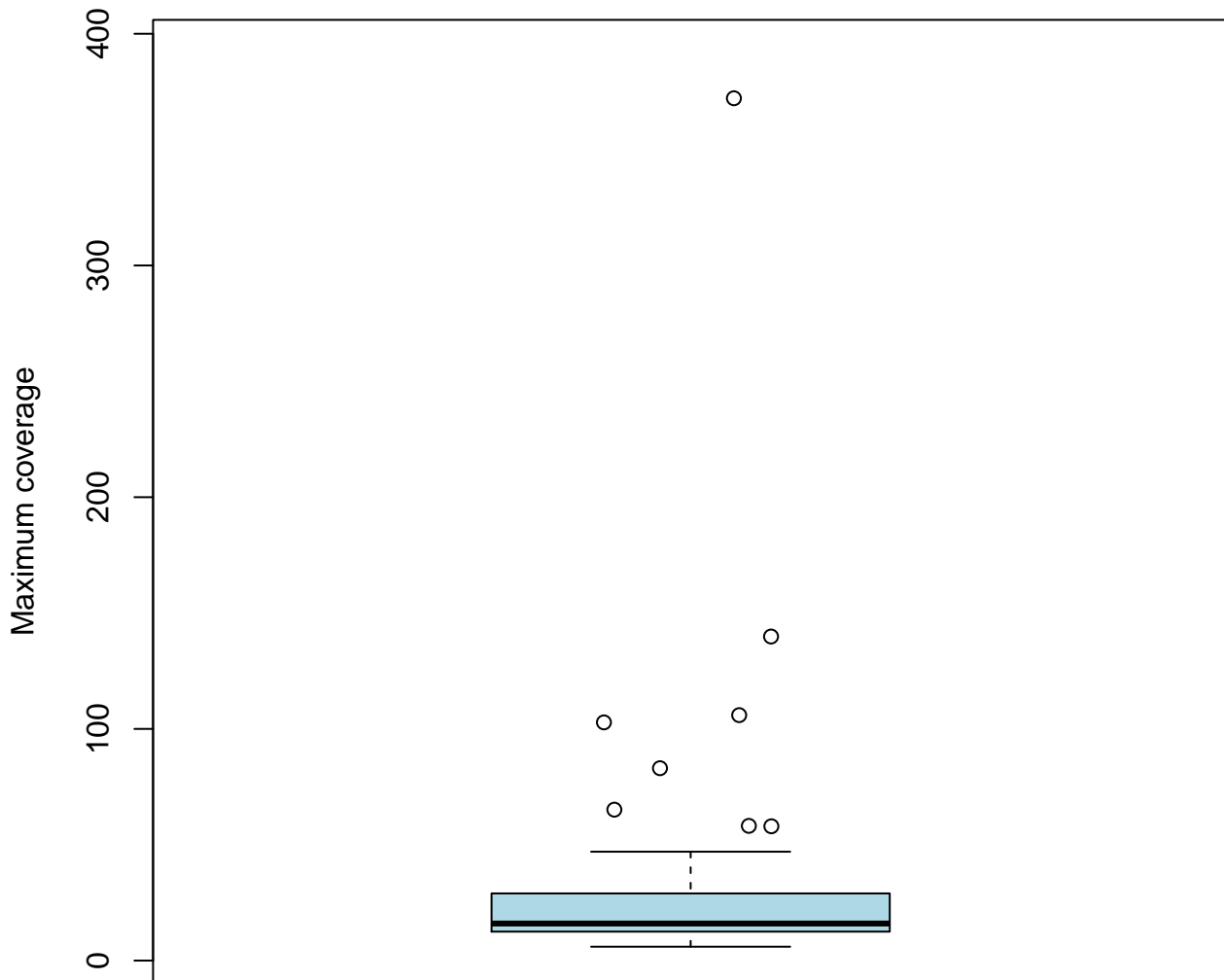
**jx in Intropolis but not "annotated" by knownGene:
136 samples; n jx: 138**



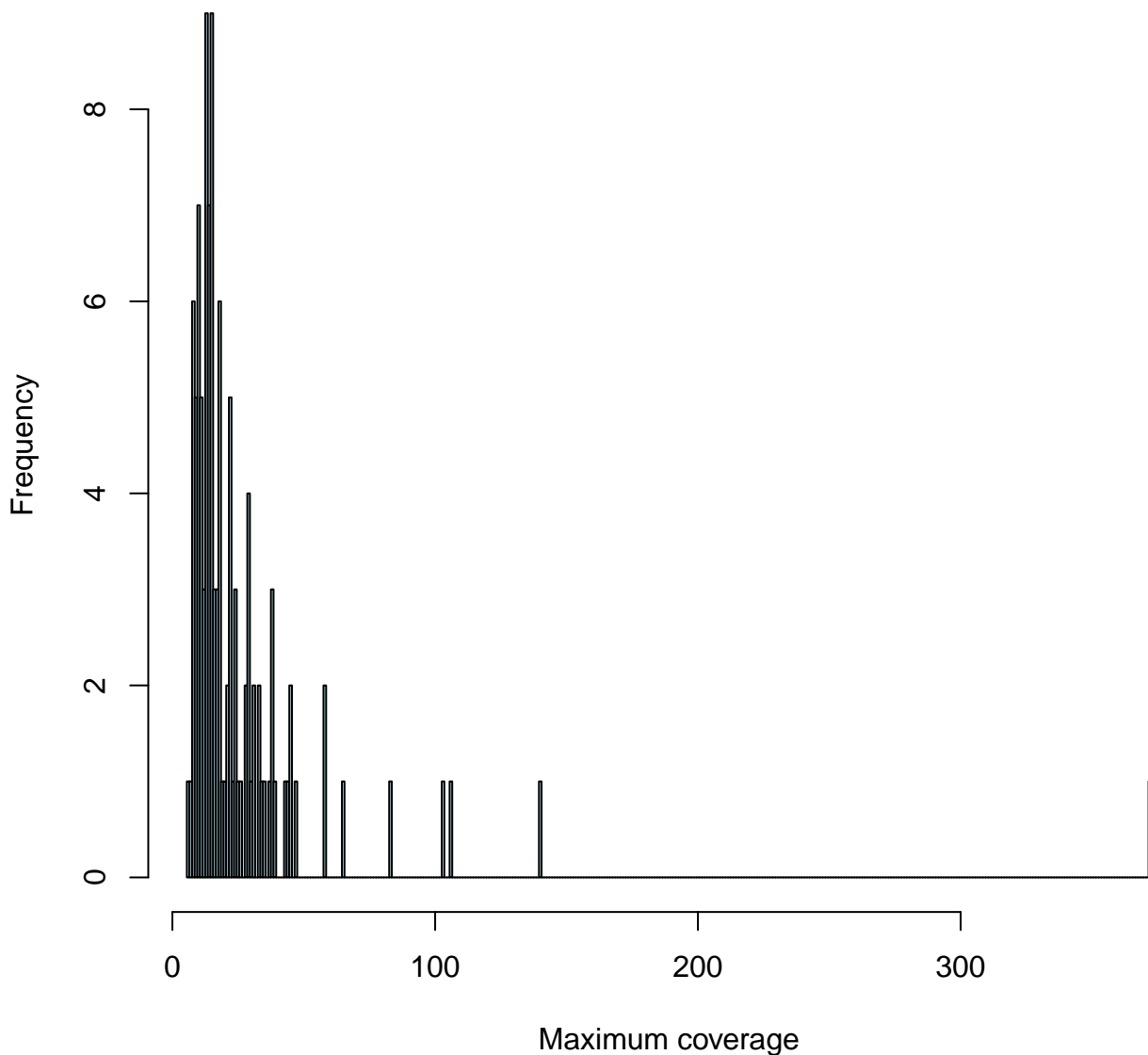
**jx in Intropolis but not "annotated" by knownGene:
136 samples; n jx: 138**



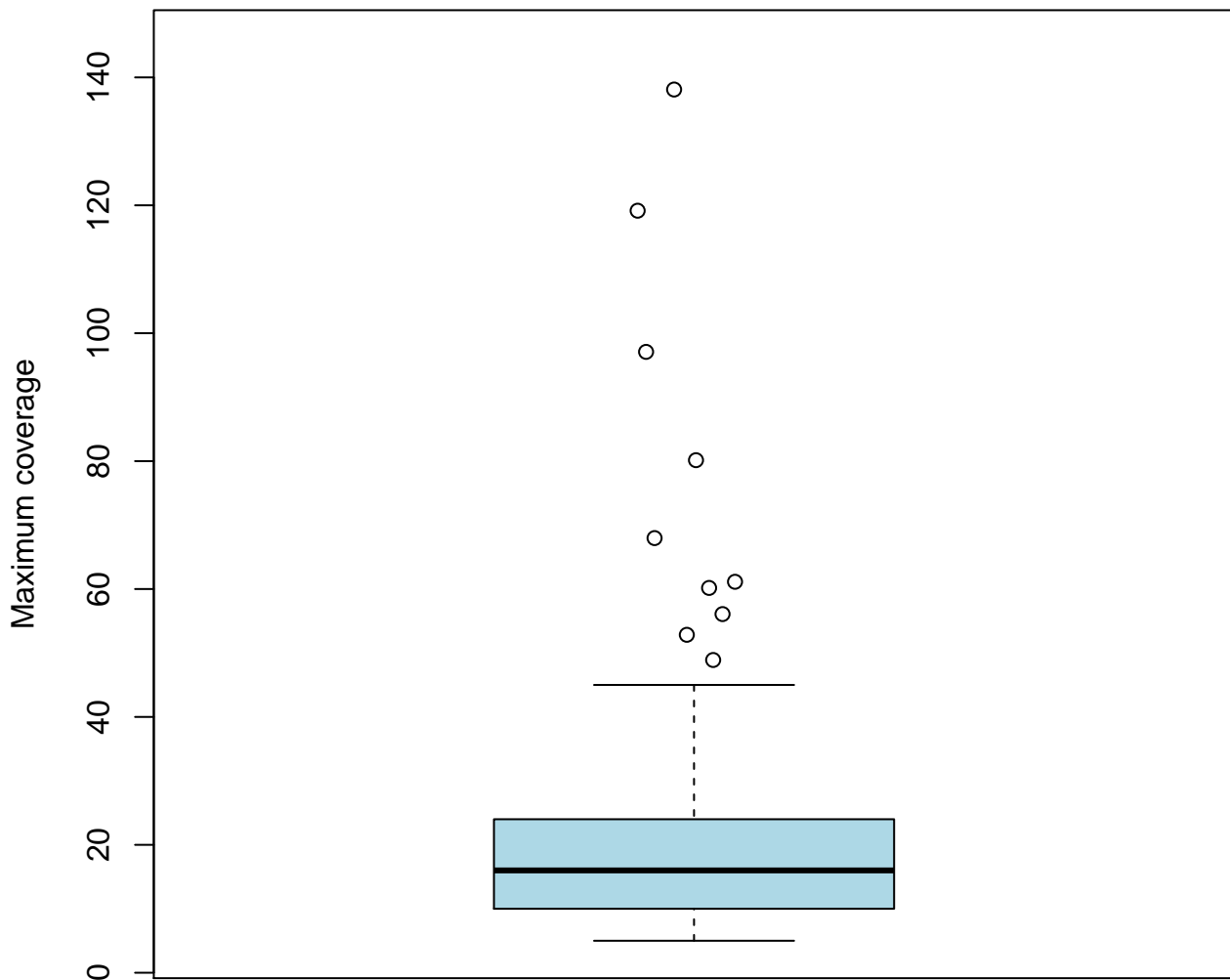
**jx in Intropolis but not "annotated" by knownGene:
137 samples; n jx: 111**



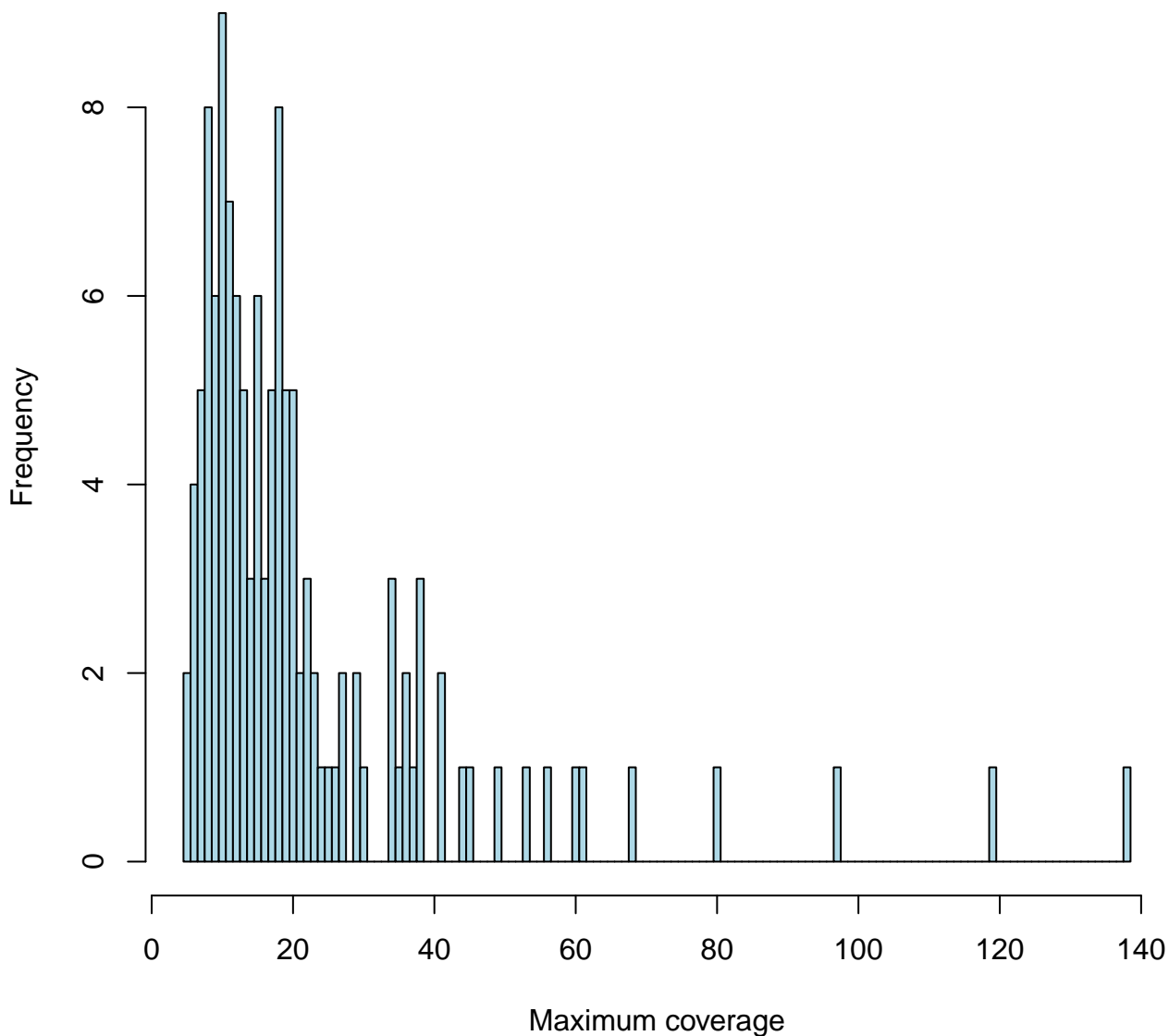
**jx in Intropolis but not "annotated" by knownGene:
137 samples; n jx: 111**



**jx in Intropolis but not "annotated" by knownGene:
138 samples; n jx: 126**



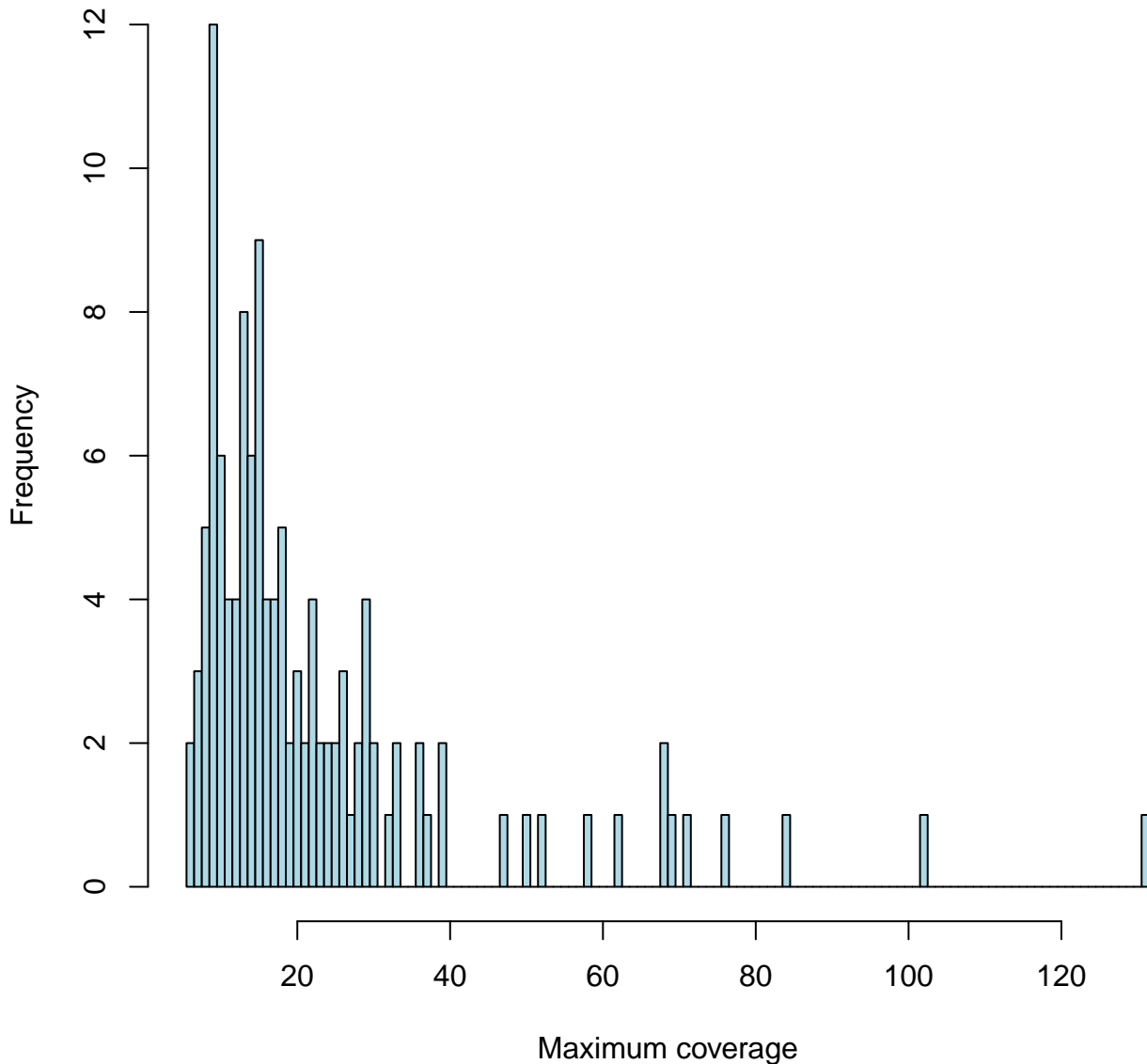
**jx in Intropolis but not "annotated" by knownGene:
138 samples; n jx: 126**



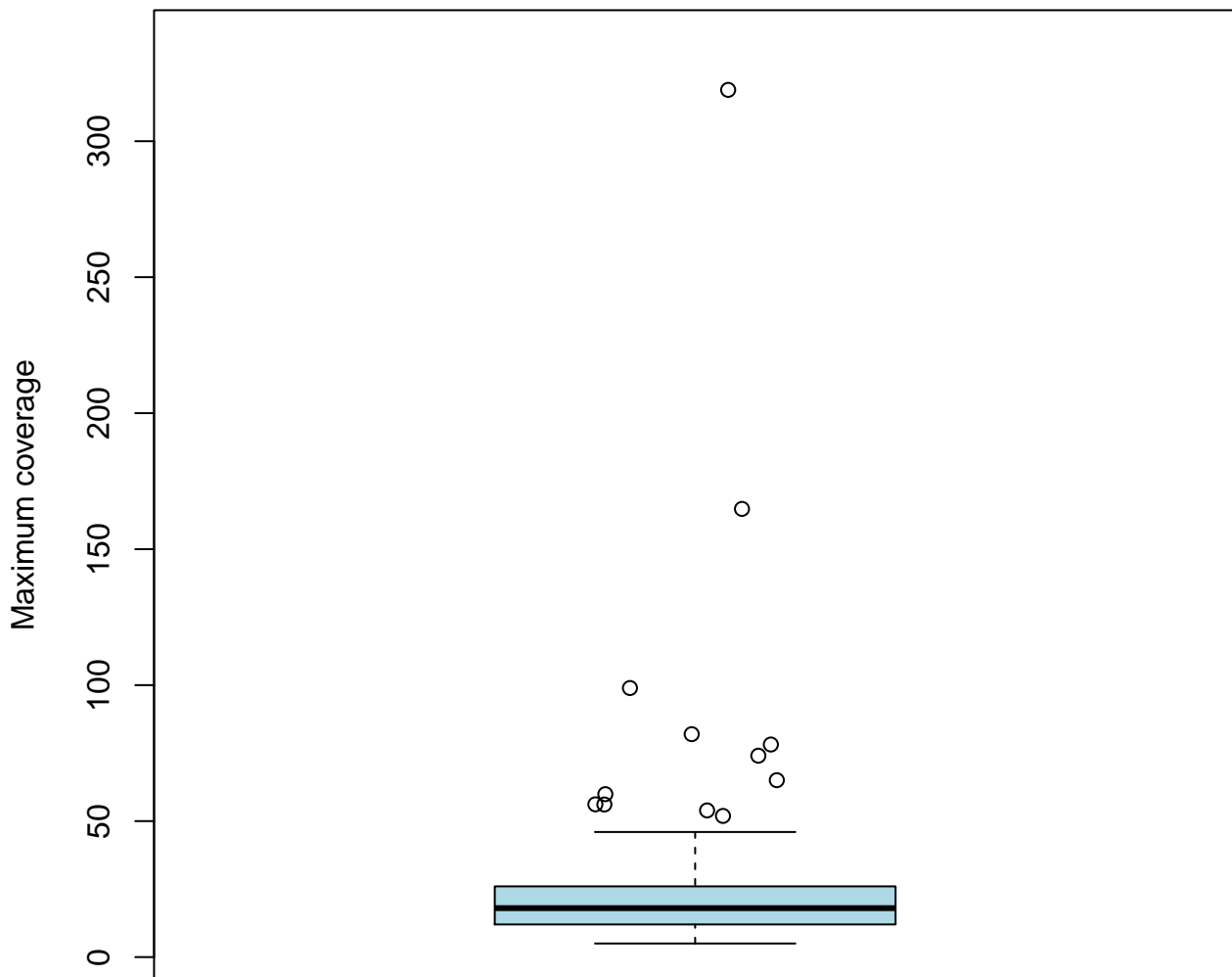
139 samples; n jx: 122



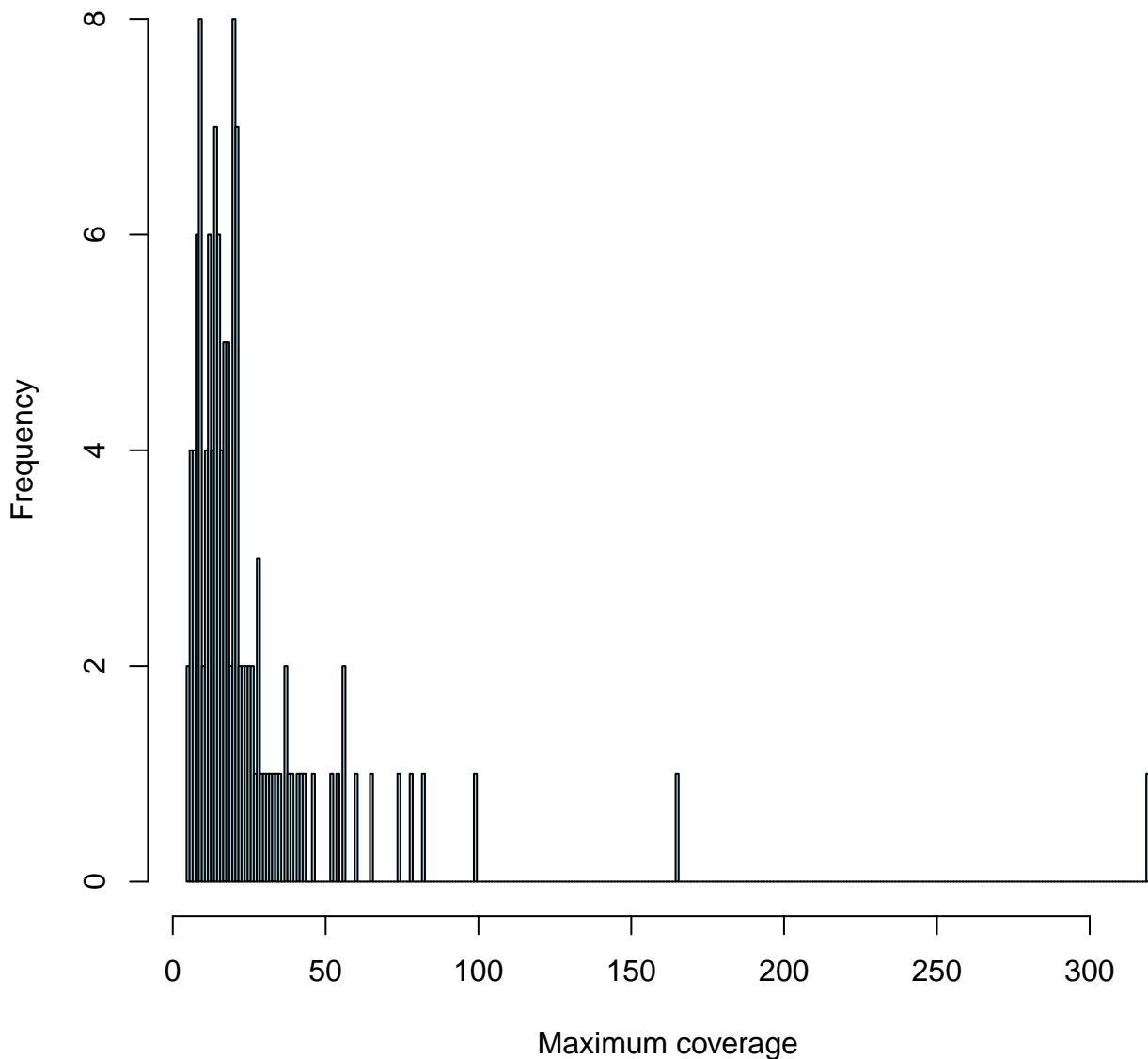
**jx in Intropolis but not "annotated" by knownGene:
139 samples; n jx: 122**



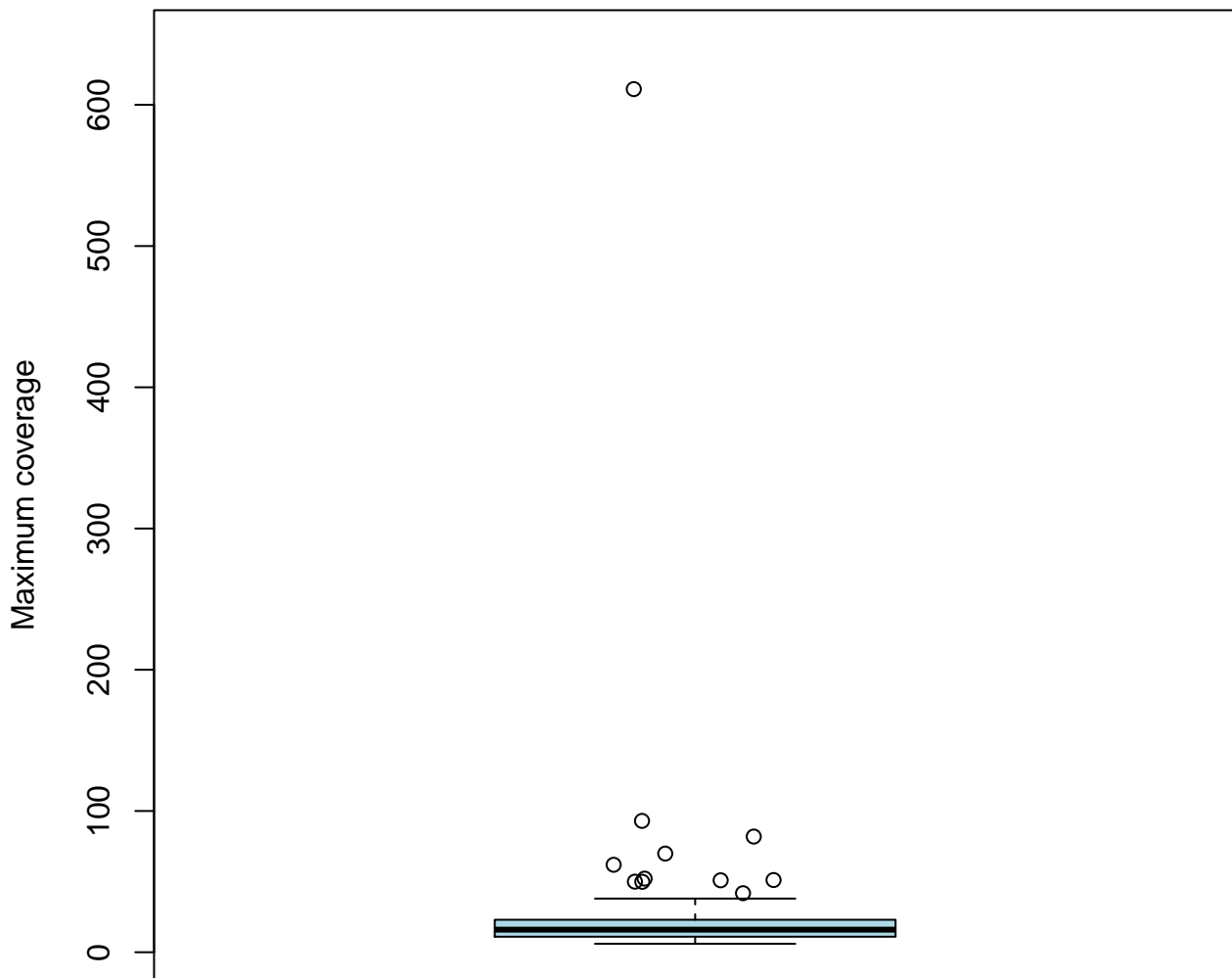
**jx in Intropolis but not "annotated" by knownGene:
140 samples; n jx: 125**



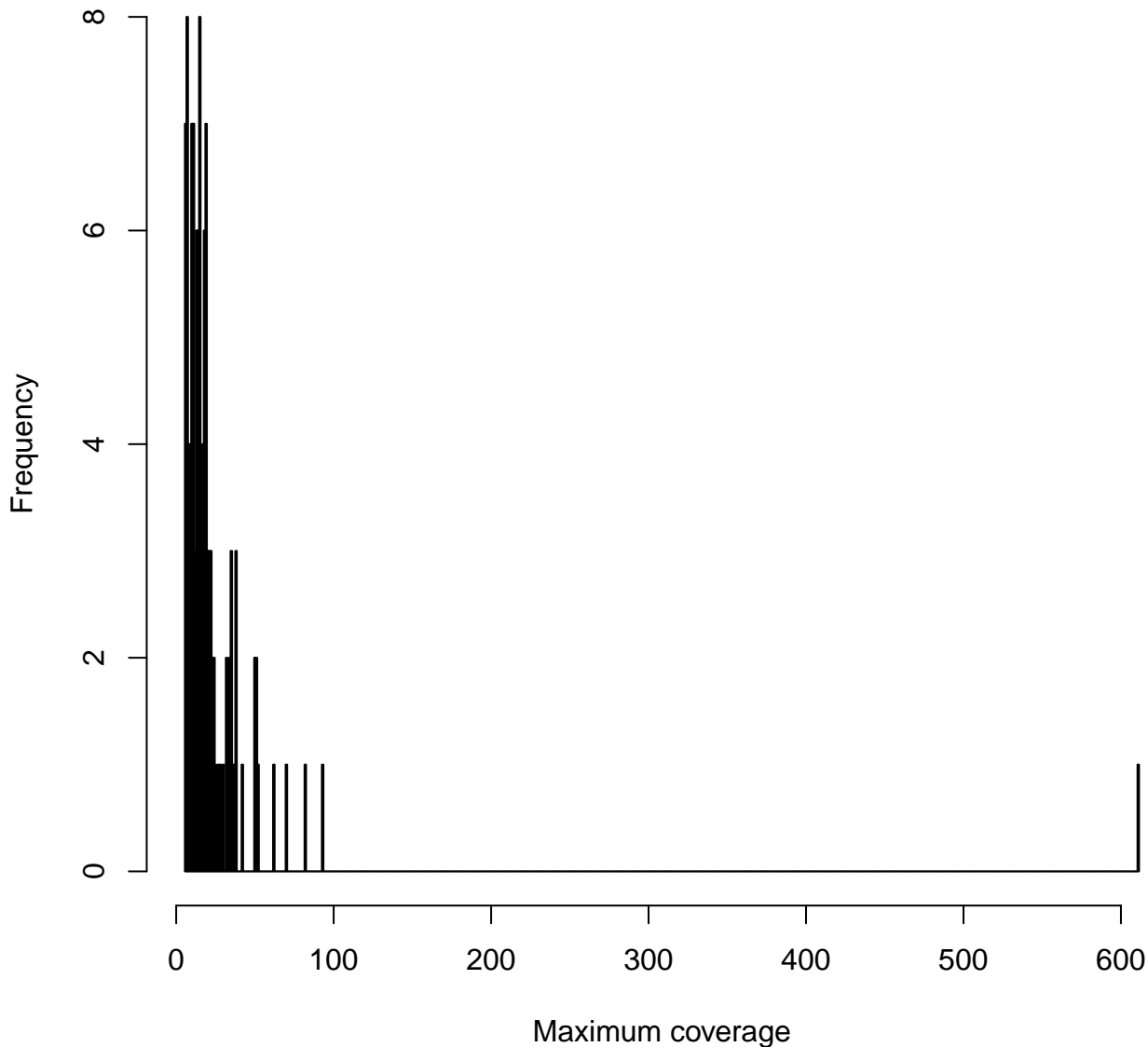
**jx in Intropolis but not "annotated" by knownGene:
140 samples; n jx: 125**



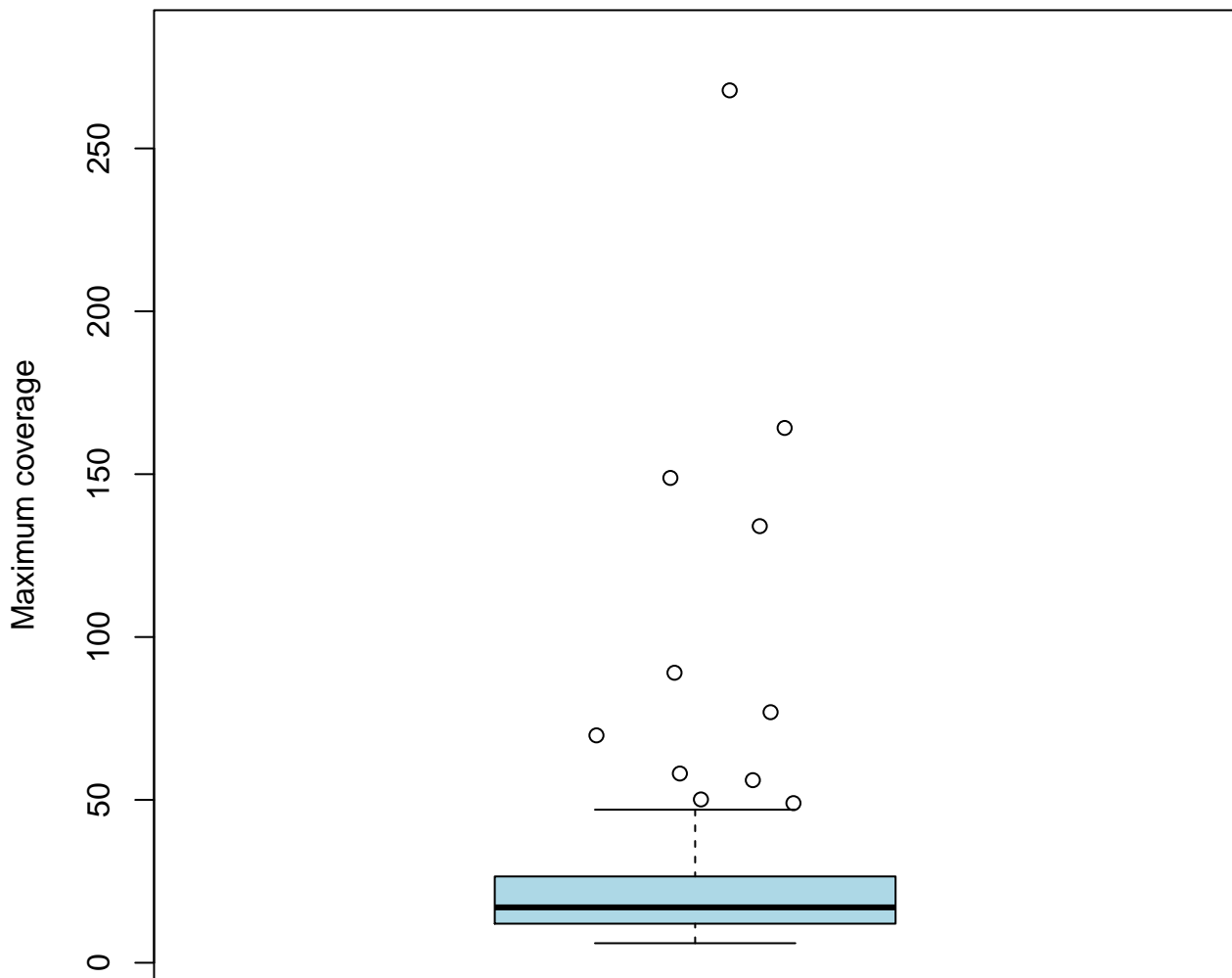
**jx in Intropolis but not "annotated" by knownGene:
141 samples; n jx: 121**



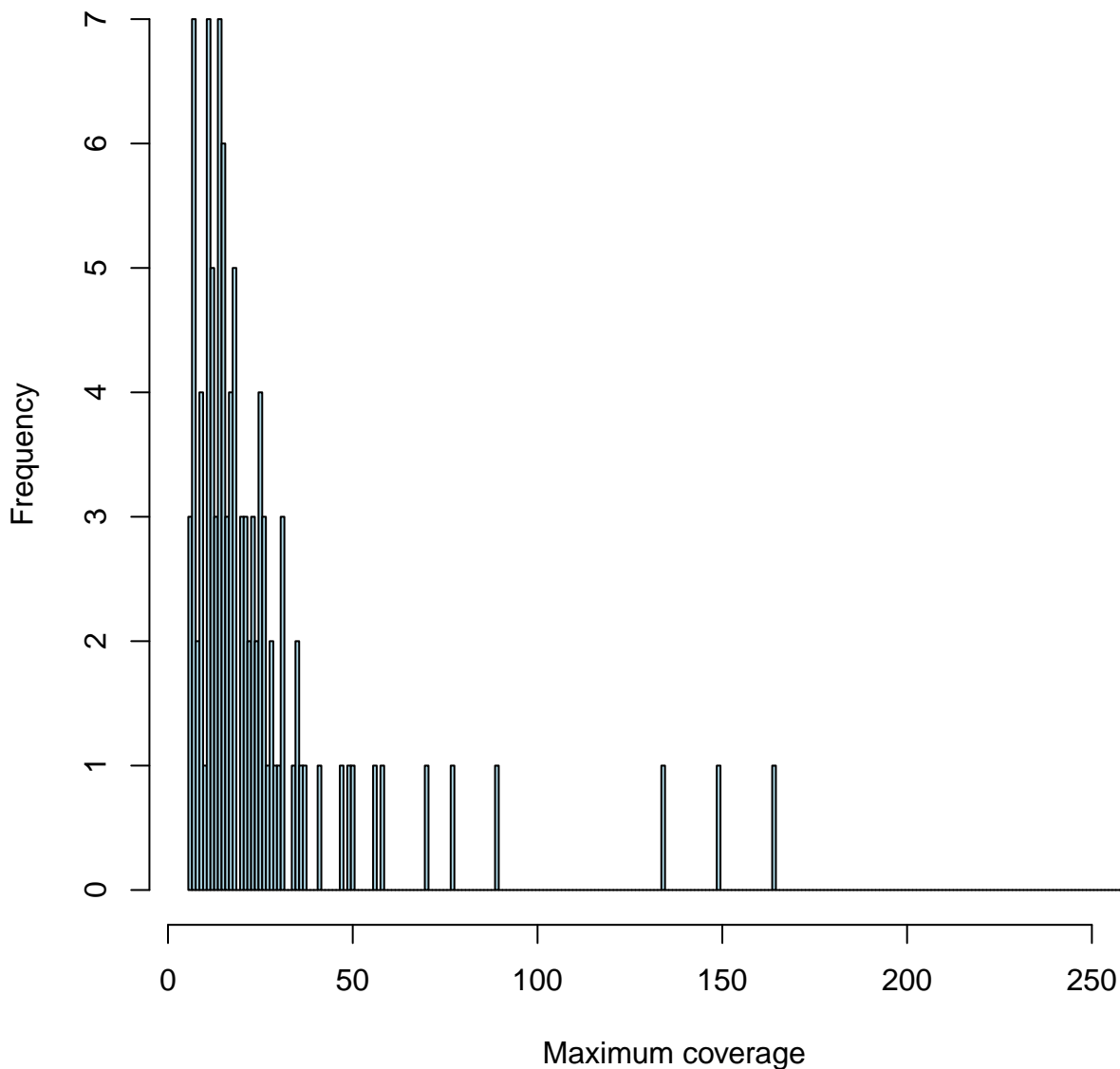
**jx in Intropolis but not "annotated" by knownGene:
141 samples; n jx: 121**



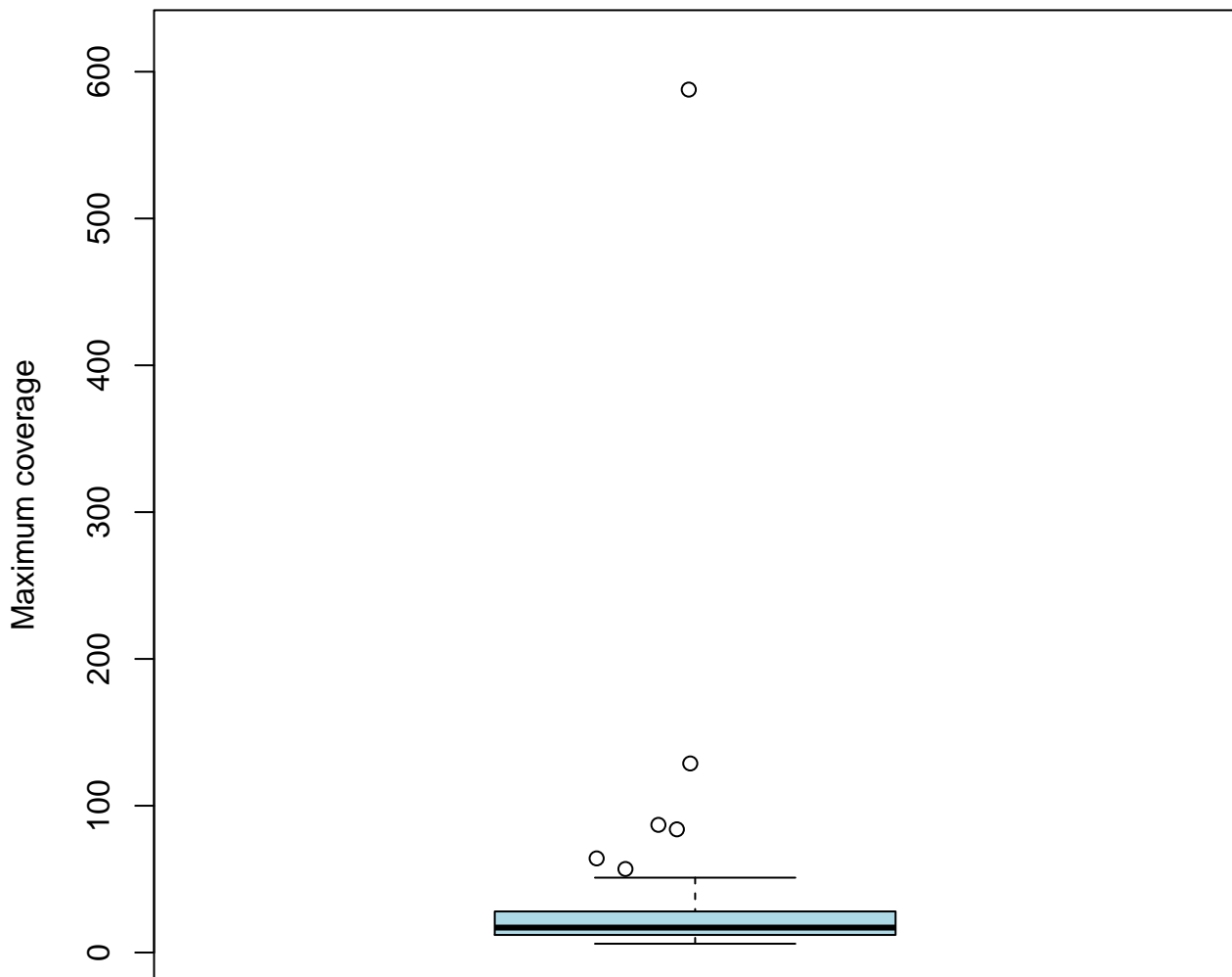
**jx in Intropolis but not "annotated" by knownGene:
142 samples; n jx: 103**



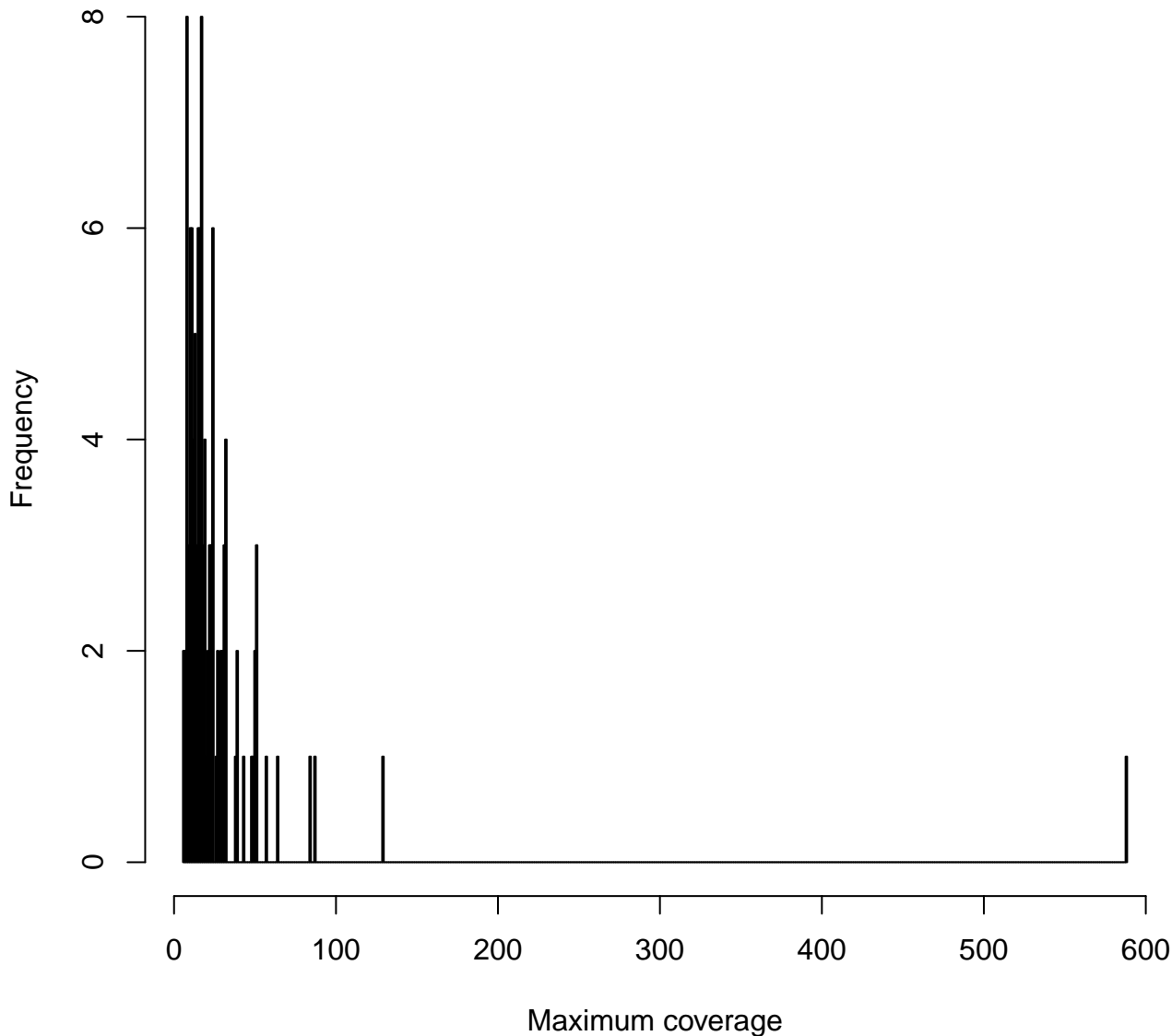
**jx in Intropolis but not "annotated" by knownGene:
142 samples; n jx: 103**



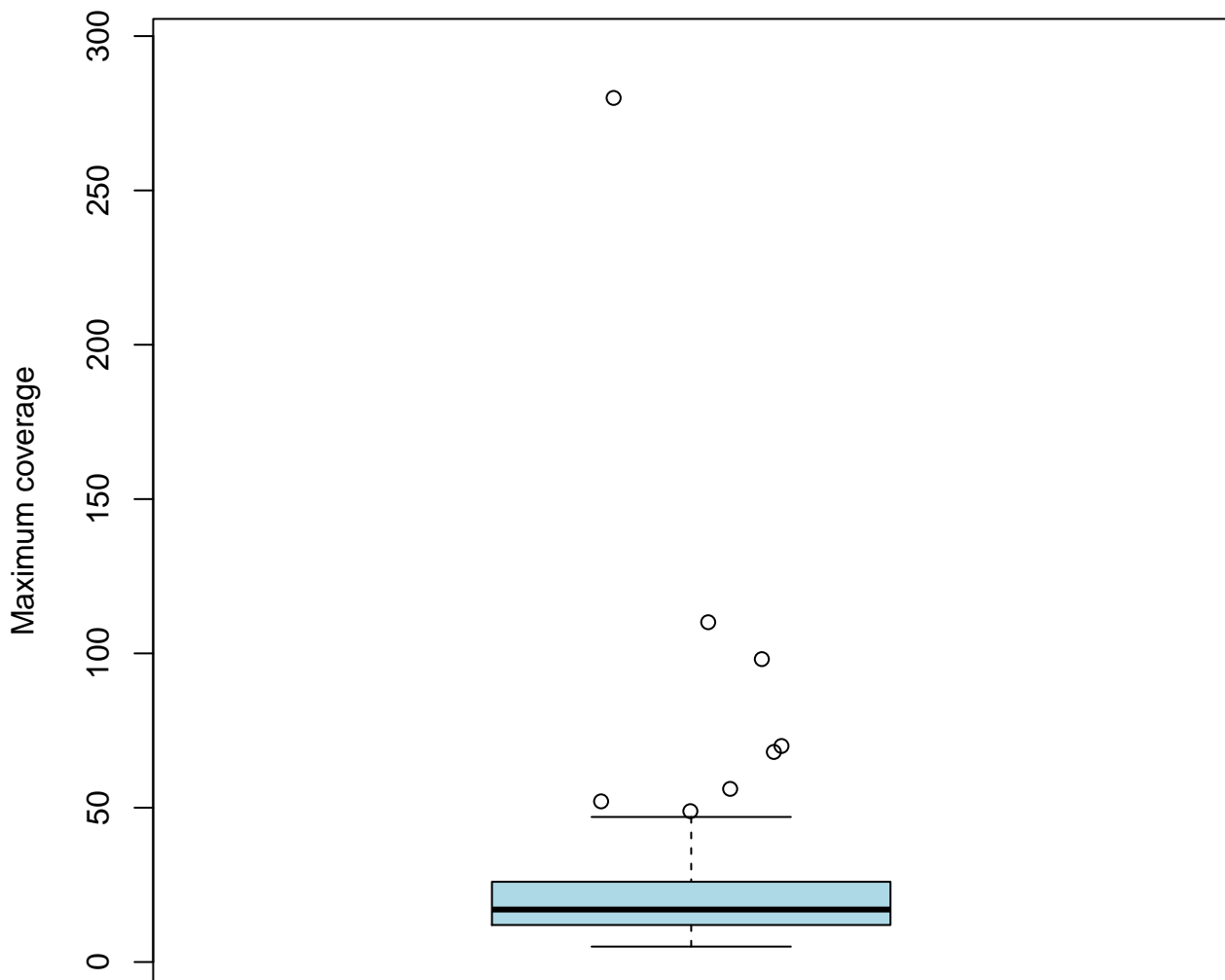
**jx in Intropolis but not "annotated" by knownGene:
143 samples; n jx: 112**



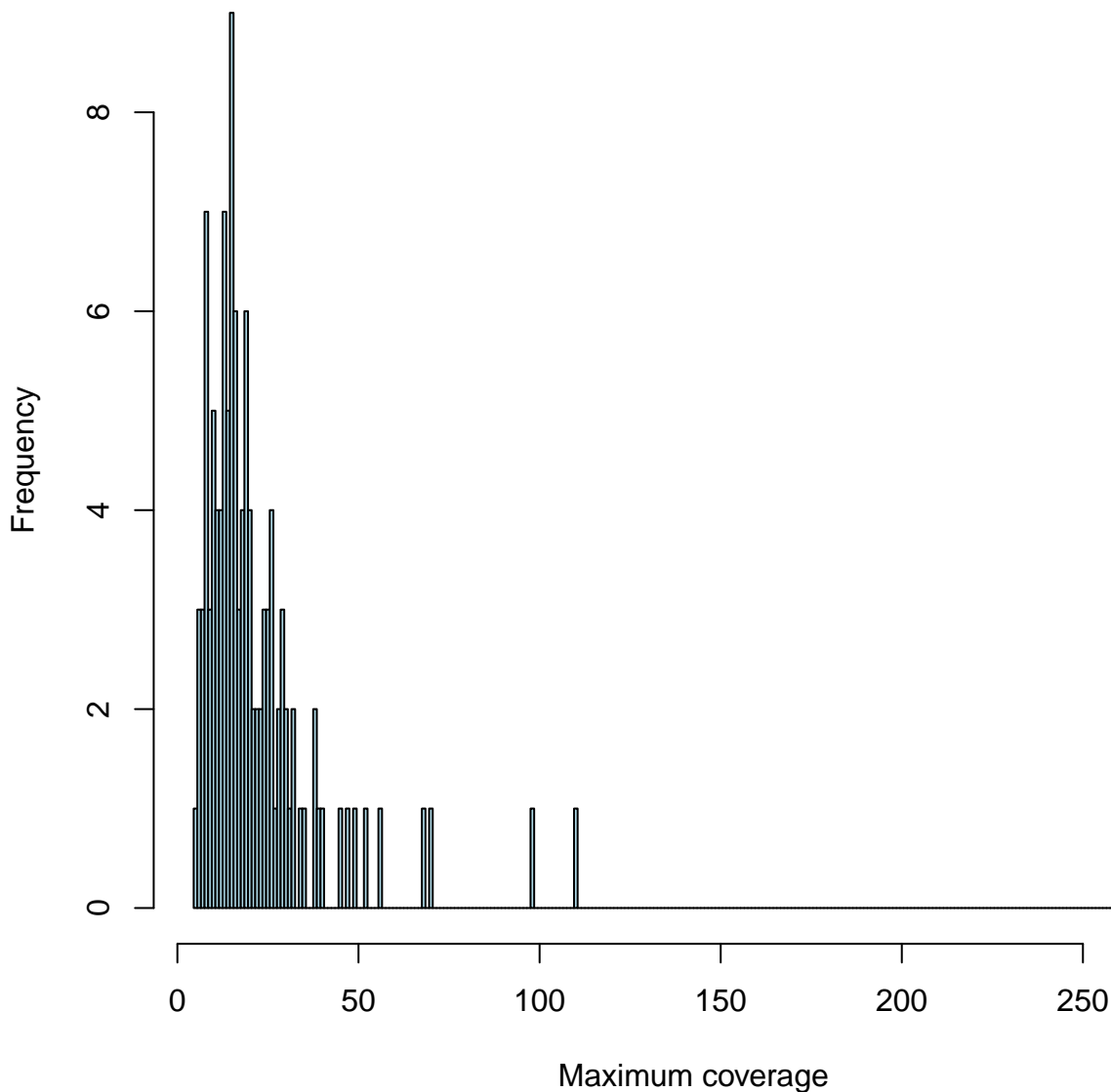
**jx in Intropolis but not "annotated" by knownGene:
143 samples; n jx: 112**



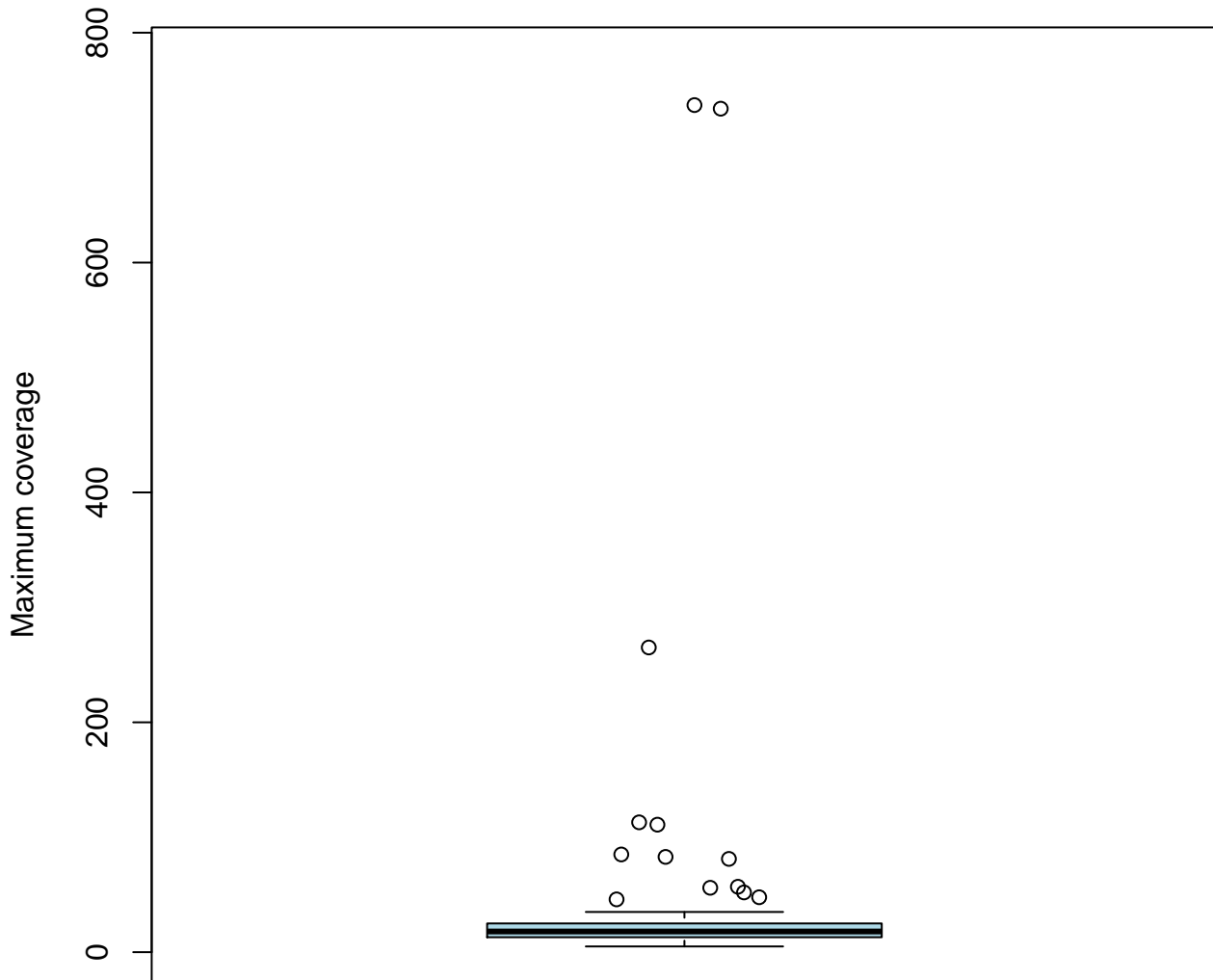
**jx in Intropolis but not "annotated" by knownGene:
144 samples; n jx: 117**



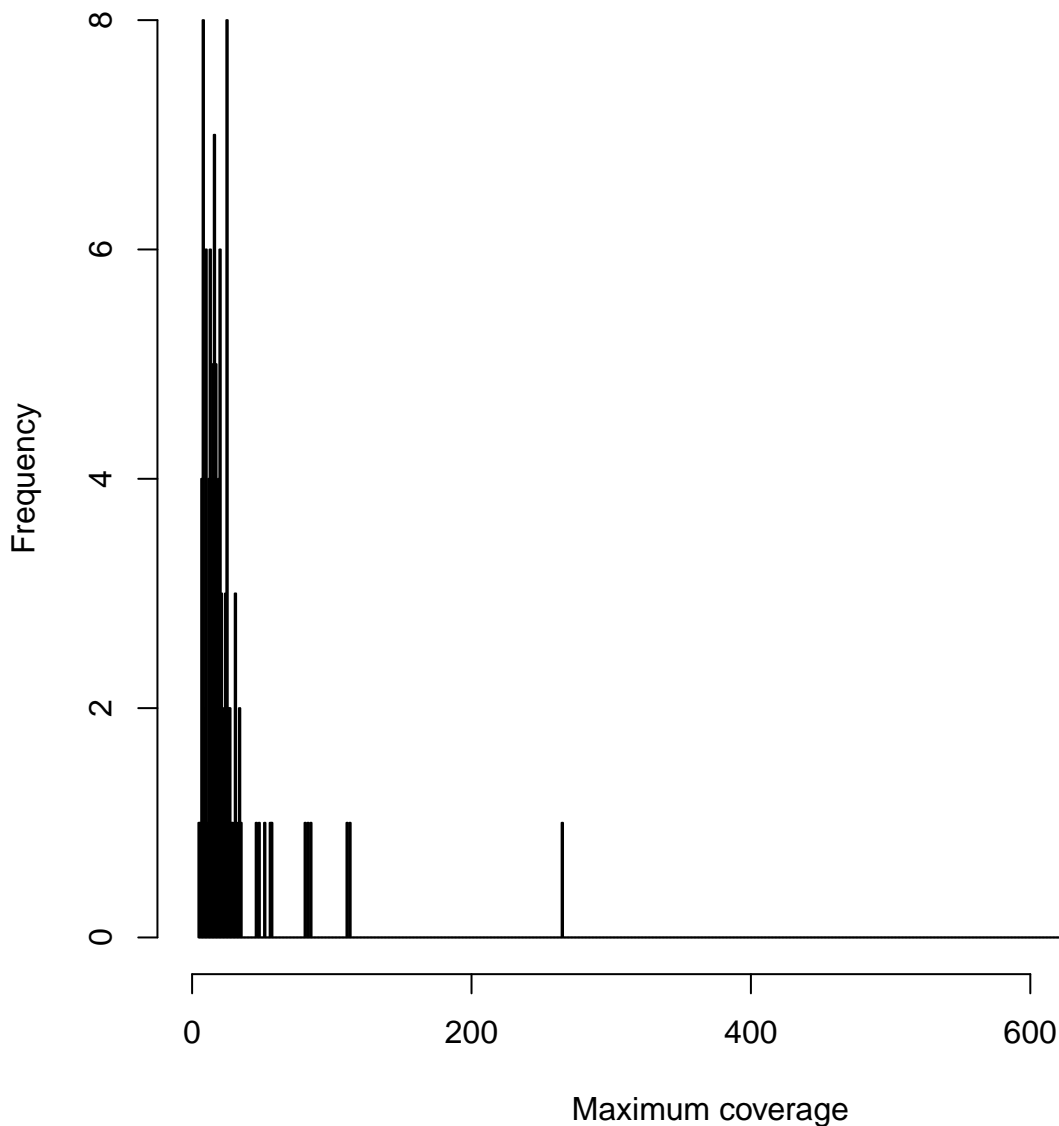
**jx in Intropolis but not "annotated" by knownGene:
144 samples; n jx: 117**



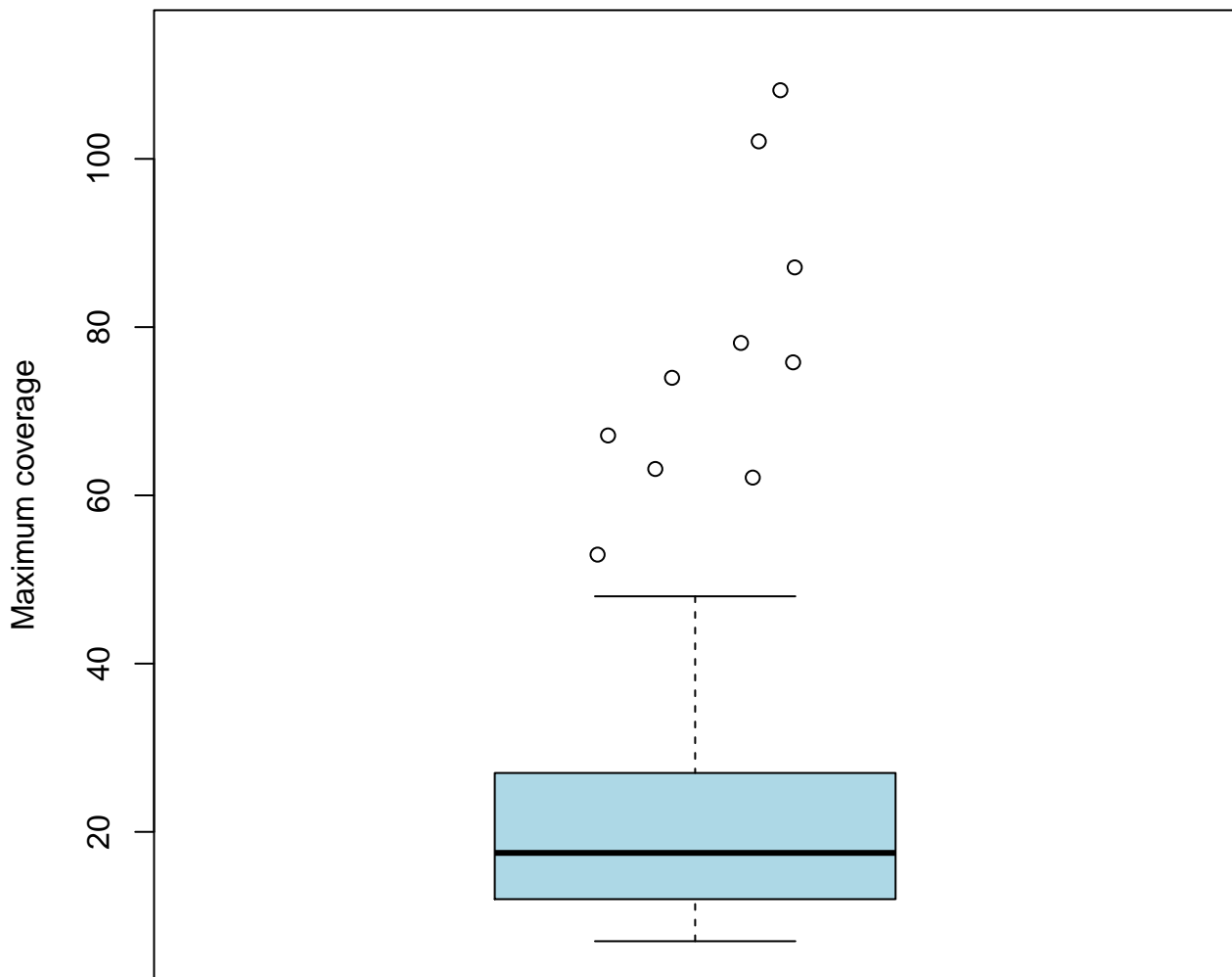
**jx in Intropolis but not "annotated" by knownGene:
145 samples; n jx: 110**



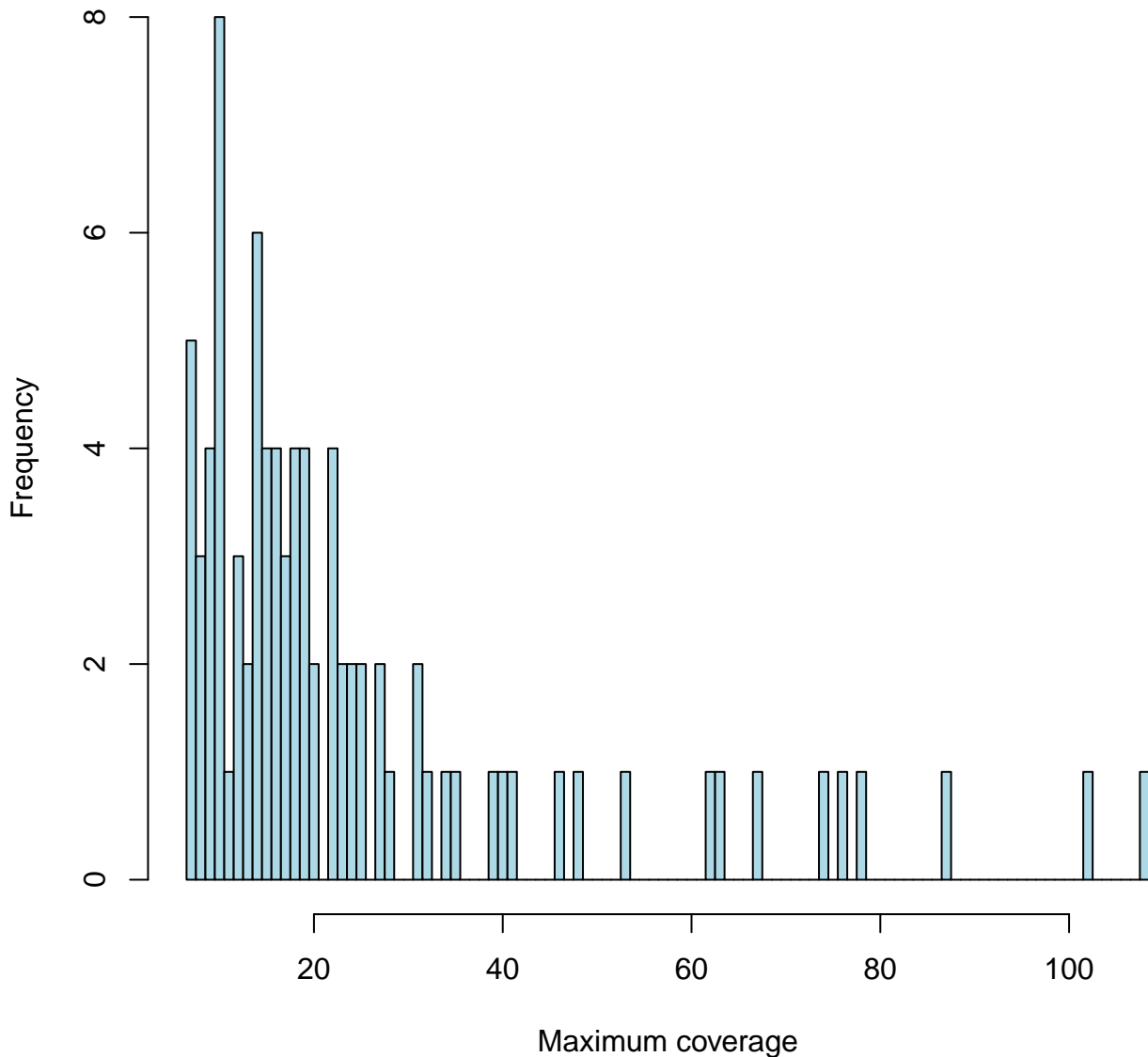
**jx in Intropolis but not "annotated" by knownGene:
145 samples; n jx: 110**



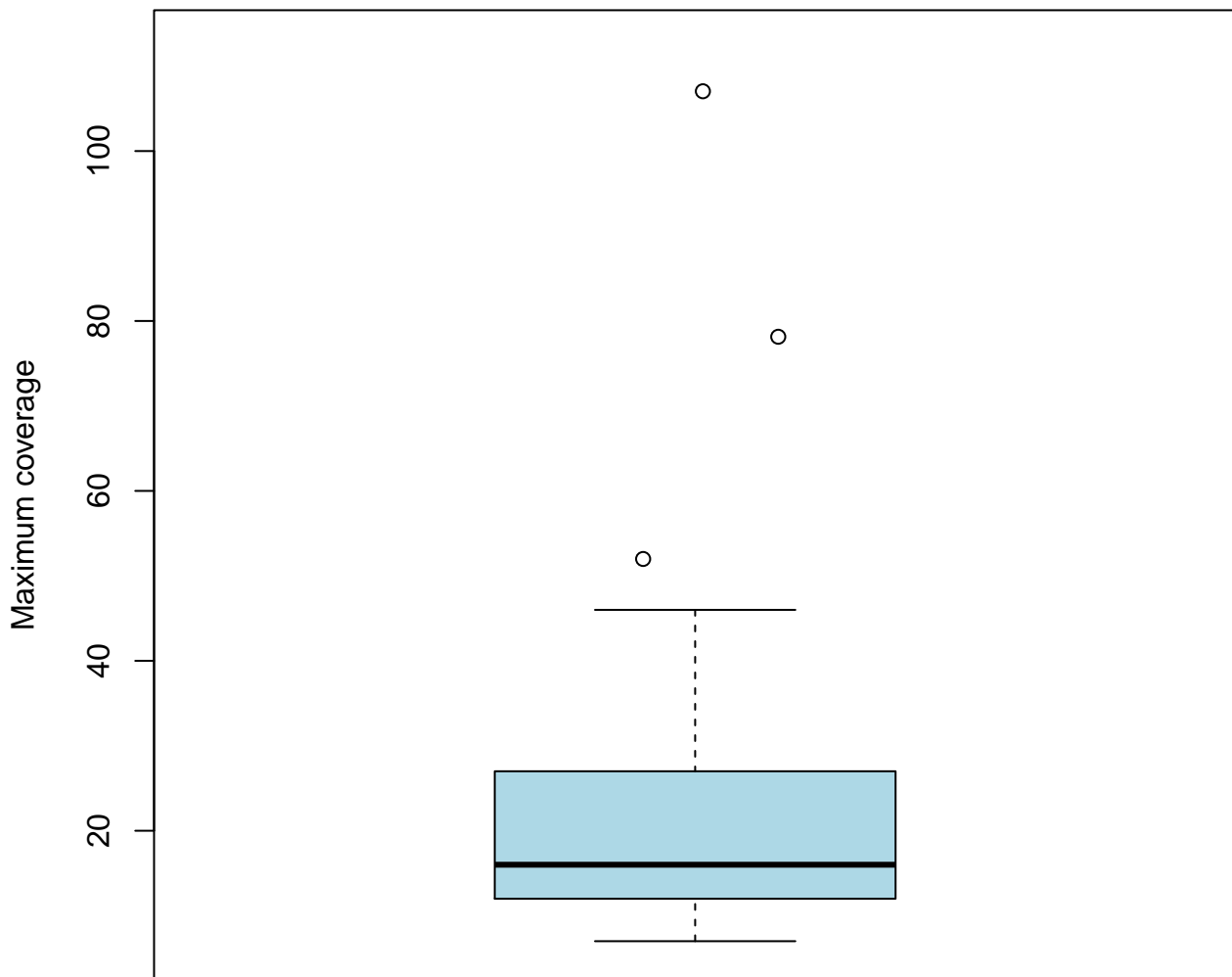
**jx in Intropolis but not "annotated" by knownGene:
146 samples; n jx: 86**



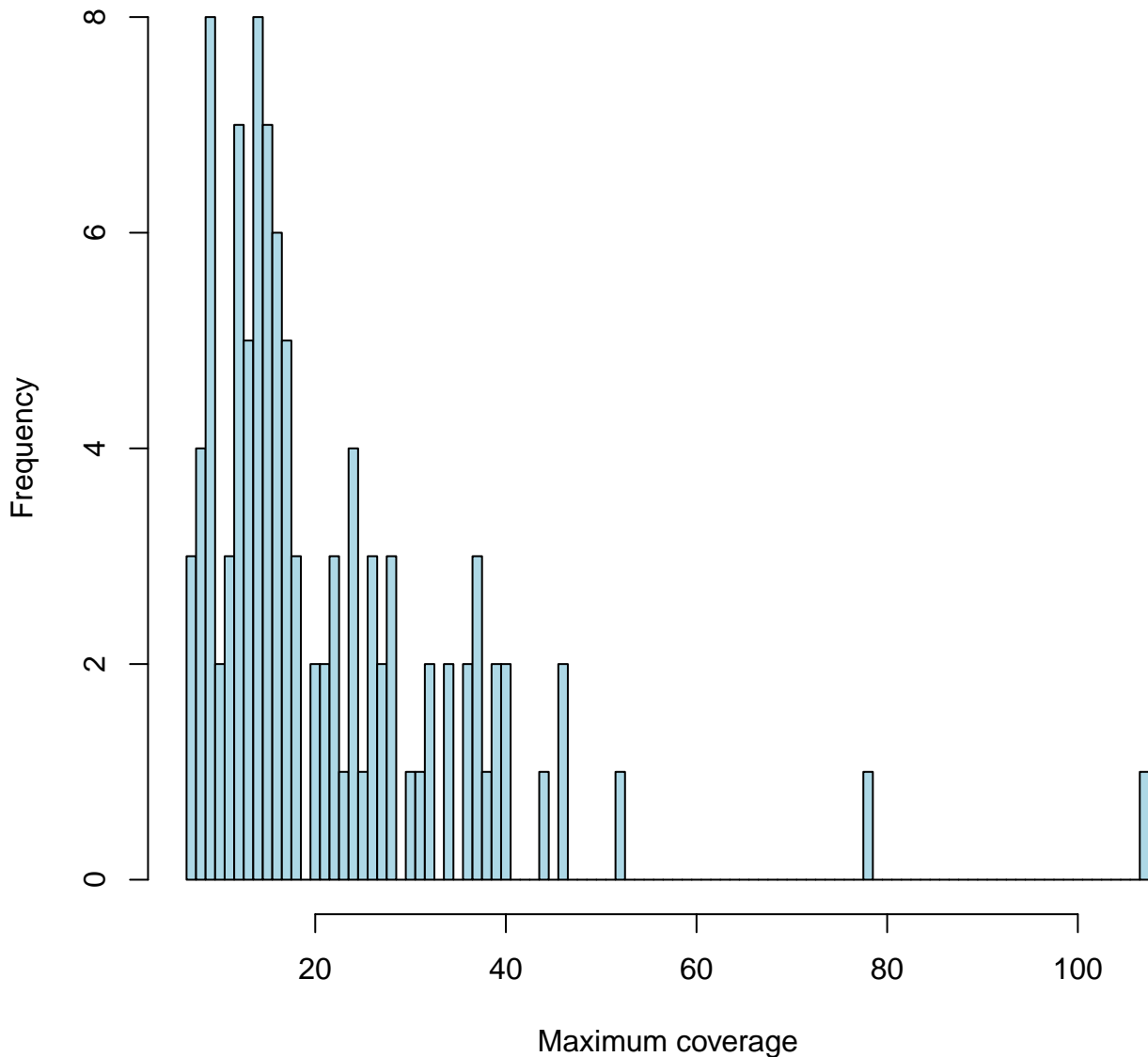
**jx in Intropolis but not "annotated" by knownGene:
146 samples; n jx: 86**



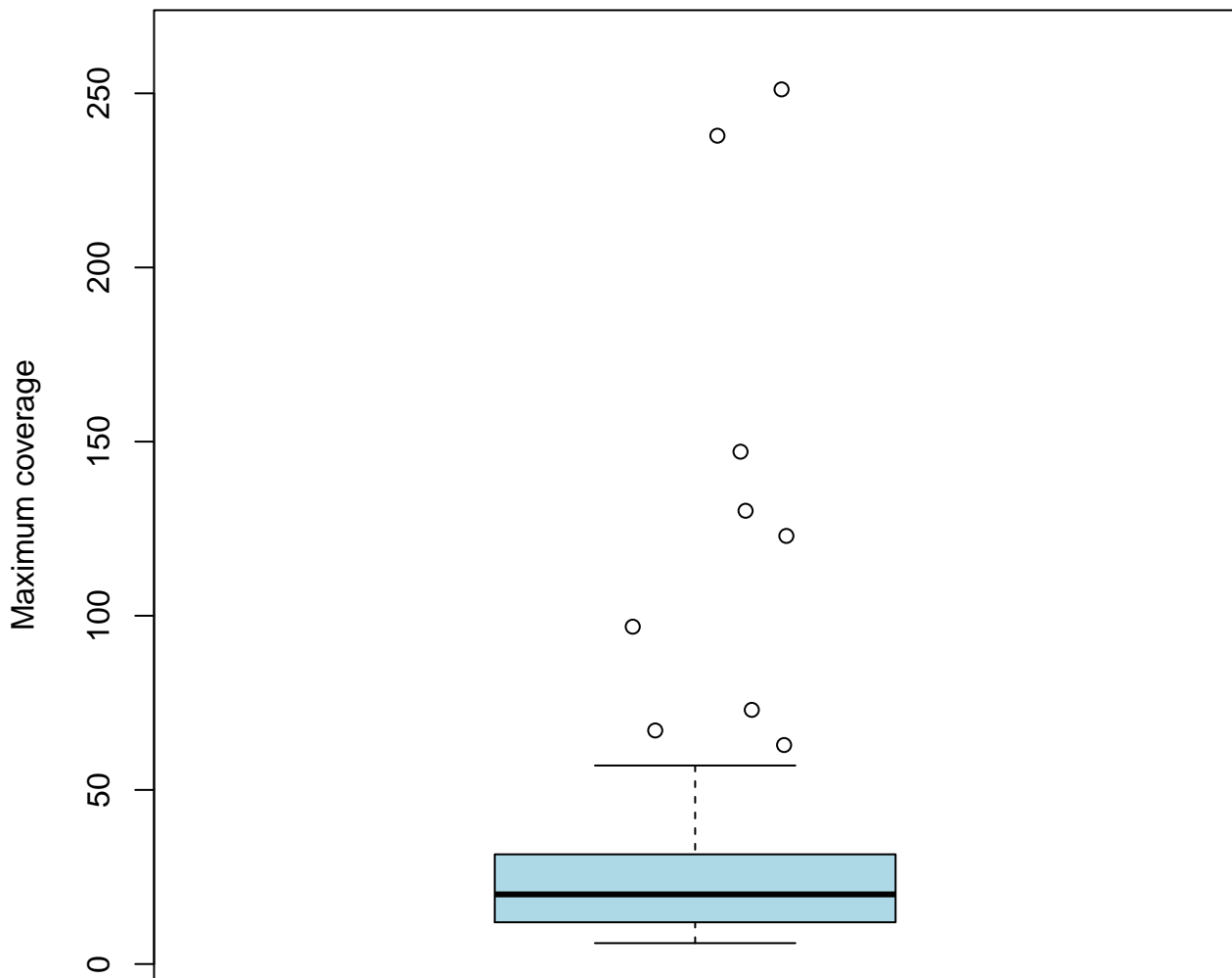
**jx in Intropolis but not "annotated" by knownGene:
147 samples; n jx: 104**



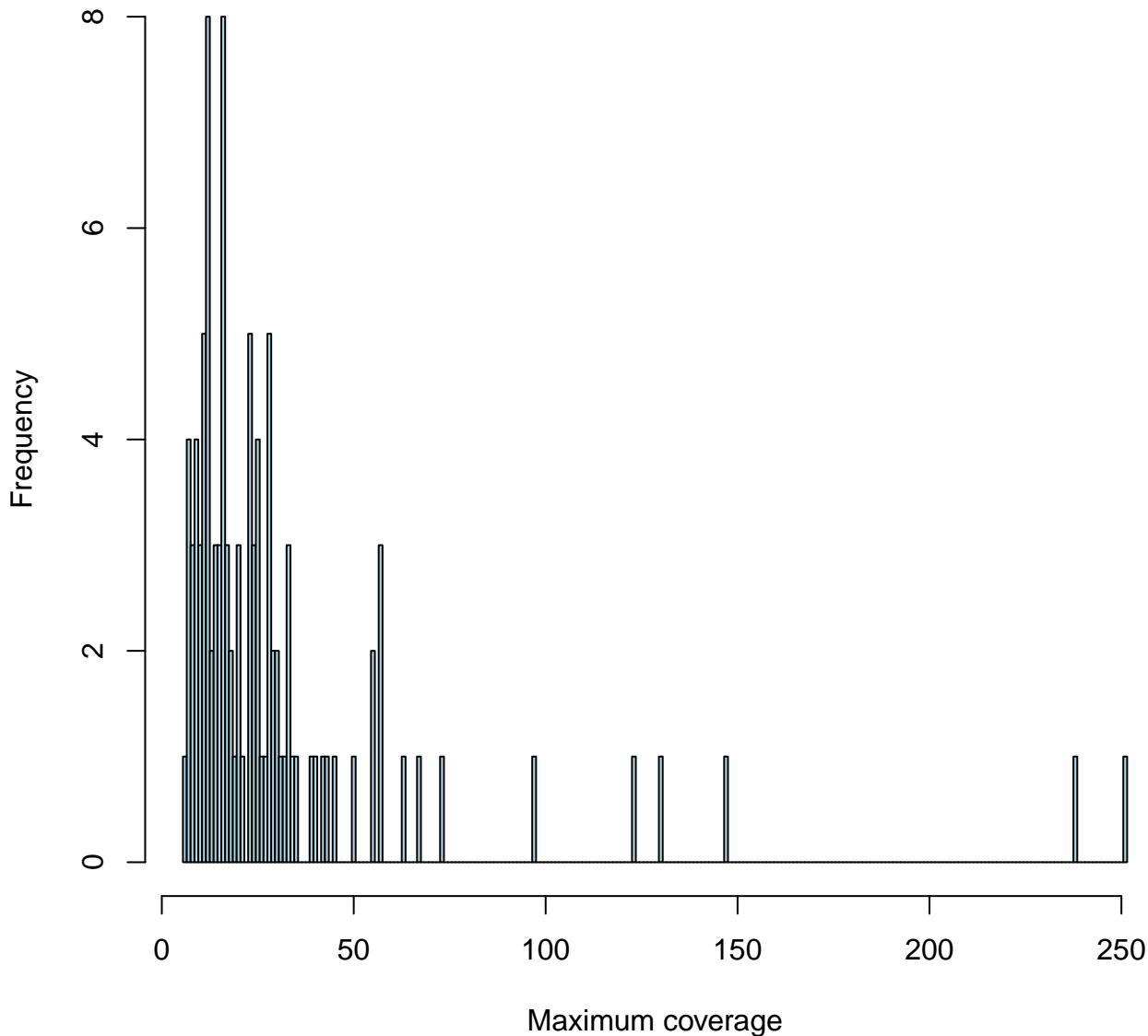
**jx in Intropolis but not "annotated" by knownGene:
147 samples; n jx: 104**



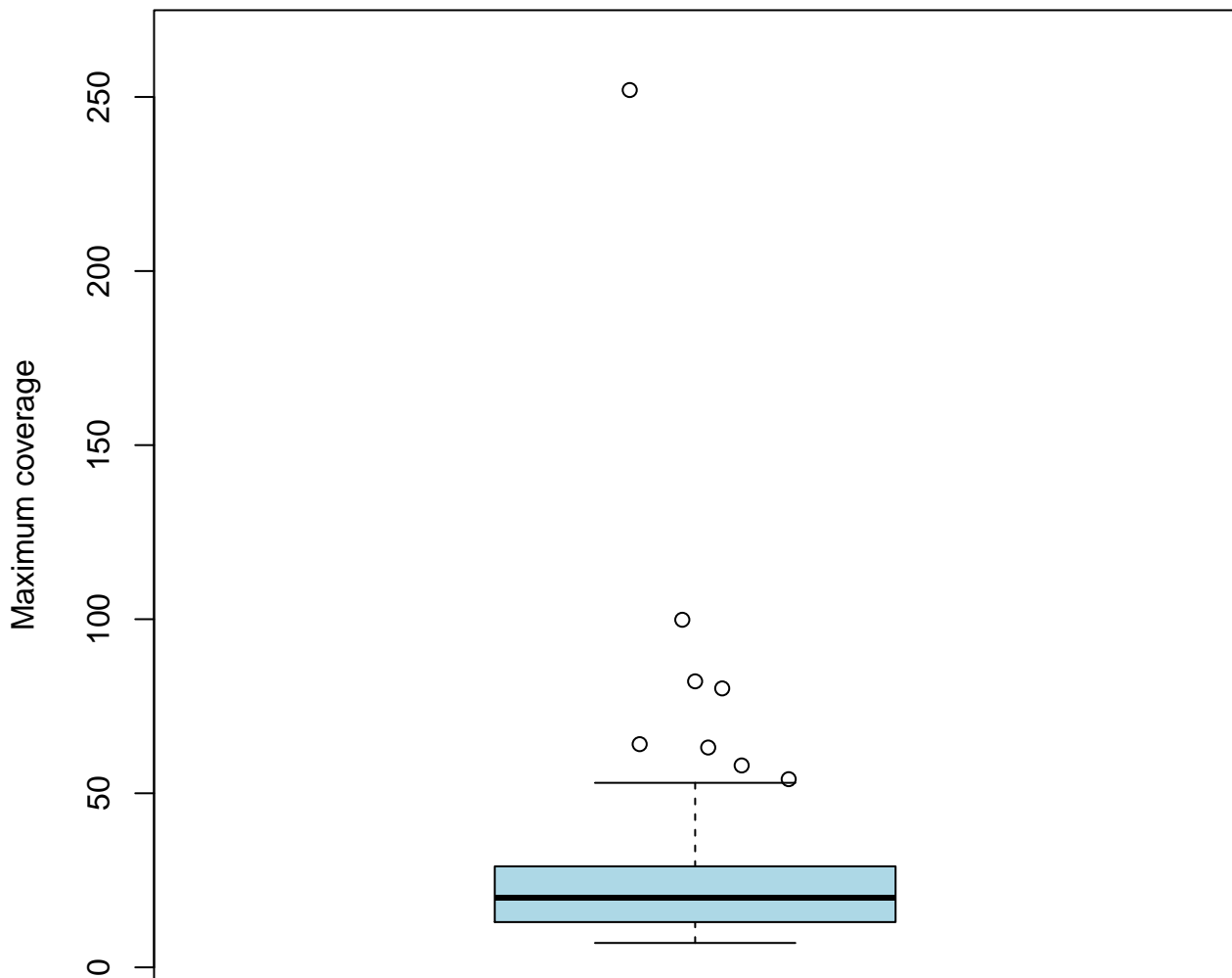
**jx in Intropolis but not "annotated" by knownGene:
148 samples; n jx: 104**



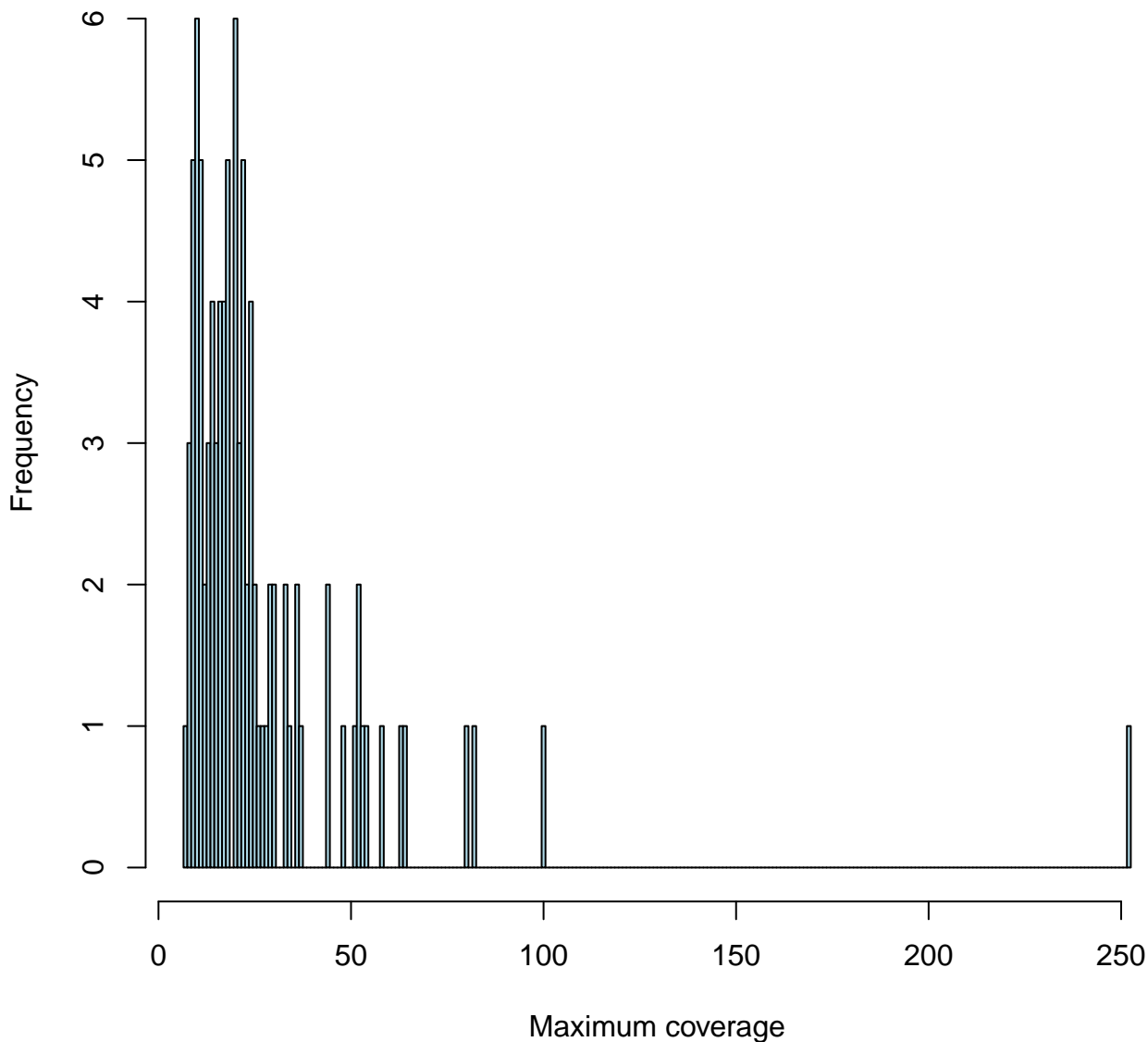
**jx in Intropolis but not "annotated" by knownGene:
148 samples; n jx: 104**



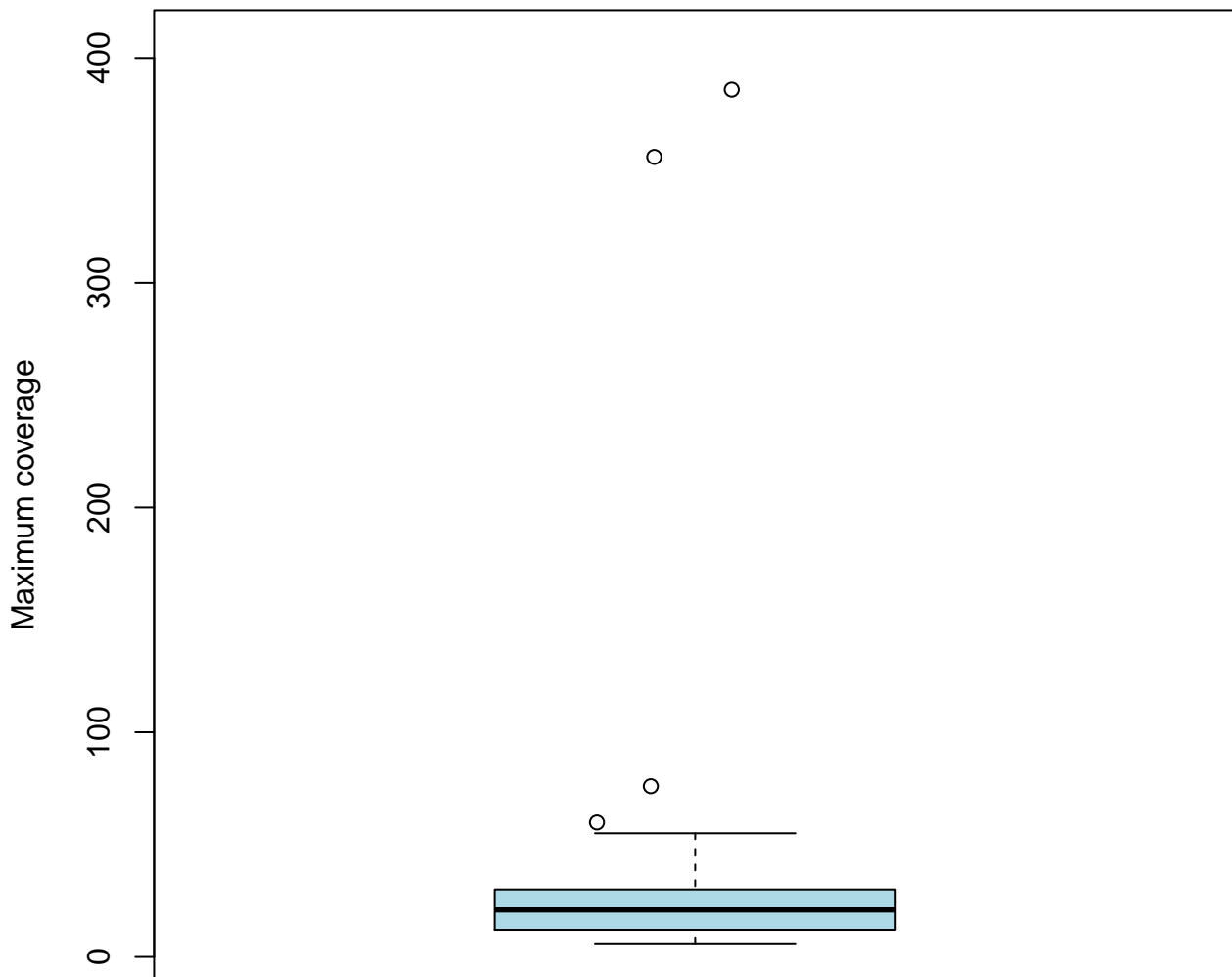
**jx in Intropolis but not "annotated" by knownGene:
149 samples; n jx: 95**



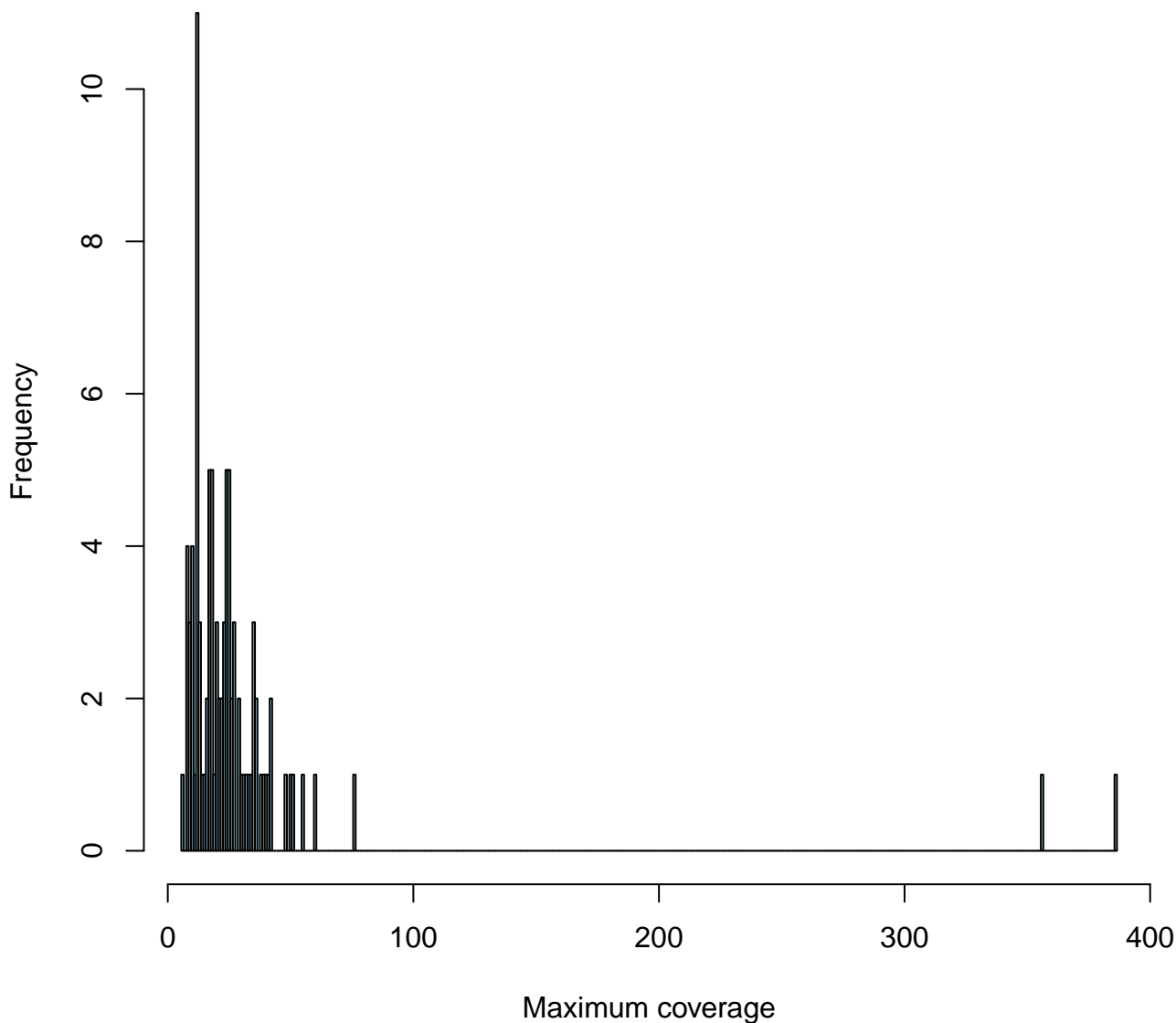
**jx in Intropolis but not "annotated" by knownGene:
149 samples; n jx: 95**



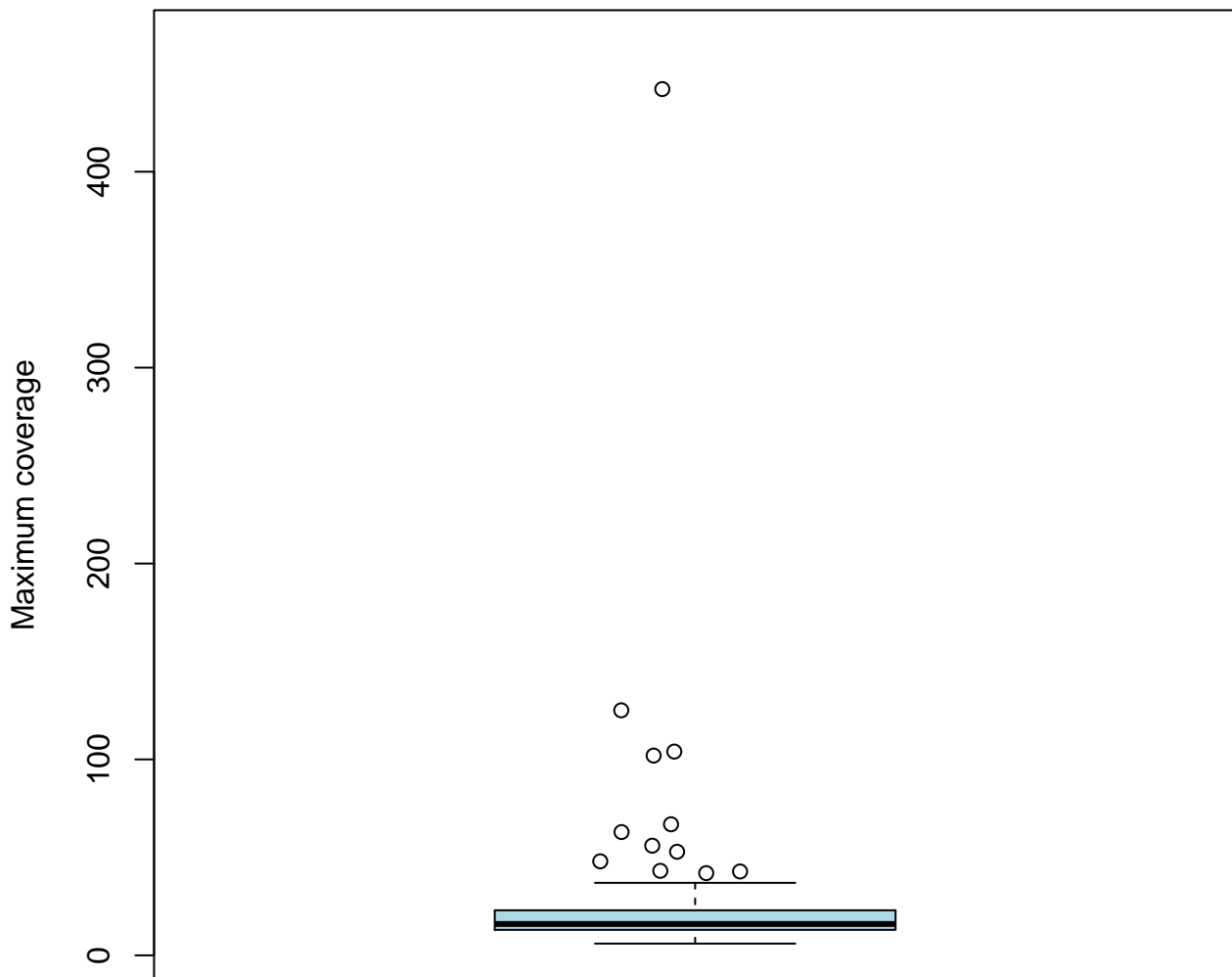
**jx in Intropolis but not "annotated" by knownGene:
150 samples; n jx: 93**



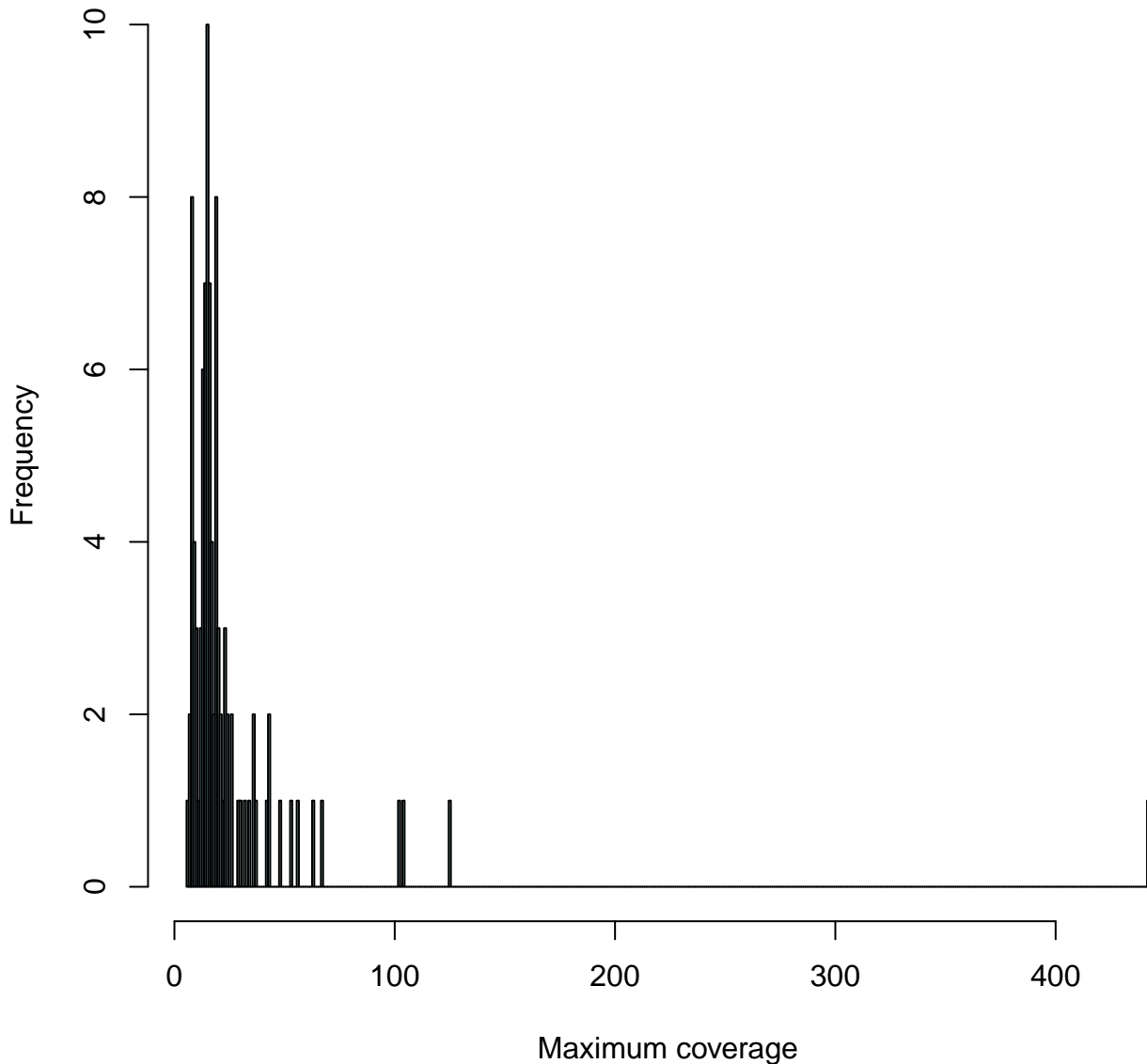
**jx in Intropolis but not "annotated" by knownGene:
150 samples; n jx: 93**



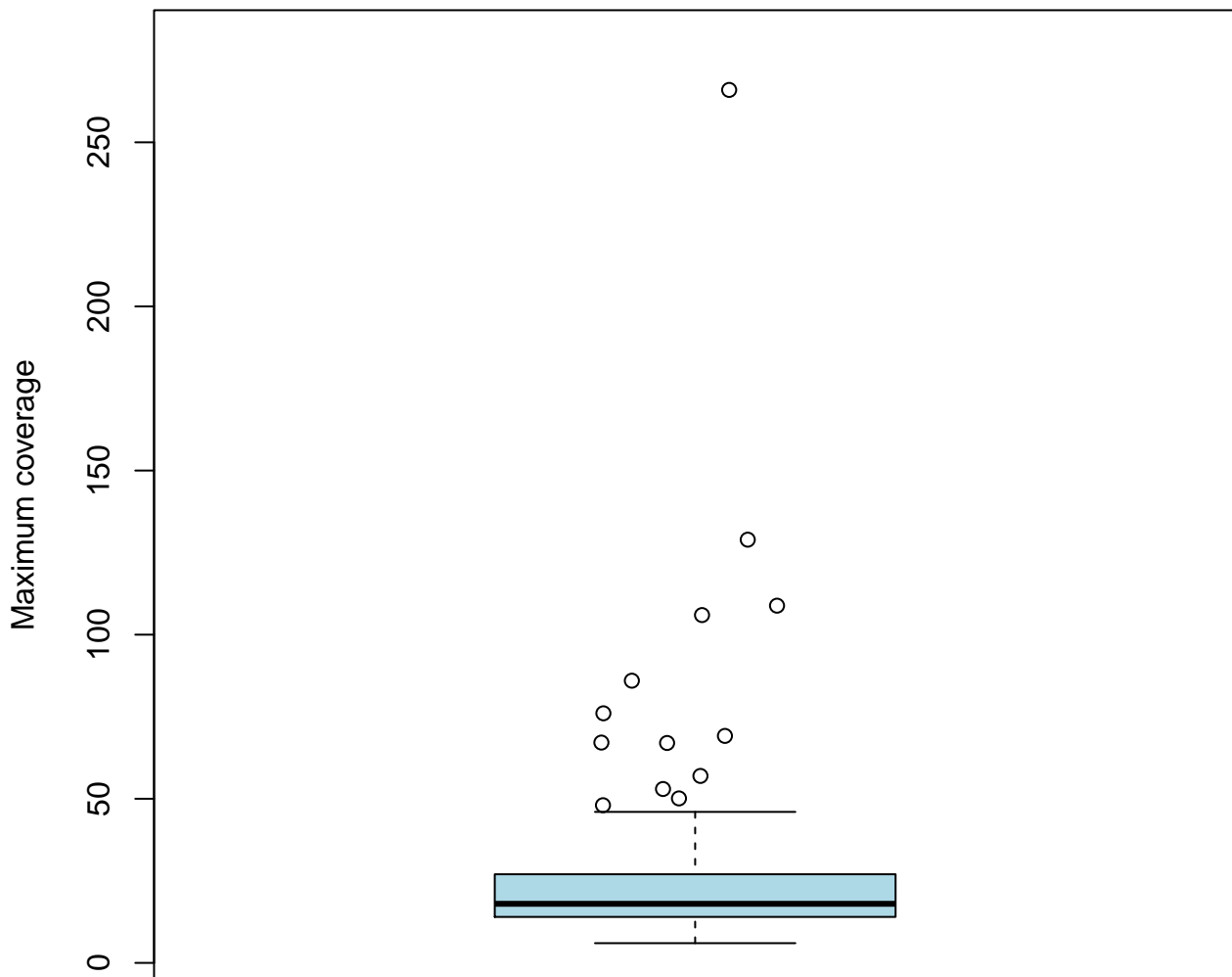
**jx in Intropolis but not "annotated" by knownGene:
151 samples; n jx: 98**



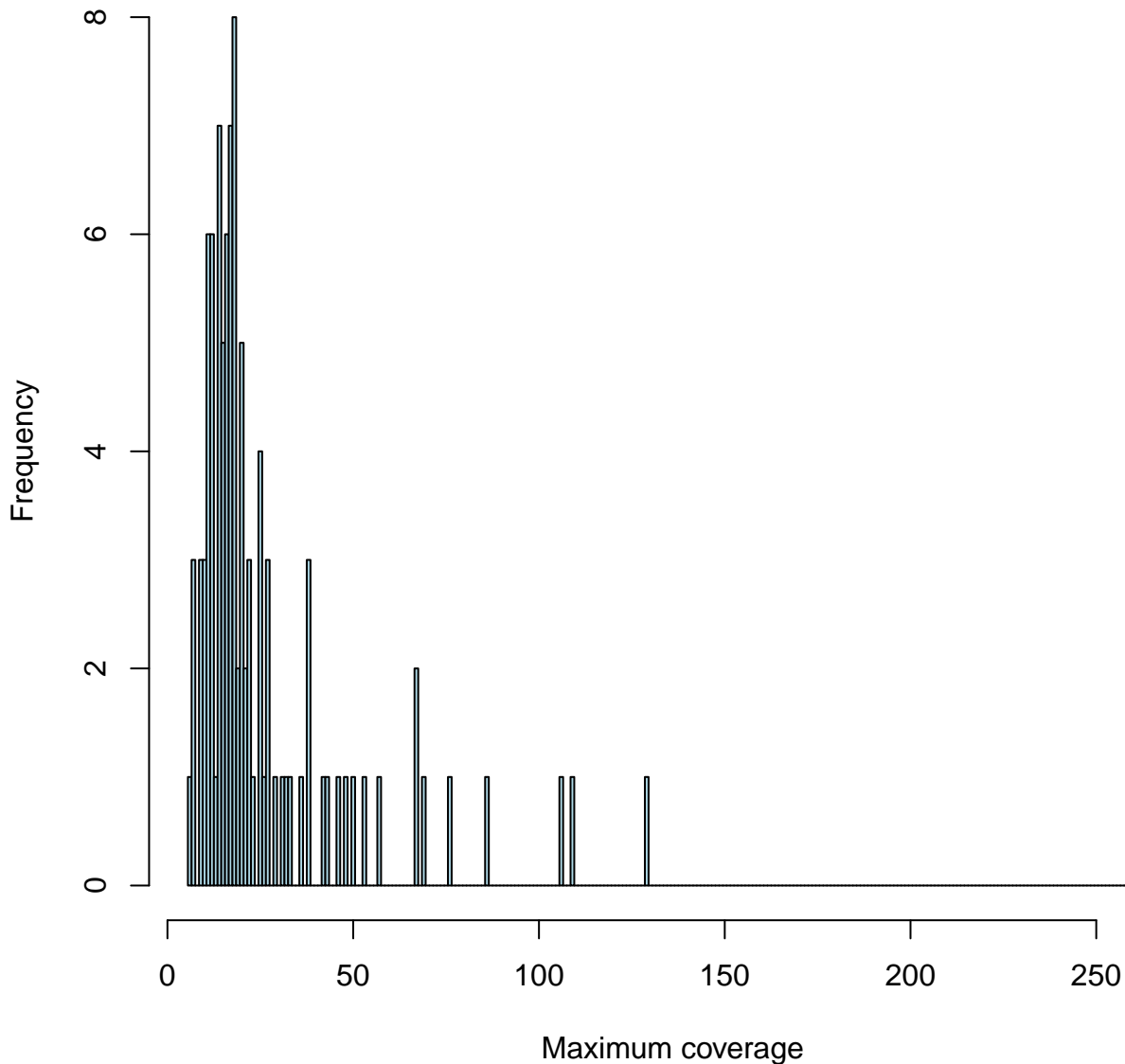
**jx in Intropolis but not "annotated" by knownGene:
151 samples; n jx: 98**



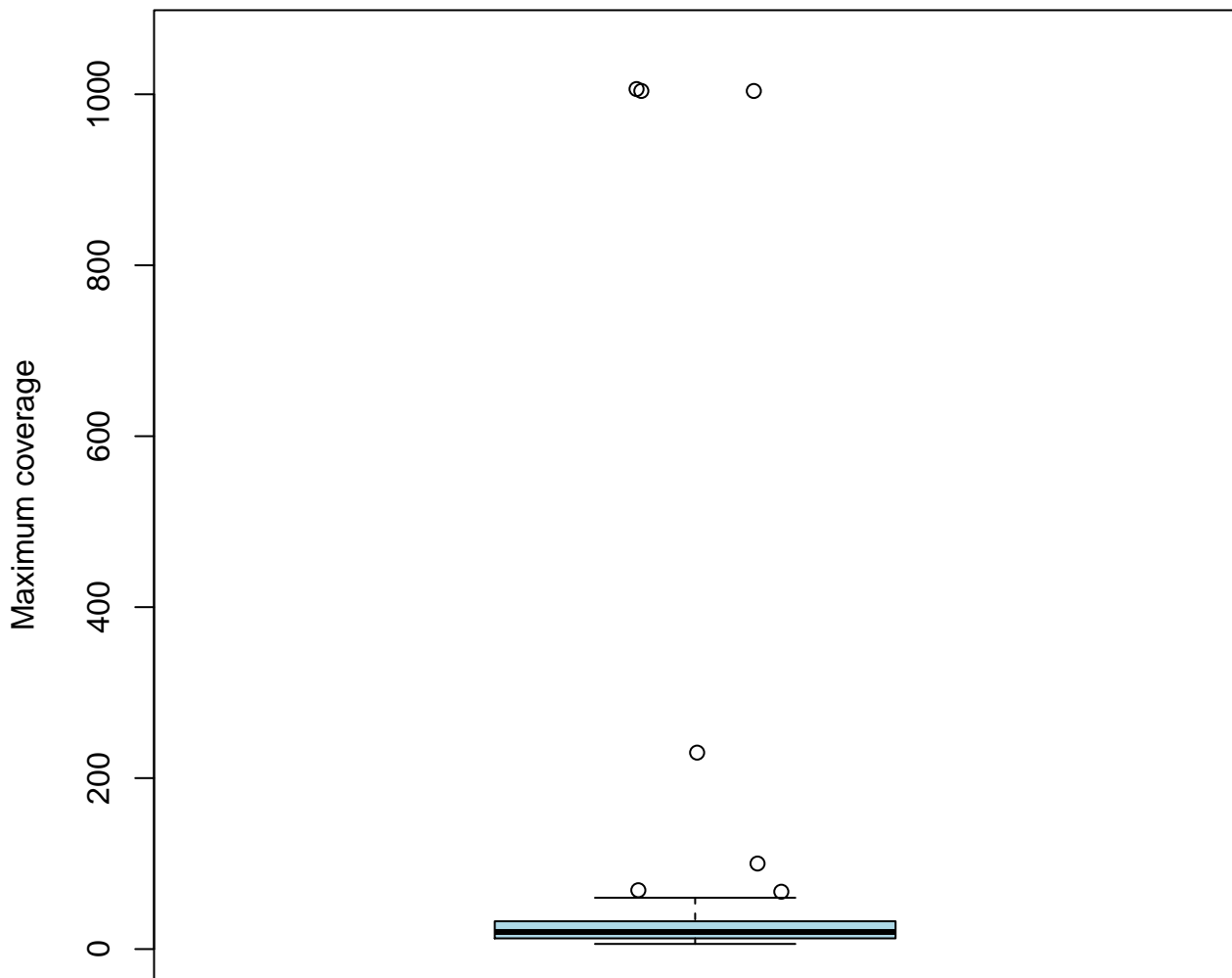
**jx in Intropolis but not "annotated" by knownGene:
152 samples; n jx: 101**



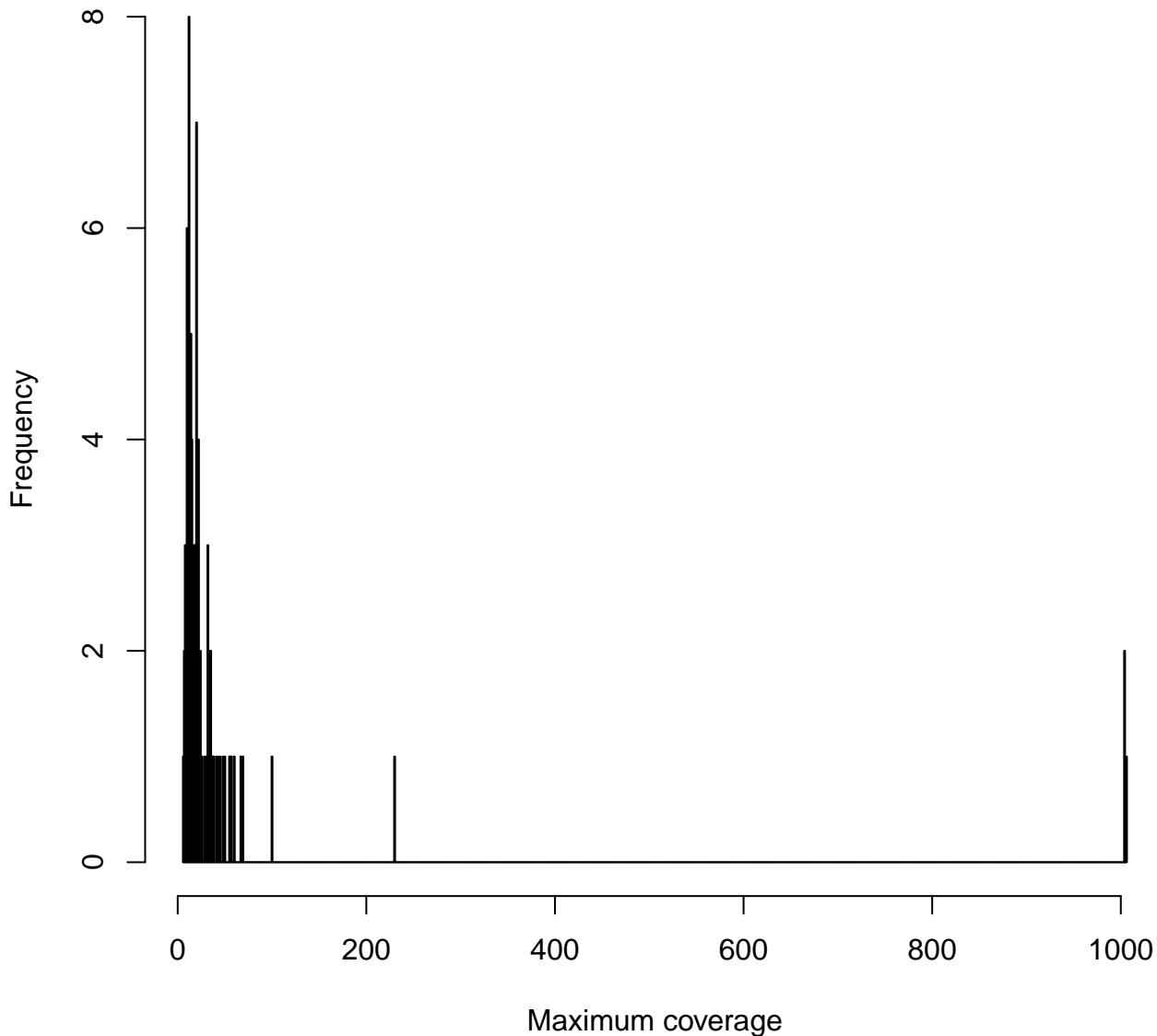
**jx in Intropolis but not "annotated" by knownGene:
152 samples; n jx: 101**



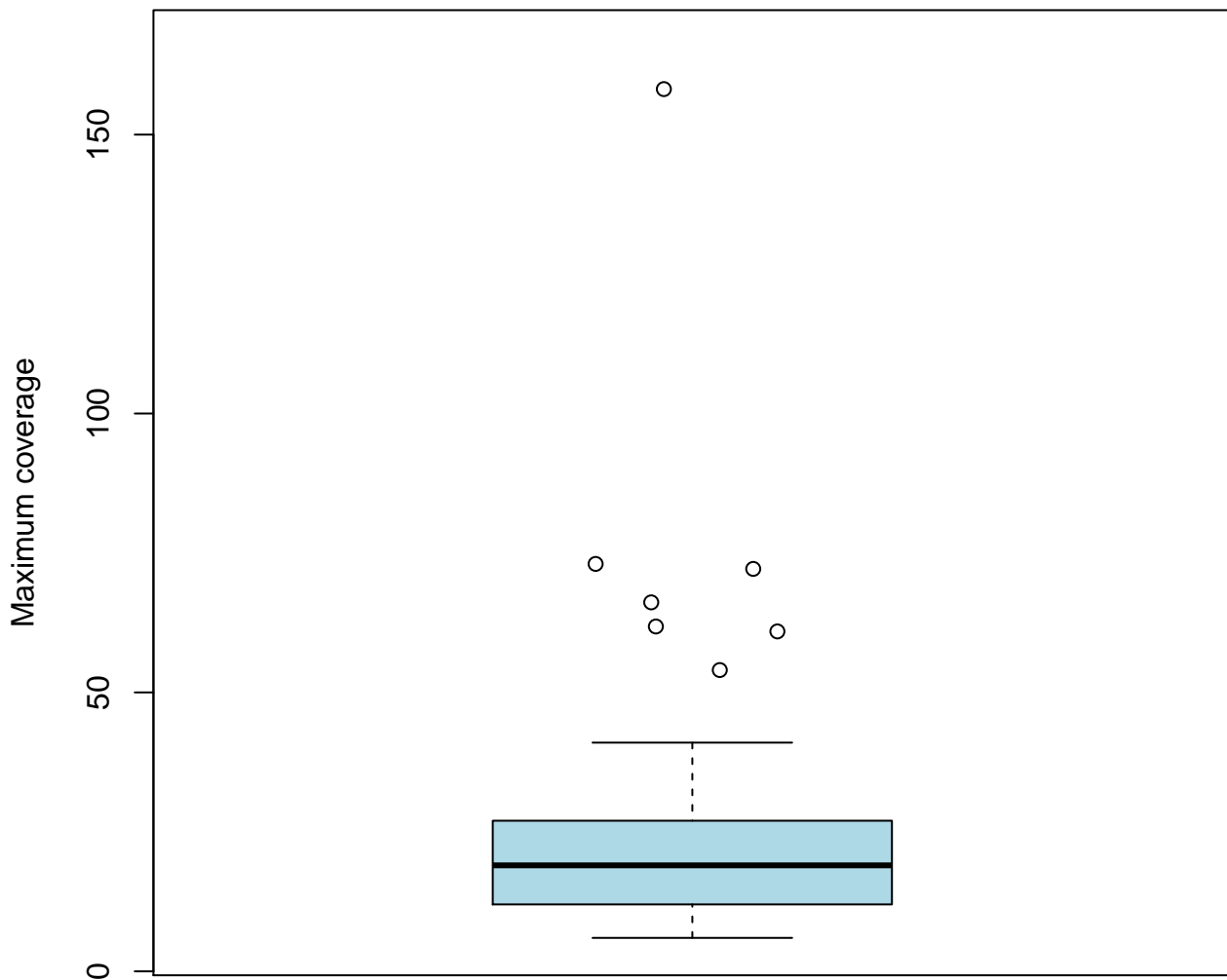
**jx in Intropolis but not "annotated" by knownGene:
153 samples; n jx: 99**



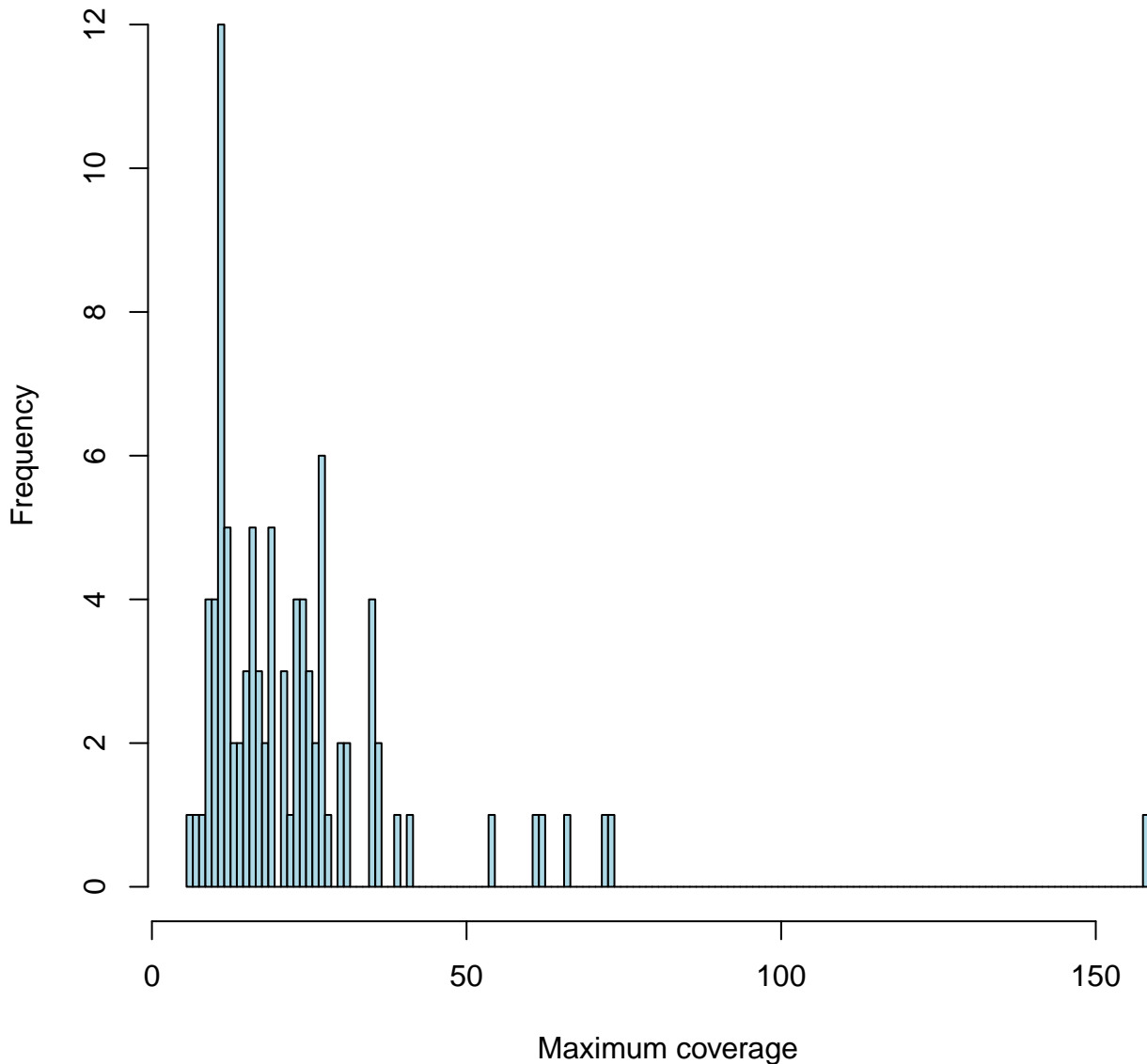
**jx in Intropolis but not "annotated" by knownGene:
153 samples; n jx: 99**



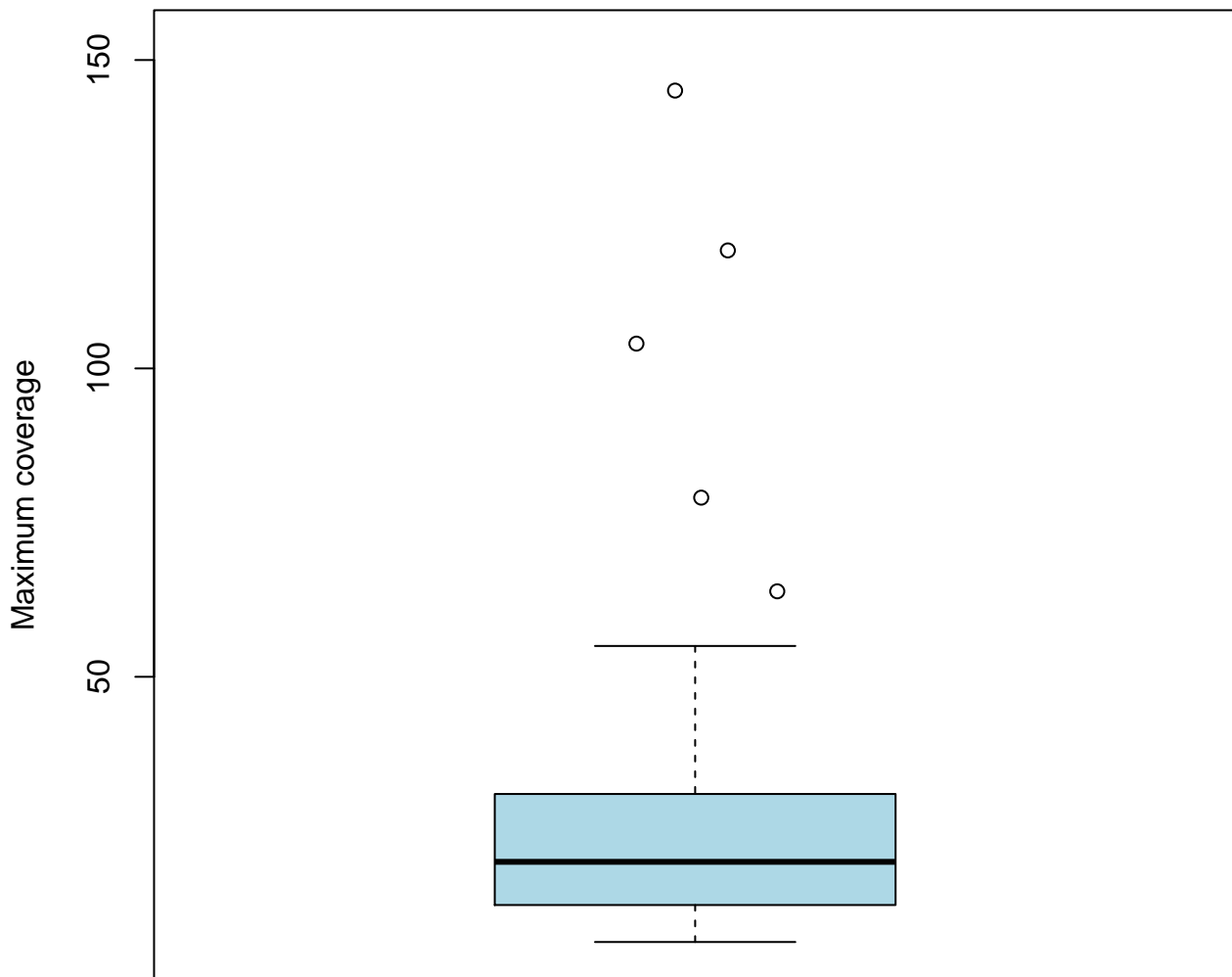
**jx in Intropolis but not "annotated" by knownGene:
154 samples; n jx: 93**



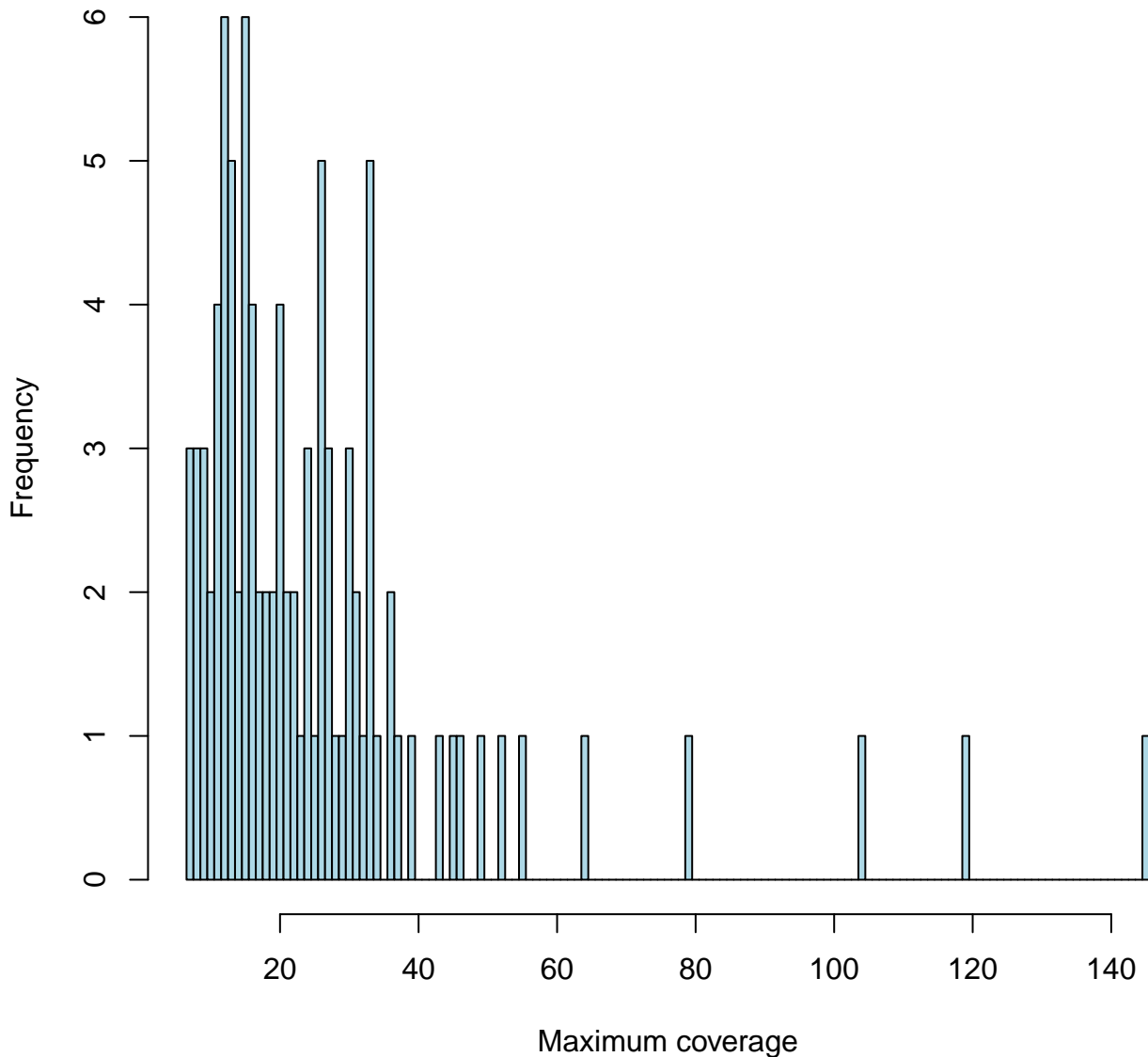
**jx in Intropolis but not "annotated" by knownGene:
154 samples; n jx: 93**



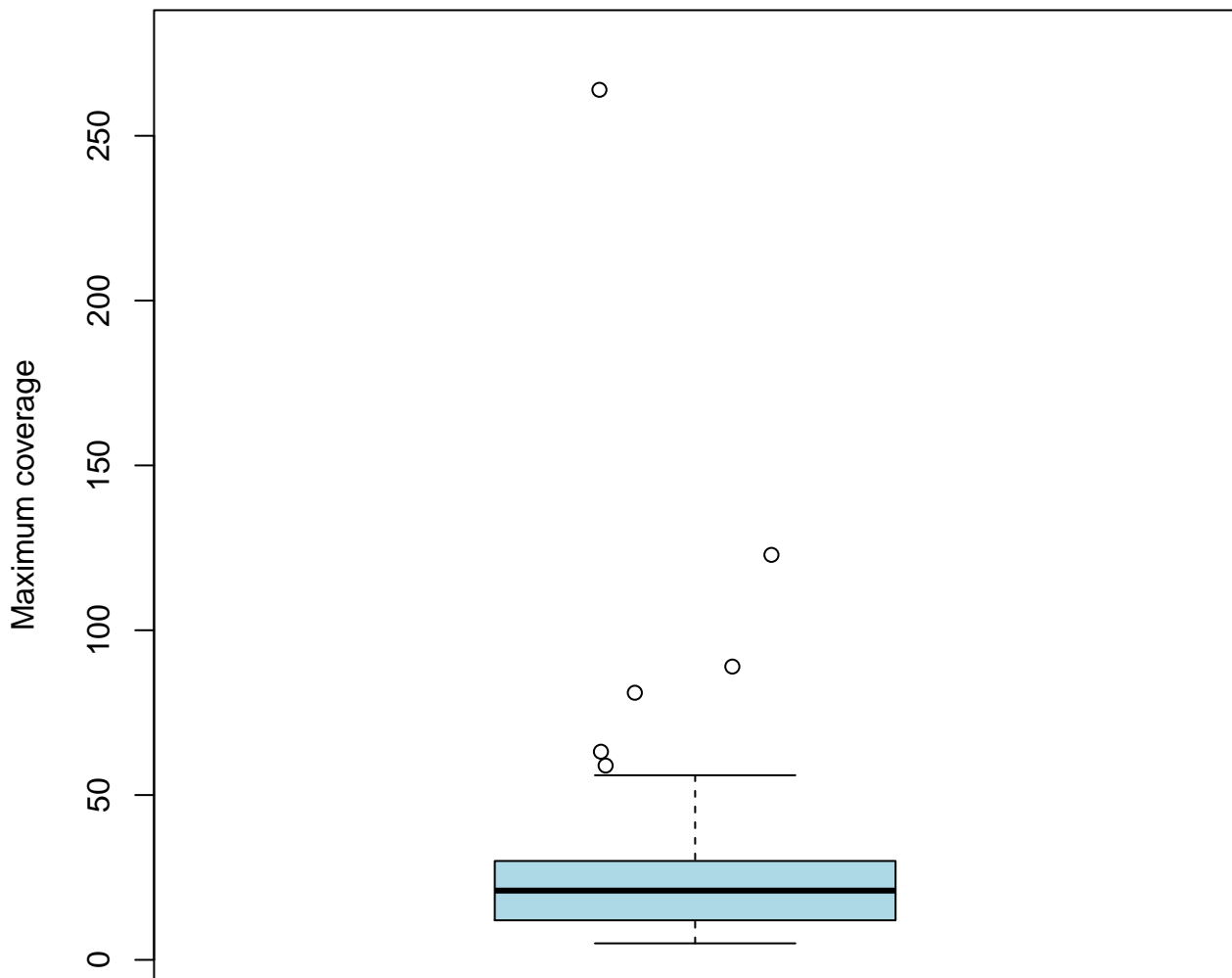
**jx in Intropolis but not "annotated" by knownGene:
155 samples; n jx: 94**



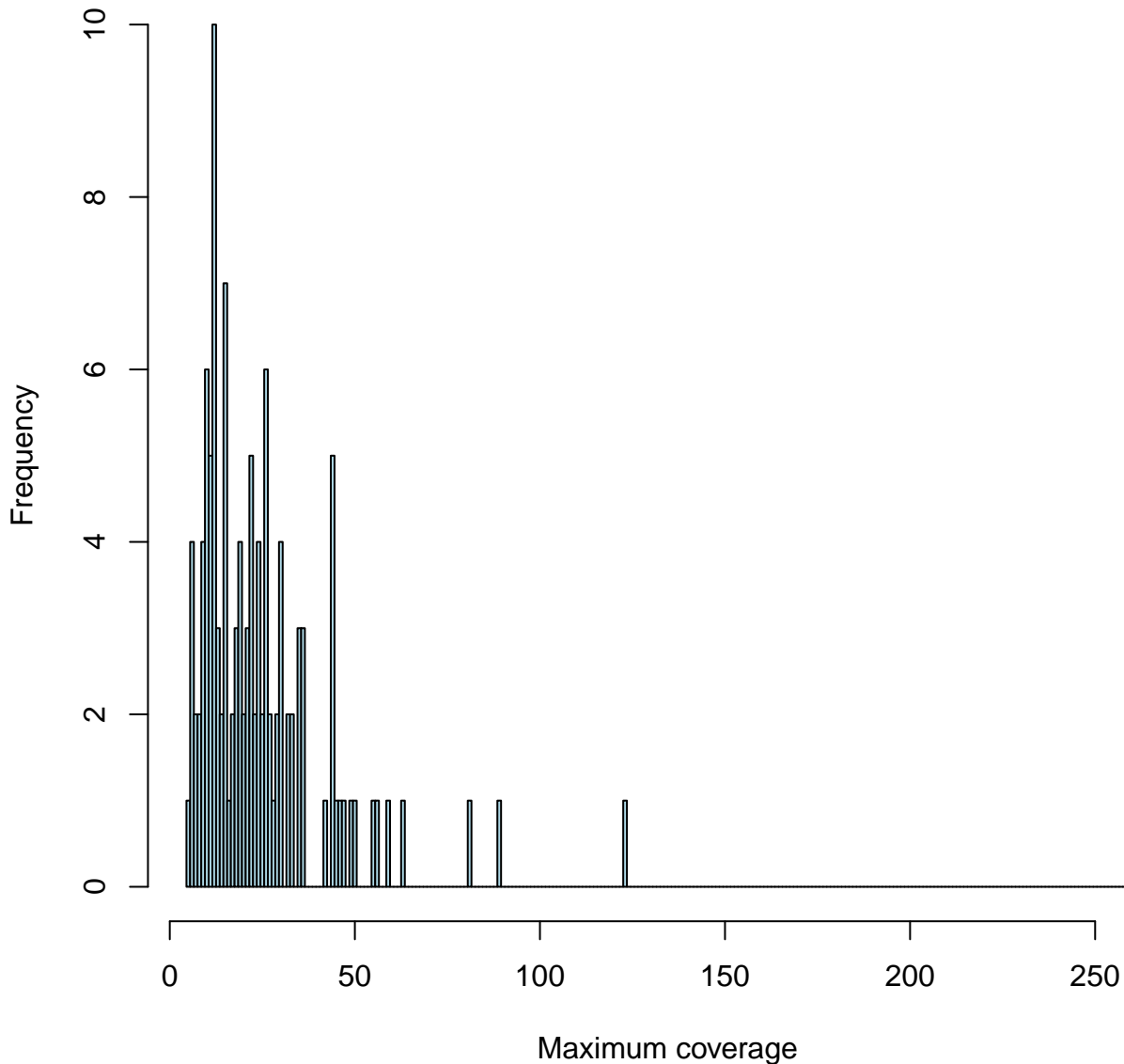
**jx in Intropolis but not "annotated" by knownGene:
155 samples; n jx: 94**



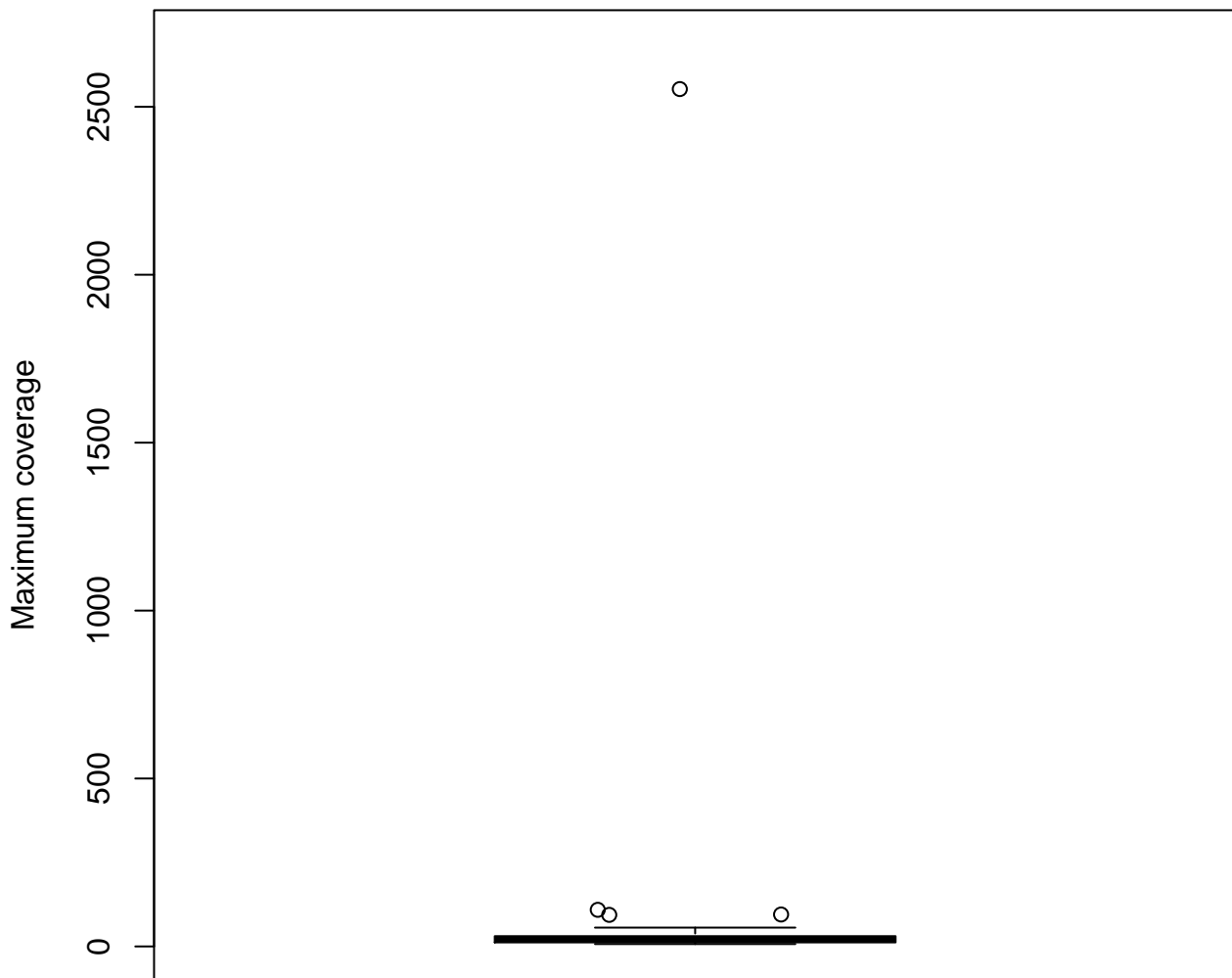
**jx in Intropolis but not "annotated" by knownGene:
156 samples; n jx: 118**



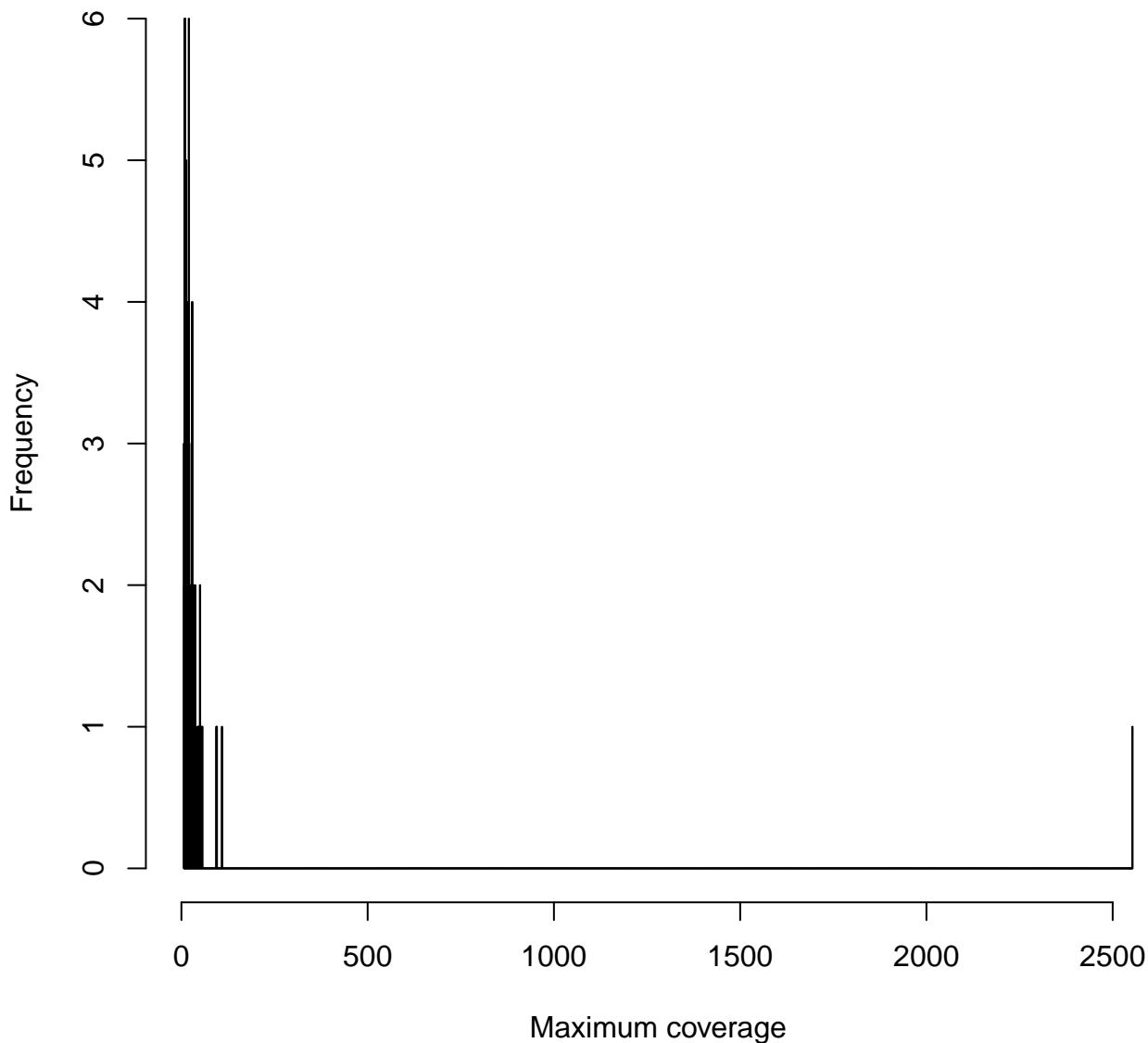
**jx in Intropolis but not "annotated" by knownGene:
156 samples; n jx: 118**



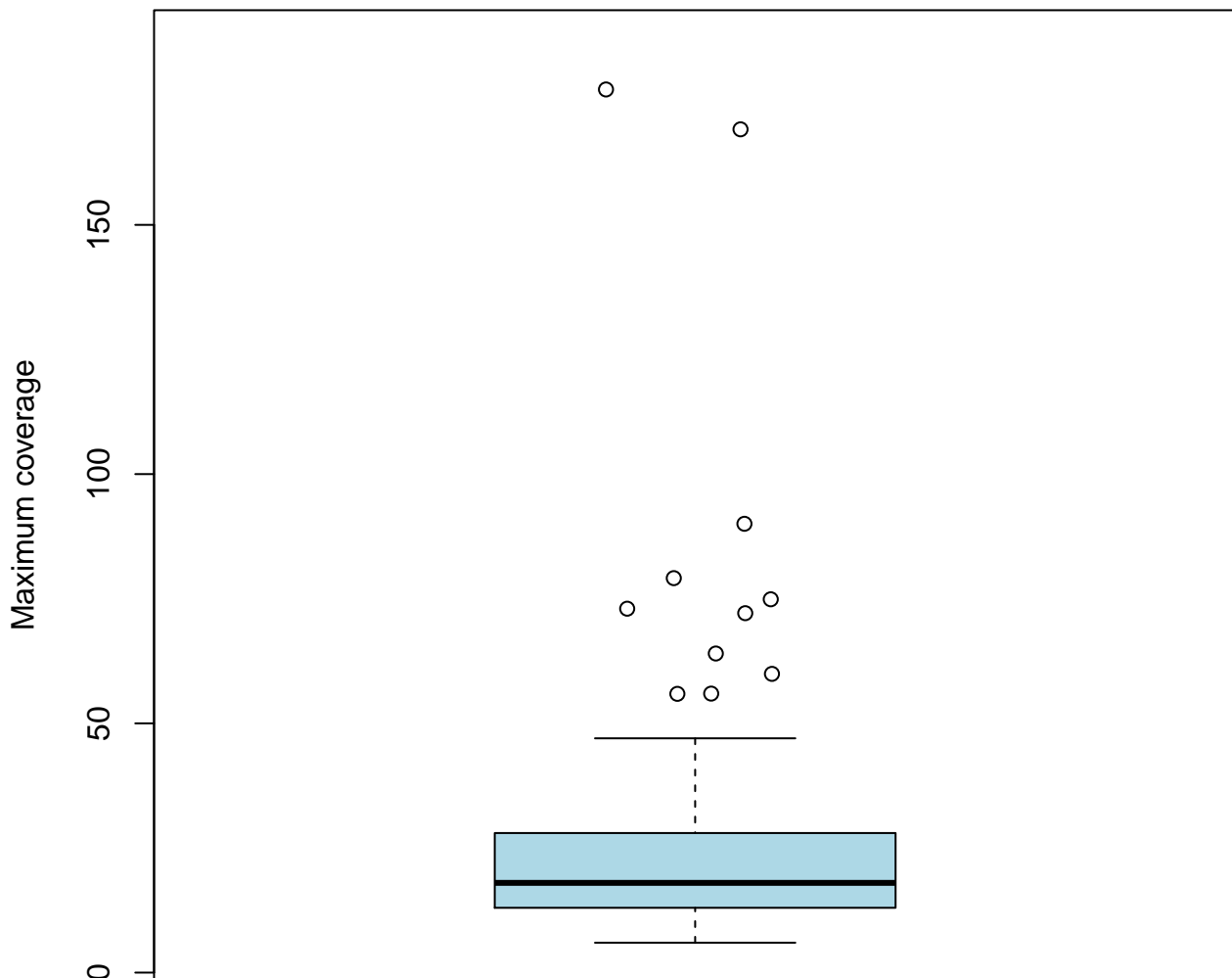
**jx in Intropolis but not "annotated" by knownGene:
157 samples; n jx: 88**



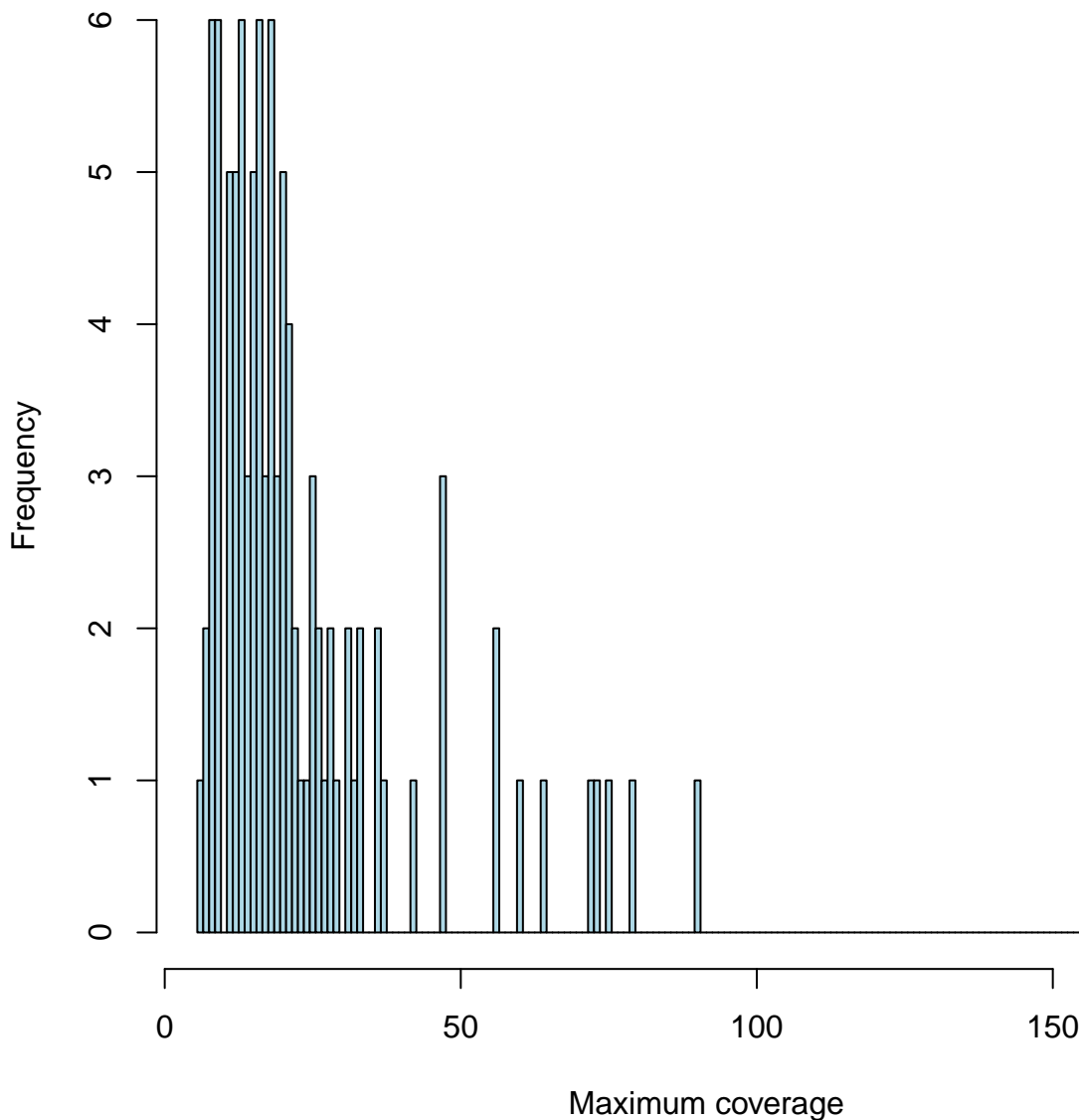
**jx in Intropolis but not "annotated" by knownGene:
157 samples; n jx: 88**



**jx in Intropolis but not "annotated" by knownGene:
158 samples; n jx: 102**



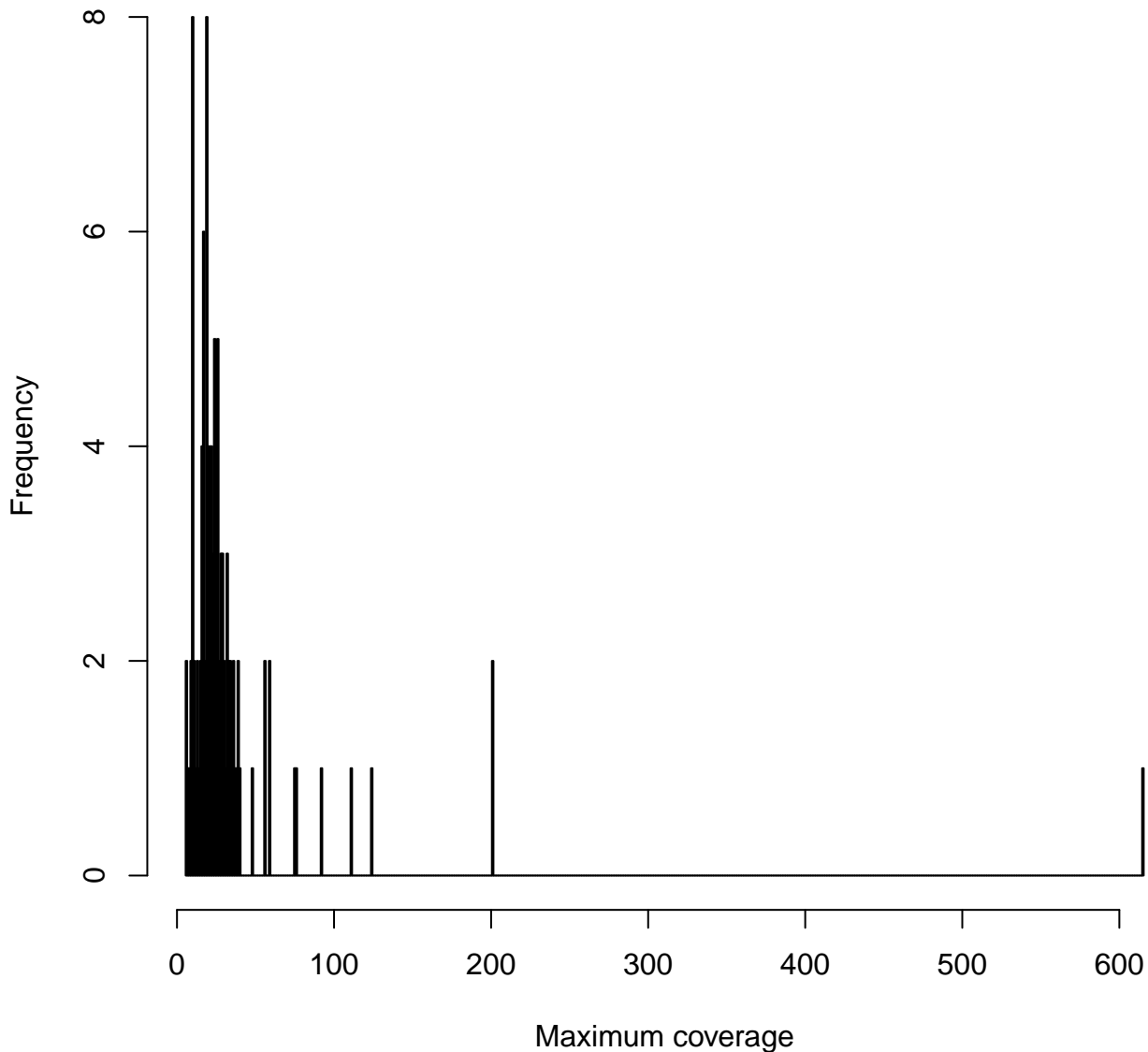
**jx in Intropolis but not "annotated" by knownGene:
158 samples; n jx: 102**



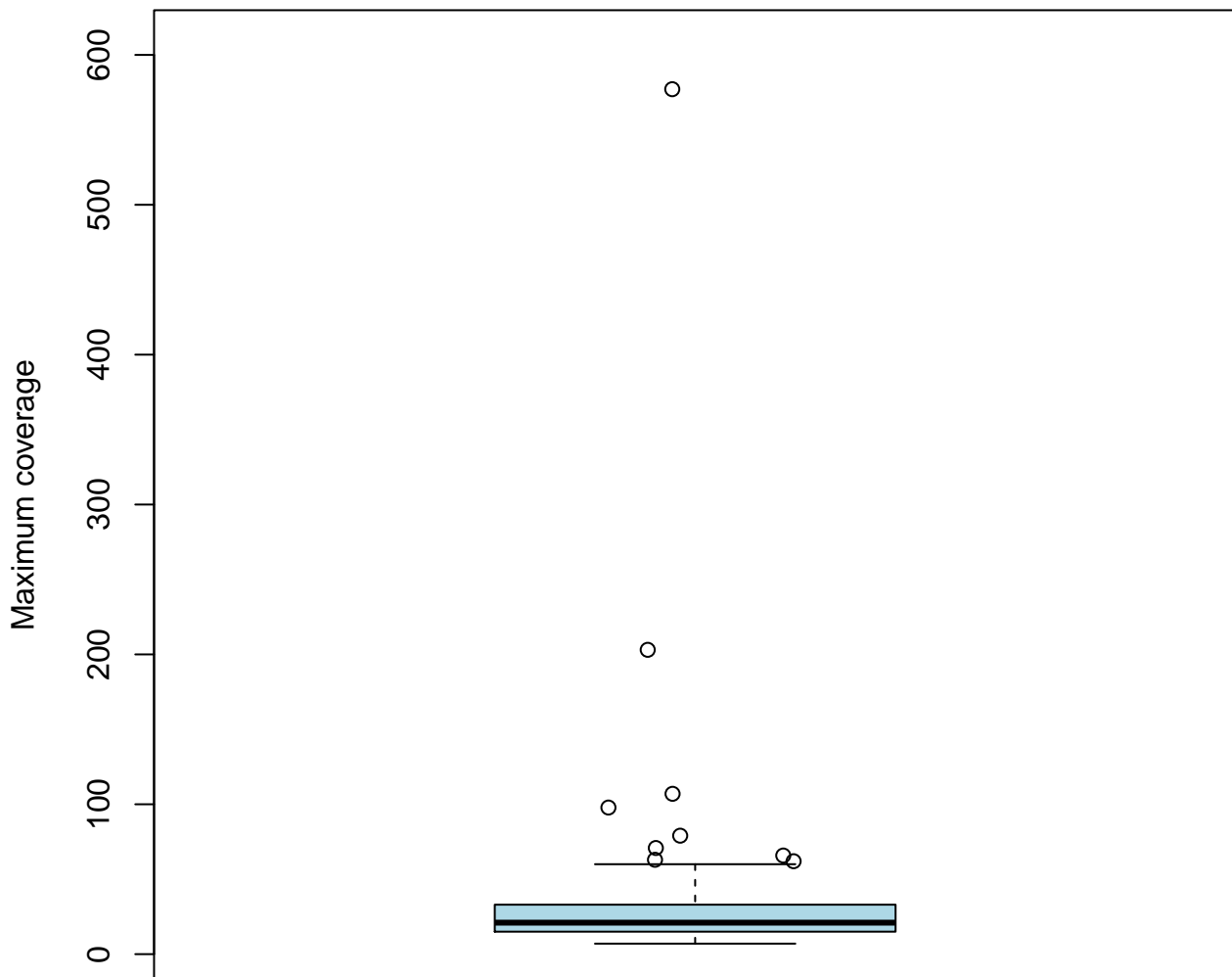
159 samples; n jx: 105



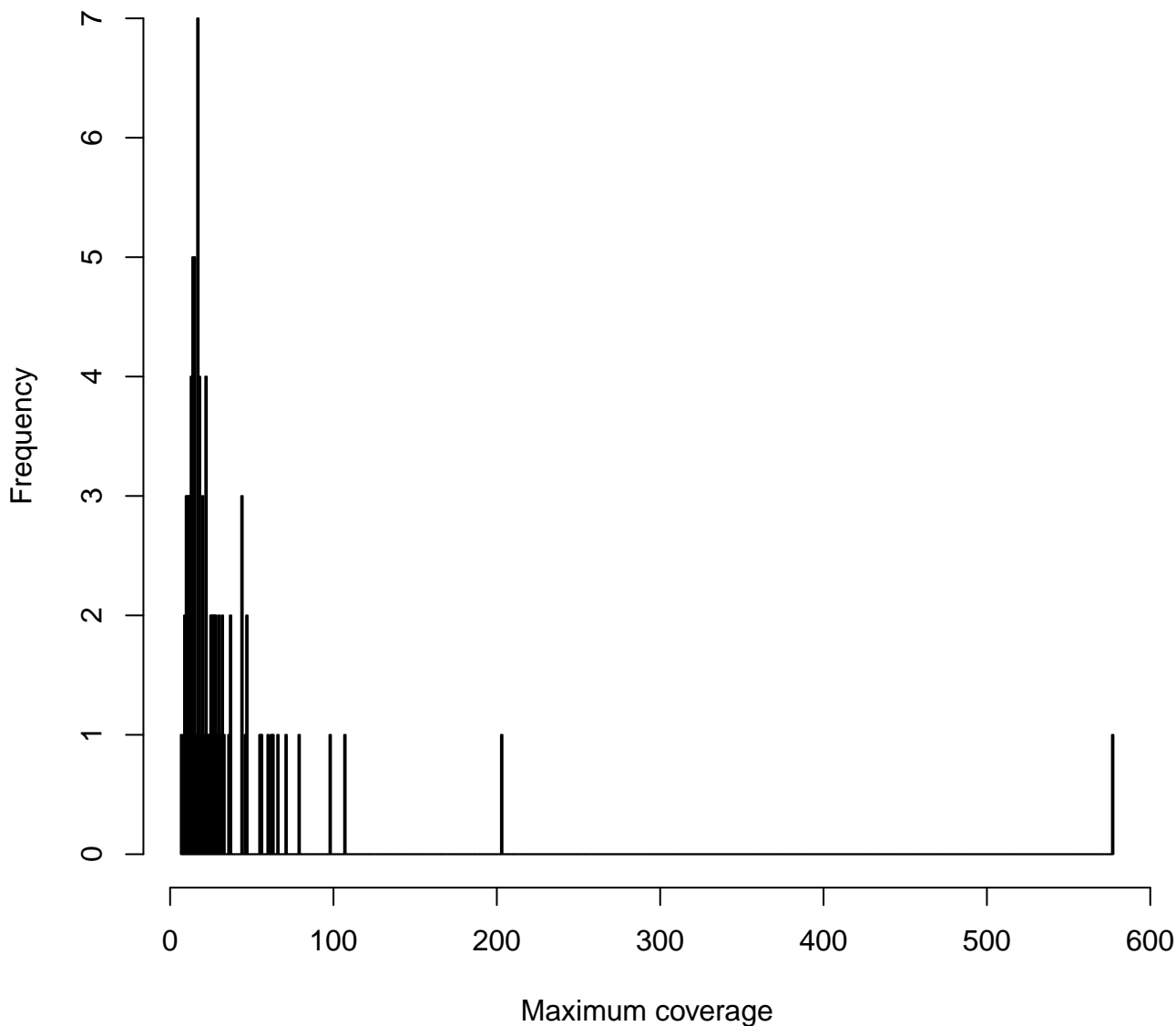
**jx in Intropolis but not "annotated" by knownGene:
159 samples; n jx: 105**



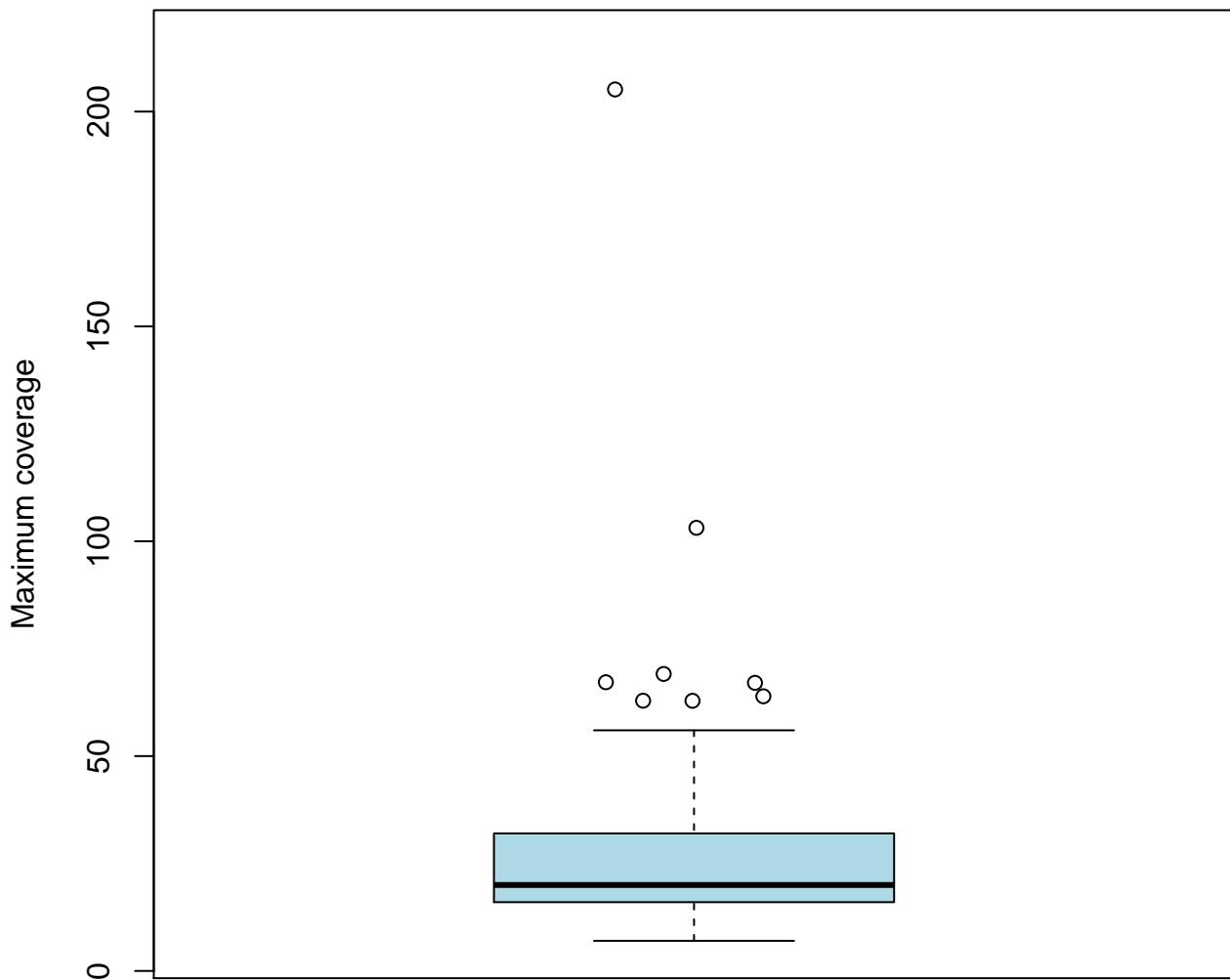
**jx in Intropolis but not "annotated" by knownGene:
160 samples; n jx: 85**



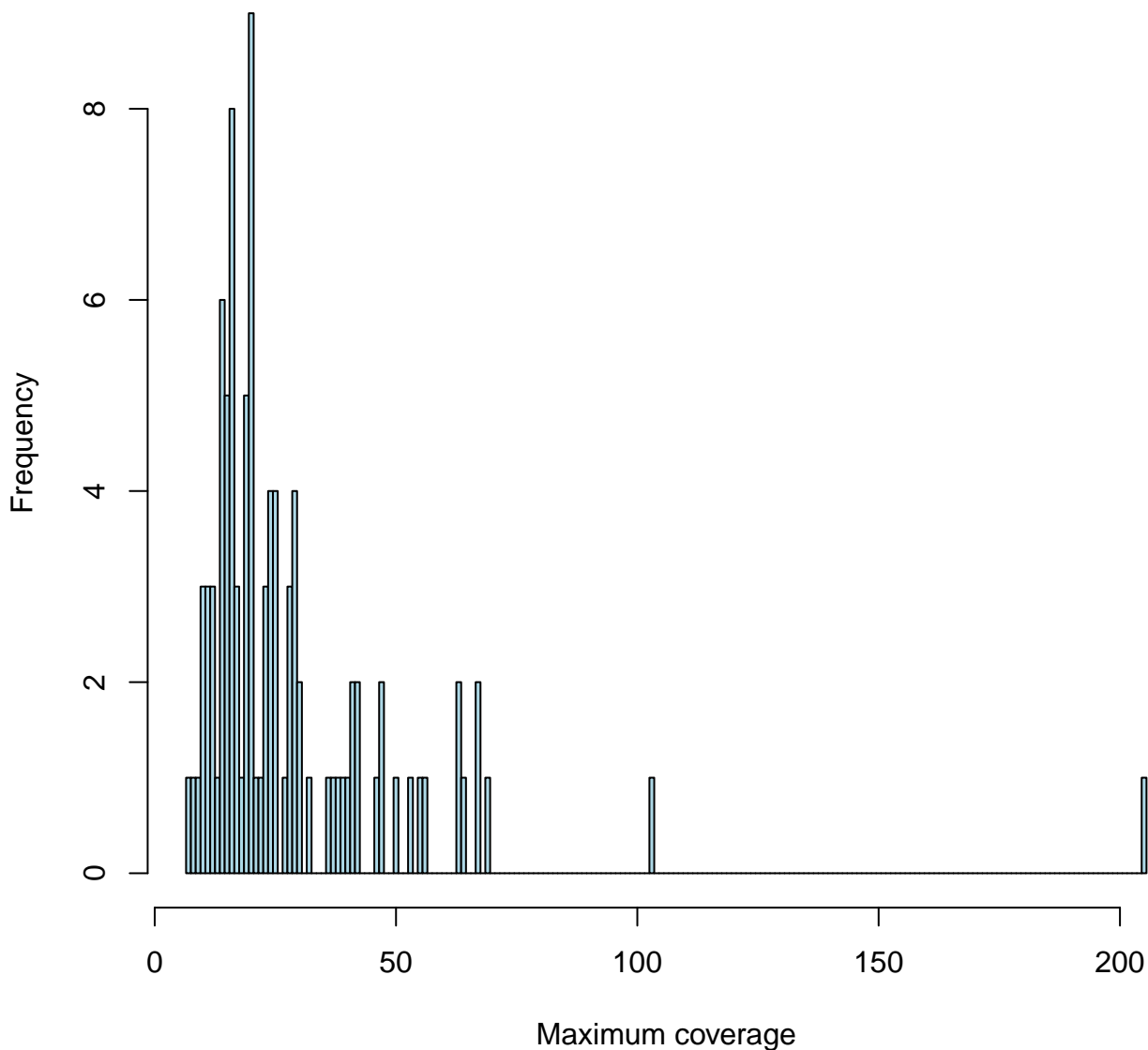
**jx in Intropolis but not "annotated" by knownGene:
160 samples; n jx: 85**



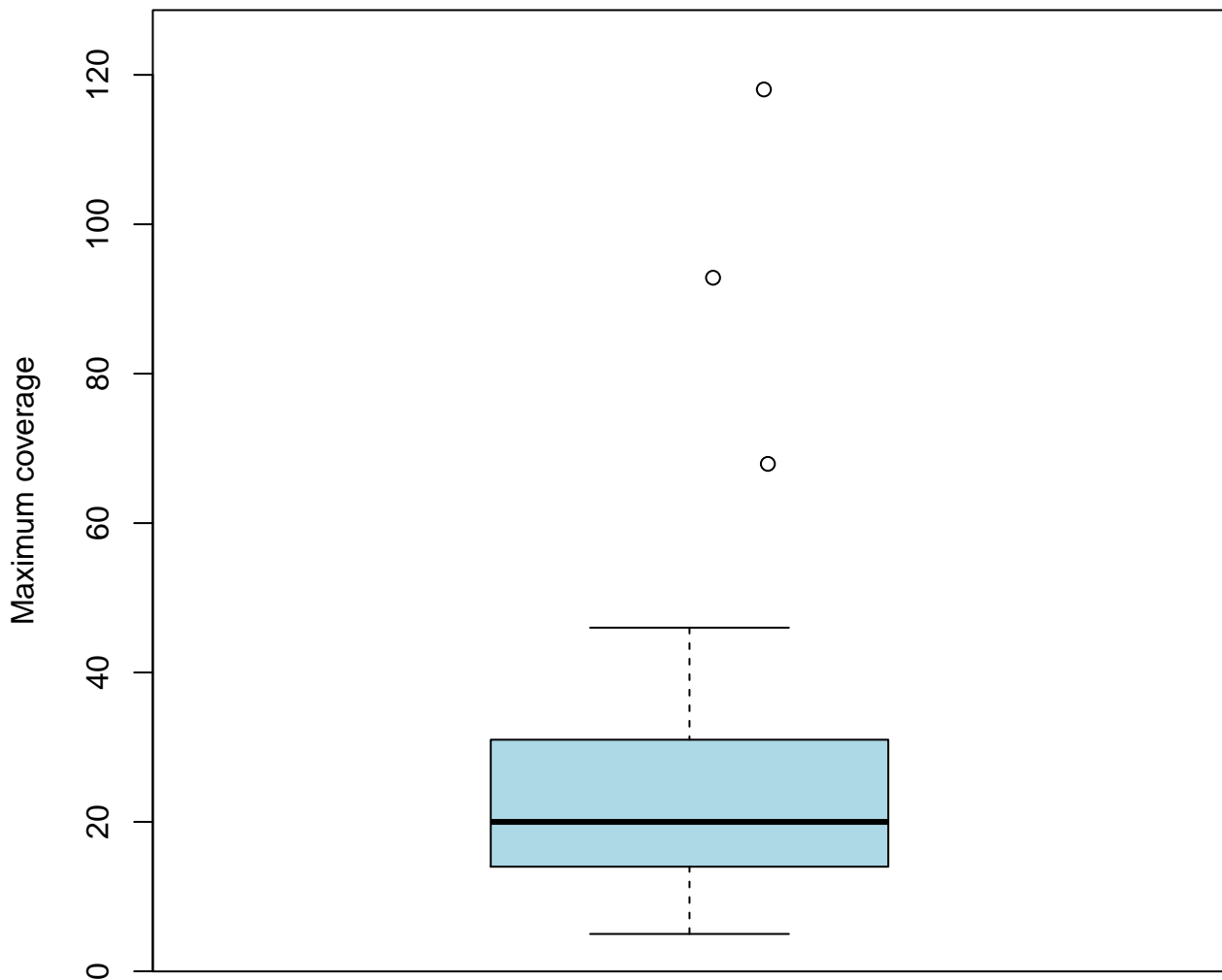
**jx in Intropolis but not "annotated" by knownGene:
161 samples; n jx: 98**



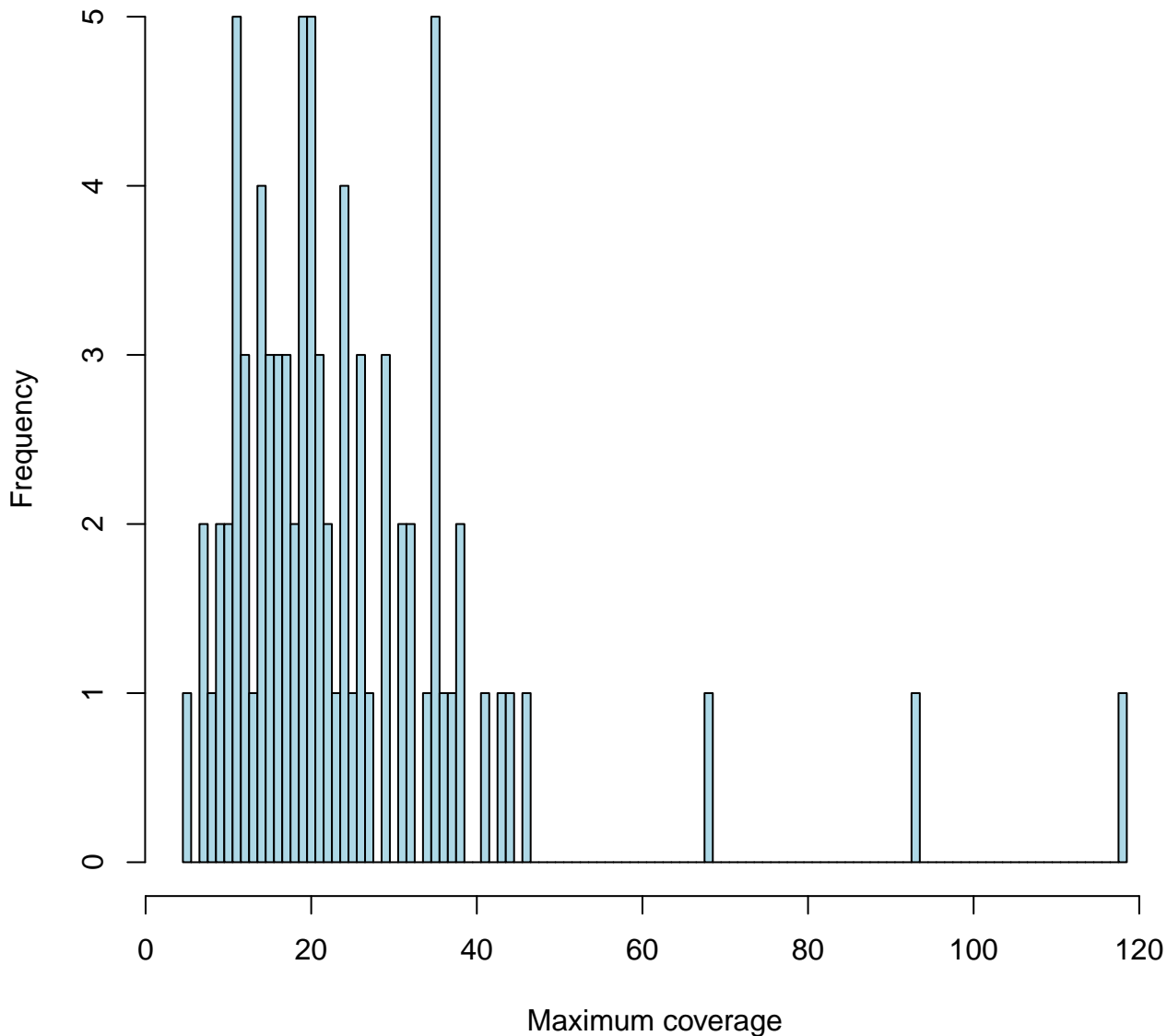
**jx in Intropolis but not "annotated" by knownGene:
161 samples; n jx: 98**



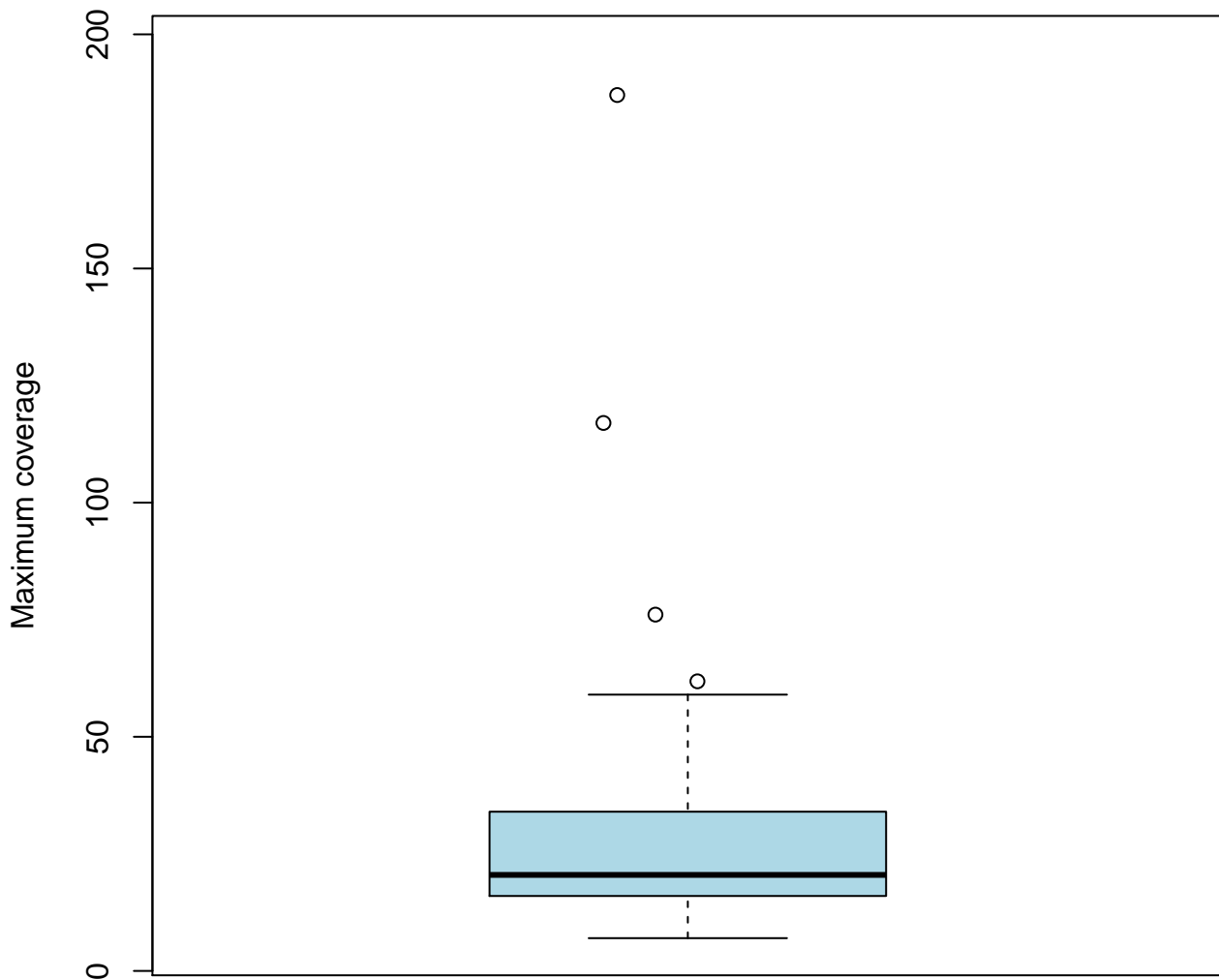
**jx in Intropolis but not "annotated" by knownGene:
162 samples; n jx: 81**



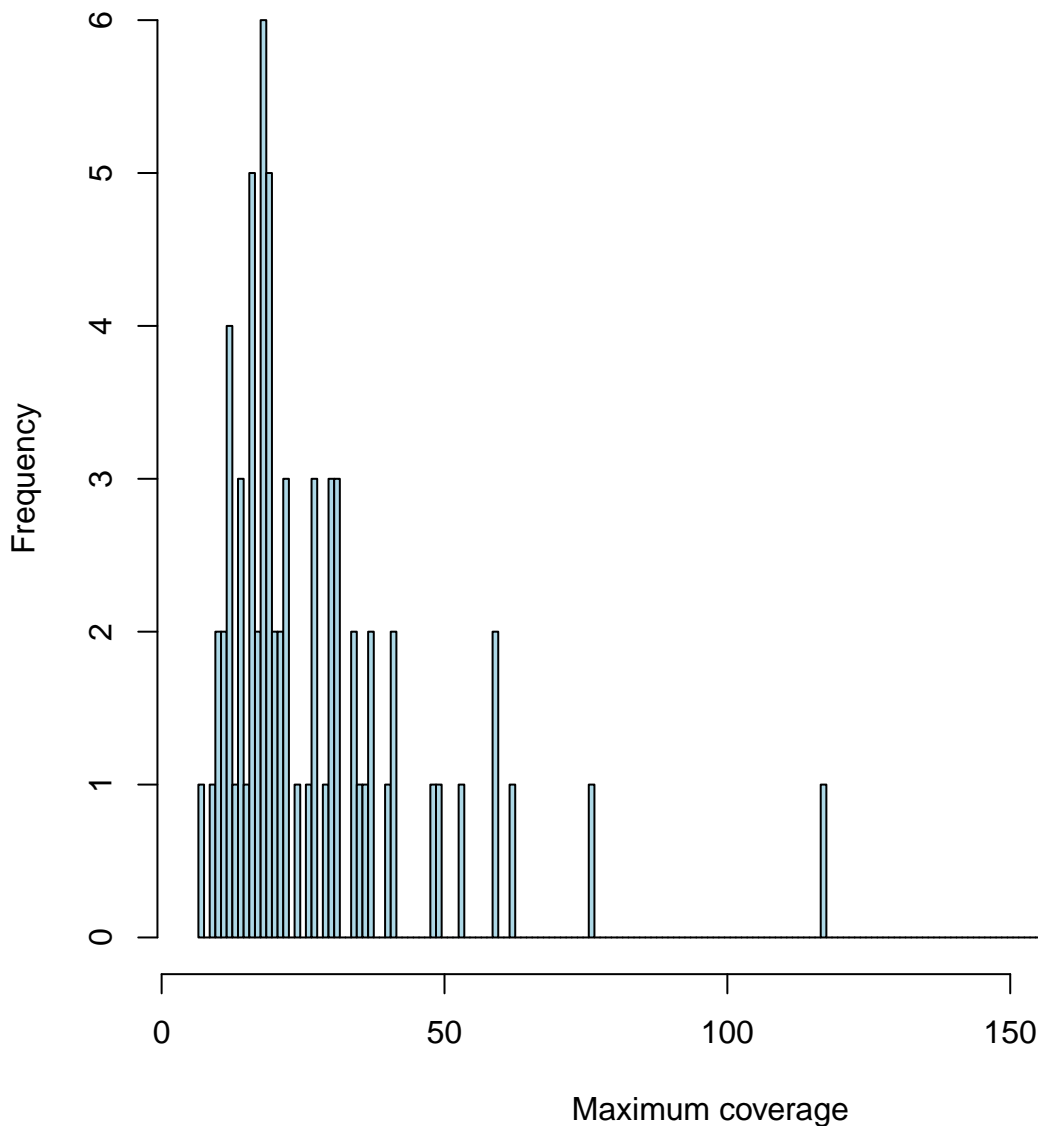
**jx in Intropolis but not "annotated" by knownGene:
162 samples; n jx: 81**



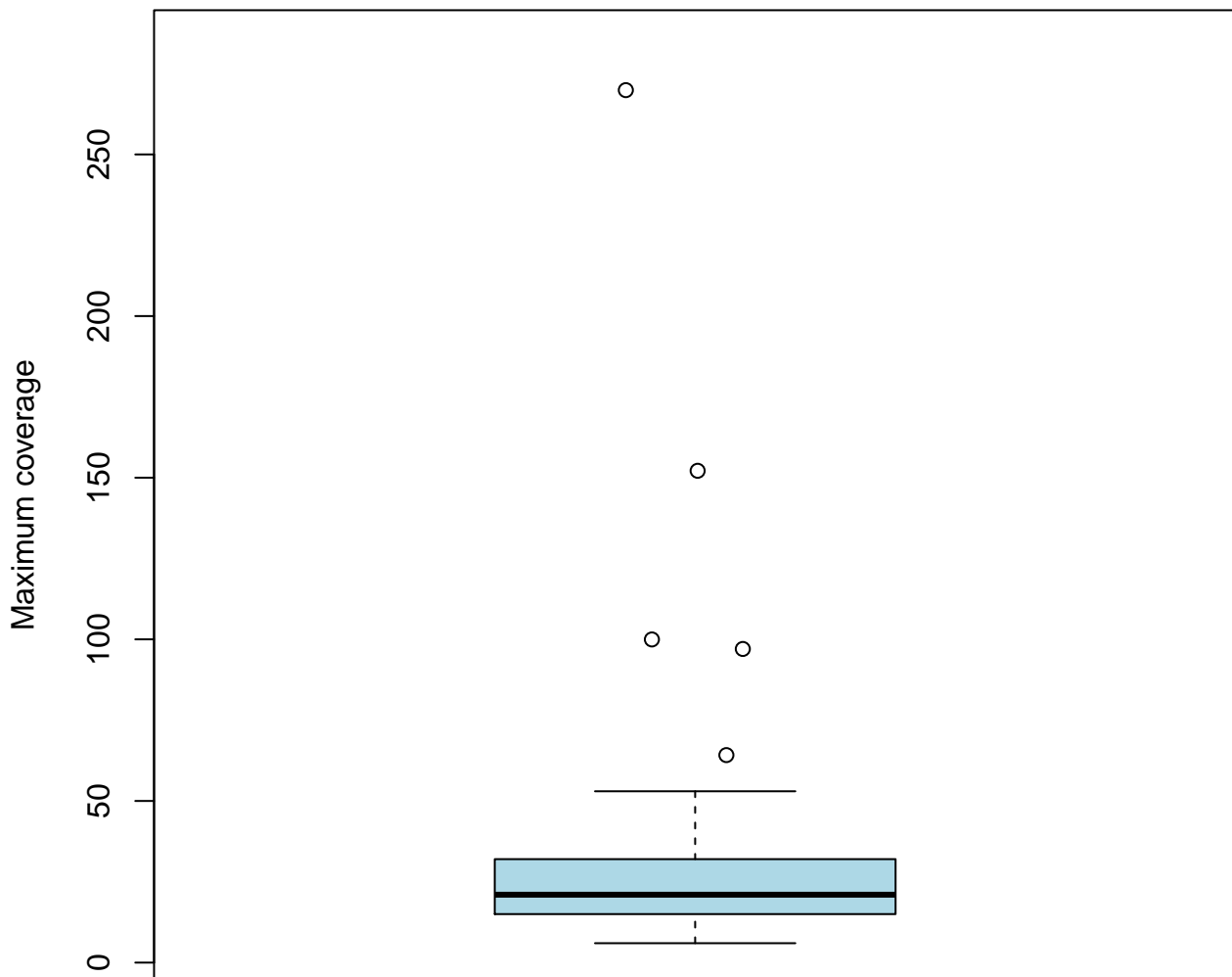
**jx in Intropolis but not "annotated" by knownGene:
163 samples; n jx: 70**



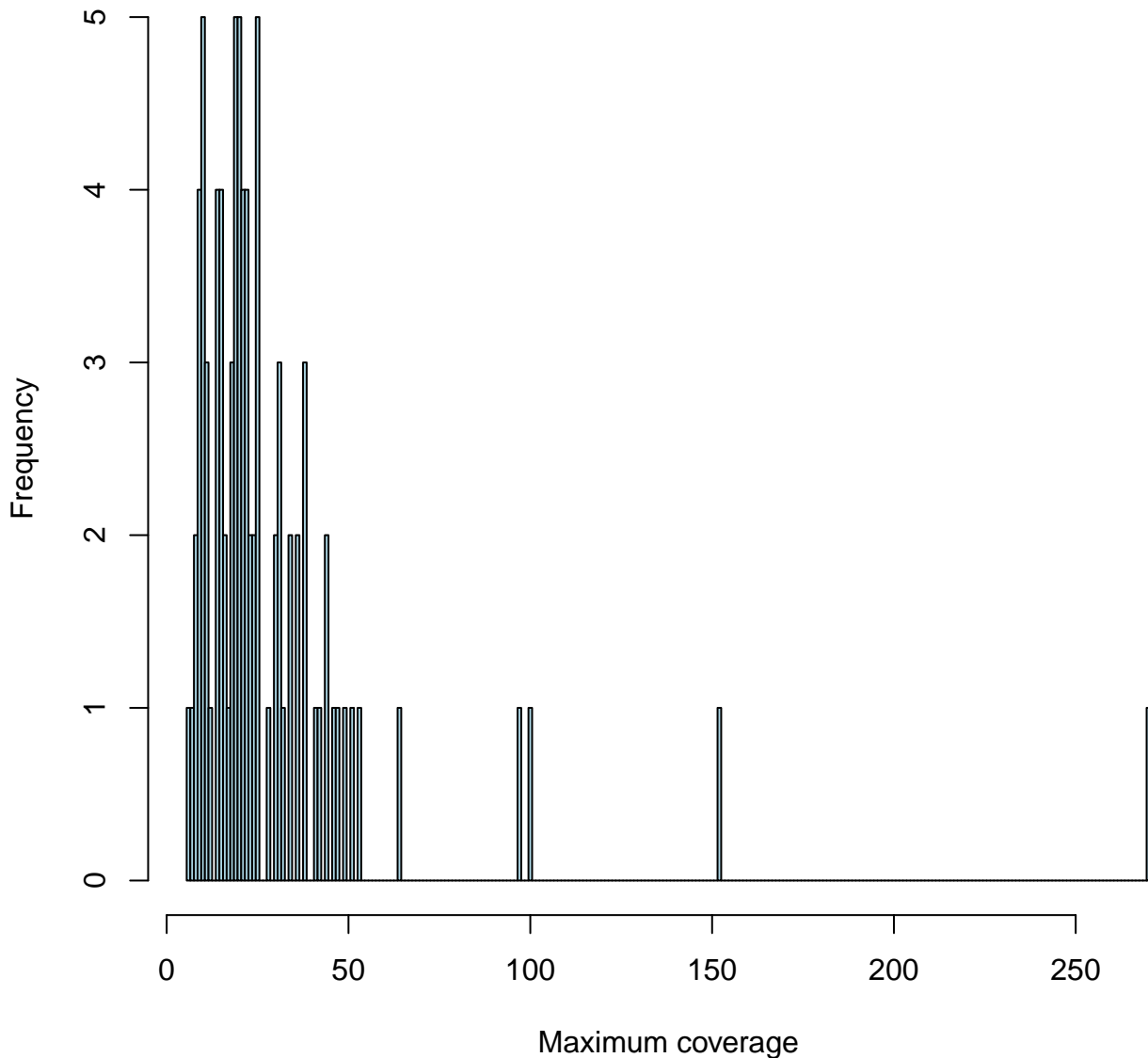
**jx in Intropolis but not "annotated" by knownGene:
163 samples; n jx: 70**



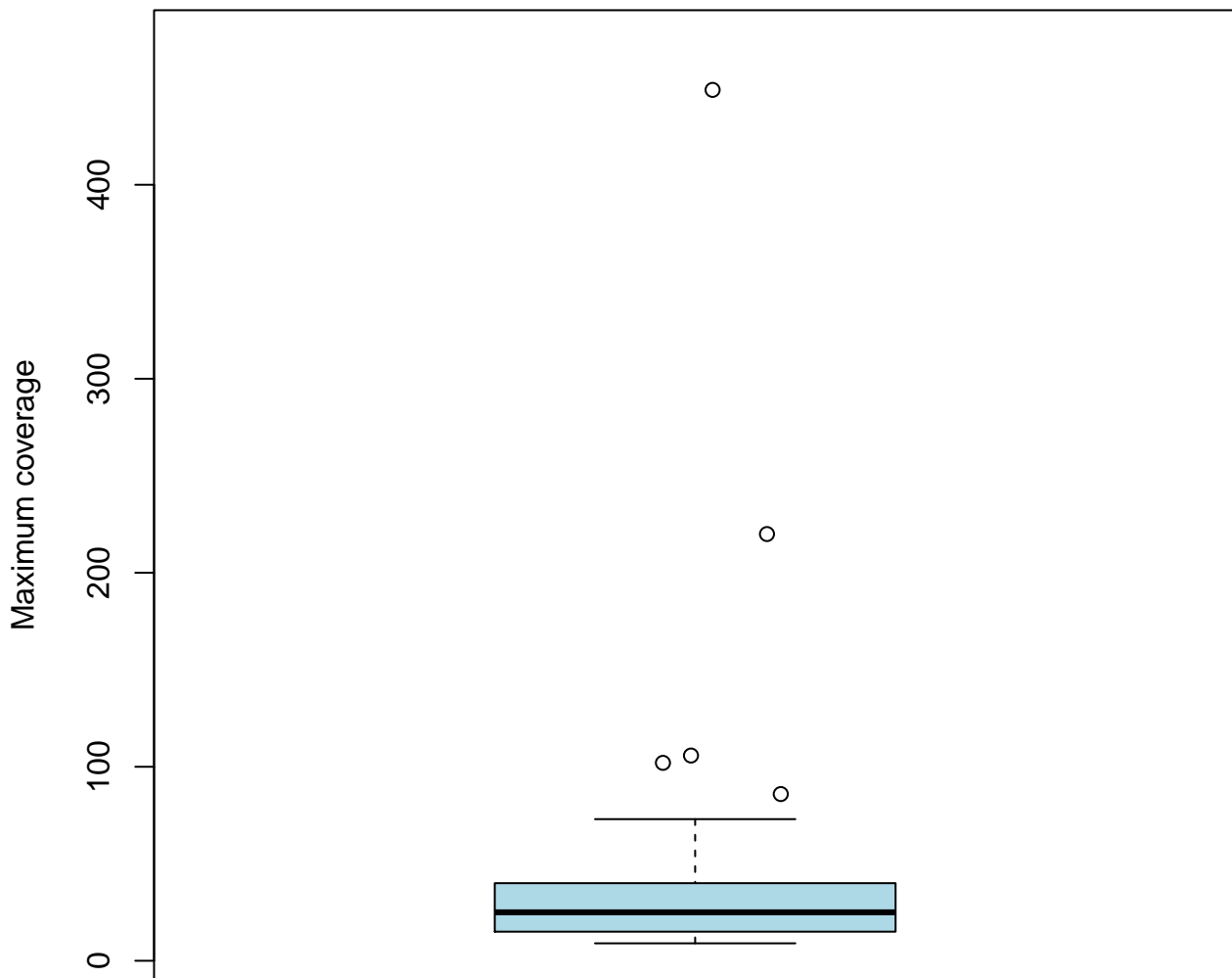
**jx in Intropolis but not "annotated" by knownGene:
164 samples; n jx: 86**



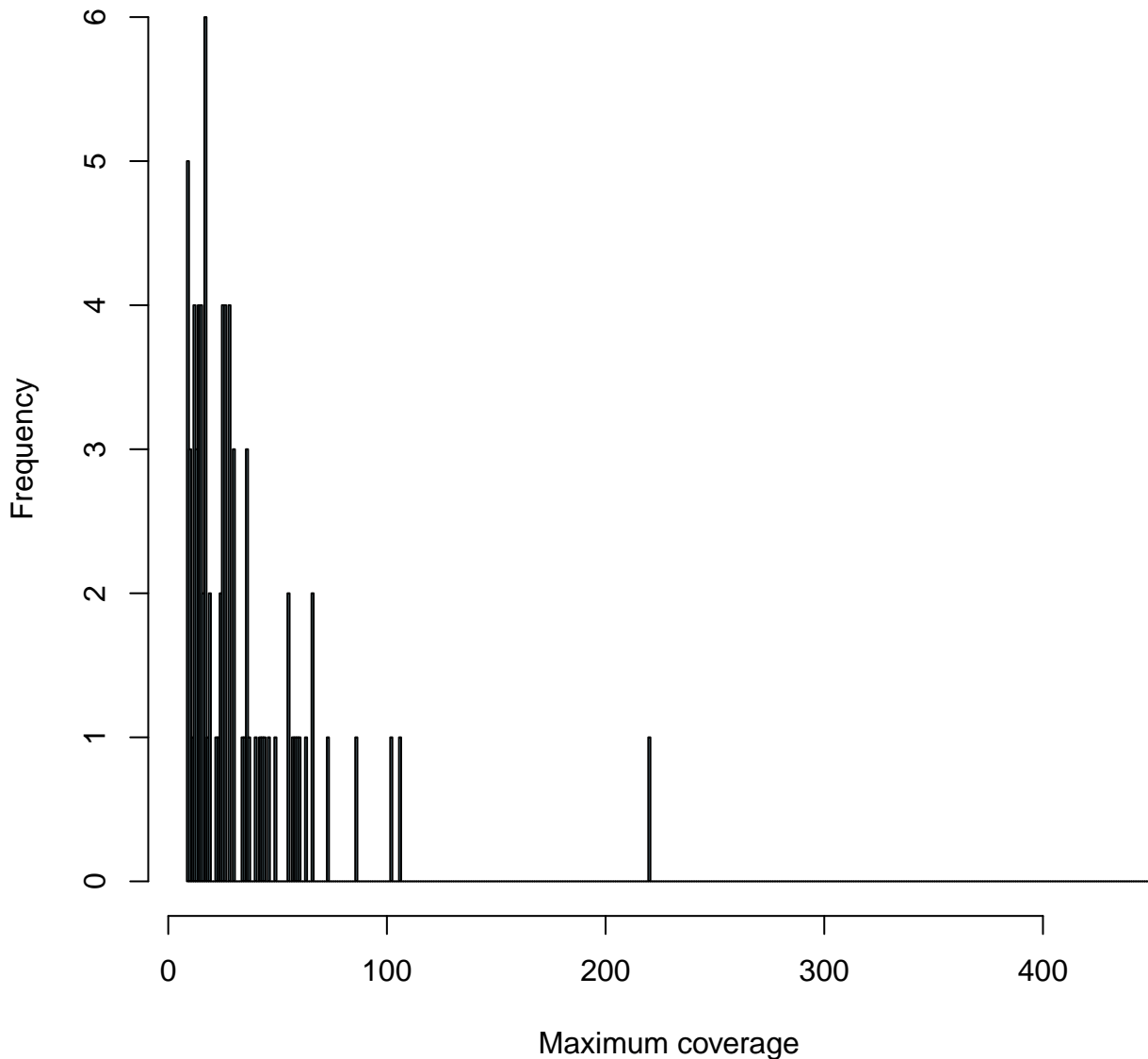
**jx in Intropolis but not "annotated" by knownGene:
164 samples; n jx: 86**



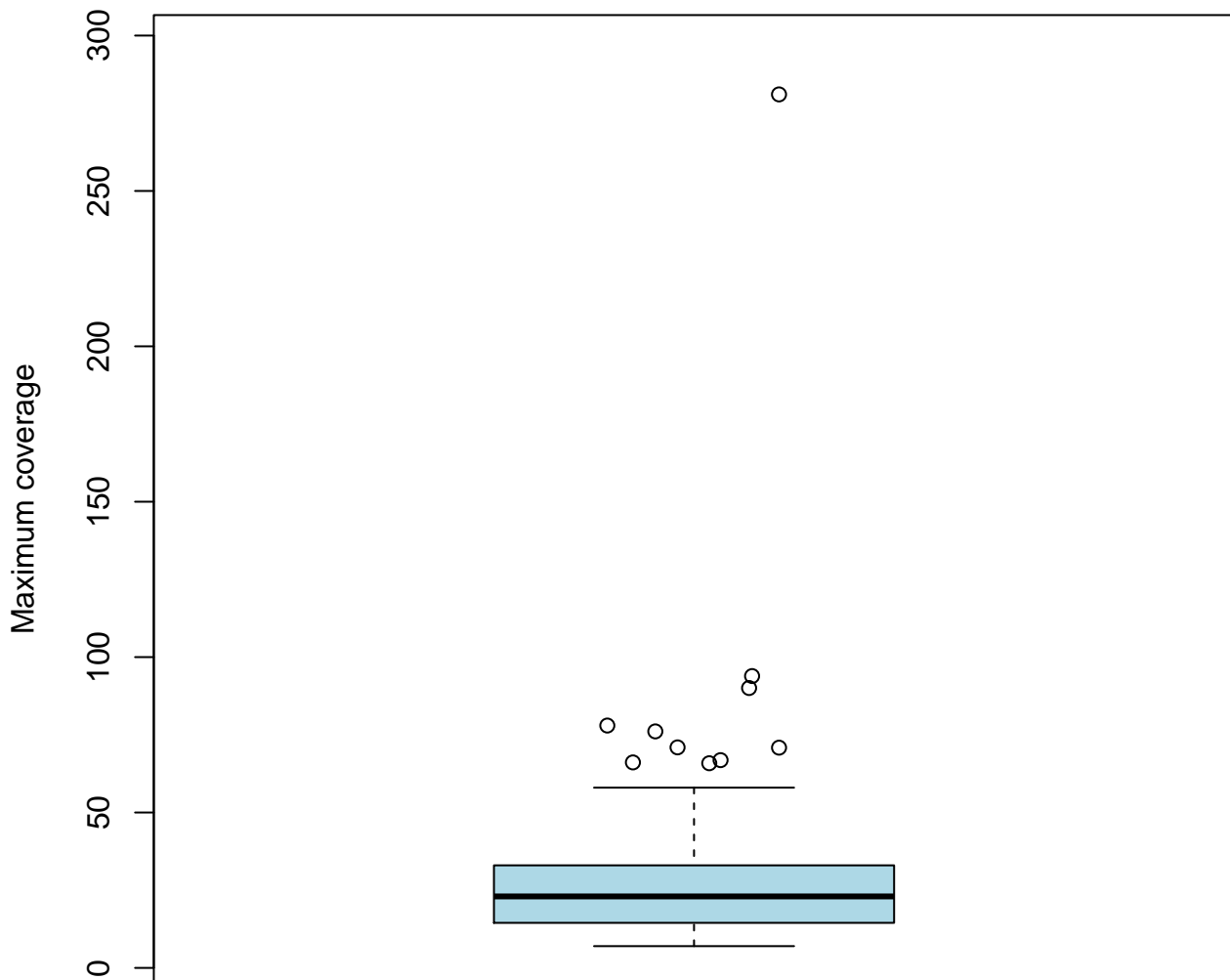
**jx in Intropolis but not "annotated" by knownGene:
165 samples; n jx: 81**



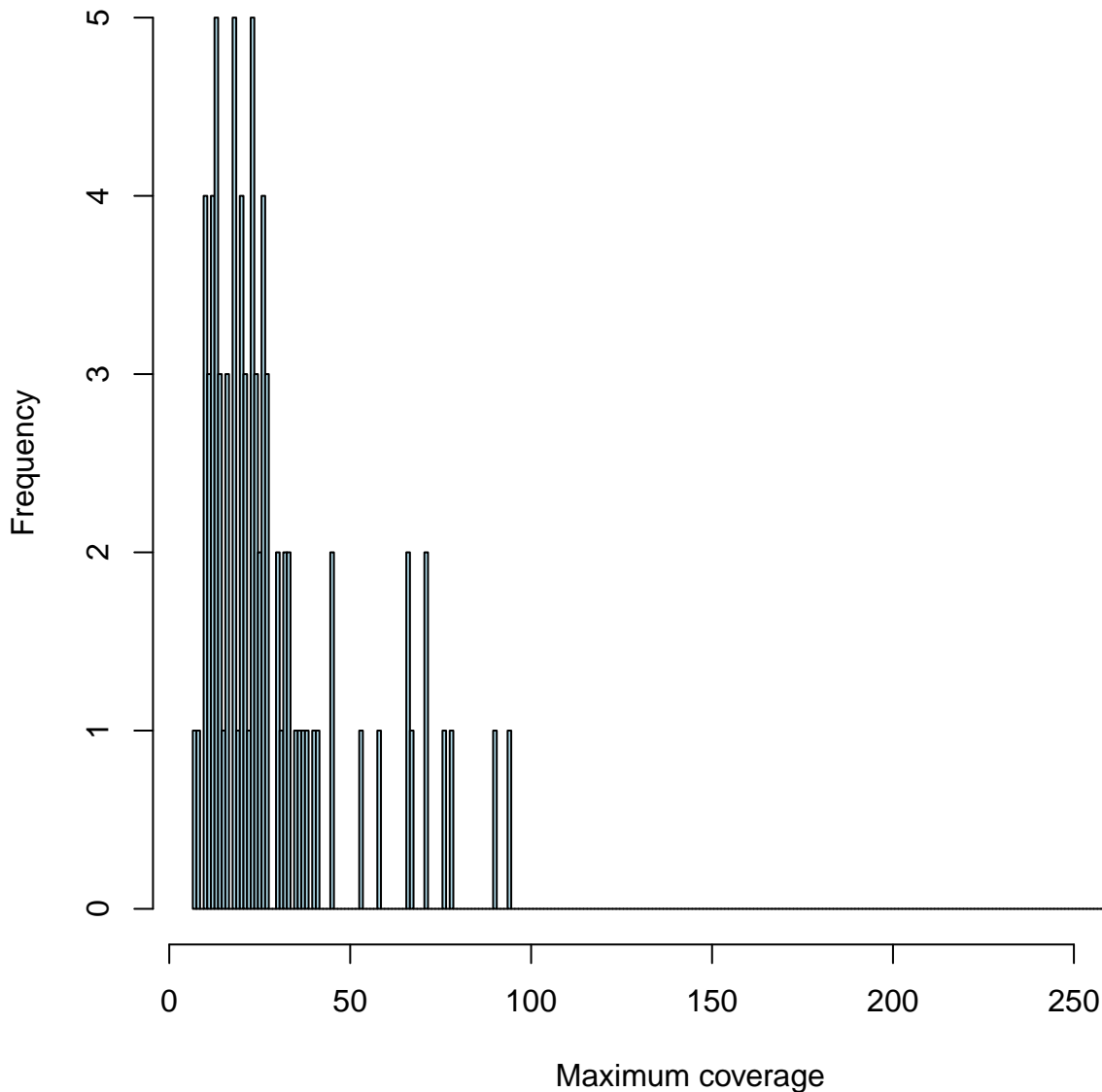
**jx in Intropolis but not "annotated" by knownGene:
165 samples; n jx: 81**



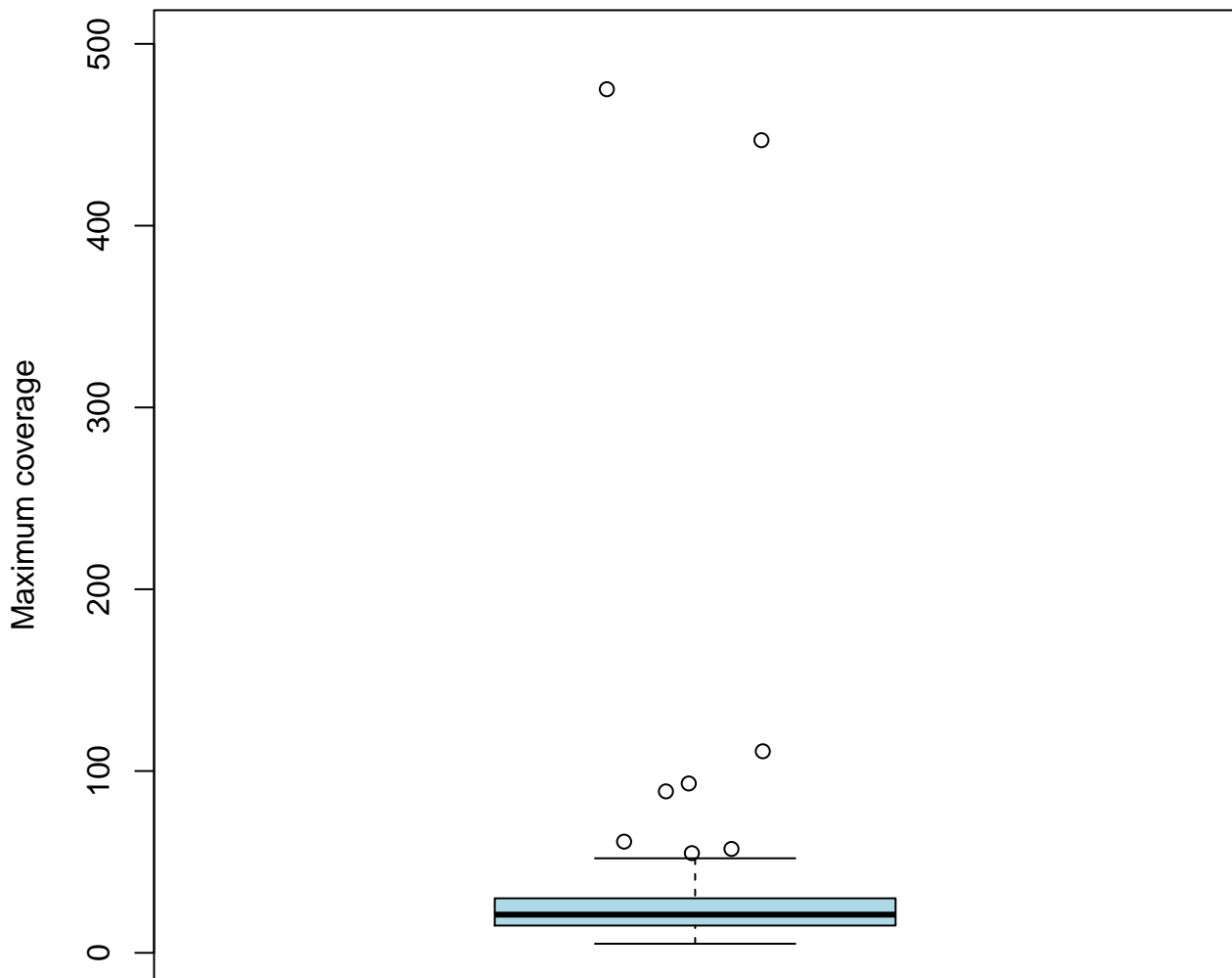
**jx in Intropolis but not "annotated" by knownGene:
166 samples; n jx: 83**



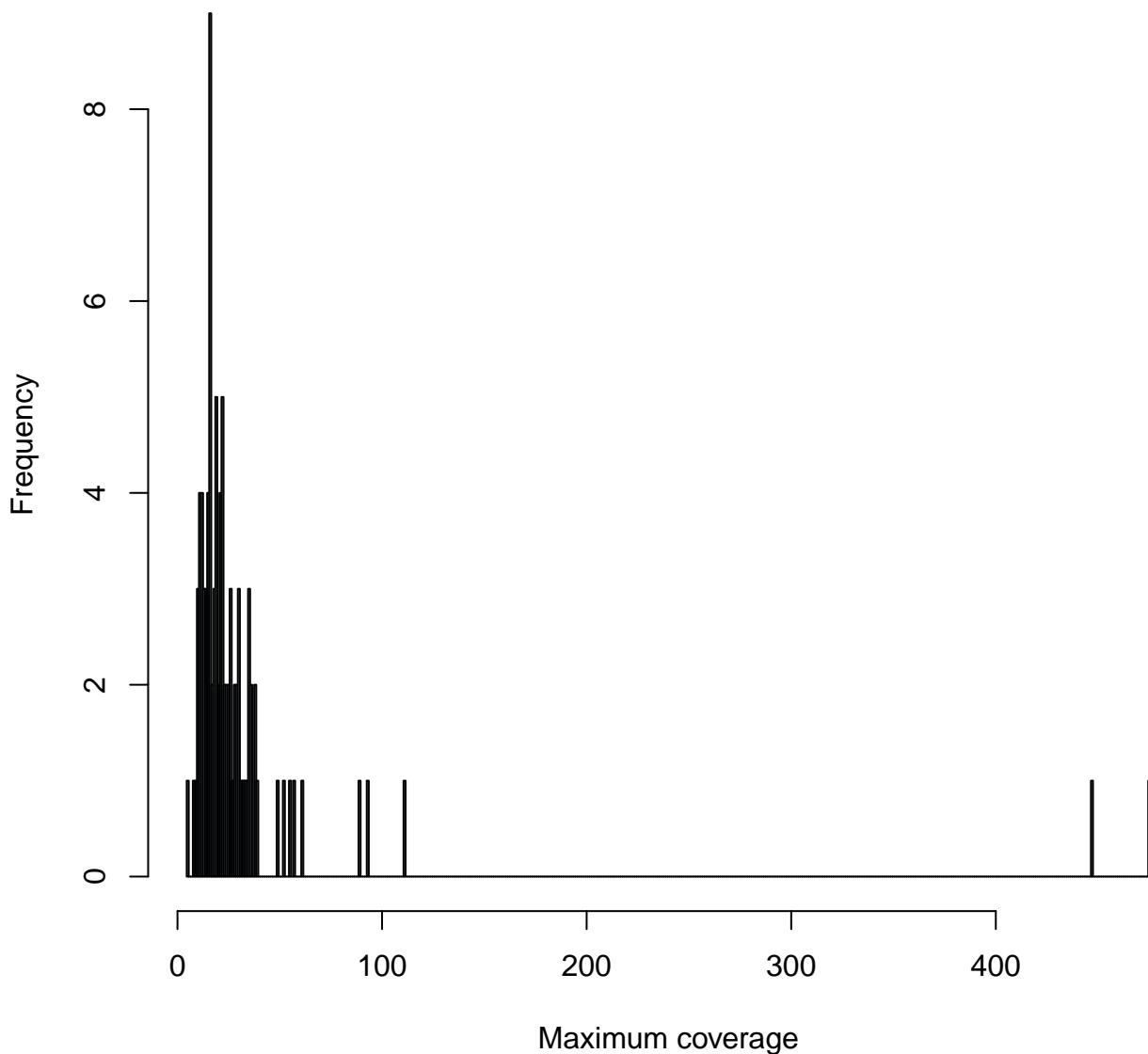
**jx in Intropolis but not "annotated" by knownGene:
166 samples; n jx: 83**



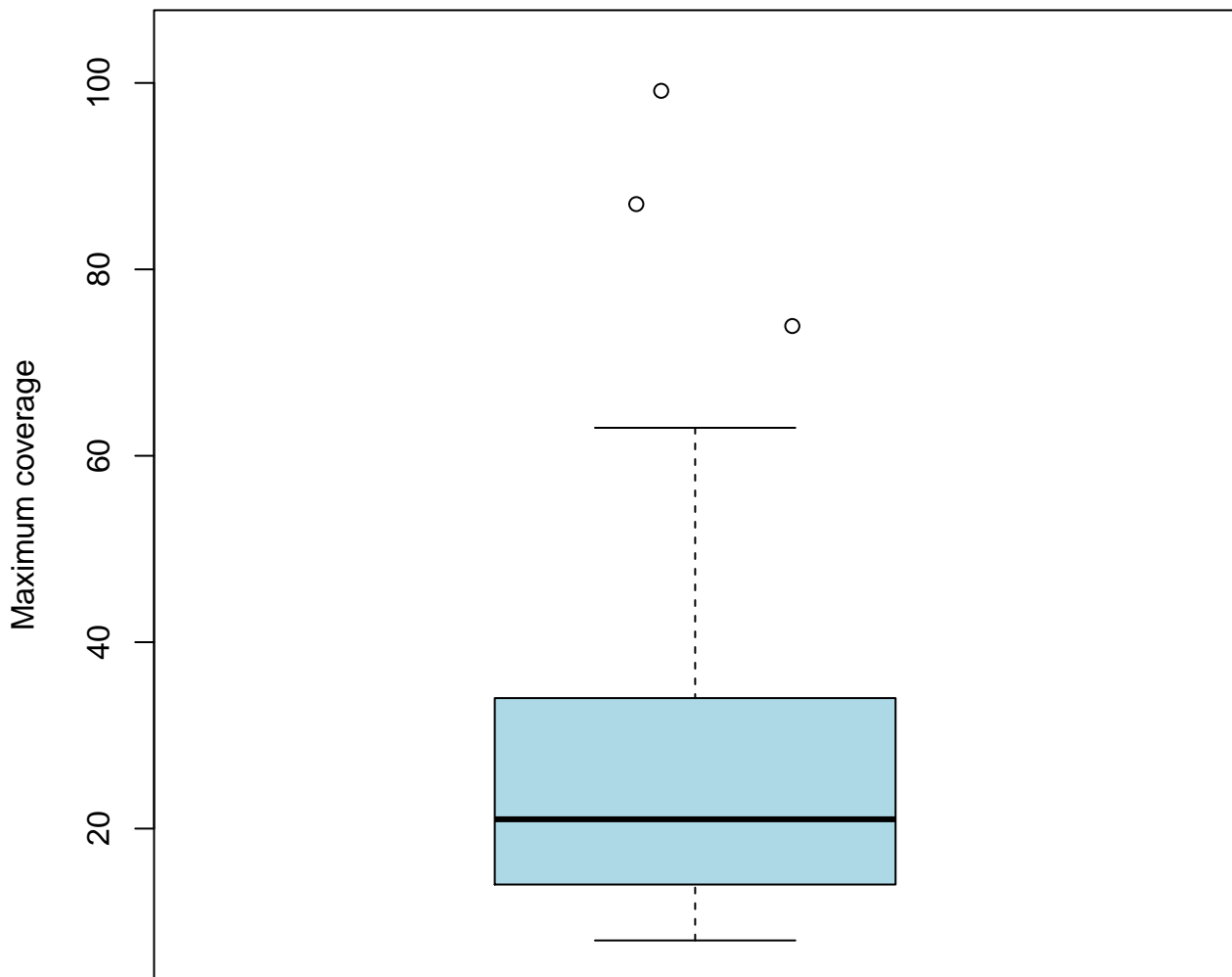
**jx in Intropolis but not "annotated" by knownGene:
167 samples; n jx: 93**



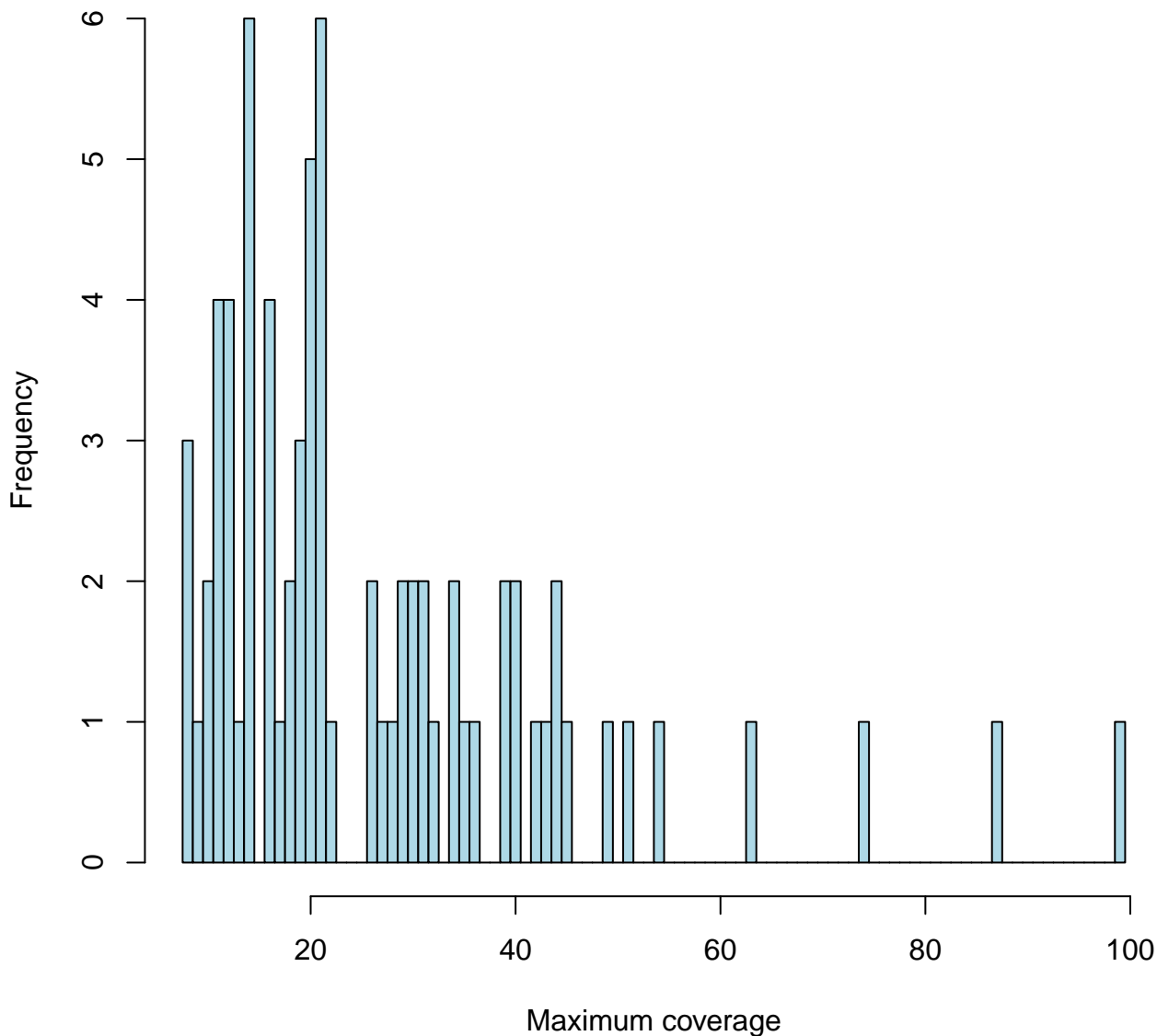
**jx in Intropolis but not "annotated" by knownGene:
167 samples; n jx: 93**



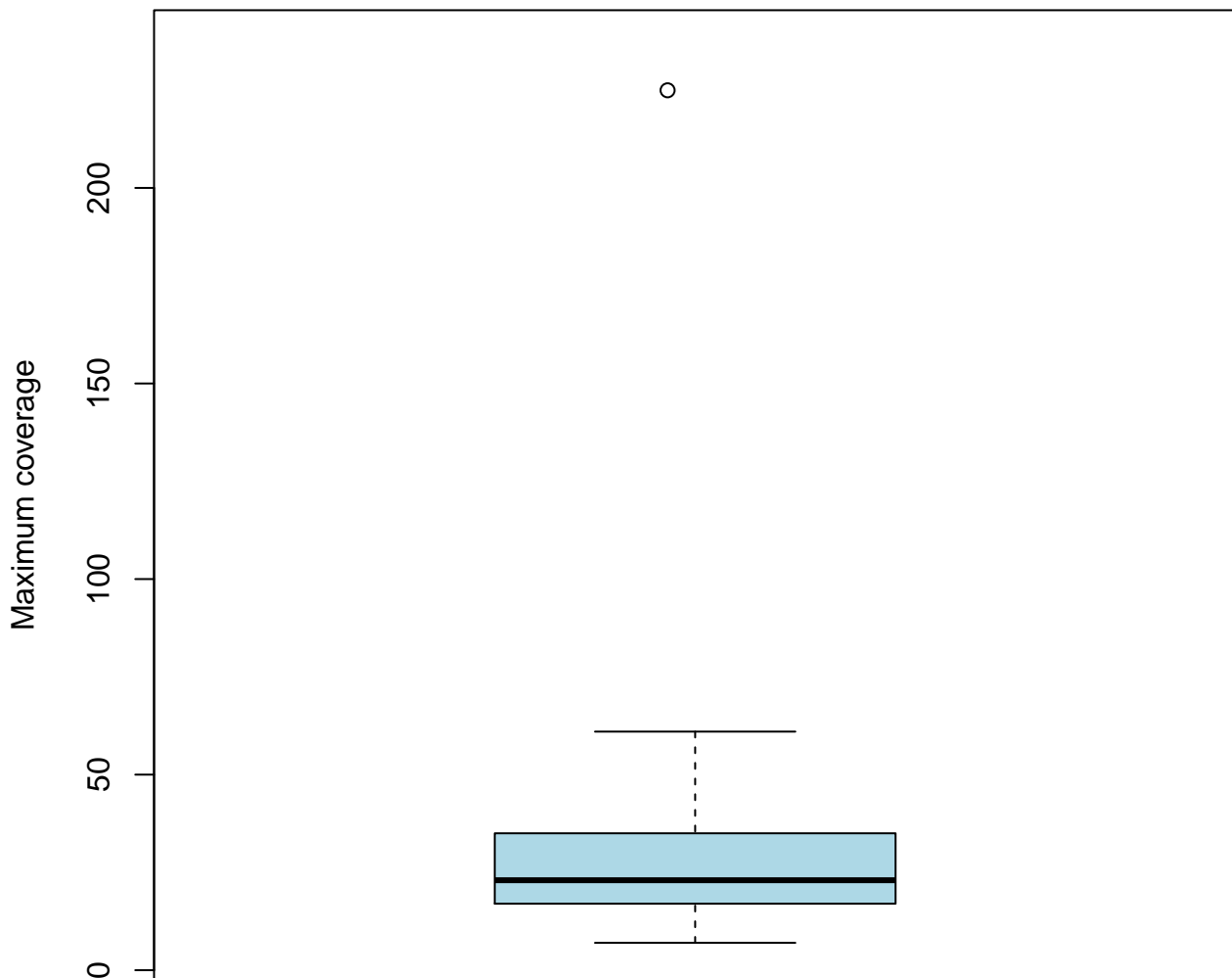
**jx in Intropolis but not "annotated" by knownGene:
168 samples; n jx: 74**



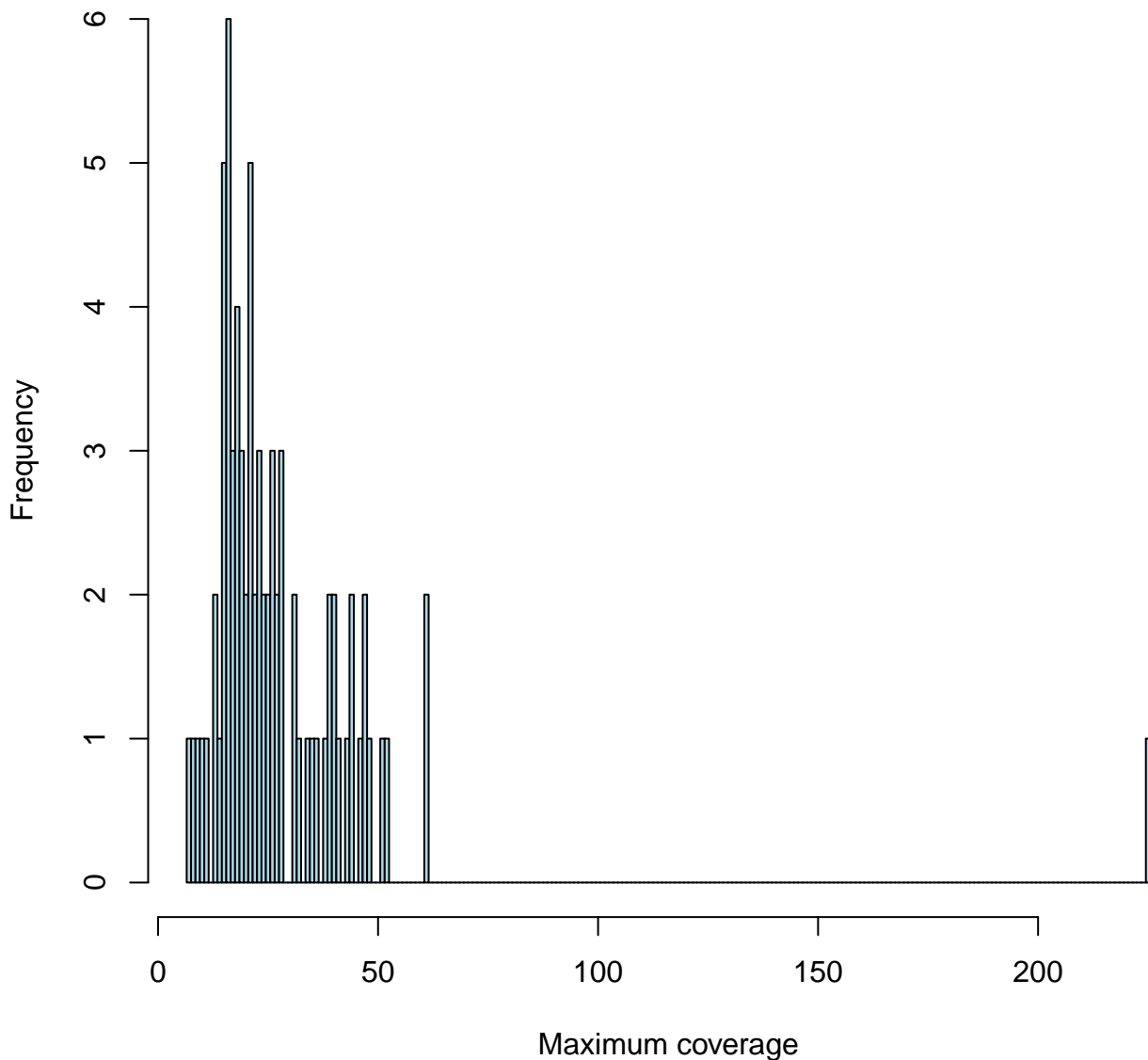
**jx in Intropolis but not "annotated" by knownGene:
168 samples; n jx: 74**



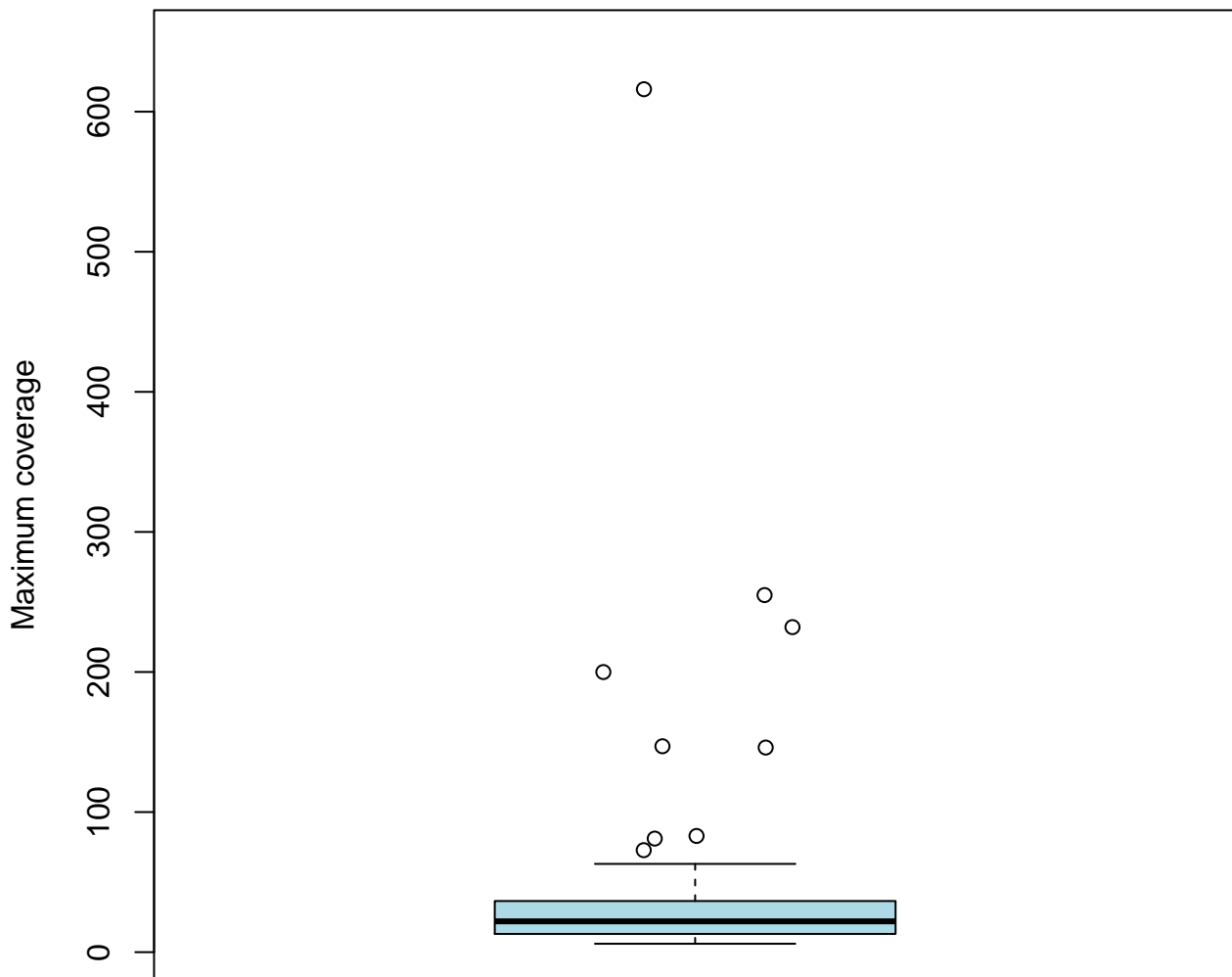
**jx in Intropolis but not "annotated" by knownGene:
169 samples; n jx: 77**



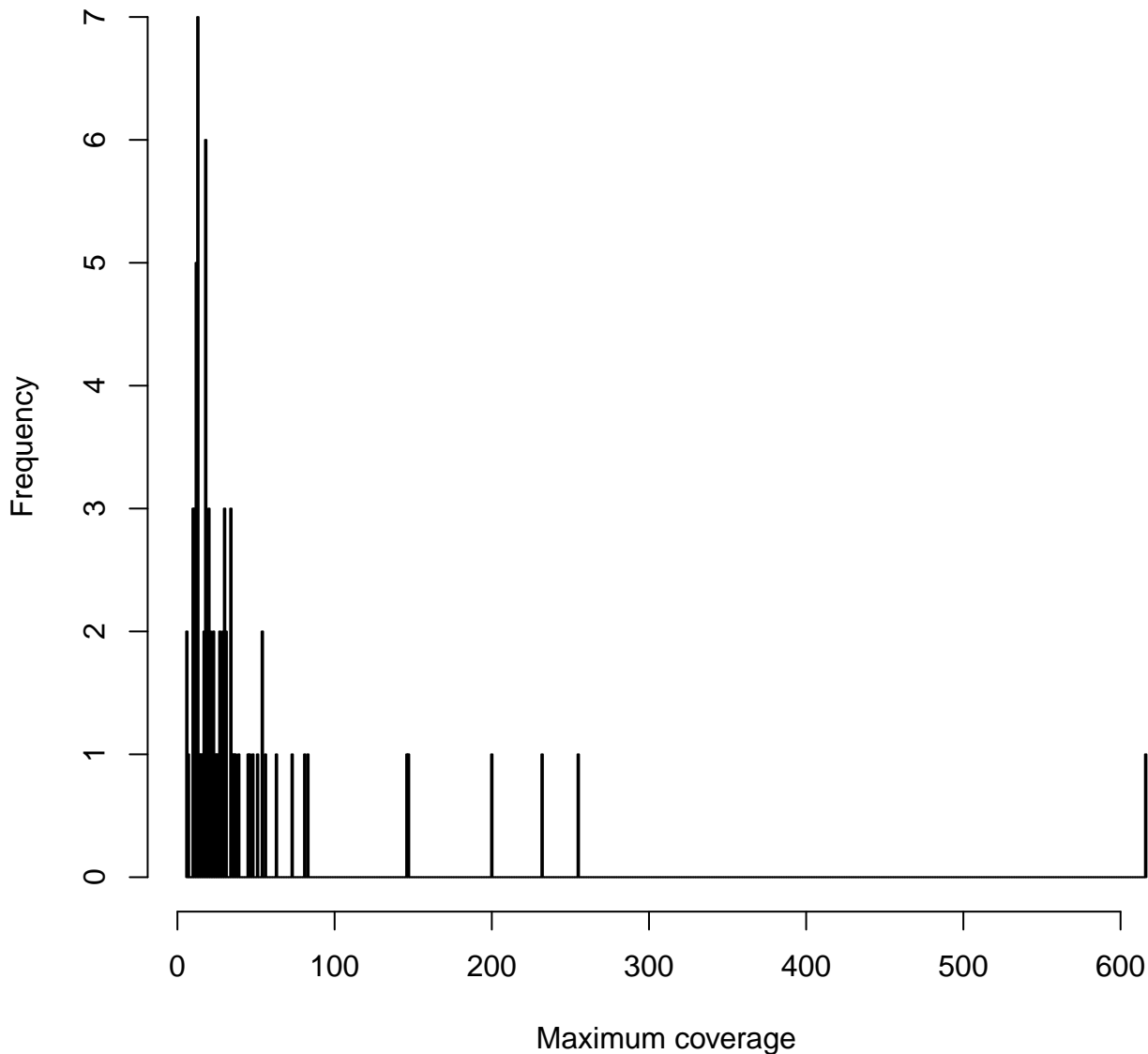
**jx in Intropolis but not "annotated" by knownGene:
169 samples; n jx: 77**



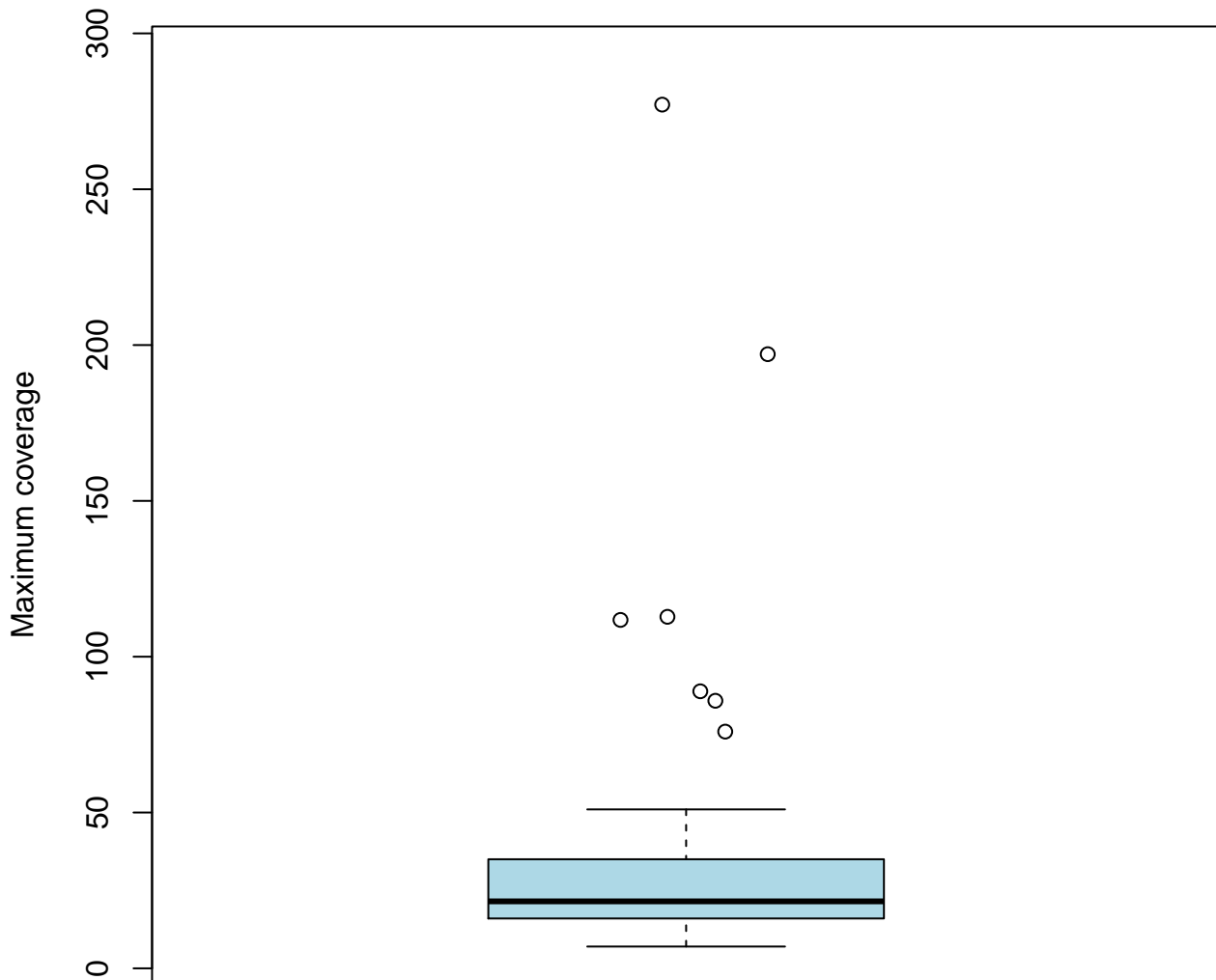
**jx in Intropolis but not "annotated" by knownGene:
170 samples; n jx: 76**



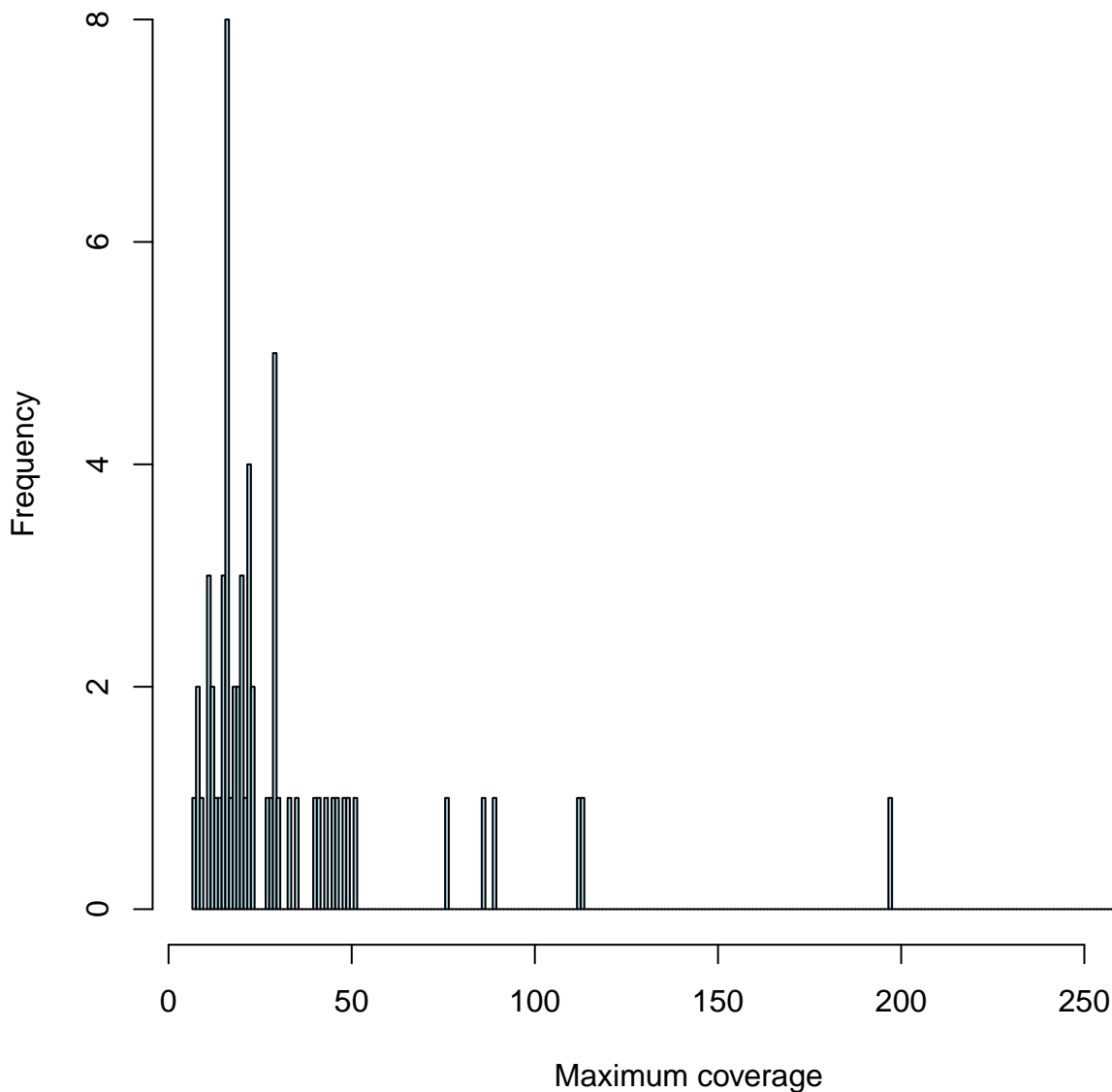
**jx in Intropolis but not "annotated" by knownGene:
170 samples; n jx: 76**



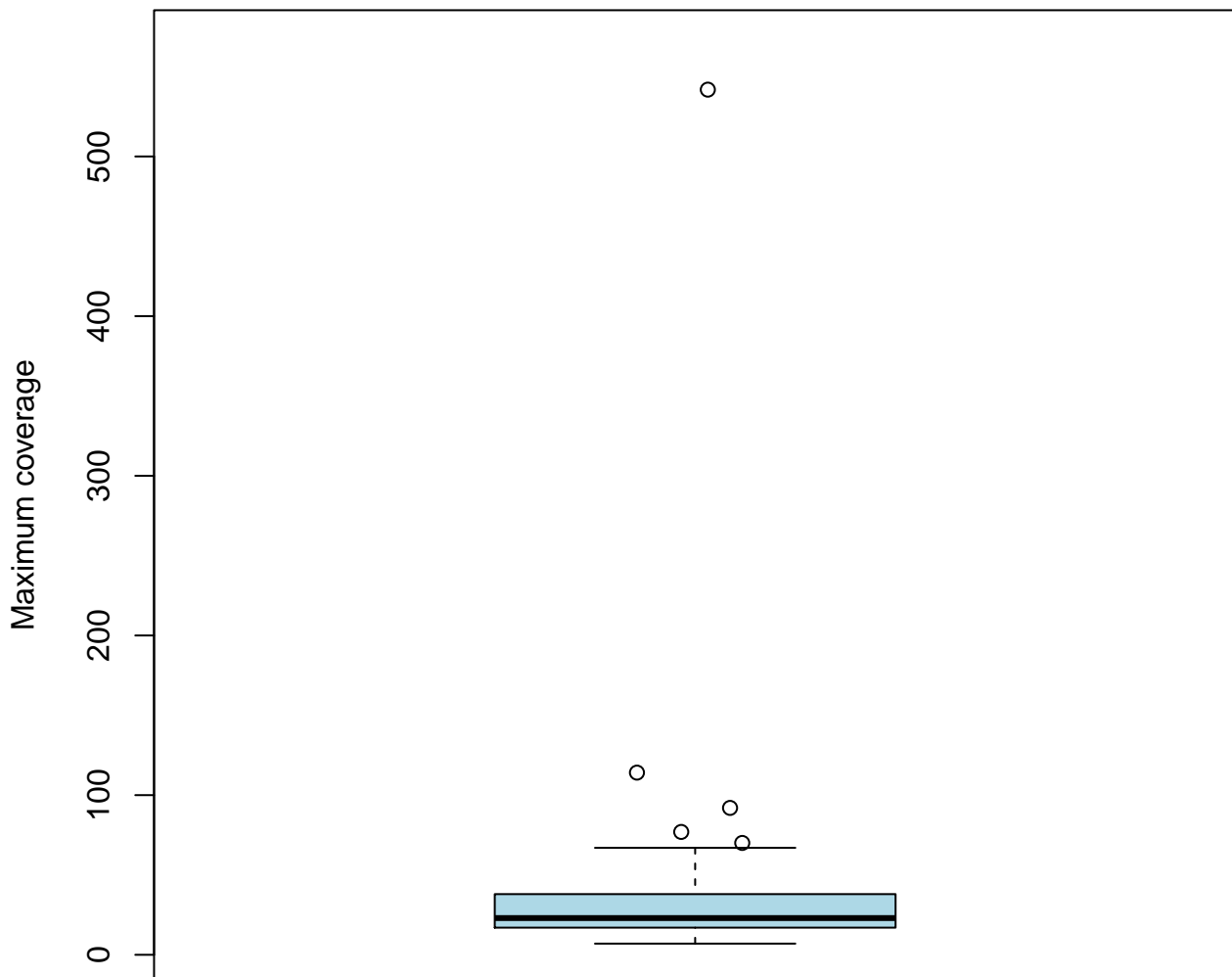
**jx in Intropolis but not "annotated" by knownGene:
171 samples; n jx: 62**



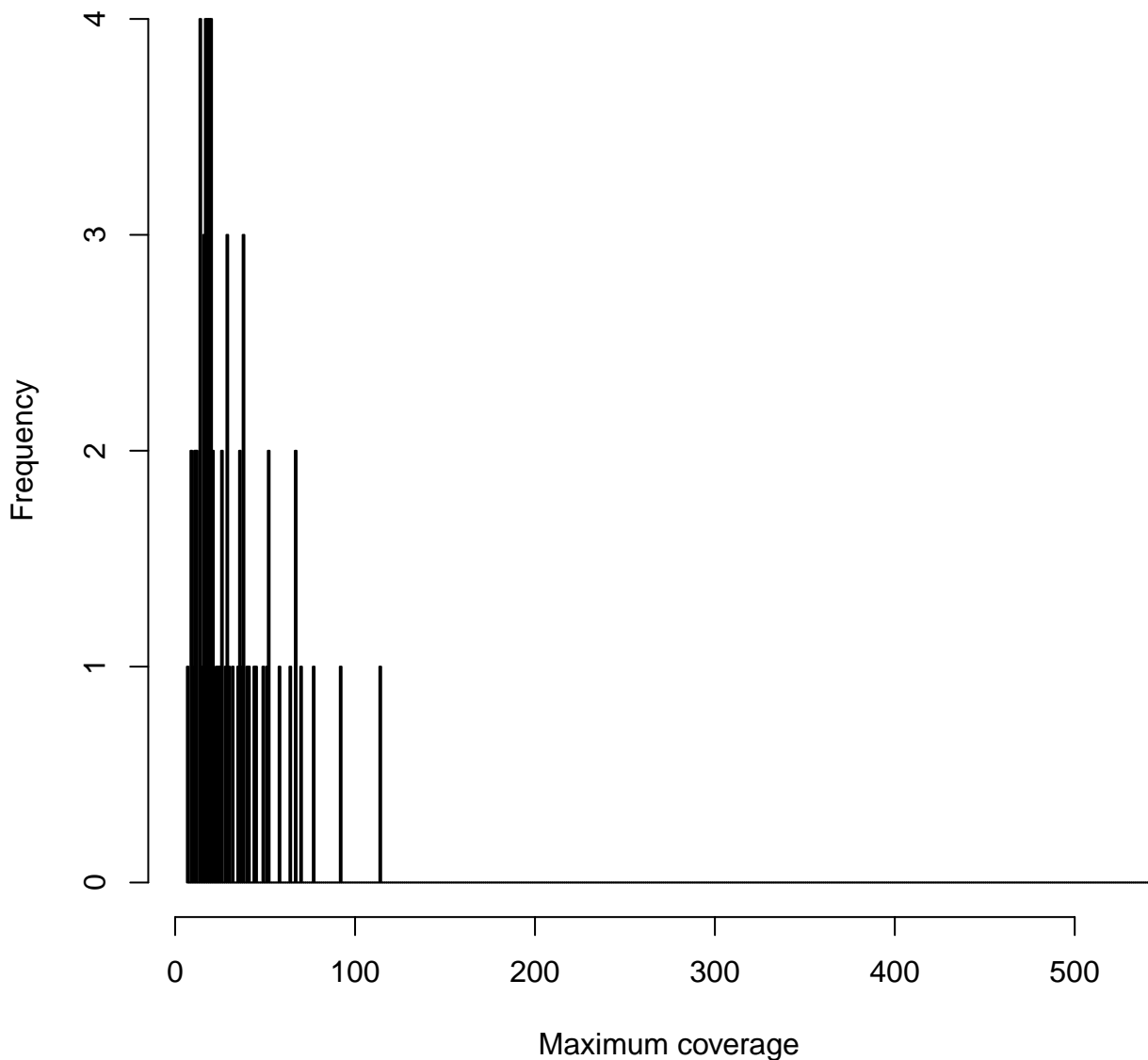
**jx in Intropolis but not "annotated" by knownGene:
171 samples; n jx: 62**



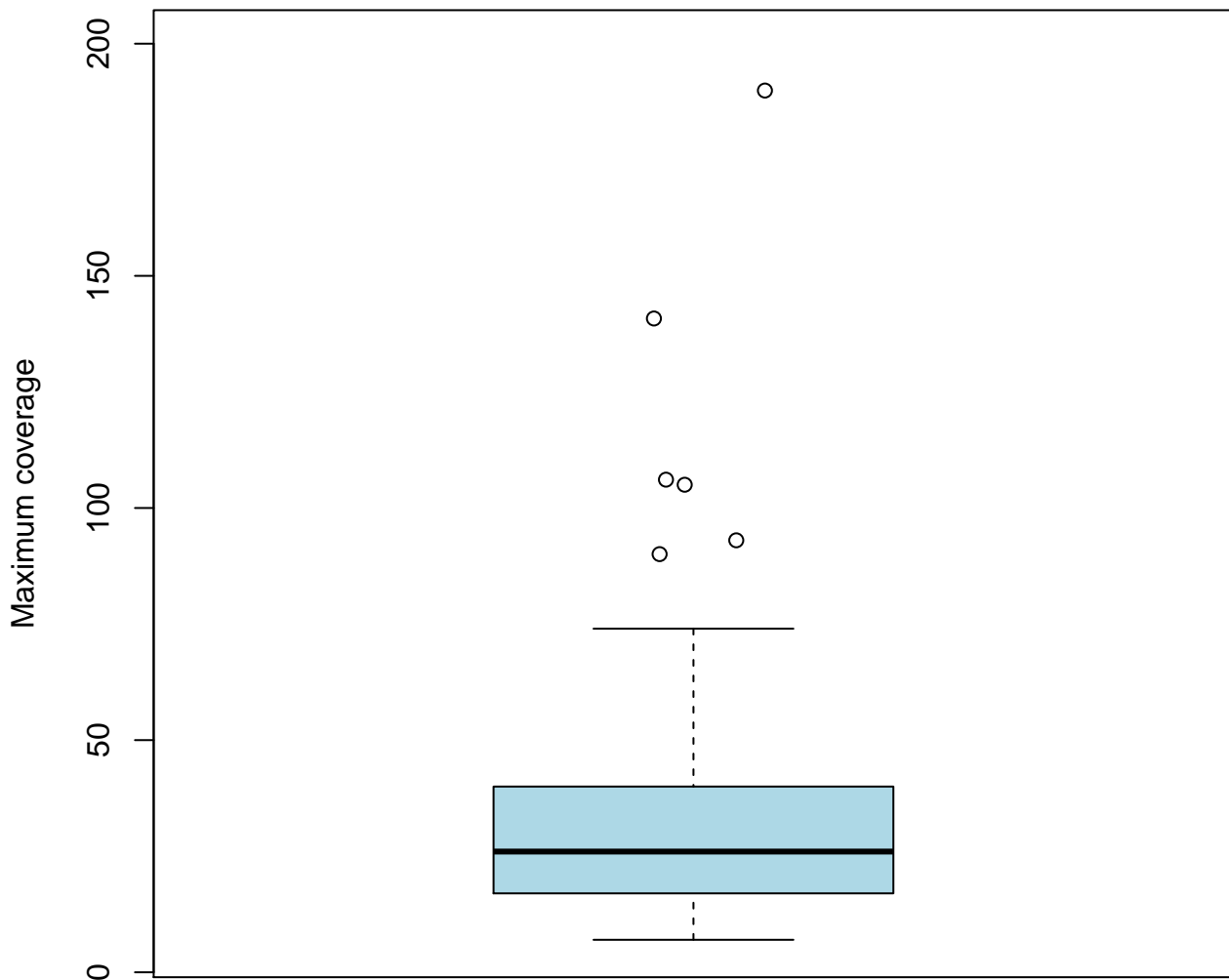
**jx in Intropolis but not "annotated" by knownGene:
172 samples; n jx: 69**



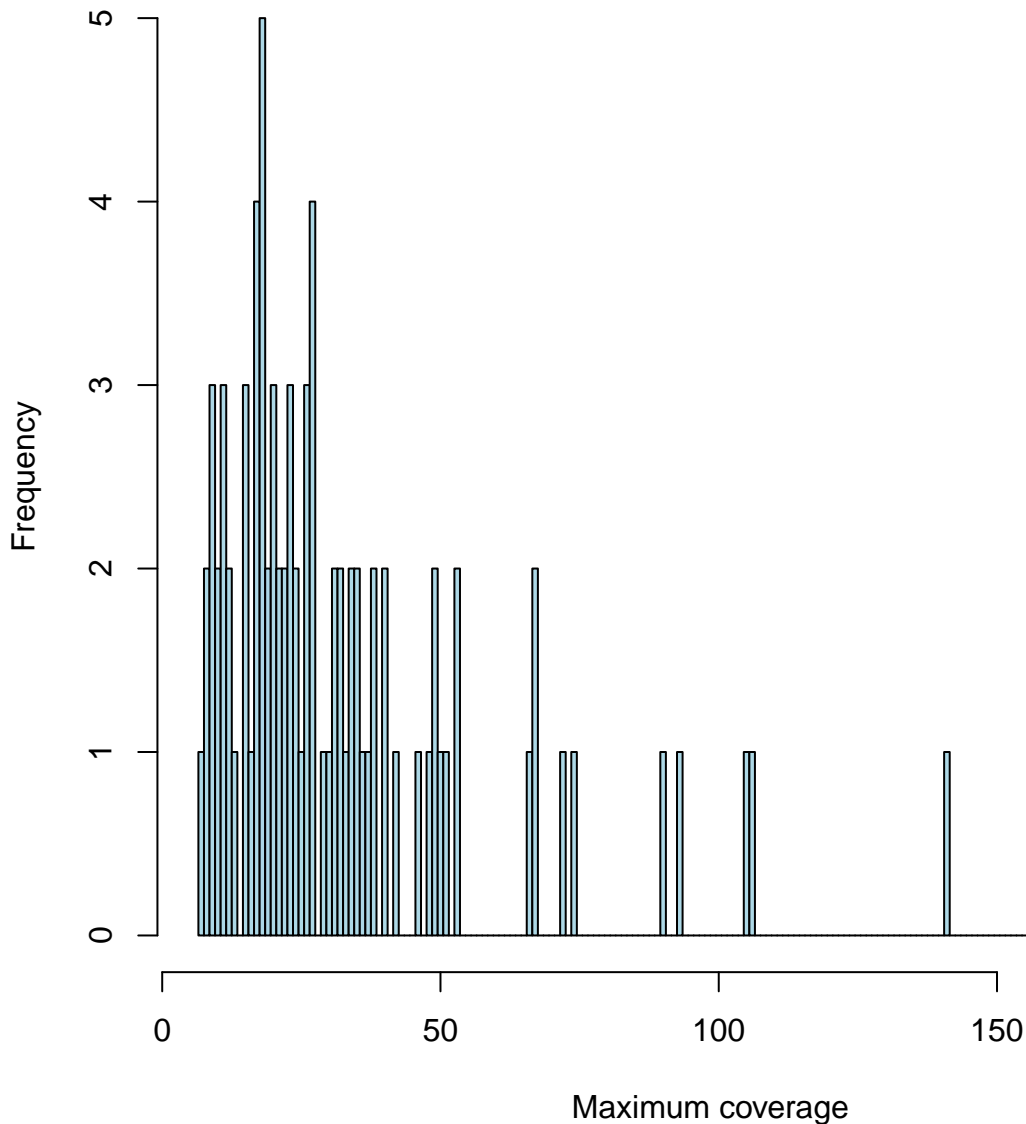
**jx in Intropolis but not "annotated" by knownGene:
172 samples; n jx: 69**



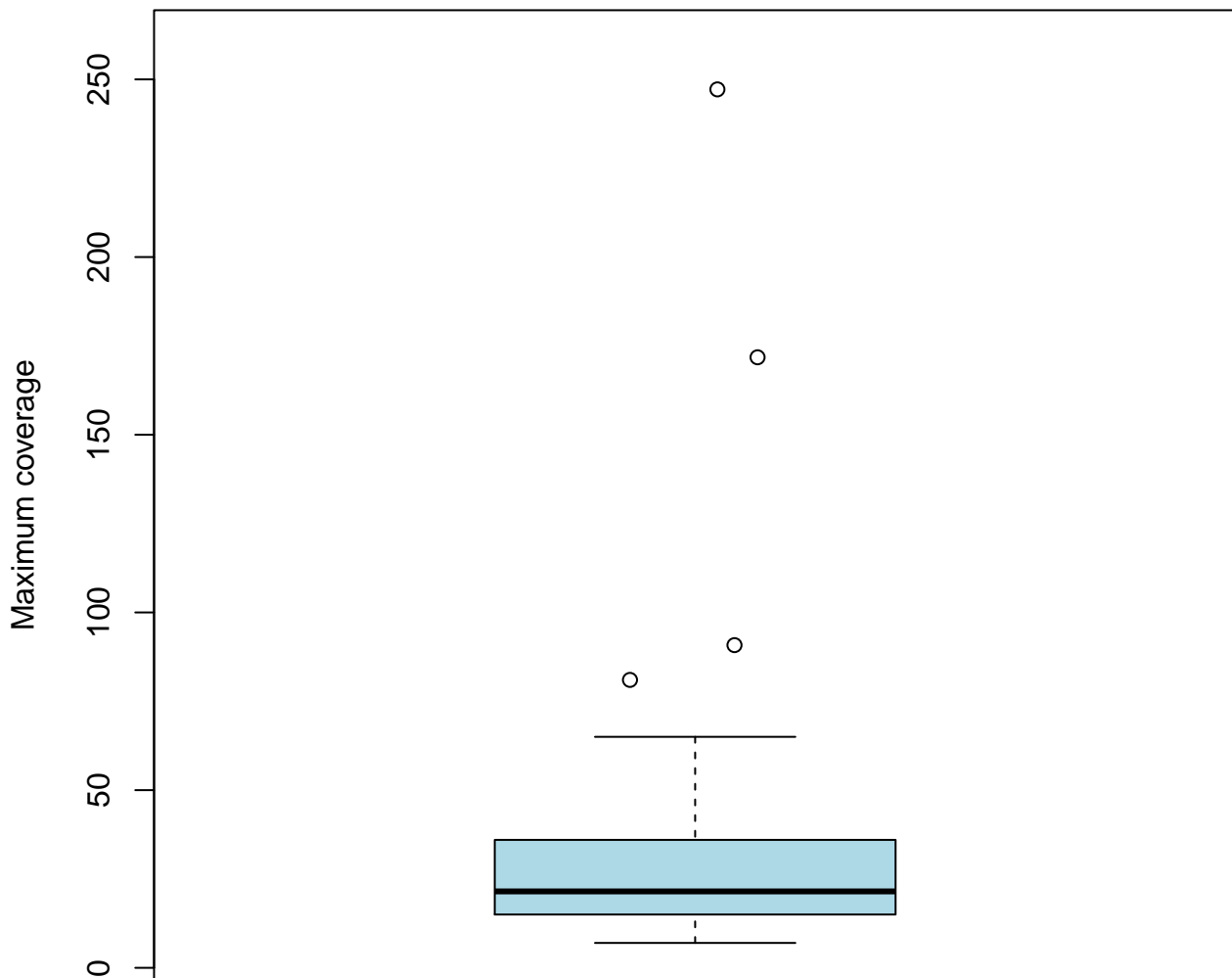
**jx in Intropolis but not "annotated" by knownGene:
173 samples; n jx: 86**



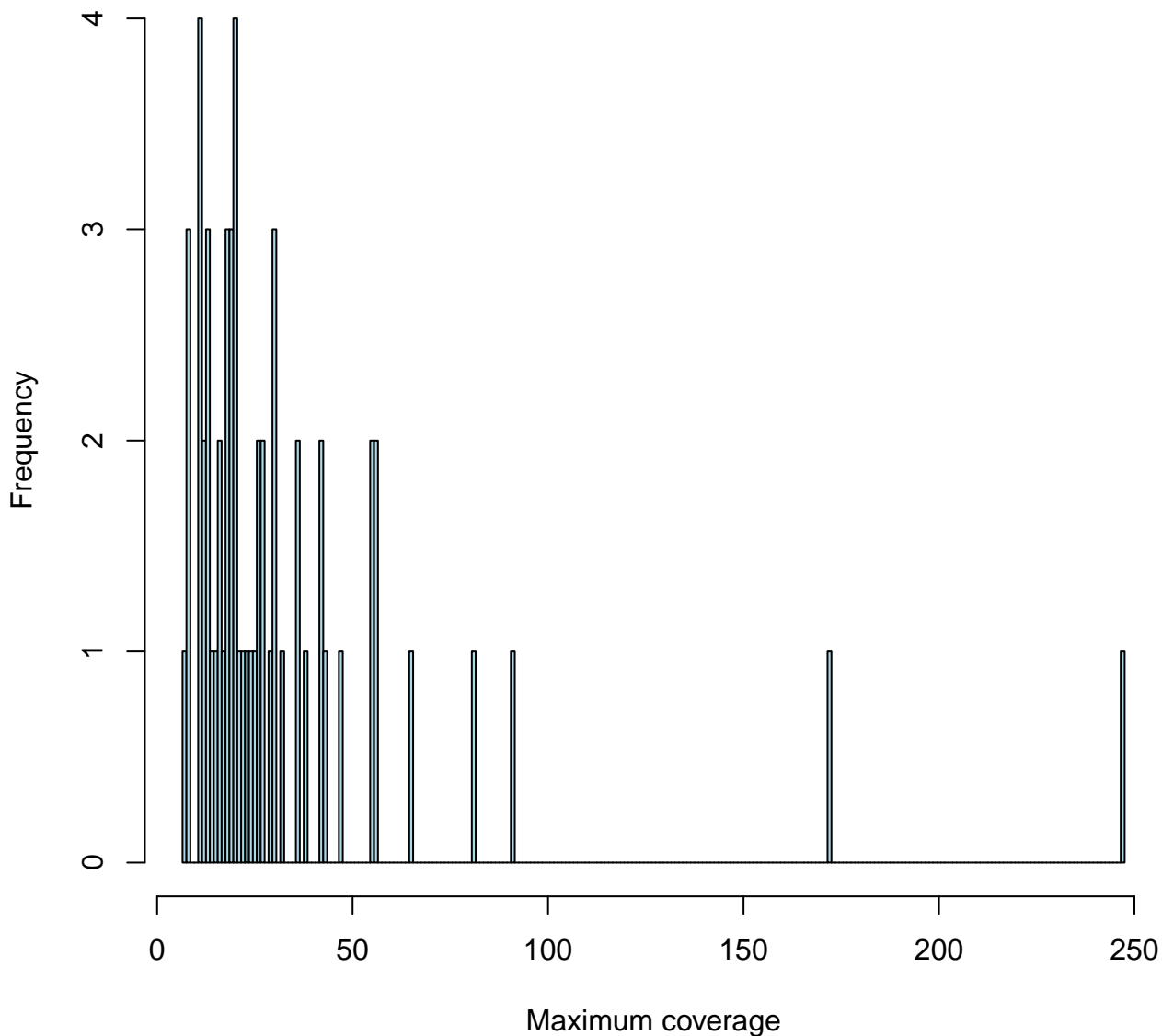
**jx in Intropolis but not "annotated" by knownGene:
173 samples; n jx: 86**



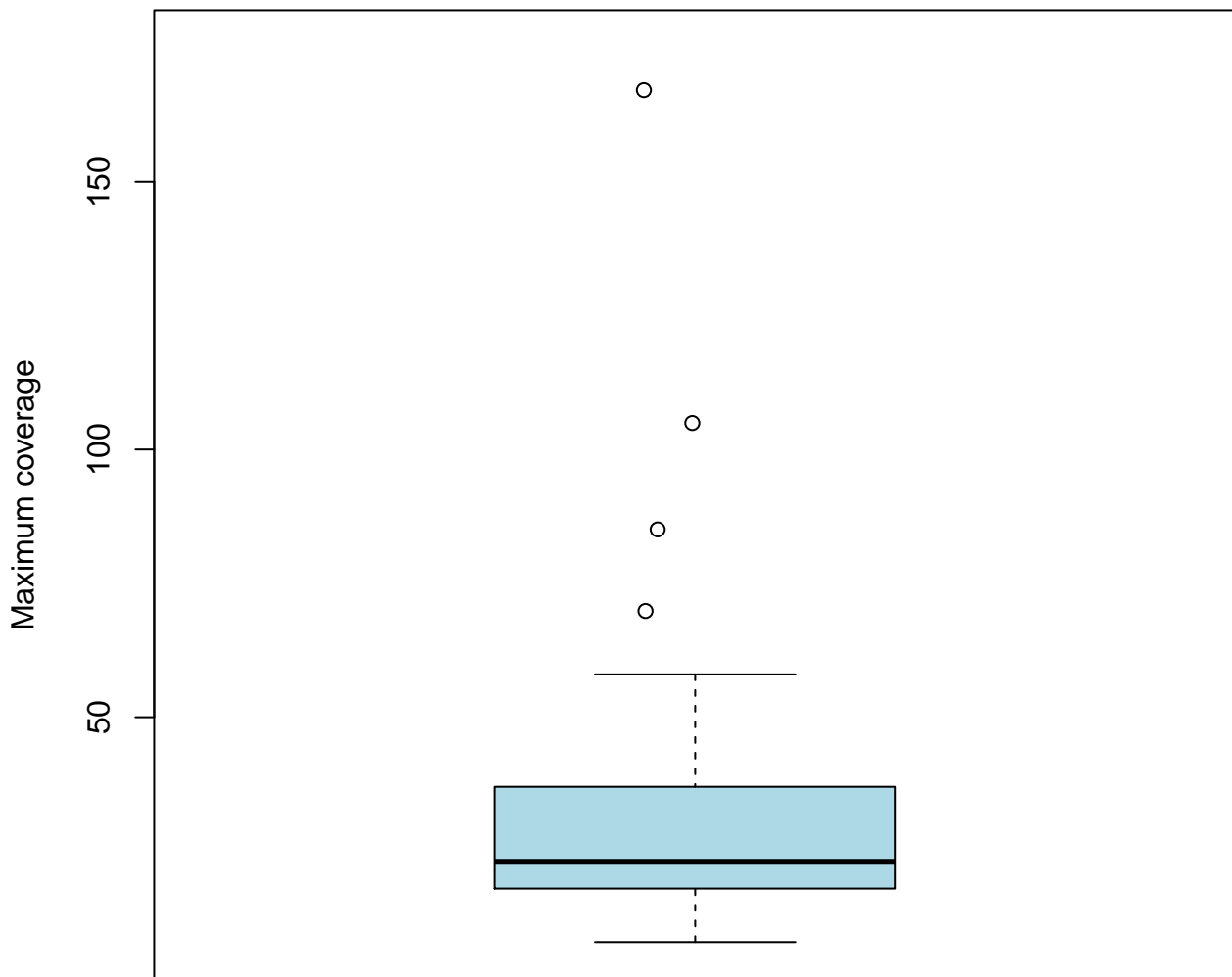
**jx in Intropolis but not "annotated" by knownGene:
174 samples; n jx: 58**



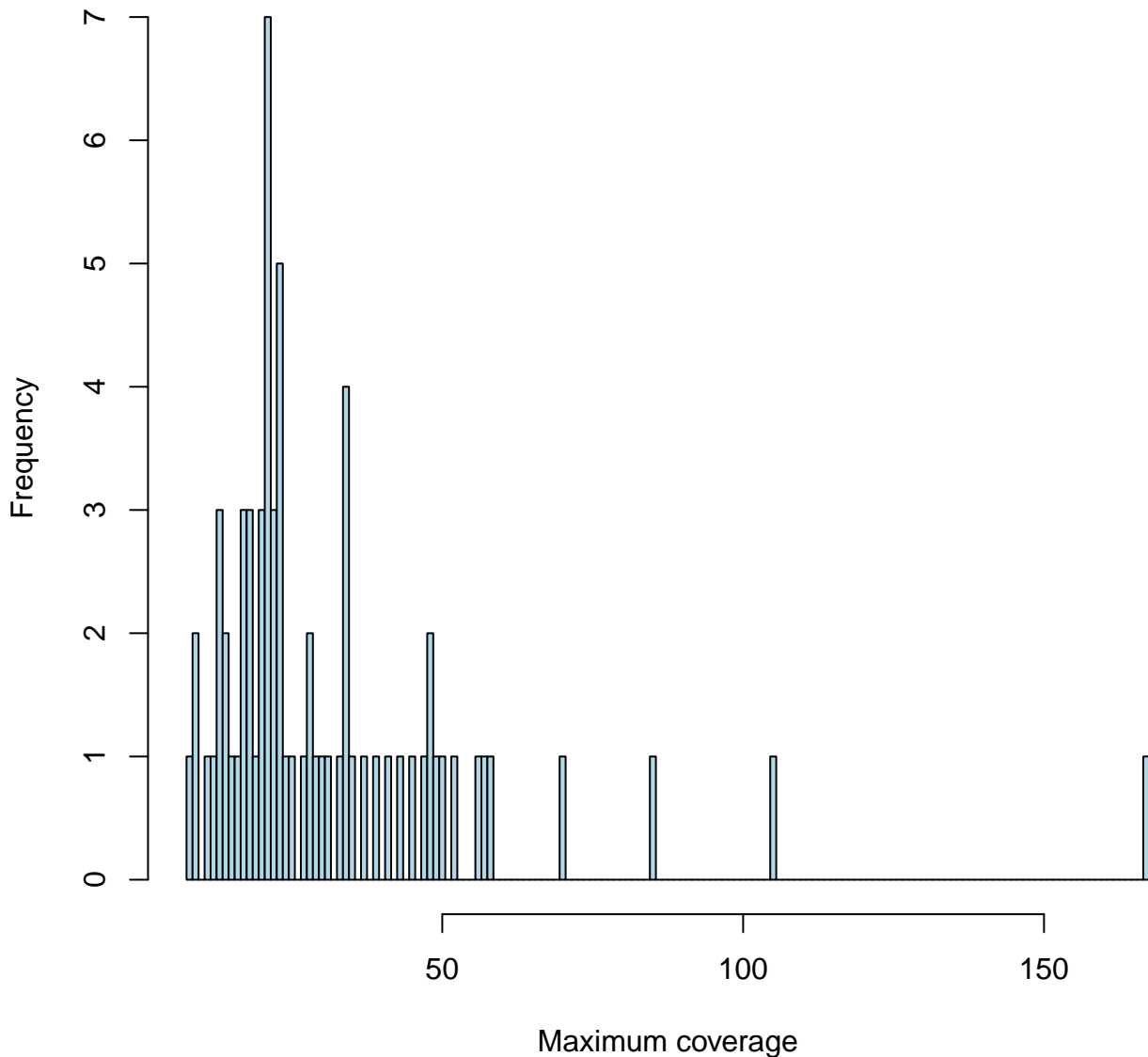
**jx in Intropolis but not "annotated" by knownGene:
174 samples; n jx: 58**



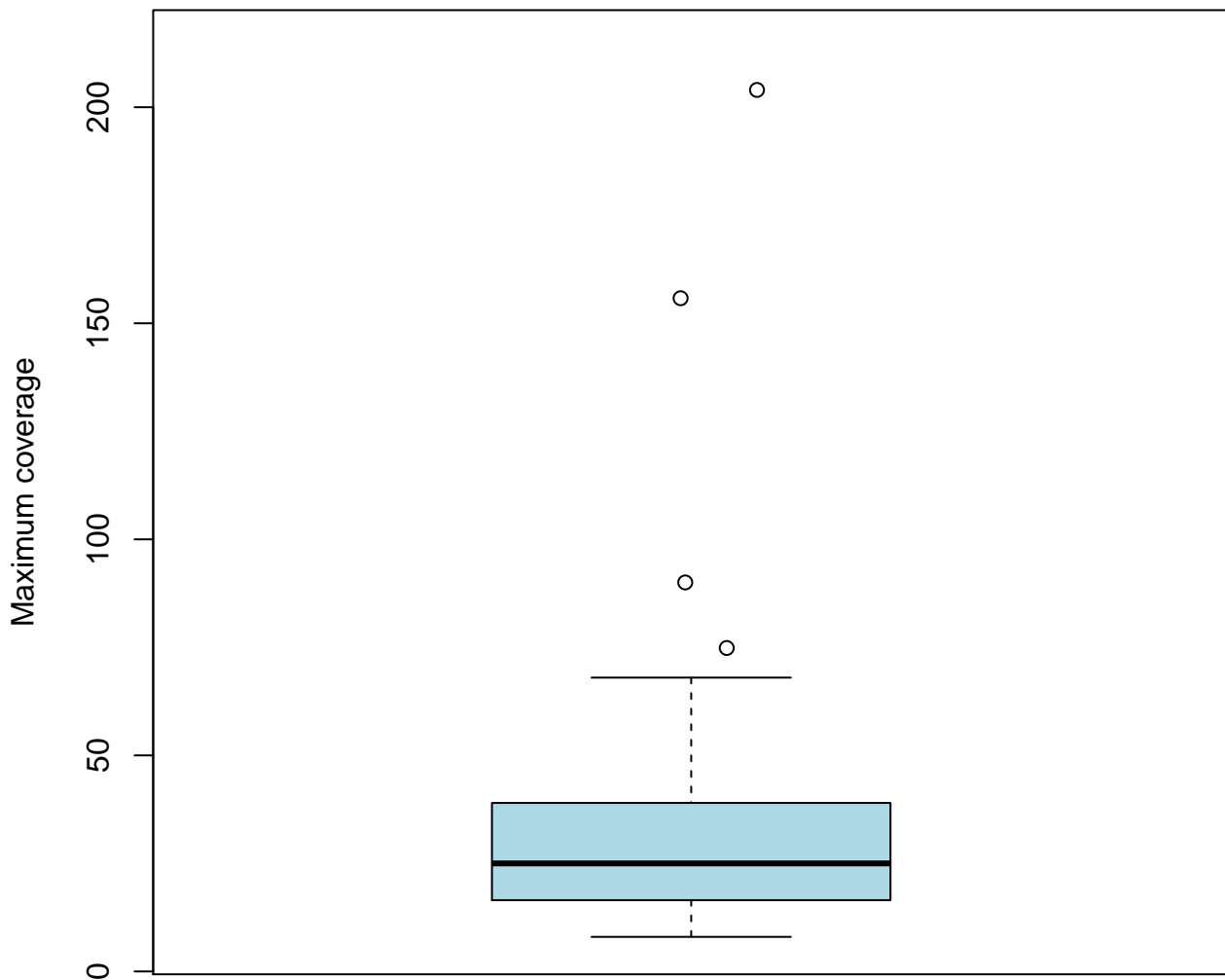
**jx in Intropolis but not "annotated" by knownGene:
175 samples; n jx: 69**



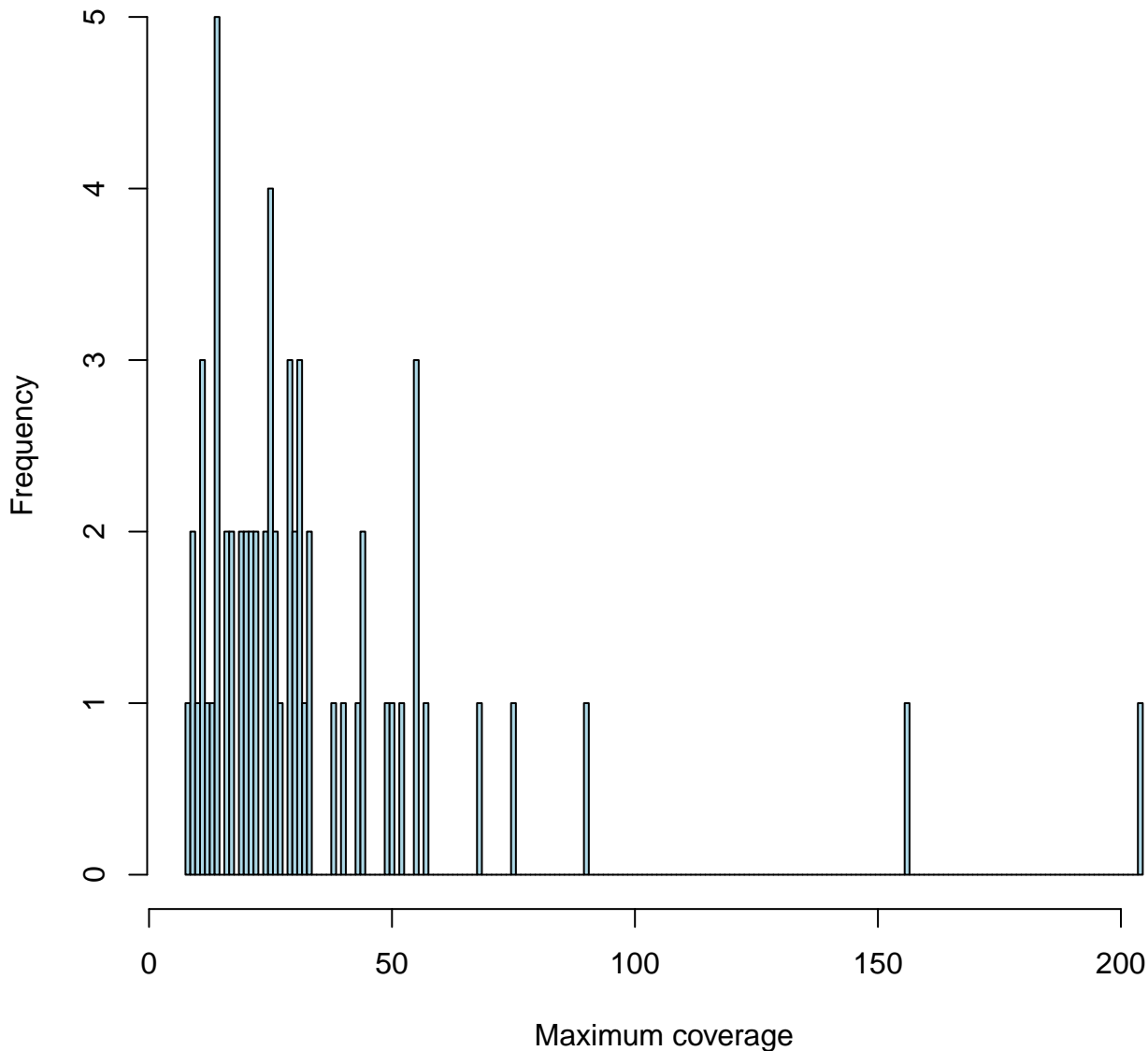
**jx in Intropolis but not "annotated" by knownGene:
175 samples; n jx: 69**



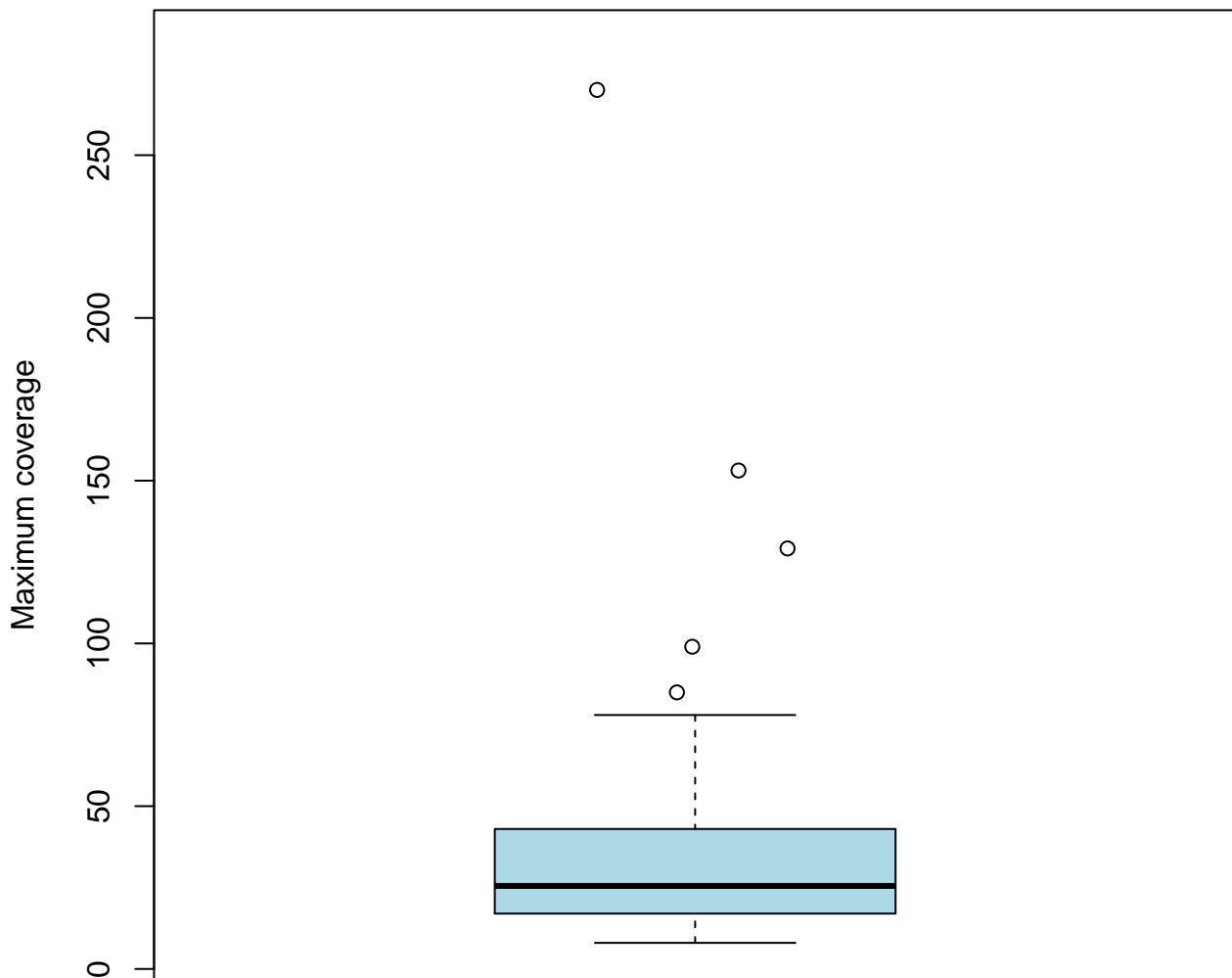
**jx in Intropolis but not "annotated" by knownGene:
176 samples; n jx: 63**



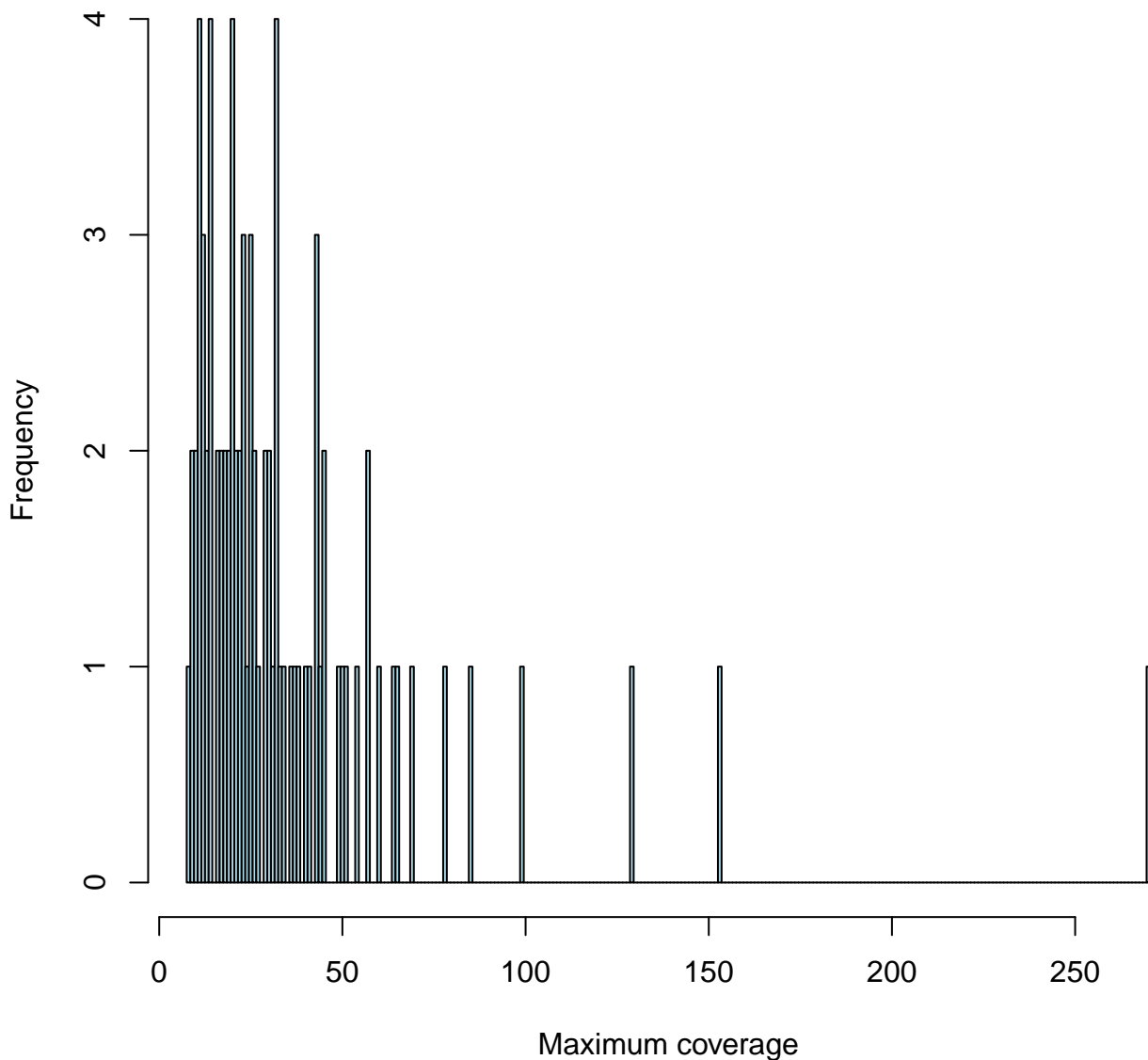
**jx in Intropolis but not "annotated" by knownGene:
176 samples; n jx: 63**



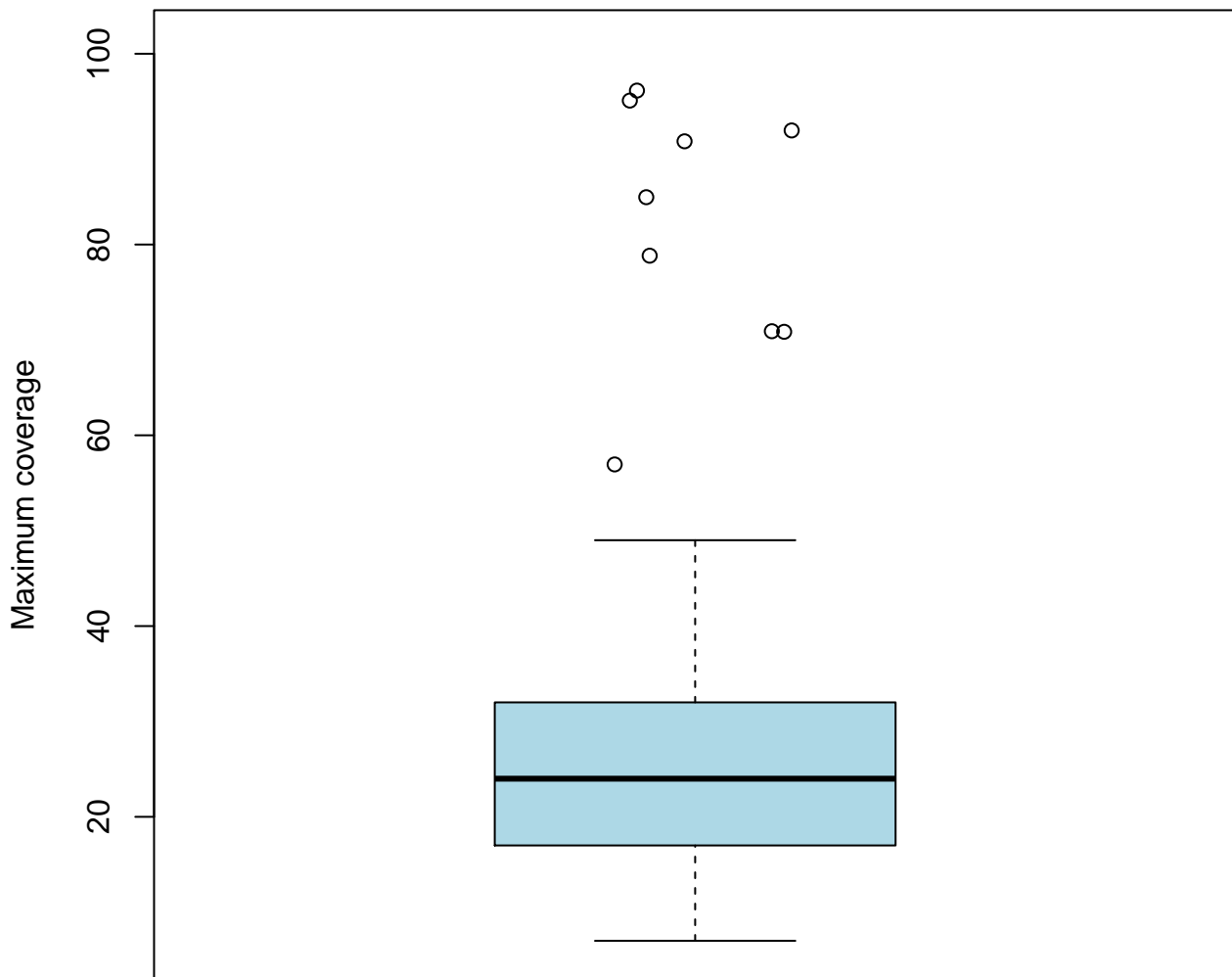
**jx in Intropolis but not "annotated" by knownGene:
177 samples; n jx: 82**



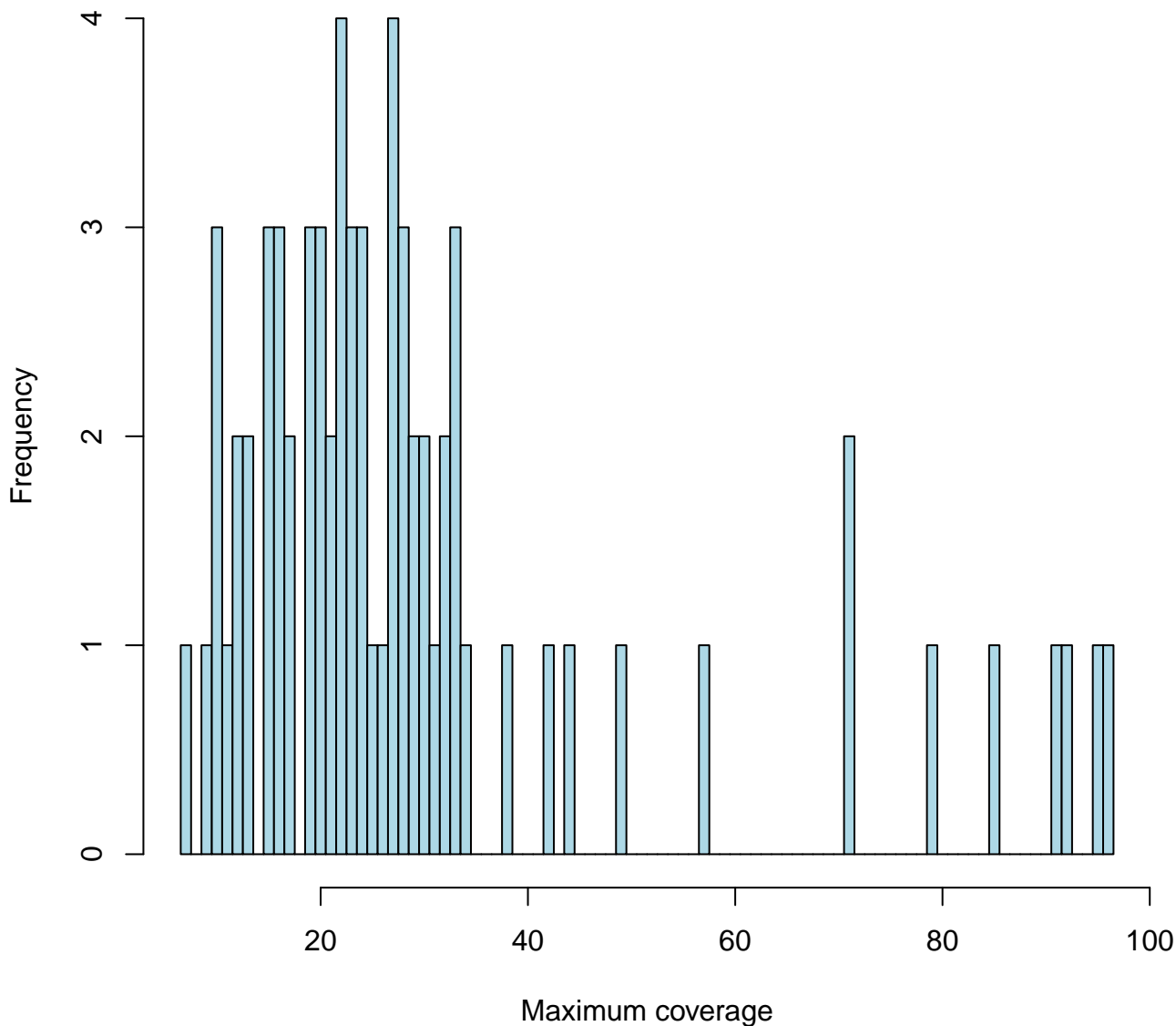
**jl in Intropolis but not "annotated" by knownGene:
177 samples; n jl: 82**



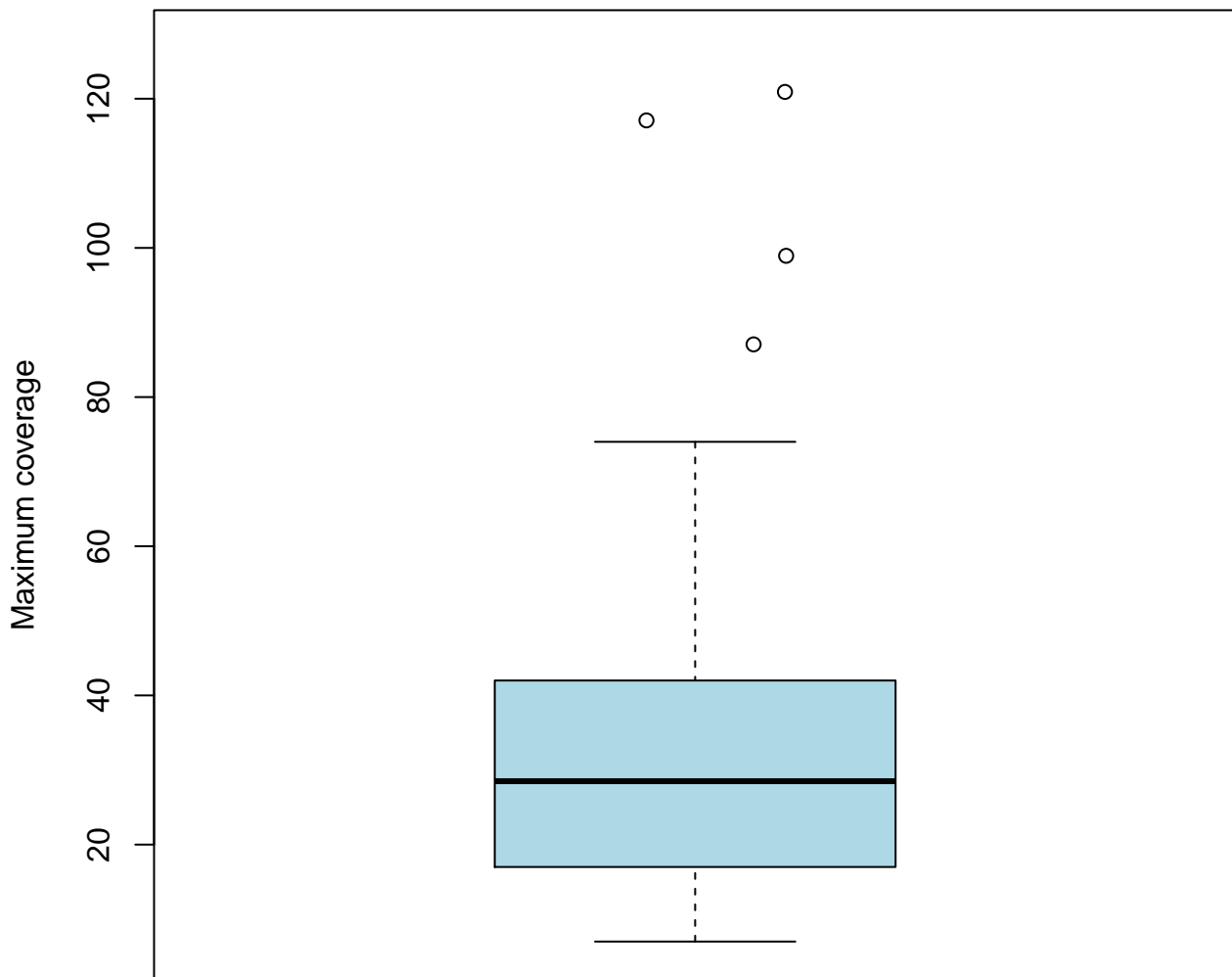
**jx in Intropolis but not "annotated" by knownGene:
178 samples; n jx: 69**



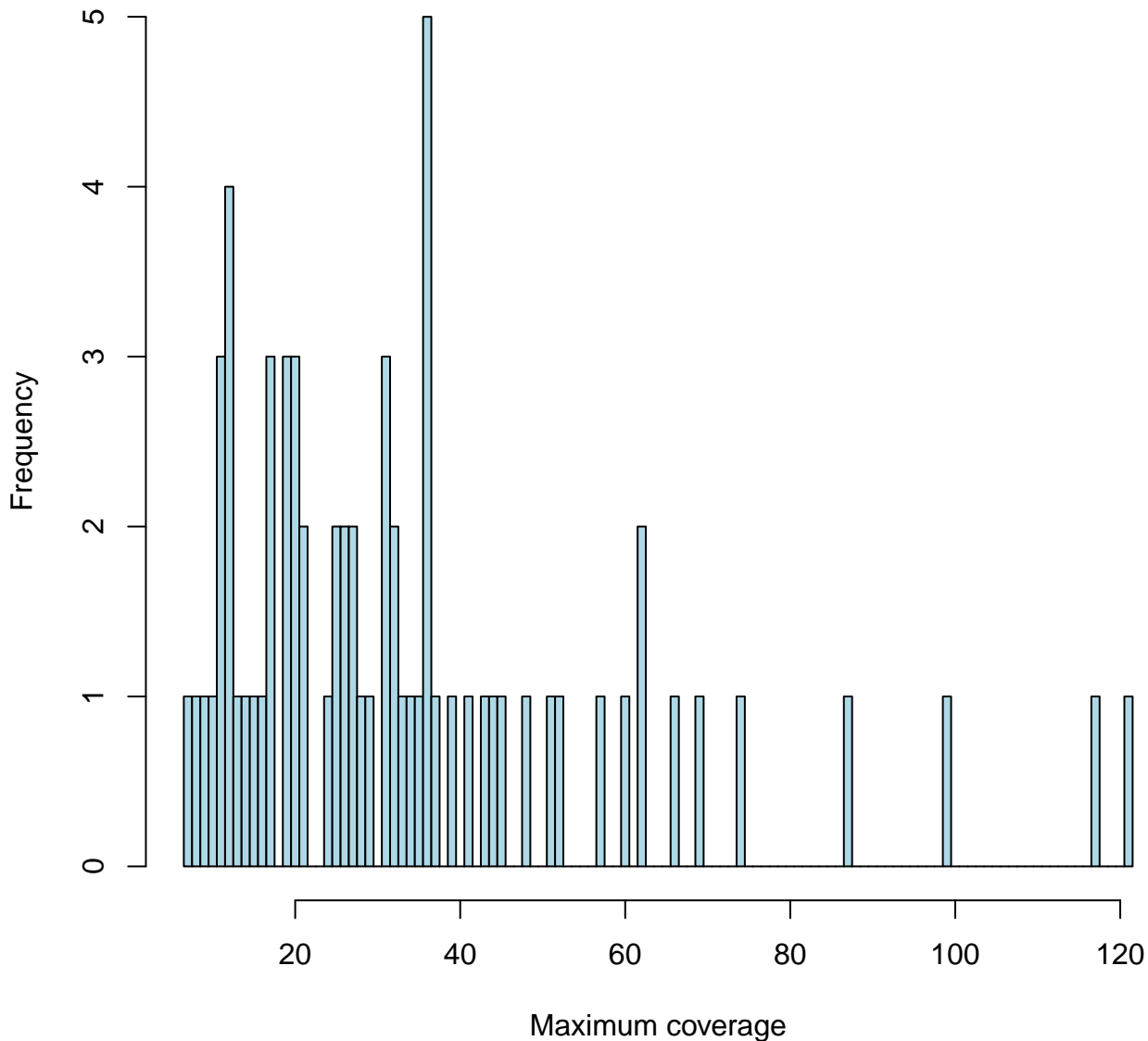
**jx in Intropolis but not "annotated" by knownGene:
178 samples; n jx: 69**



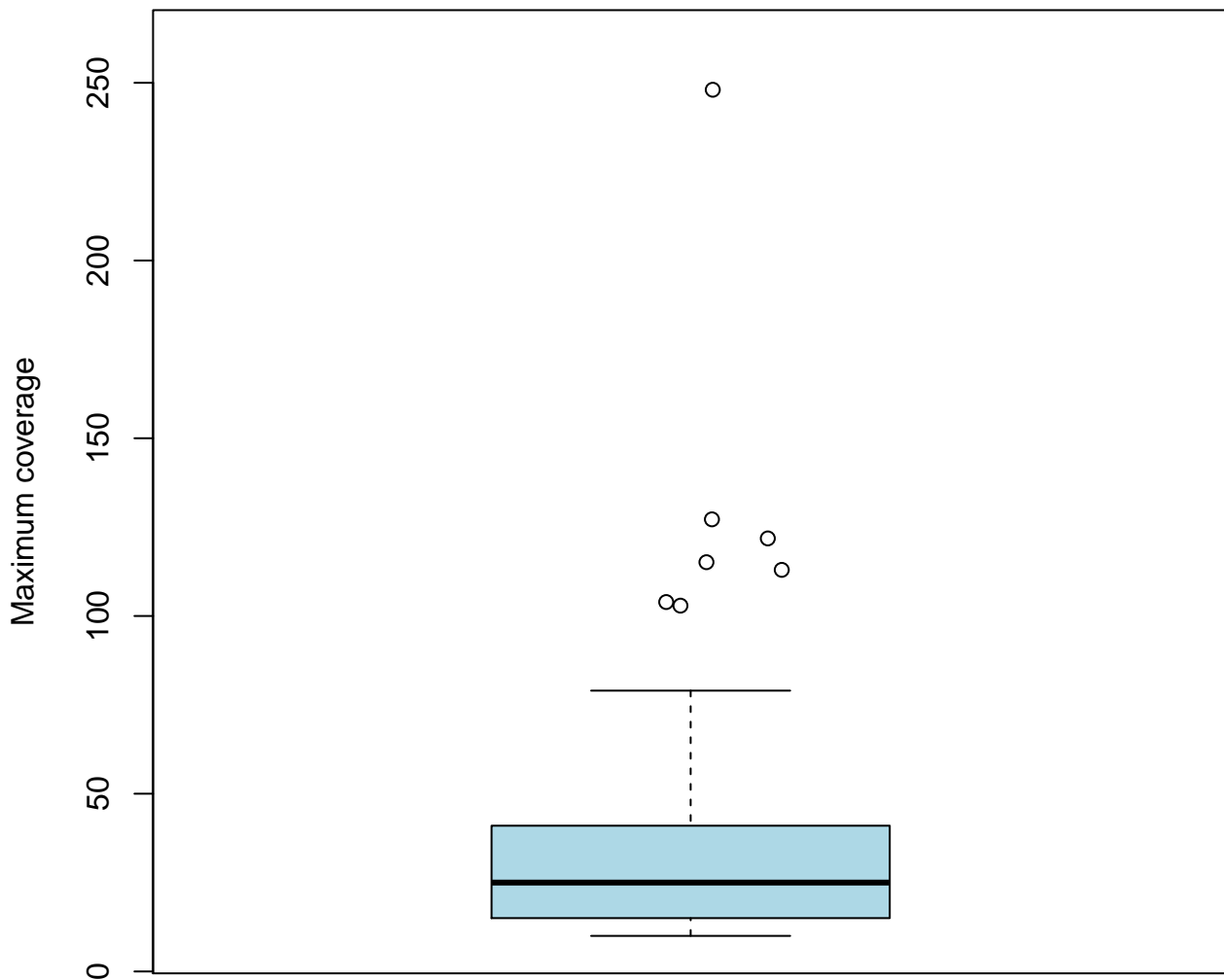
**jx in Intropolis but not "annotated" by knownGene:
179 samples; n jx: 68**



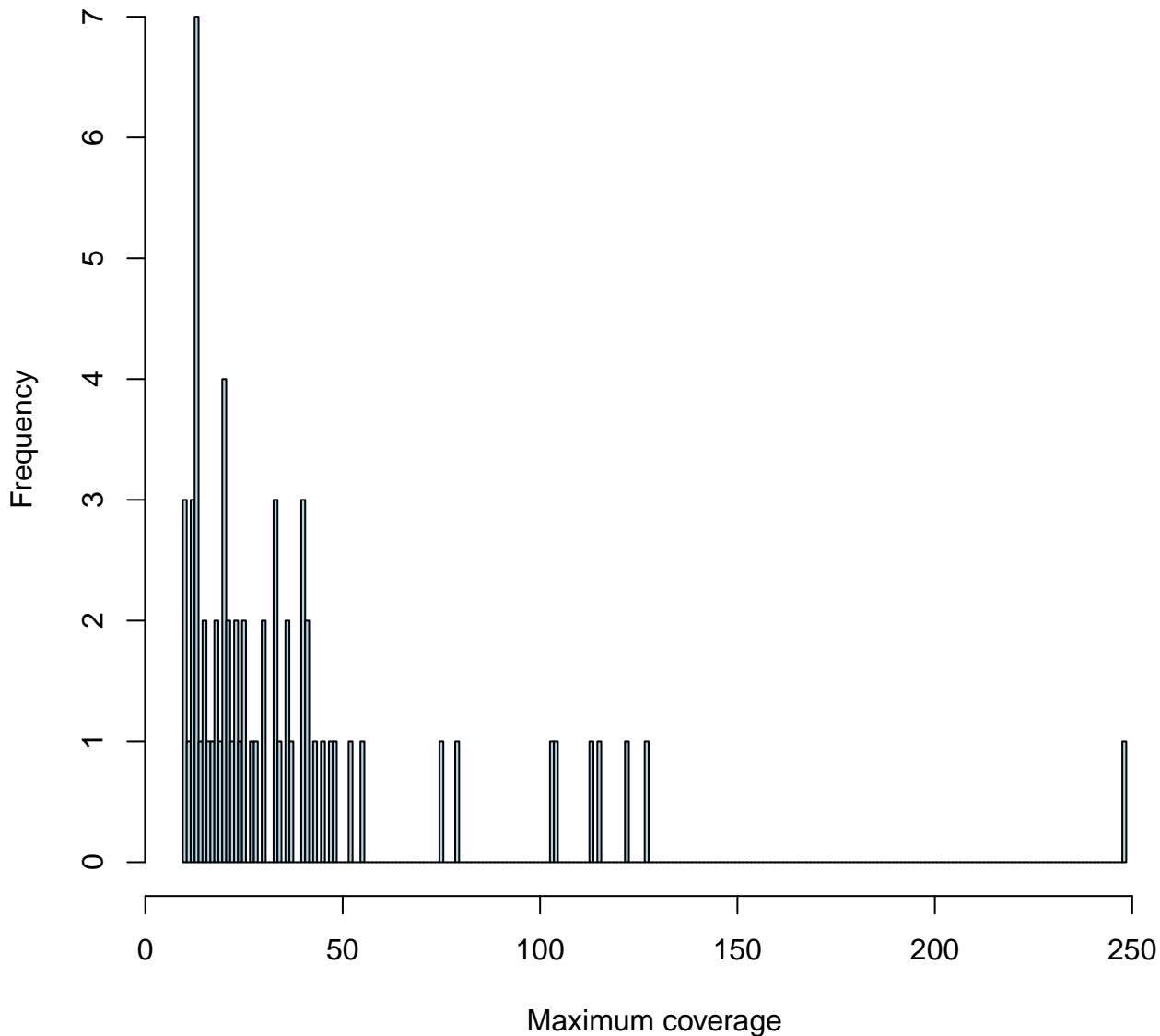
**jx in Intropolis but not "annotated" by knownGene:
179 samples; n jx: 68**



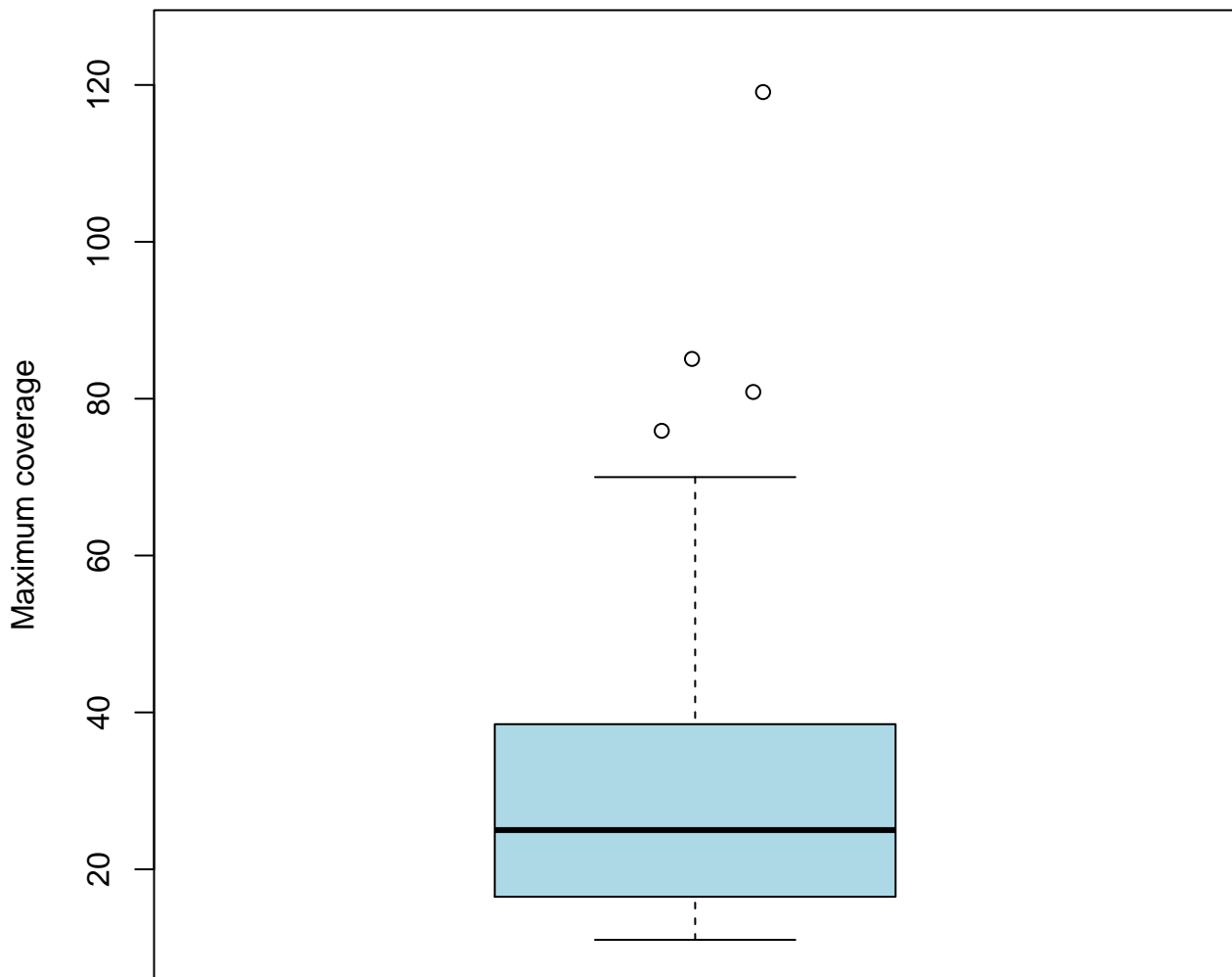
**jx in Intropolis but not "annotated" by knownGene:
180 samples; n jx: 65**



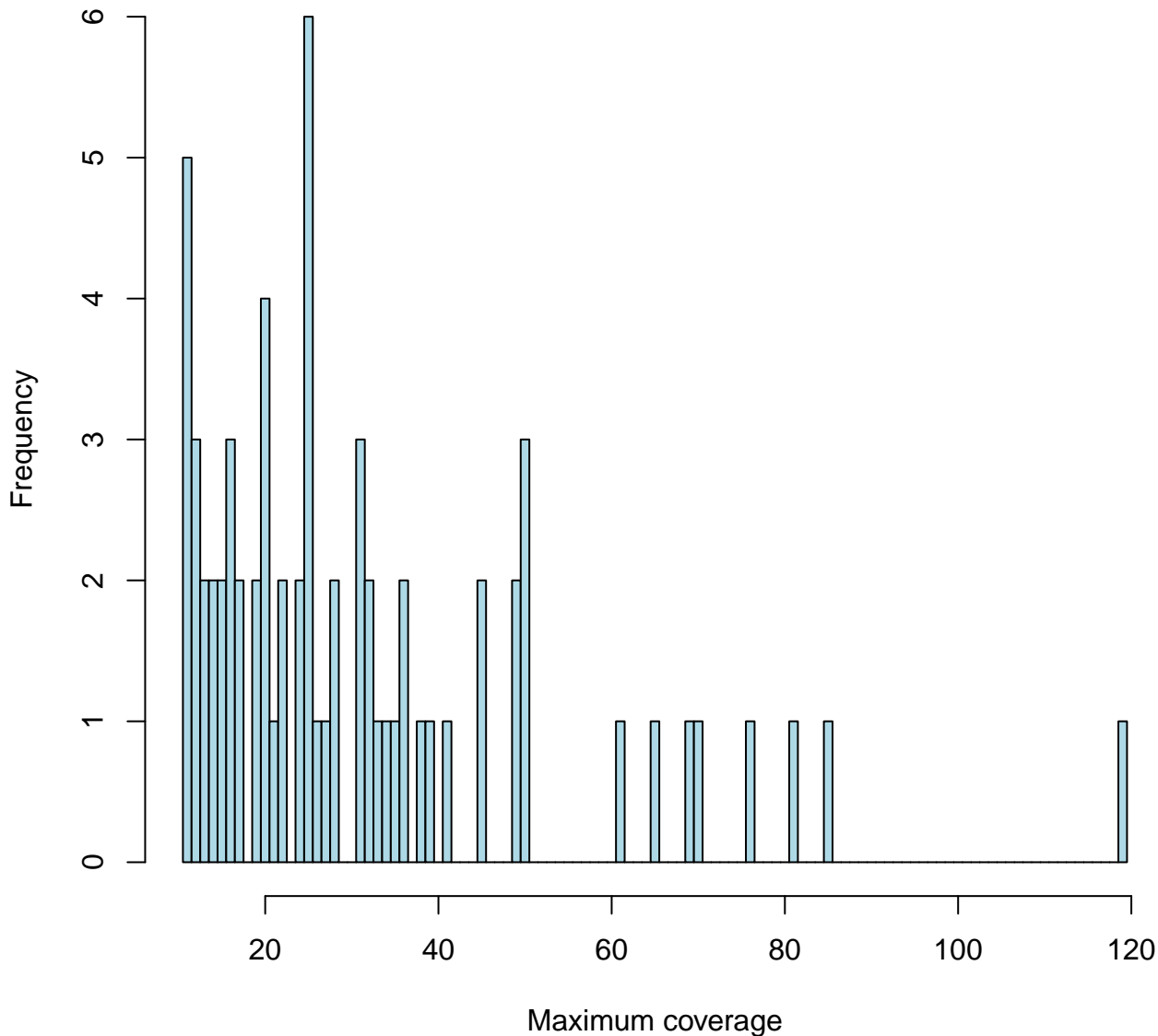
**jx in Intropolis but not "annotated" by knownGene:
180 samples; n jx: 65**



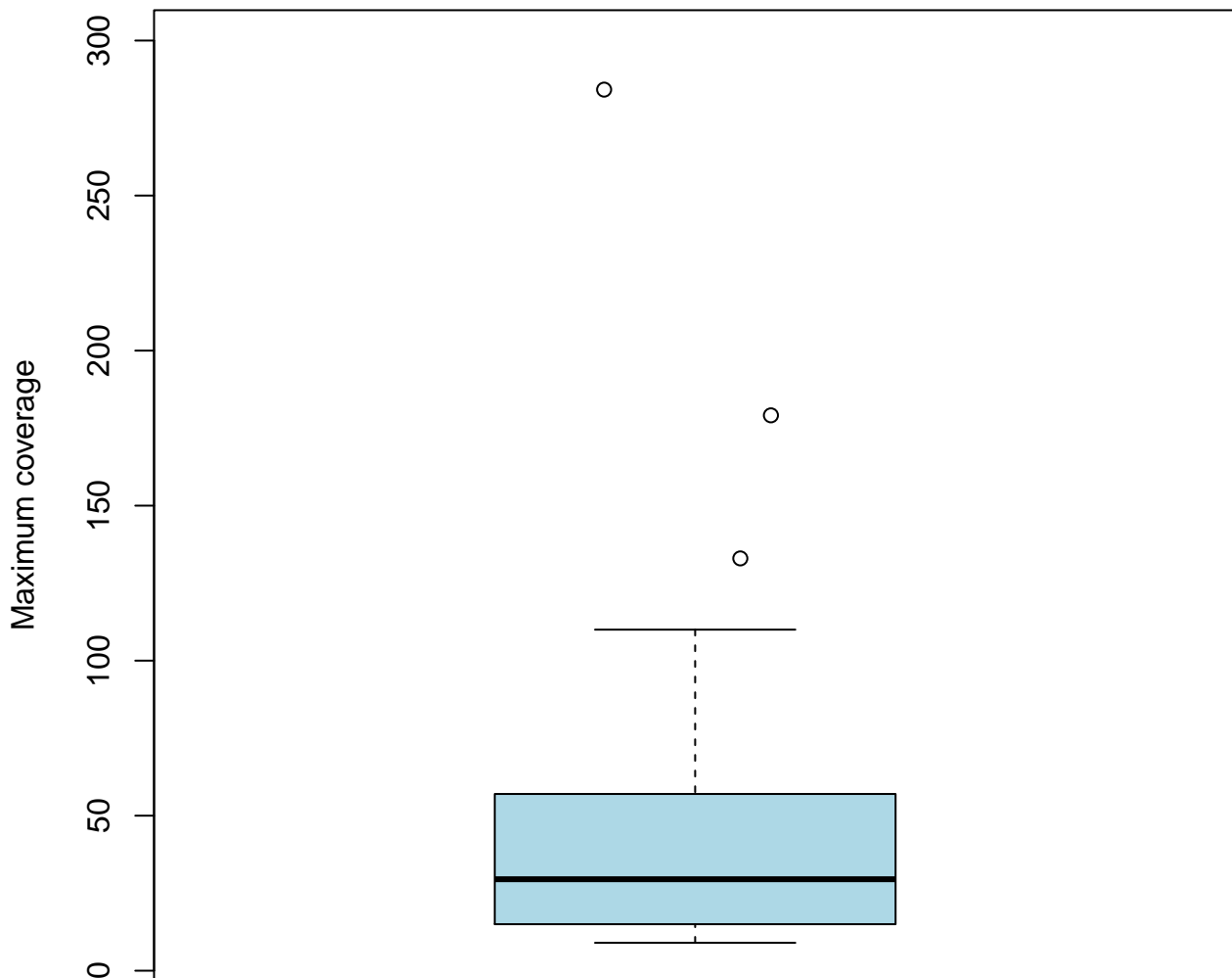
**jx in Intropolis but not "annotated" by knownGene:
181 samples; n jx: 68**



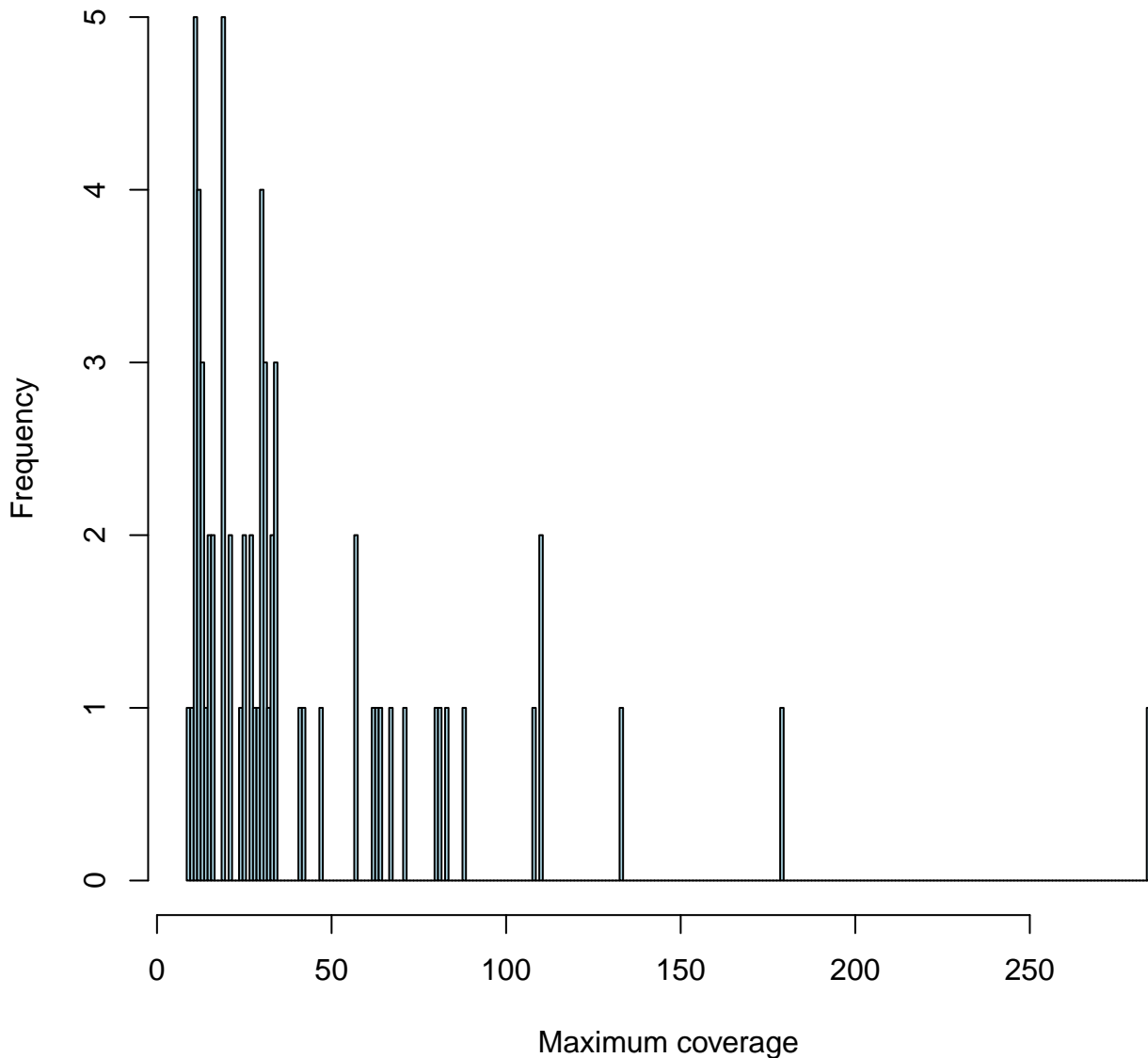
**jx in Intropolis but not "annotated" by knownGene:
181 samples; n jx: 68**



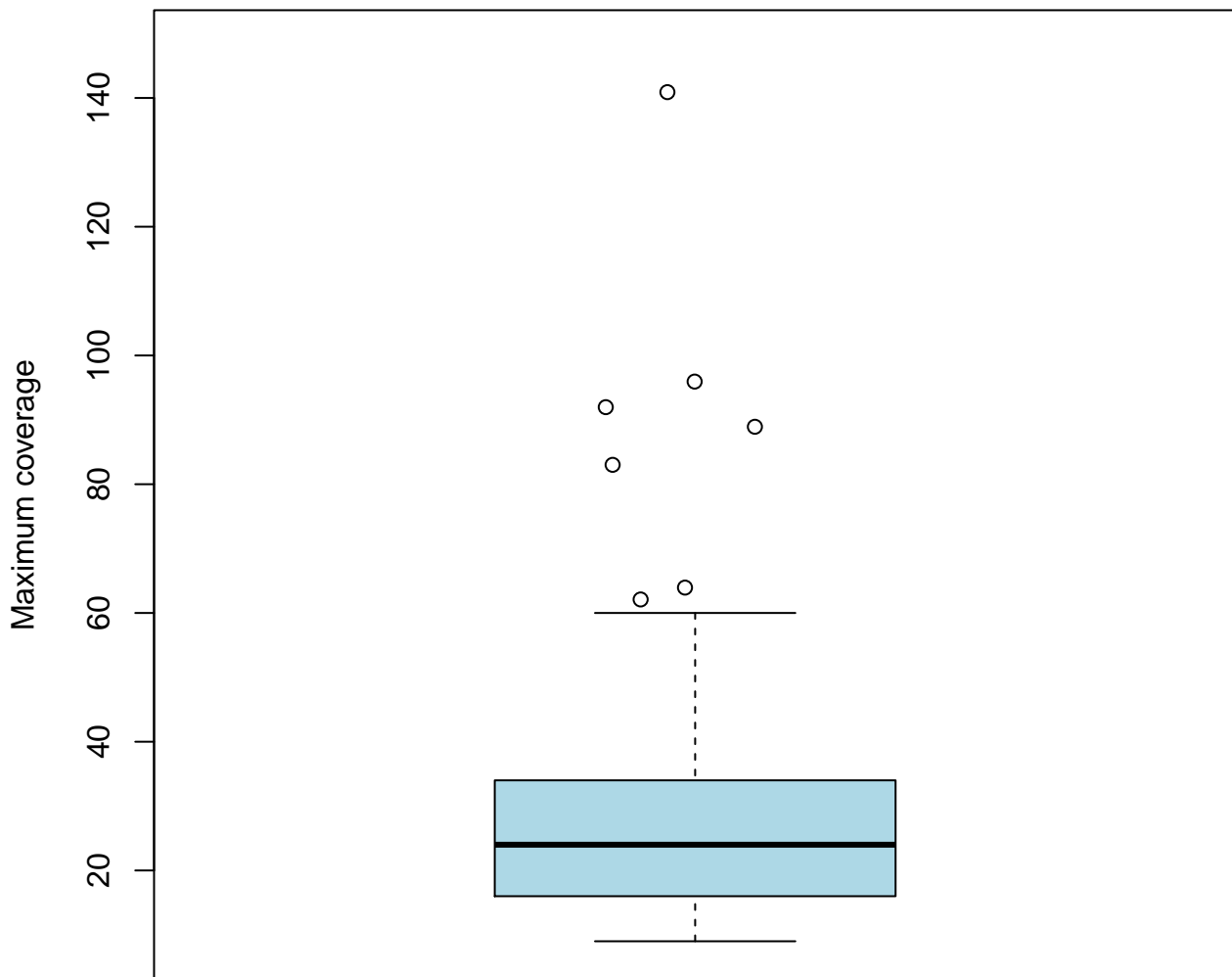
**jx in Intropolis but not "annotated" by knownGene:
182 samples; n jx: 66**



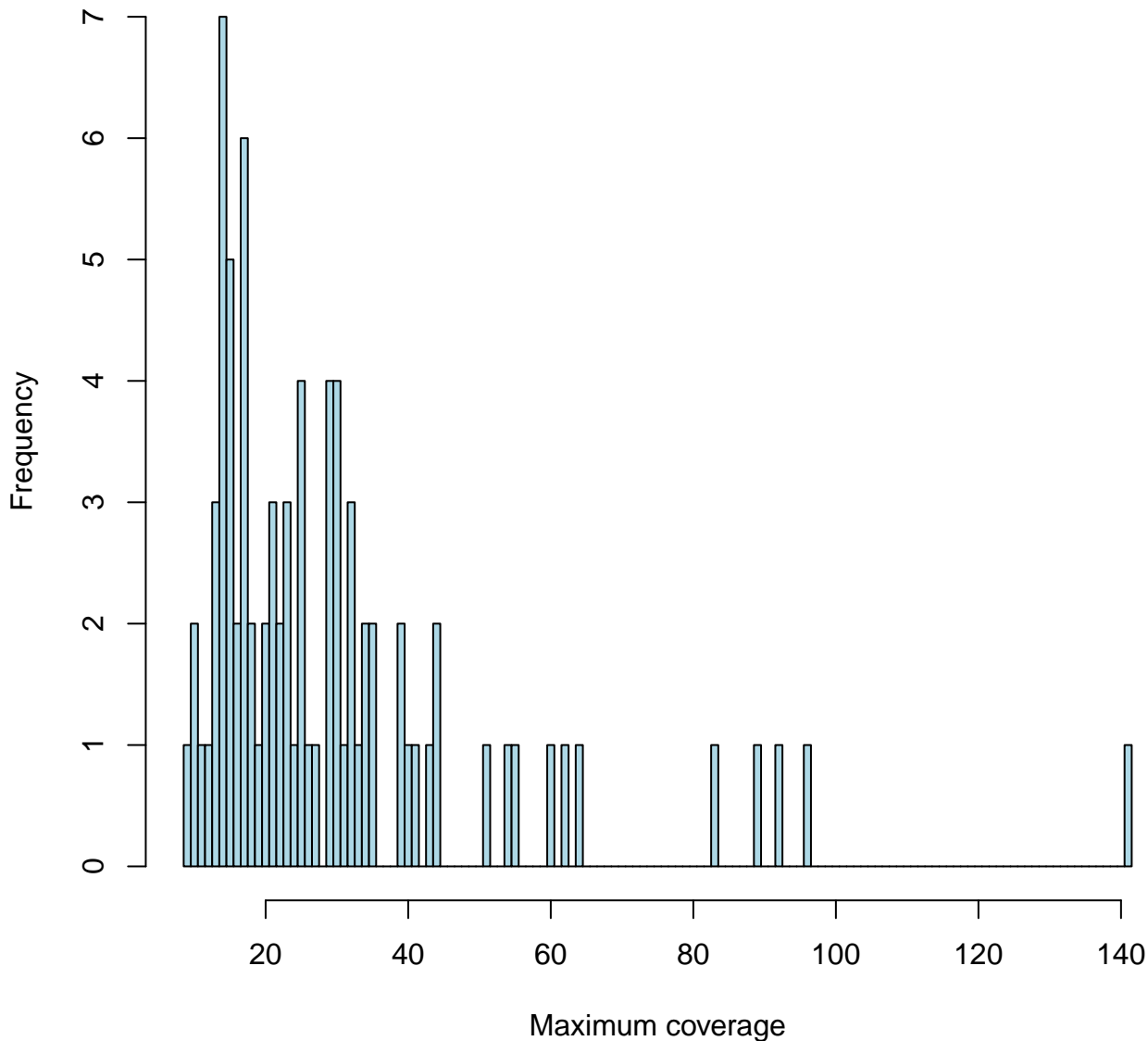
**jx in Intropolis but not "annotated" by knownGene:
182 samples; n jx: 66**



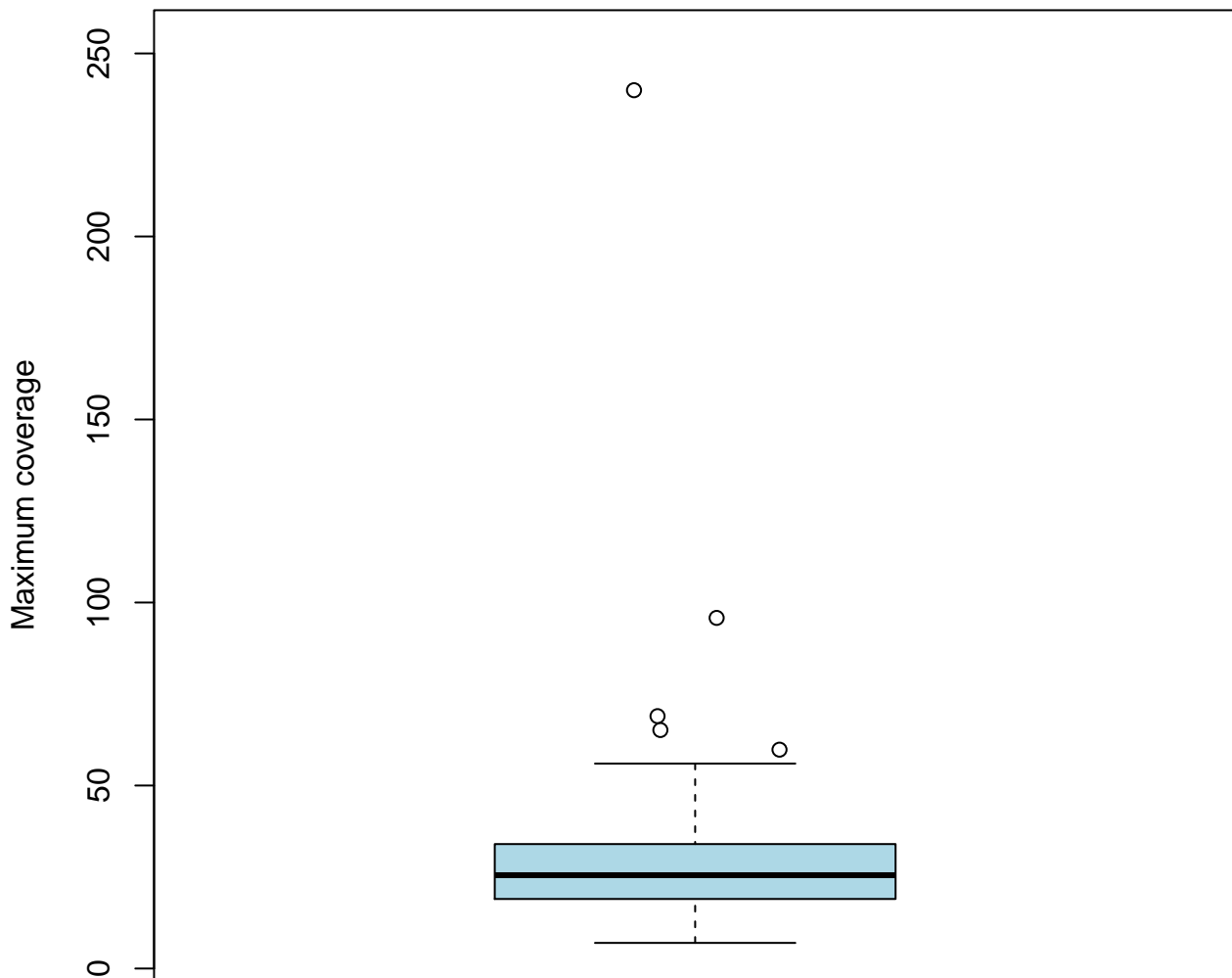
**jx in Intropolis but not "annotated" by knownGene:
183 samples; n jx: 83**



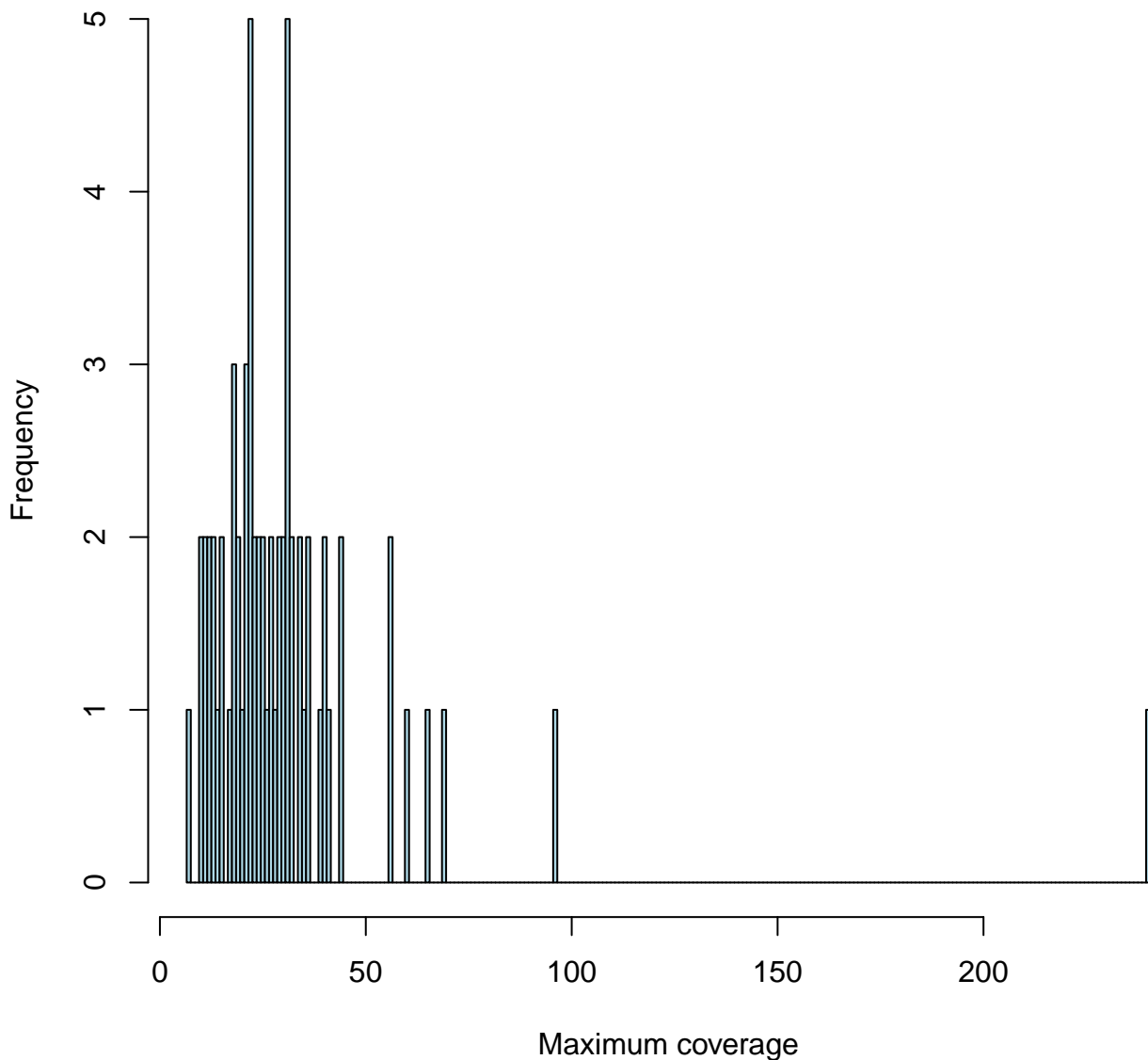
**jx in Intropolis but not "annotated" by knownGene:
183 samples; n jx: 83**



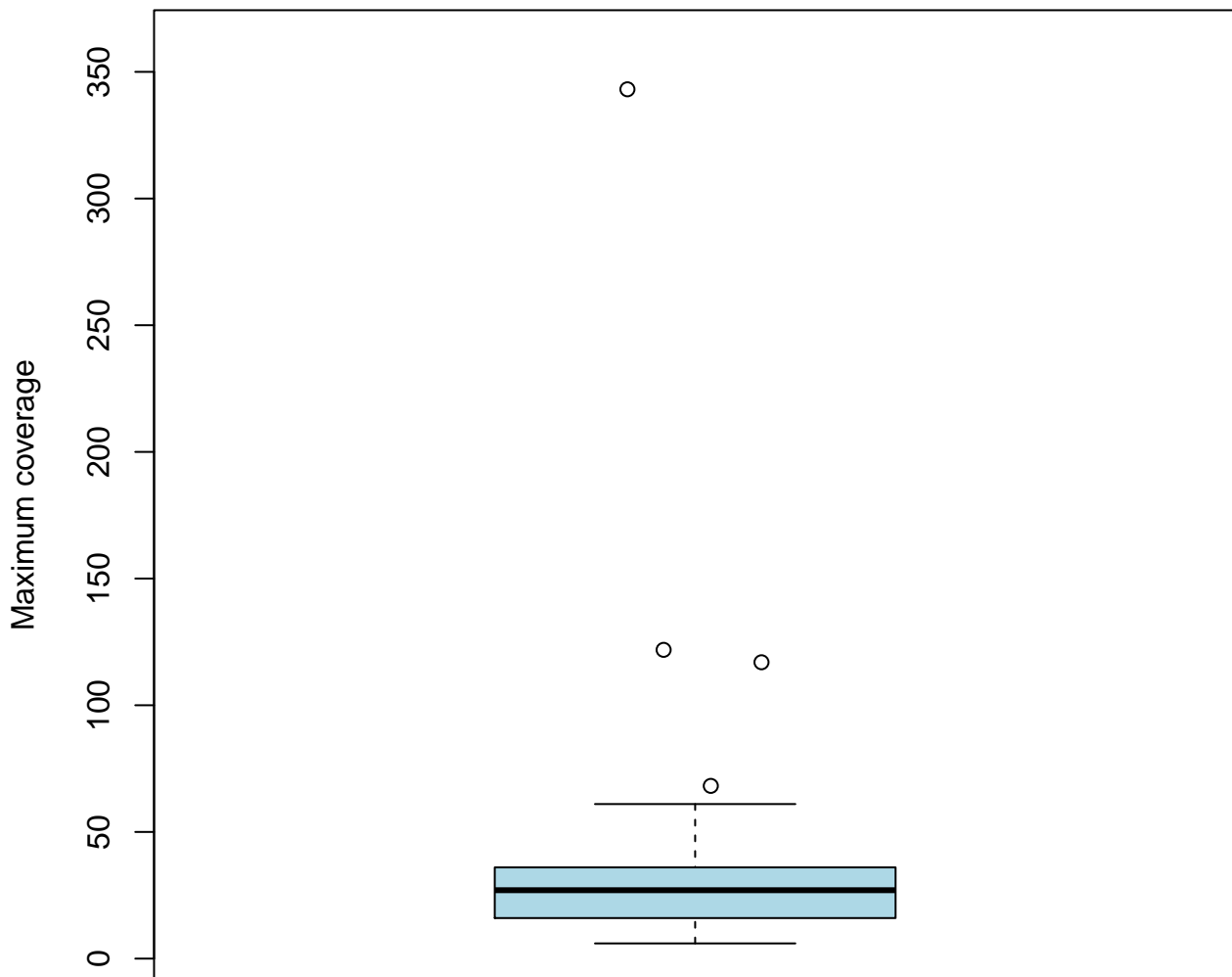
**jx in Intropolis but not "annotated" by knownGene:
184 samples; n jx: 66**



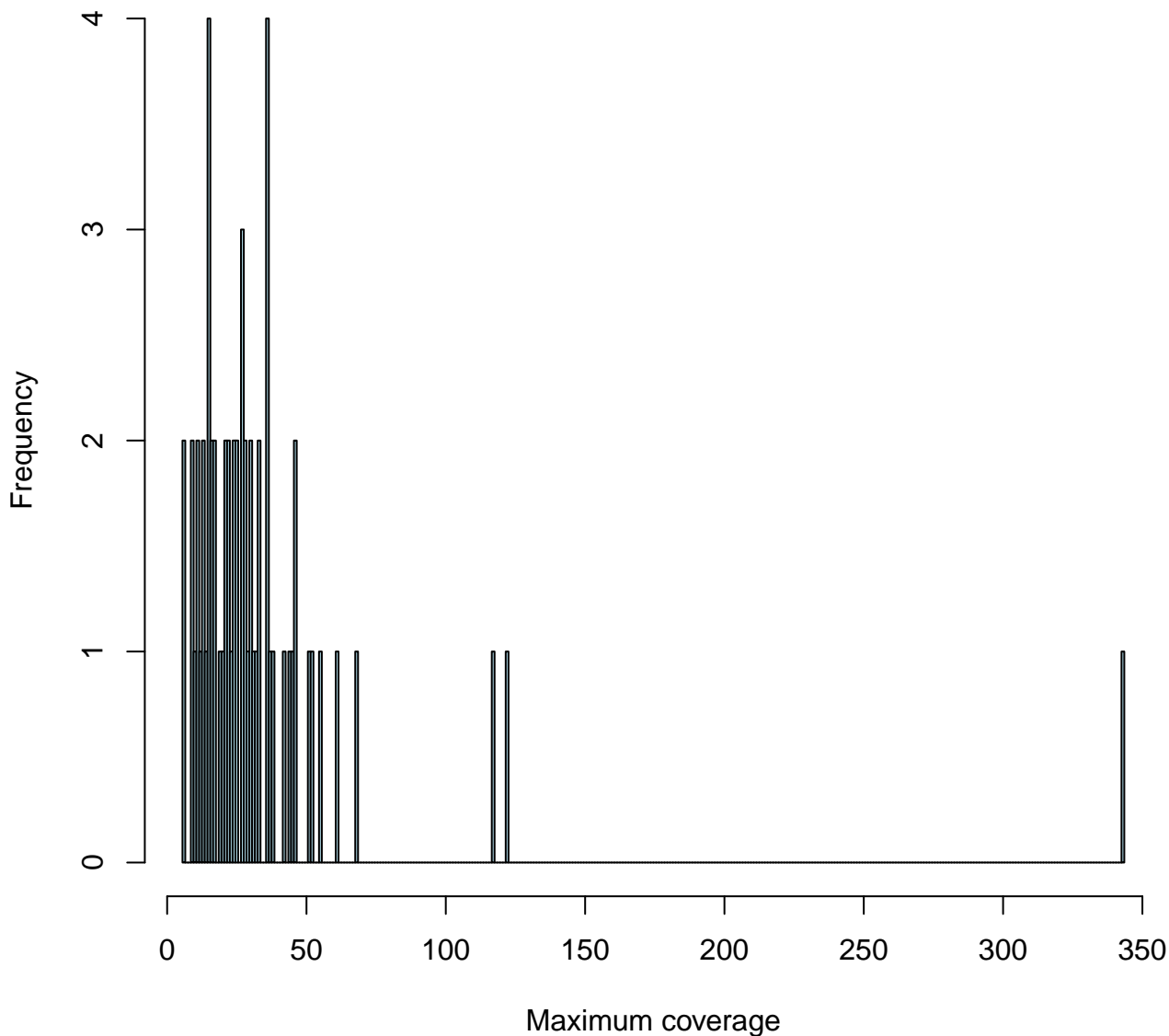
**jx in Intropolis but not "annotated" by knownGene:
184 samples; n jx: 66**



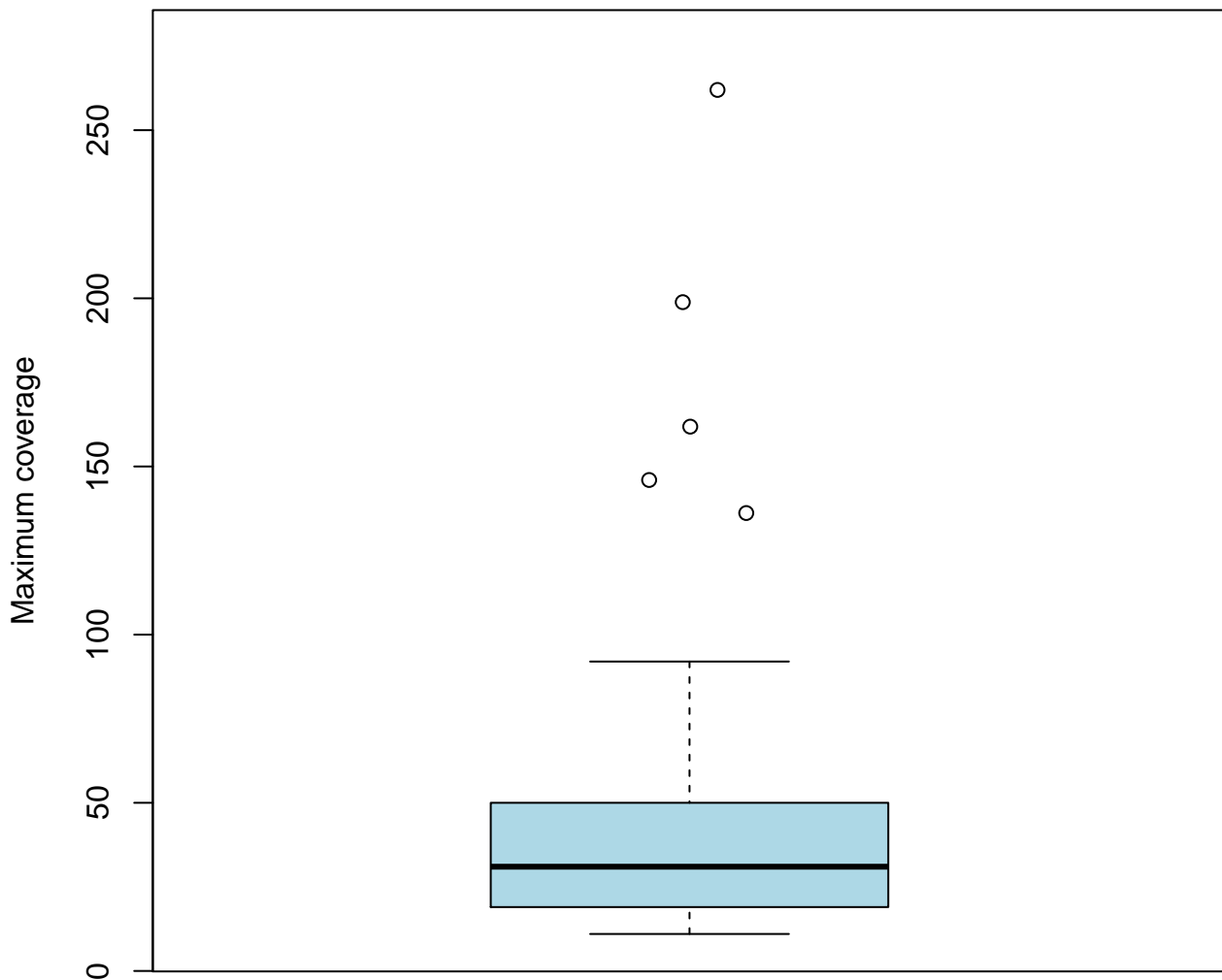
**jx in Intropolis but not "annotated" by knownGene:
185 samples; n jx: 61**



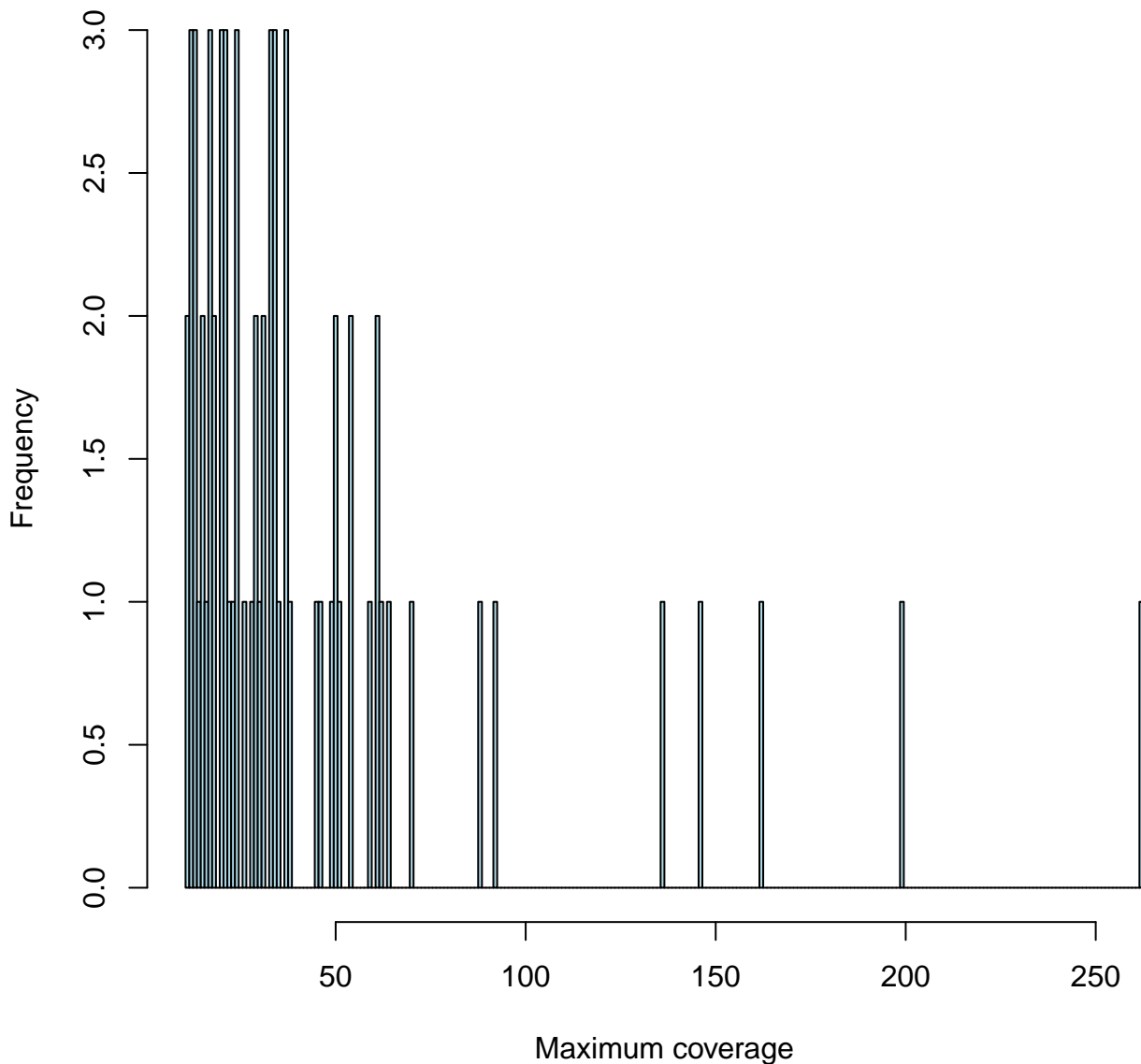
**jx in Intropolis but not "annotated" by knownGene:
185 samples; n jx: 61**



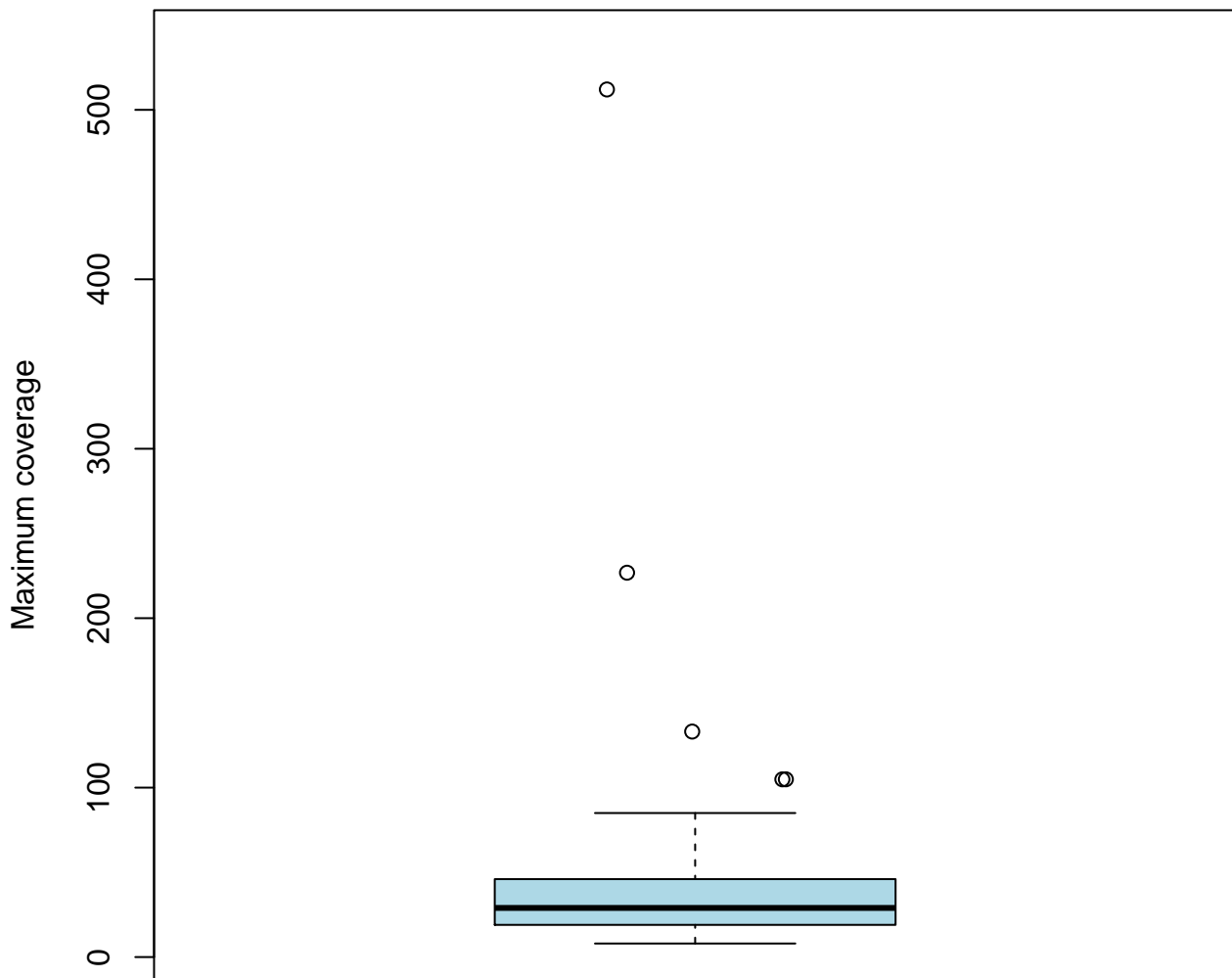
**jx in Intropolis but not "annotated" by knownGene:
186 samples; n jx: 67**



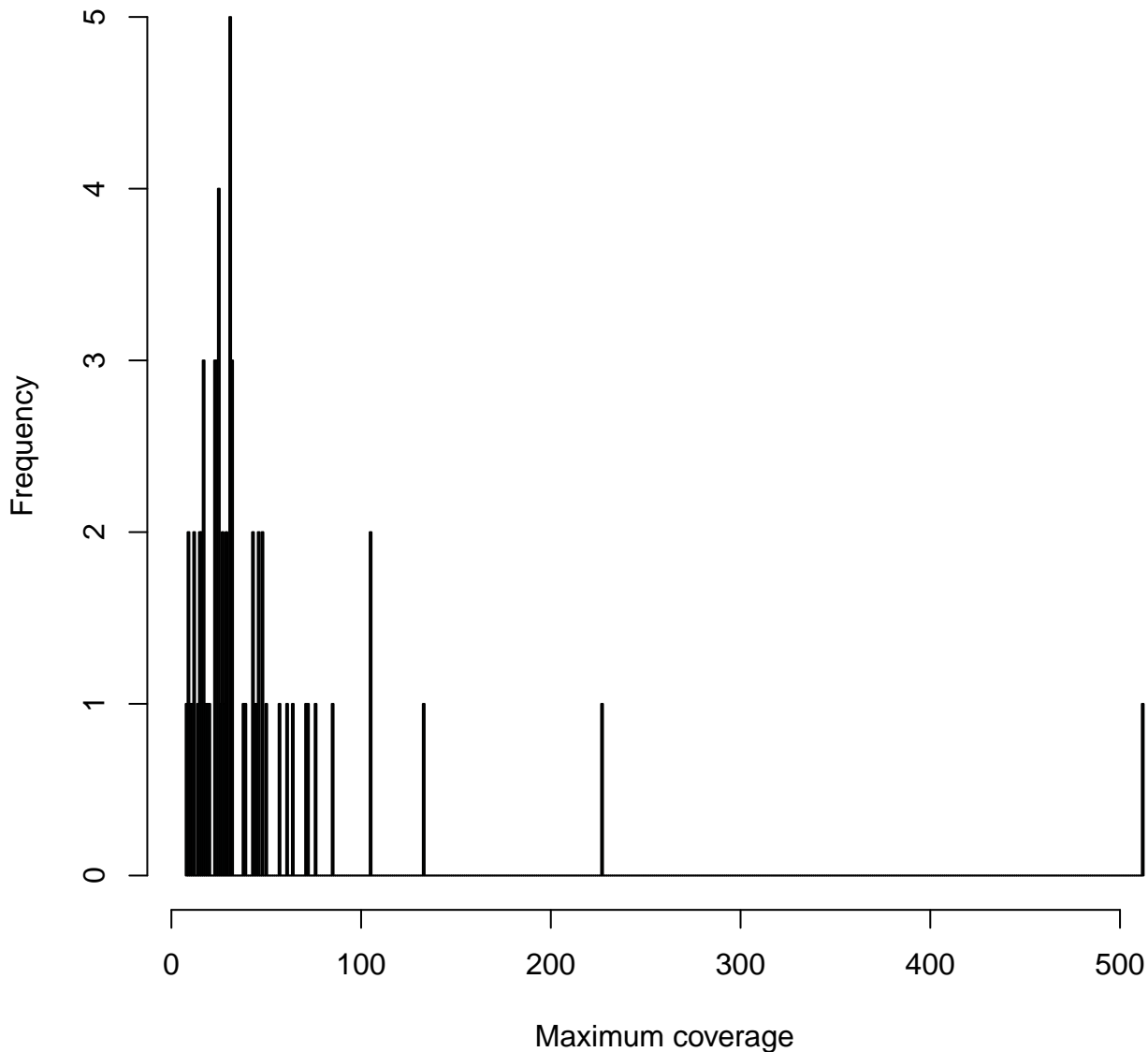
**jx in Intropolis but not "annotated" by knownGene:
186 samples; n jx: 67**



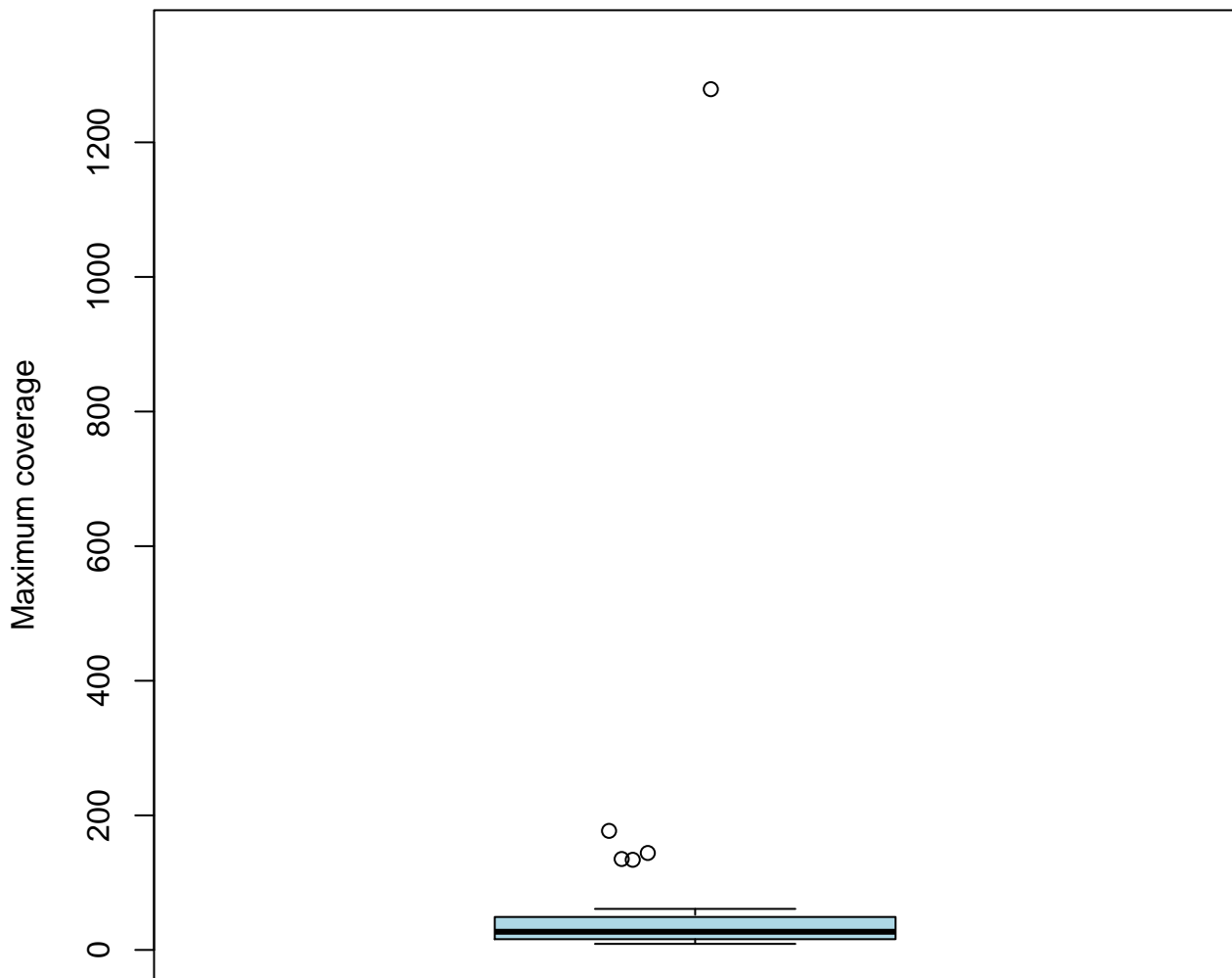
**jx in Intropolis but not "annotated" by knownGene:
187 samples; n jx: 65**



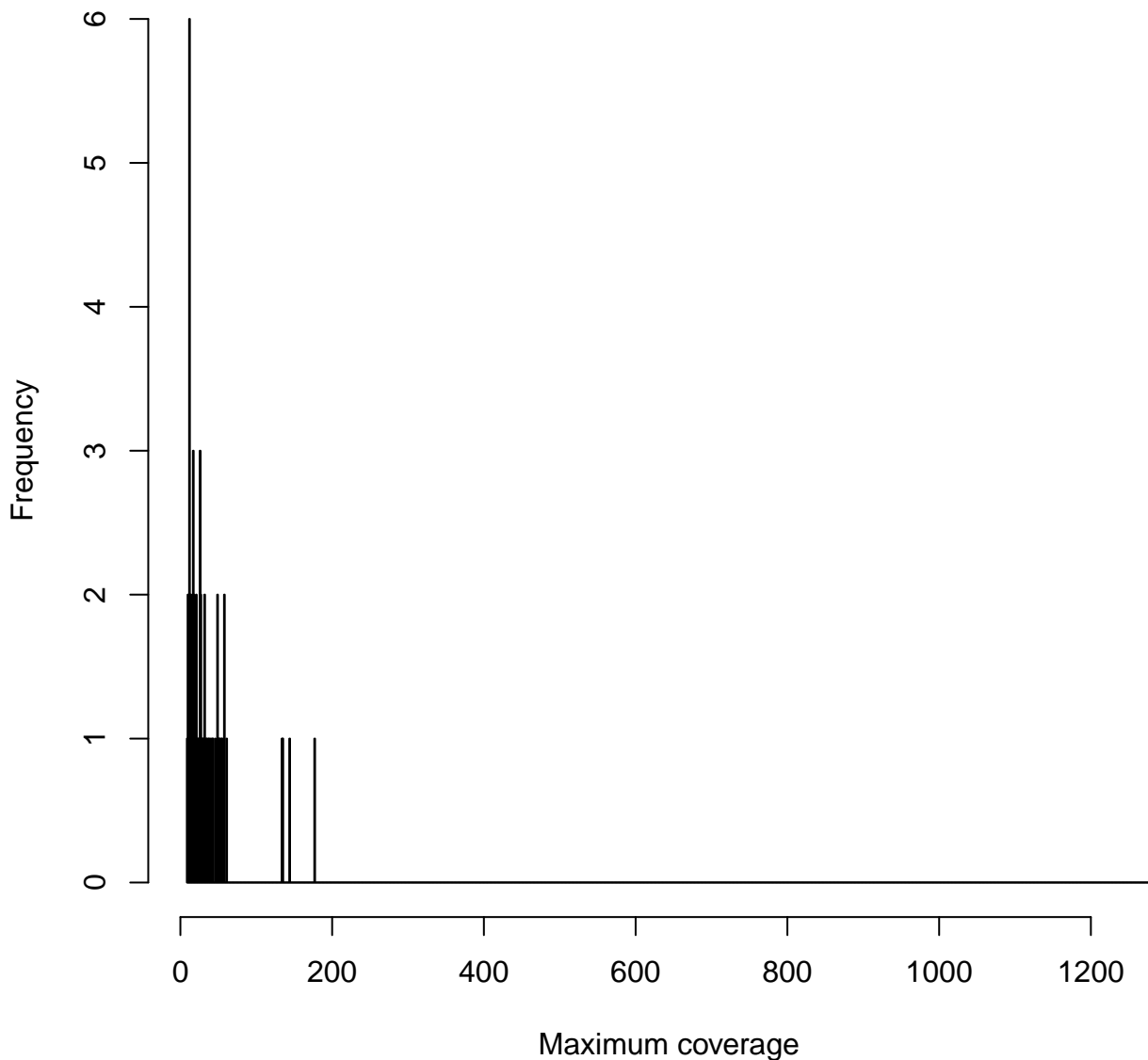
**jx in Intropolis but not "annotated" by knownGene:
187 samples; n jx: 65**



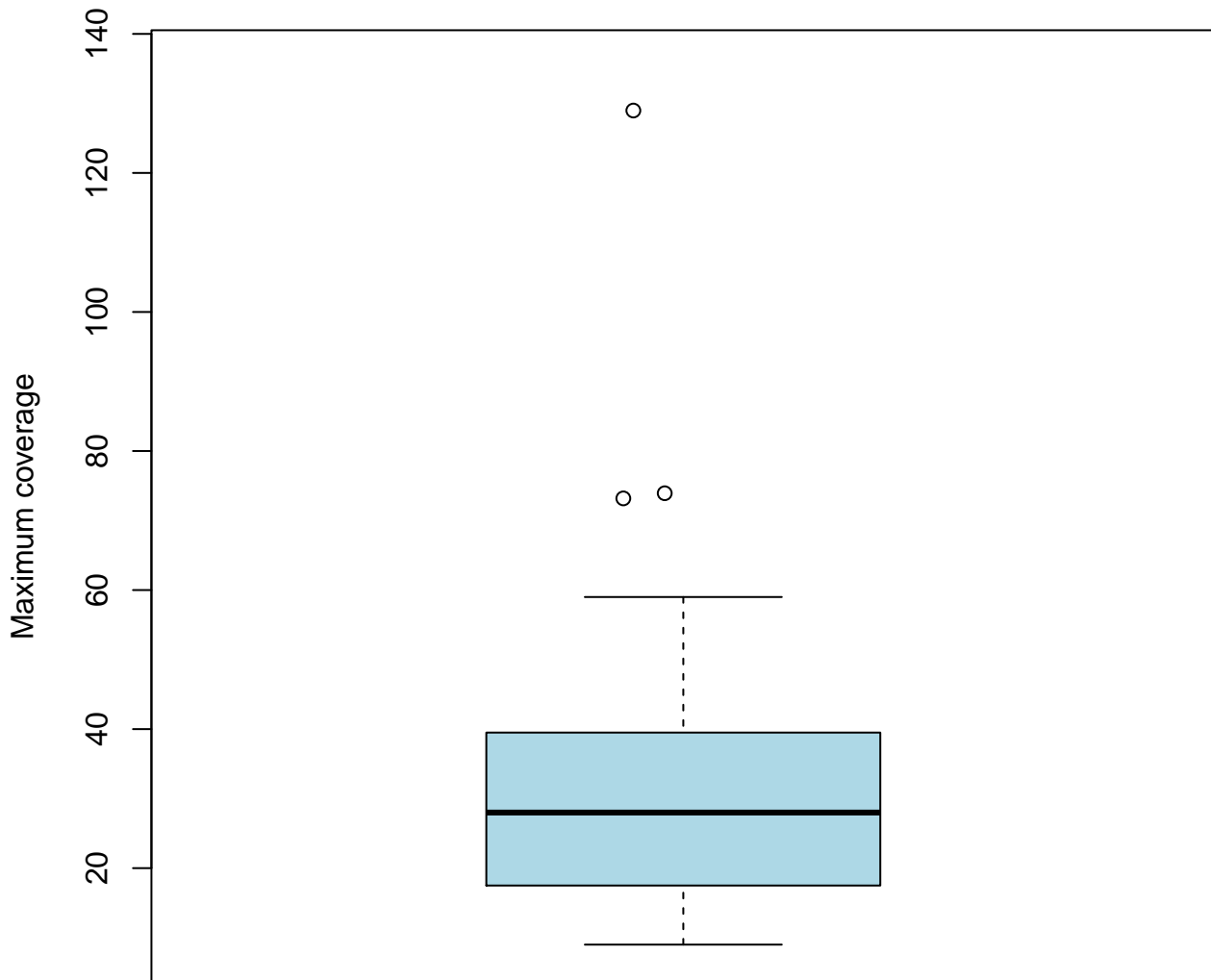
**jx in Intropolis but not "annotated" by knownGene:
188 samples; n jx: 57**



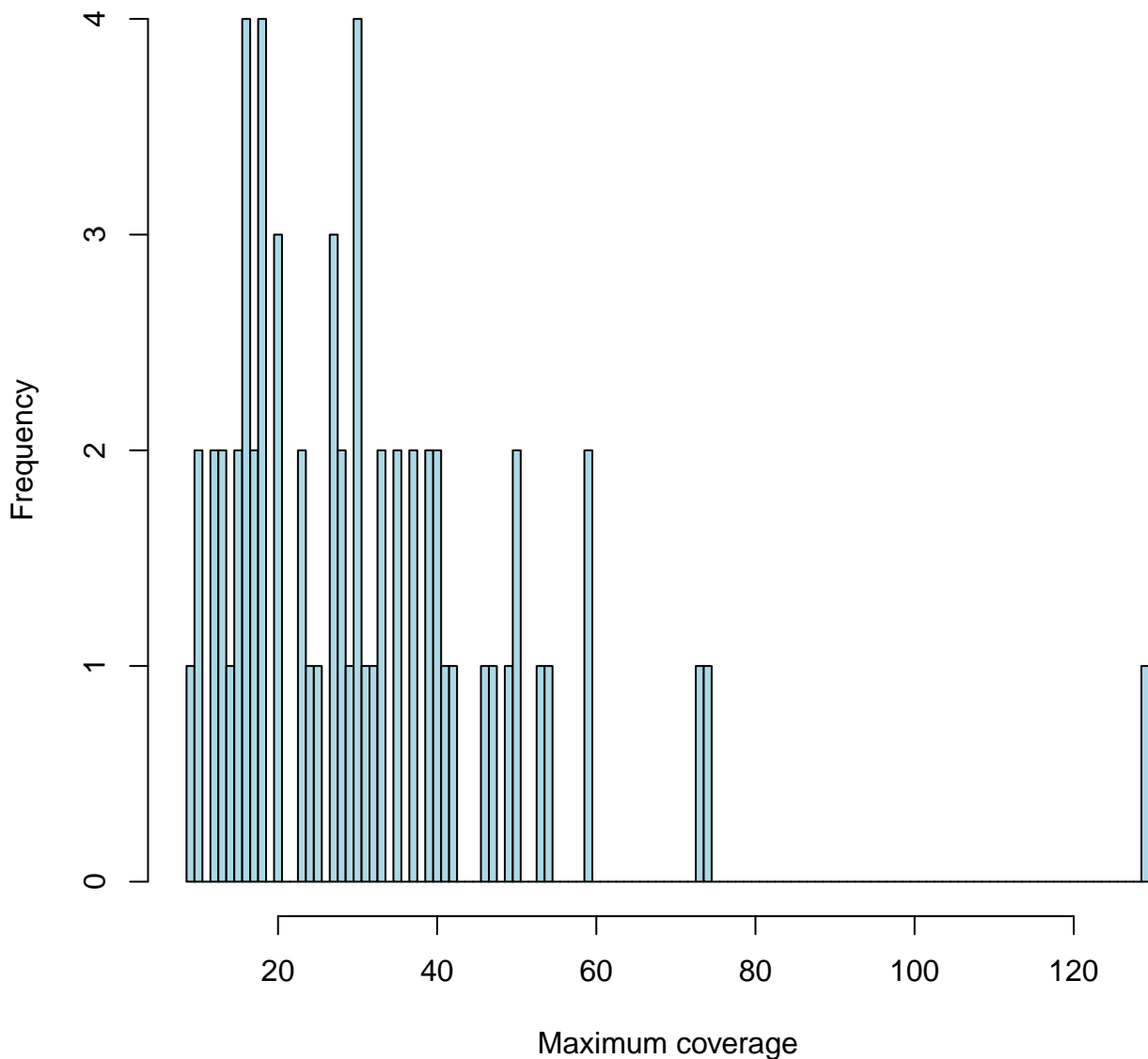
**jx in Intropolis but not "annotated" by knownGene:
188 samples; n jx: 57**



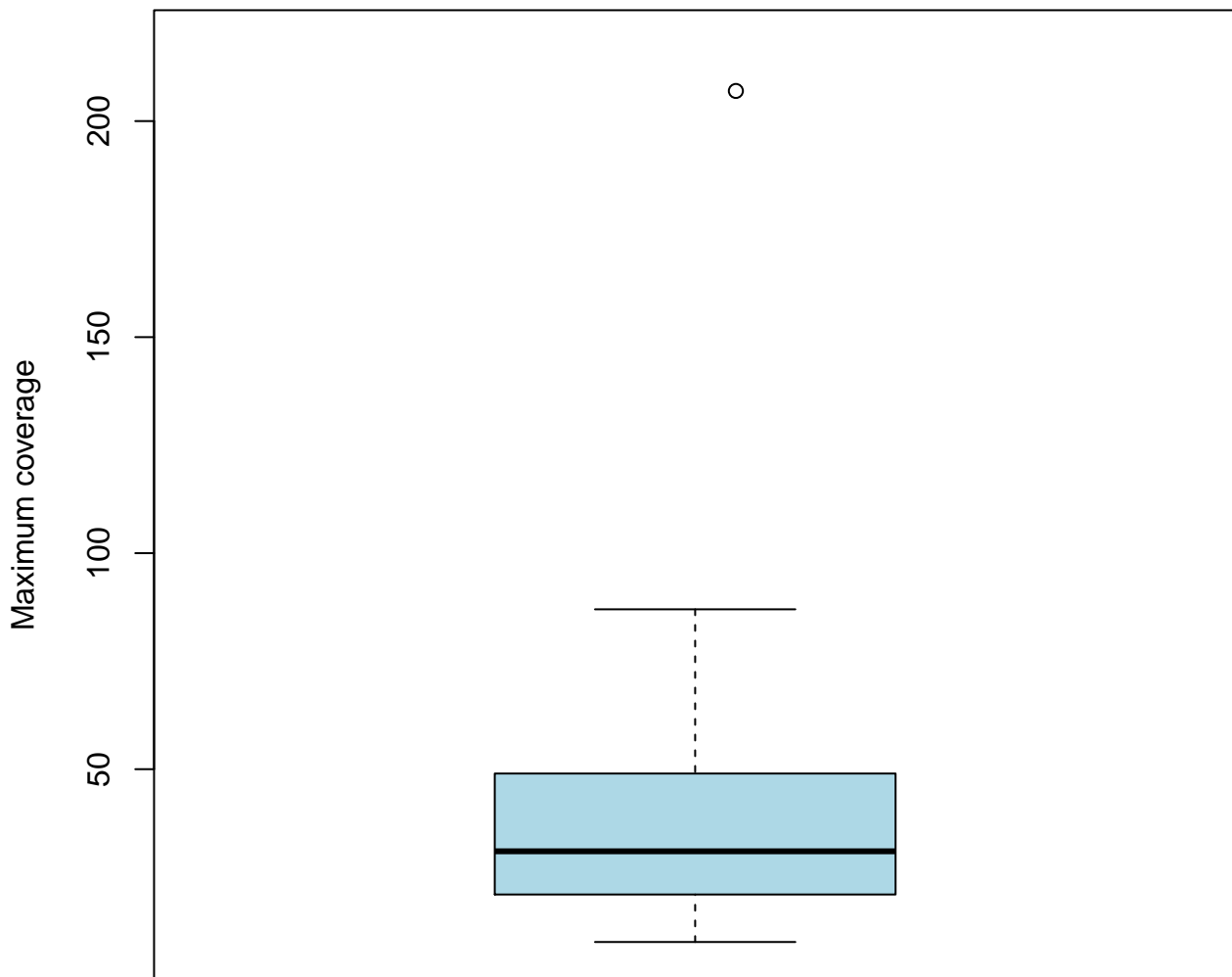
**jx in Intropolis but not "annotated" by knownGene:
189 samples; n jx: 63**



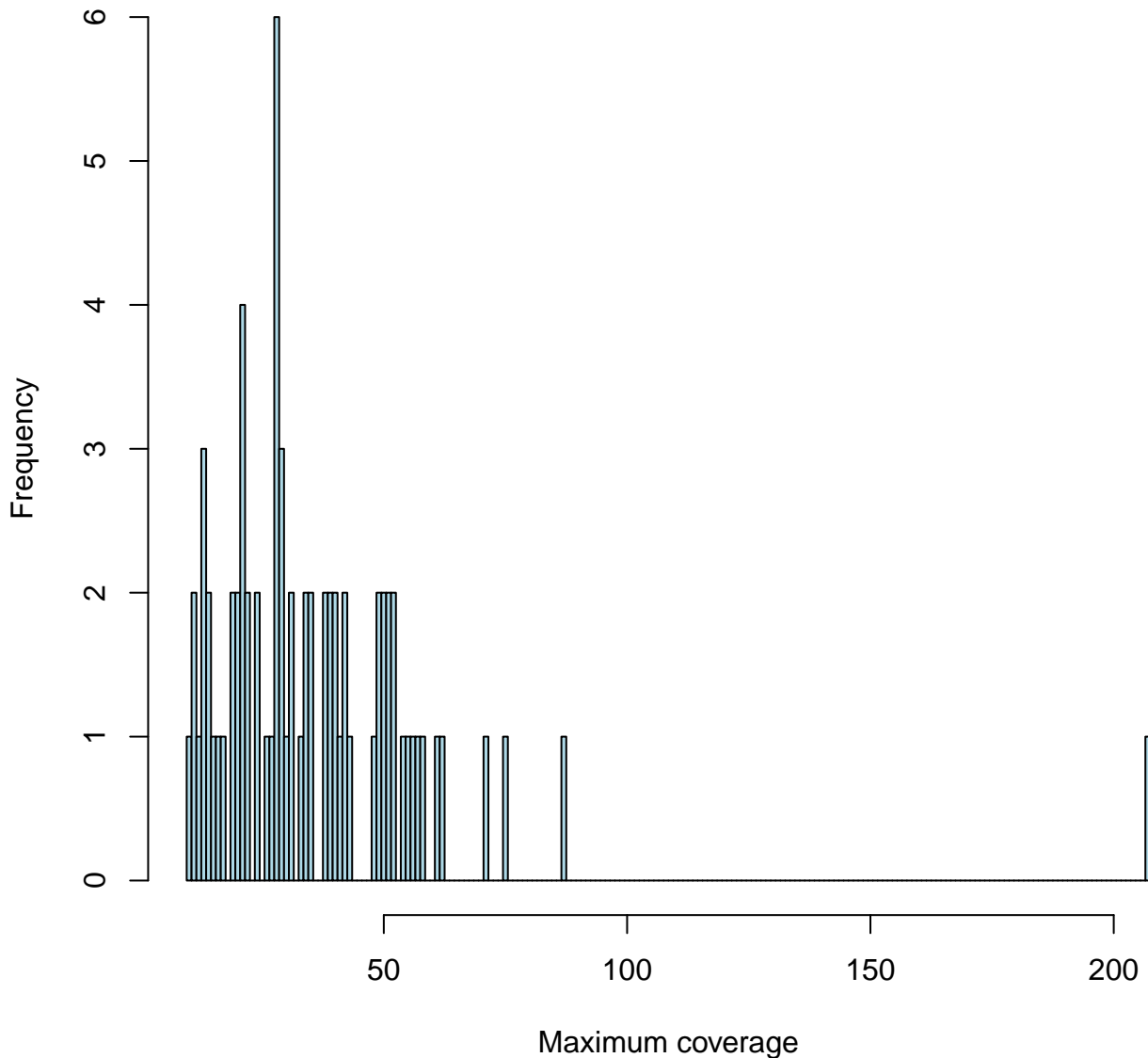
**jx in Intropolis but not "annotated" by knownGene:
189 samples; n jx: 63**



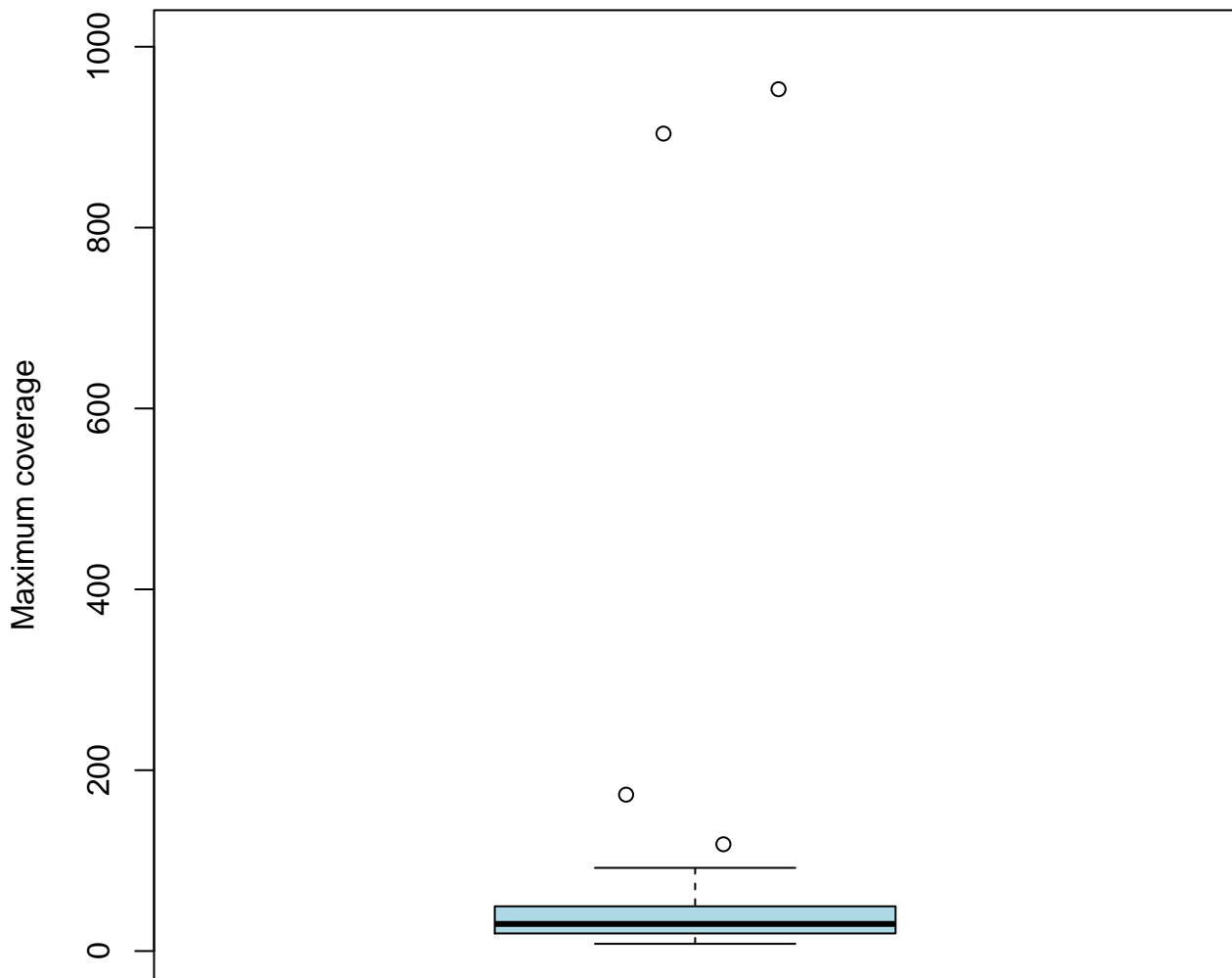
**jx in Intropolis but not "annotated" by knownGene:
190 samples; n jx: 73**



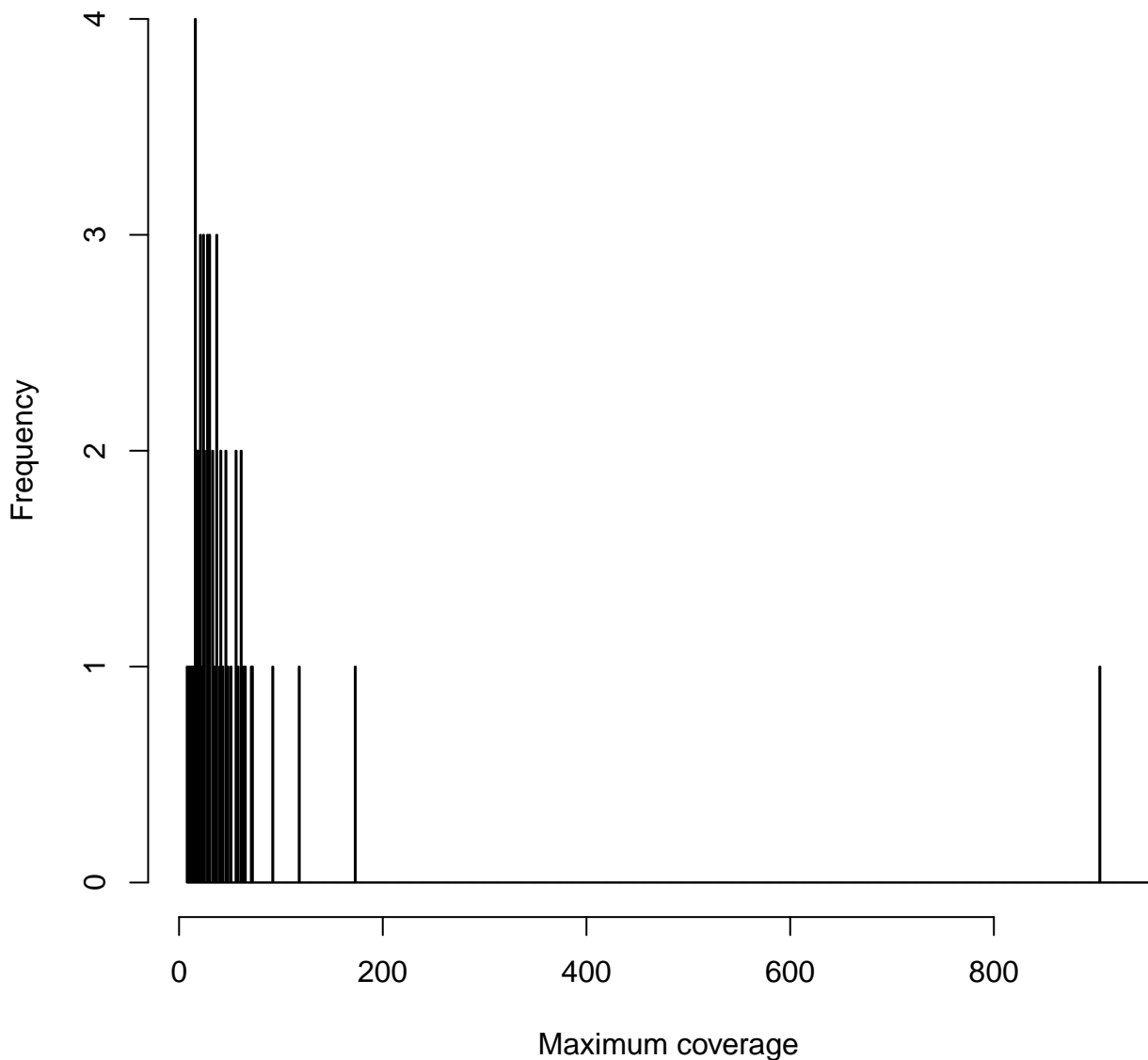
**jx in Intropolis but not "annotated" by knownGene:
190 samples; n jx: 73**



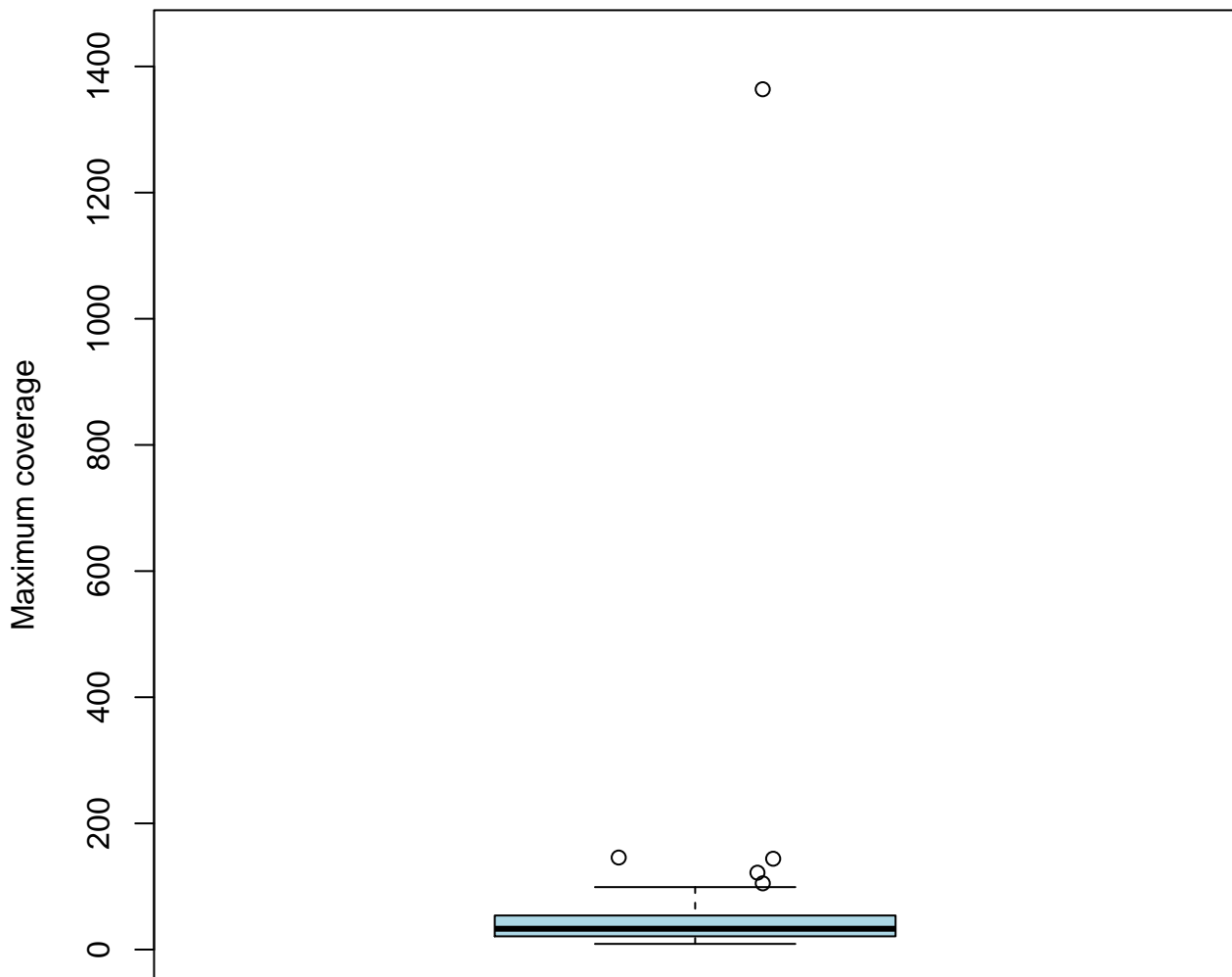
**jx in Intropolis but not "annotated" by knownGene:
191 samples; n jx: 60**



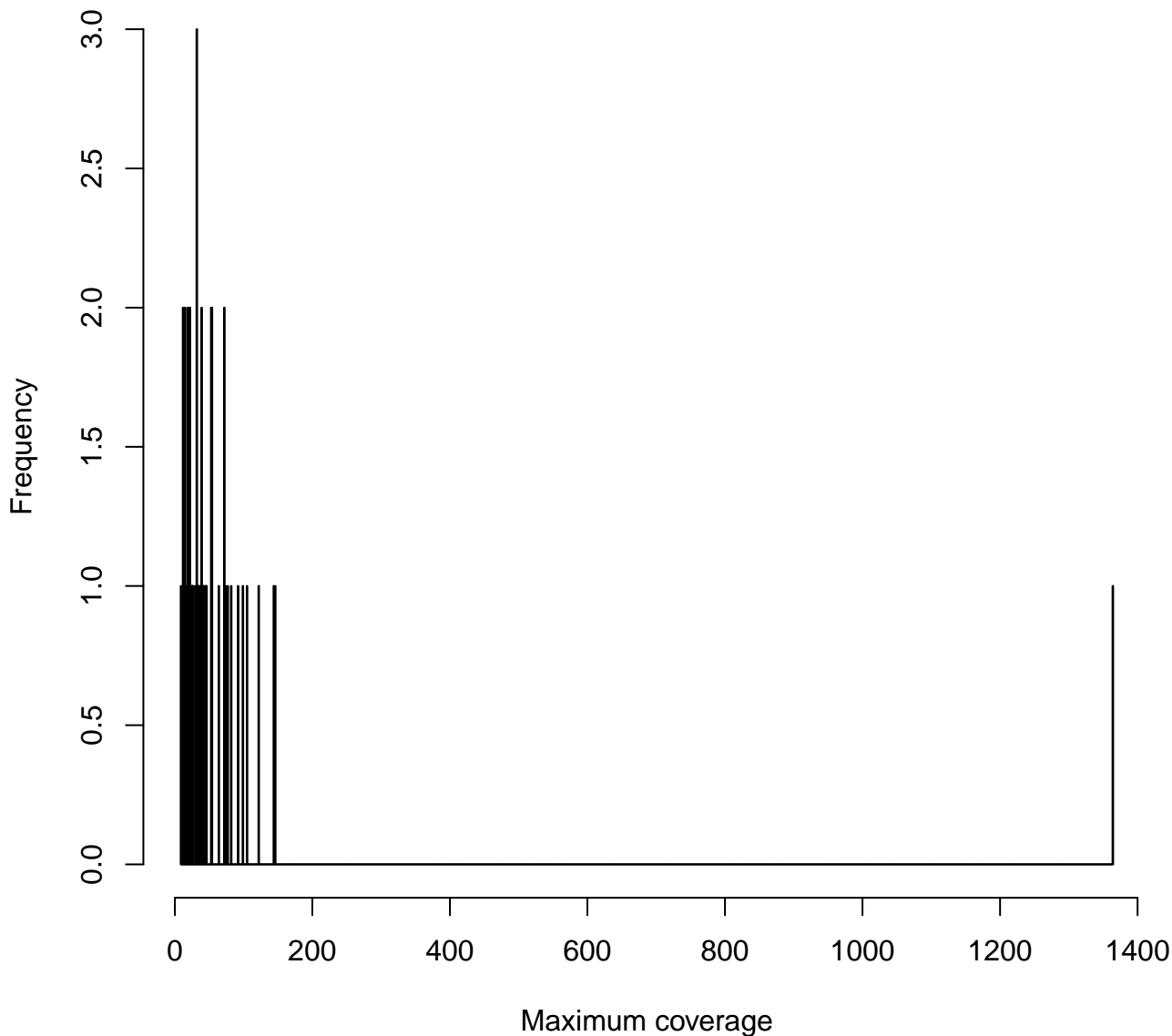
**jx in Intropolis but not "annotated" by knownGene:
191 samples; n jx: 60**



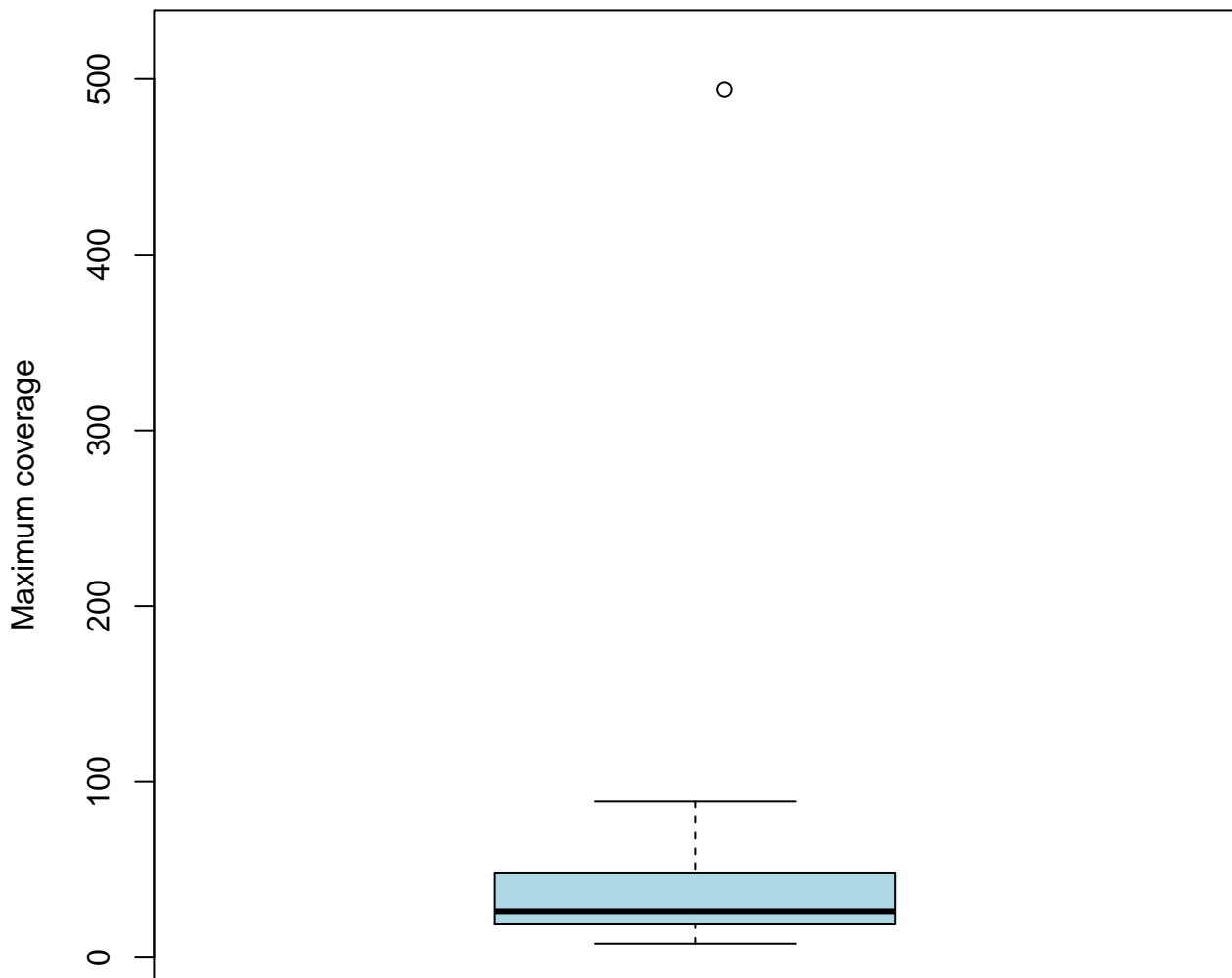
**jx in Intropolis but not "annotated" by knownGene:
192 samples; n jx: 57**



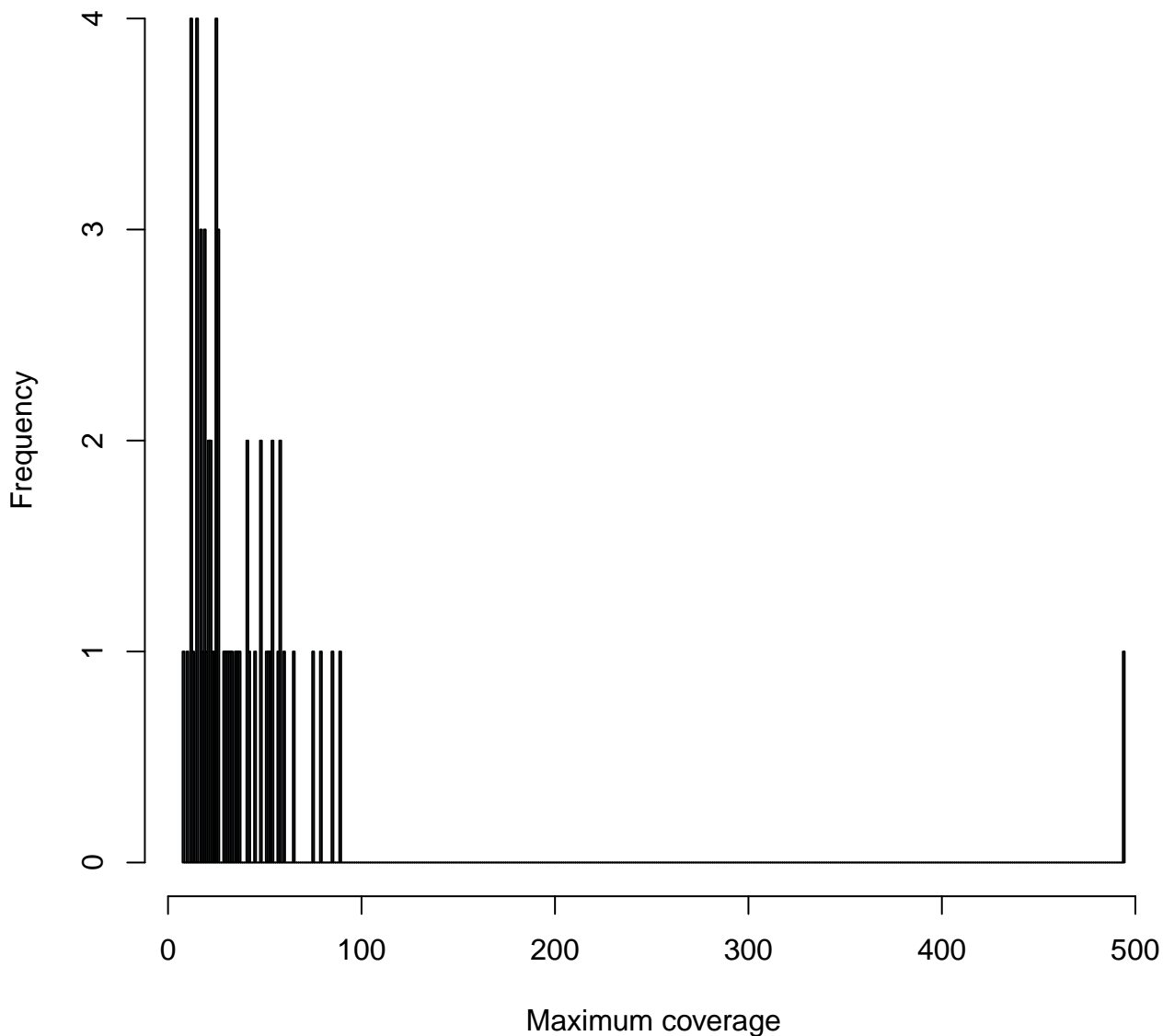
**jx in Intropolis but not "annotated" by knownGene:
192 samples; n jx: 57**



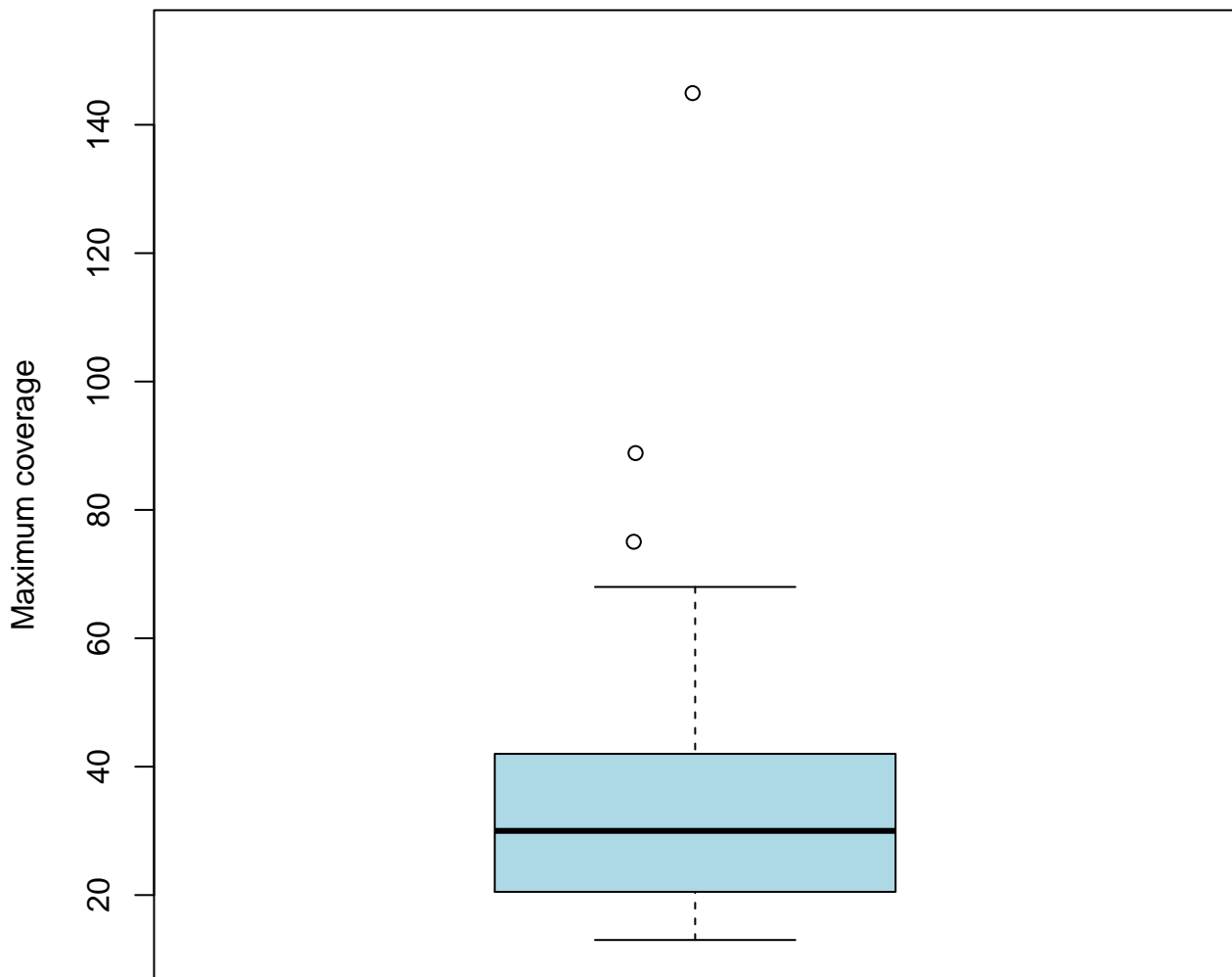
**jx in Intropolis but not "annotated" by knownGene:
193 samples; n jx: 61**



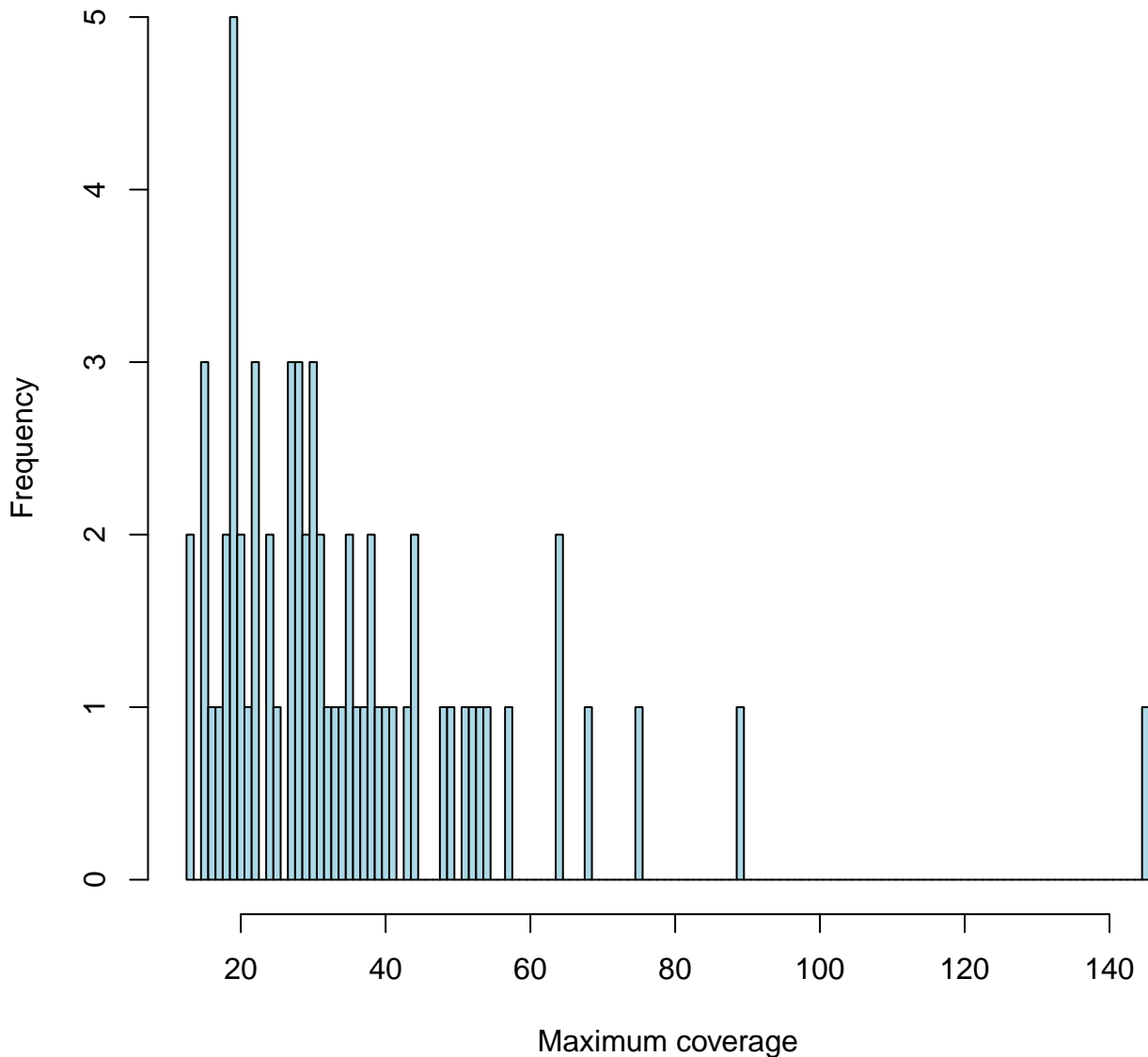
**jx in Intropolis but not "annotated" by knownGene:
193 samples; n jx: 61**



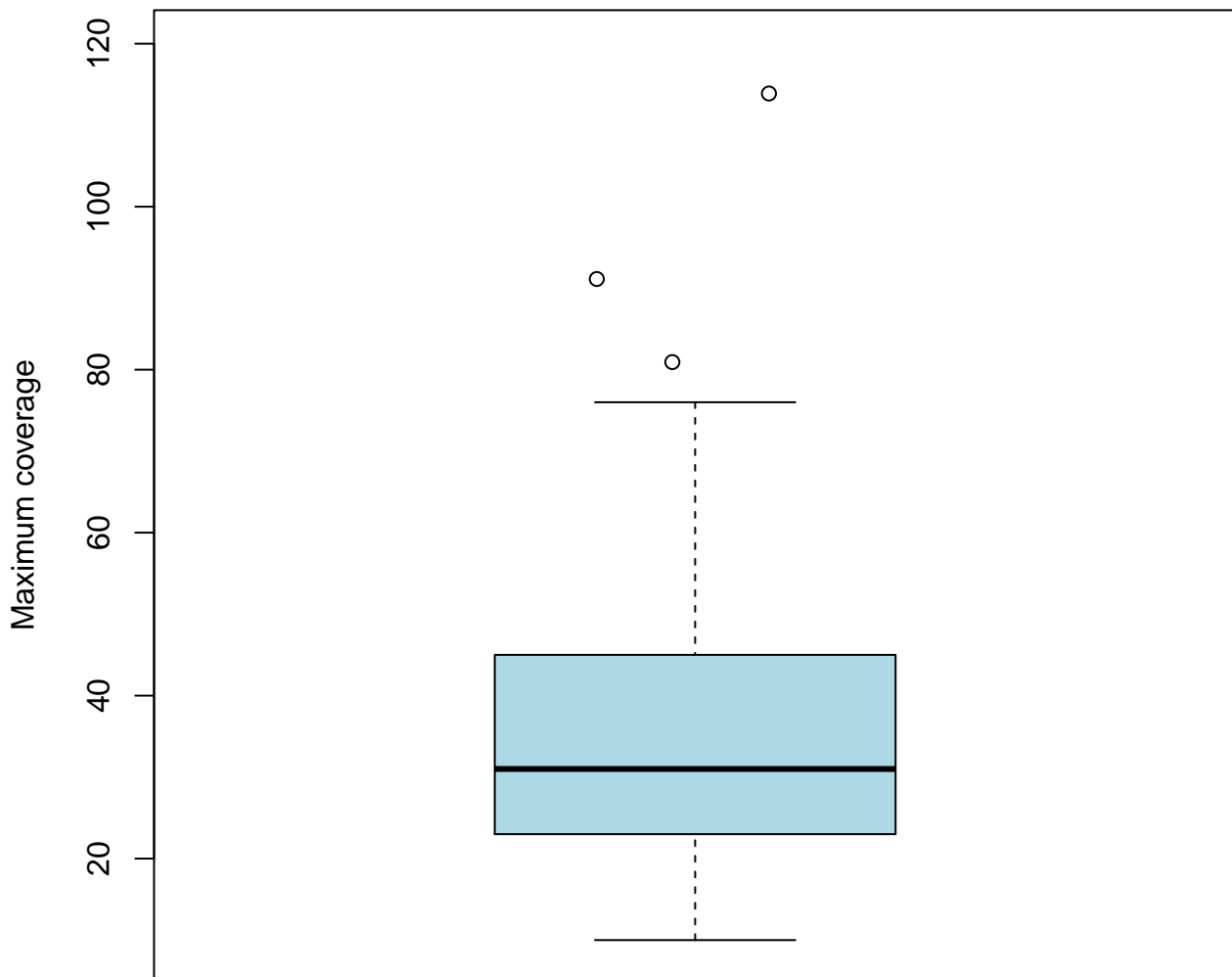
**jx in Intropolis but not "annotated" by knownGene:
194 samples; n jx: 64**



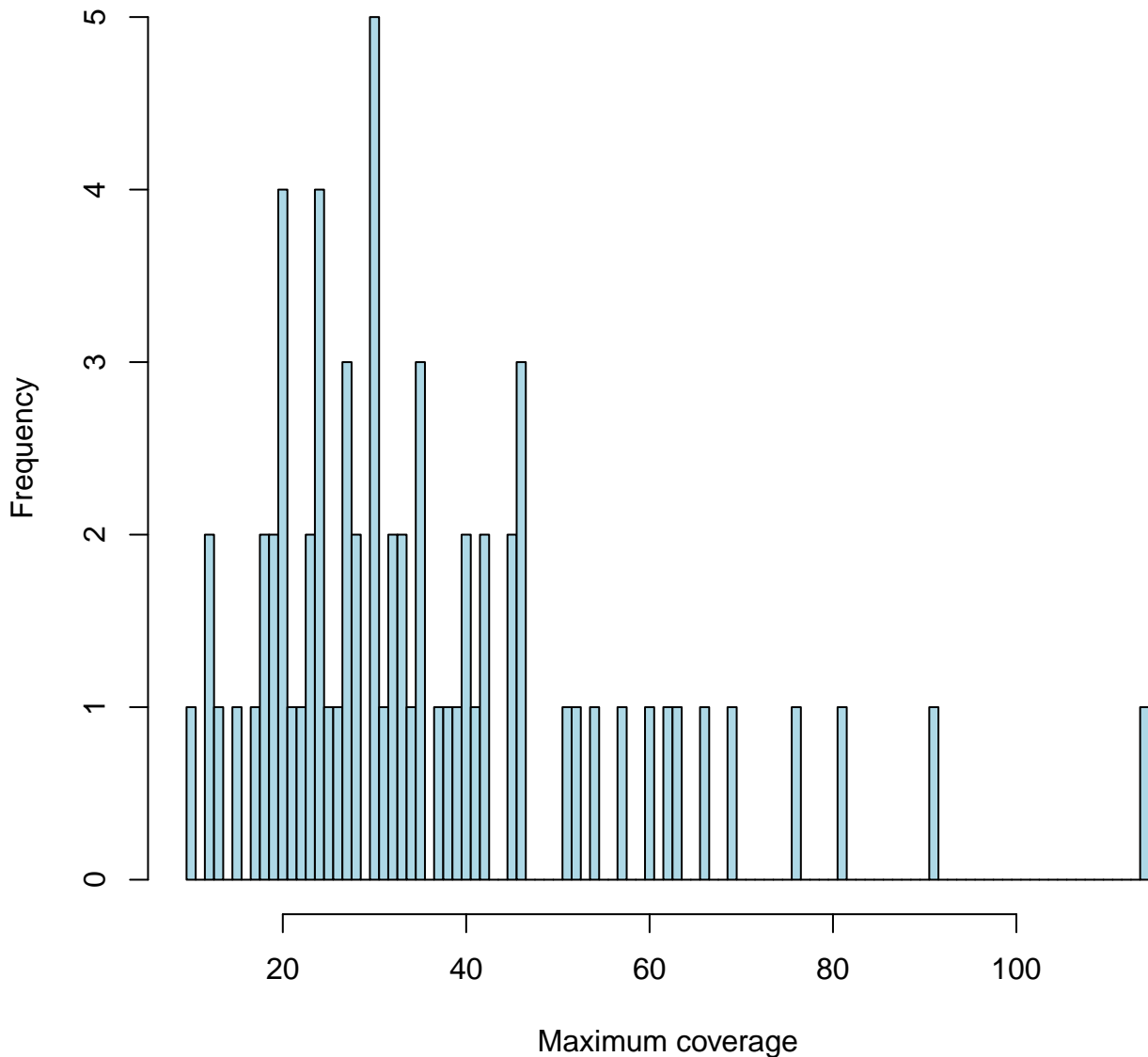
**jx in Intropolis but not "annotated" by knownGene:
194 samples; n jx: 64**



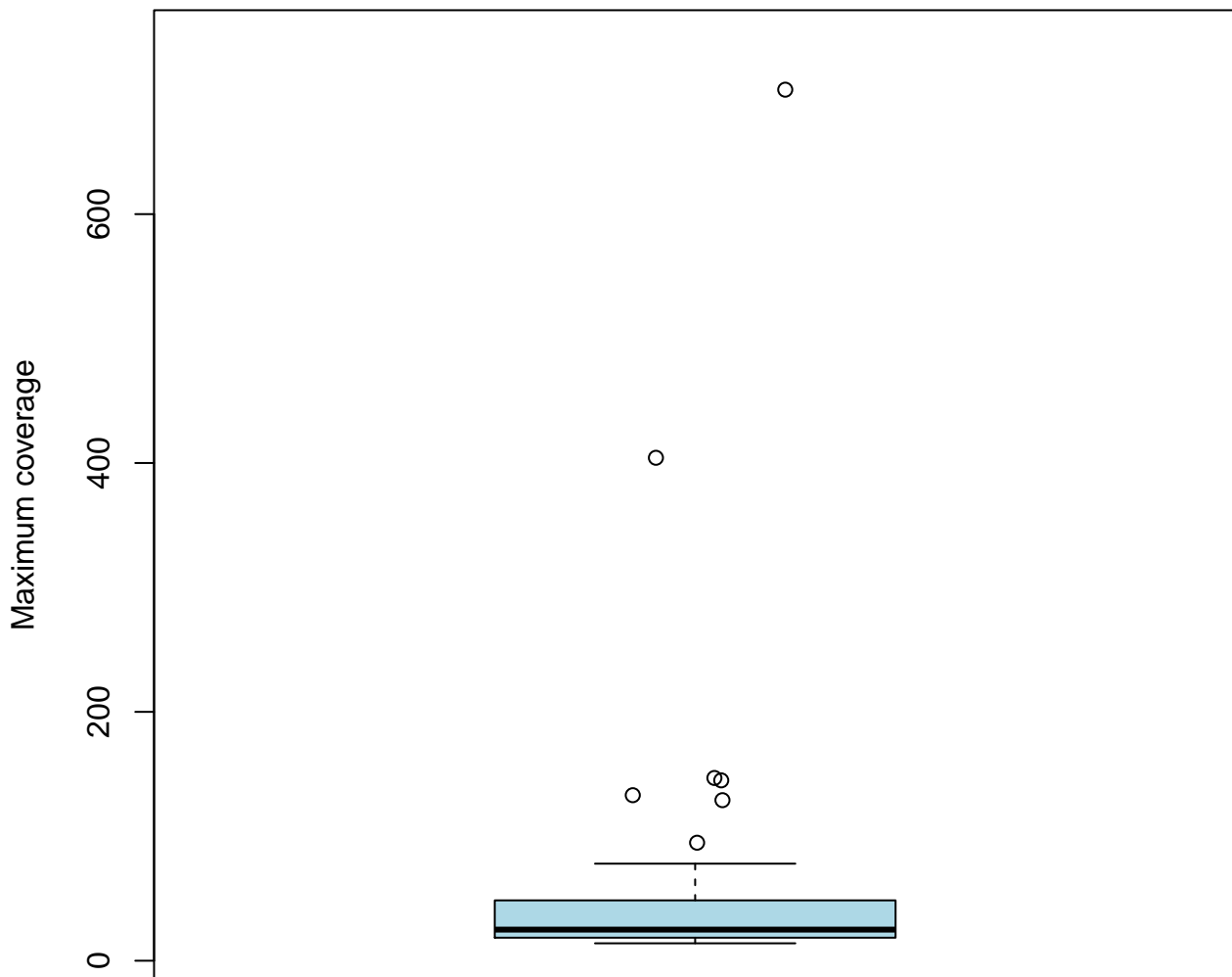
**jx in Intropolis but not "annotated" by knownGene:
195 samples; n jx: 69**



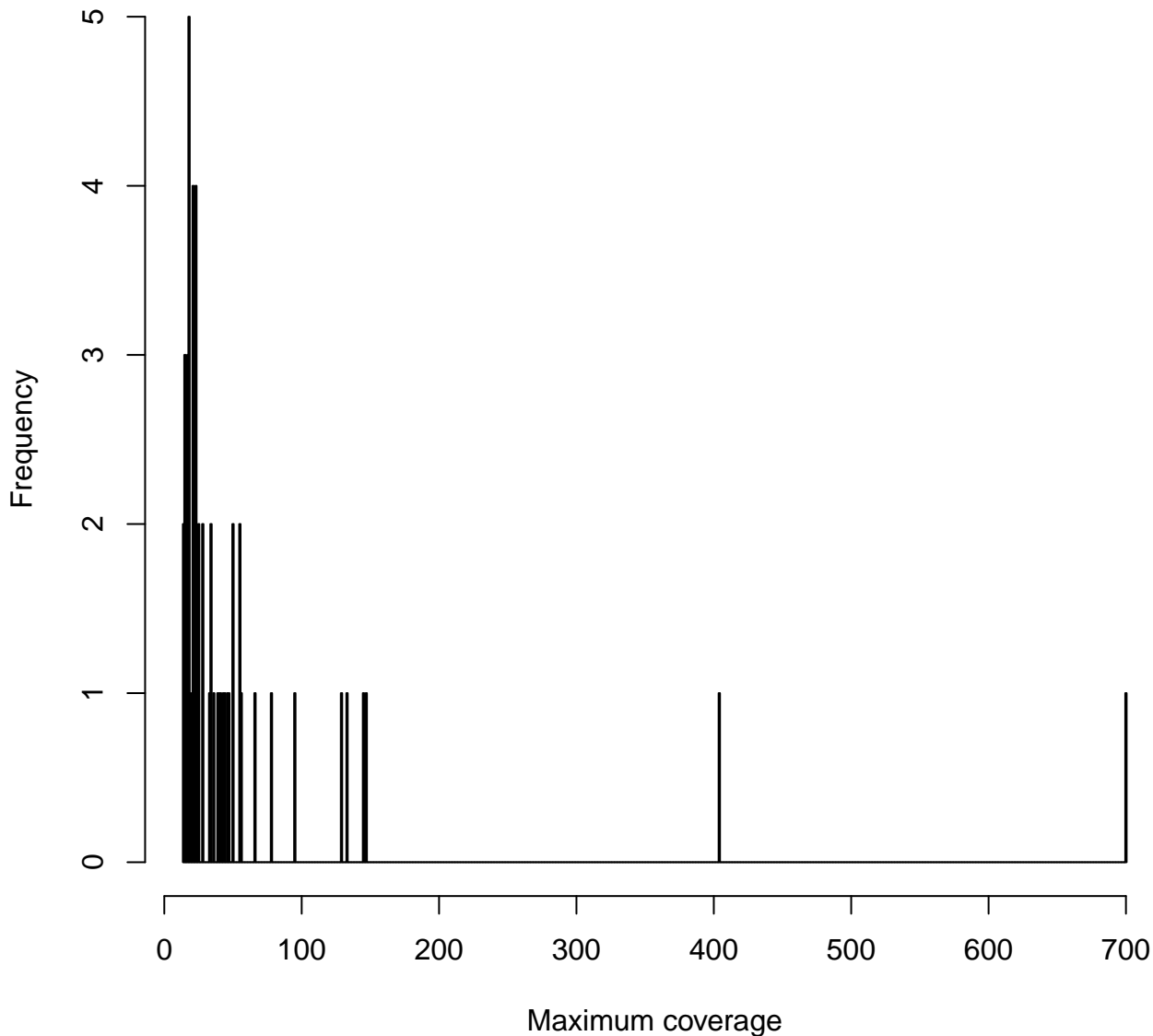
**jx in Intropolis but not "annotated" by knownGene:
195 samples; n jx: 69**



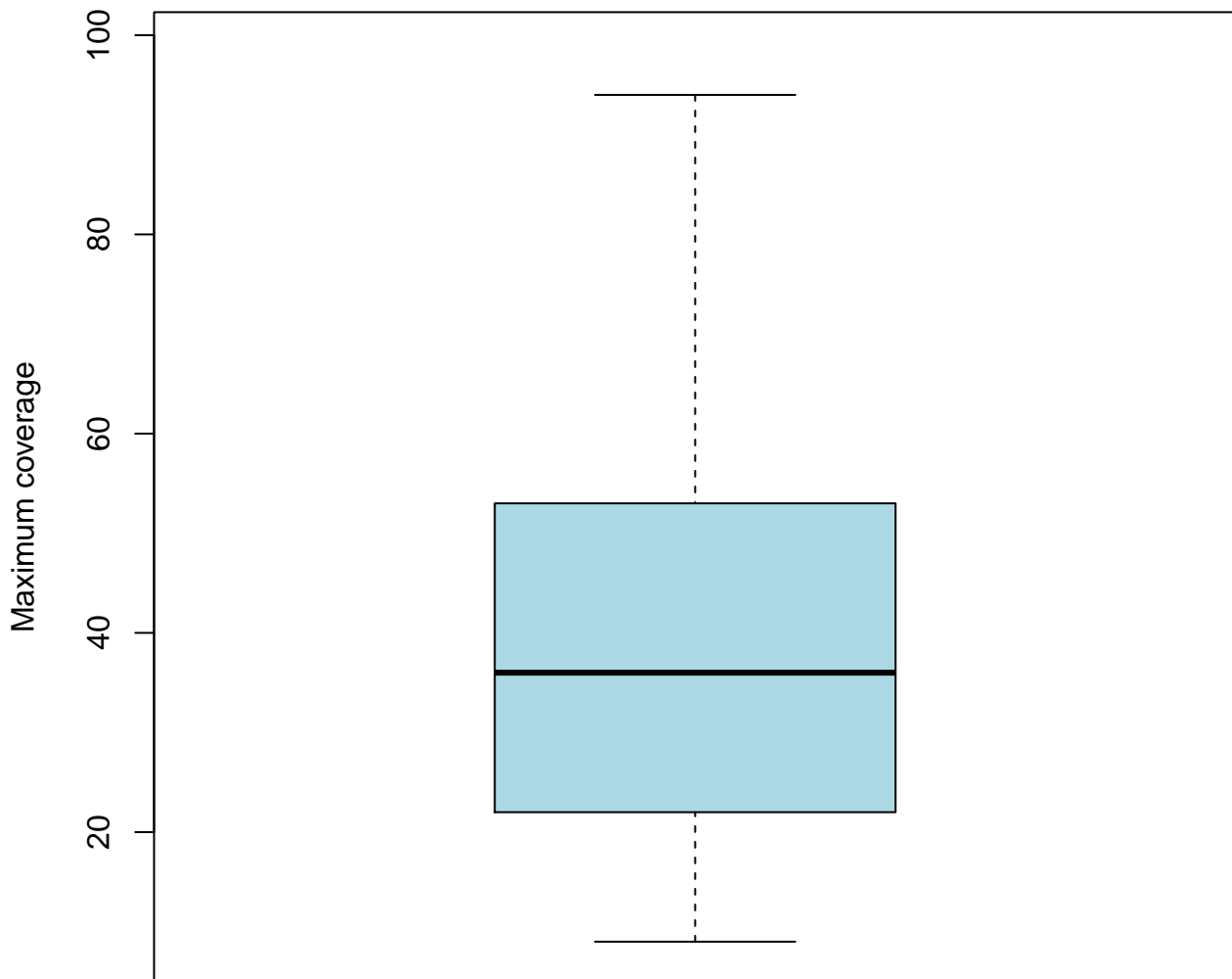
**jx in Intropolis but not "annotated" by knownGene:
196 samples; n jx: 56**



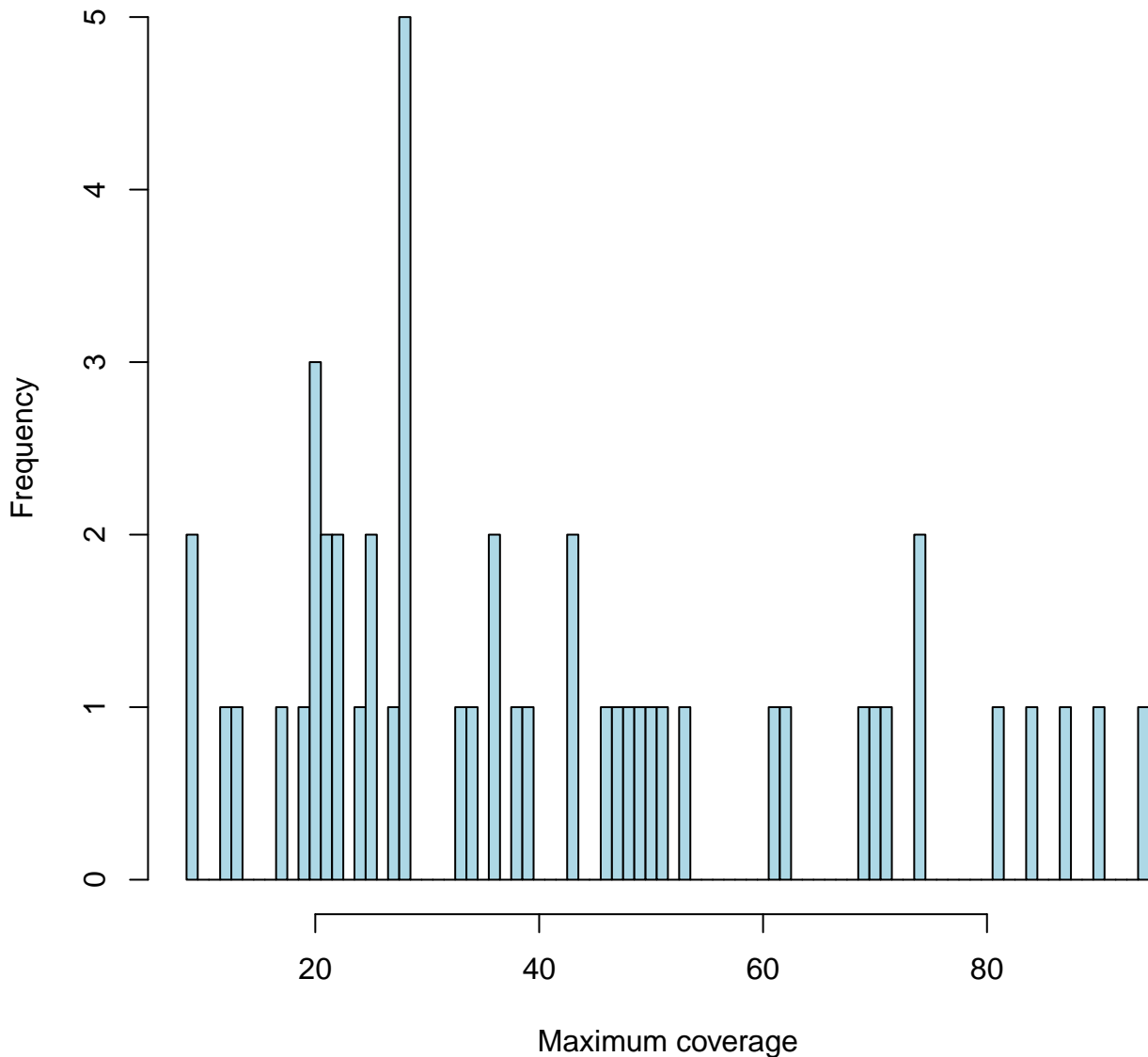
**jx in Intropolis but not "annotated" by knownGene:
196 samples; n jx: 56**



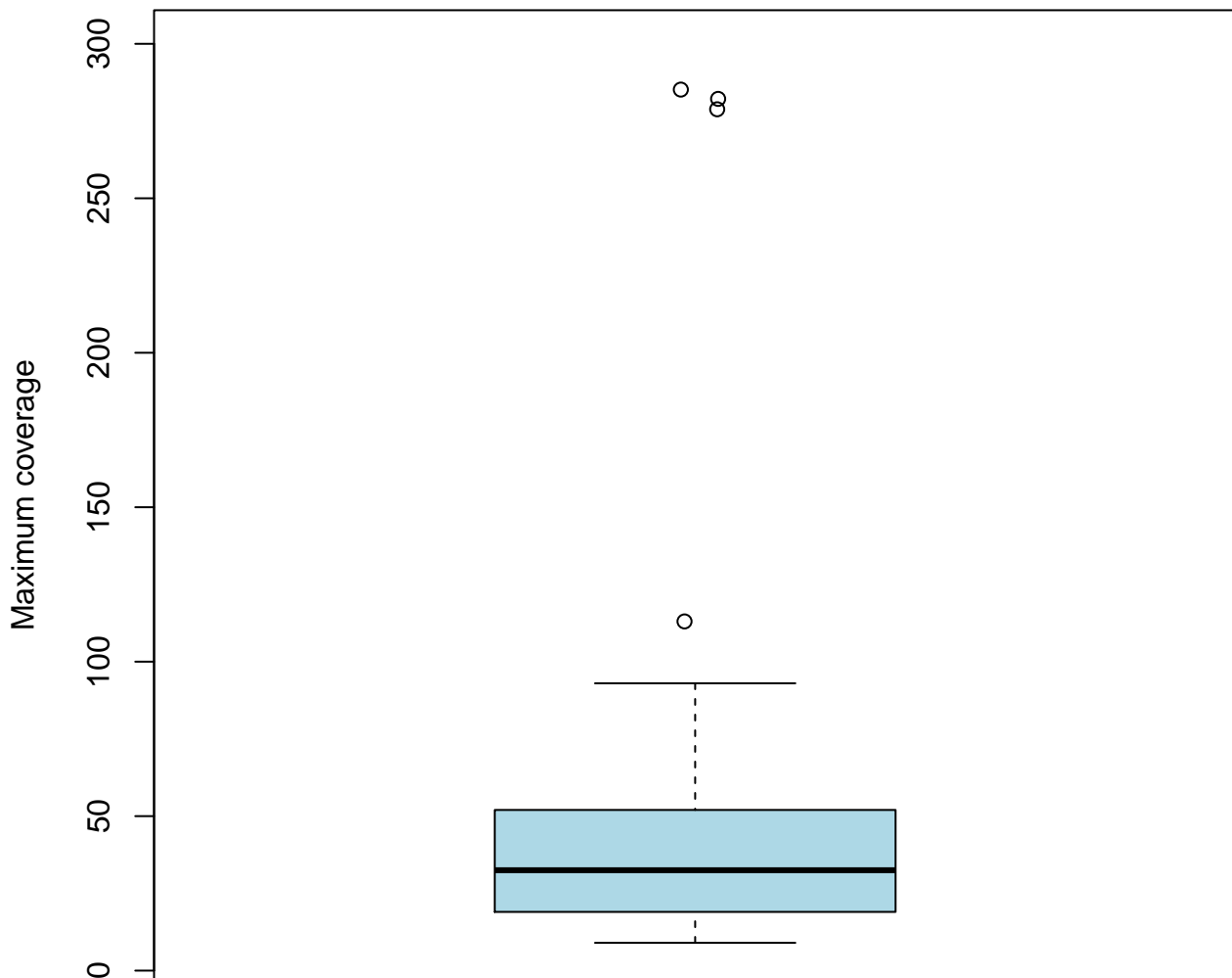
**jx in Intropolis but not "annotated" by knownGene:
197 samples; n jx: 49**



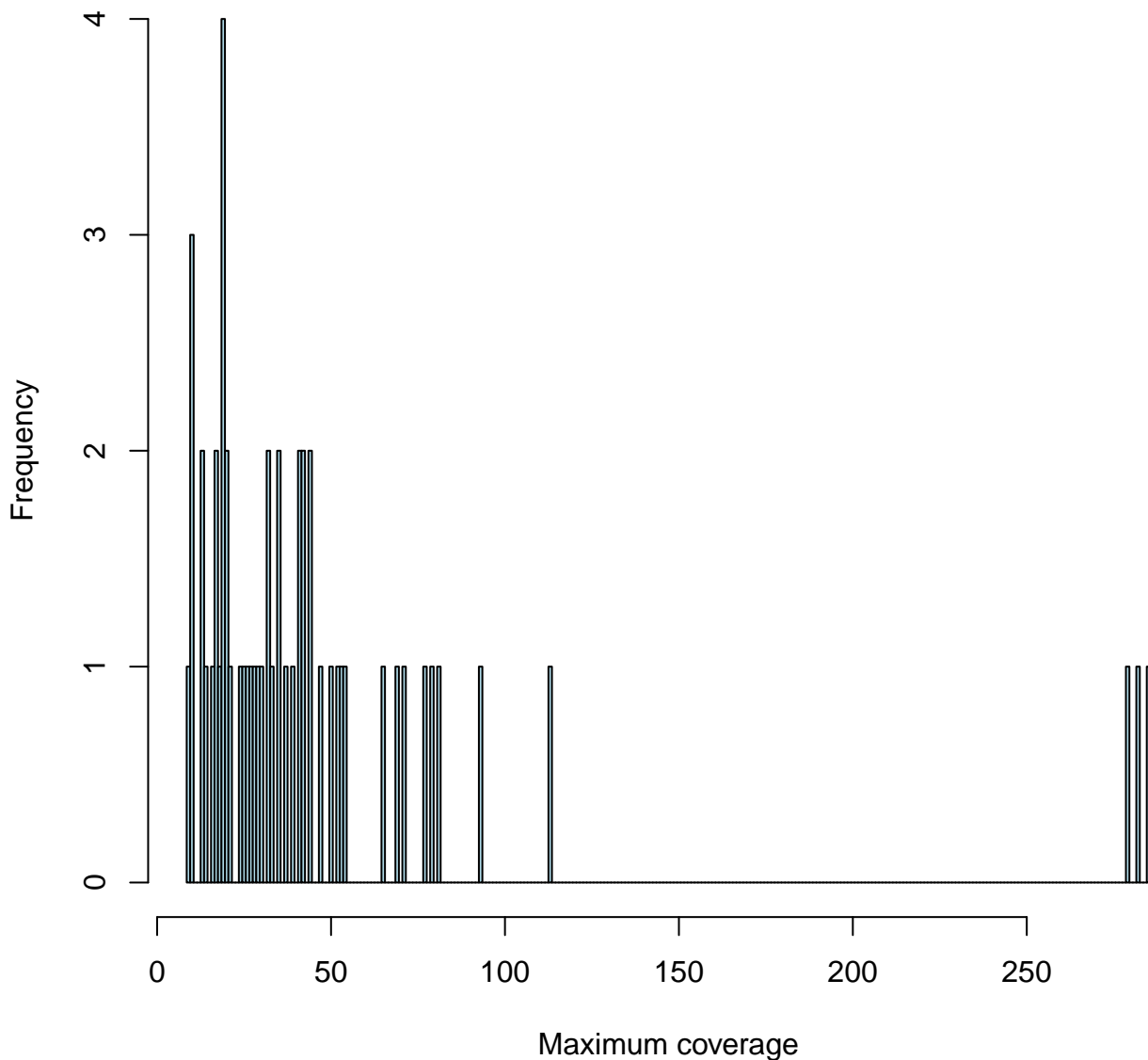
**jx in Intropolis but not "annotated" by knownGene:
197 samples; n jx: 49**



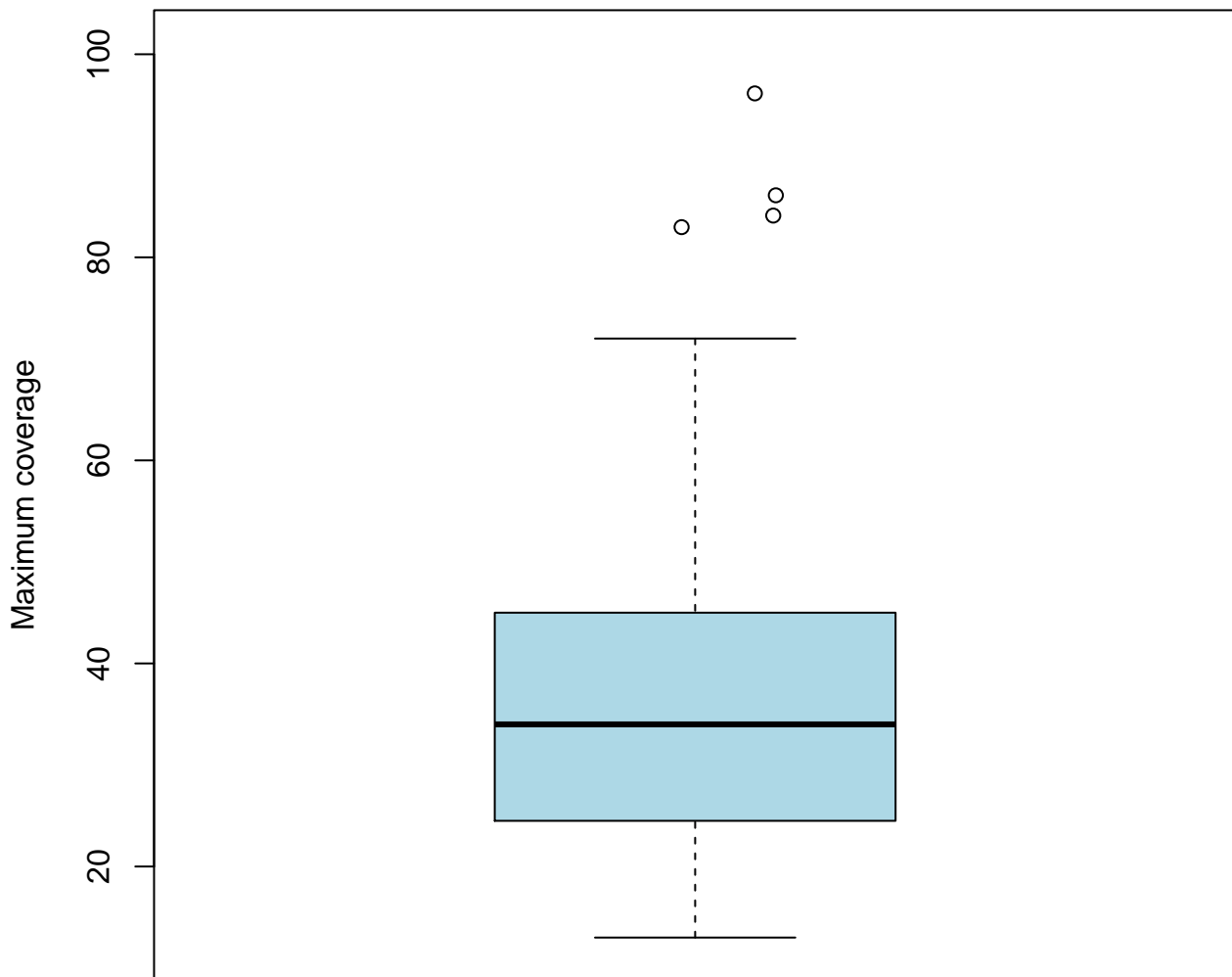
**jx in Intropolis but not "annotated" by knownGene:
198 samples; n jx: 54**



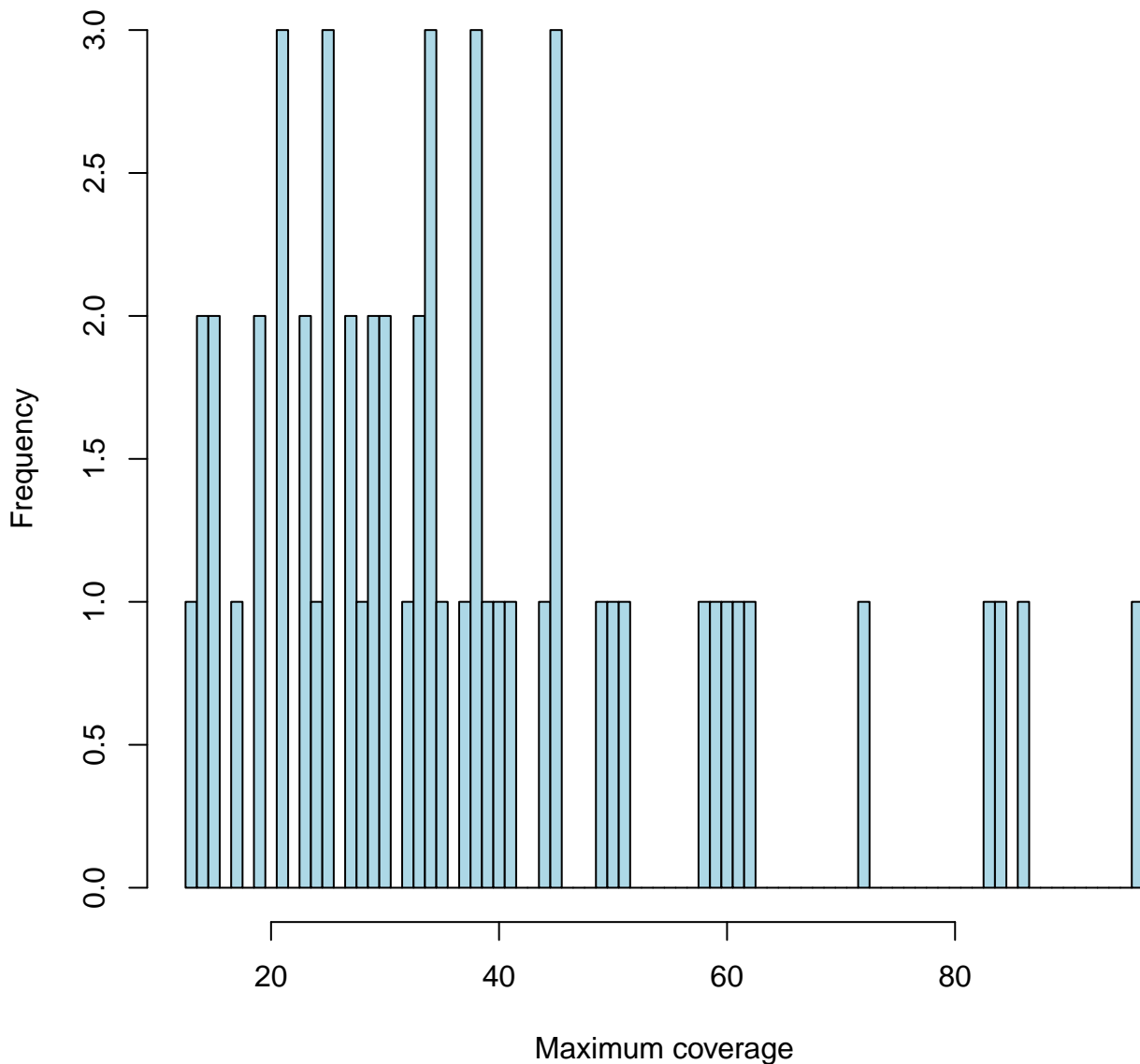
**jx in Intropolis but not "annotated" by knownGene:
198 samples; n jx: 54**



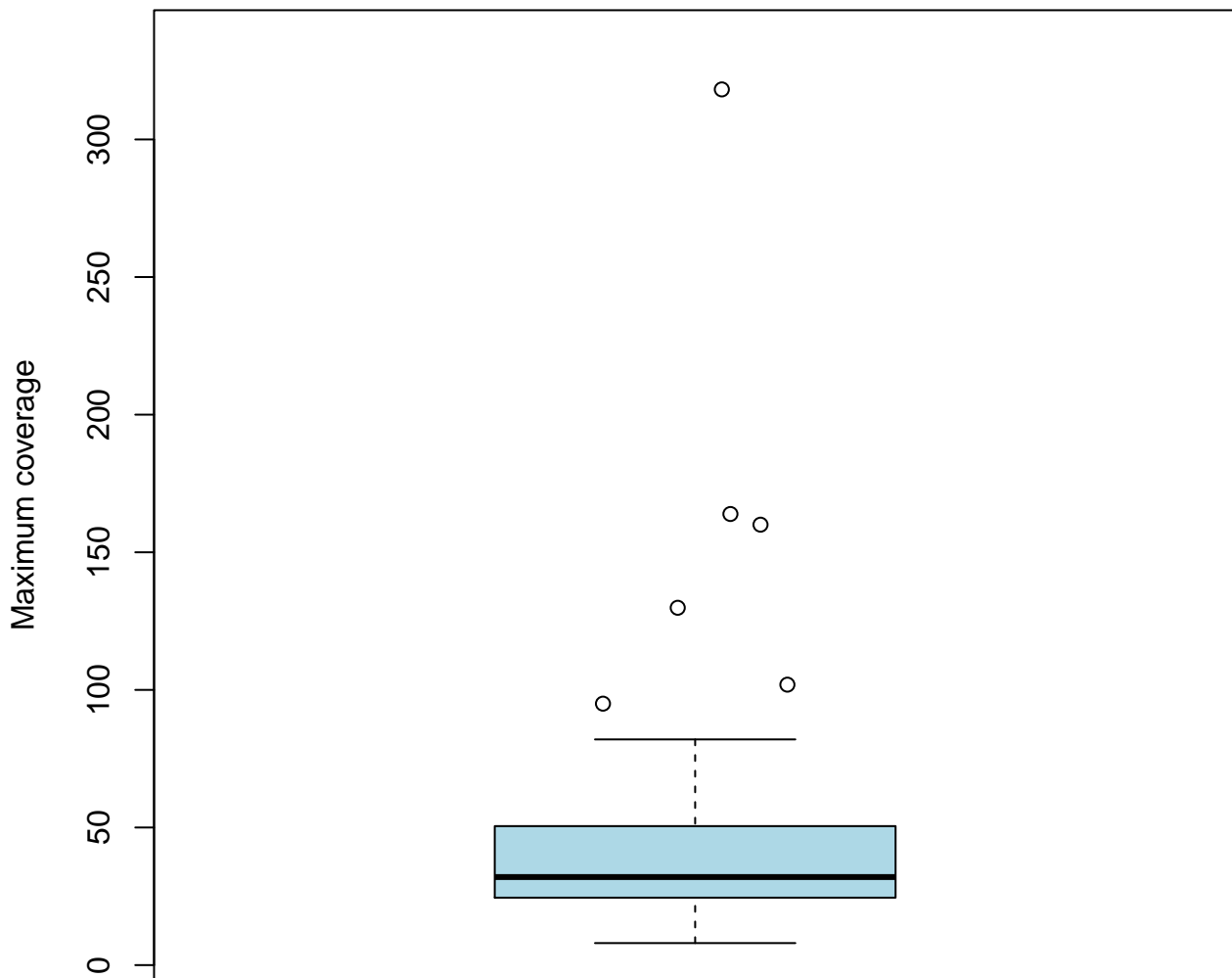
**jx in Intropolis but not "annotated" by knownGene:
199 samples; n jx: 55**



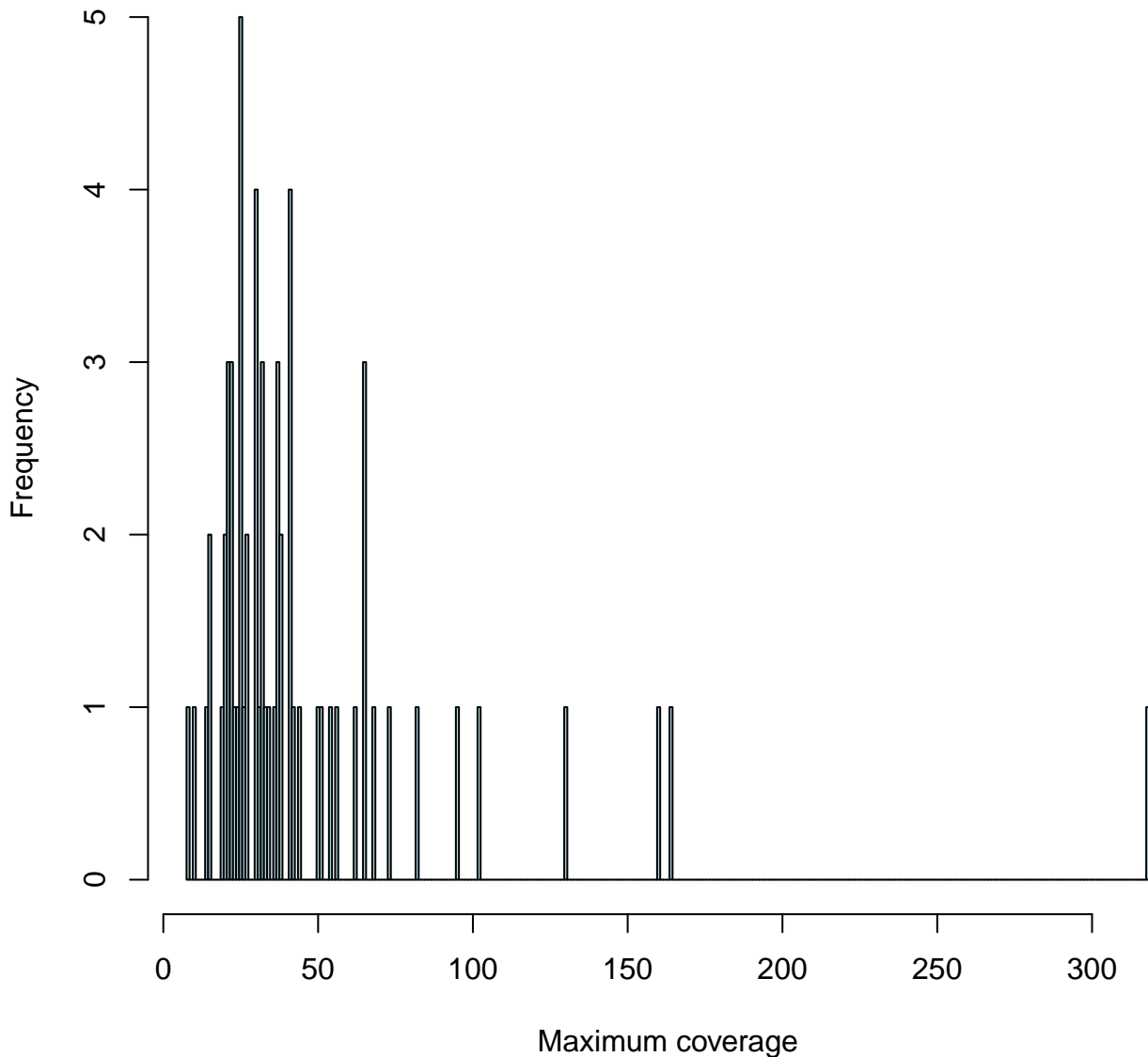
**jx in Intropolis but not "annotated" by knownGene:
199 samples; n jx: 55**



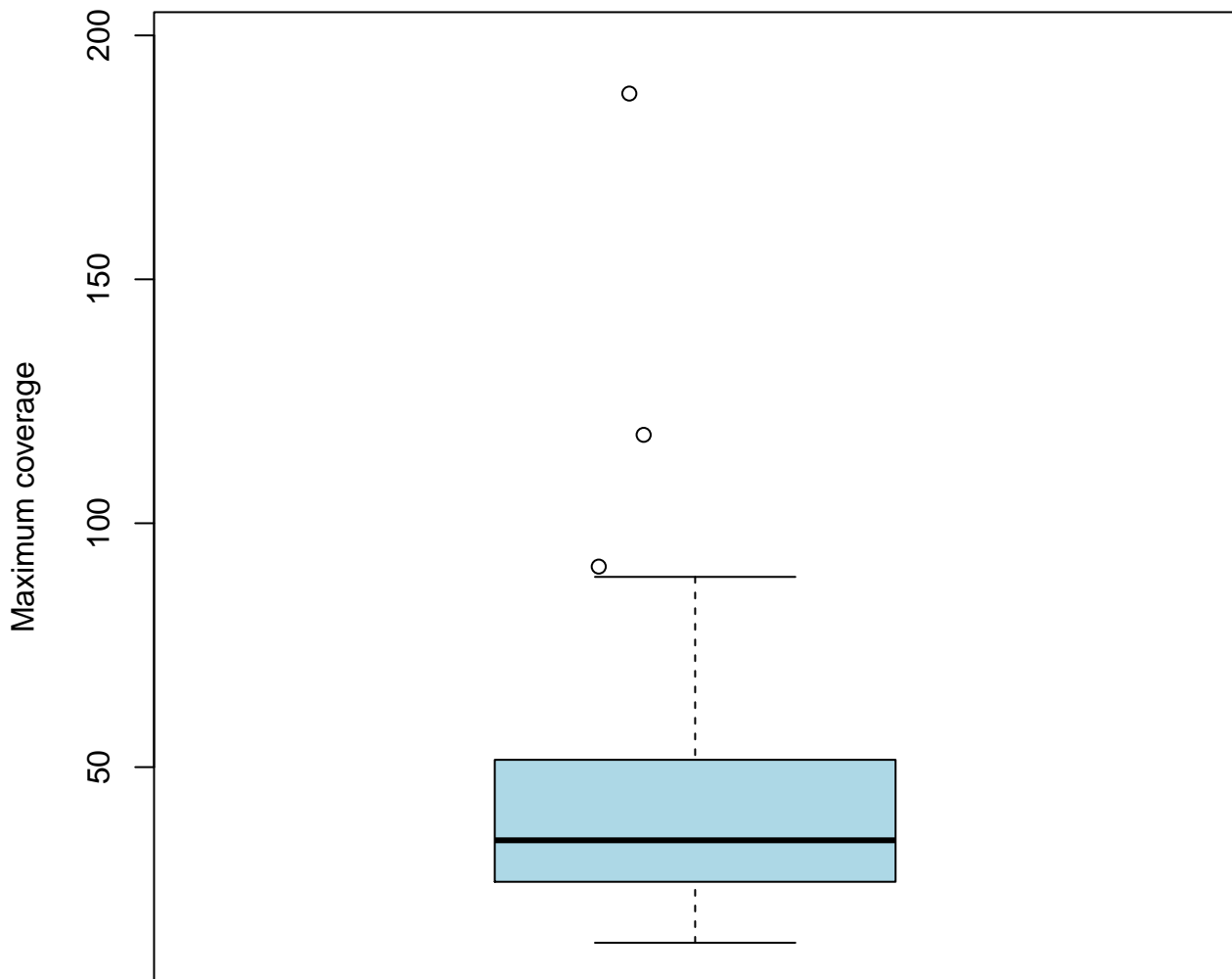
**jx in Intropolis but not "annotated" by knownGene:
200 samples; n jx: 63**



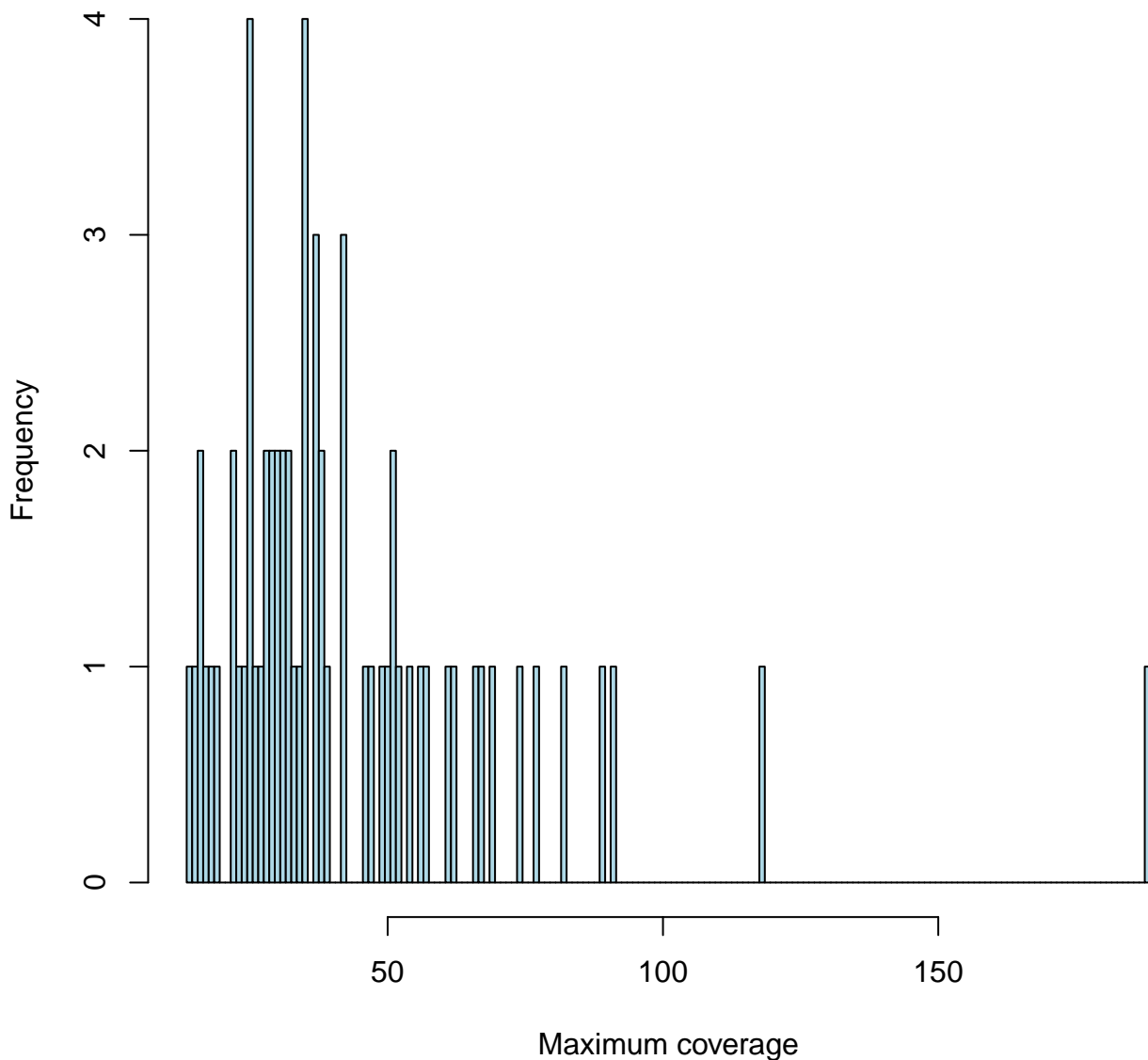
**jx in Intropolis but not "annotated" by knownGene:
200 samples; n jx: 63**



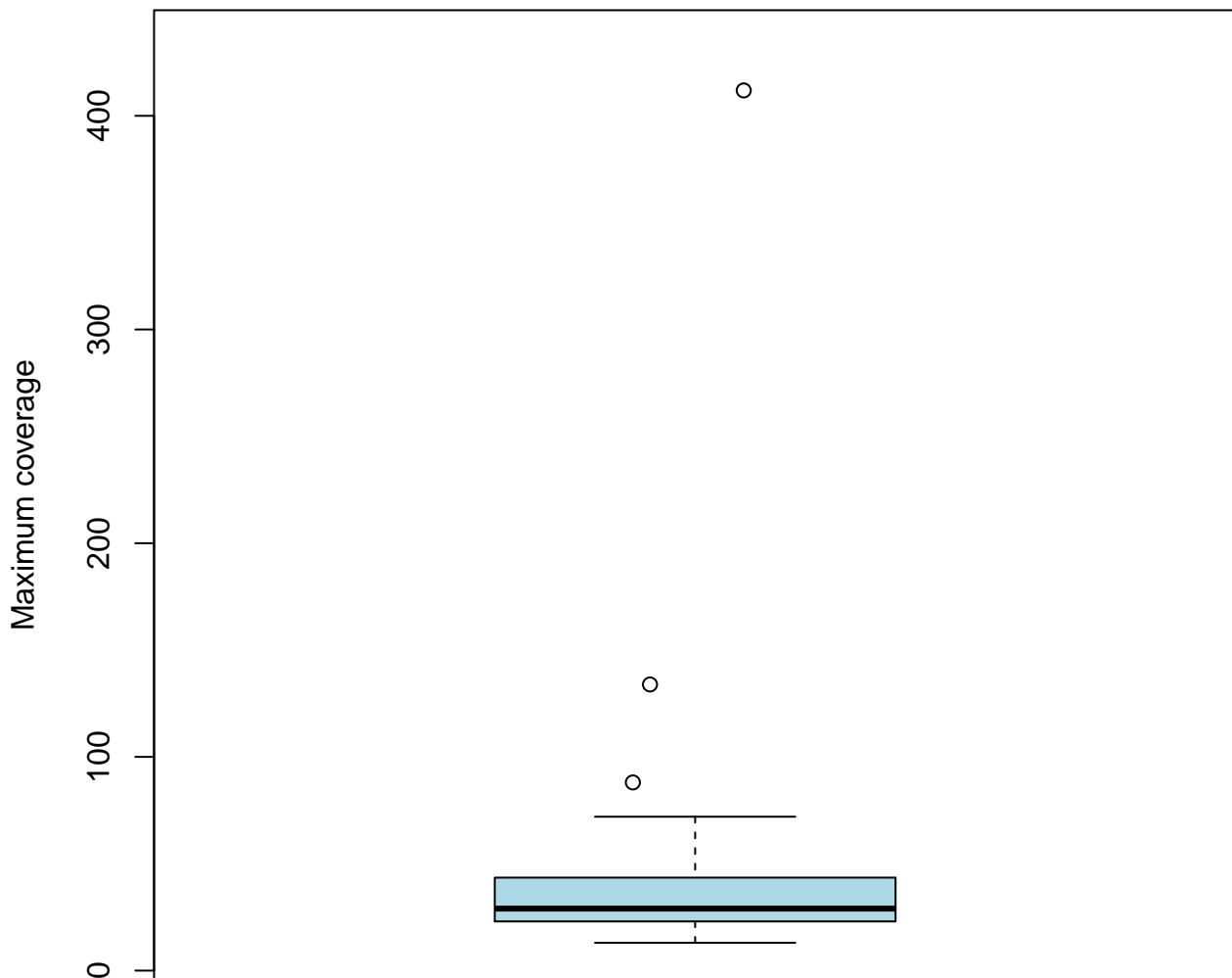
**jx in Intropolis but not "annotated" by knownGene:
201 samples; n jx: 64**



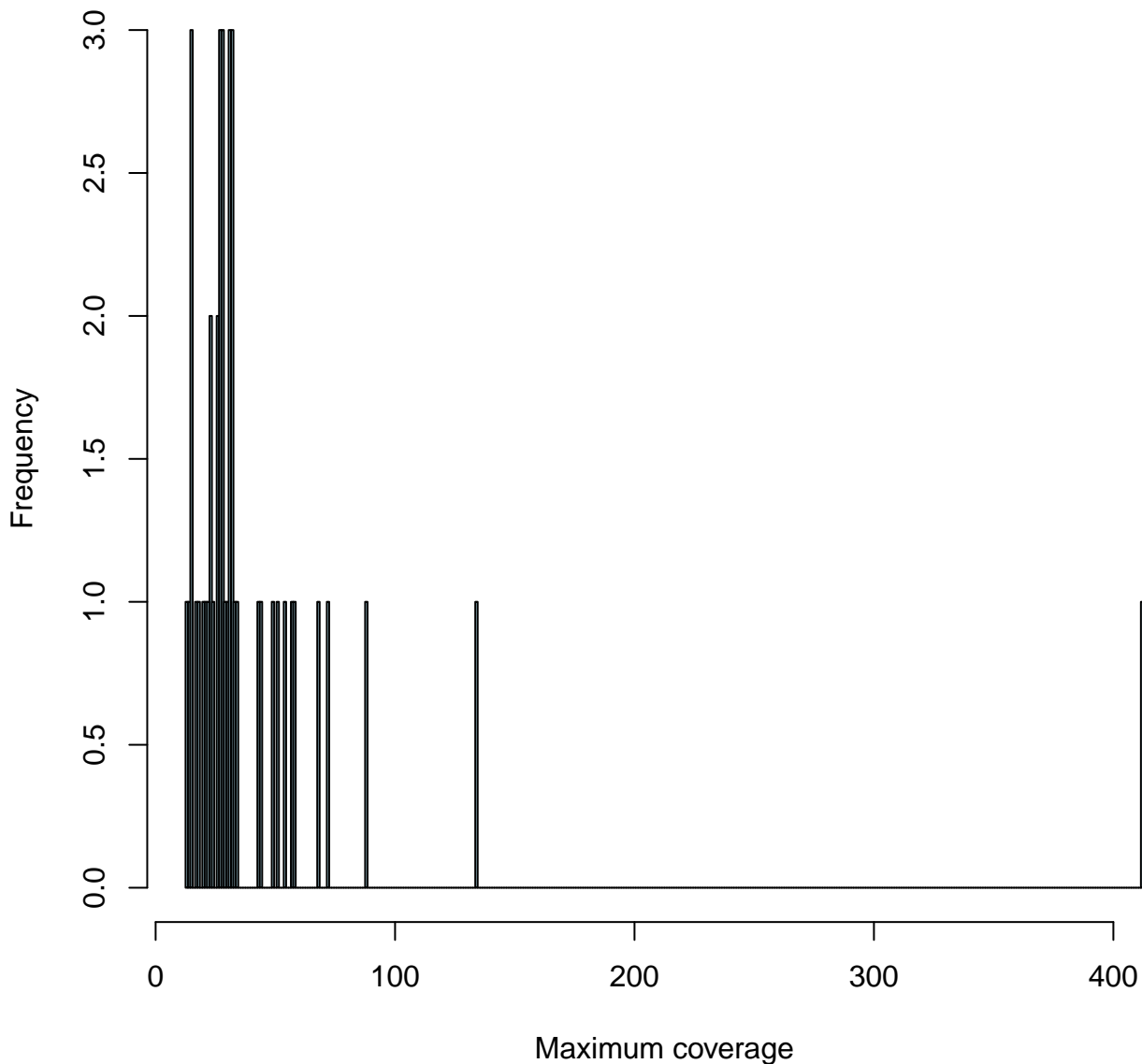
**jx in Intropolis but not "annotated" by knownGene:
201 samples; n jx: 64**



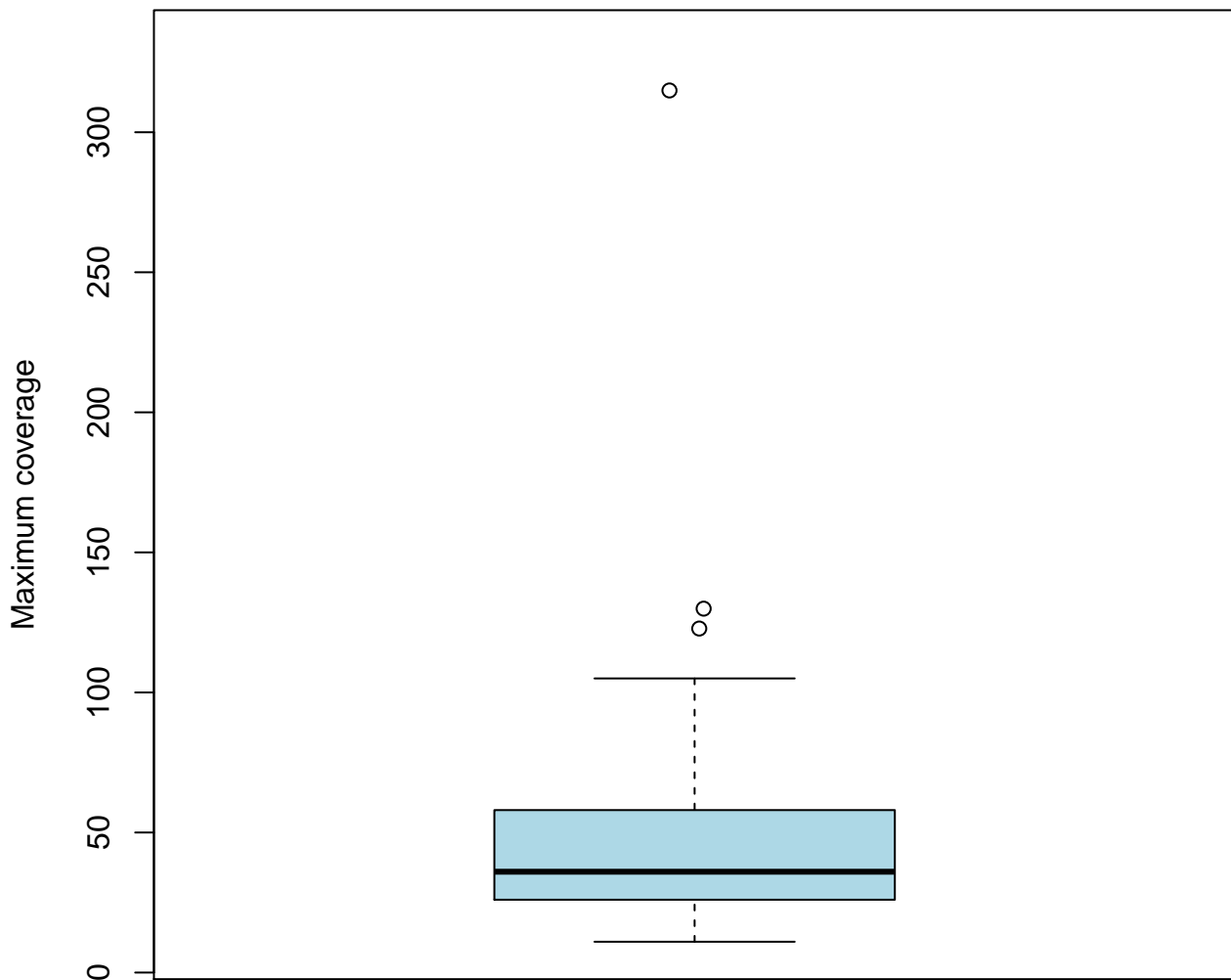
**jx in Intropolis but not "annotated" by knownGene:
202 samples; n jx: 43**



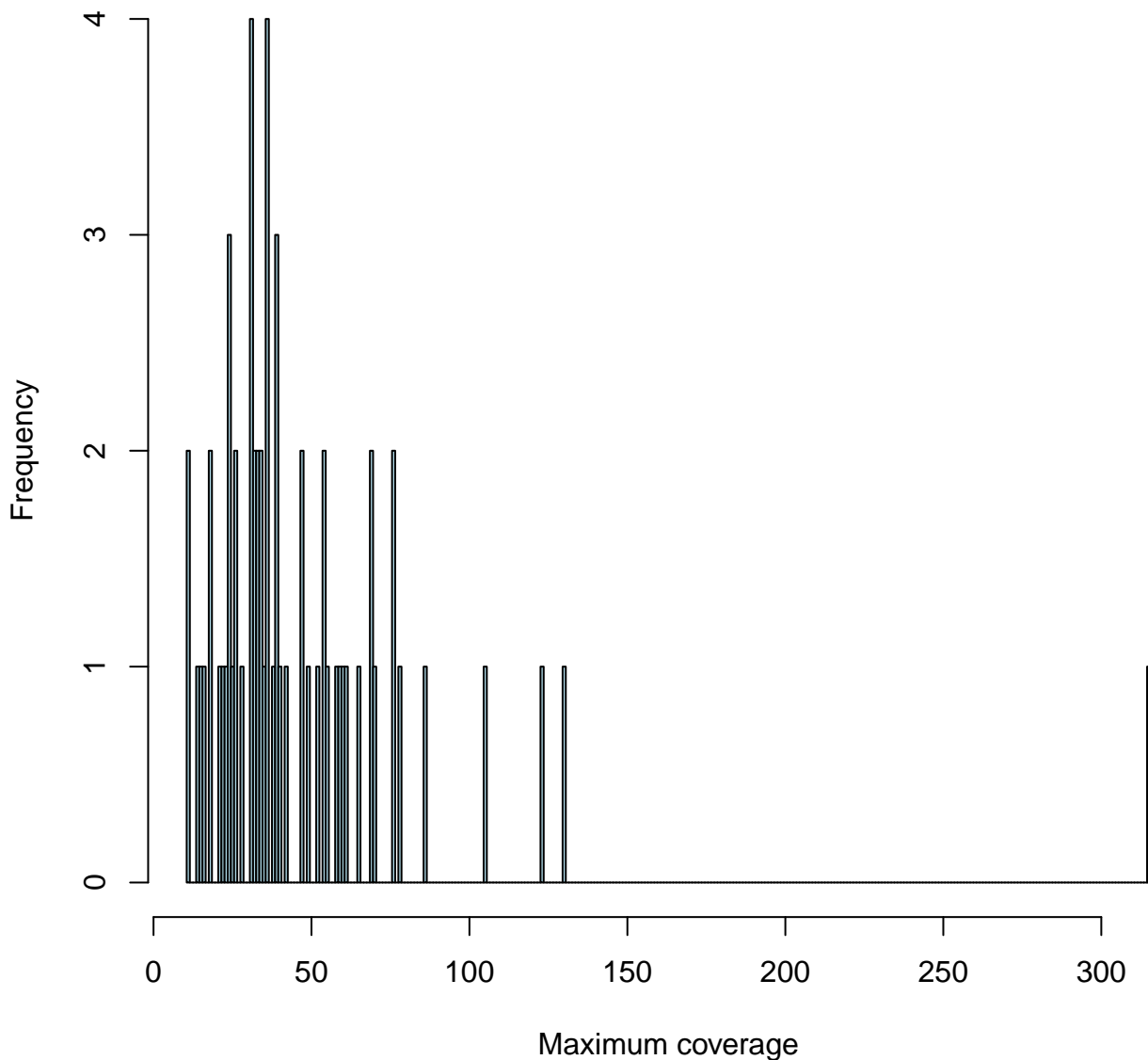
**jx in Intropolis but not "annotated" by knownGene:
202 samples; n jx: 43**



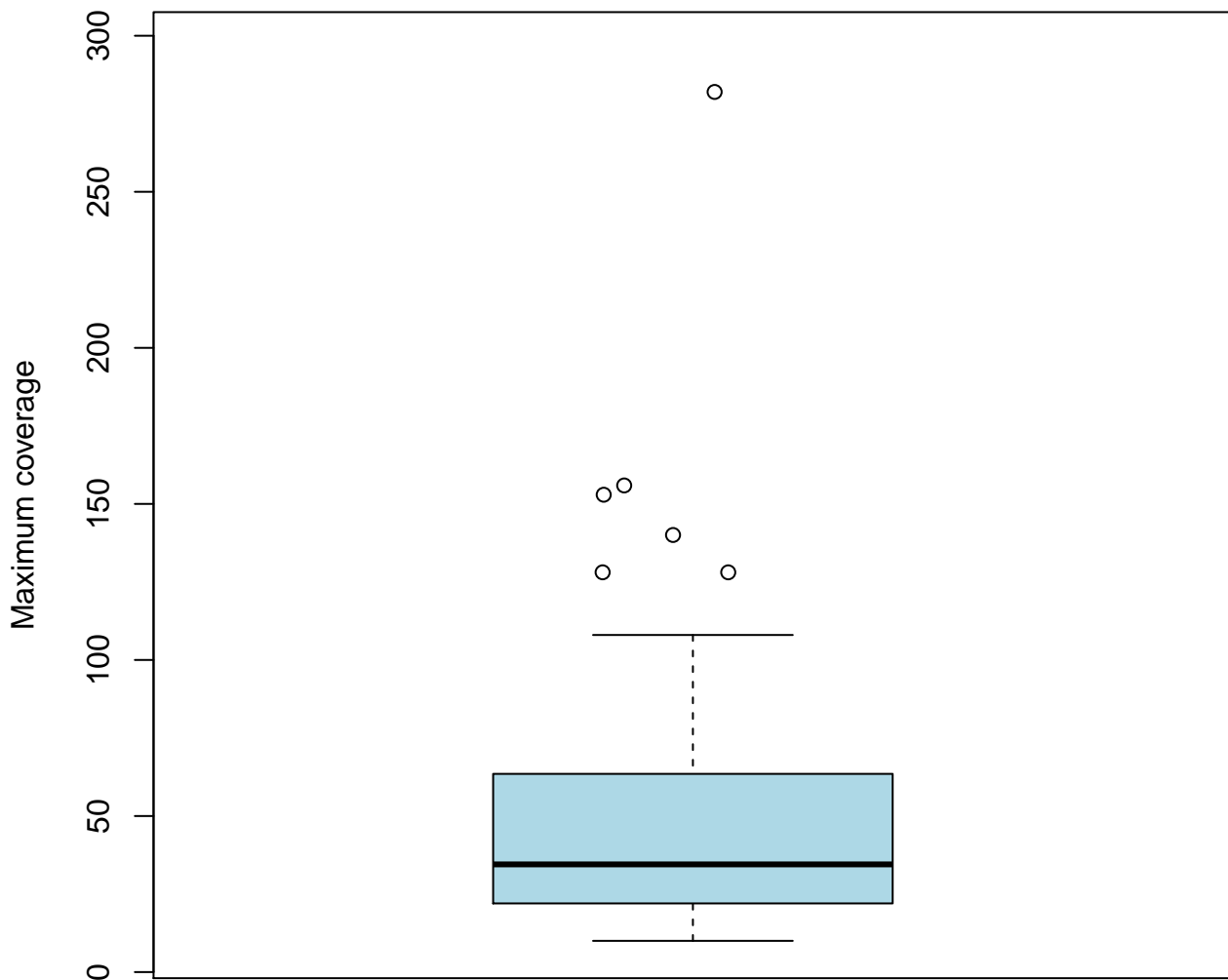
**jx in Intropolis but not "annotated" by knownGene:
203 samples; n jx: 61**



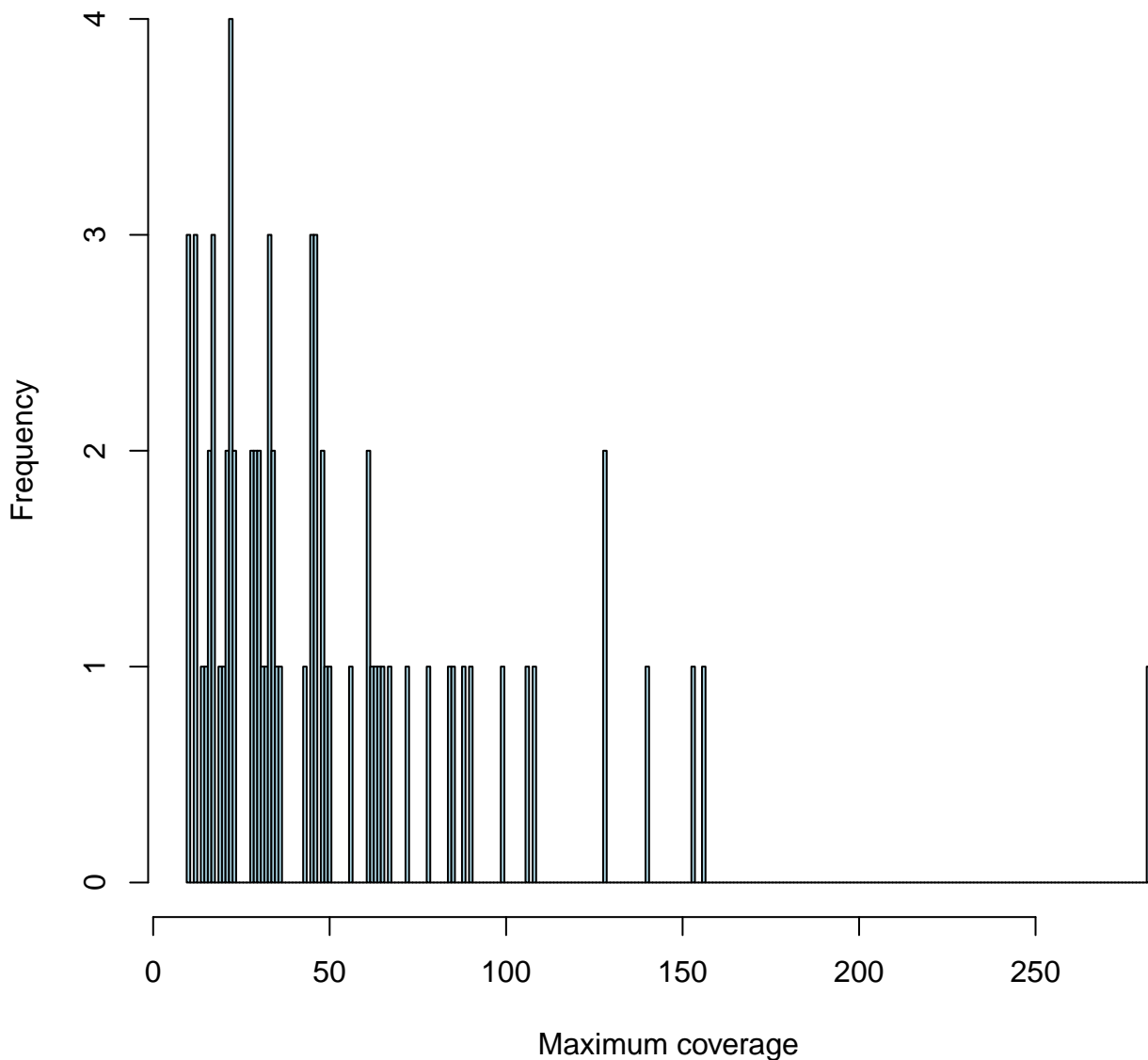
**jx in Intropolis but not "annotated" by knownGene:
203 samples; n jx: 61**



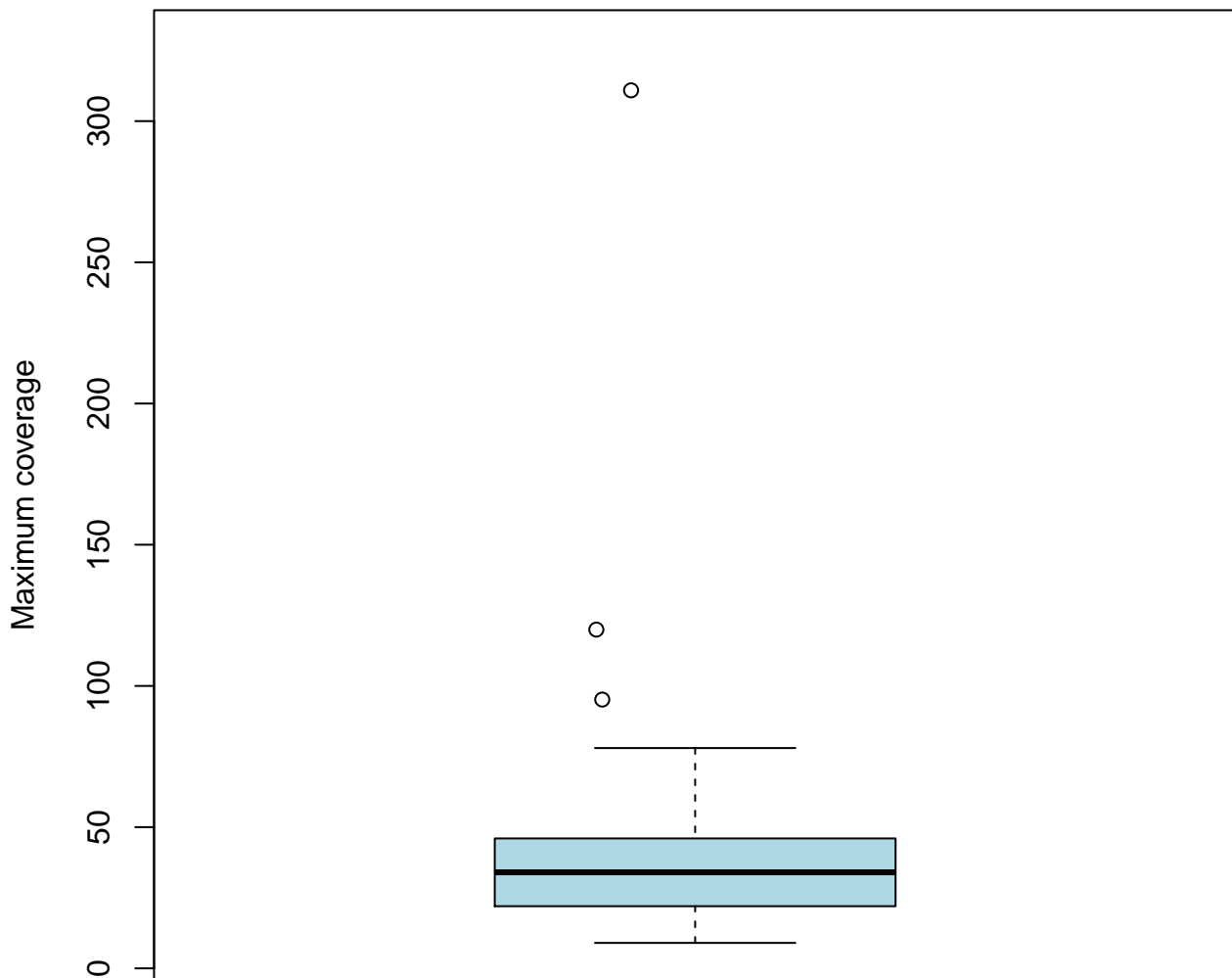
**jx in Intropolis but not "annotated" by knownGene:
204 samples; n jx: 72**



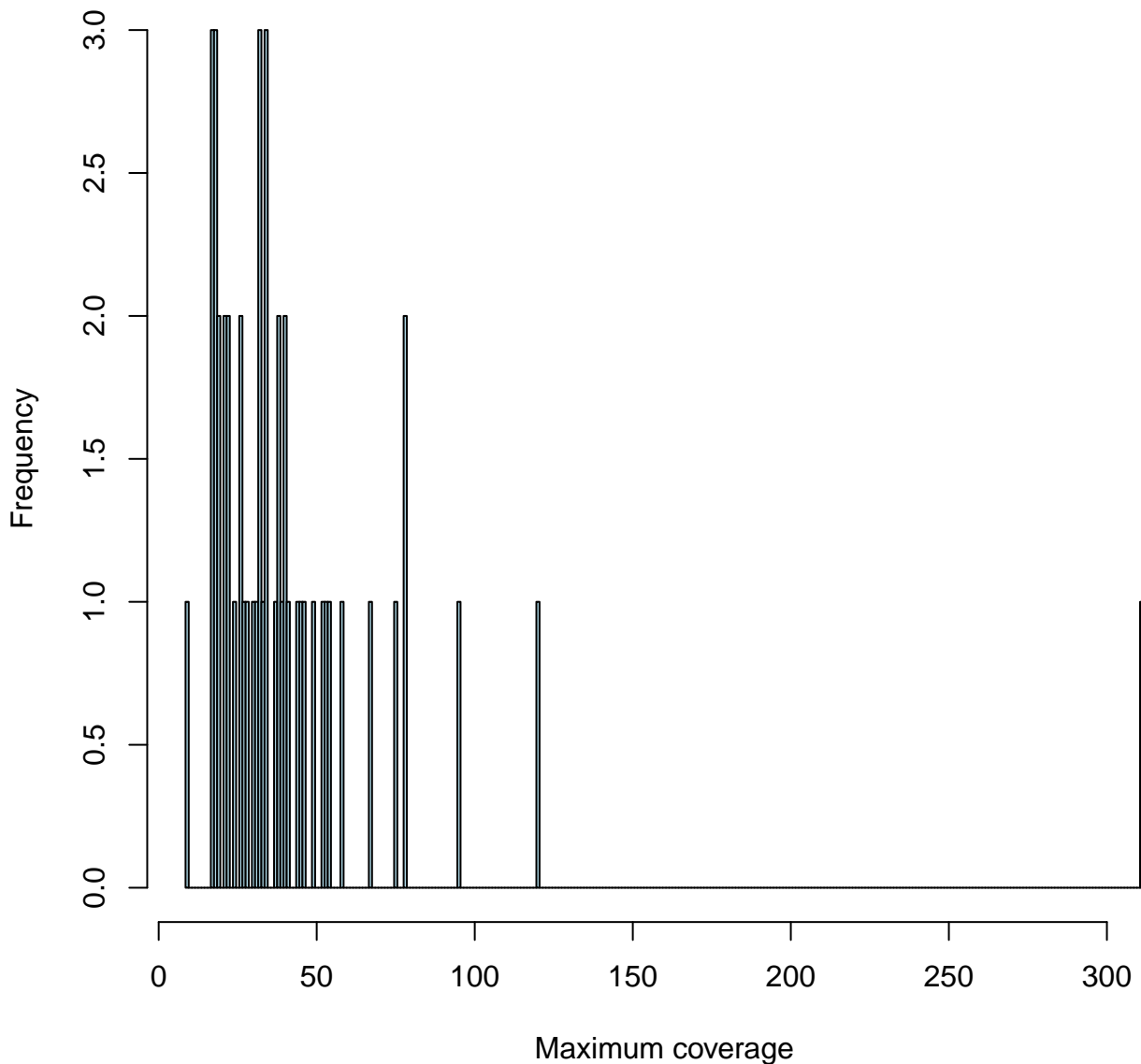
**jx in Intropolis but not "annotated" by knownGene:
204 samples; n jx: 72**



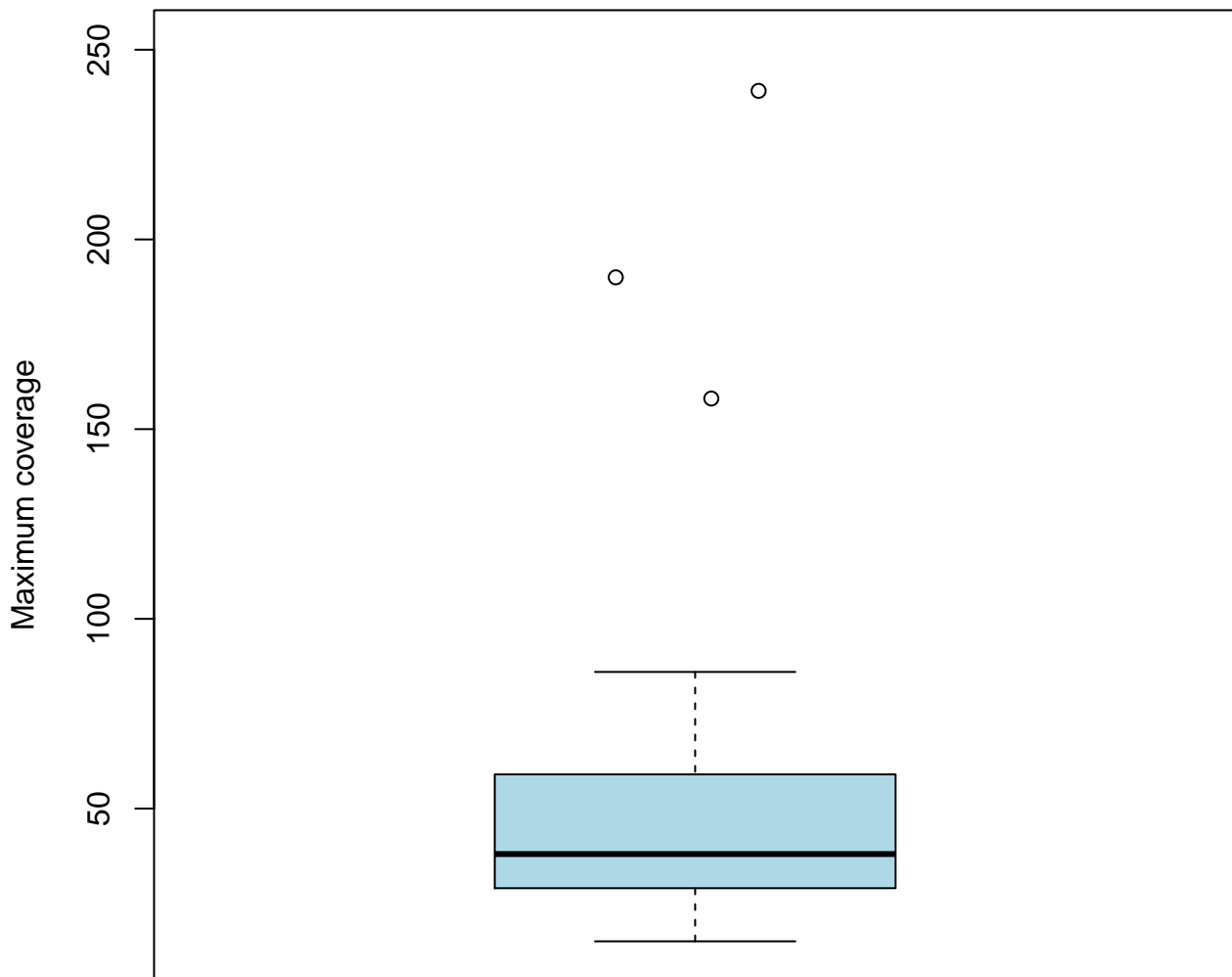
**jx in Intropolis but not "annotated" by knownGene:
205 samples; n jx: 49**



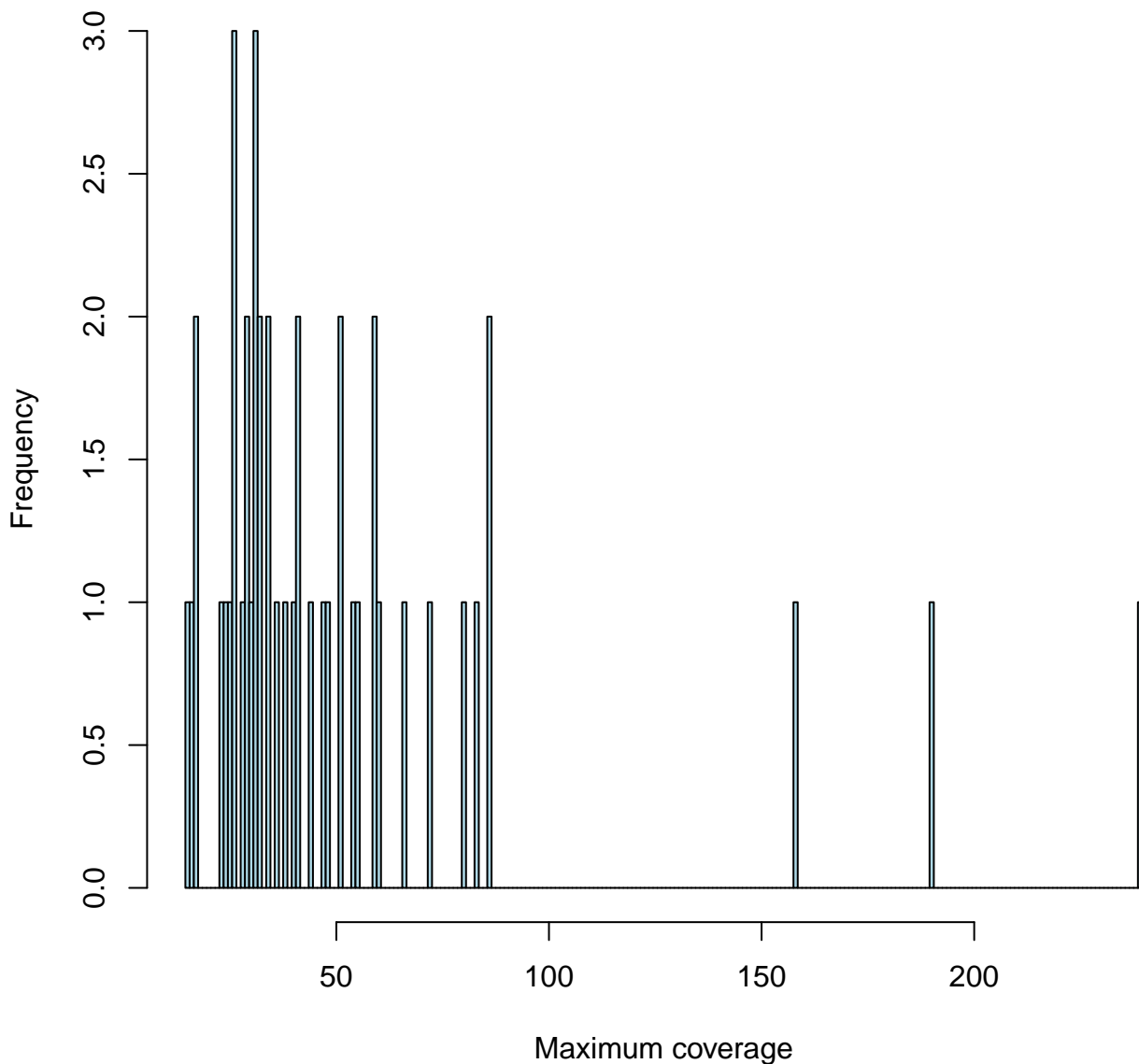
**jx in Intropolis but not "annotated" by knownGene:
205 samples; n jx: 49**



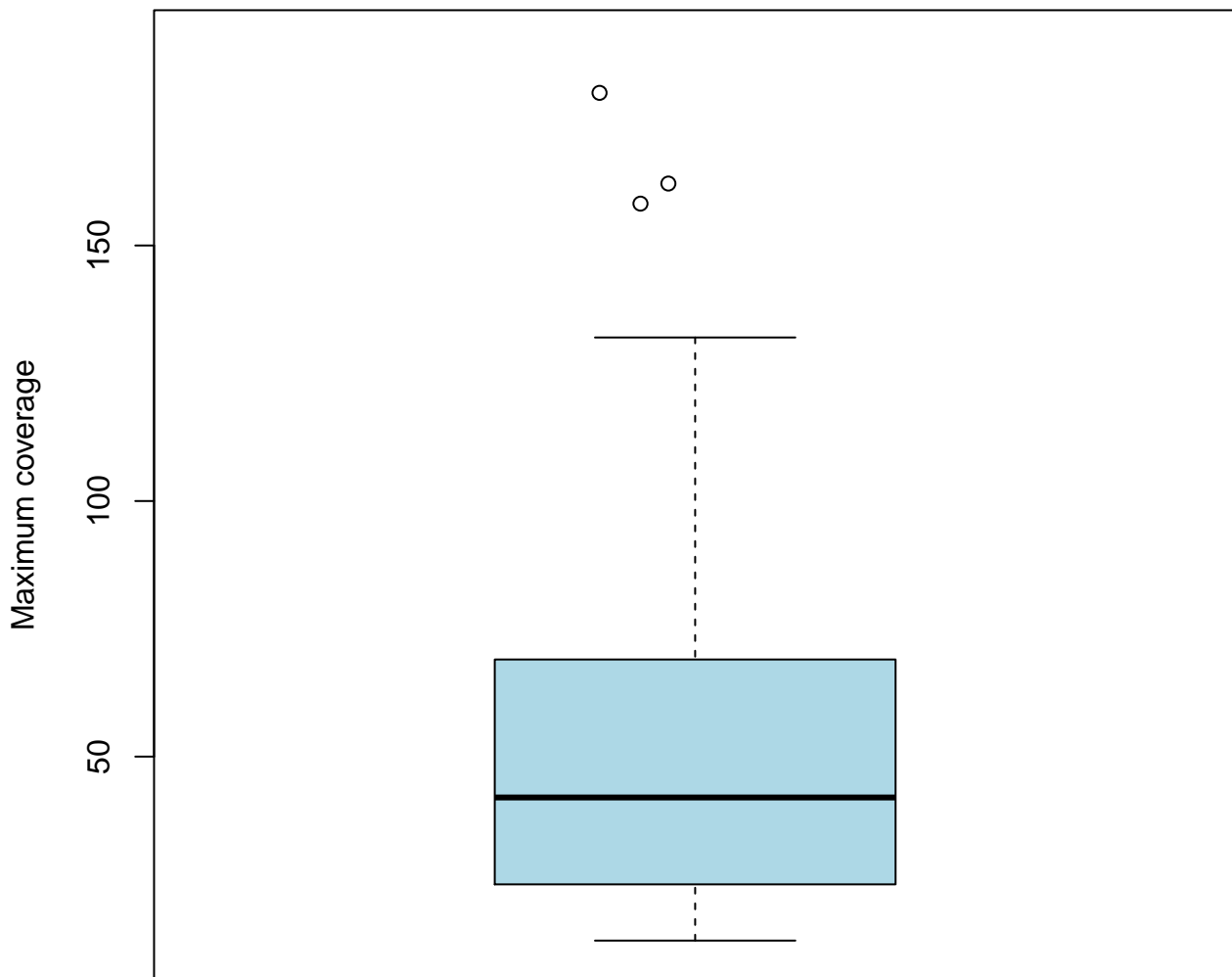
**jx in Intropolis but not "annotated" by knownGene:
206 samples; n jx: 45**



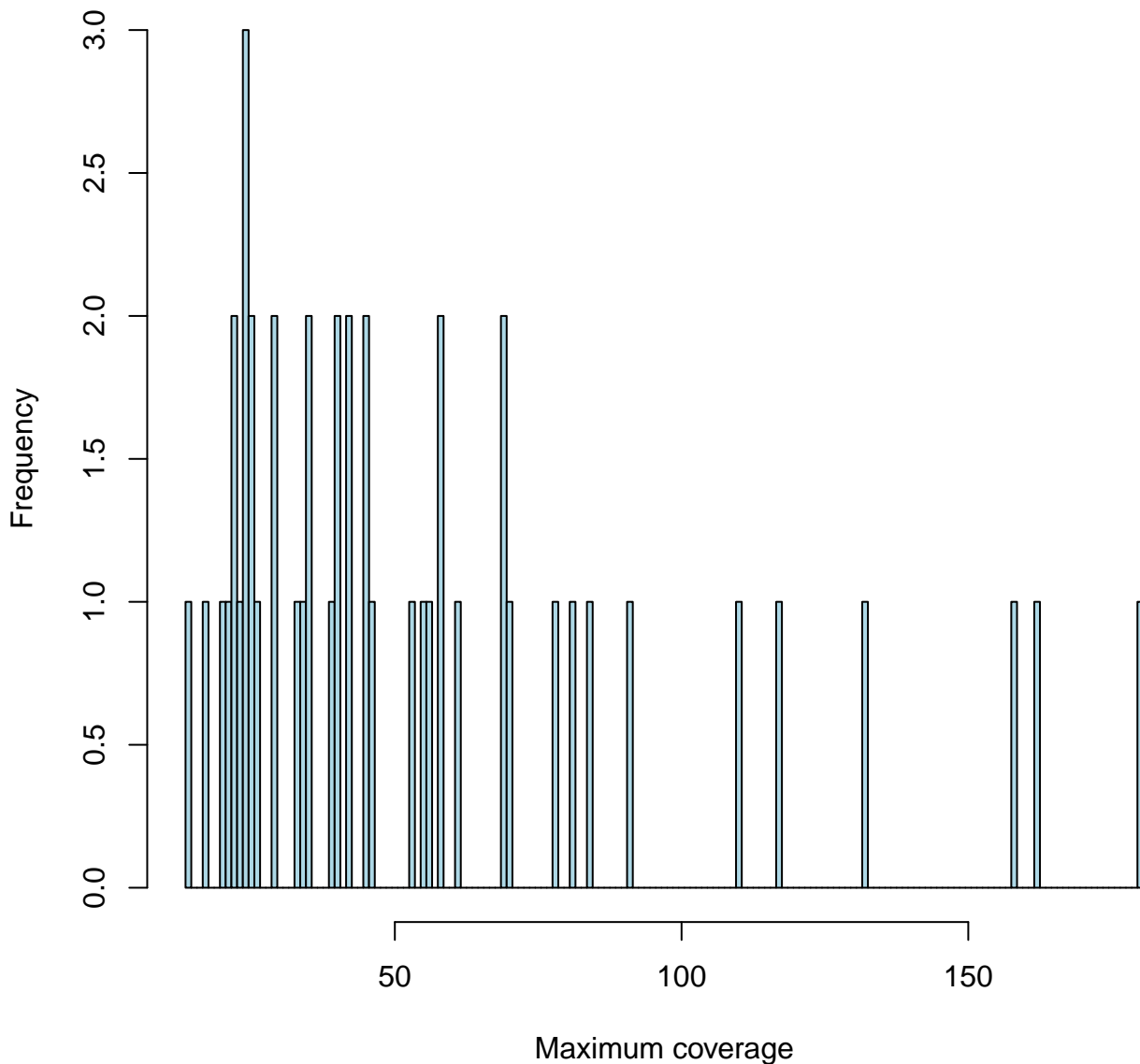
**jx in Intropolis but not "annotated" by knownGene:
206 samples; n jx: 45**



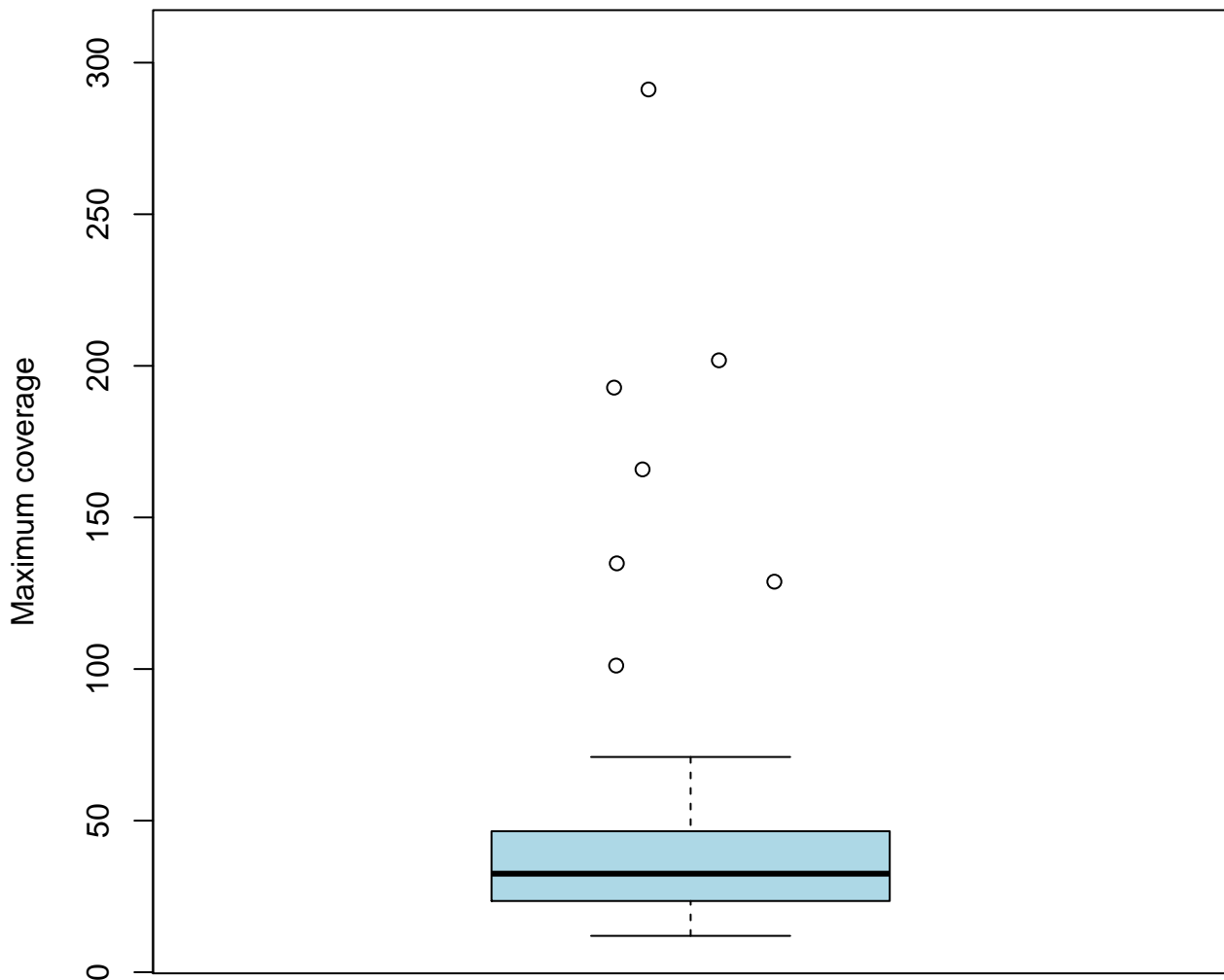
**jx in Intropolis but not "annotated" by knownGene:
207 samples; n jx: 46**



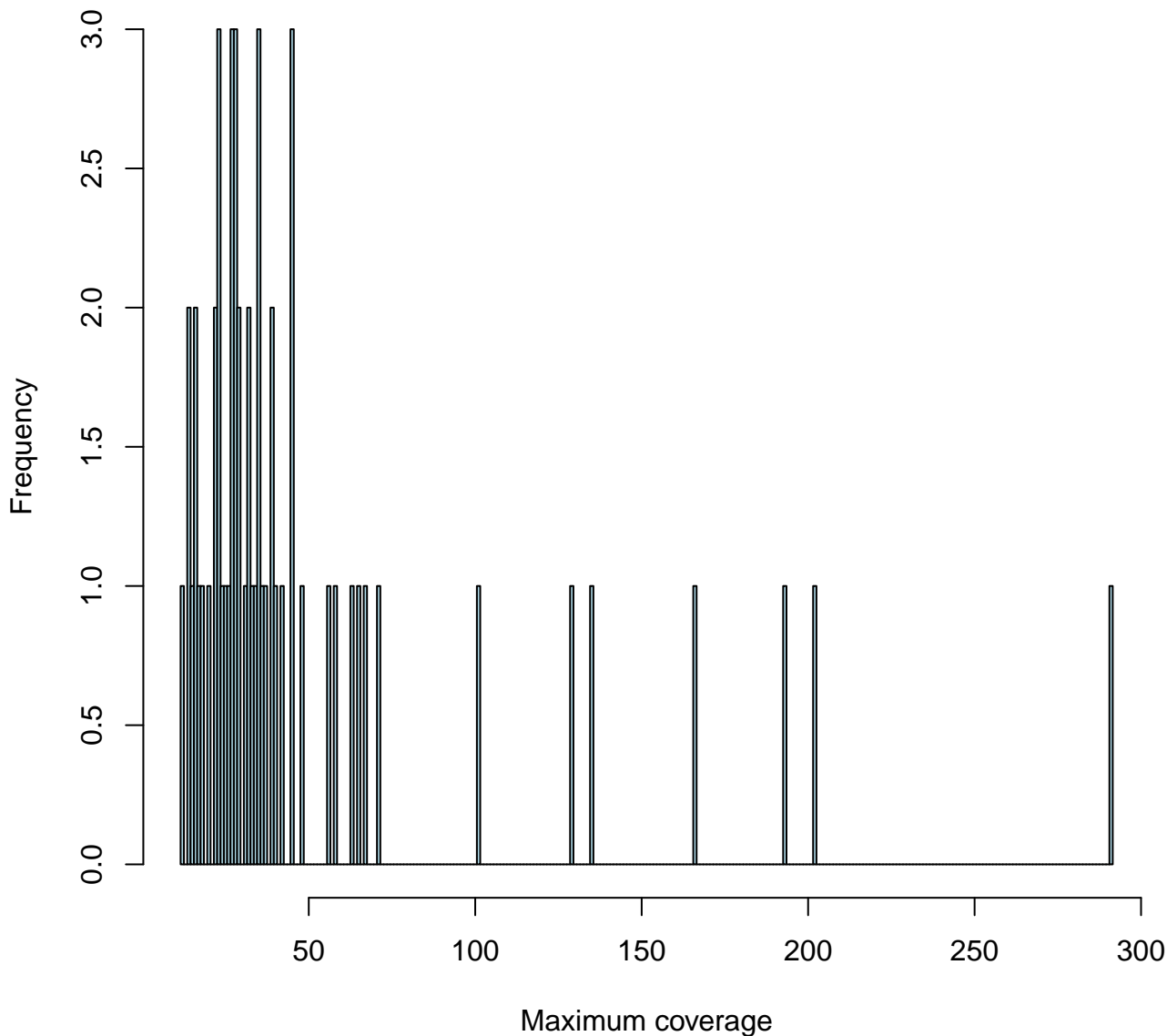
**jx in Intropolis but not "annotated" by knownGene:
207 samples; n jx: 46**



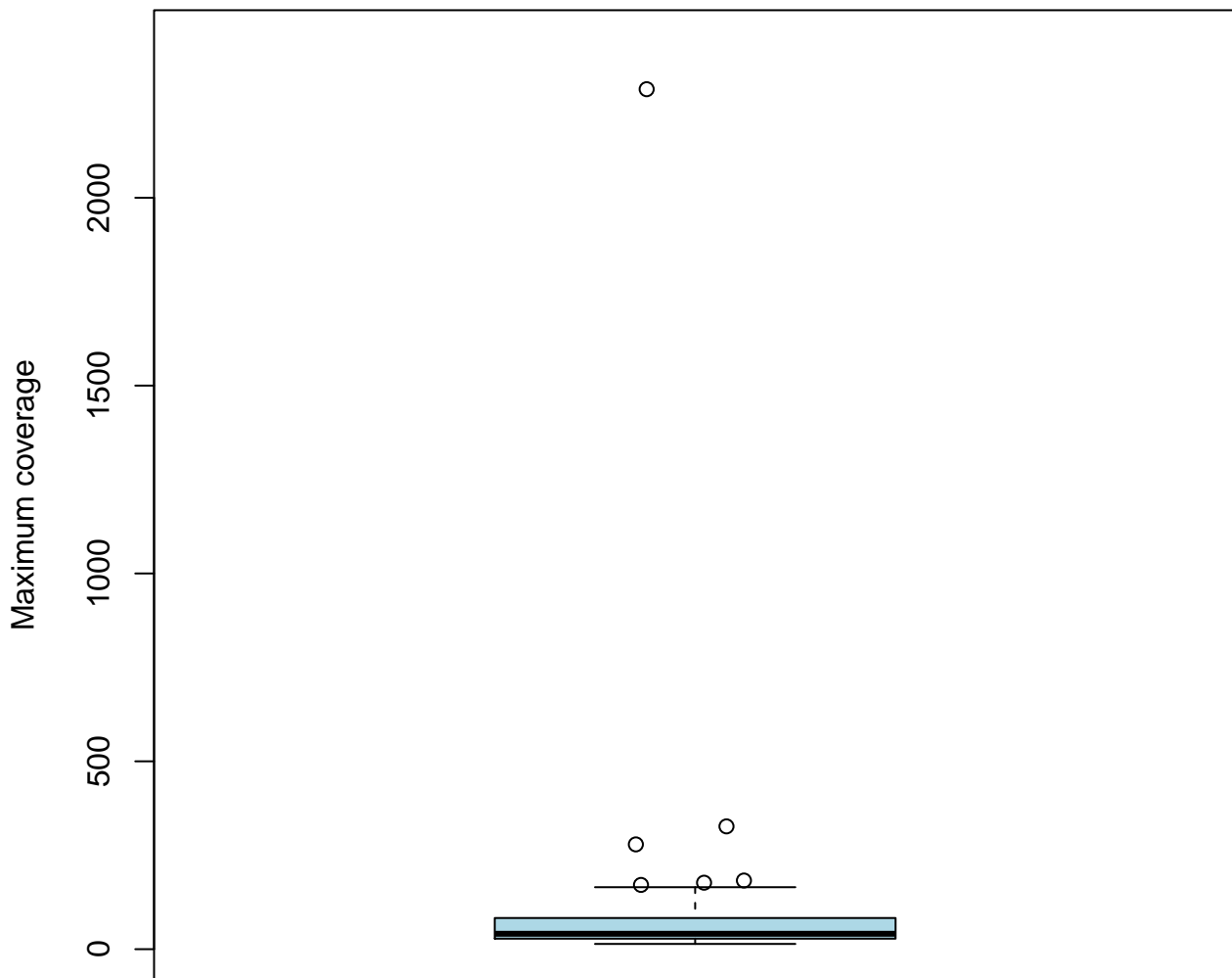
**jx in Intropolis but not "annotated" by knownGene:
208 samples; n jx: 56**



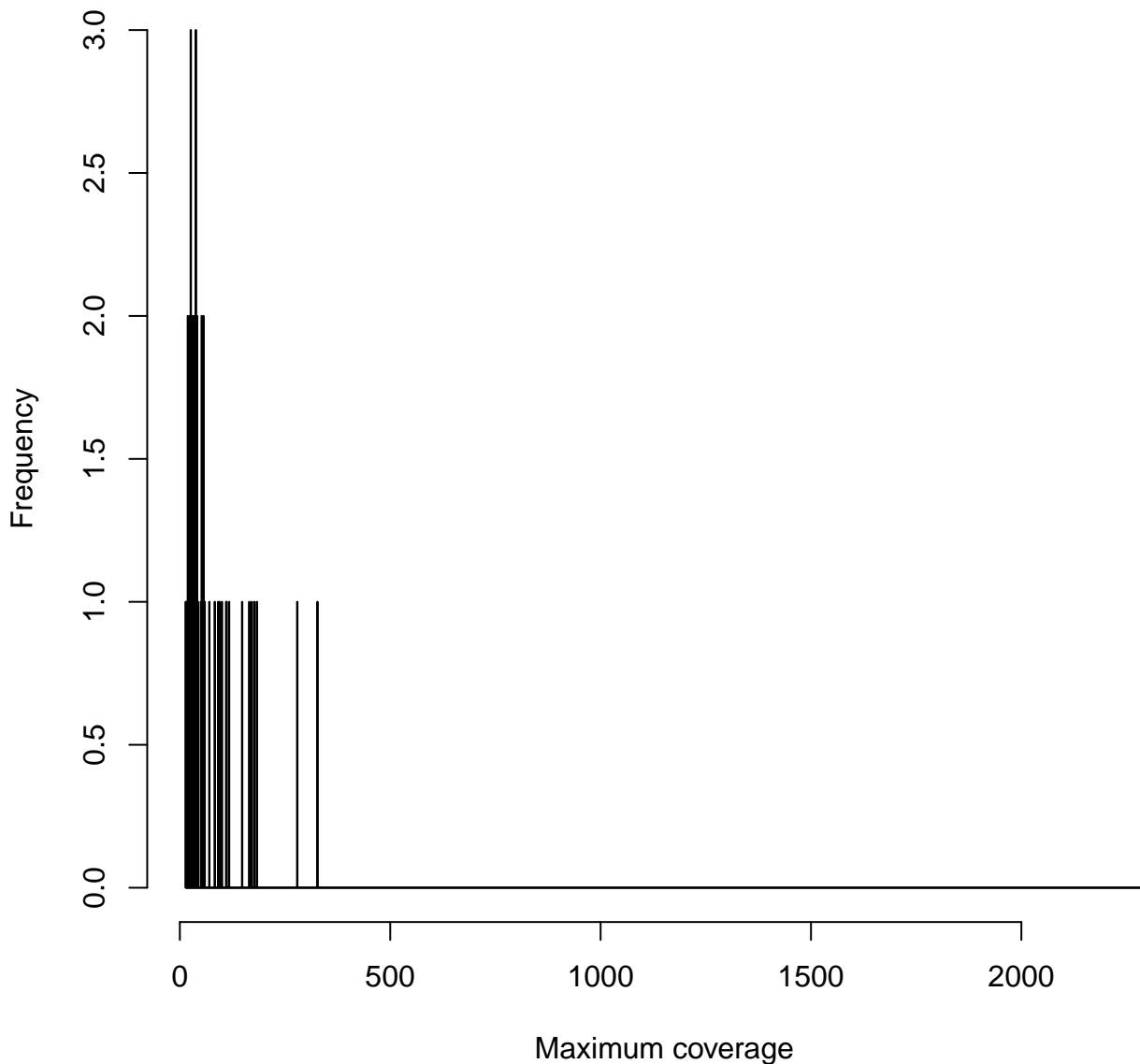
**jx in Intropolis but not "annotated" by knownGene:
208 samples; n jx: 56**



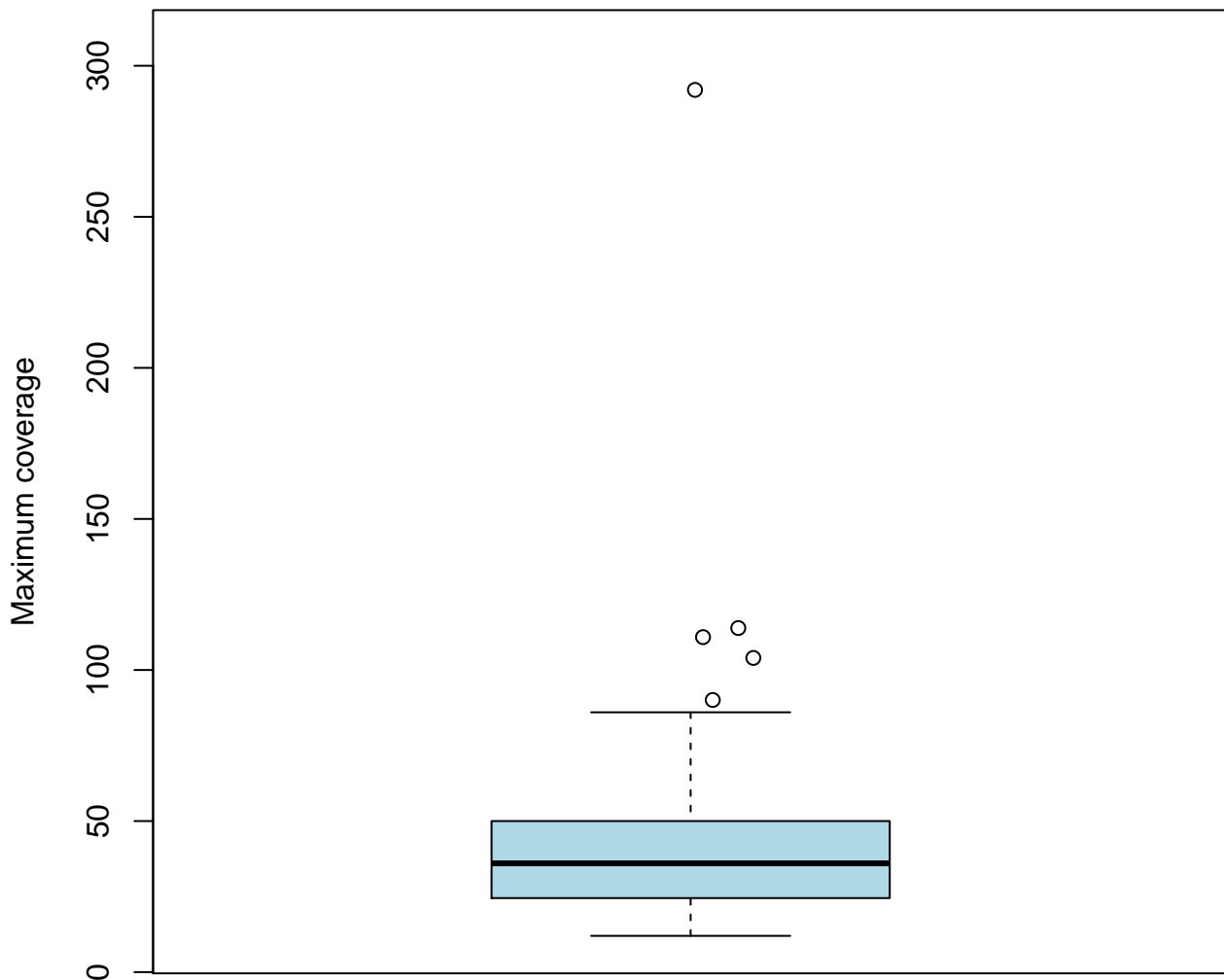
**jx in Intropolis but not "annotated" by knownGene:
209 samples; n jx: 58**



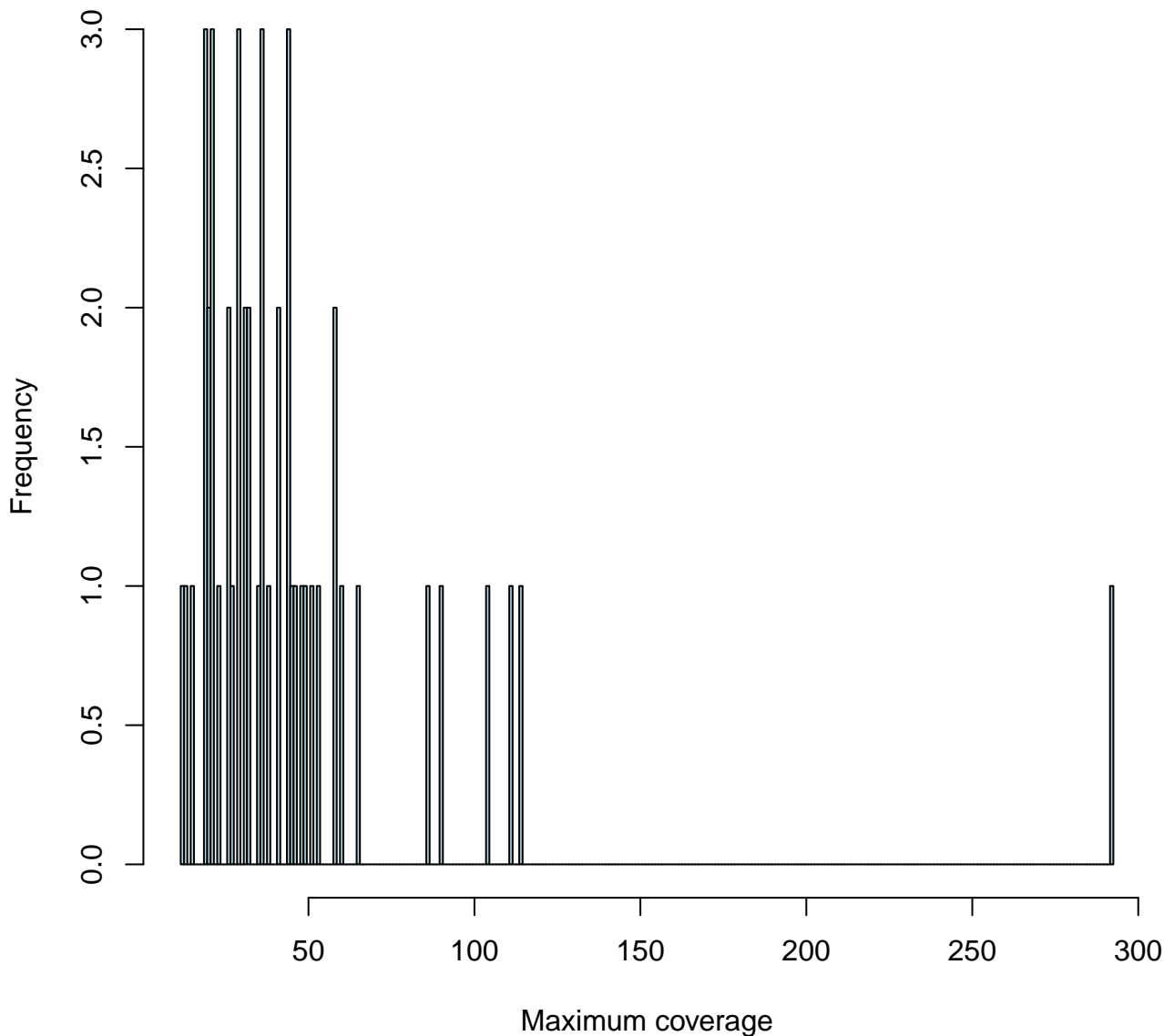
**jx in Intropolis but not "annotated" by knownGene:
209 samples; n jx: 58**



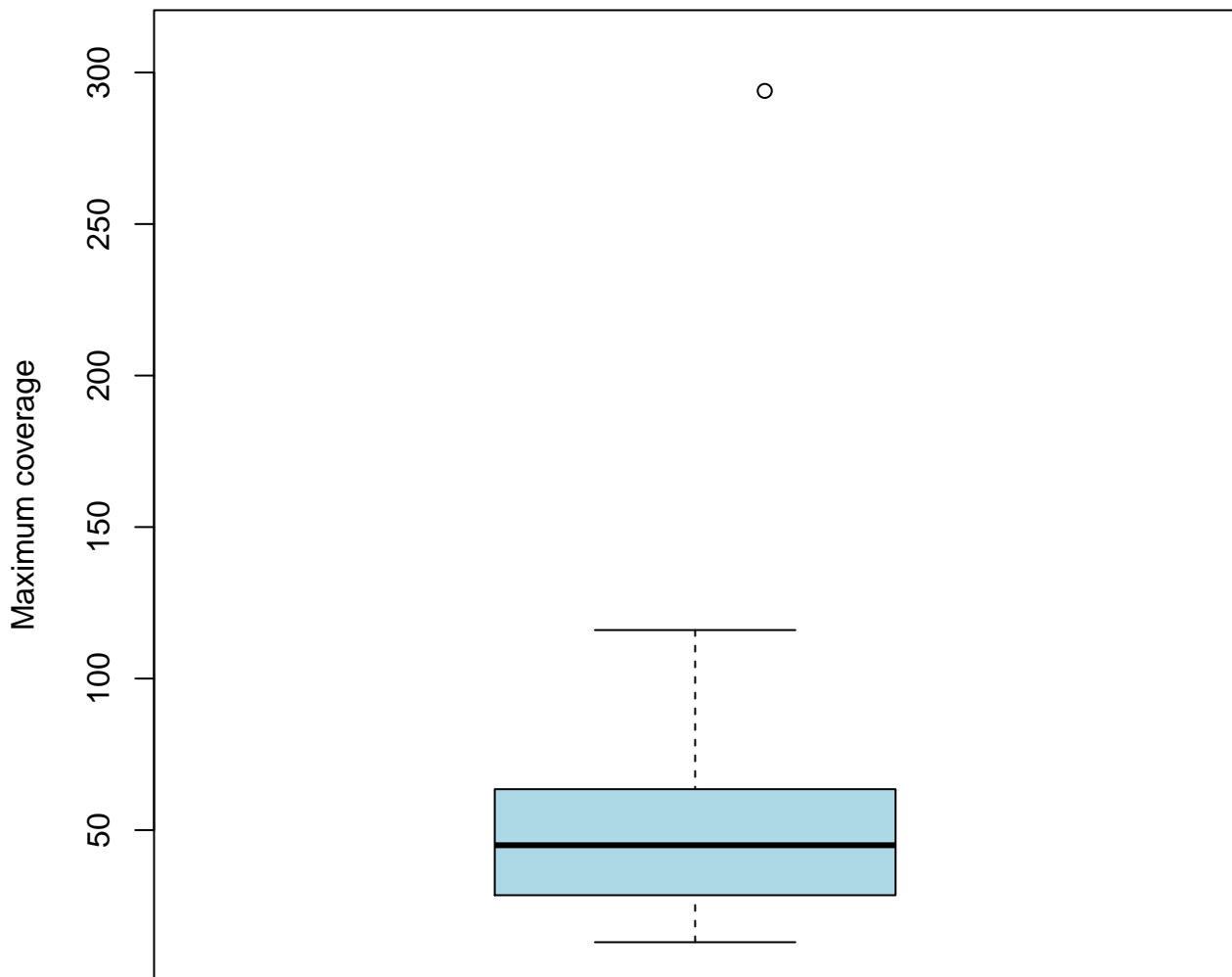
**jx in Intropolis but not "annotated" by knownGene:
210 samples; n jx: 48**



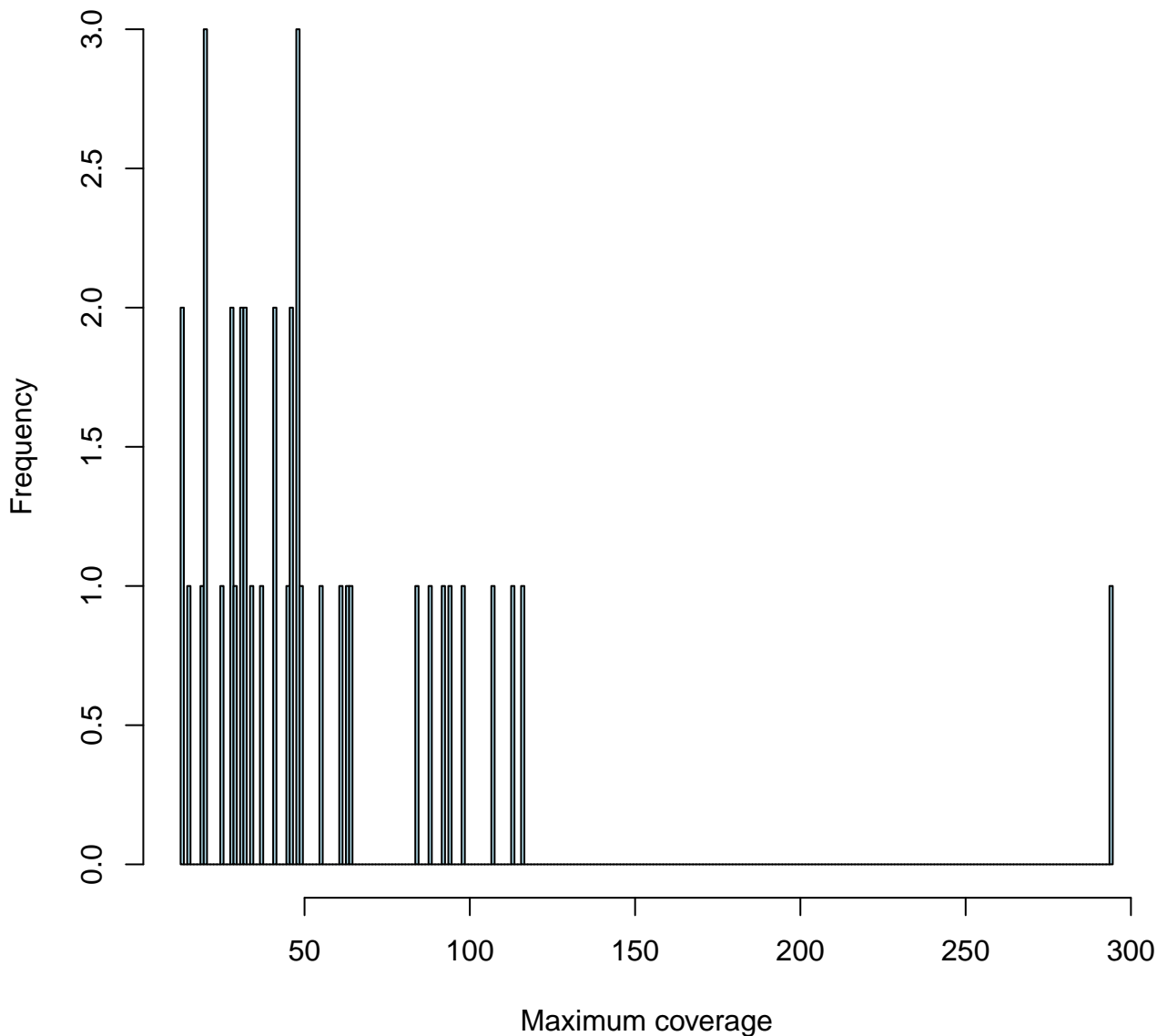
**jx in Intropolis but not "annotated" by knownGene:
210 samples; n jx: 48**



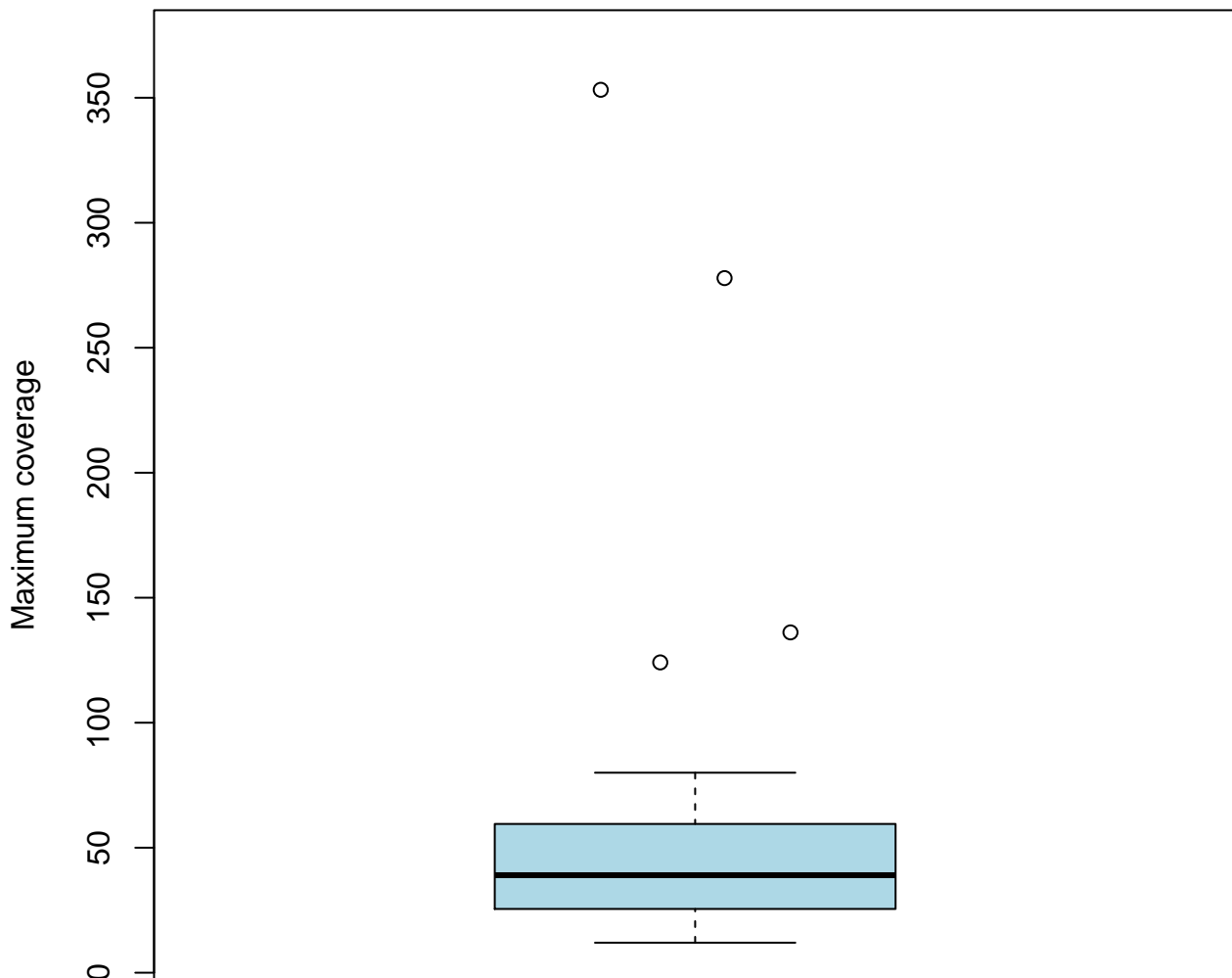
**jx in Intropolis but not "annotated" by knownGene:
211 samples; n jx: 39**



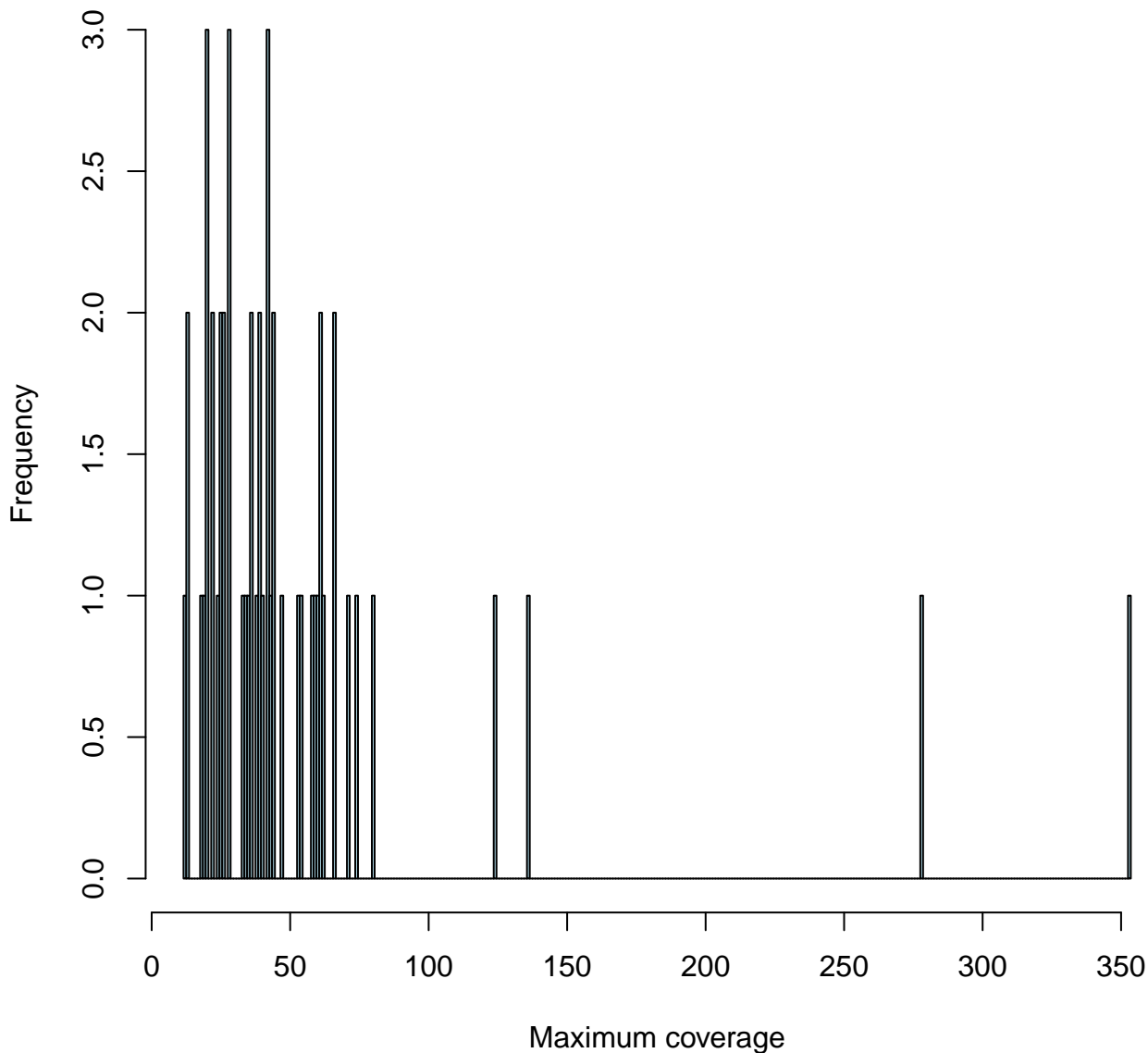
**jx in Intropolis but not "annotated" by knownGene:
211 samples; n jx: 39**



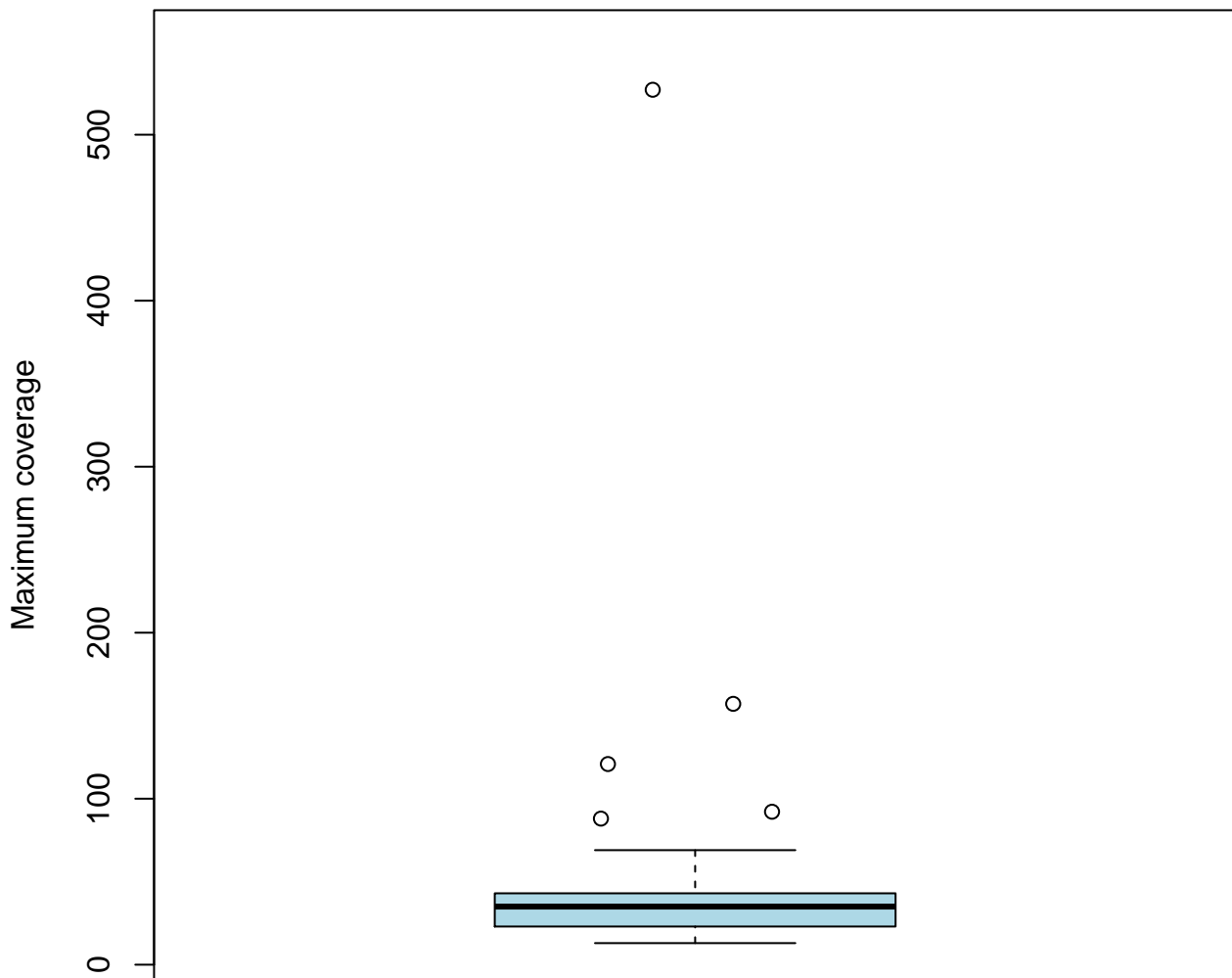
**jx in Intropolis but not "annotated" by knownGene:
212 samples; n jx: 51**



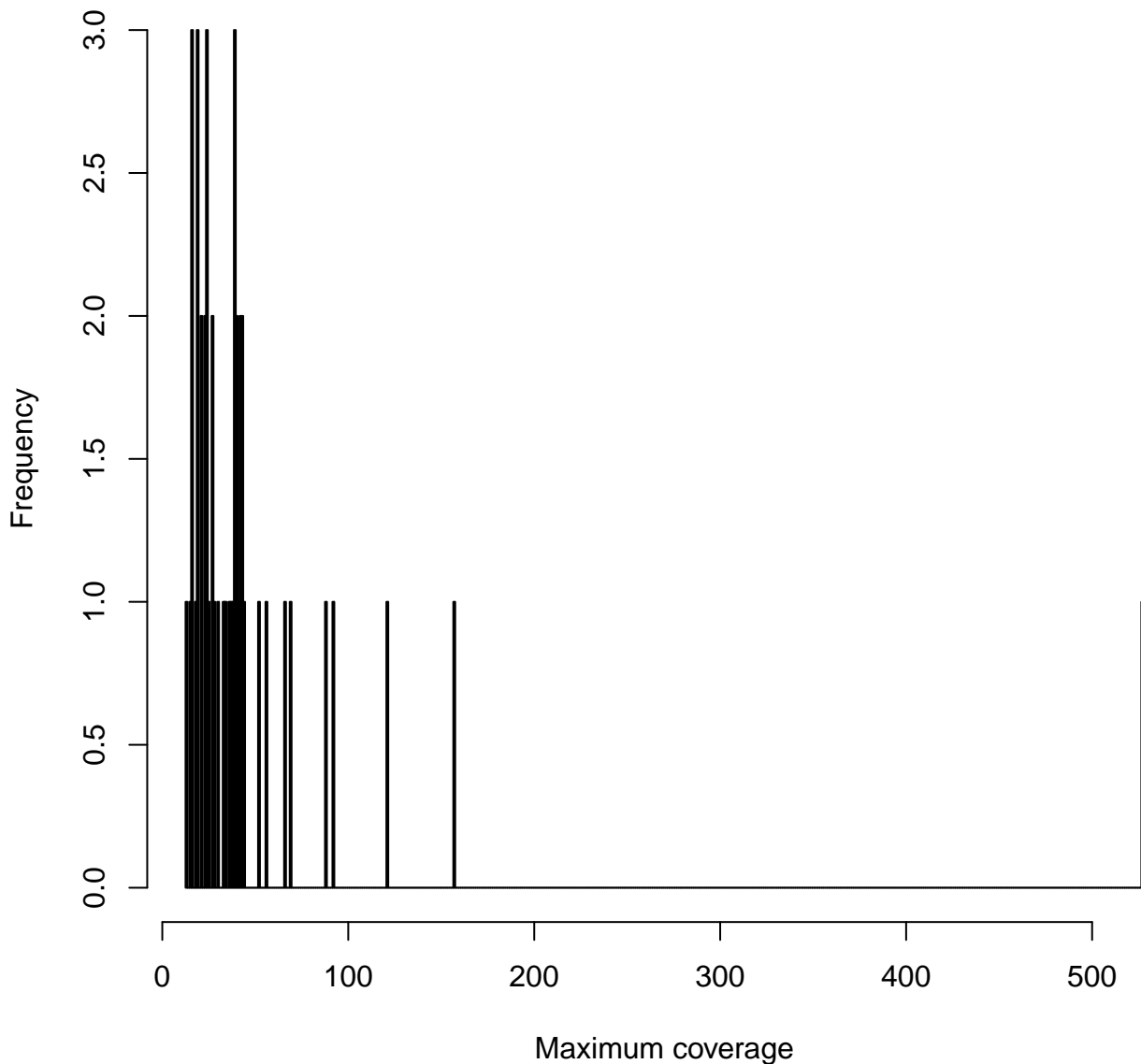
**jl in Intropolis but not "annotated" by knownGene:
212 samples; n jl: 51**



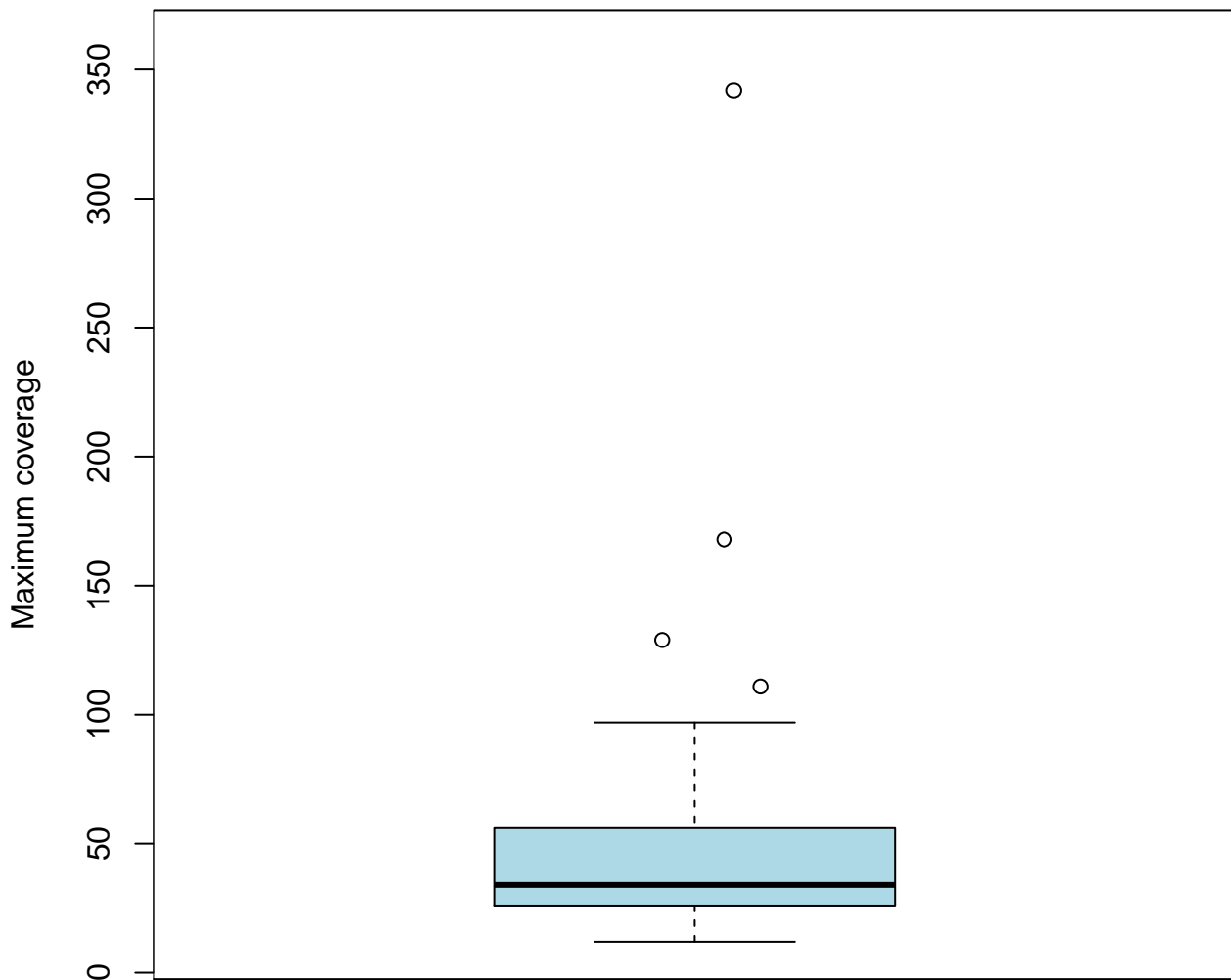
**jx in Intropolis but not "annotated" by knownGene:
213 samples; n jx: 46**



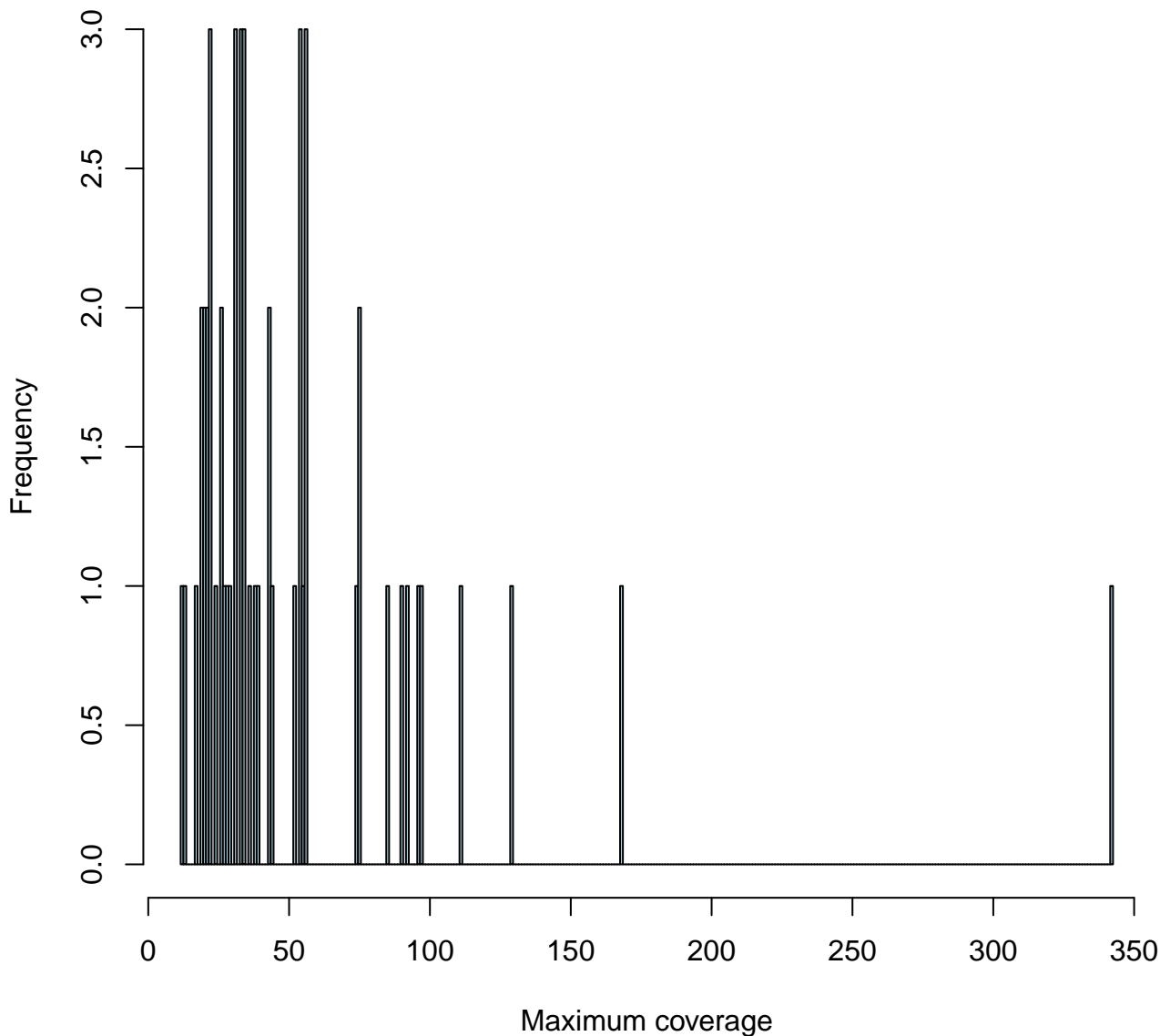
**jsx in Intropolis but not "annotated" by knownGene:
213 samples; n jsx: 46**



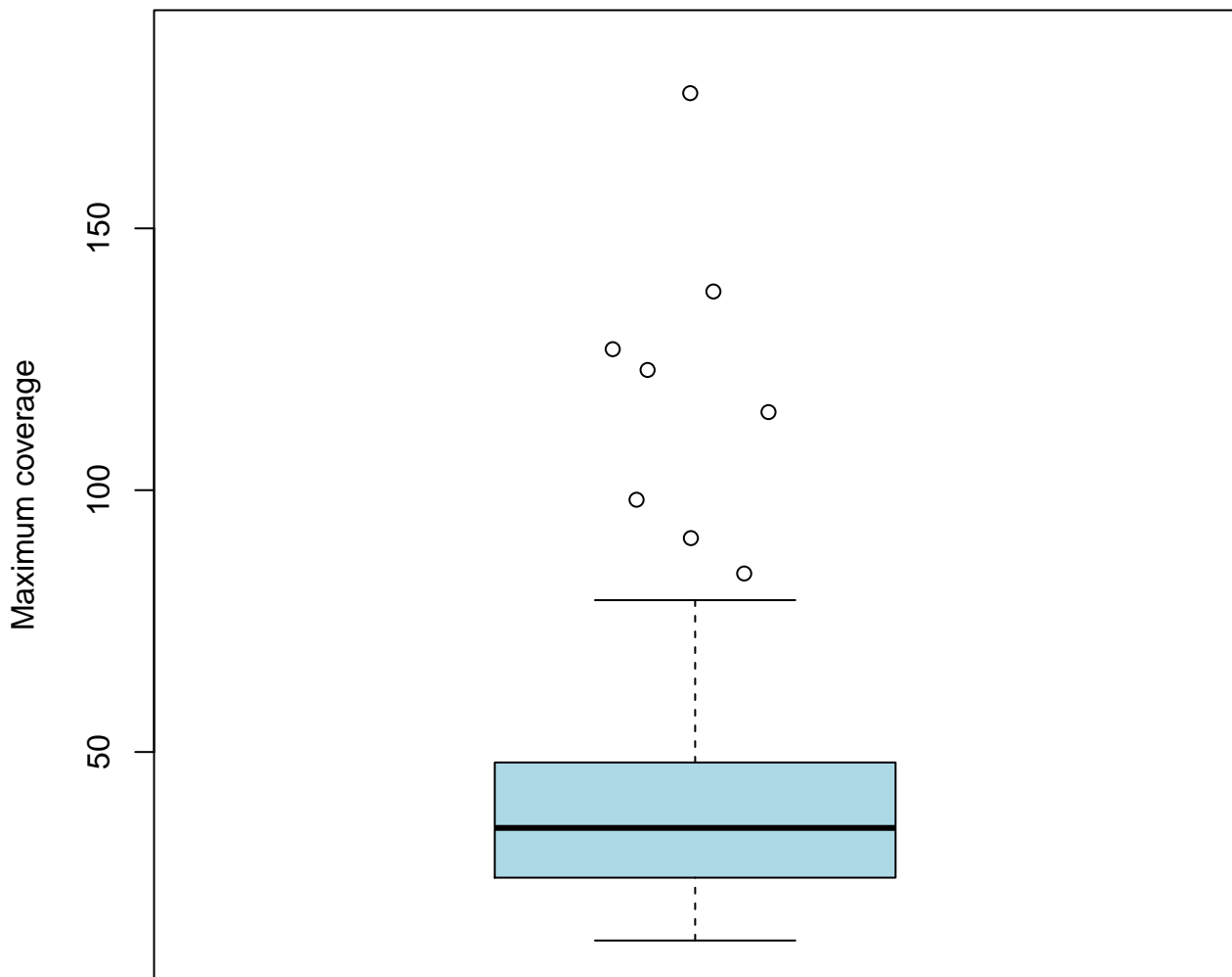
**jx in Intropolis but not "annotated" by knownGene:
214 samples; n jx: 53**



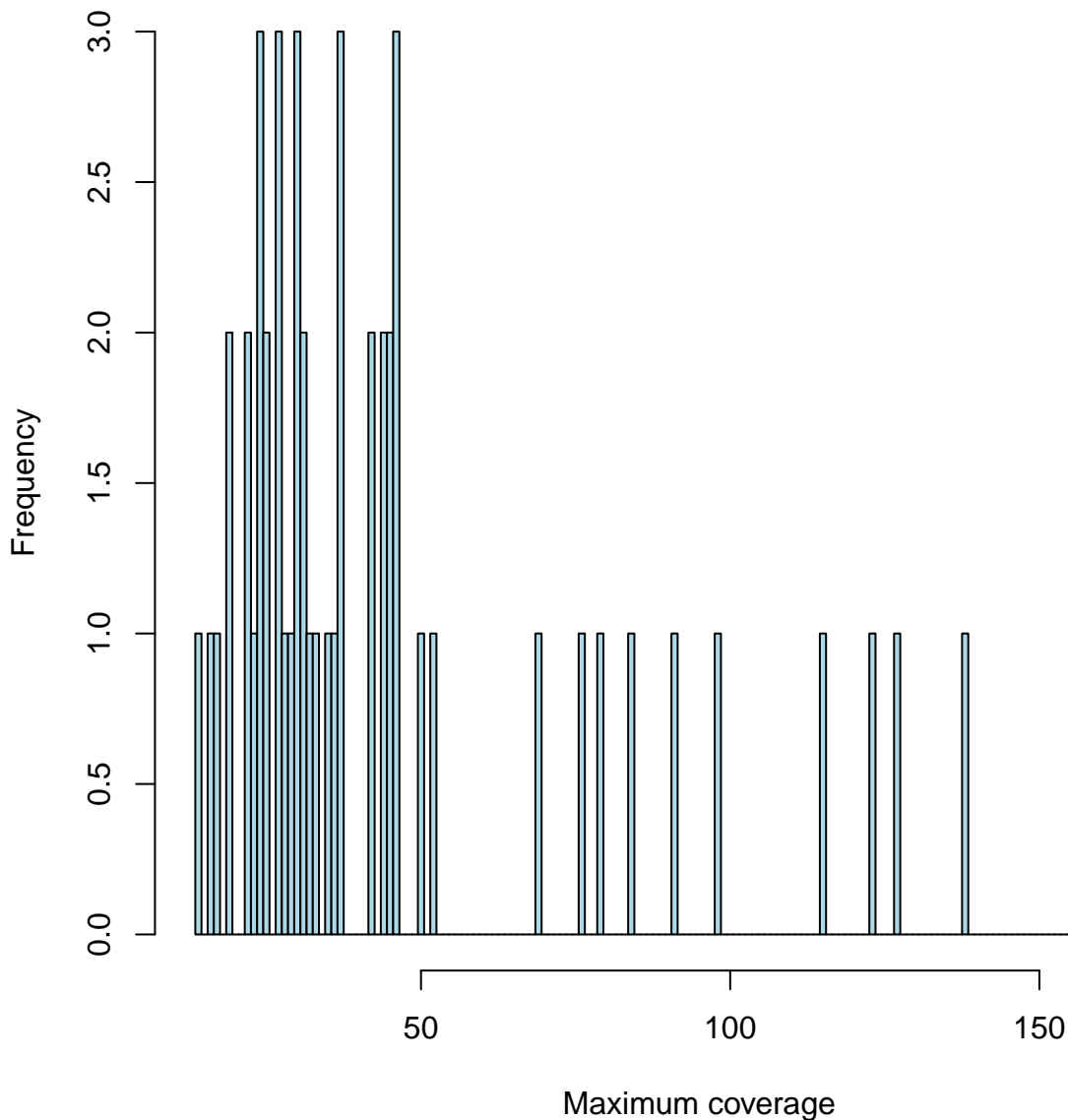
**jx in Intropolis but not "annotated" by knownGene:
214 samples; n jx: 53**



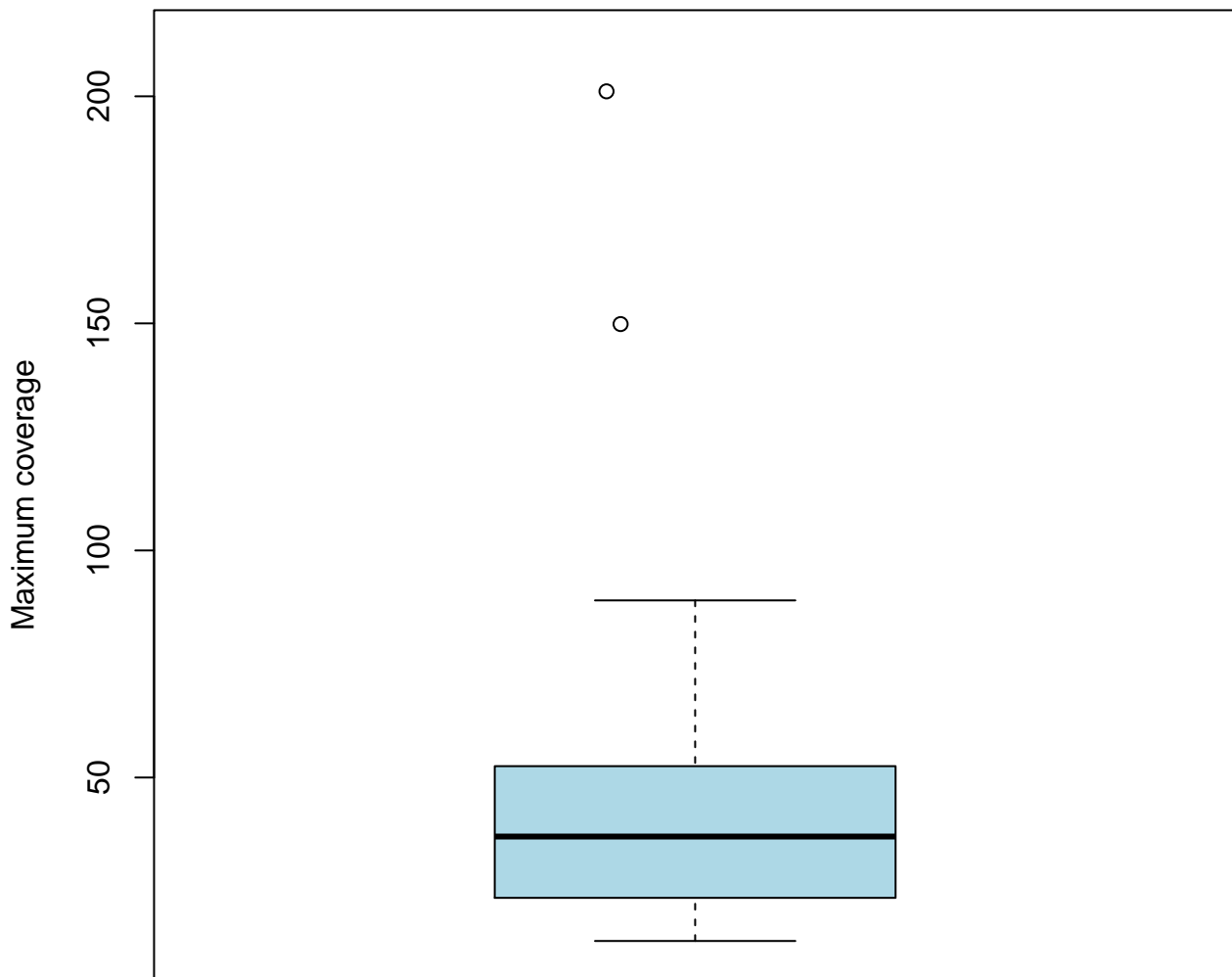
**jx in Intropolis but not "annotated" by knownGene:
215 samples; n jx: 52**



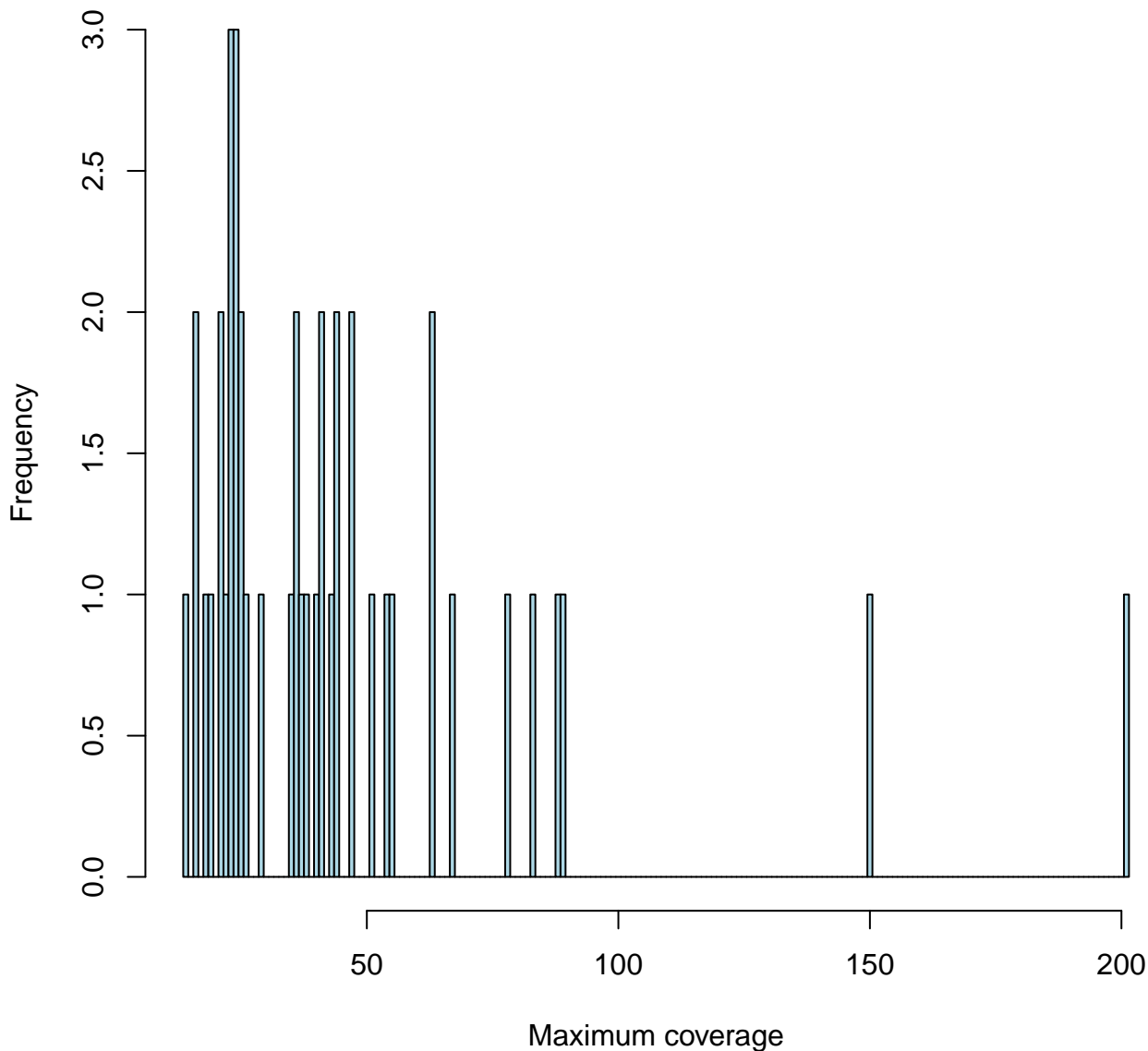
**jx in Intropolis but not "annotated" by knownGene:
215 samples; n jx: 52**



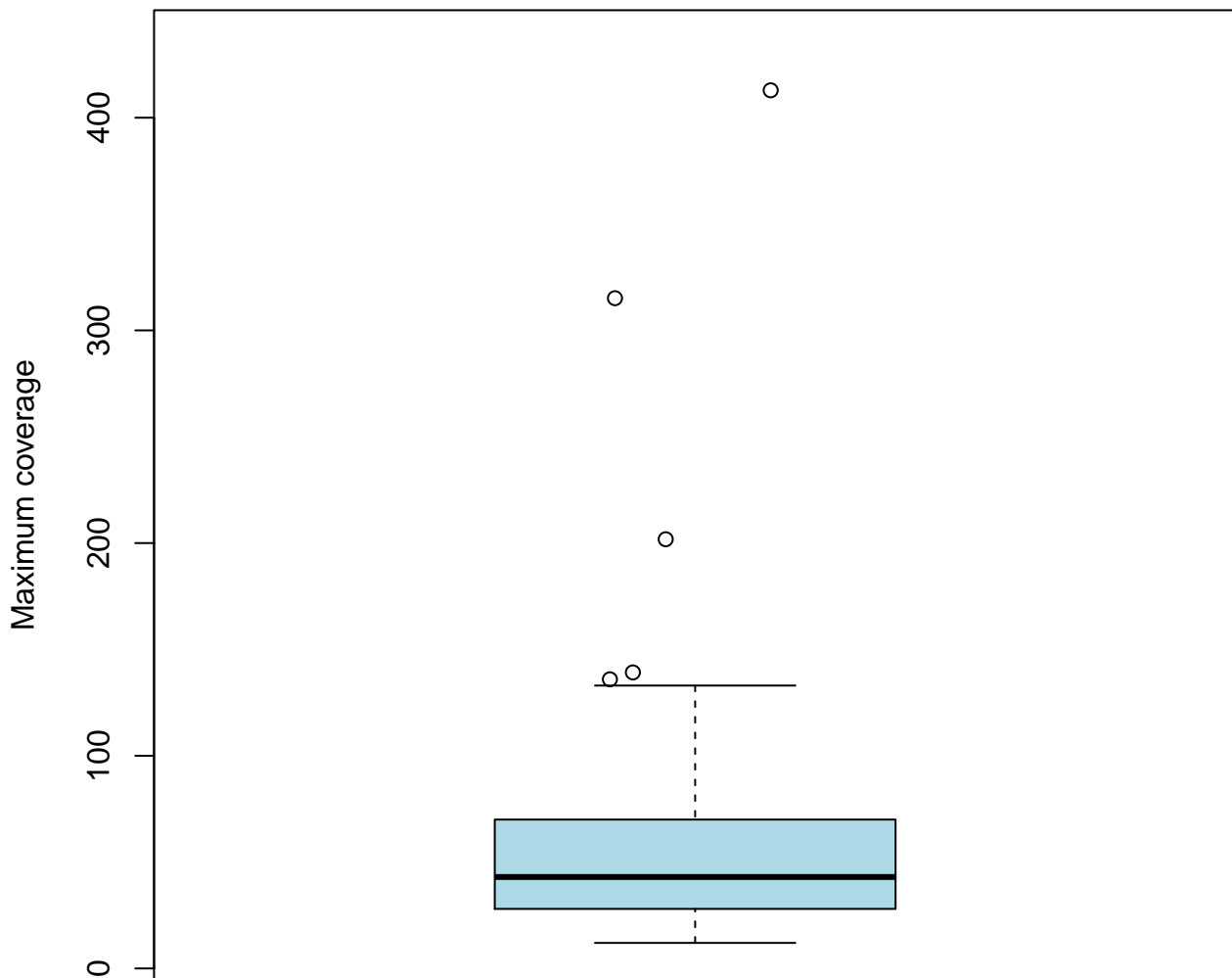
**jx in Intropolis but not "annotated" by knownGene:
216 samples; n jx: 43**



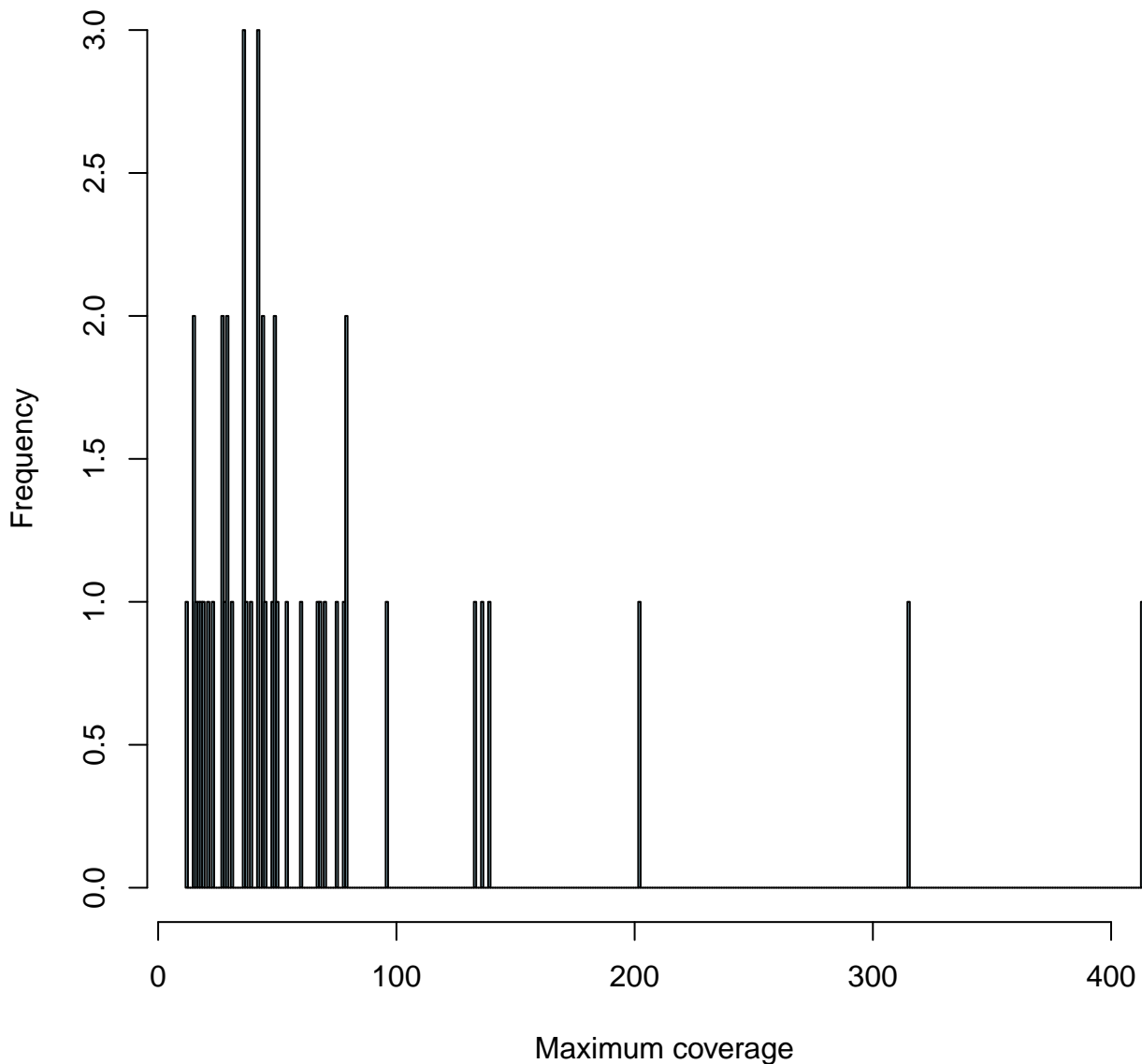
**jx in Intropolis but not "annotated" by knownGene:
216 samples; n jx: 43**



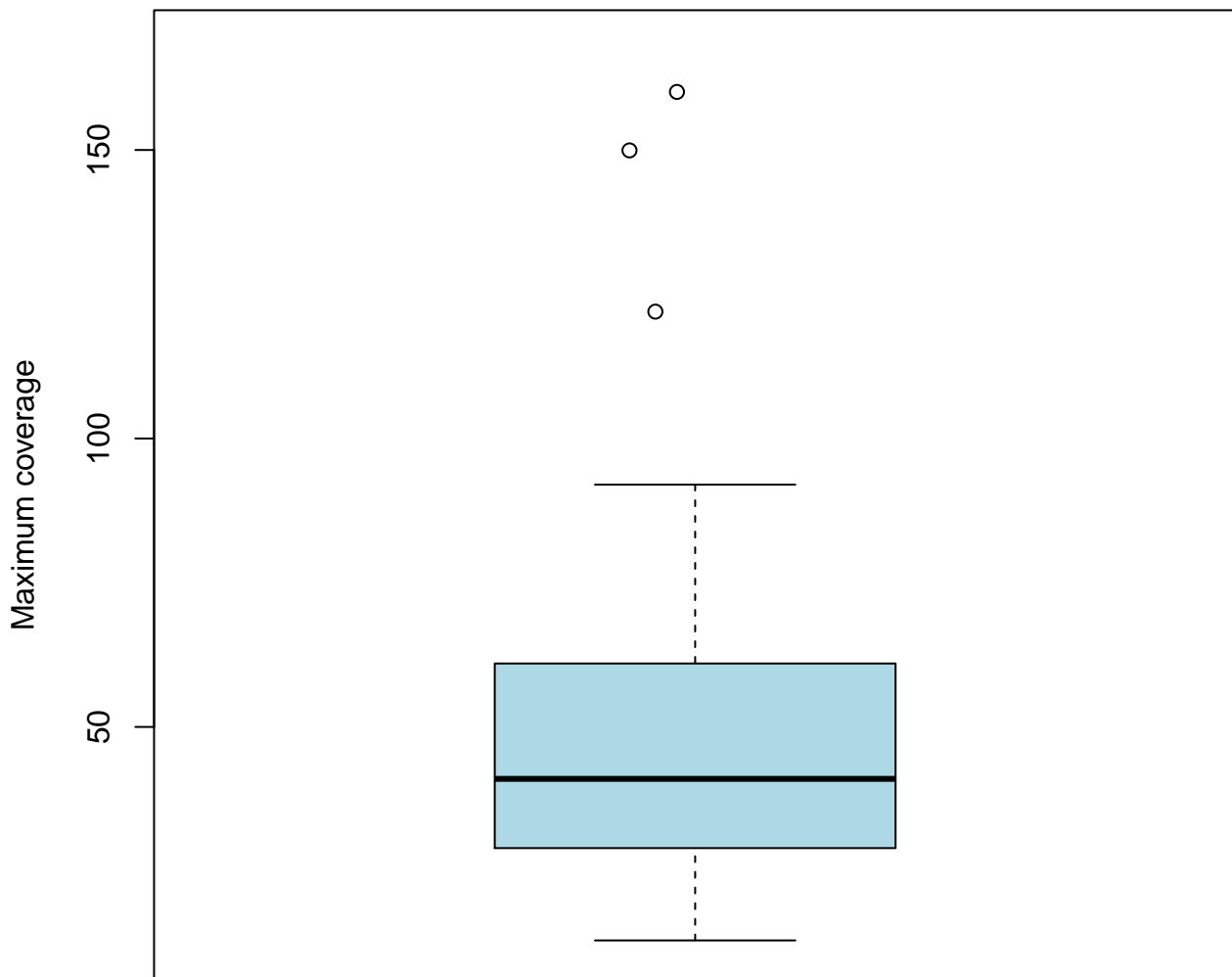
**jx in Intropolis but not "annotated" by knownGene:
217 samples; n jx: 46**



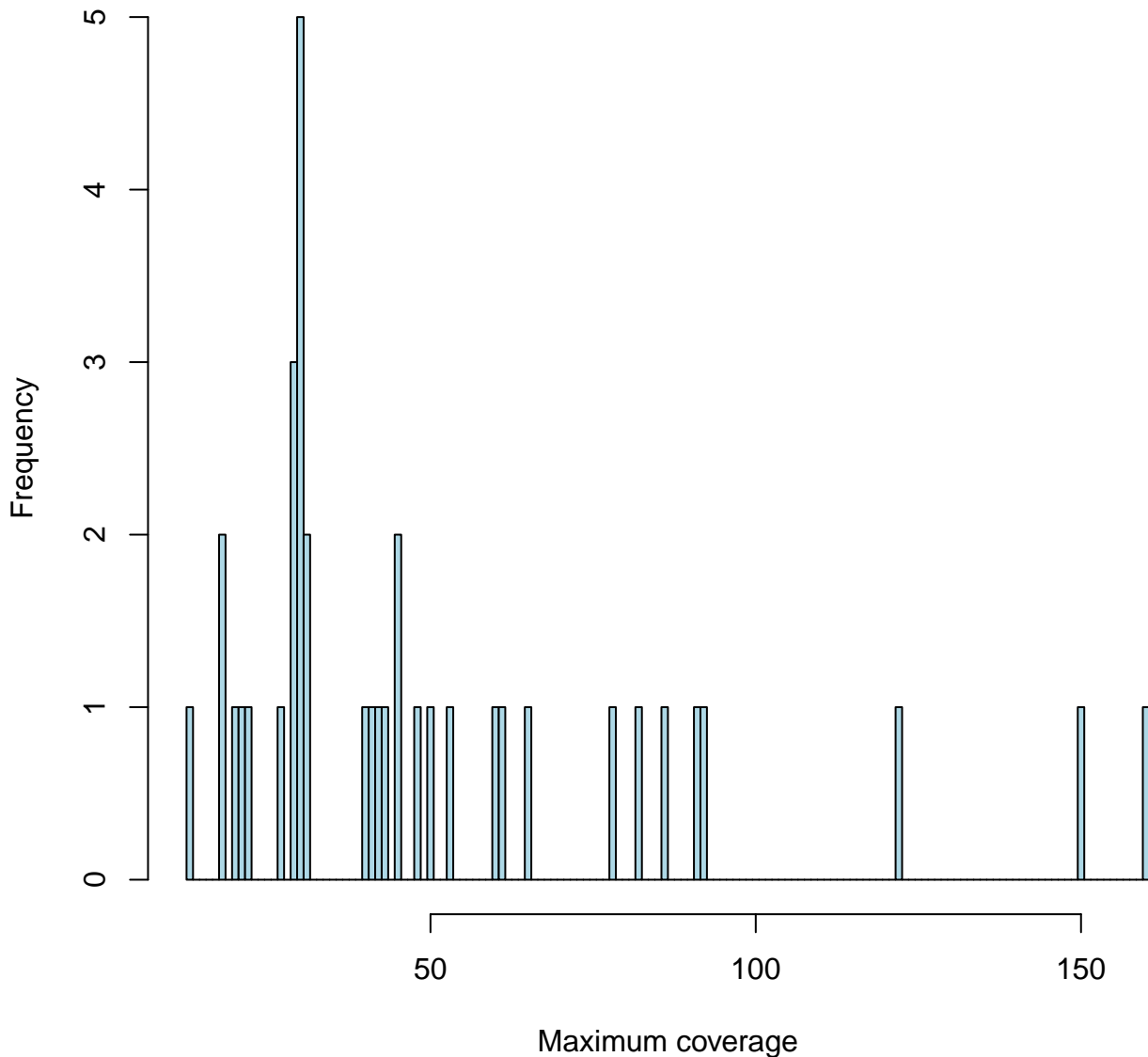
**jx in Intropolis but not "annotated" by knownGene:
217 samples; n jx: 46**



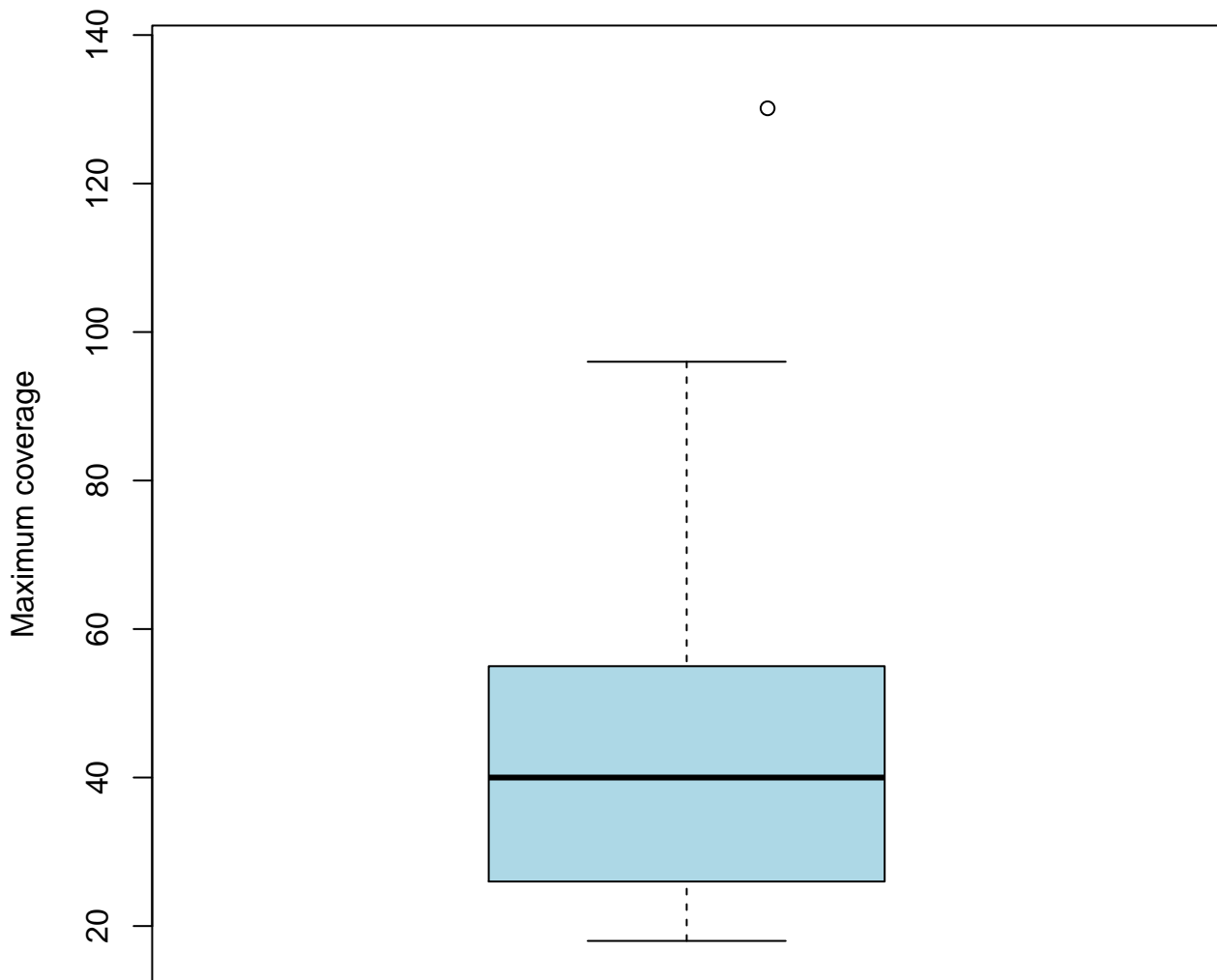
**jx in Intropolis but not "annotated" by knownGene:
218 samples; n jx: 37**



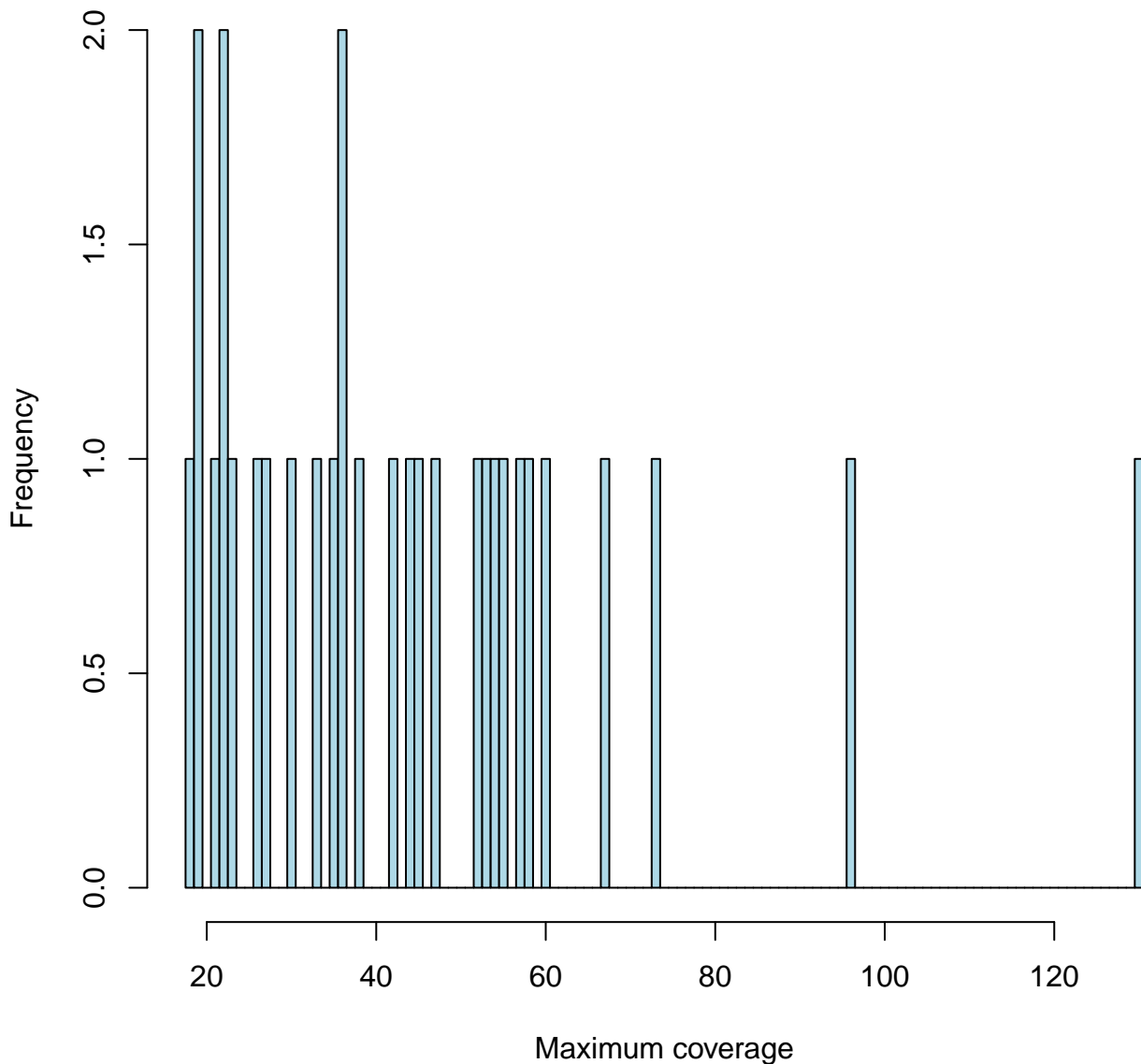
**jx in Intropolis but not "annotated" by knownGene:
218 samples; n jx: 37**



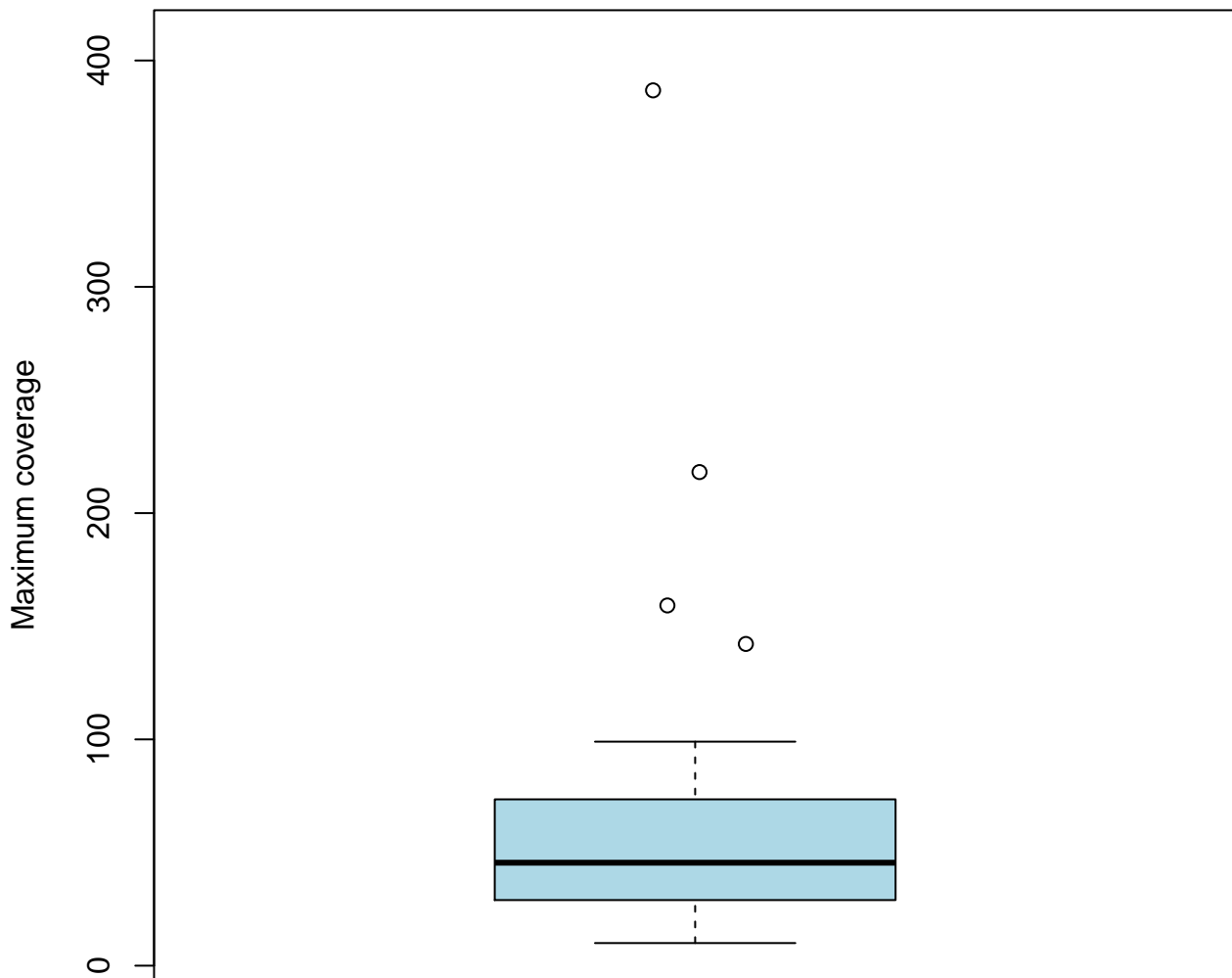
**jx in Intropolis but not "annotated" by knownGene:
219 samples; n jx: 30**



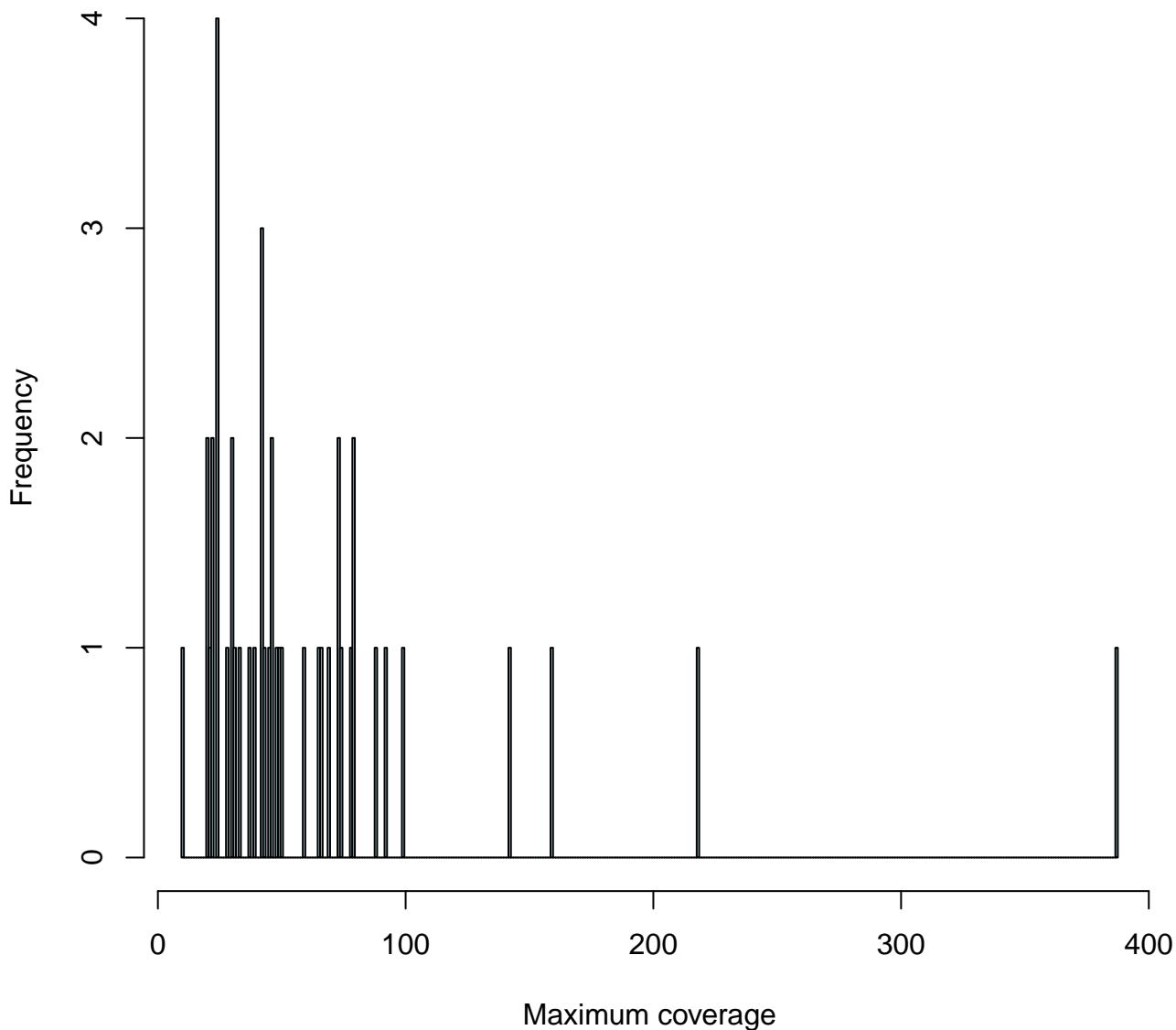
**jx in Intropolis but not "annotated" by knownGene:
219 samples; n jx: 30**



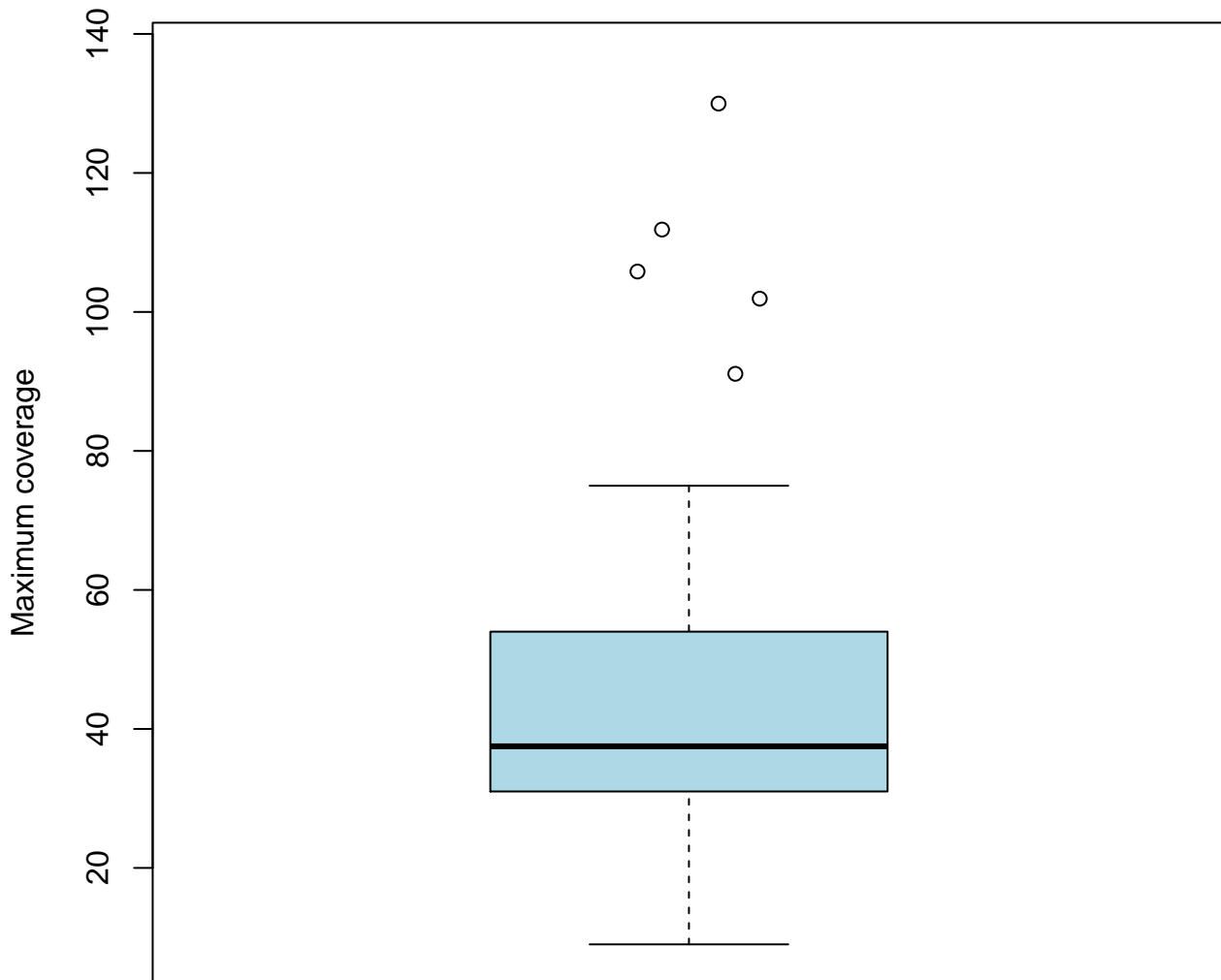
**jx in Intropolis but not "annotated" by knownGene:
220 samples; n jx: 44**



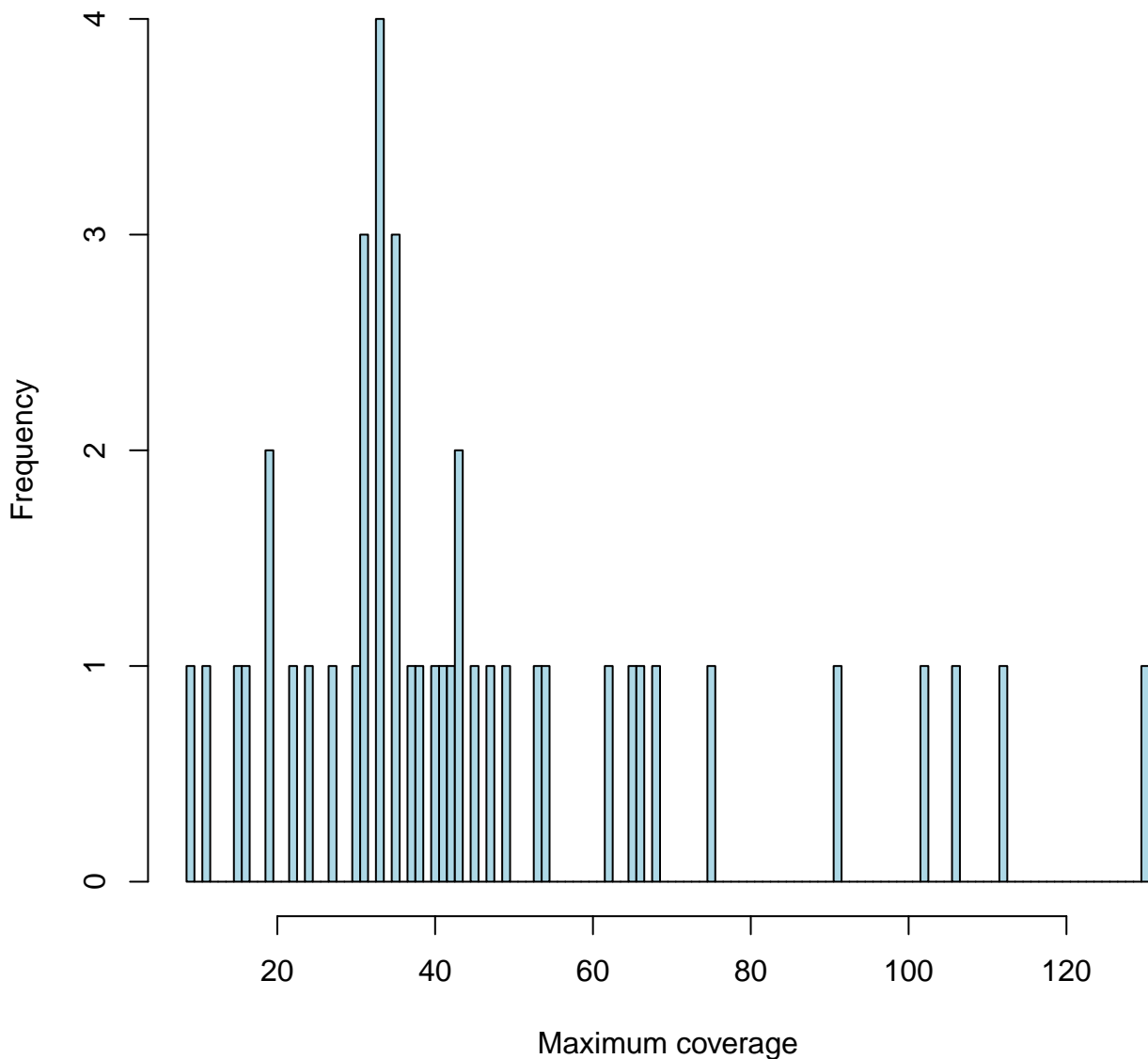
**jx in Intropolis but not "annotated" by knownGene:
220 samples; n jx: 44**



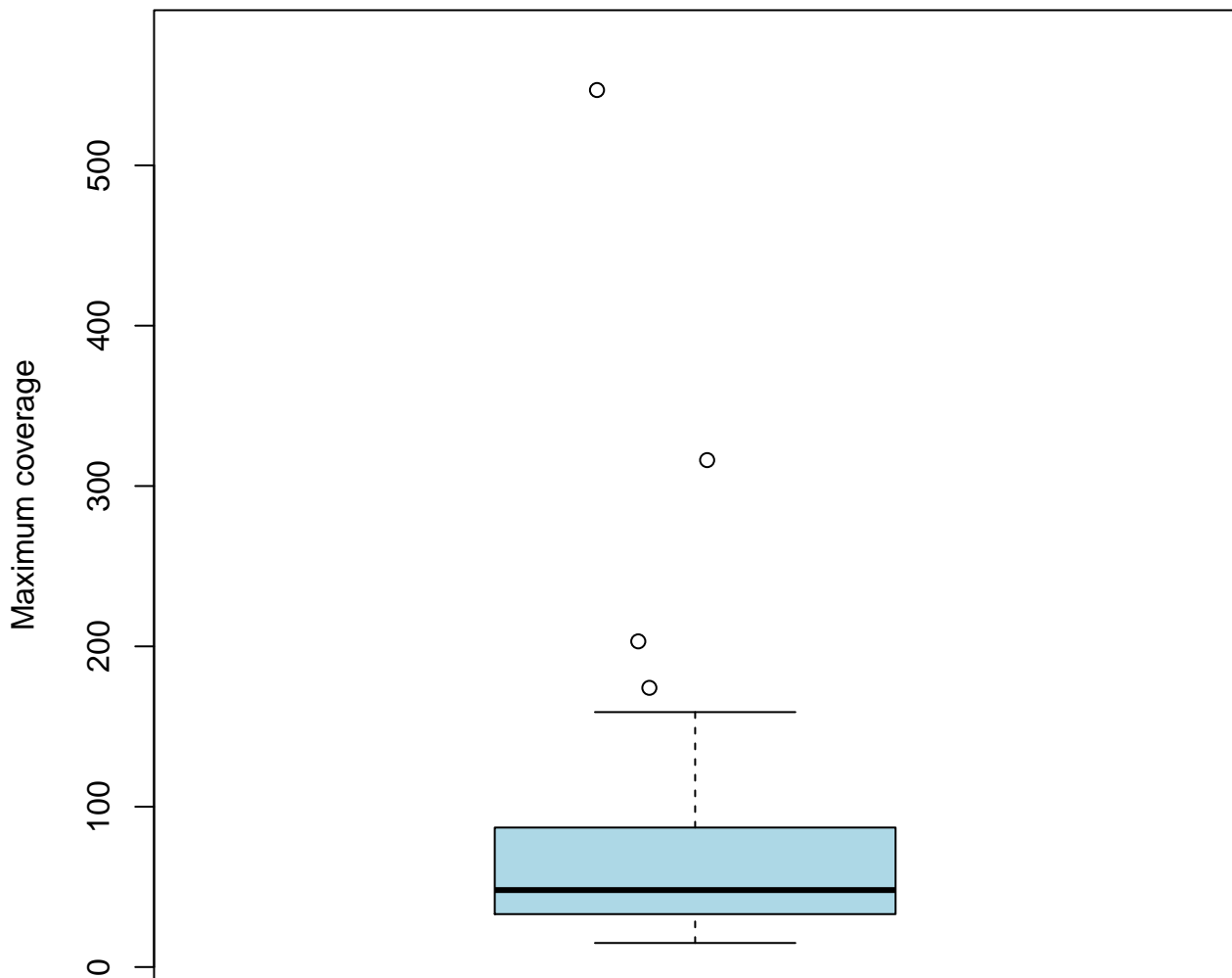
**jx in Intropolis but not "annotated" by knownGene:
221 samples; n jx: 42**



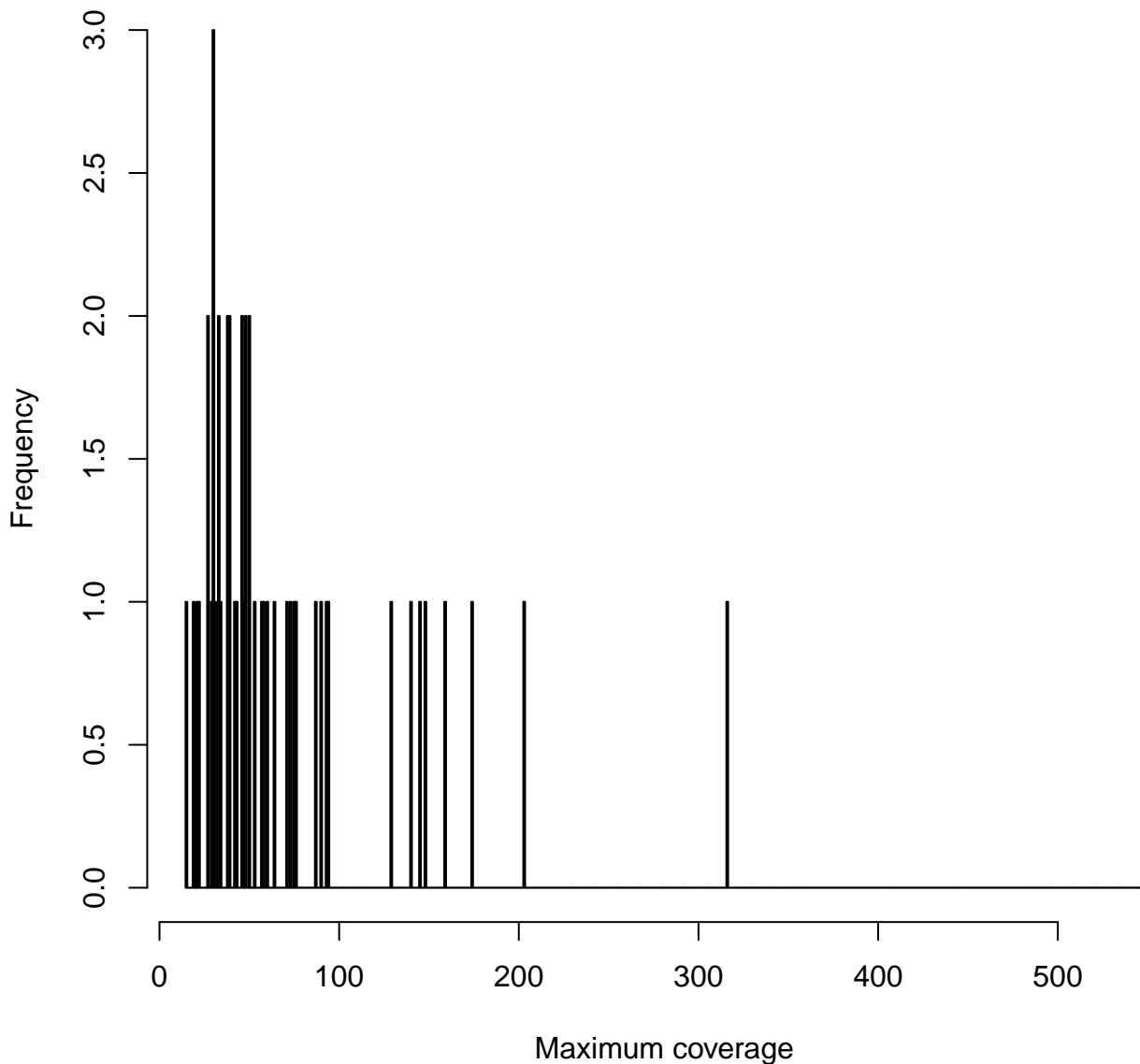
**jx in Intropolis but not "annotated" by knownGene:
221 samples; n jx: 42**



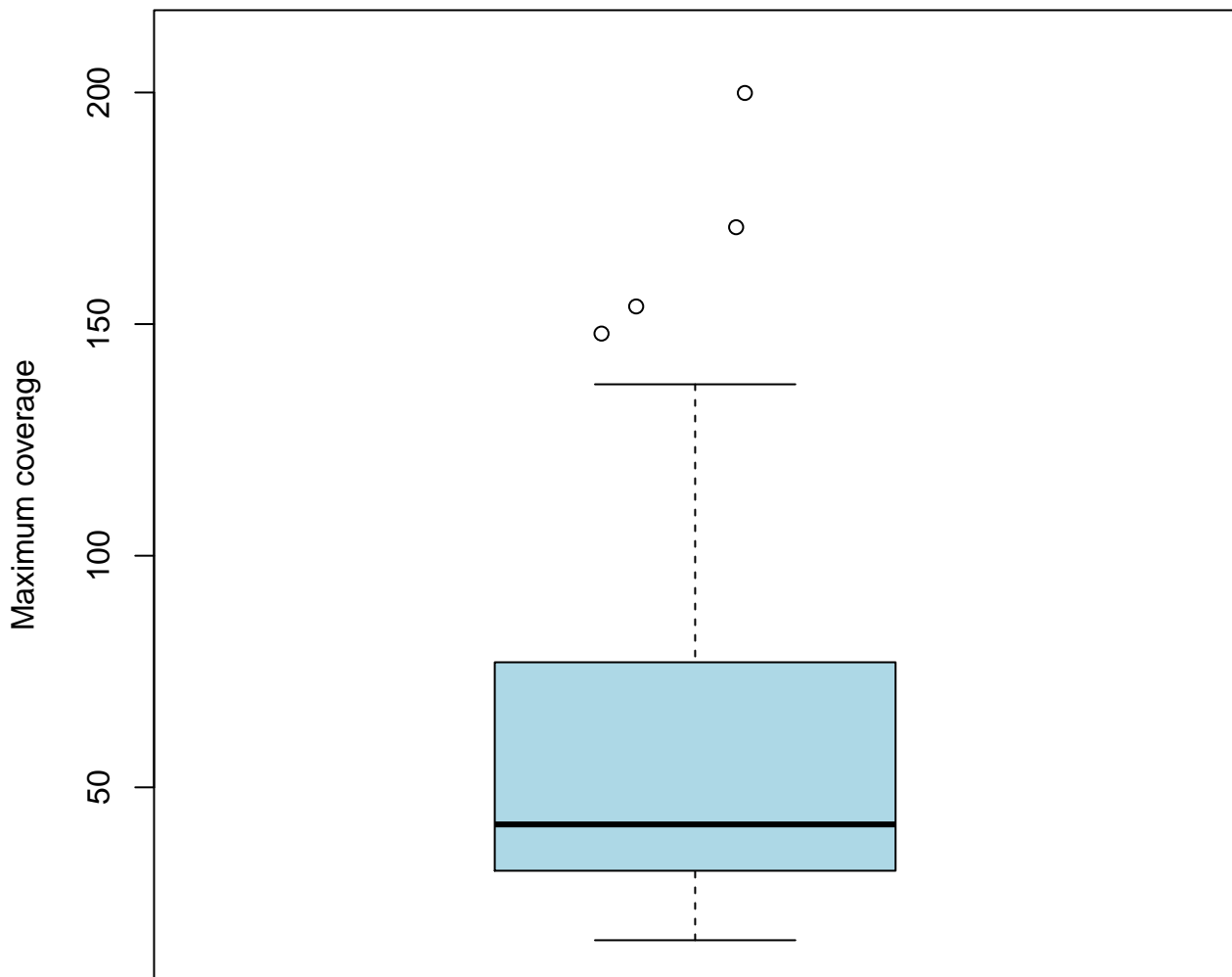
**jx in Intropolis but not "annotated" by knownGene:
222 samples; n jx: 49**



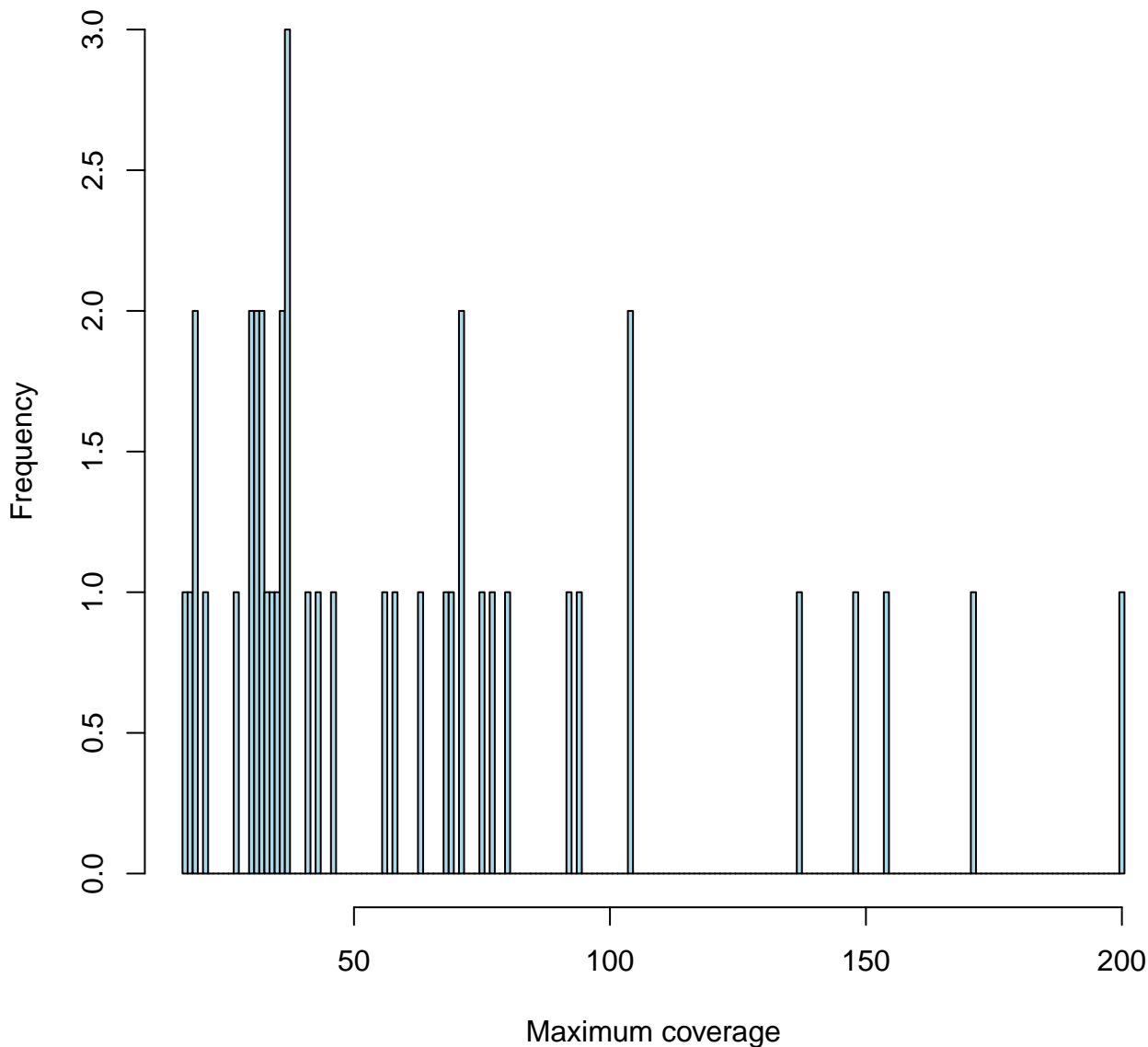
**jx in Intropolis but not "annotated" by knownGene:
222 samples; n jx: 49**



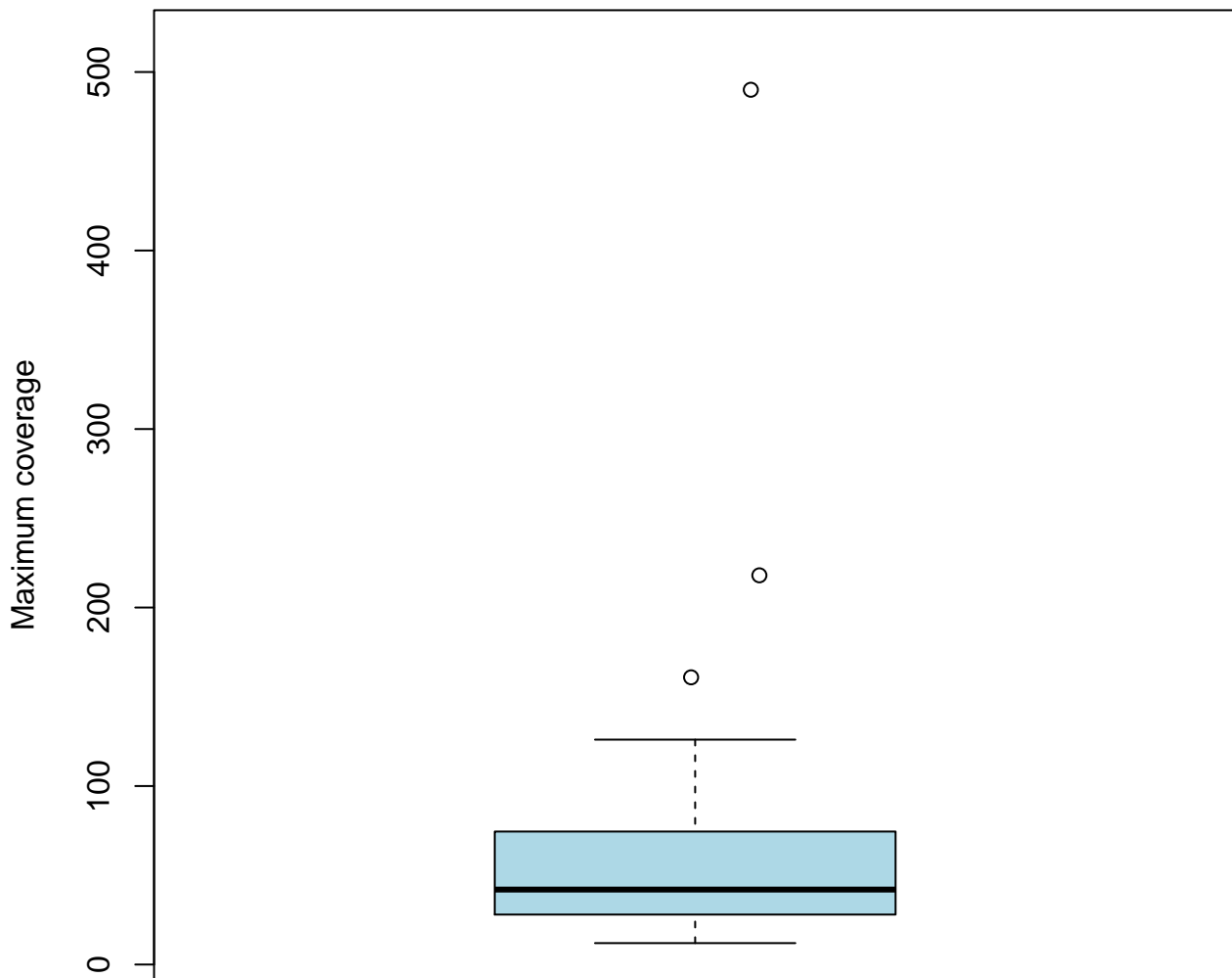
**jx in Intropolis but not "annotated" by knownGene:
223 samples; n jx: 42**



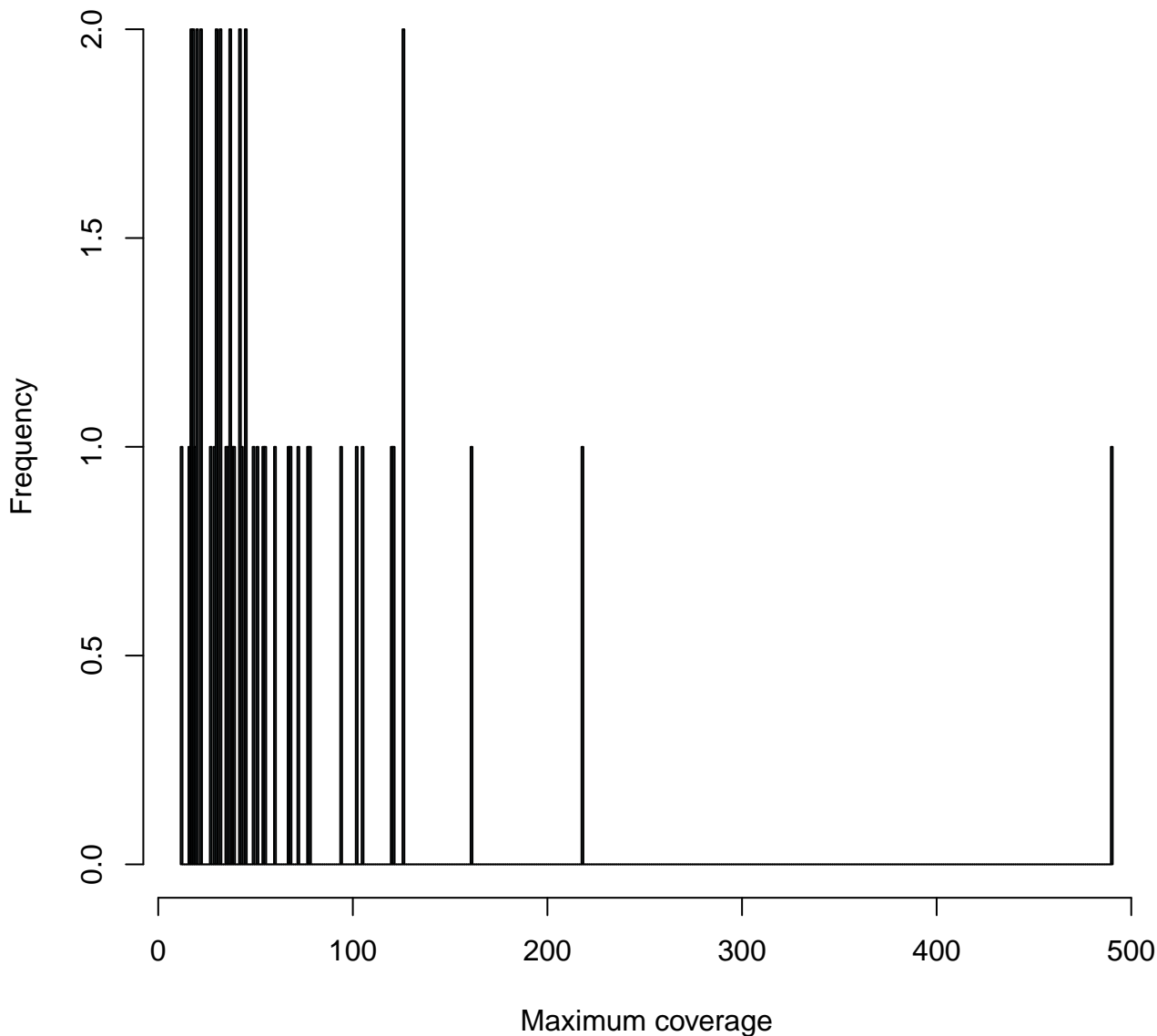
**jx in Intropolis but not "annotated" by knownGene:
223 samples; n jx: 42**



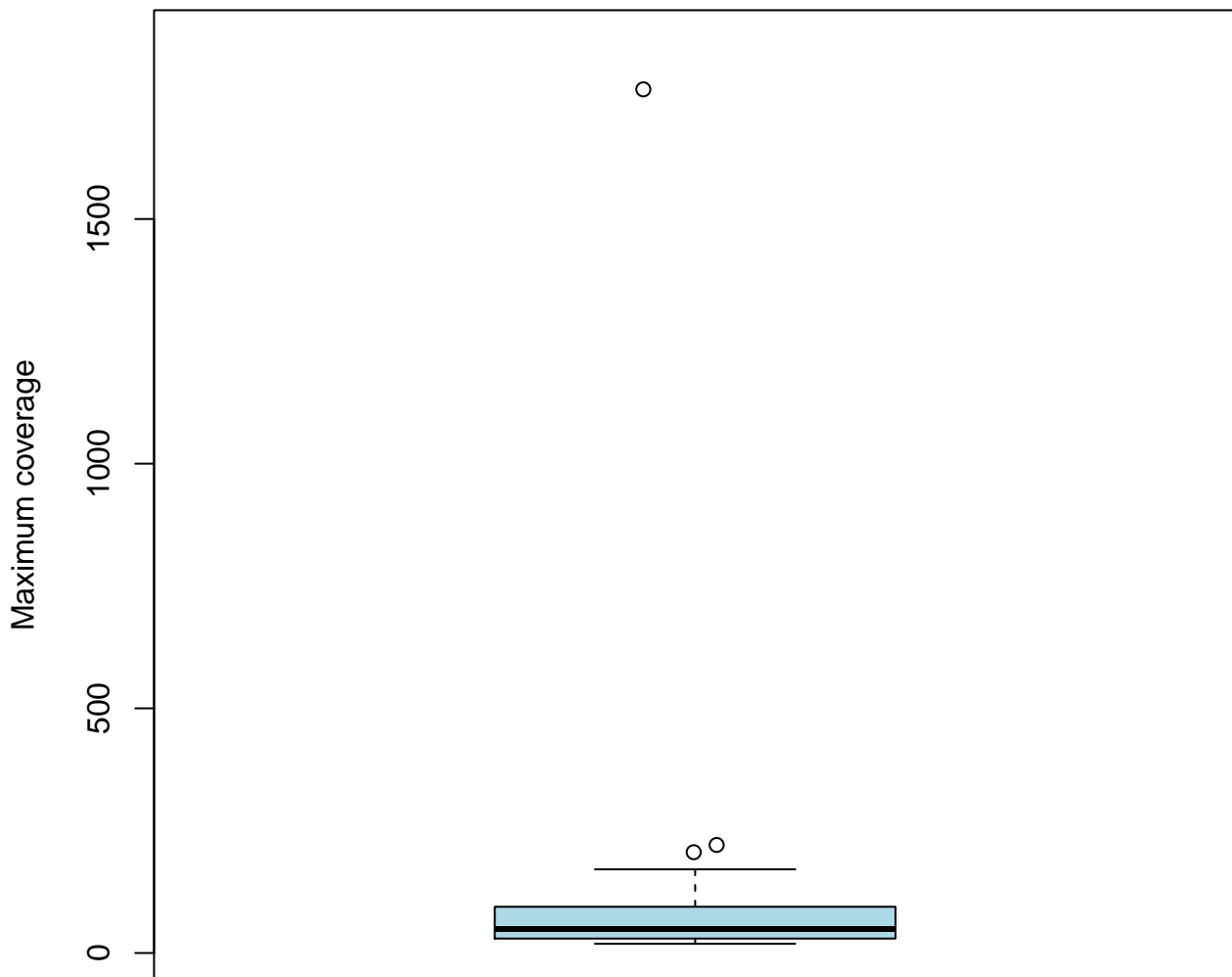
**jx in Intropolis but not "annotated" by knownGene:
224 samples; n jx: 48**



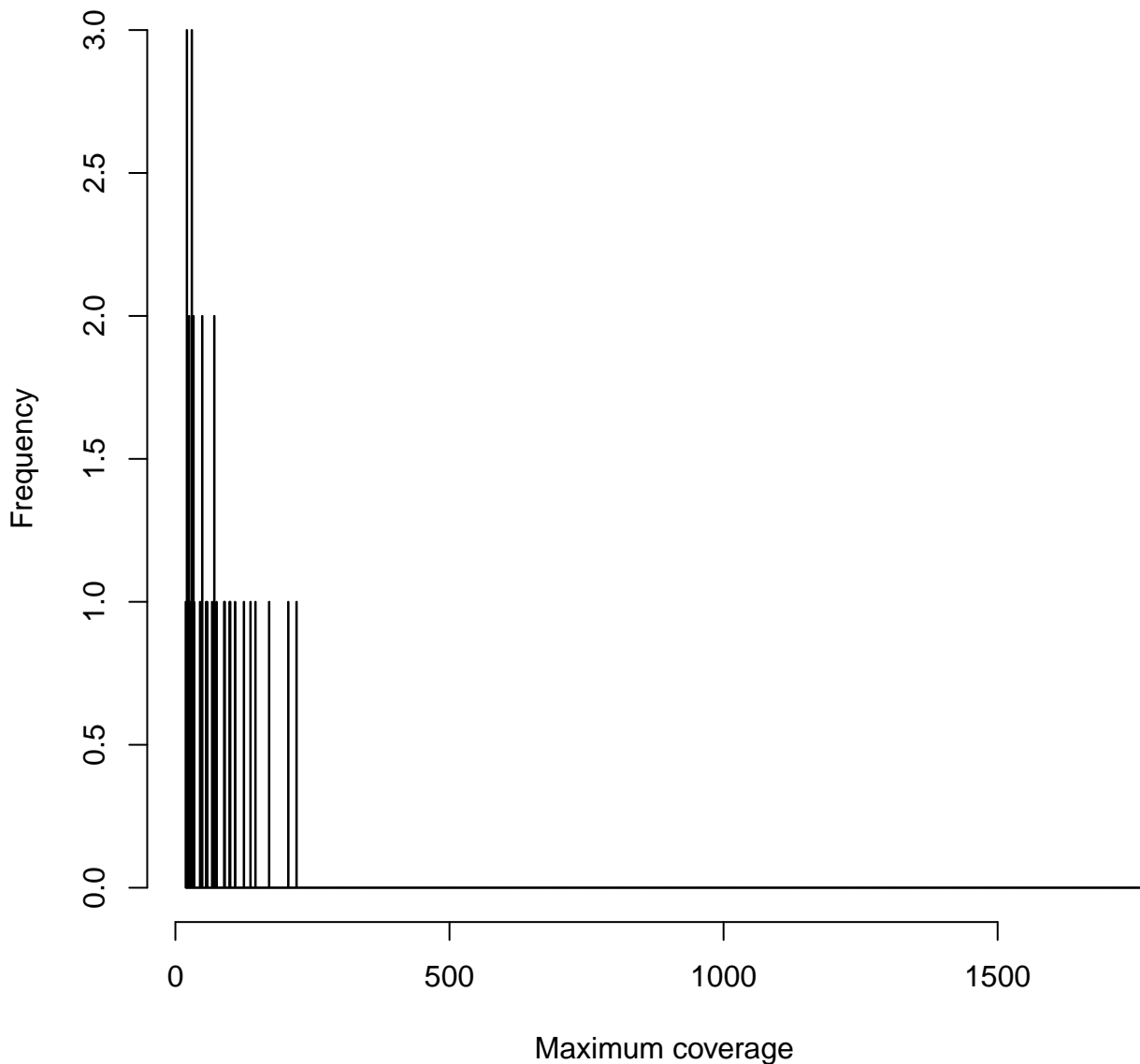
**jx in Intropolis but not "annotated" by knownGene:
224 samples; n jx: 48**



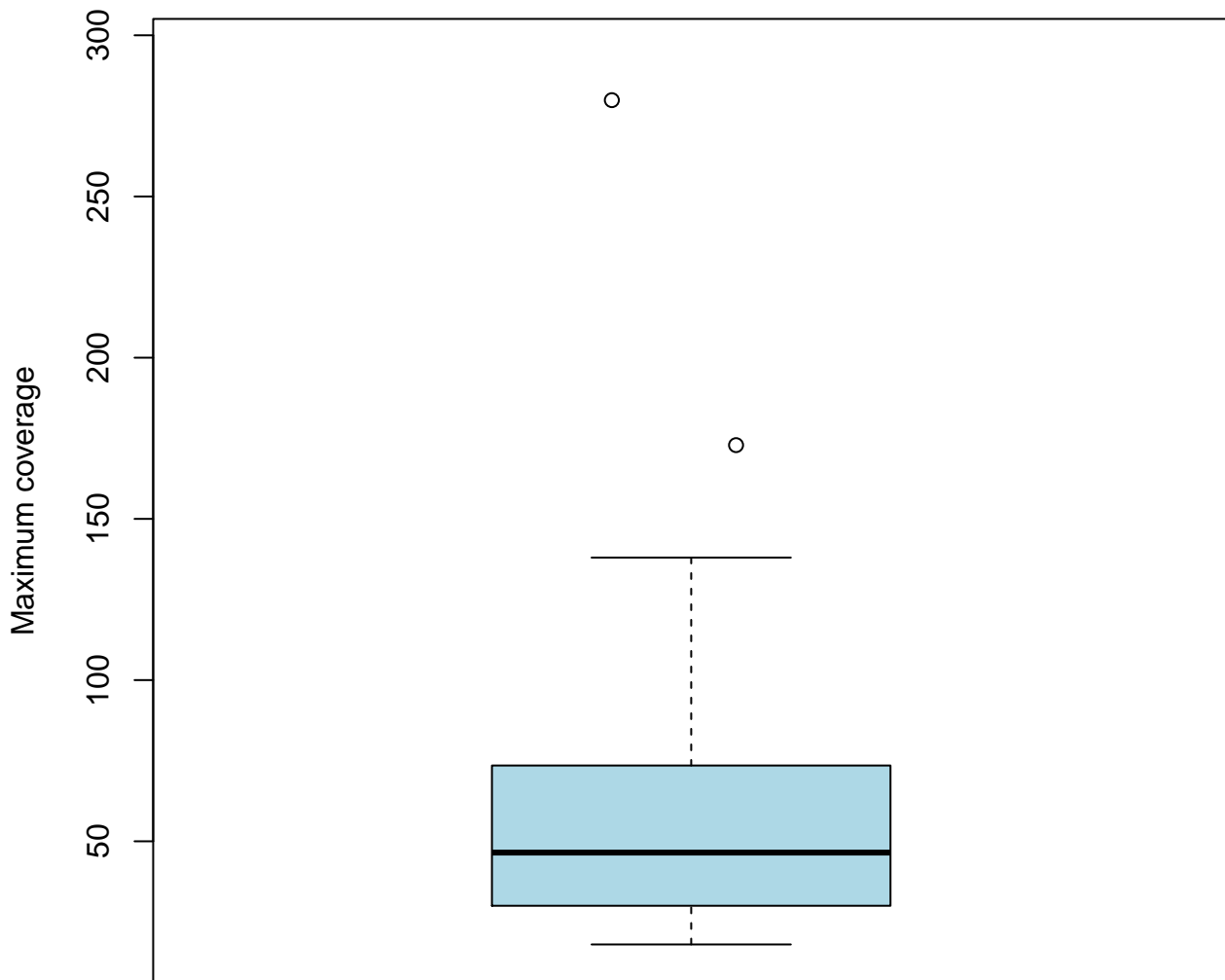
**jx in Intropolis but not "annotated" by knownGene:
225 samples; n jx: 39**



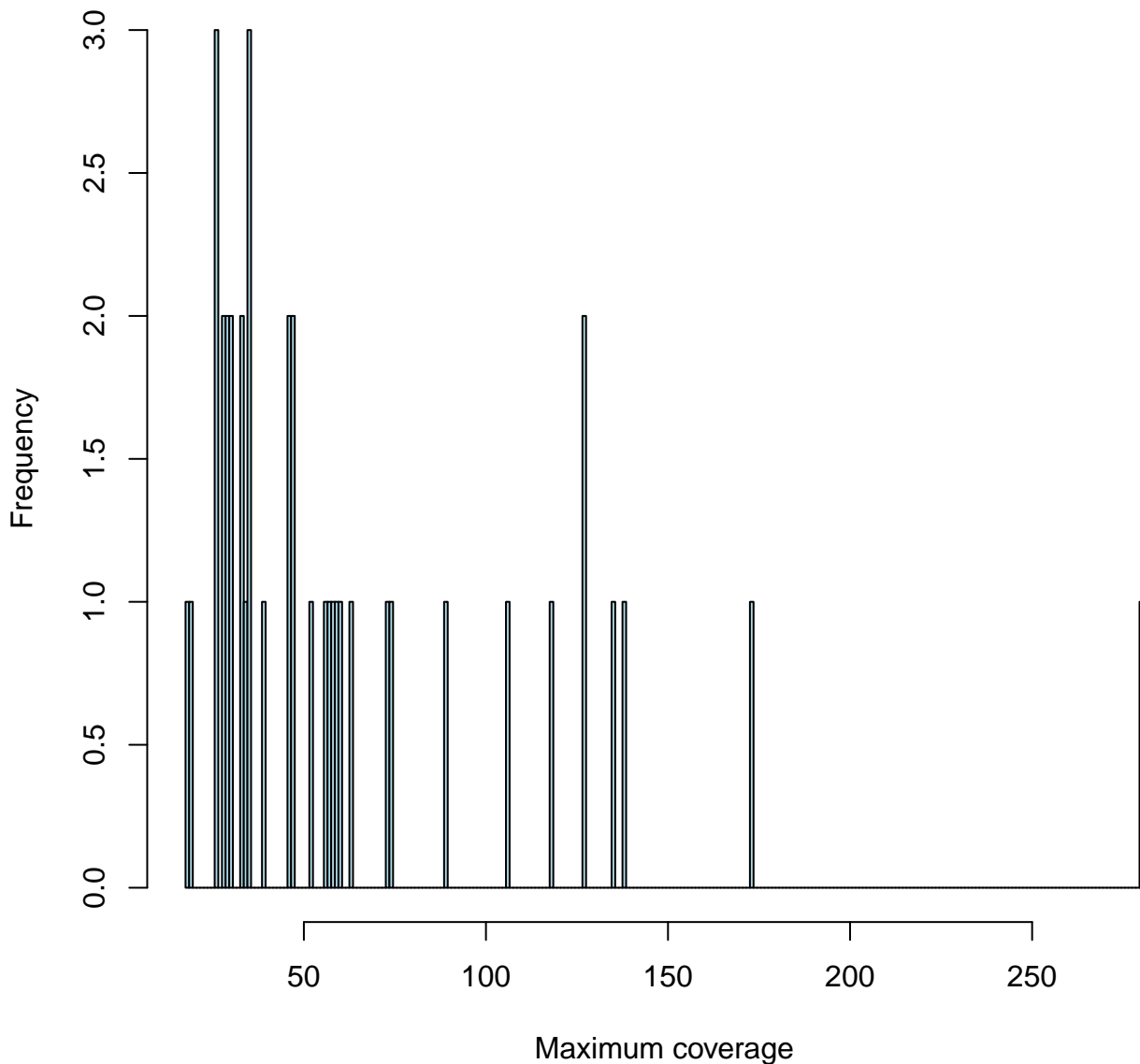
**jx in Intropolis but not "annotated" by knownGene:
225 samples; n jx: 39**



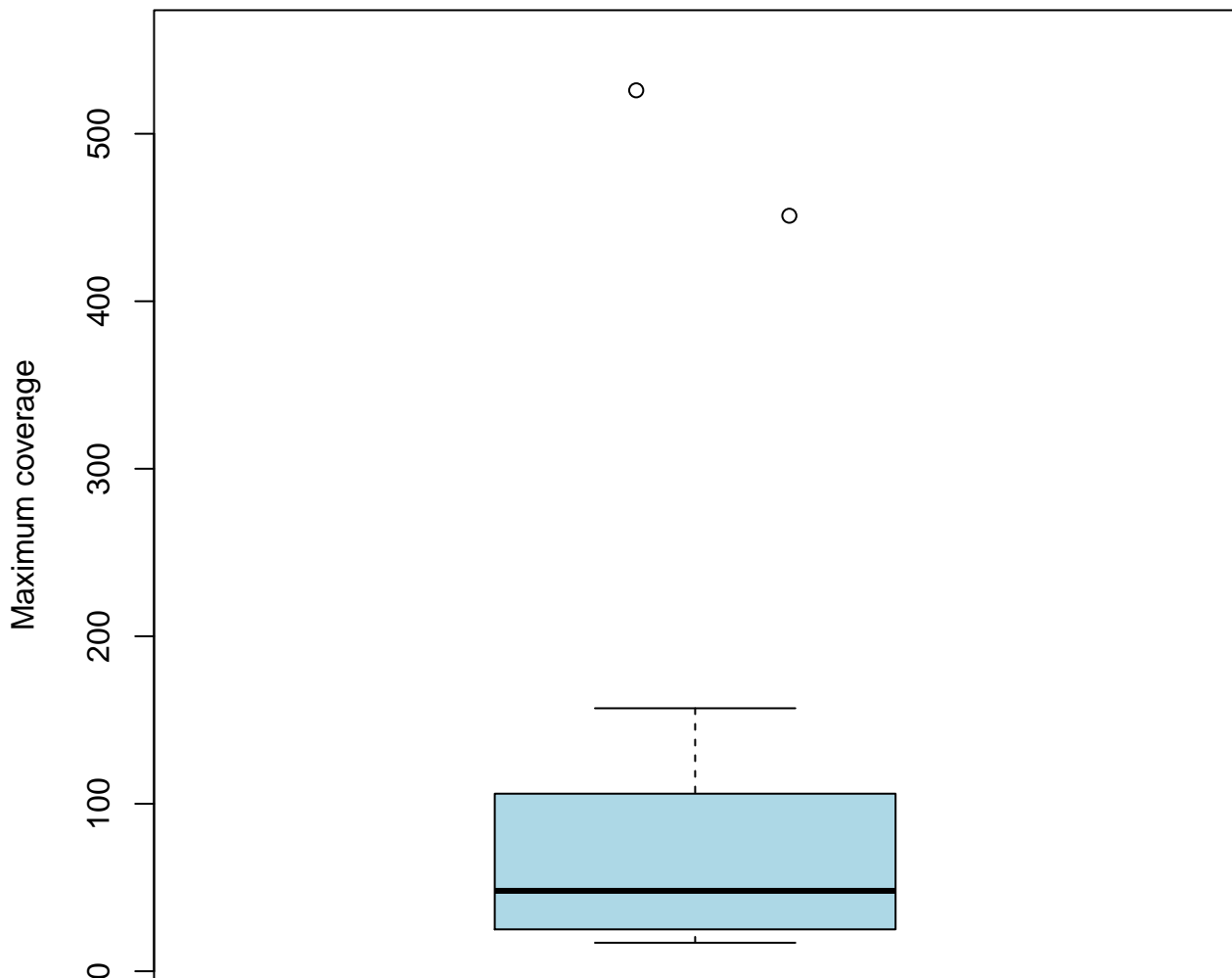
**jx in Intropolis but not "annotated" by knownGene:
226 samples; n jx: 40**



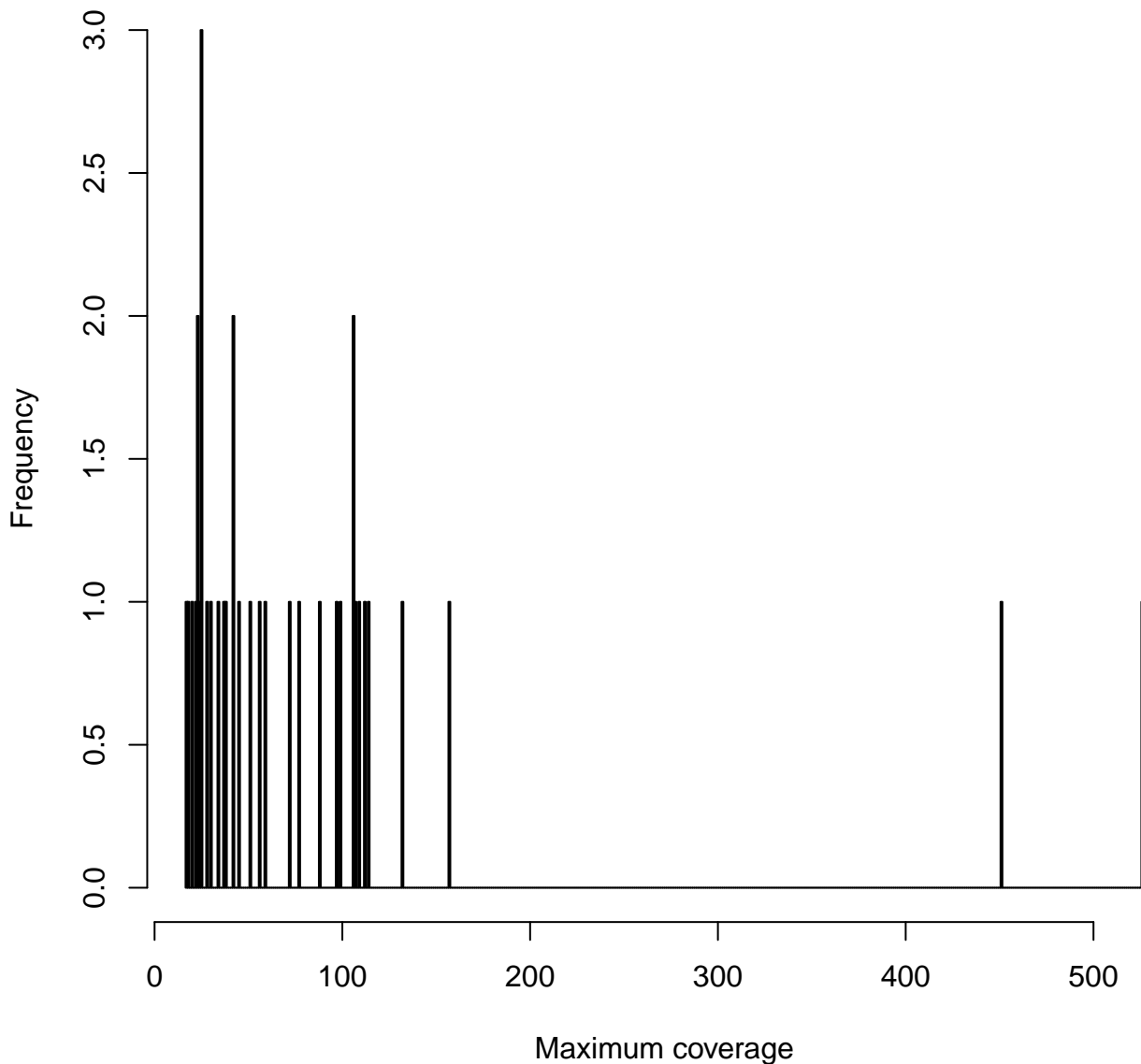
**jx in Intropolis but not "annotated" by knownGene:
226 samples; n jx: 40**



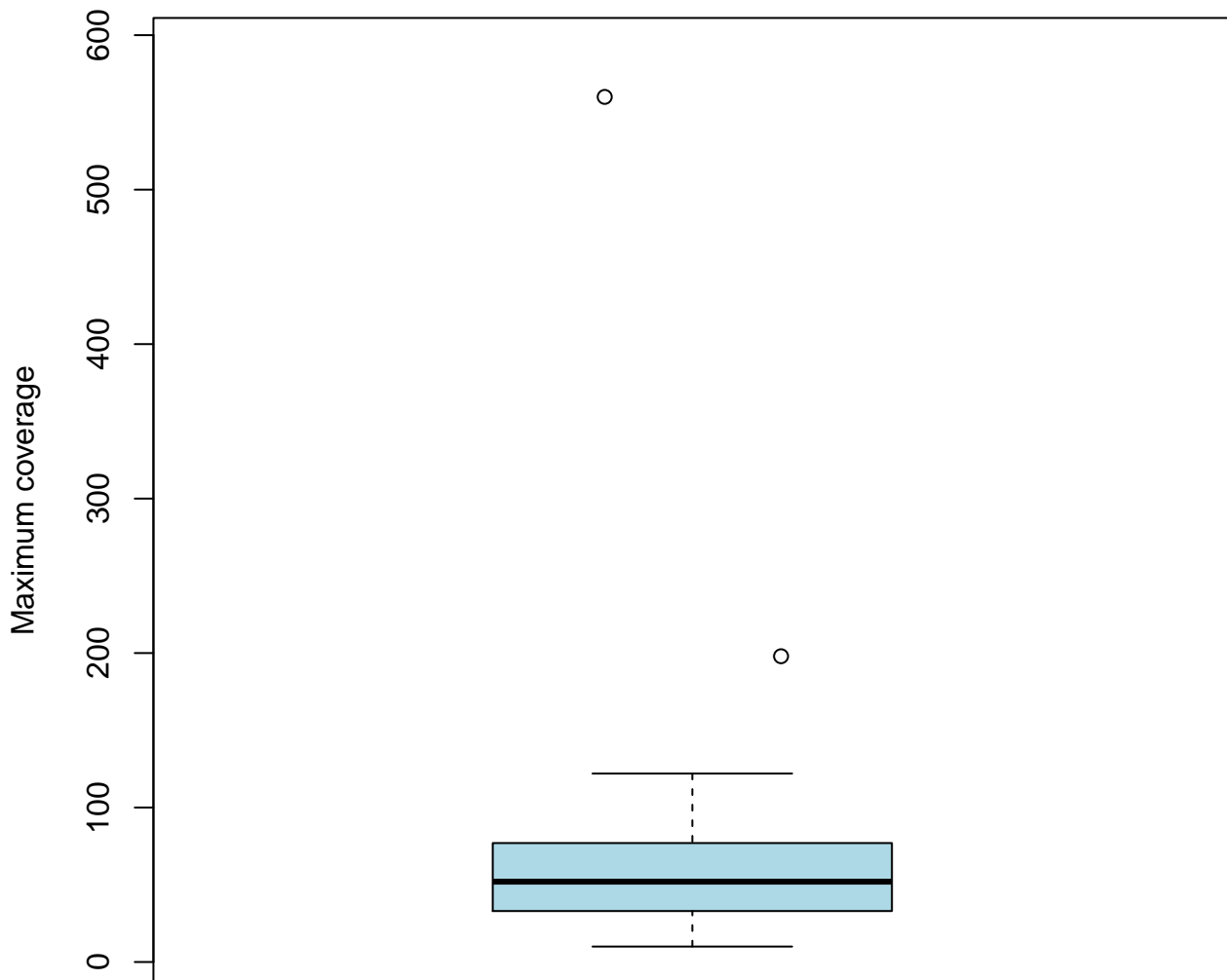
**jx in Intropolis but not "annotated" by knownGene:
227 samples; n jx: 36**



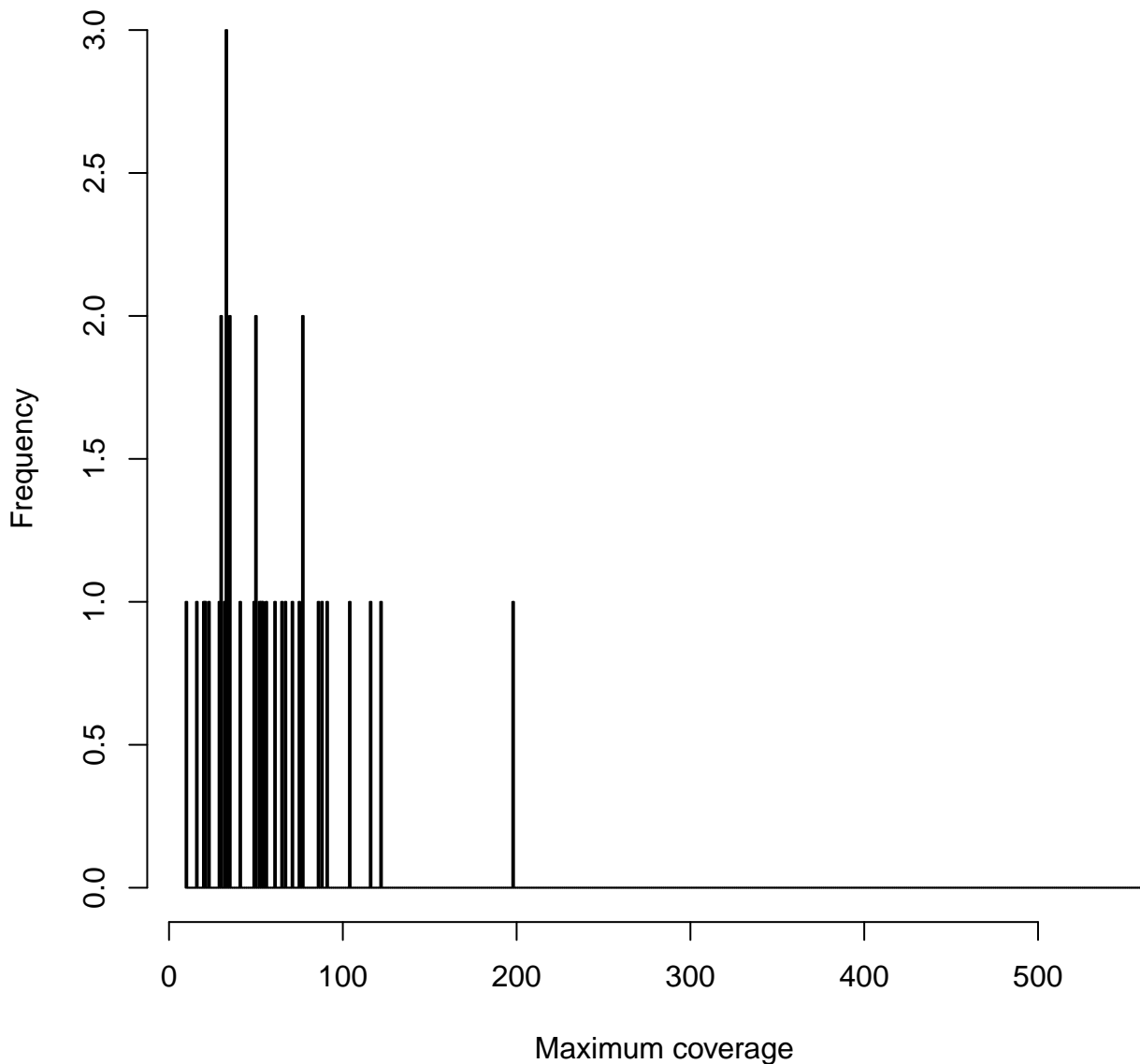
**jx in Intropolis but not "annotated" by knownGene:
227 samples; n jx: 36**



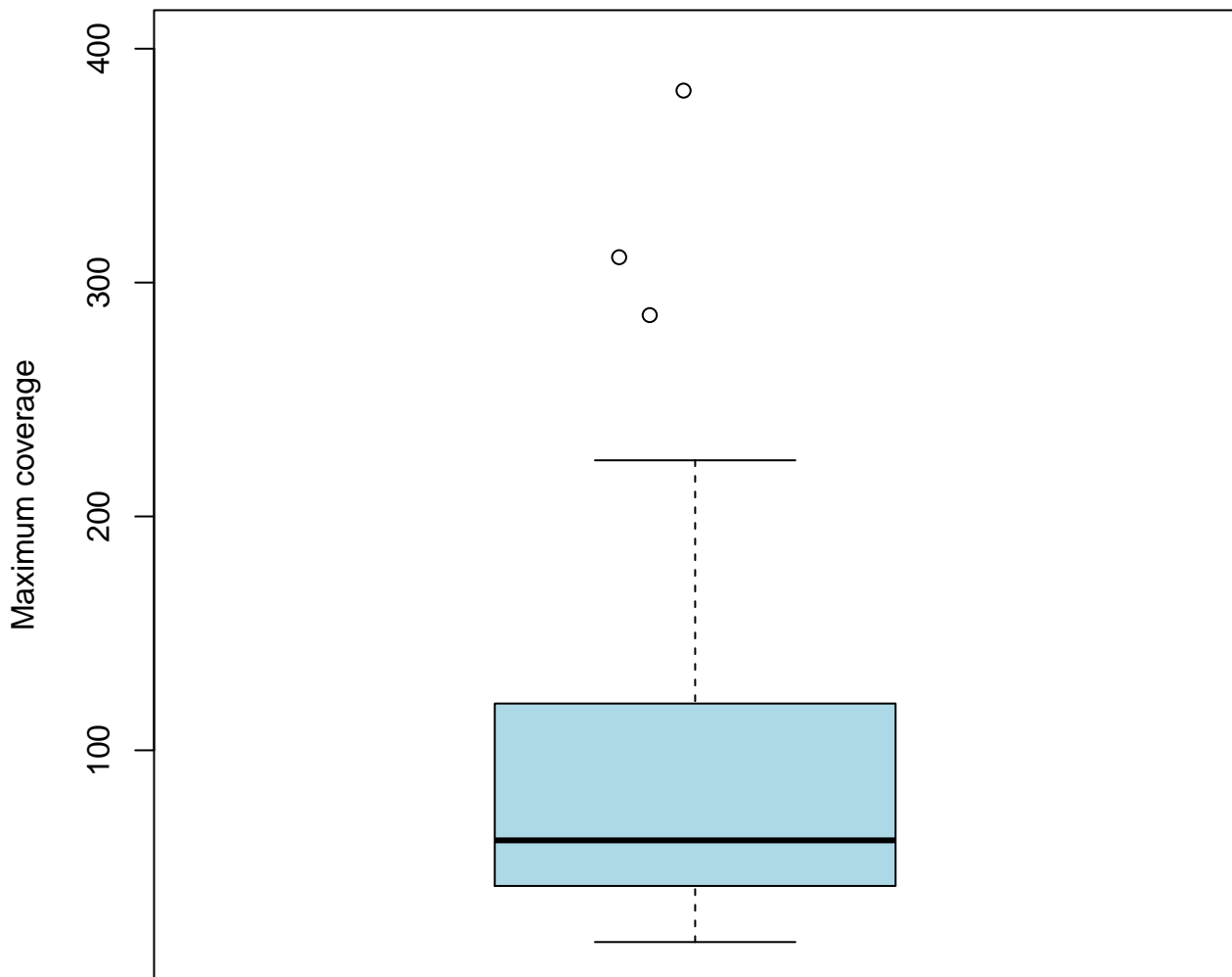
**jx in Intropolis but not "annotated" by knownGene:
228 samples; n jx: 37**



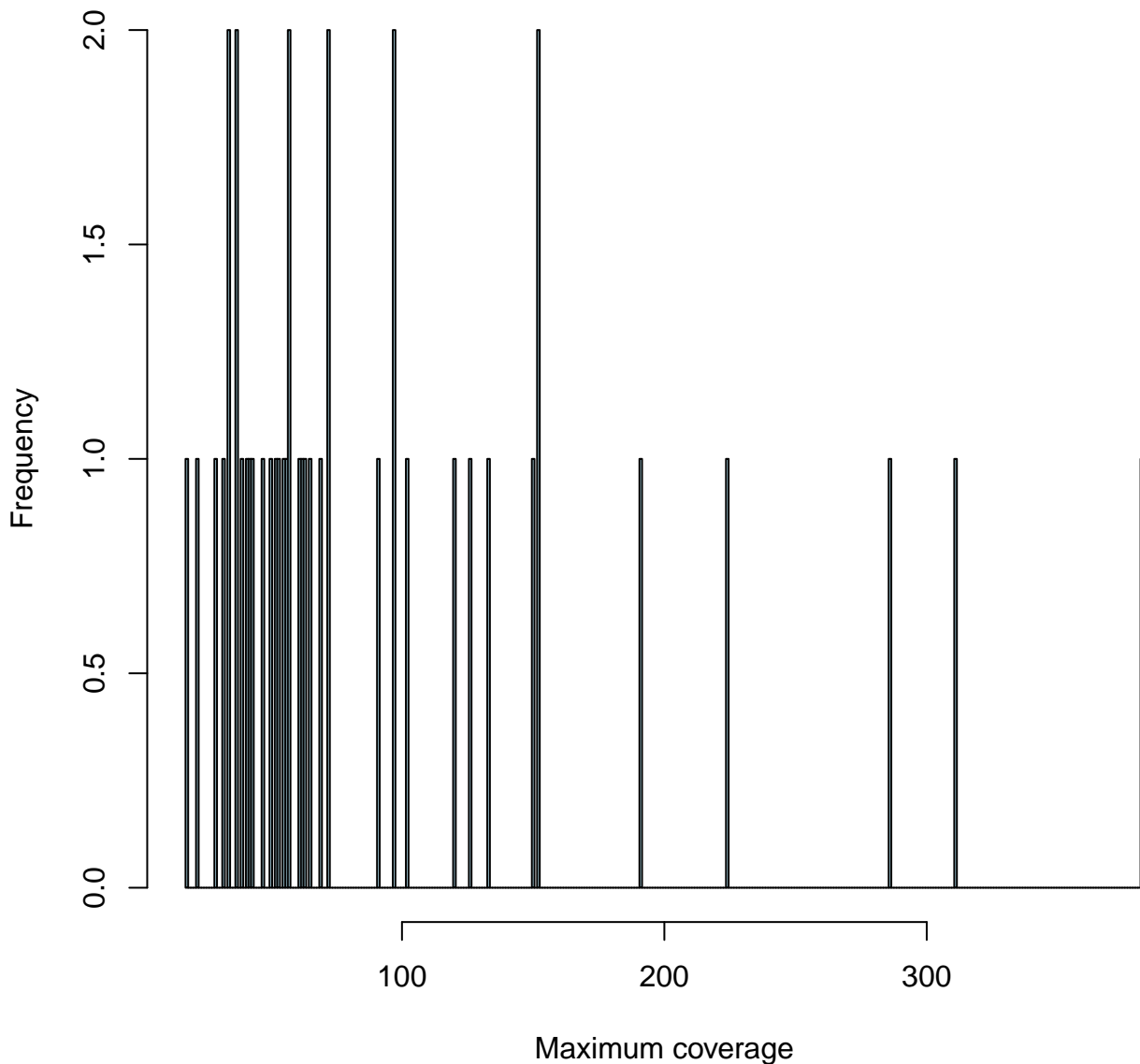
**jx in Intropolis but not "annotated" by knownGene:
228 samples; n jx: 37**



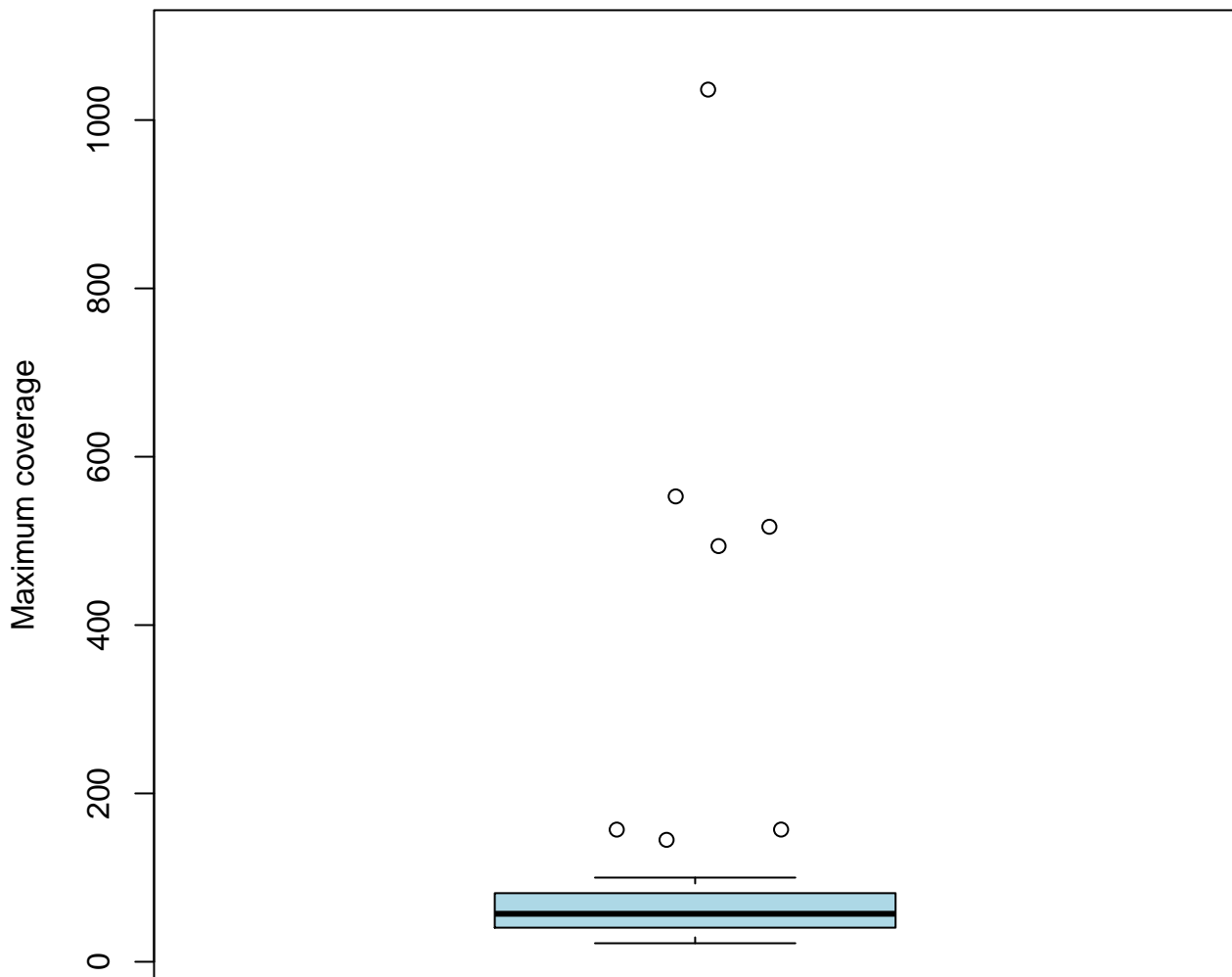
**jx in Intropolis but not "annotated" by knownGene:
229 samples; n jx: 42**



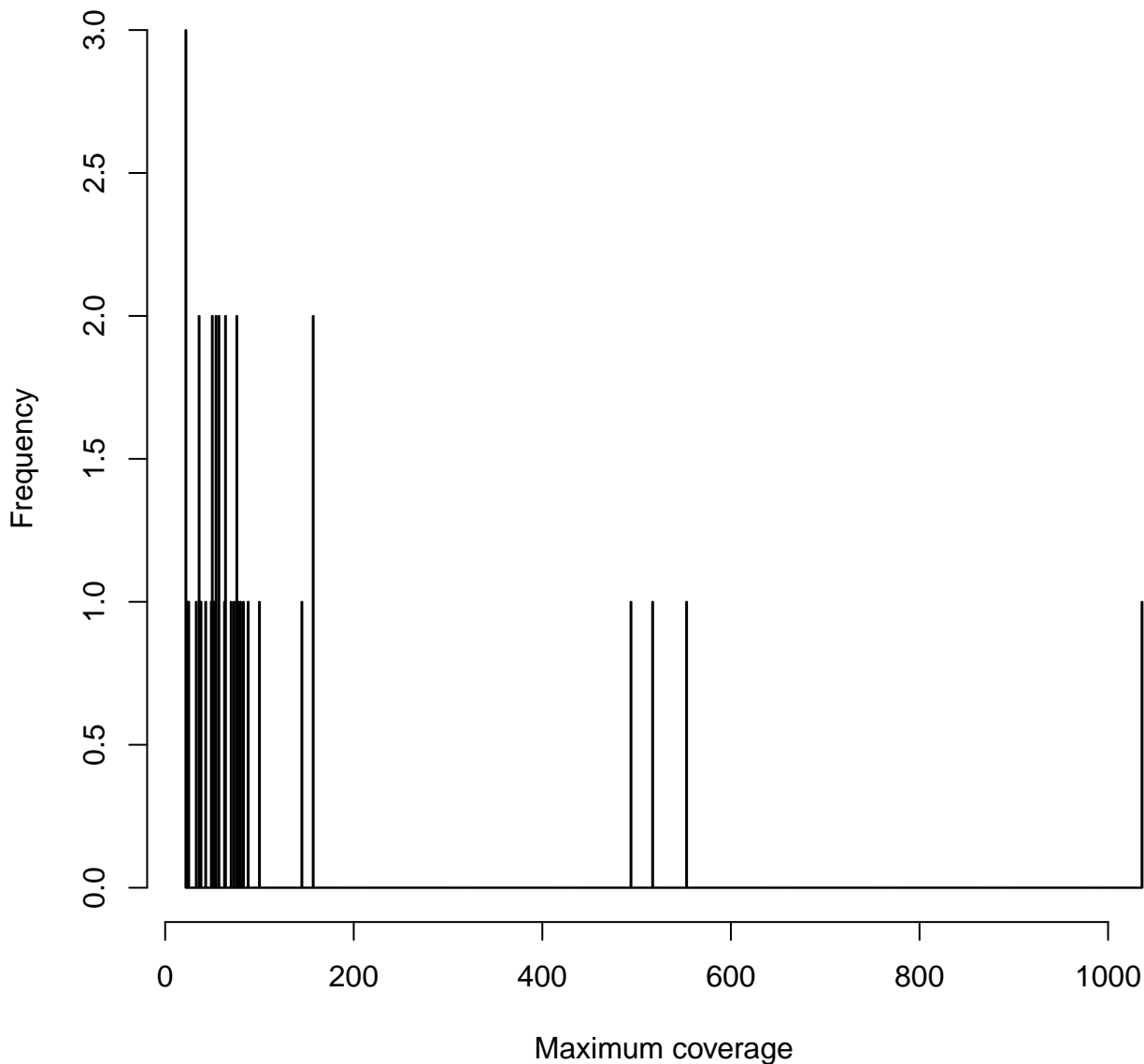
**jx in Intropolis but not "annotated" by knownGene:
229 samples; n jx: 42**



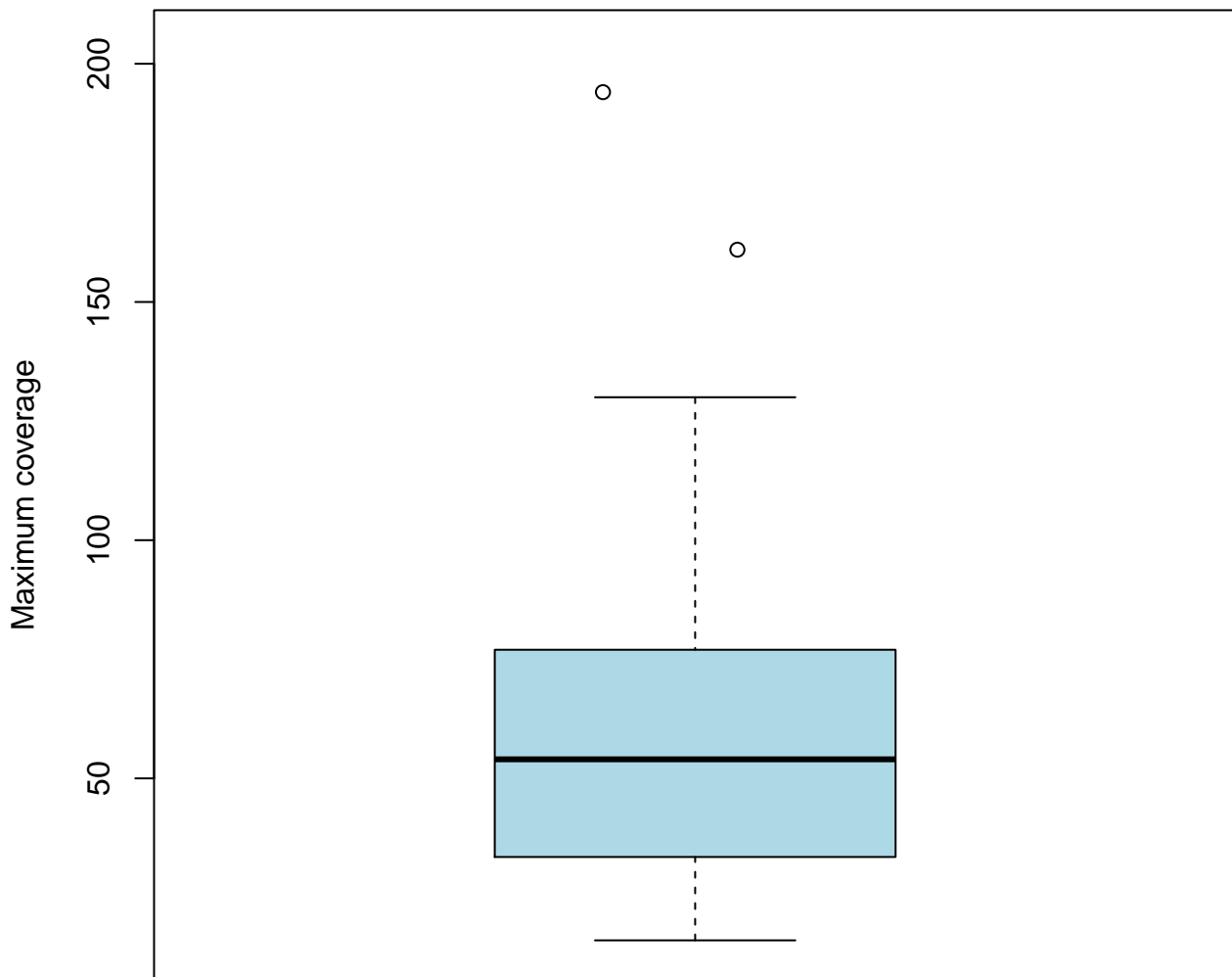
**jx in Intropolis but not "annotated" by knownGene:
230 samples; n jx: 39**



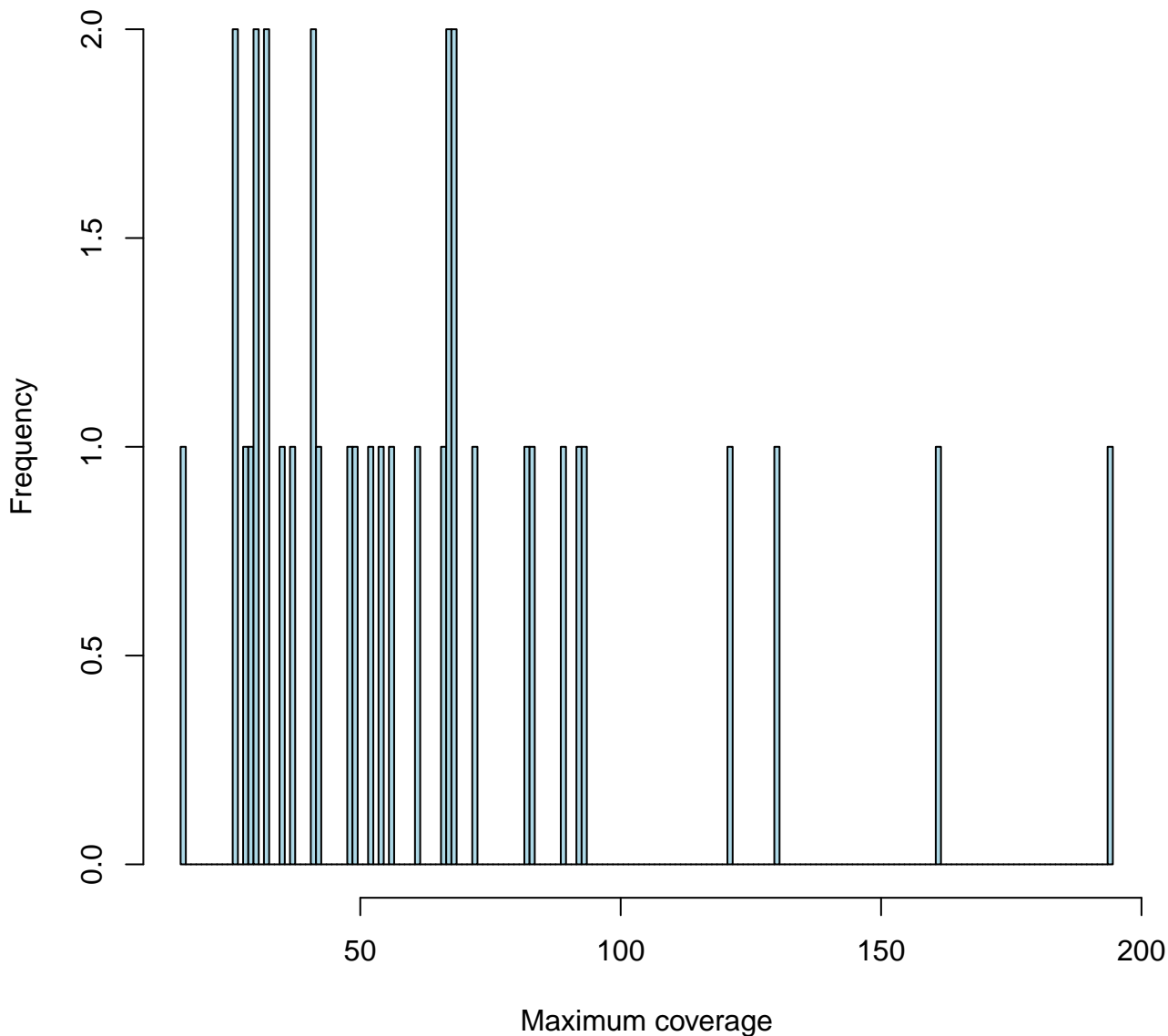
**jx in Intropolis but not "annotated" by knownGene:
230 samples; n jx: 39**



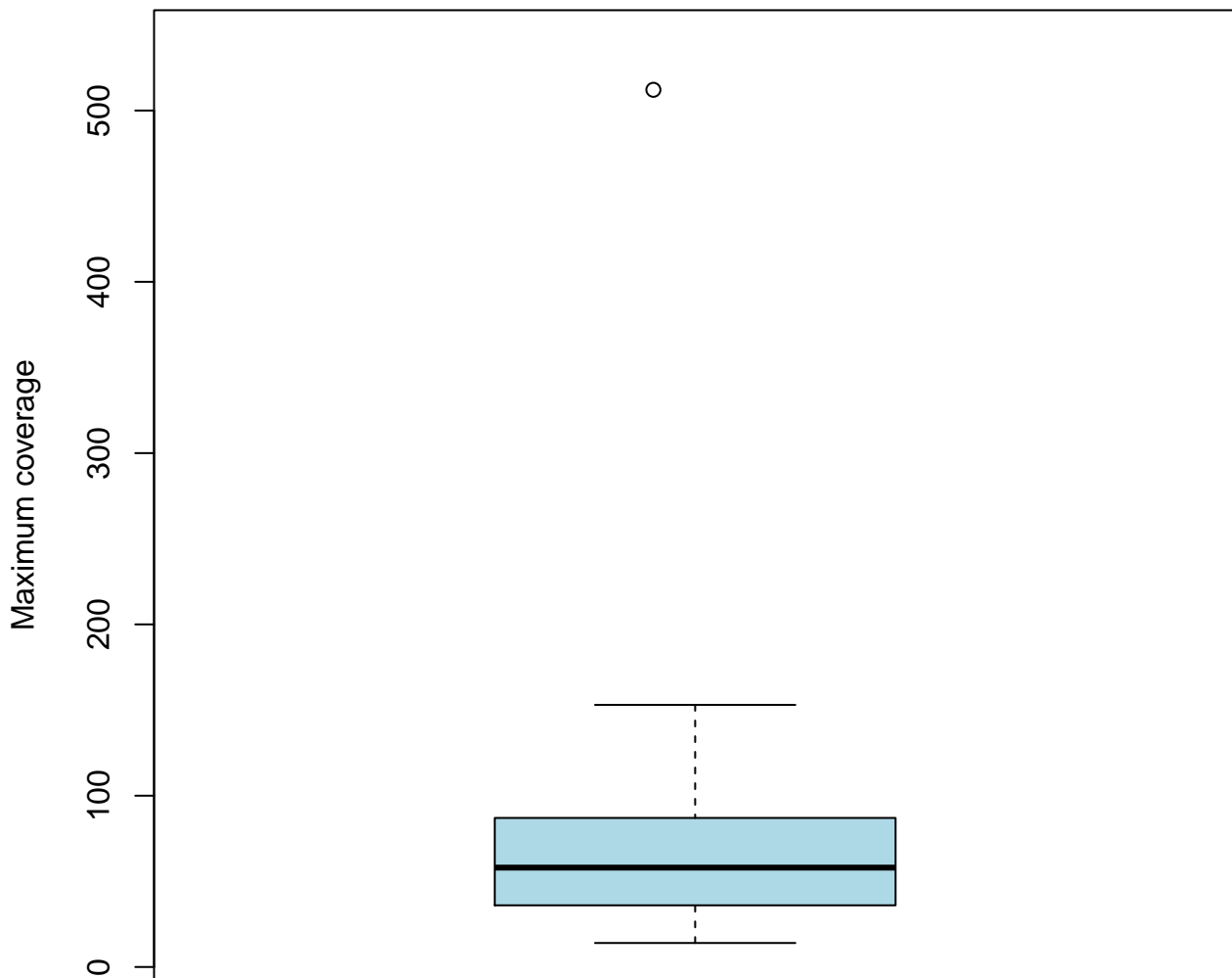
**jx in Intropolis but not "annotated" by knownGene:
231 samples; n jx: 35**



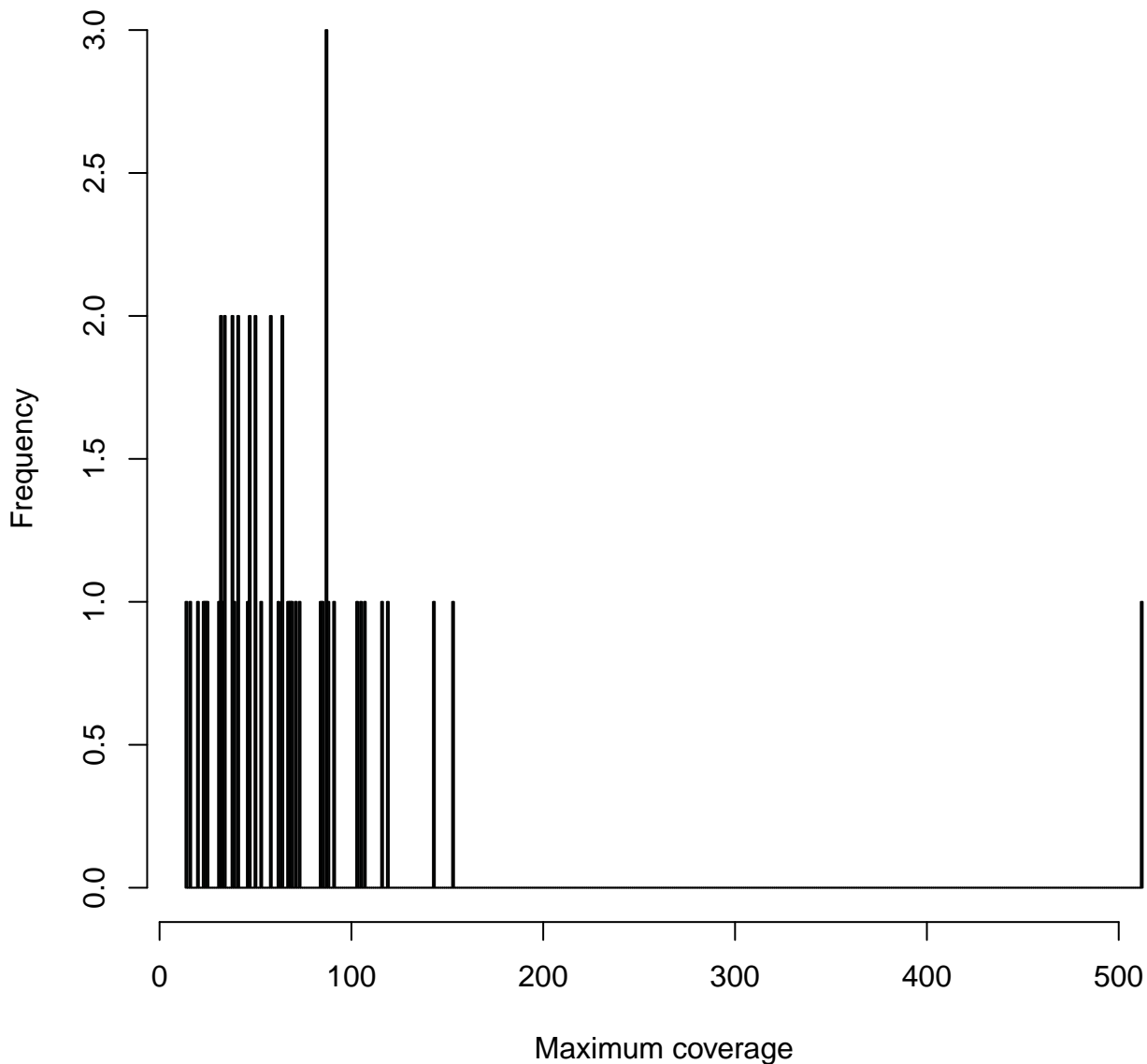
**jx in Intropolis but not "annotated" by knownGene:
231 samples; n jx: 35**



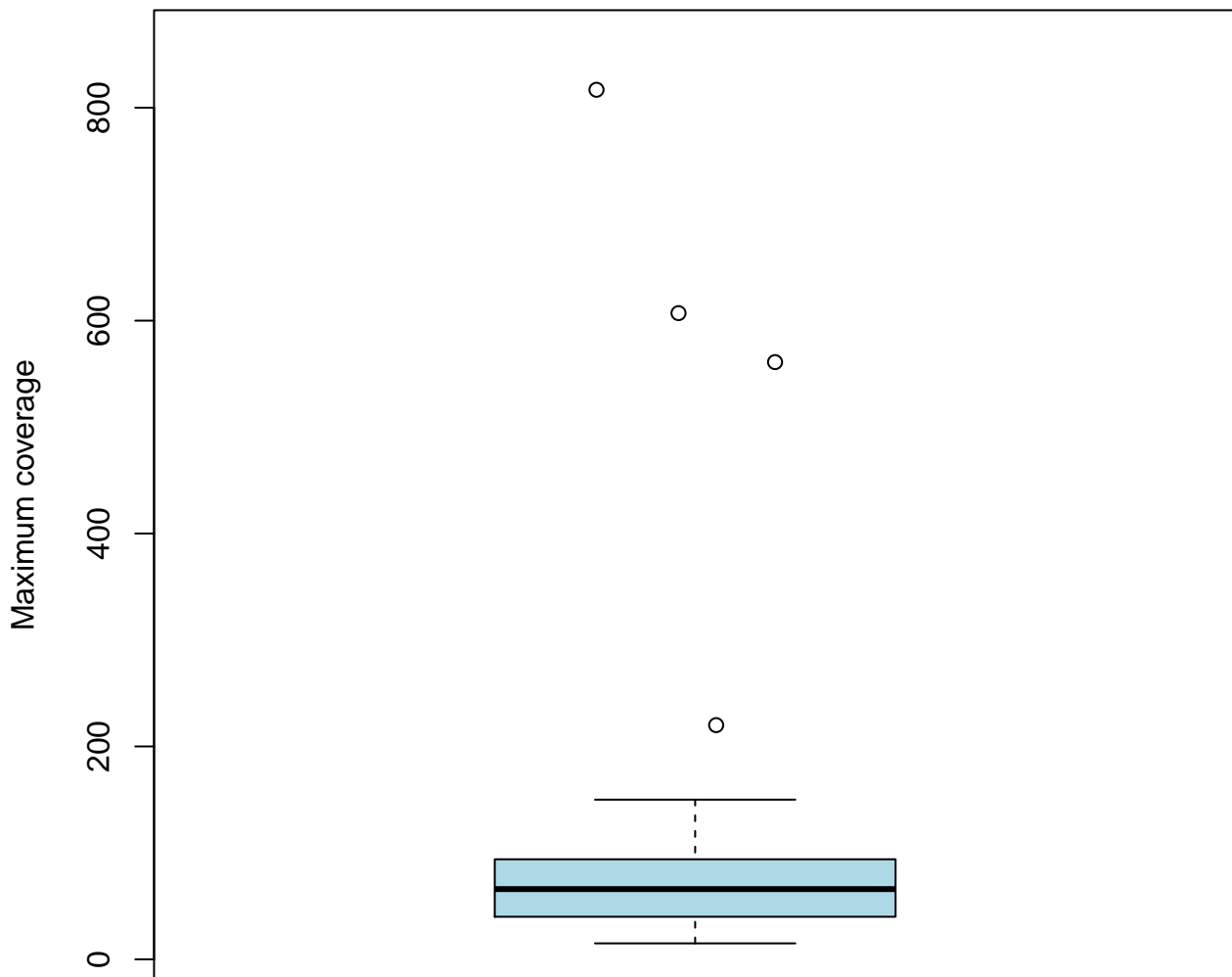
**jx in Intropolis but not "annotated" by knownGene:
232 samples; n jx: 48**



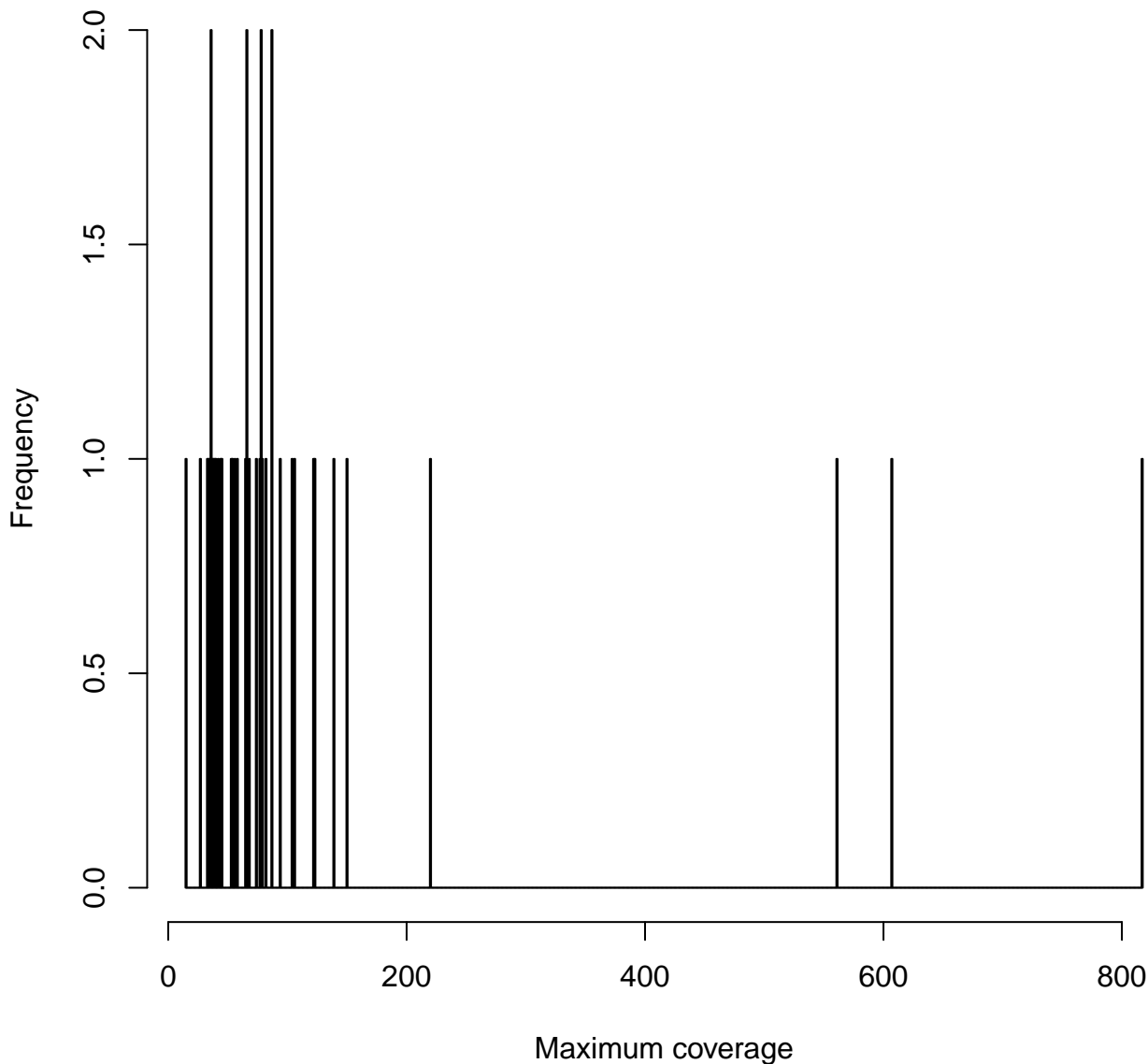
**jx in Intropolis but not "annotated" by knownGene:
232 samples; n jx: 48**



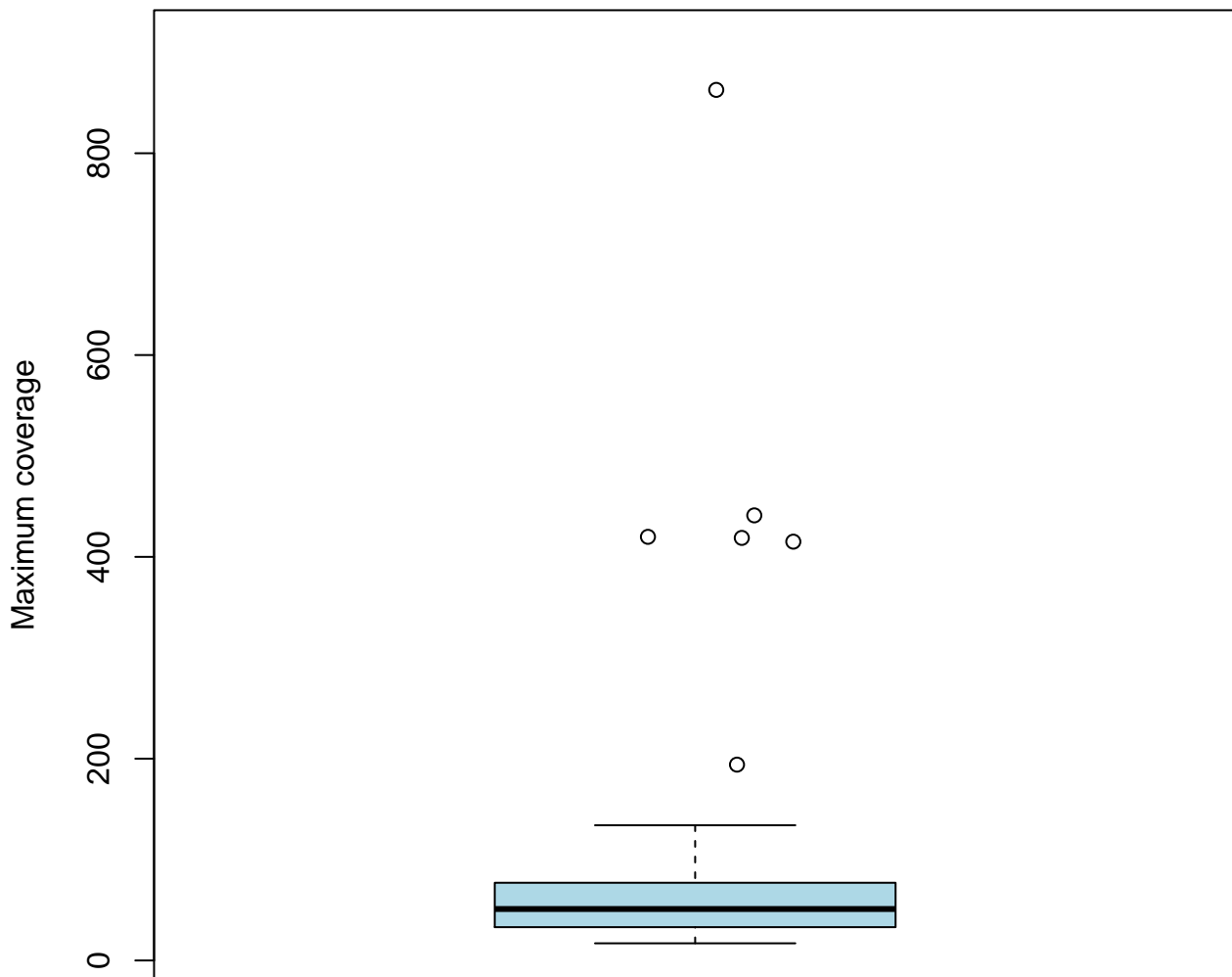
**jx in Intropolis but not "annotated" by knownGene:
233 samples; n jx: 41**



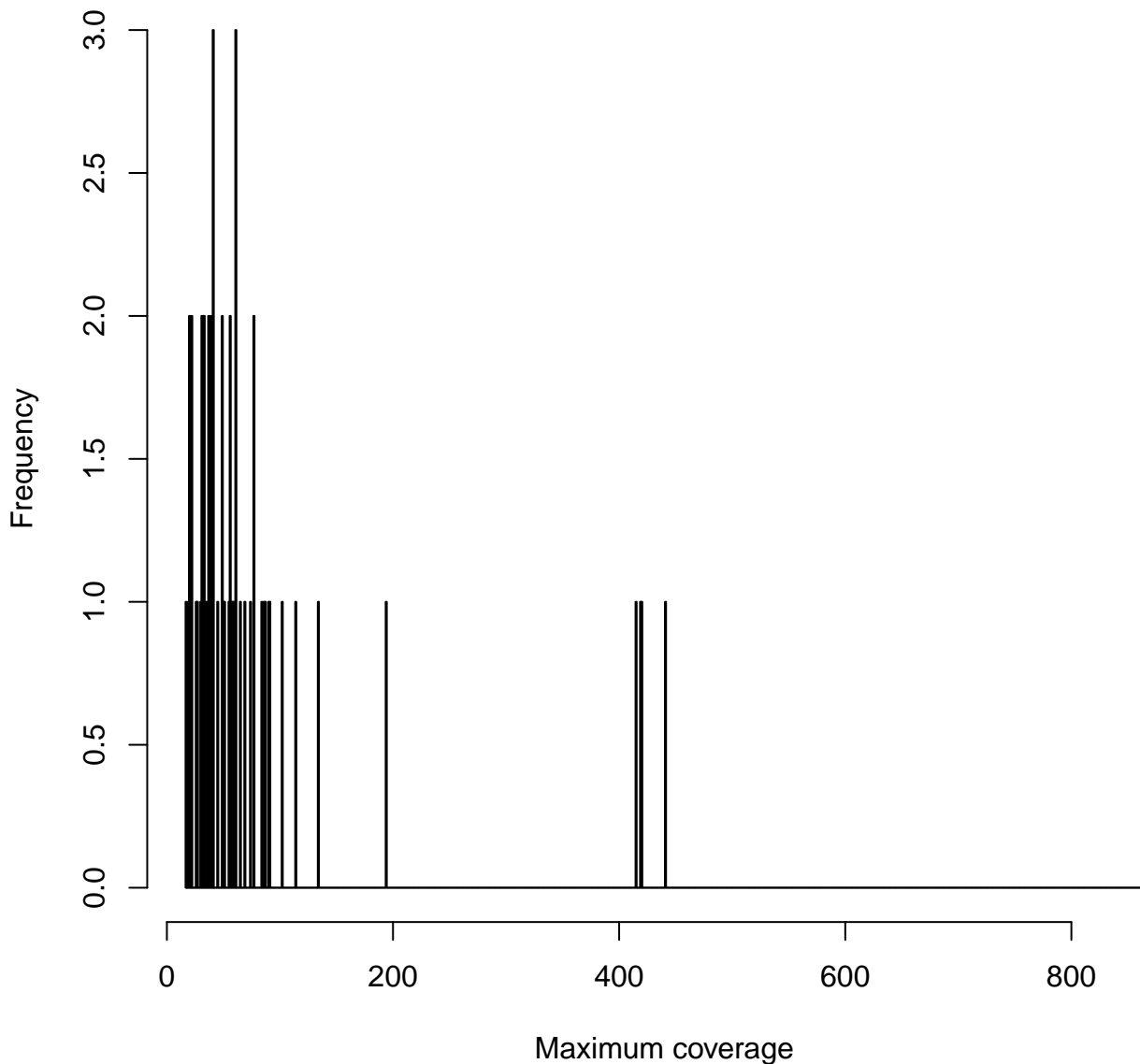
**jx in Intropolis but not "annotated" by knownGene:
233 samples; n jx: 41**



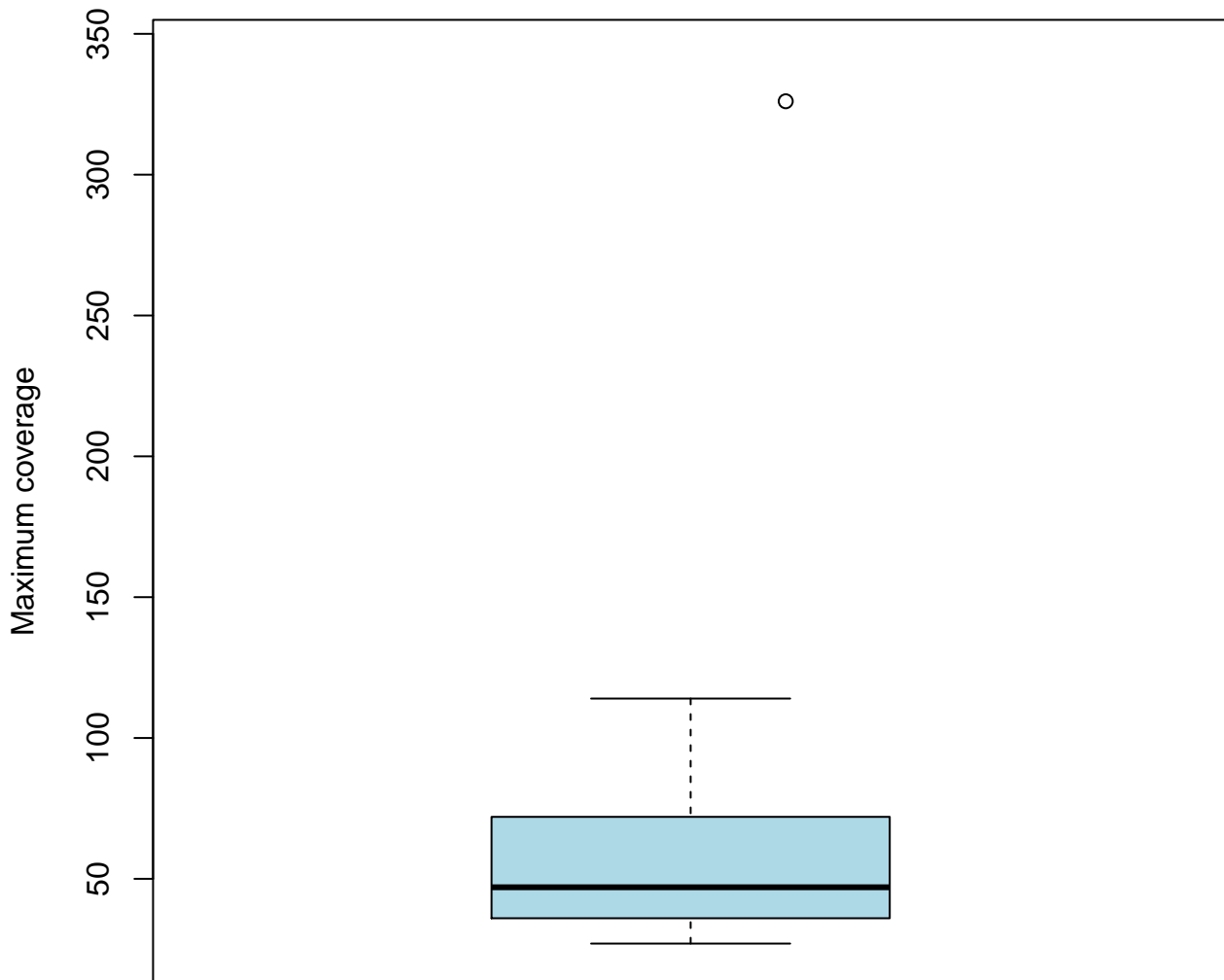
**jx in Intropolis but not "annotated" by knownGene:
234 samples; n jx: 57**



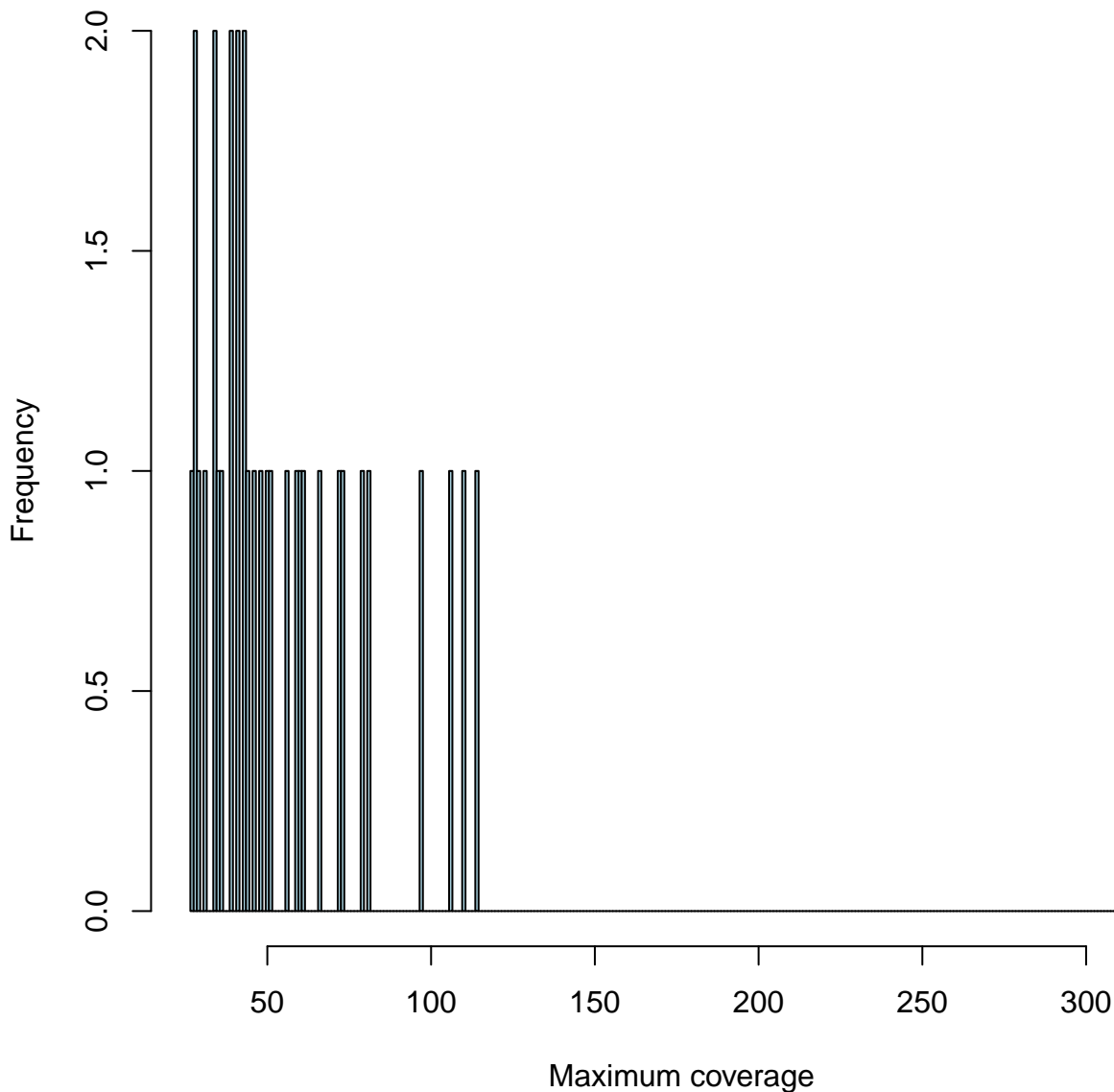
**jx in Intropolis but not "annotated" by knownGene:
234 samples; n jx: 57**



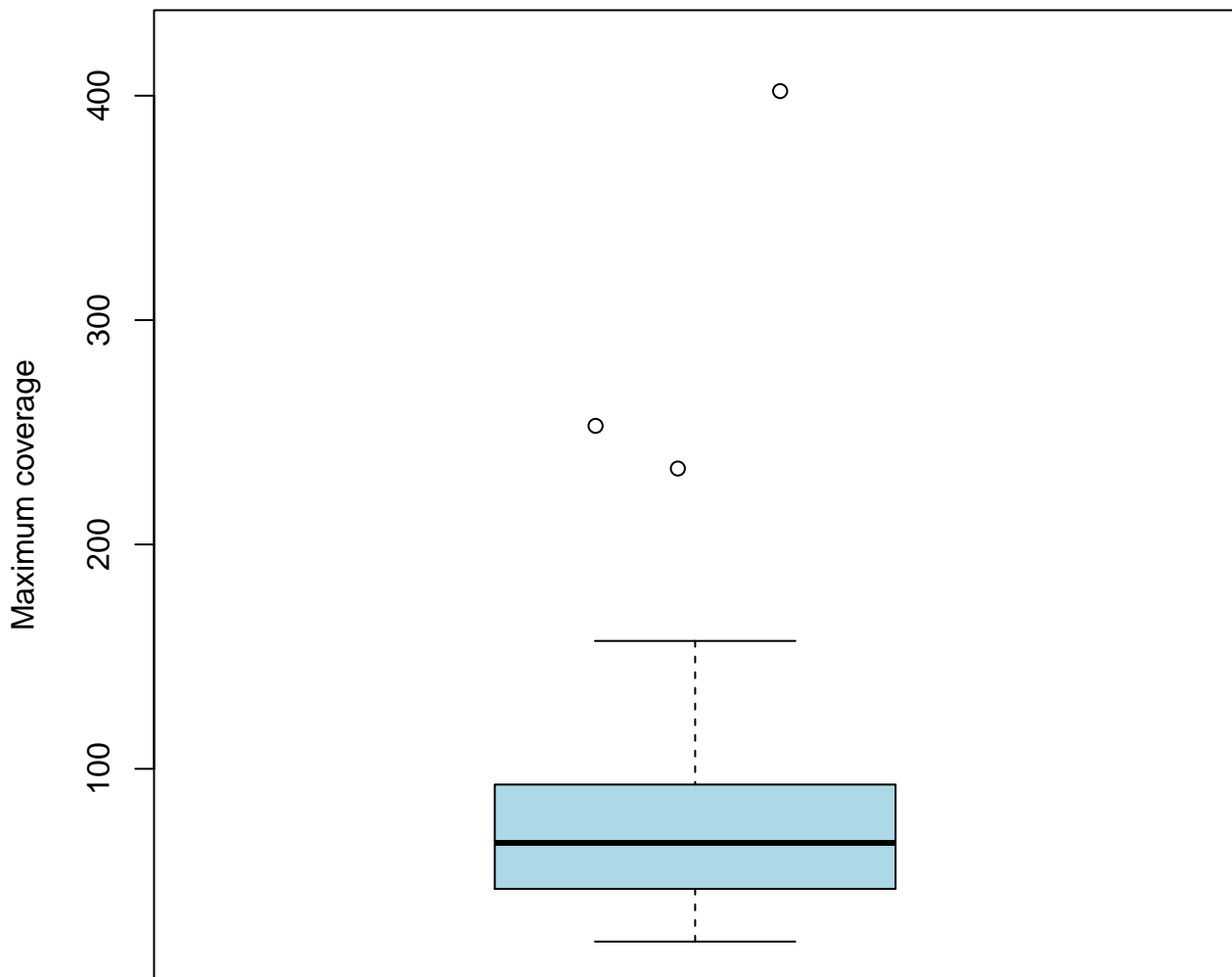
**jx in Intropolis but not "annotated" by knownGene:
235 samples; n jx: 34**



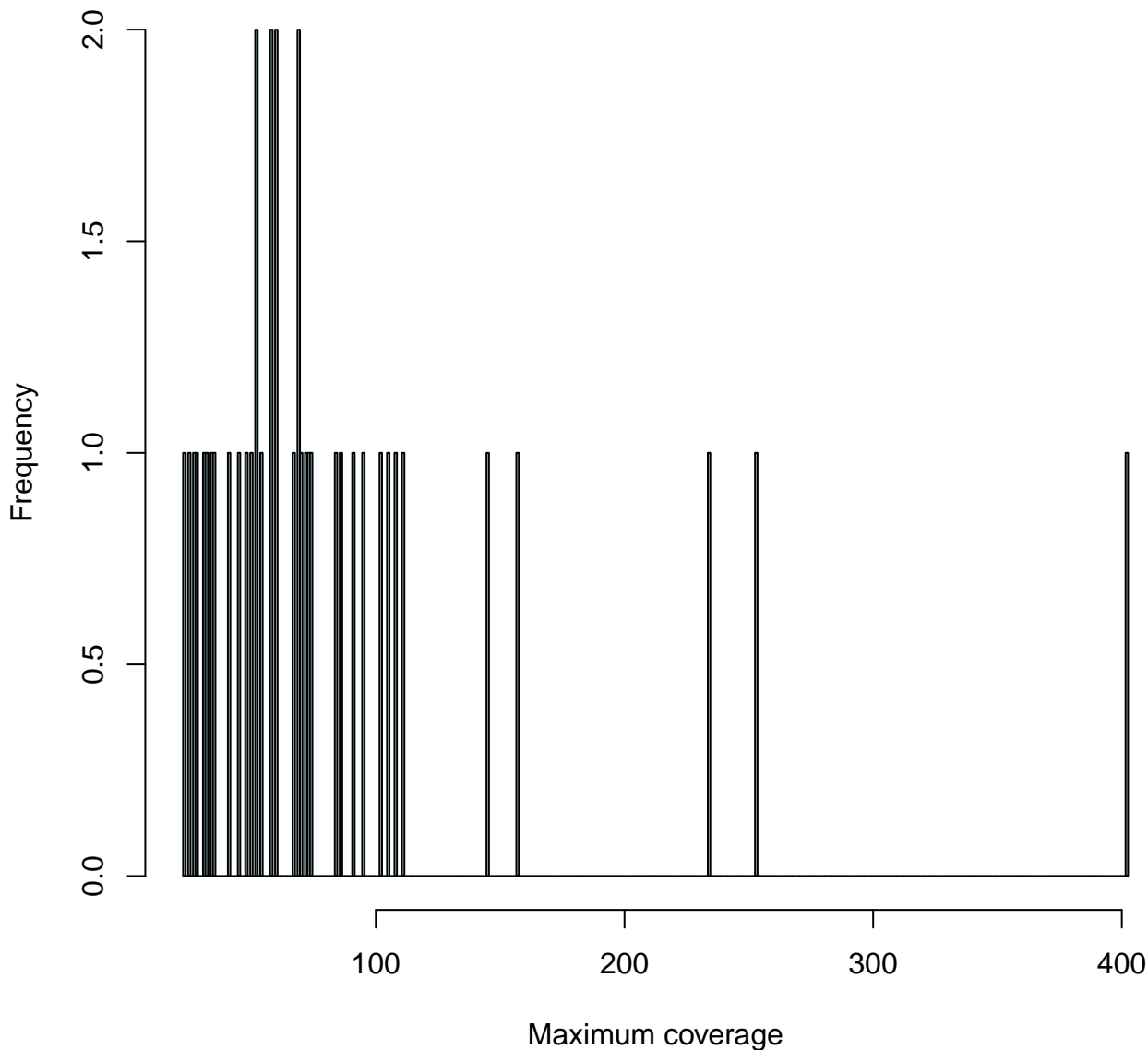
**jx in Intropolis but not "annotated" by knownGene:
235 samples; n jx: 34**



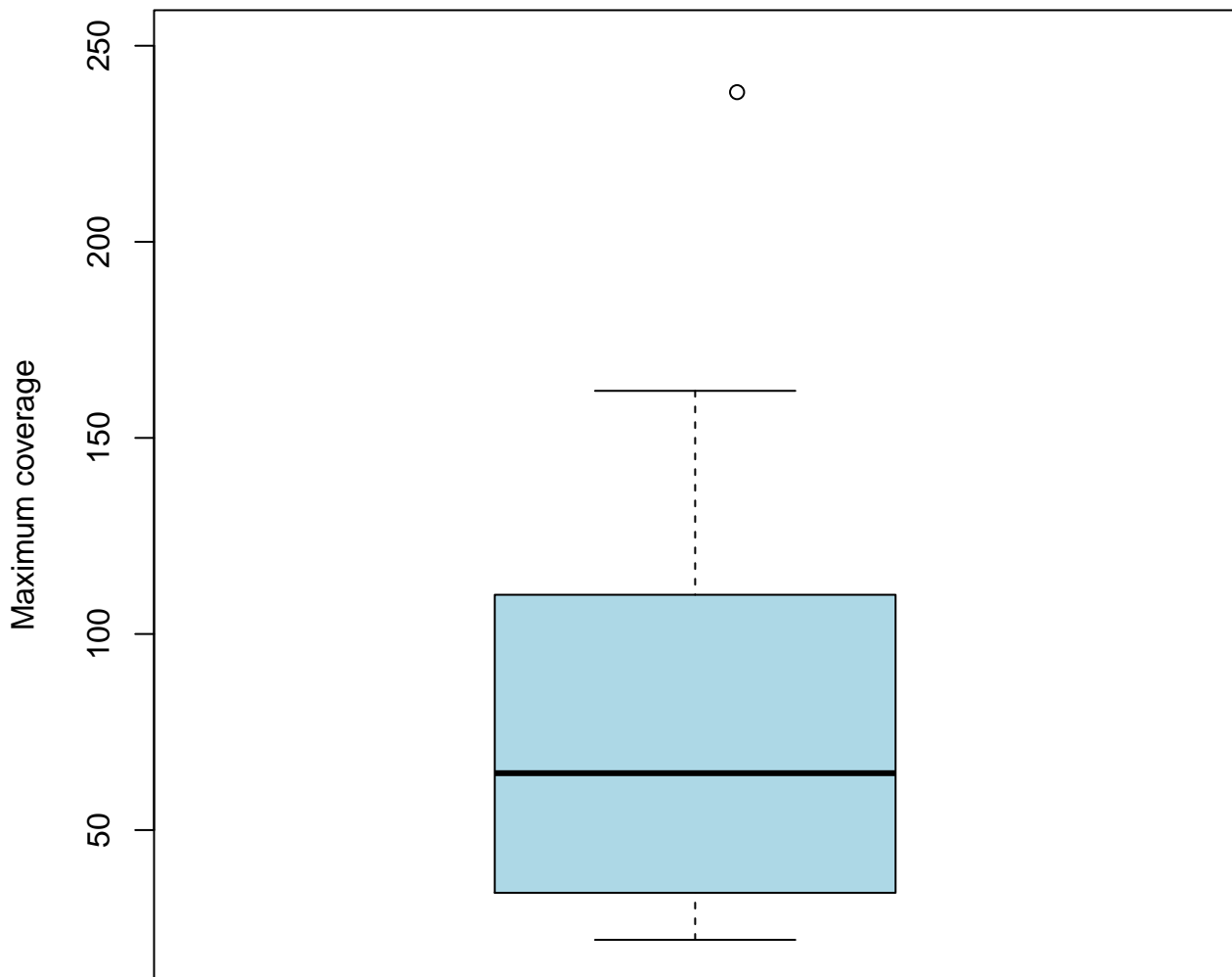
**jx in Intropolis but not "annotated" by knownGene:
236 samples; n jx: 39**



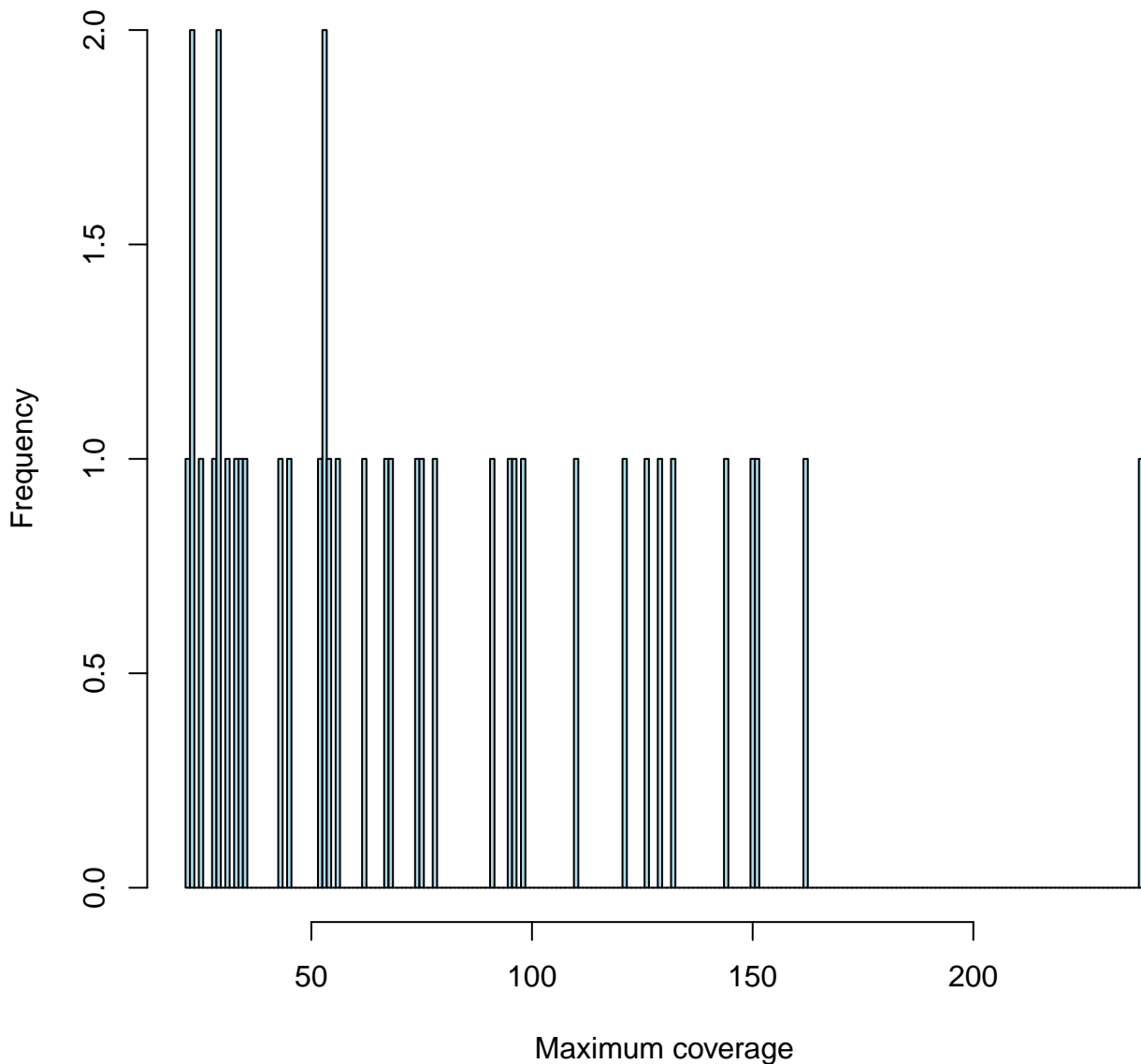
**jx in Intropolis but not "annotated" by knownGene:
236 samples; n jx: 39**



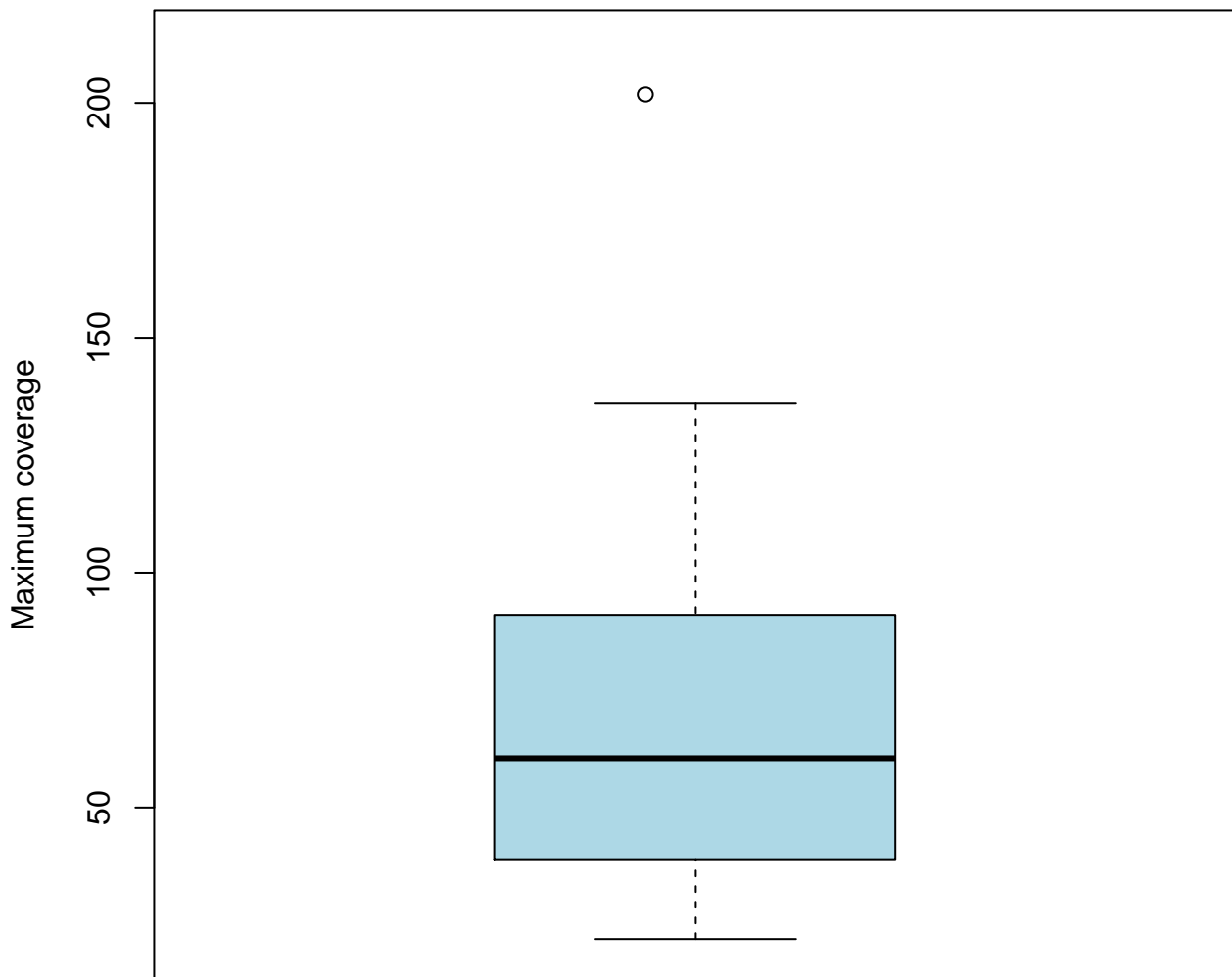
**jx in Intropolis but not "annotated" by knownGene:
237 samples; n jx: 38**



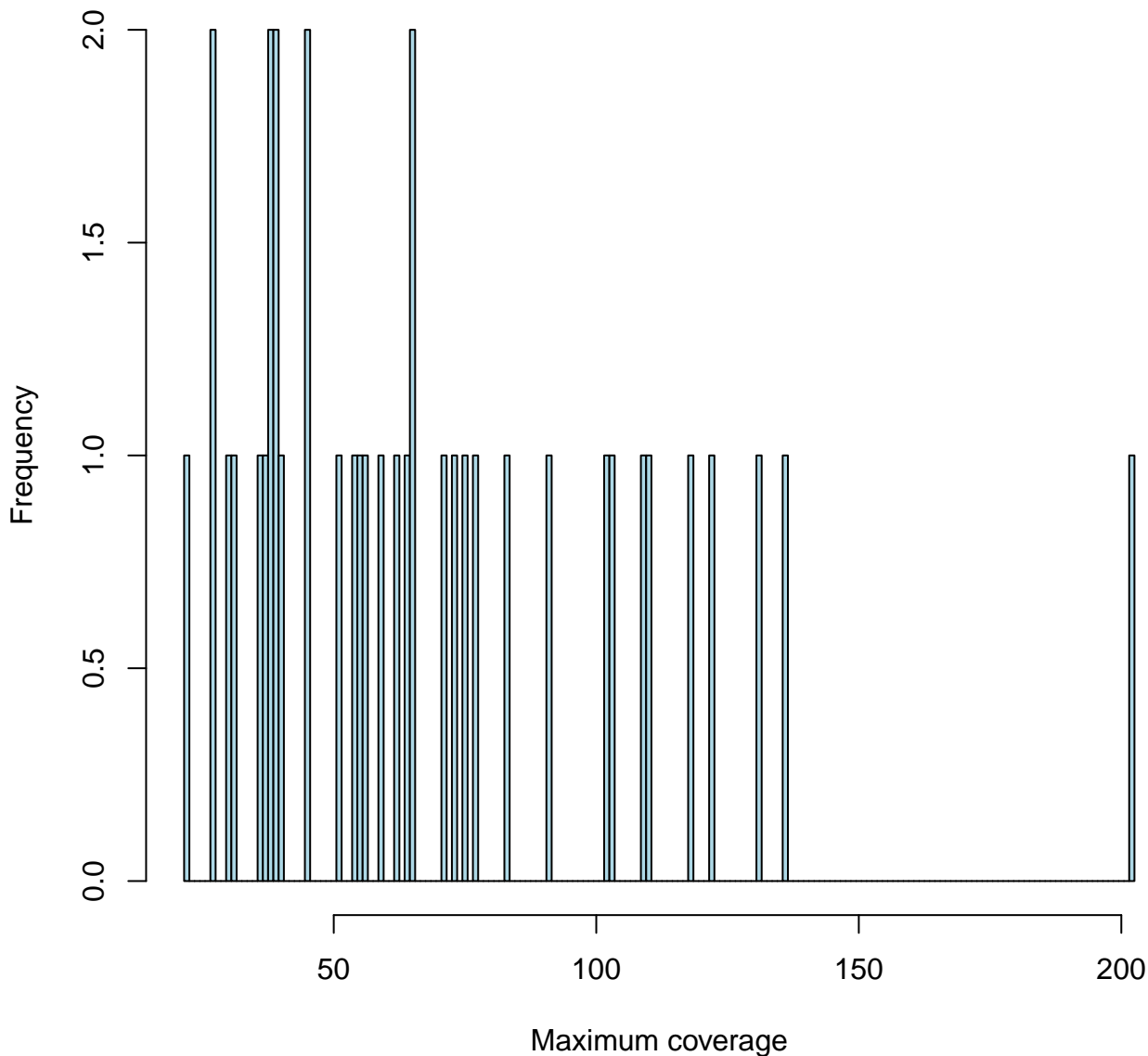
**jx in Intropolis but not "annotated" by knownGene:
237 samples; n jx: 38**



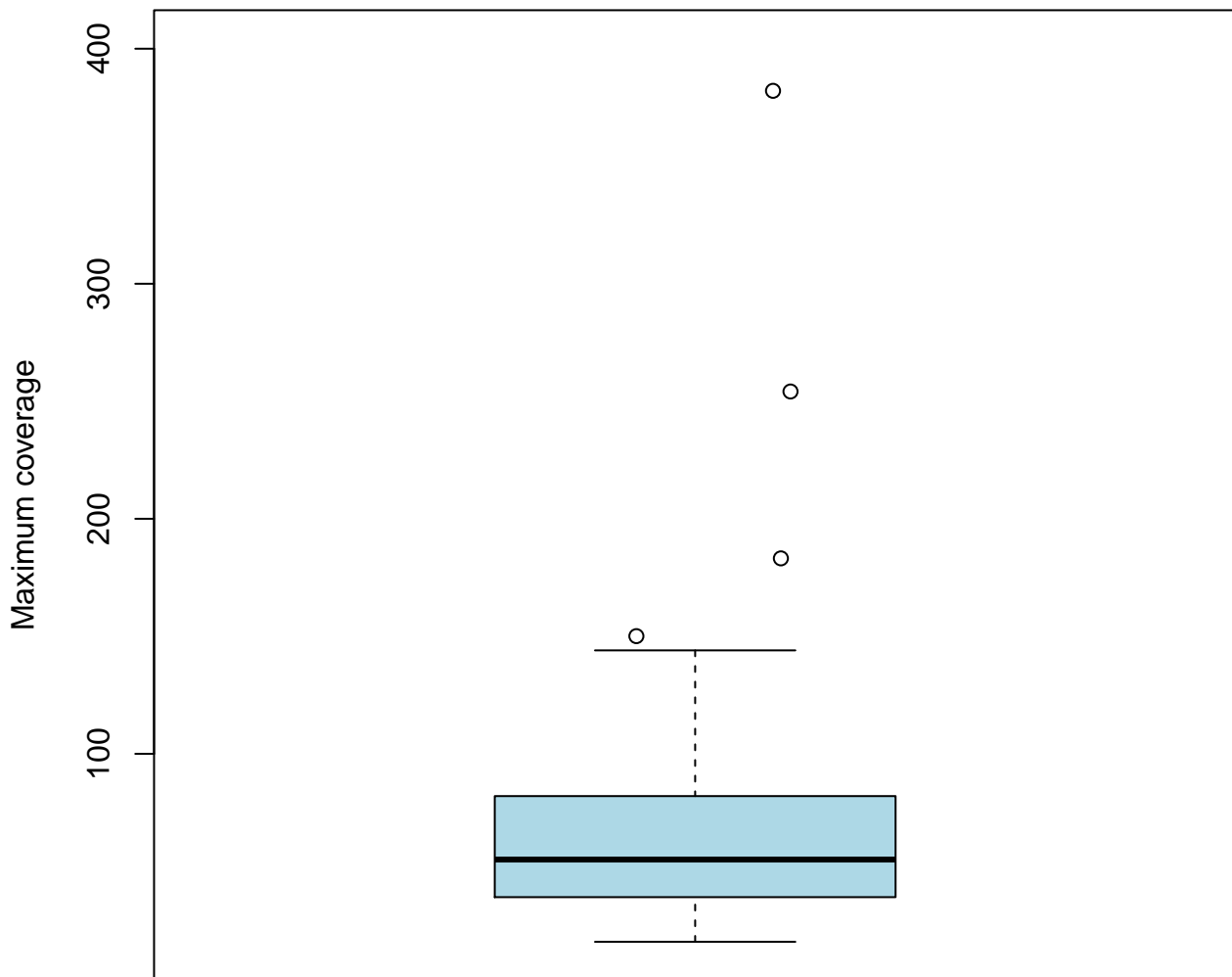
**jx in Intropolis but not "annotated" by knownGene:
238 samples; n jx: 38**



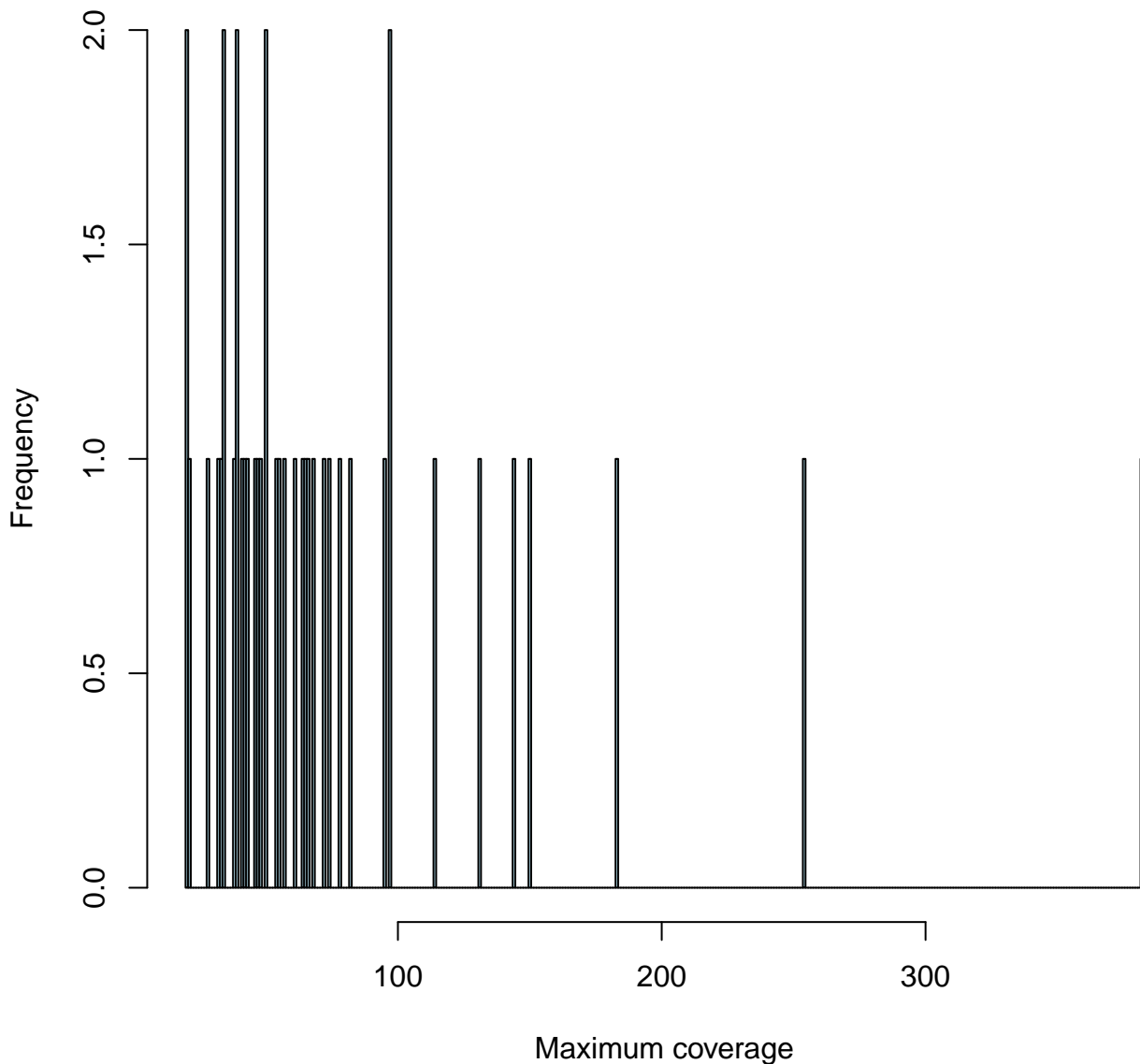
**jx in Intropolis but not "annotated" by knownGene:
238 samples; n jx: 38**



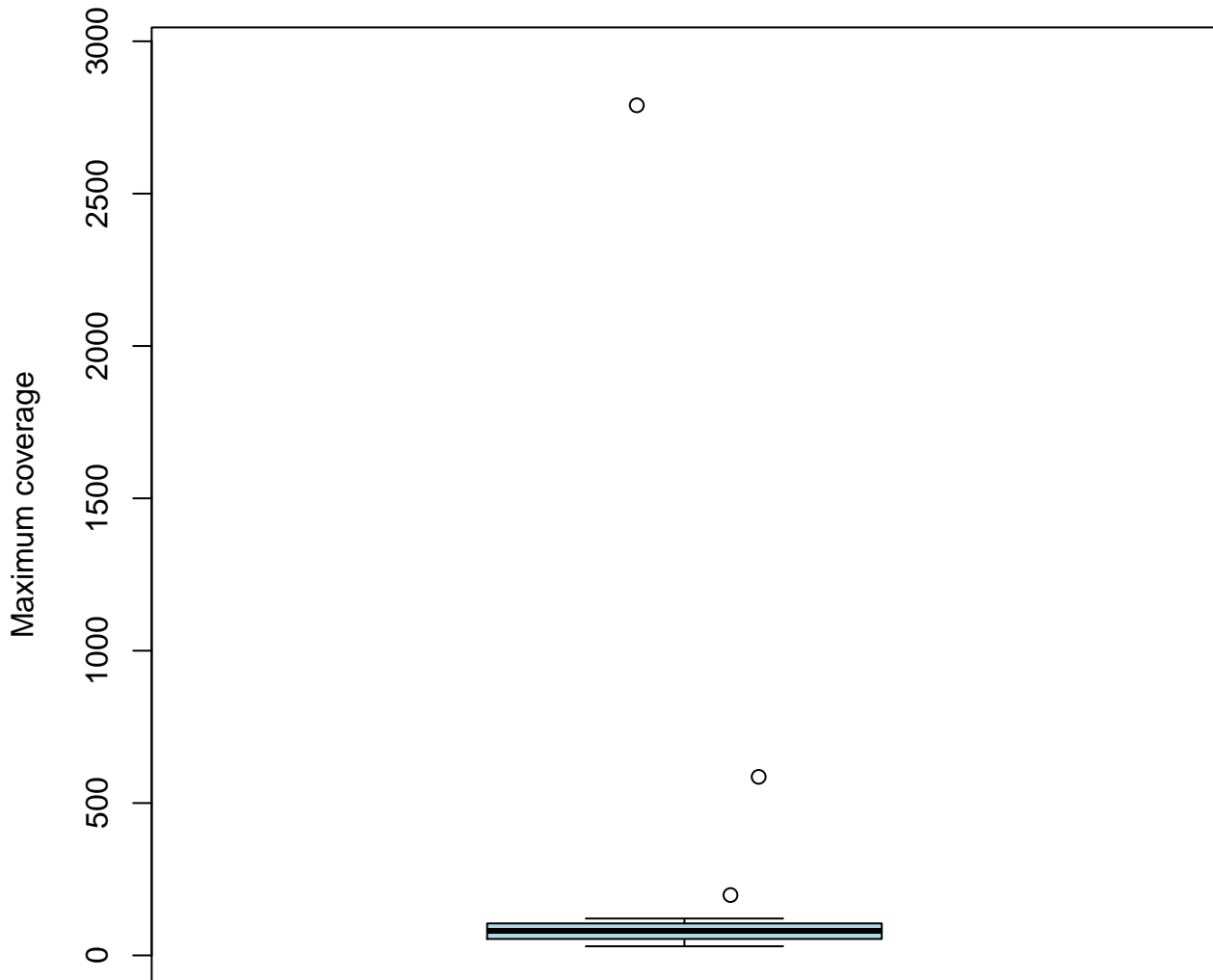
**jx in Intropolis but not "annotated" by knownGene:
239 samples; n jx: 41**



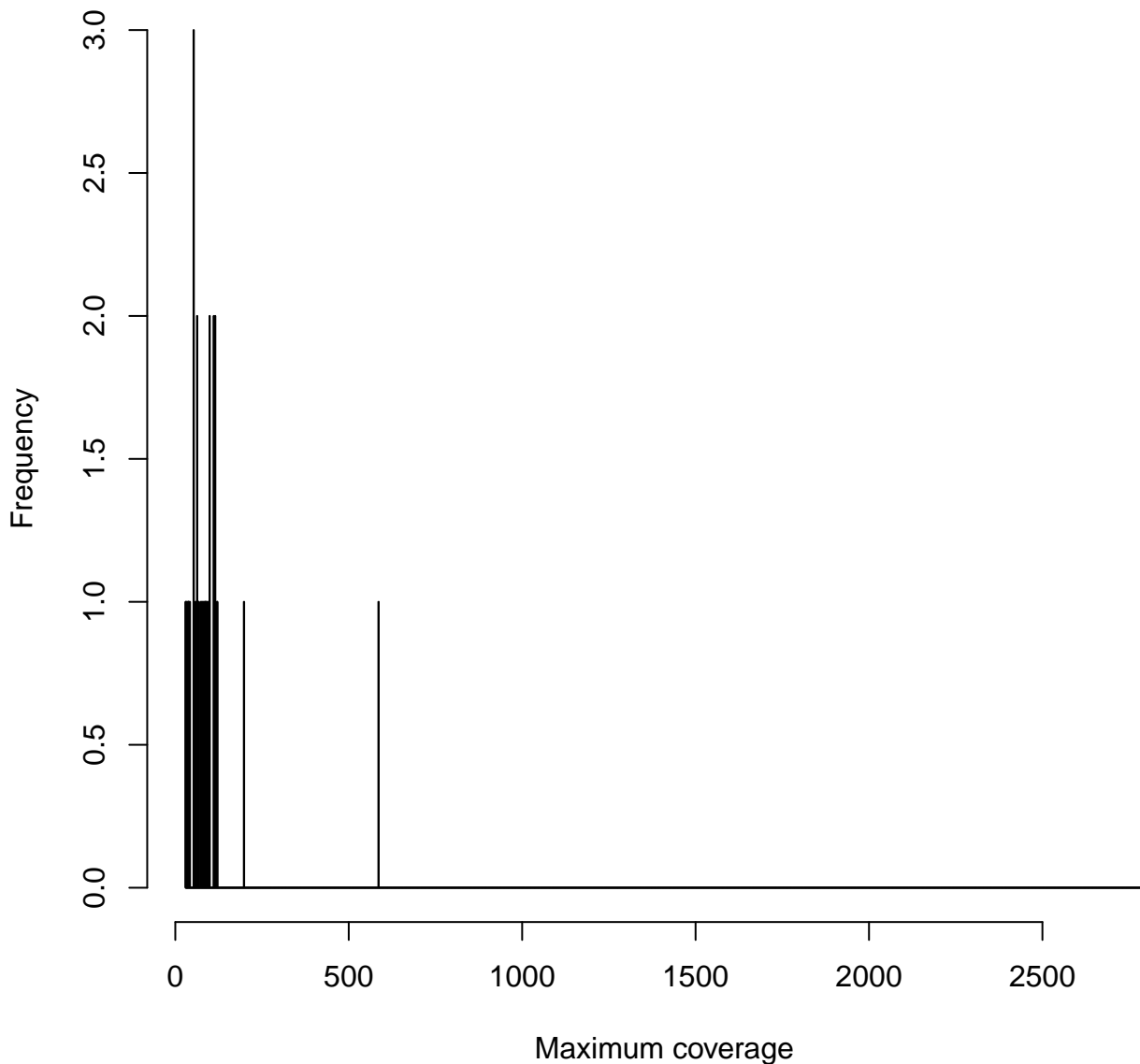
**jx in Intropolis but not "annotated" by knownGene:
239 samples; n jx: 41**



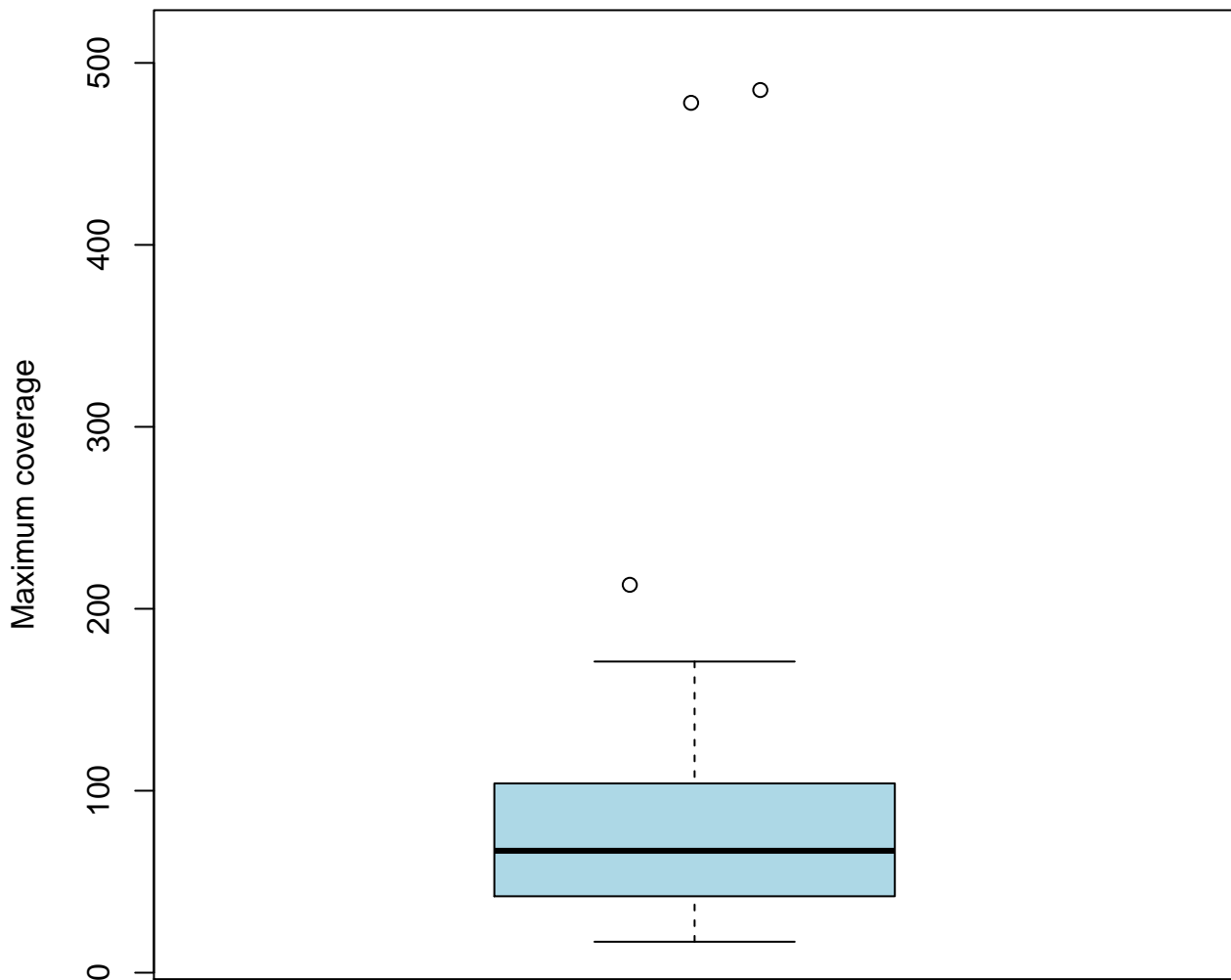
**jx in Intropolis but not "annotated" by knownGene:
240 samples; n jx: 40**



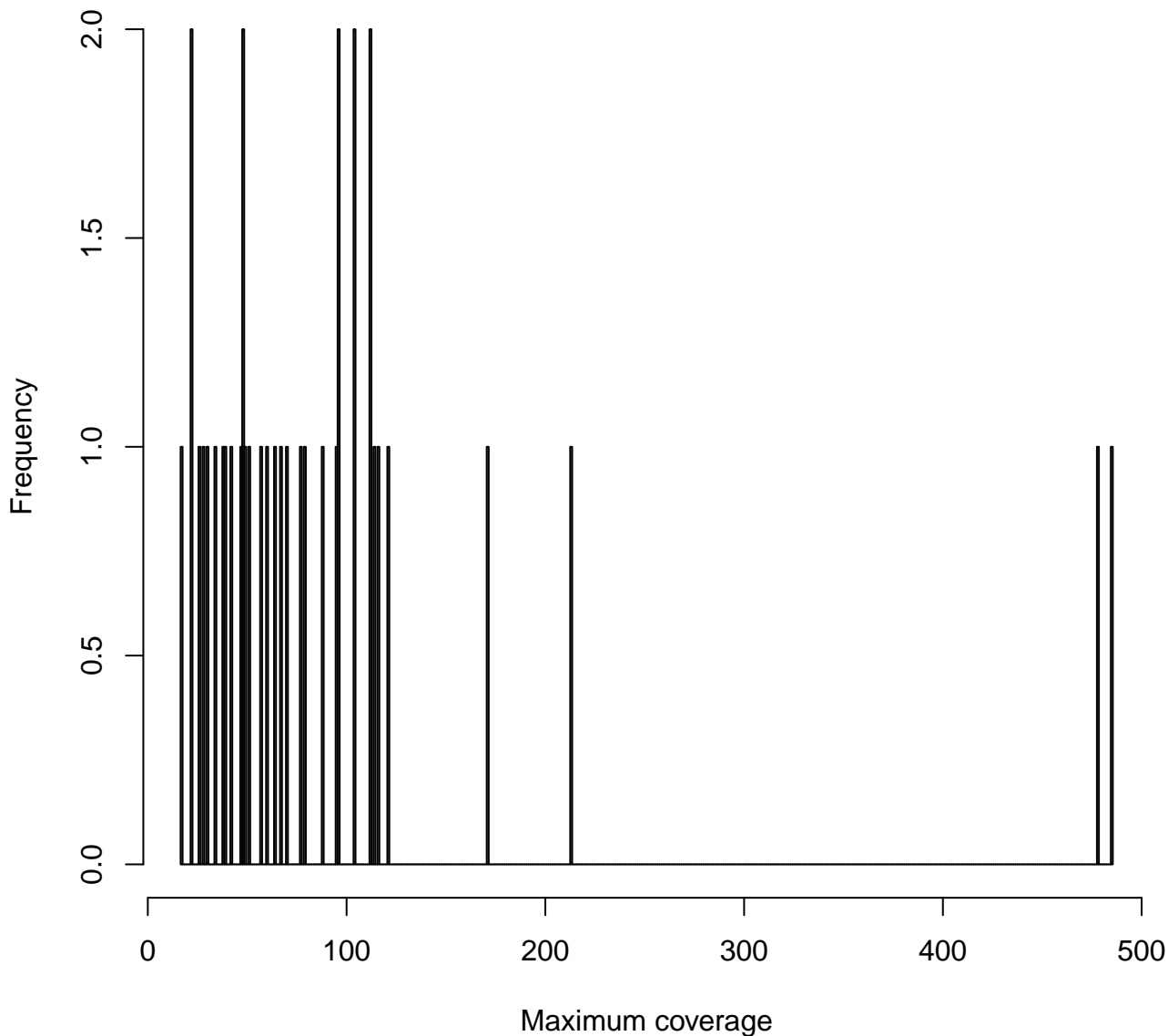
**jx in Intropolis but not "annotated" by knownGene:
240 samples; n jx: 40**



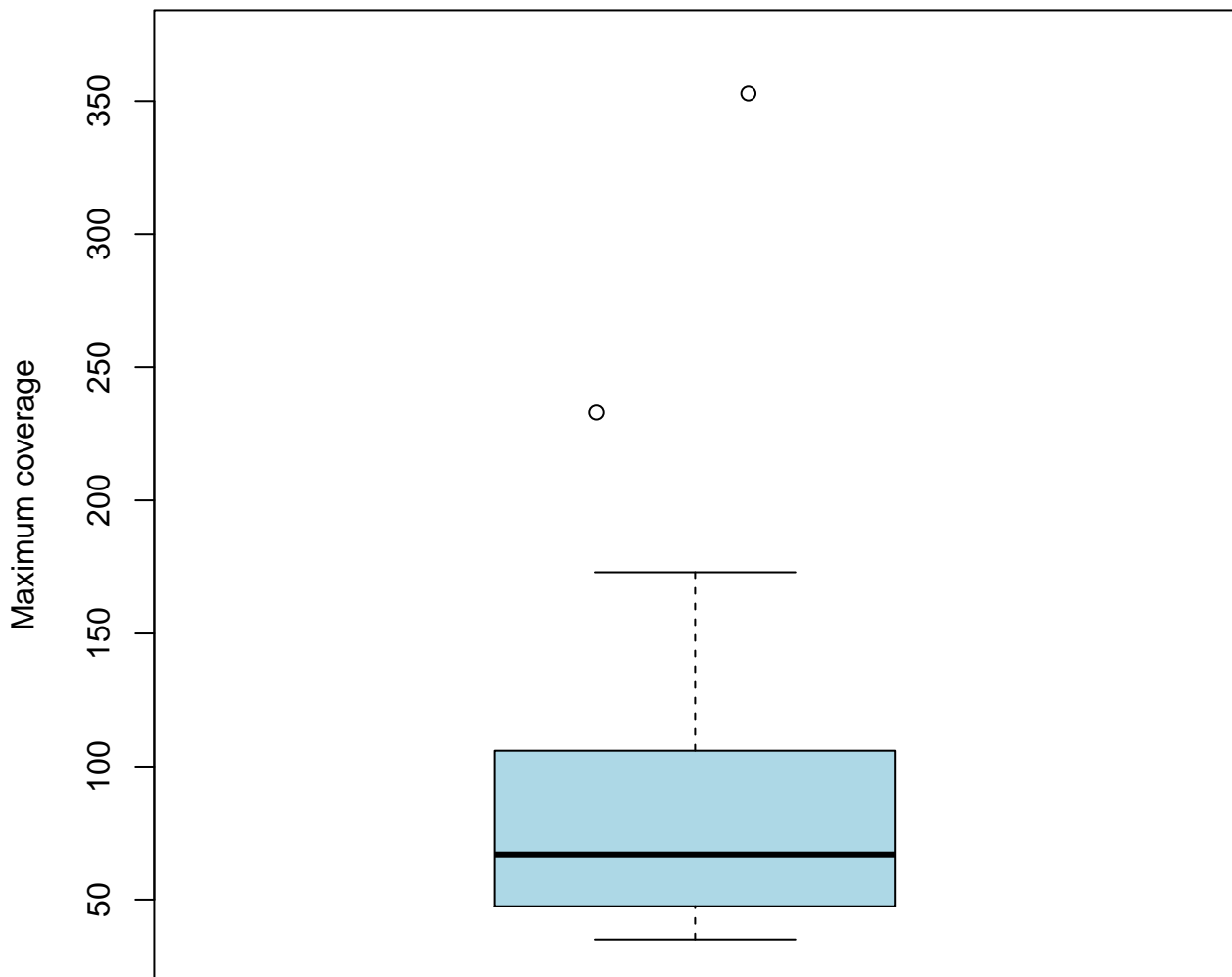
**jx in Intropolis but not "annotated" by knownGene:
241 samples; n jx: 37**



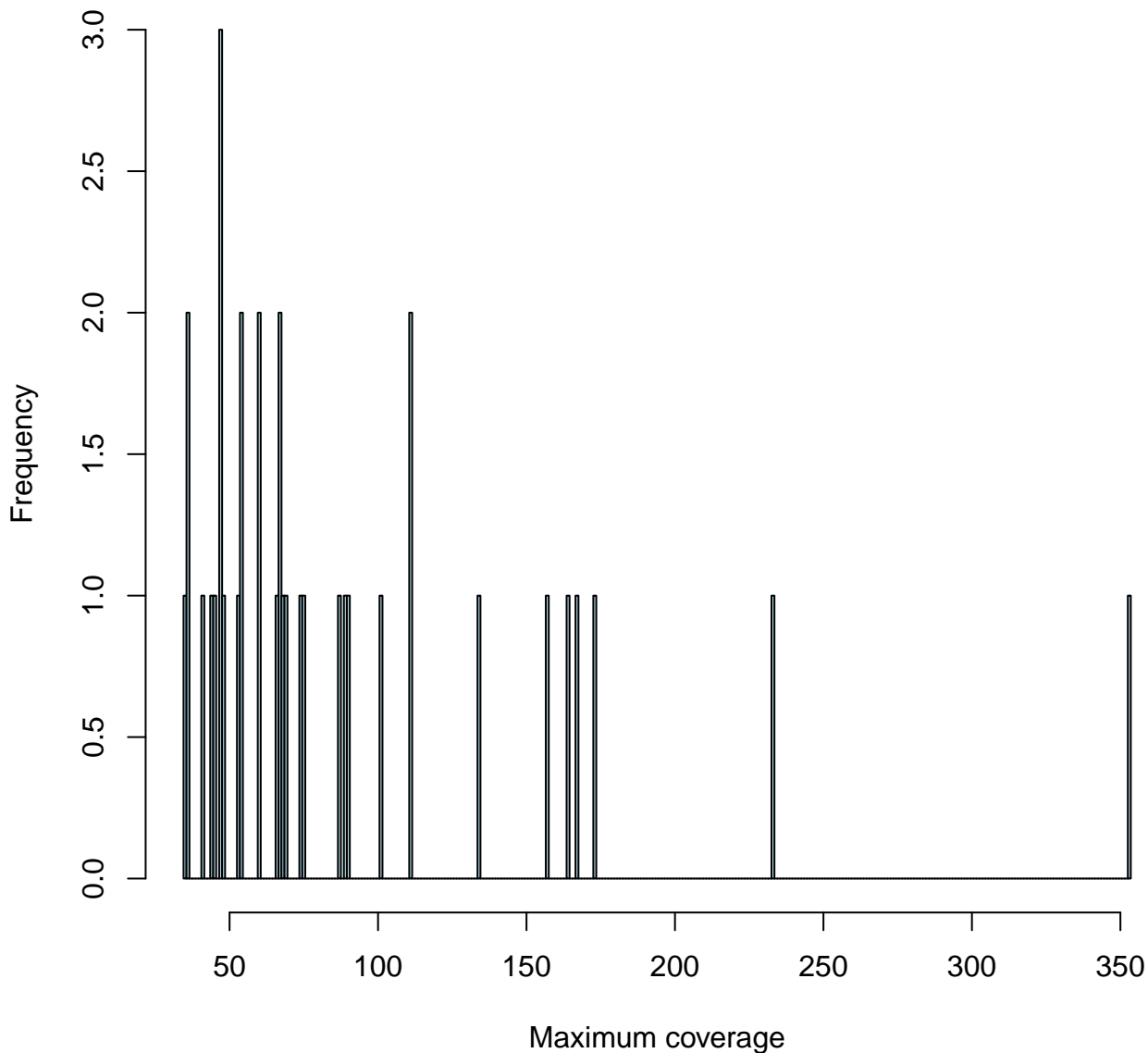
**jx in Intropolis but not "annotated" by knownGene:
241 samples; n jx: 37**



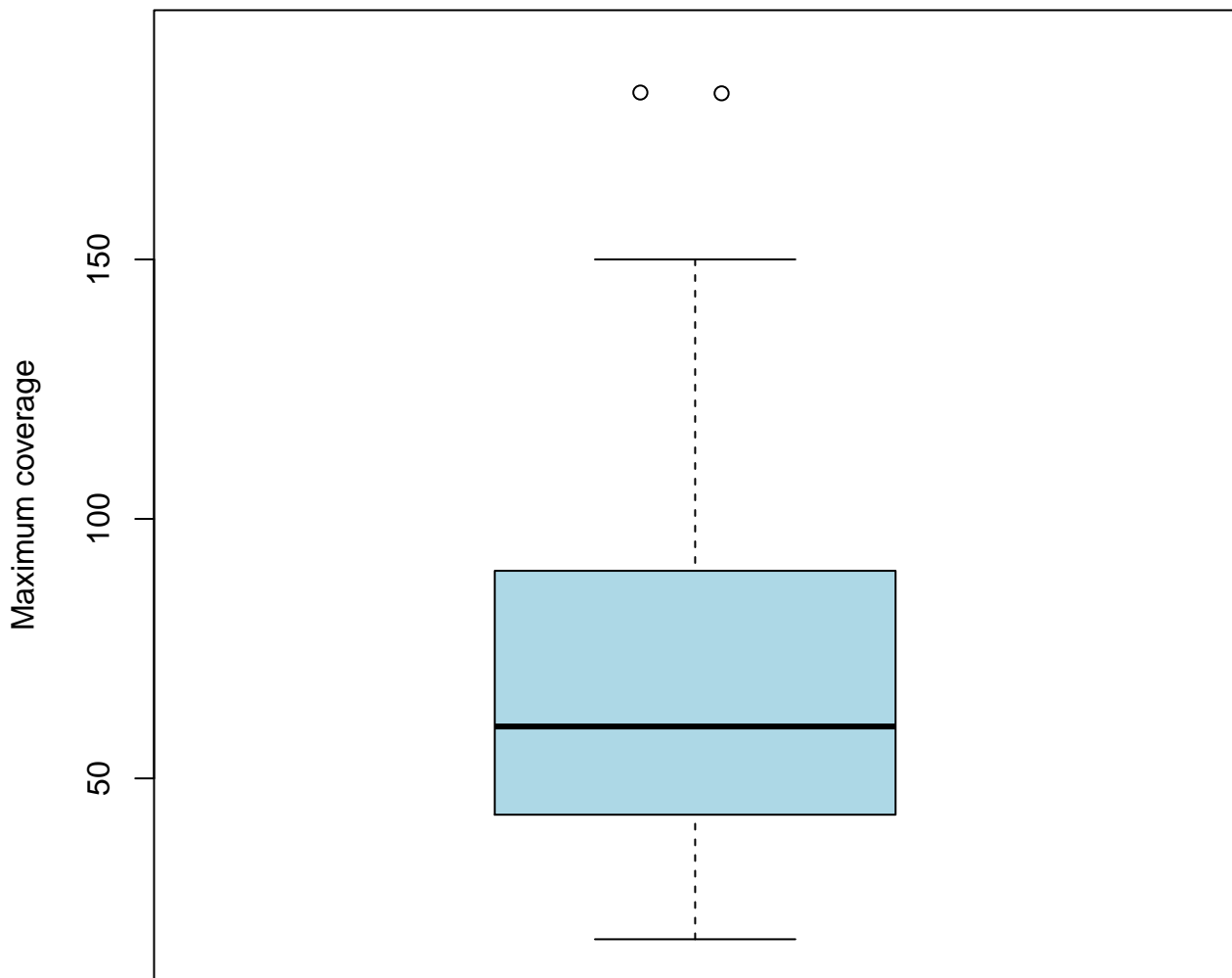
**jx in Intropolis but not "annotated" by knownGene:
242 samples; n jx: 35**



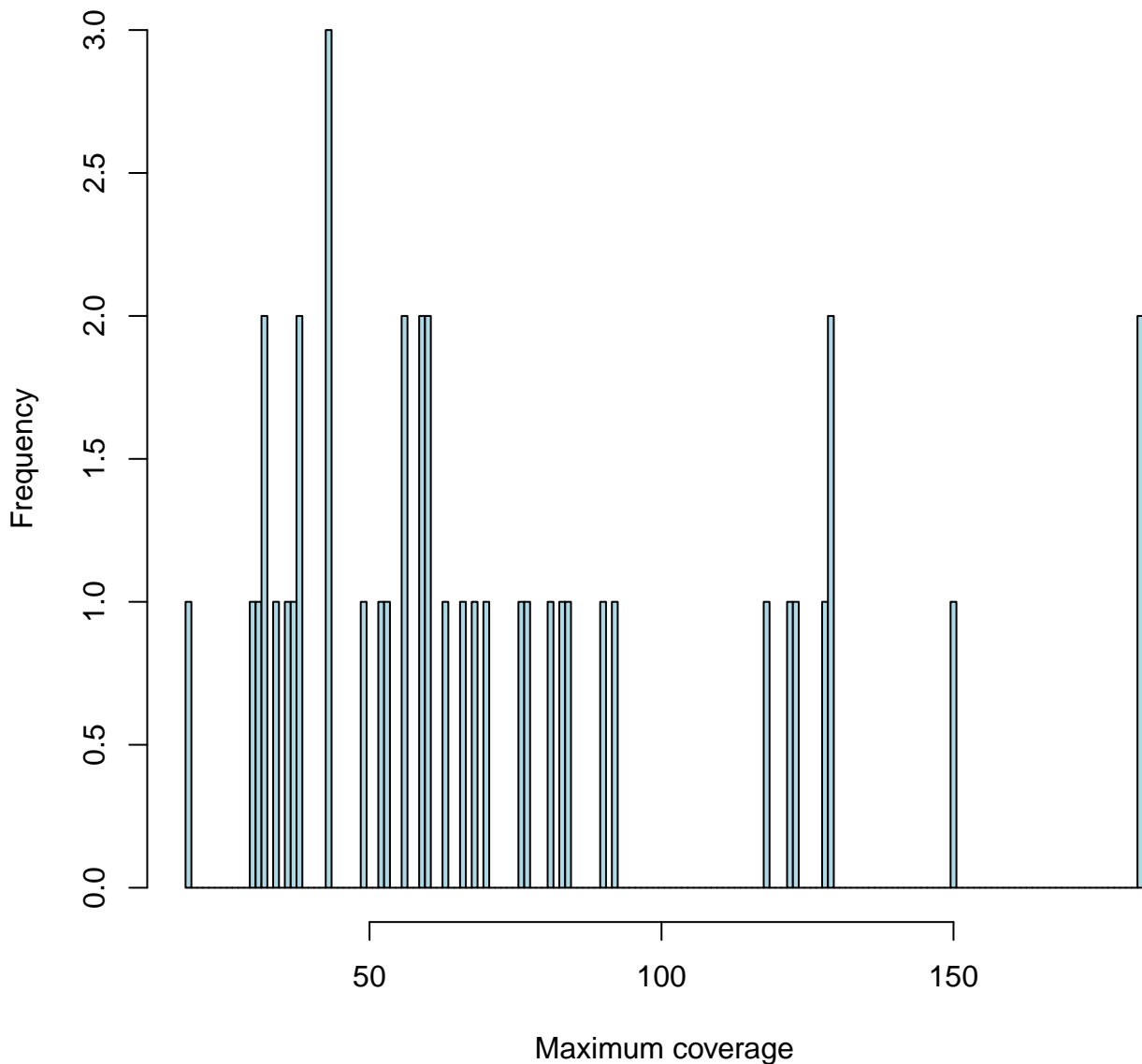
**jx in Intropolis but not "annotated" by knownGene:
242 samples; n jx: 35**



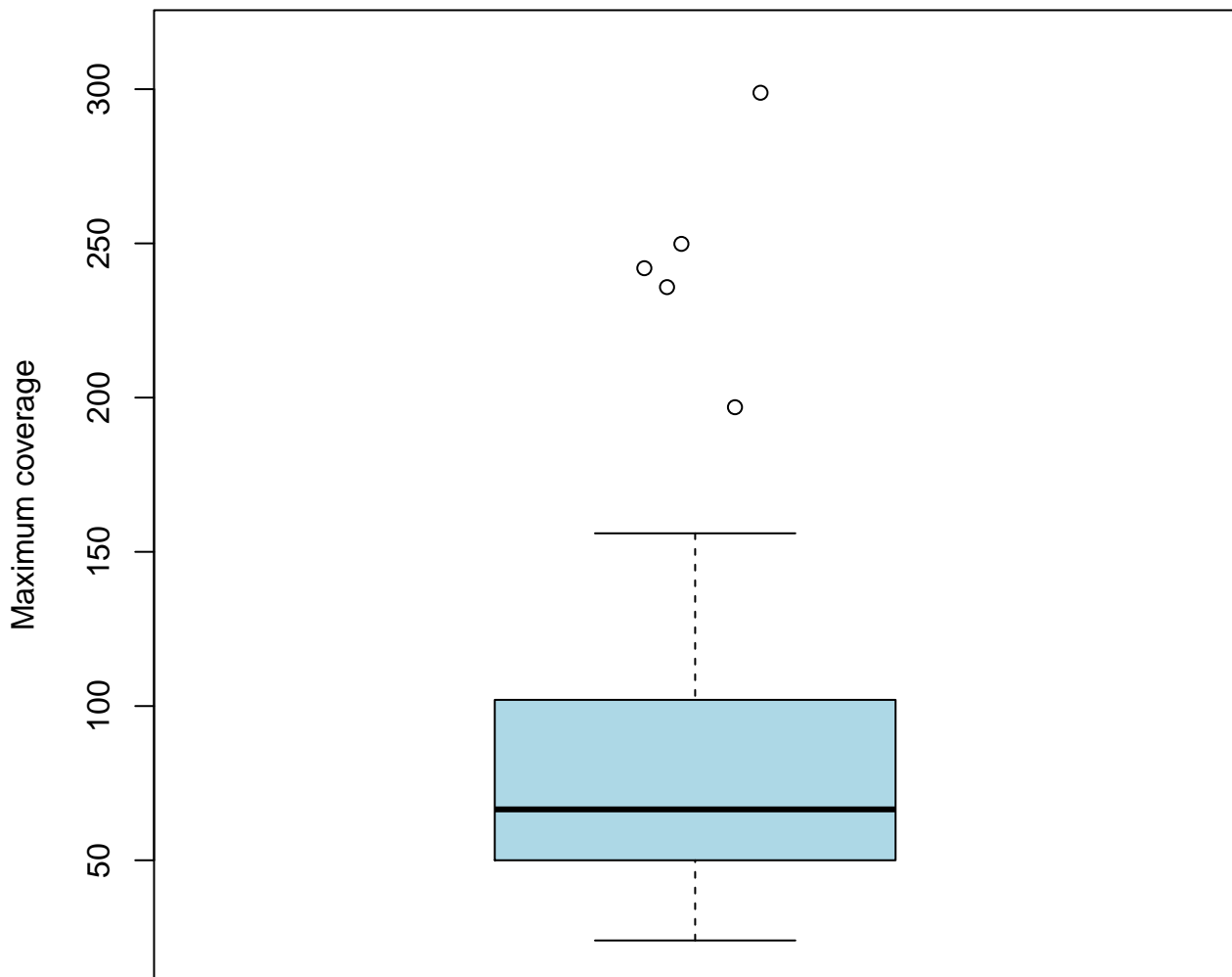
**jx in Intropolis but not "annotated" by knownGene:
243 samples; n jx: 42**



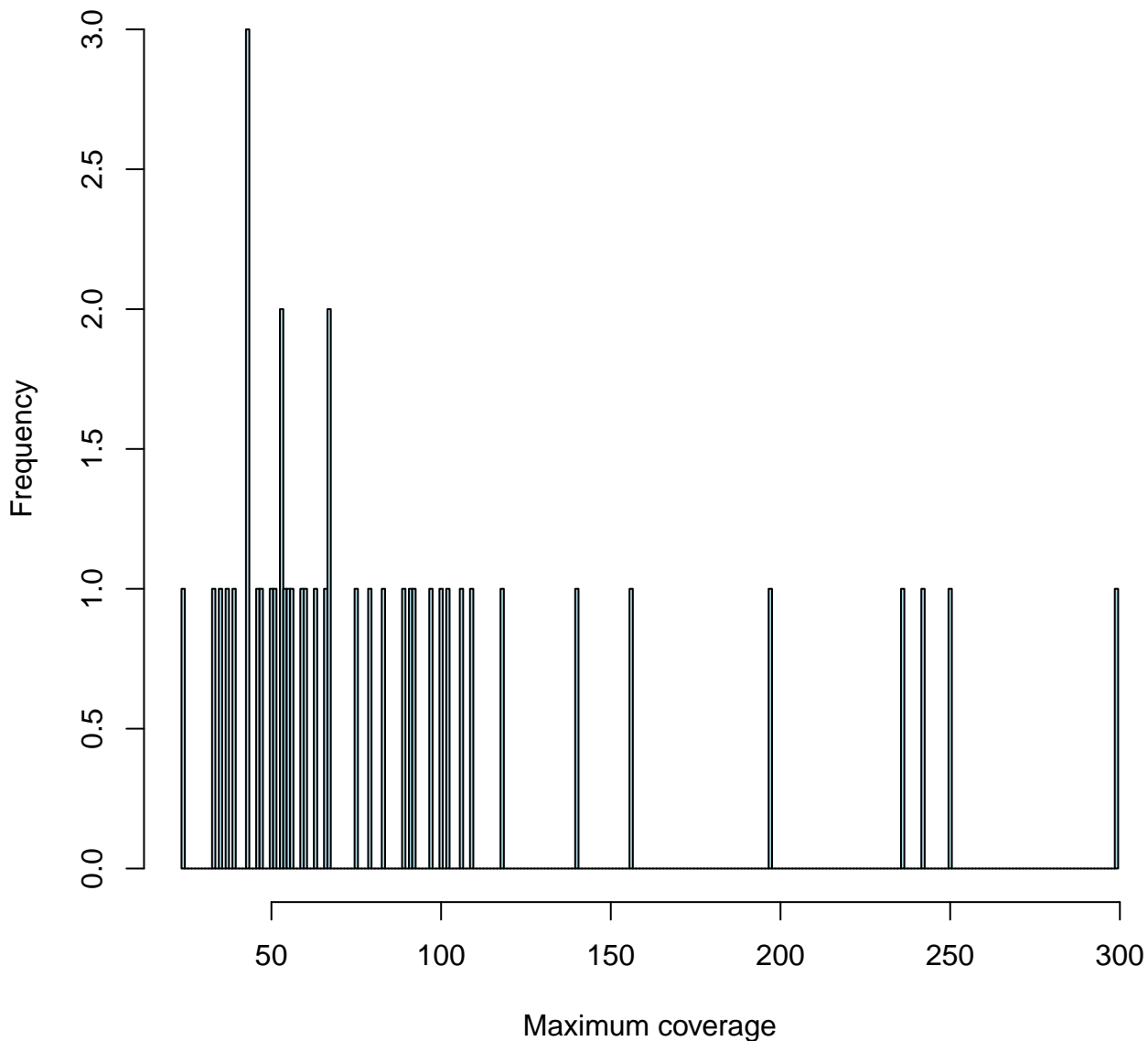
**jx in Intropolis but not "annotated" by knownGene:
243 samples; n jx: 42**



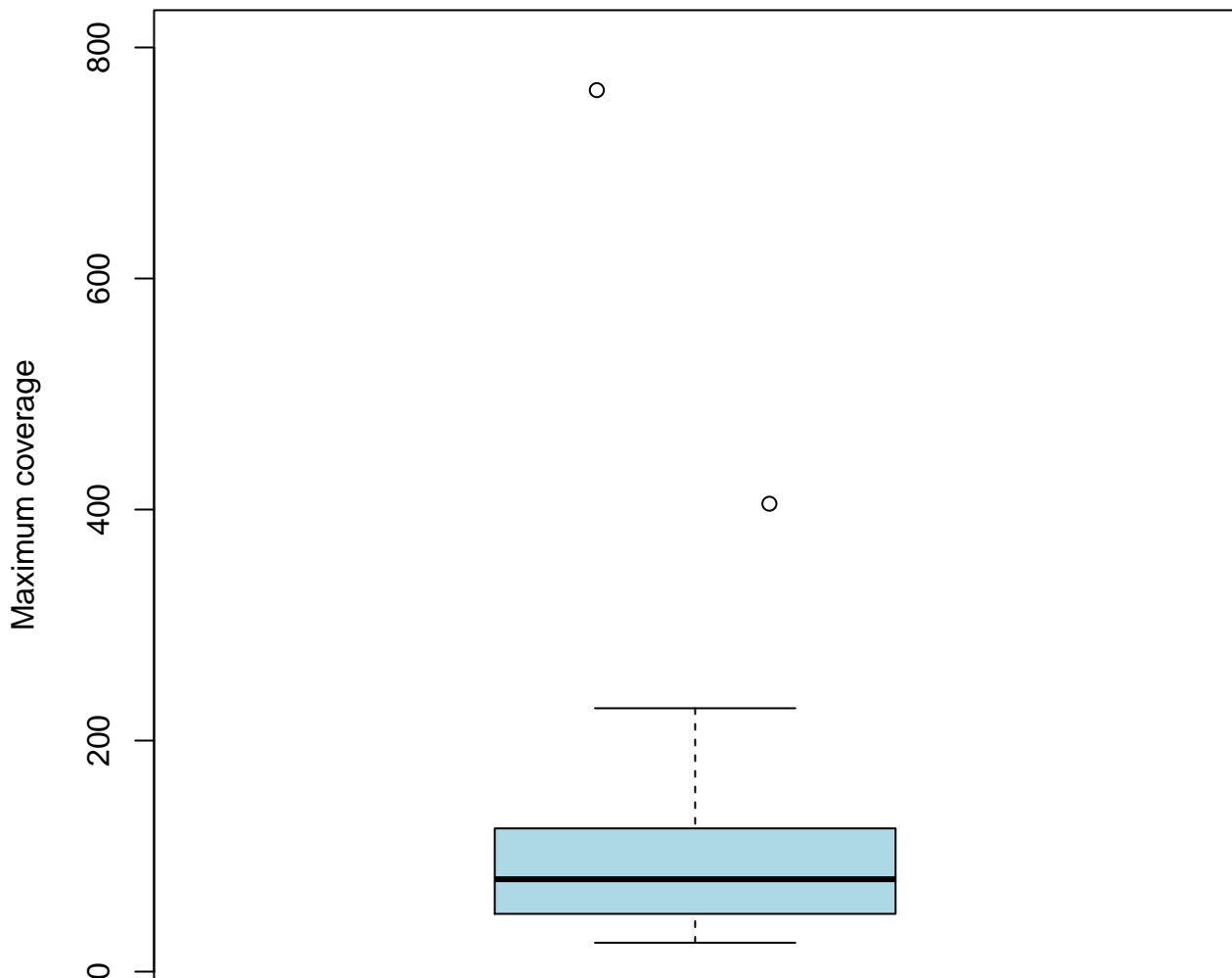
**jx in Intropolis but not "annotated" by knownGene:
244 samples; n jx: 42**



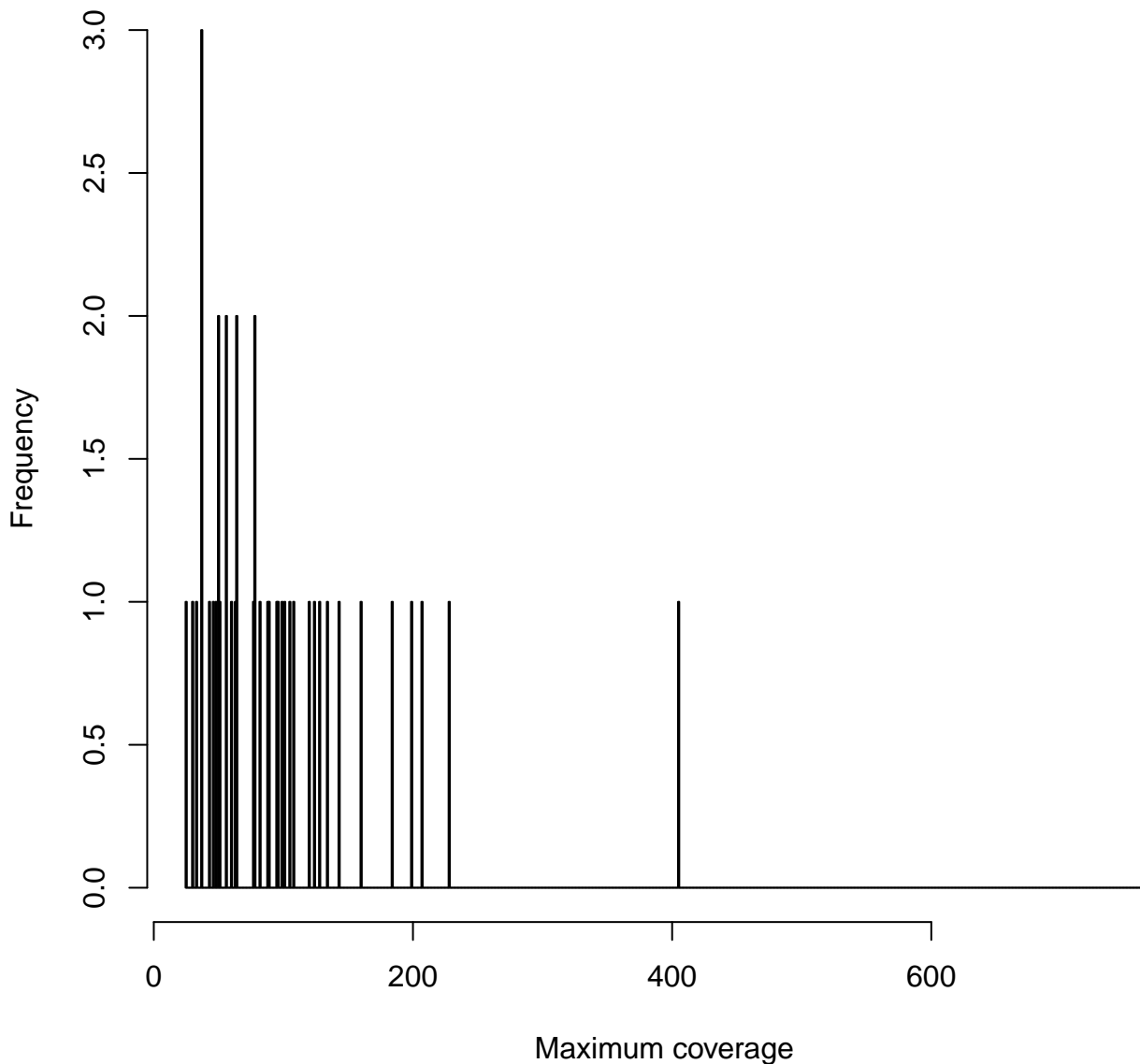
**jx in Intropolis but not "annotated" by knownGene:
244 samples; n jx: 42**



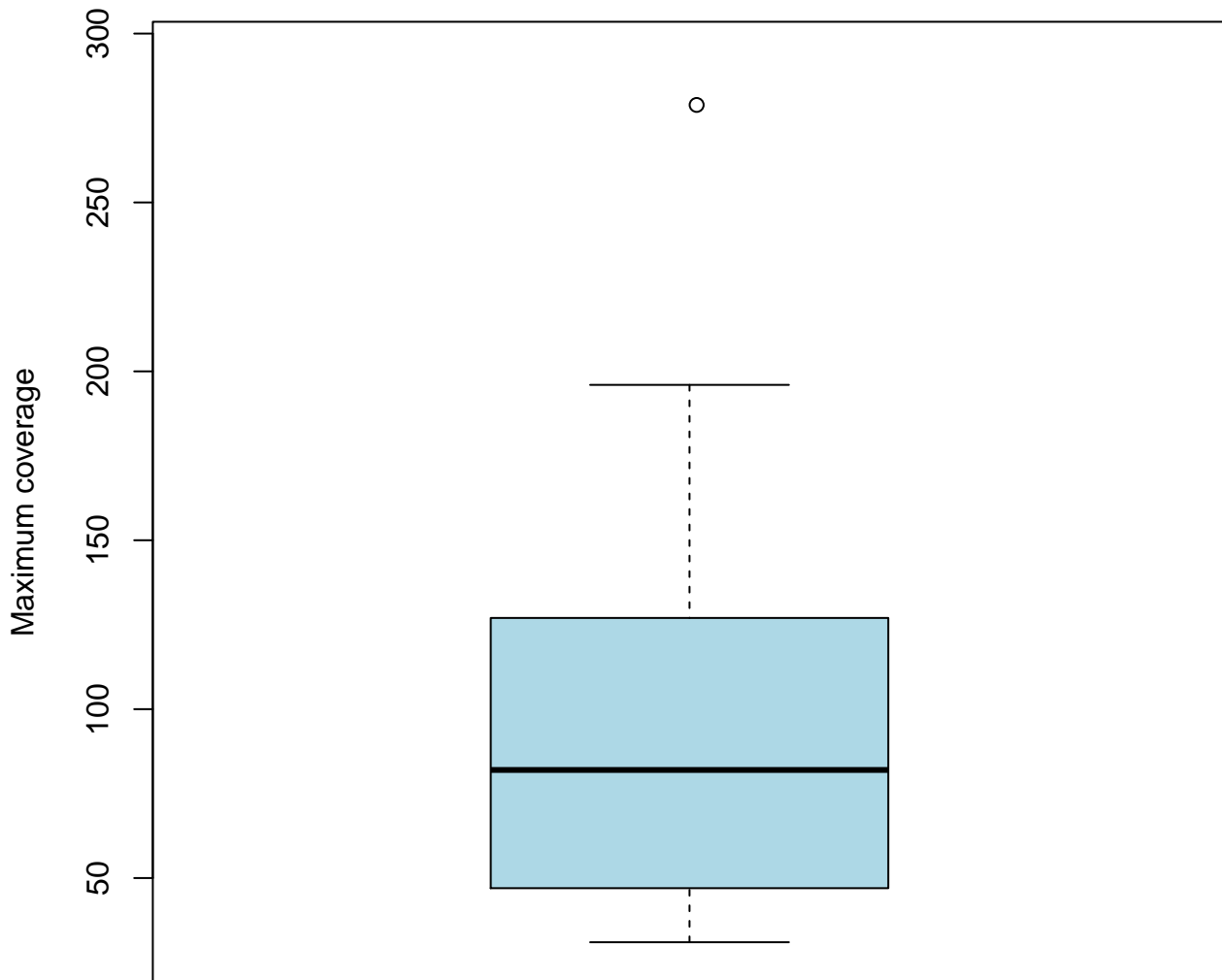
**jx in Intropolis but not "annotated" by knownGene:
245 samples; n jx: 42**



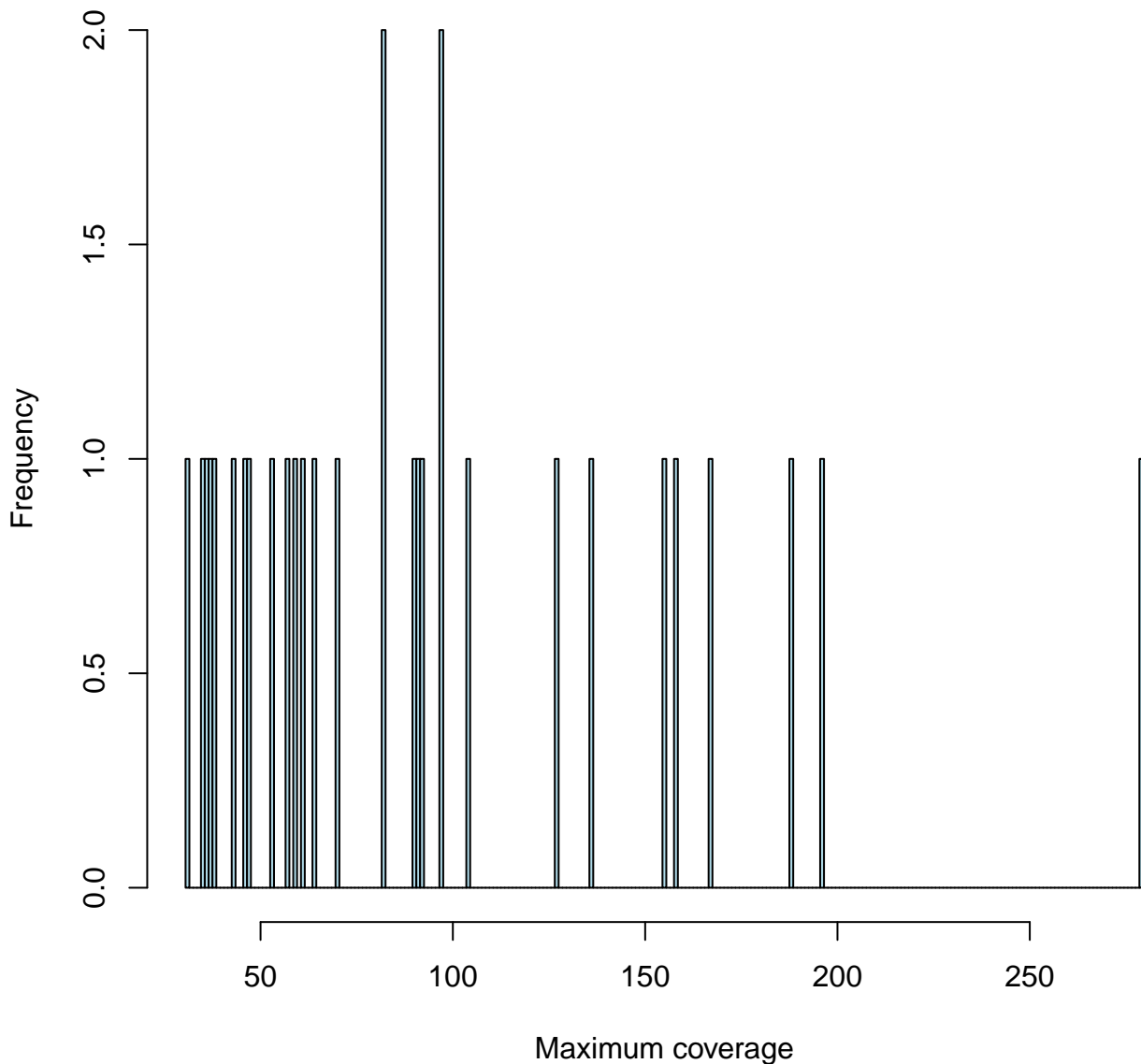
**jx in Intropolis but not "annotated" by knownGene:
245 samples; n jx: 42**



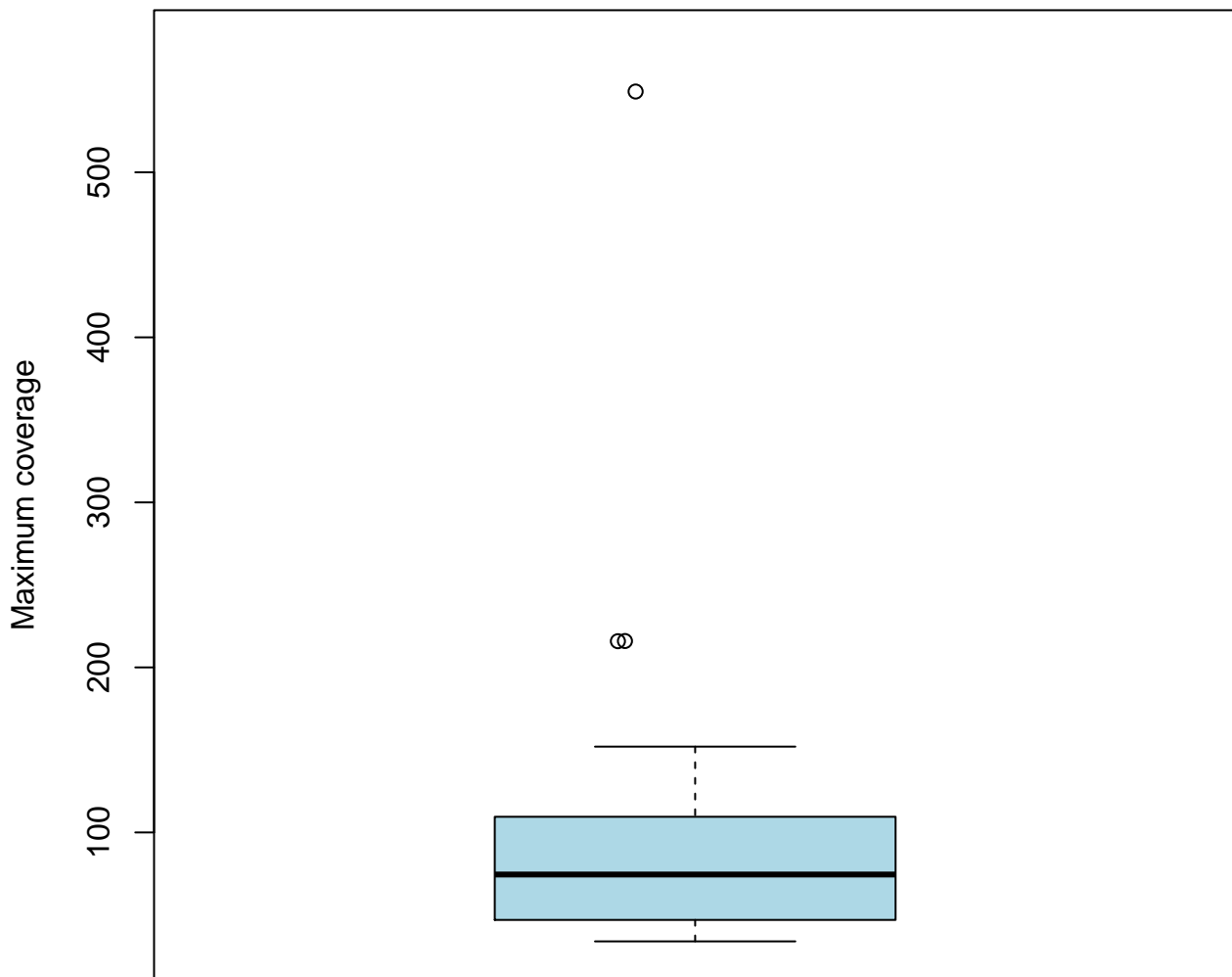
**jx in Intropolis but not "annotated" by knownGene:
246 samples; n jx: 30**



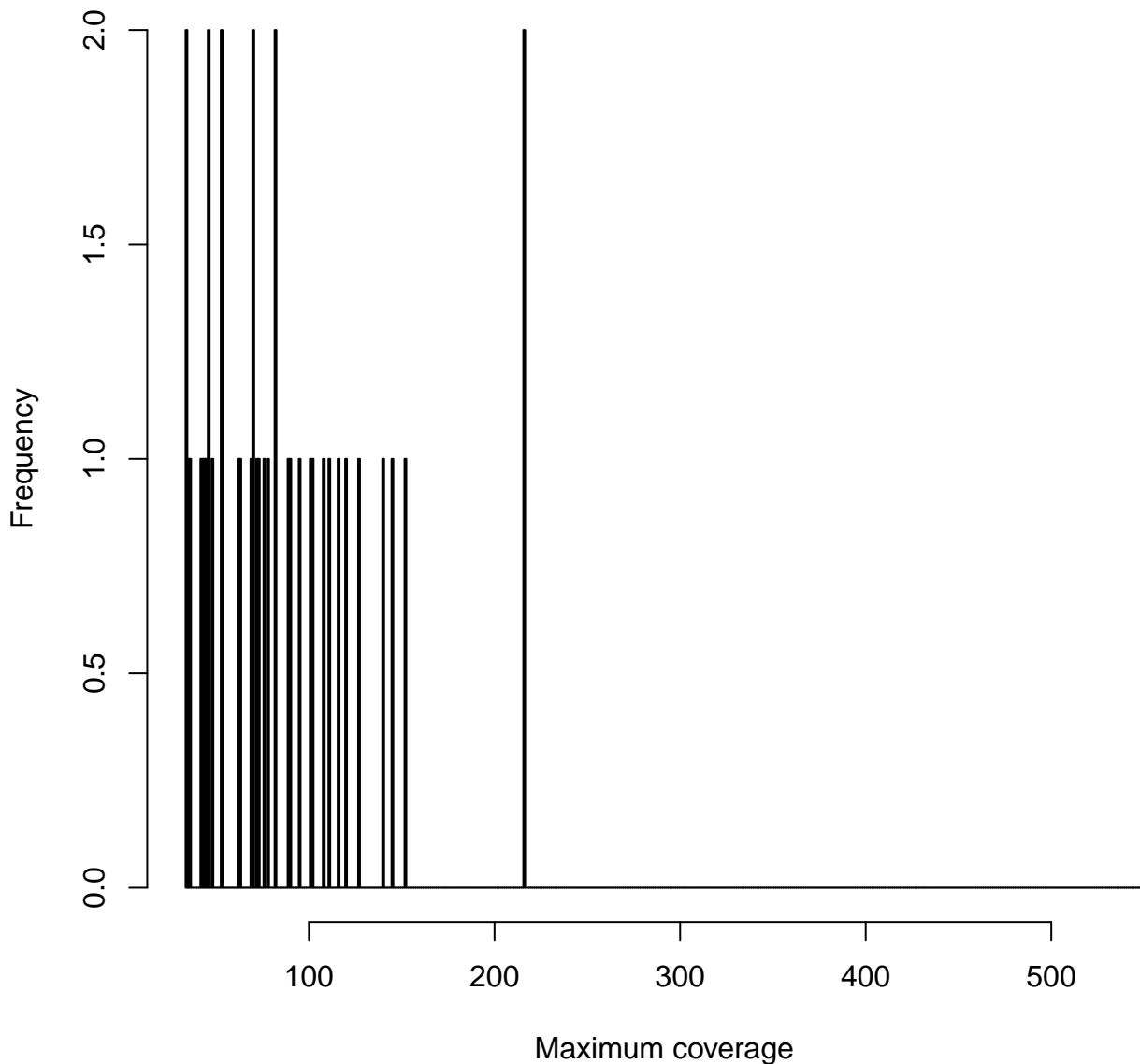
**jx in Intropolis but not "annotated" by knownGene:
246 samples; n jx: 30**



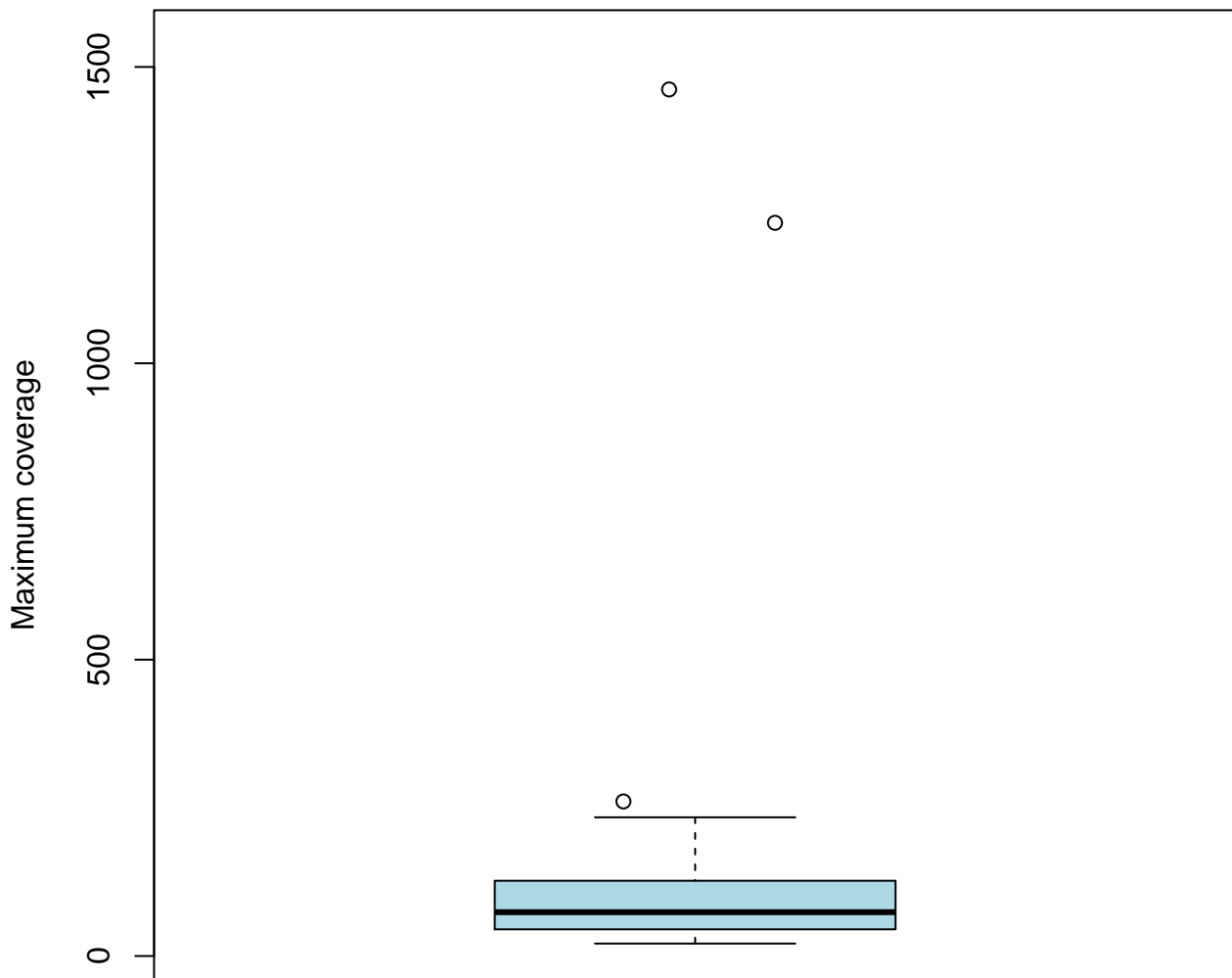
**jx in Intropolis but not "annotated" by knownGene:
247 samples; n jx: 40**



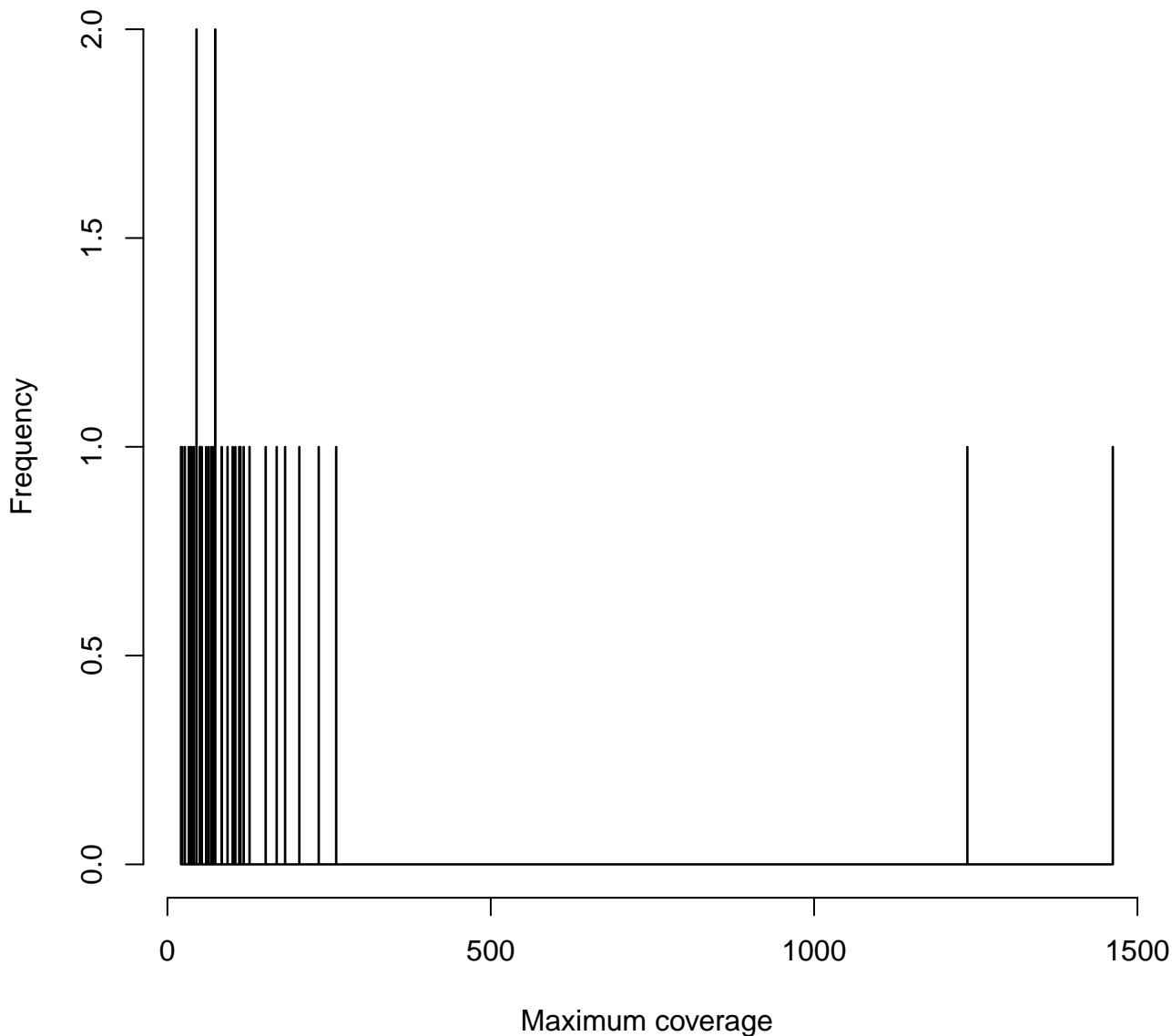
**jx in Intropolis but not "annotated" by knownGene:
247 samples; n jx: 40**



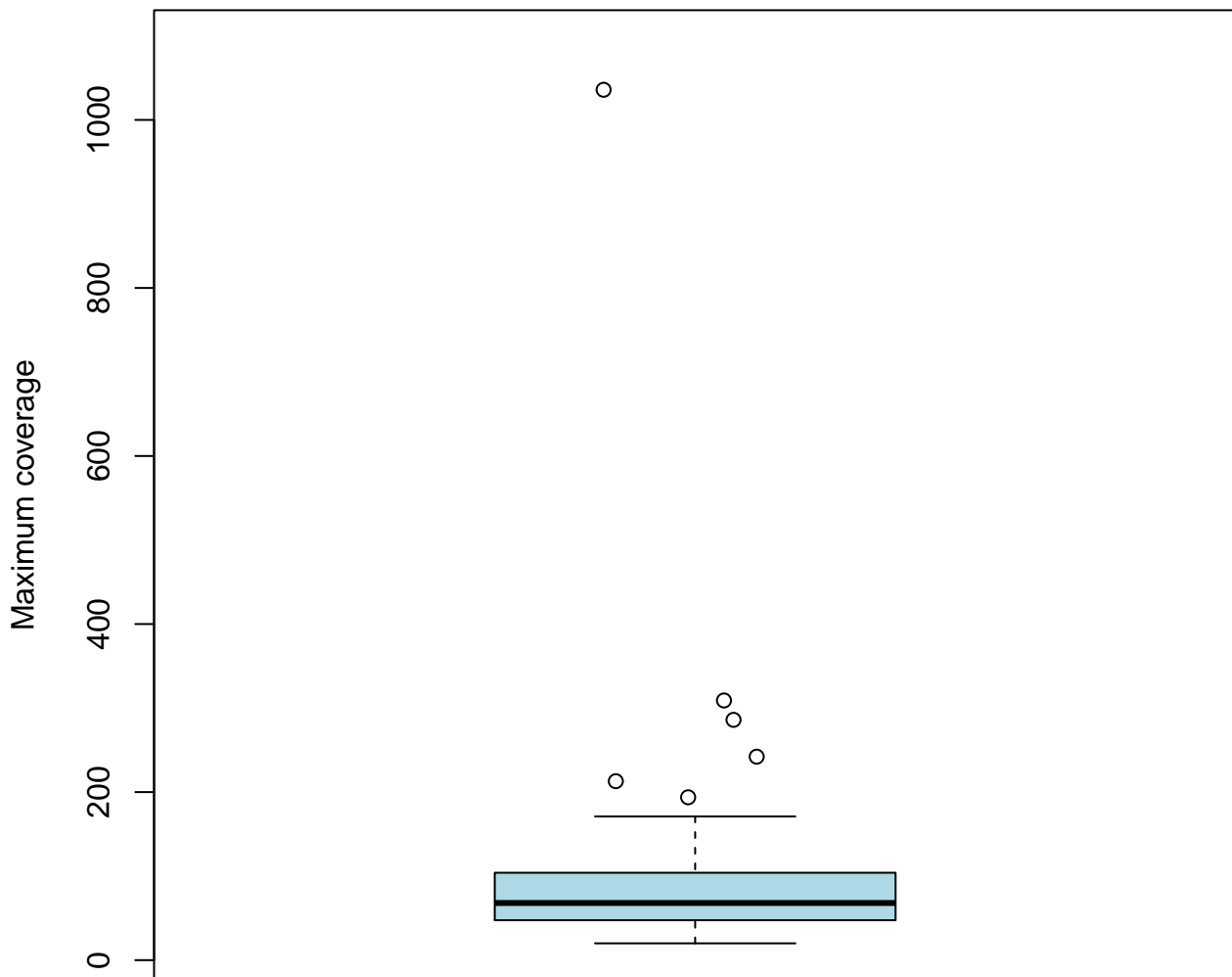
**jx in Intropolis but not "annotated" by knownGene:
248 samples; n jx: 34**



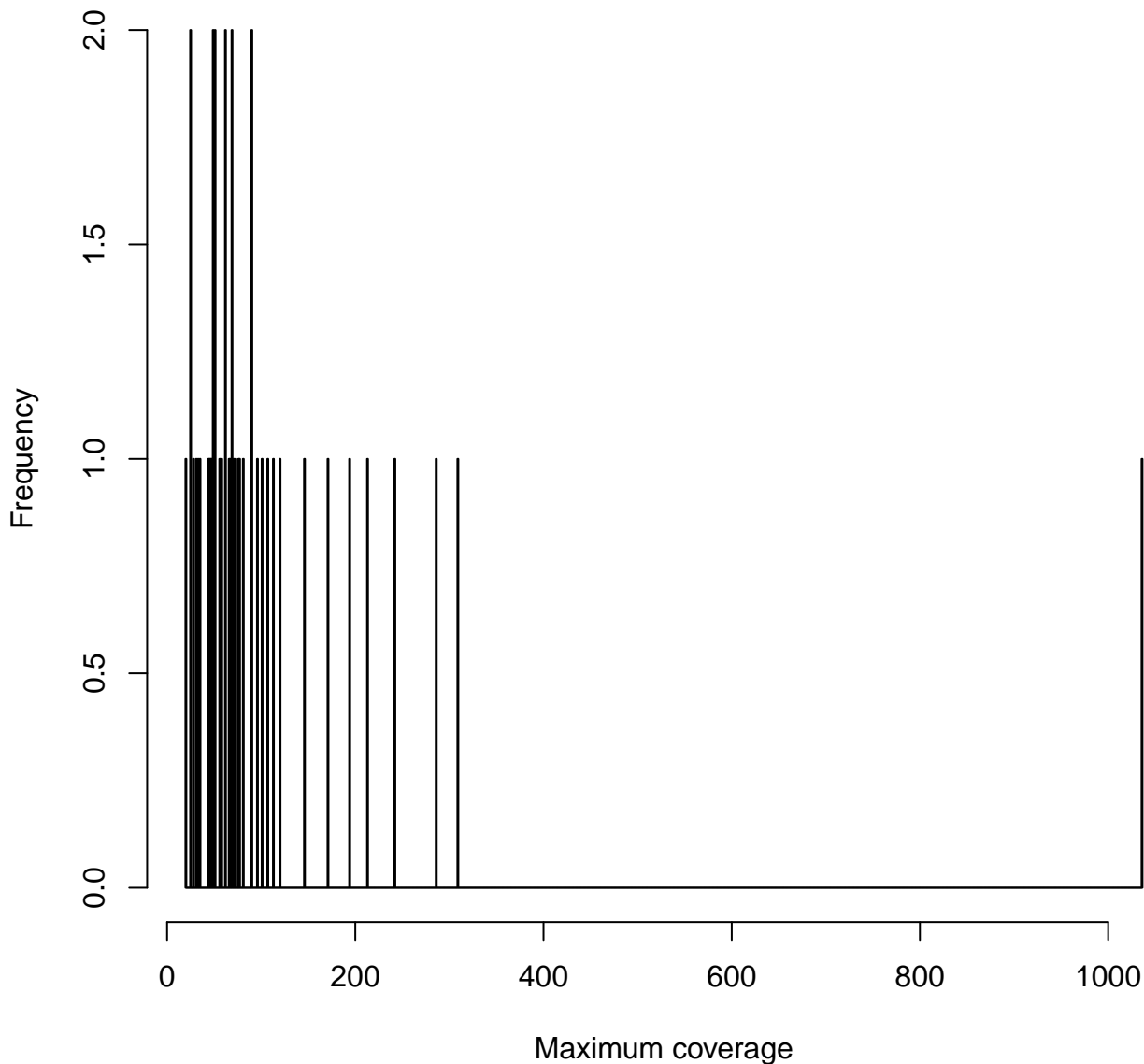
**jx in Intropolis but not "annotated" by knownGene:
248 samples; n jx: 34**



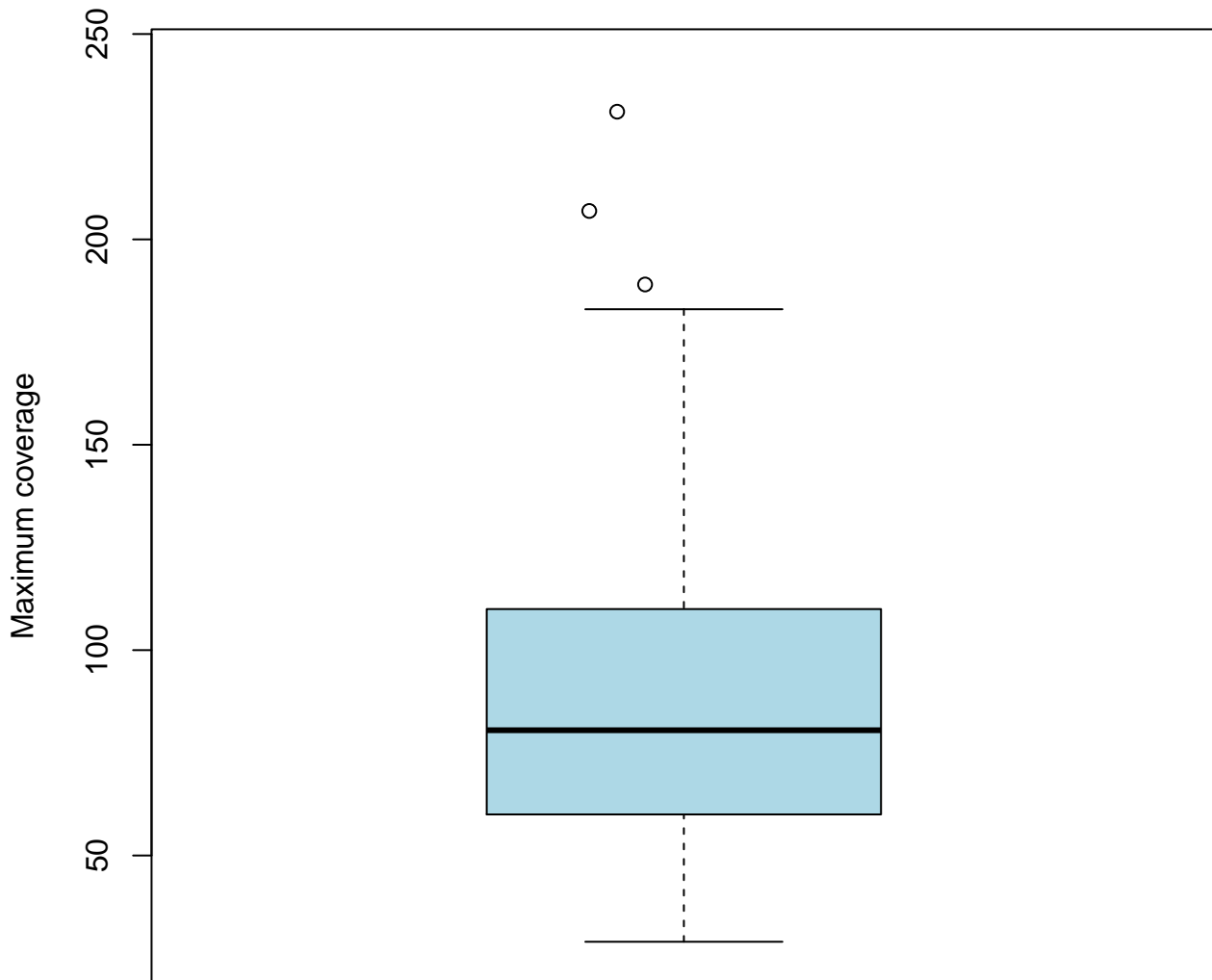
249 samples; n jx: 44



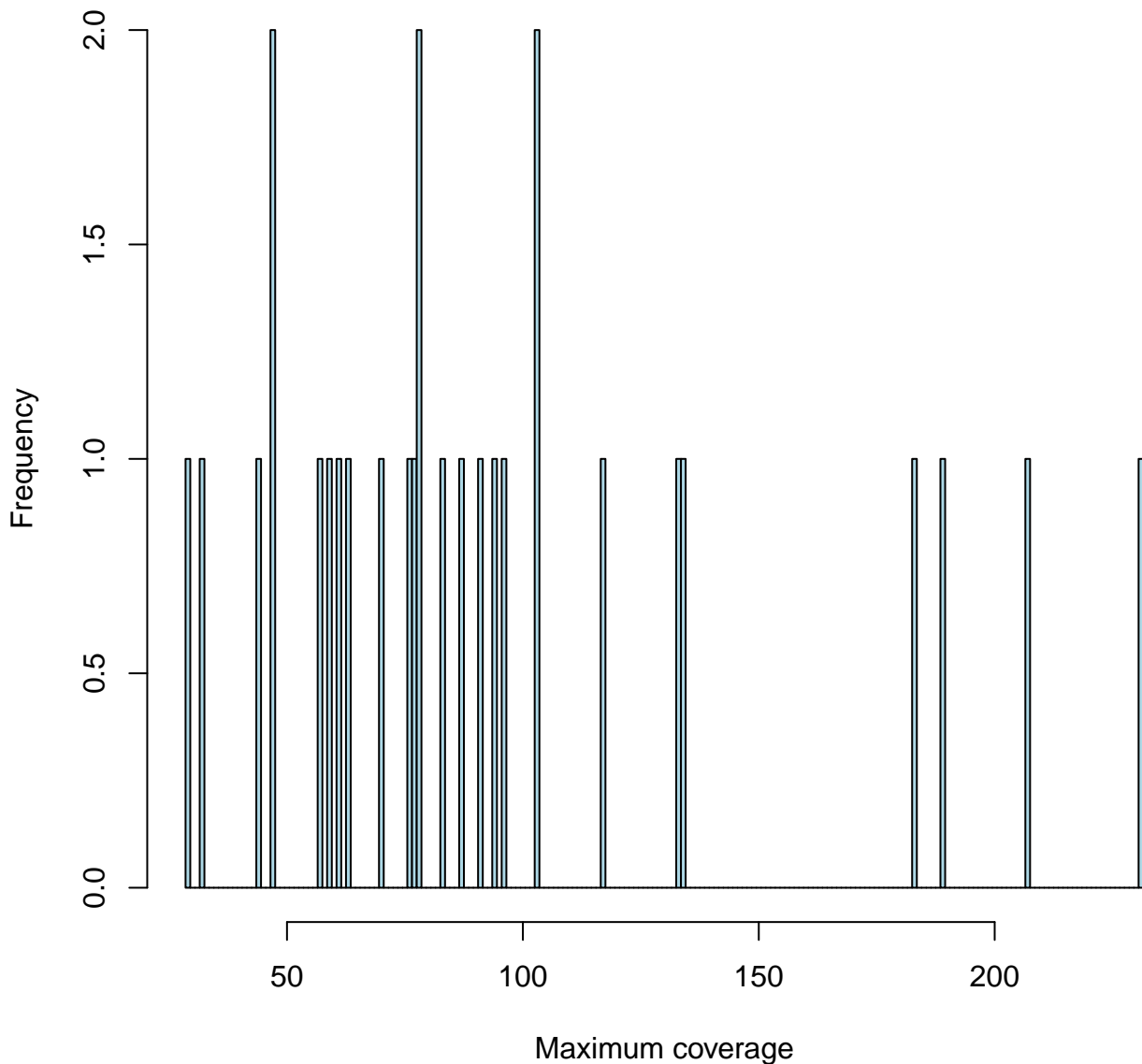
**jx in Intropolis but not "annotated" by knownGene:
249 samples; n jx: 44**



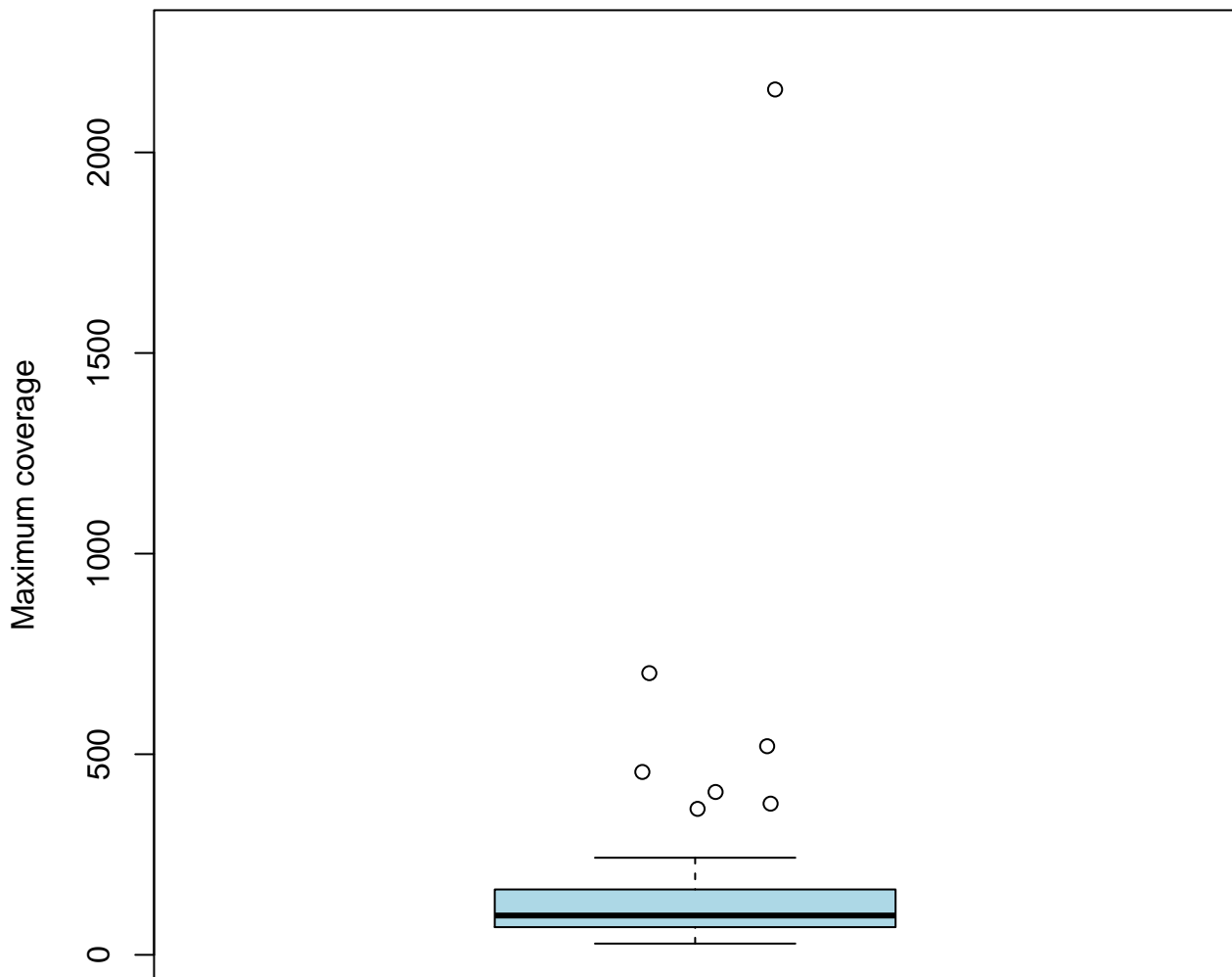
**jx in Intropolis but not "annotated" by knownGene:
250 samples; n jx: 28**



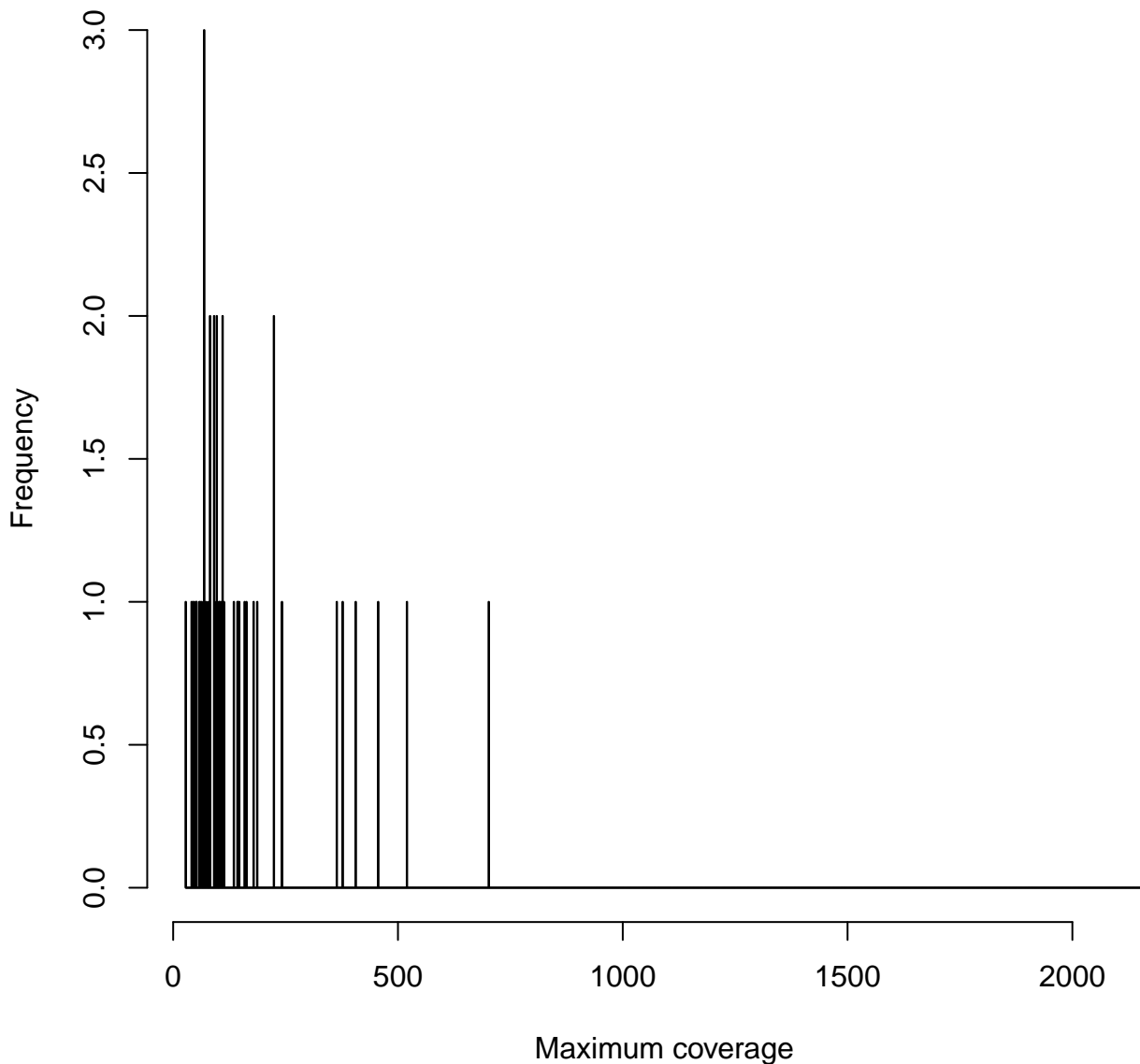
**jx in Intropolis but not "annotated" by knownGene:
250 samples; n jx: 28**



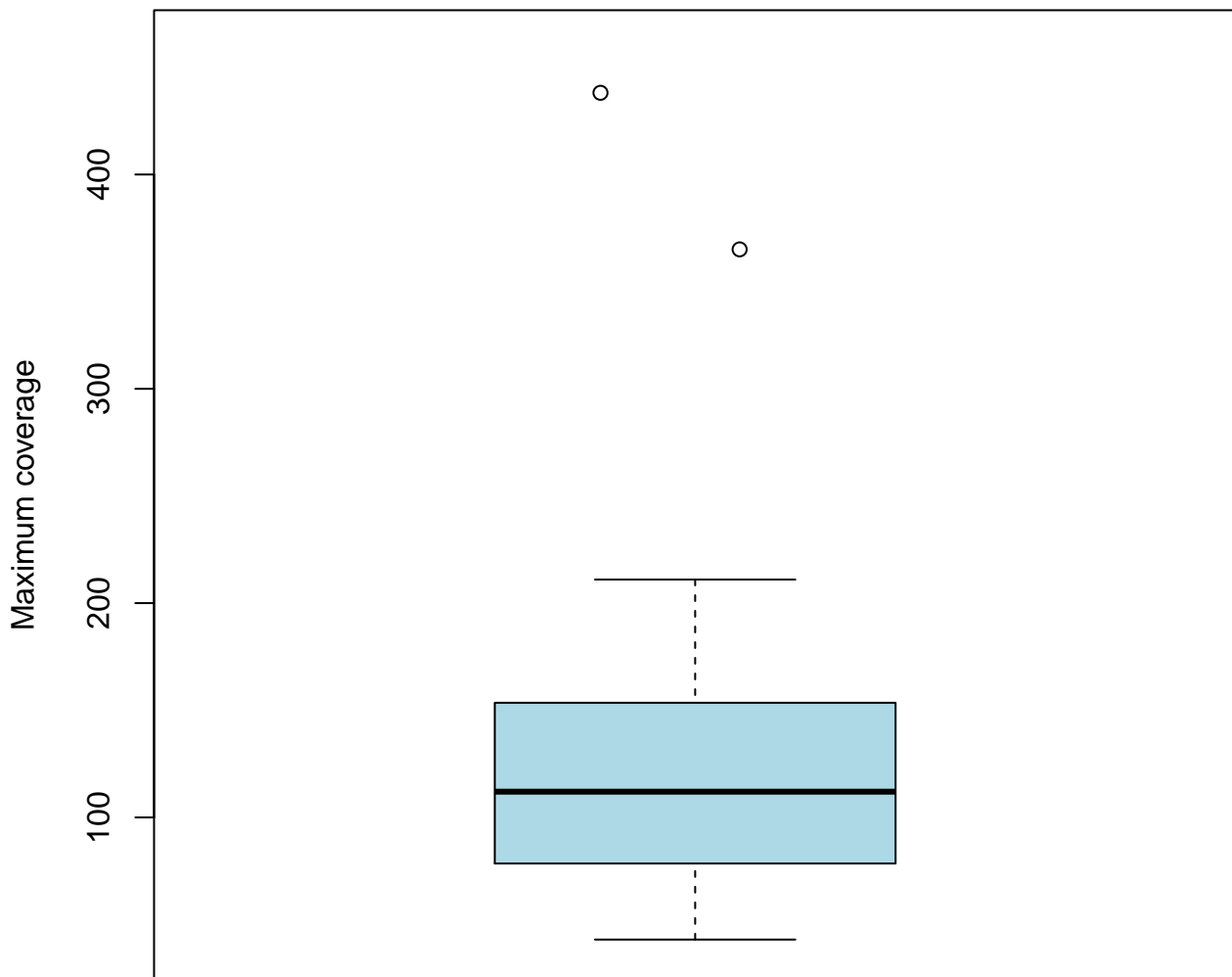
**jx in Intropolis but not "annotated" by knownGene:
251 samples; n jx: 49**



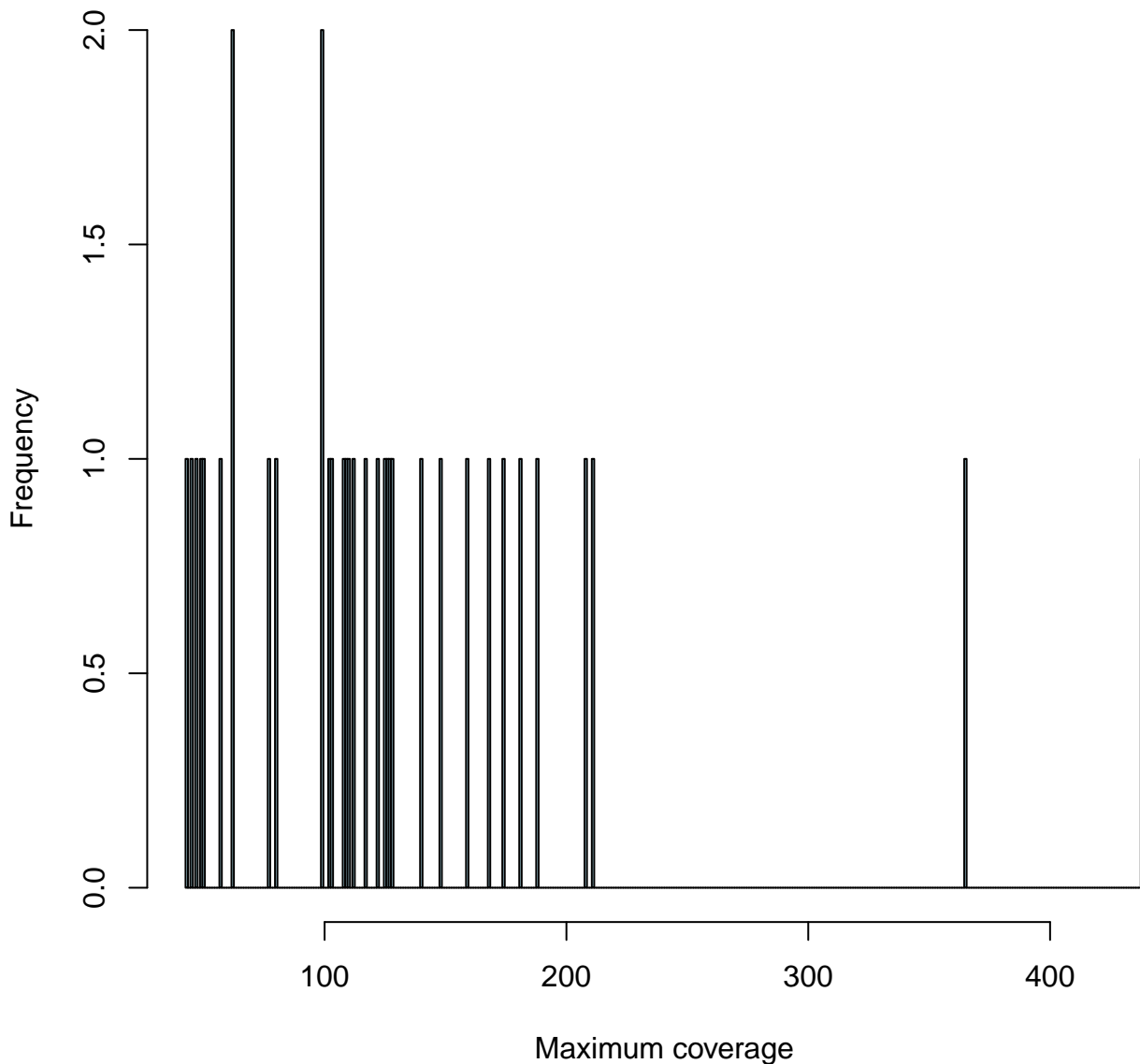
**jx in Intropolis but not "annotated" by knownGene:
251 samples; n jx: 49**



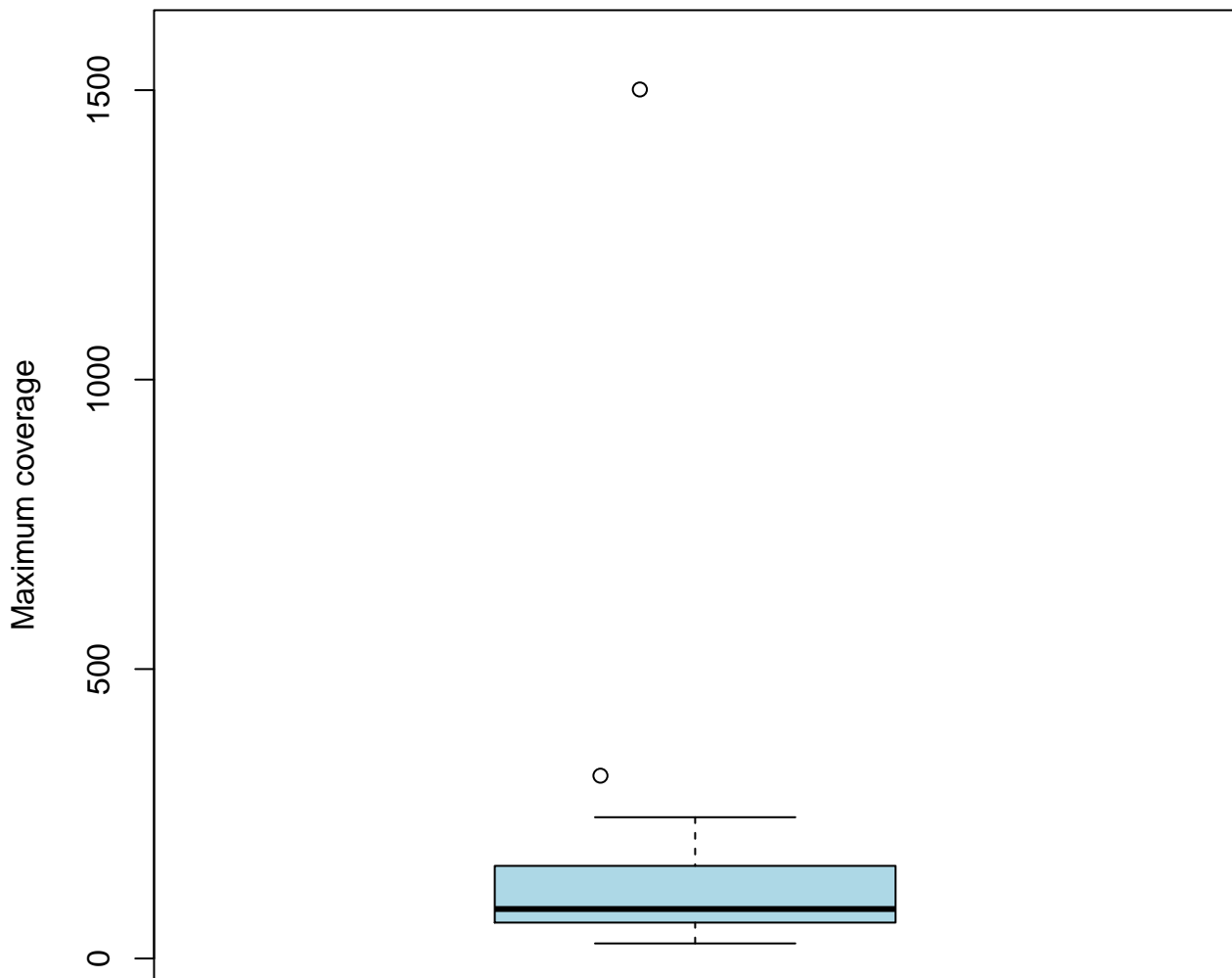
**jx in Intropolis but not "annotated" by knownGene:
252 samples; n jx: 35**



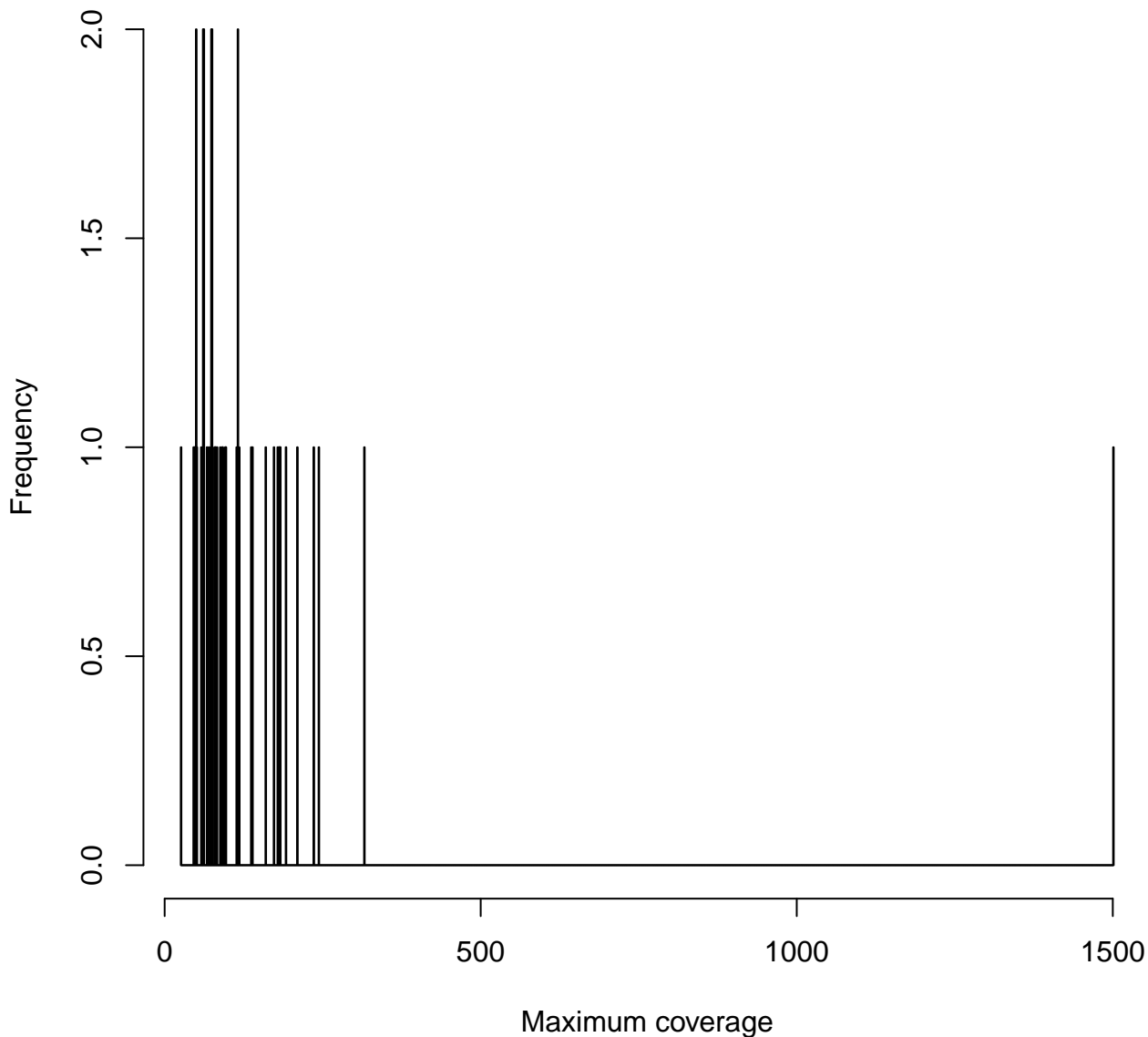
**jx in Intropolis but not "annotated" by knownGene:
252 samples; n jx: 35**



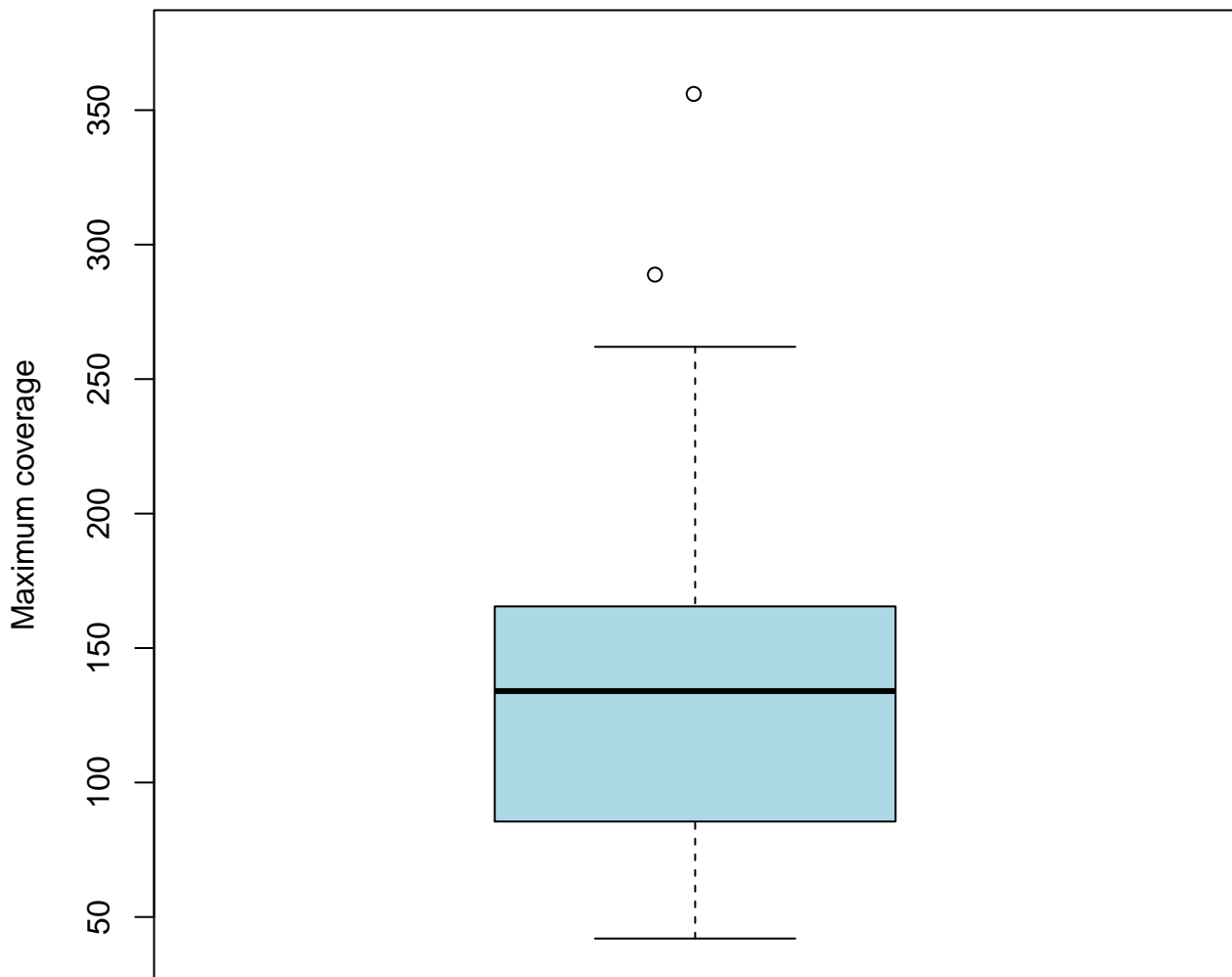
**jx in Intropolis but not "annotated" by knownGene:
253 samples; n jx: 46**



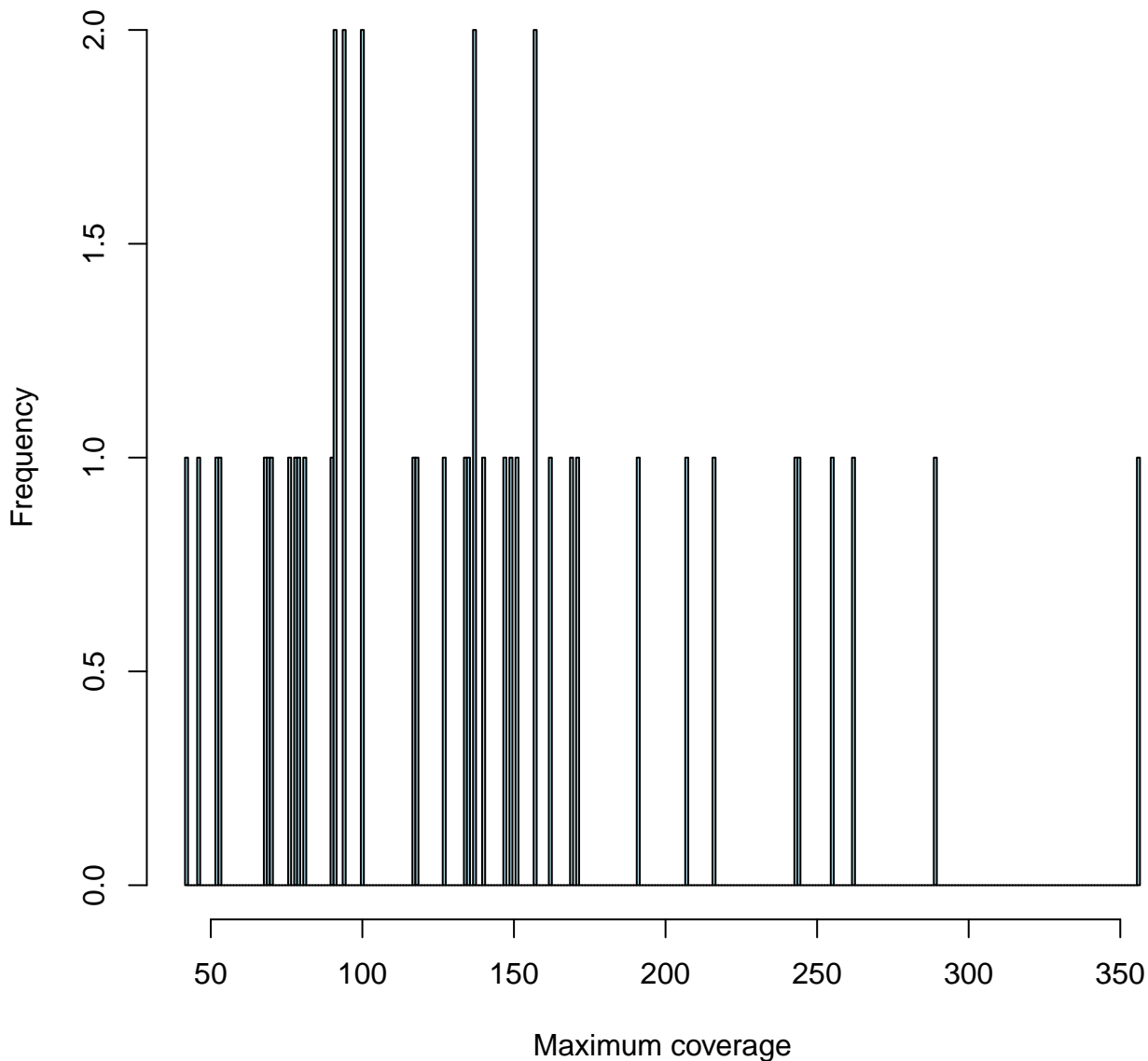
**jx in Intropolis but not "annotated" by knownGene:
253 samples; n jx: 46**



**jx in Intropolis but not "annotated" by knownGene:
254 samples; n jx: 43**



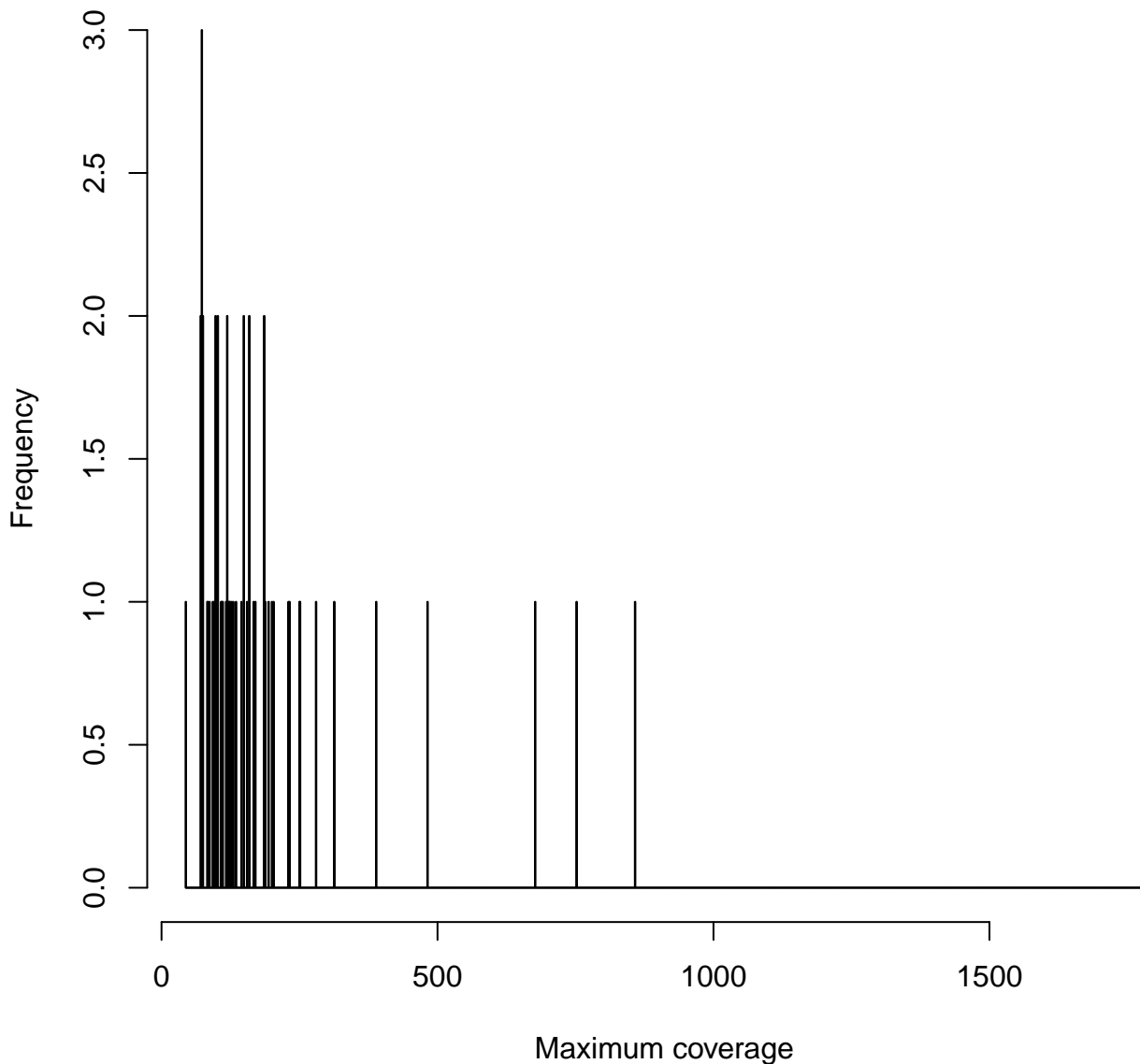
**jx in Intropolis but not "annotated" by knownGene:
254 samples; n jx: 43**



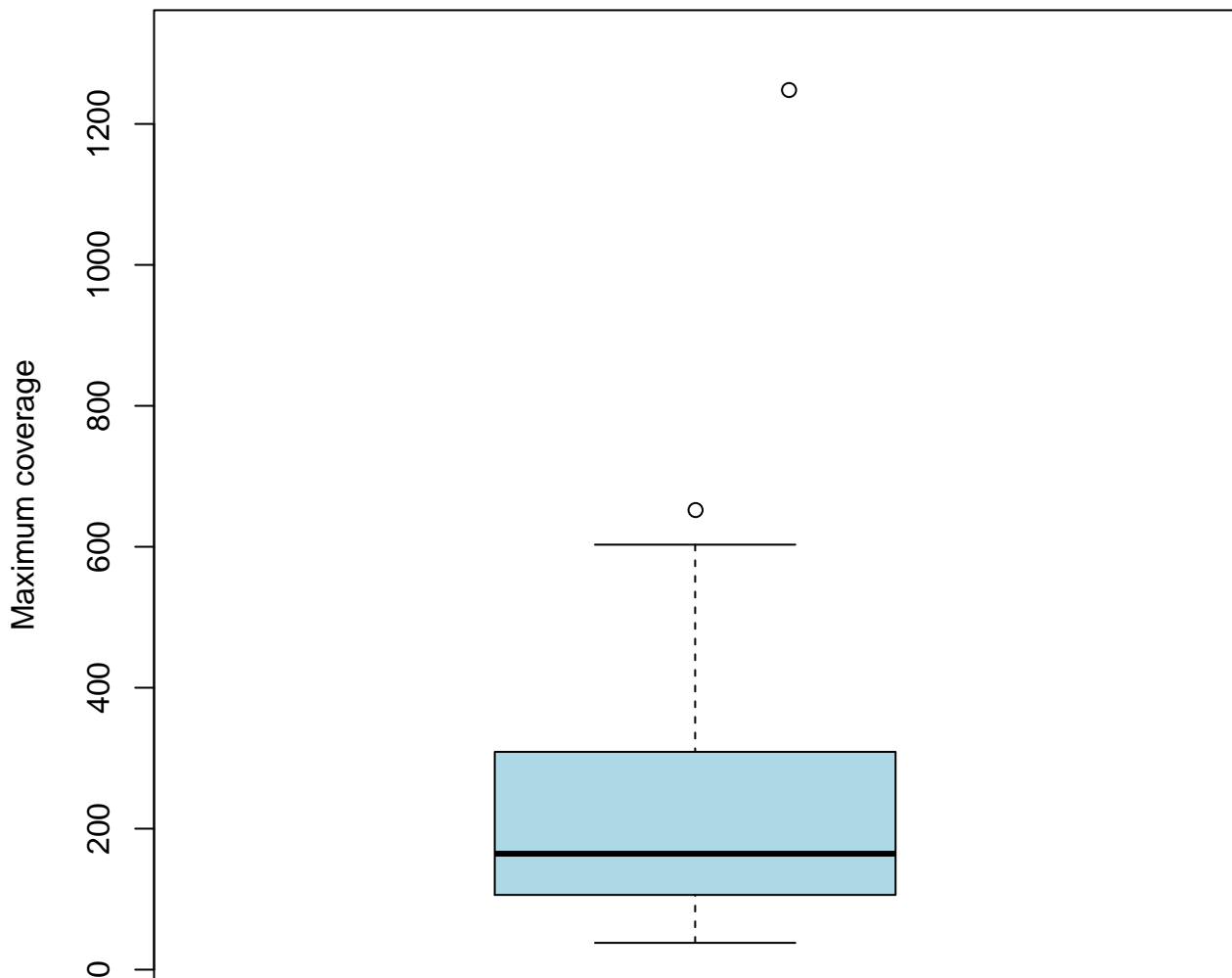
255 samples; n jx: 58



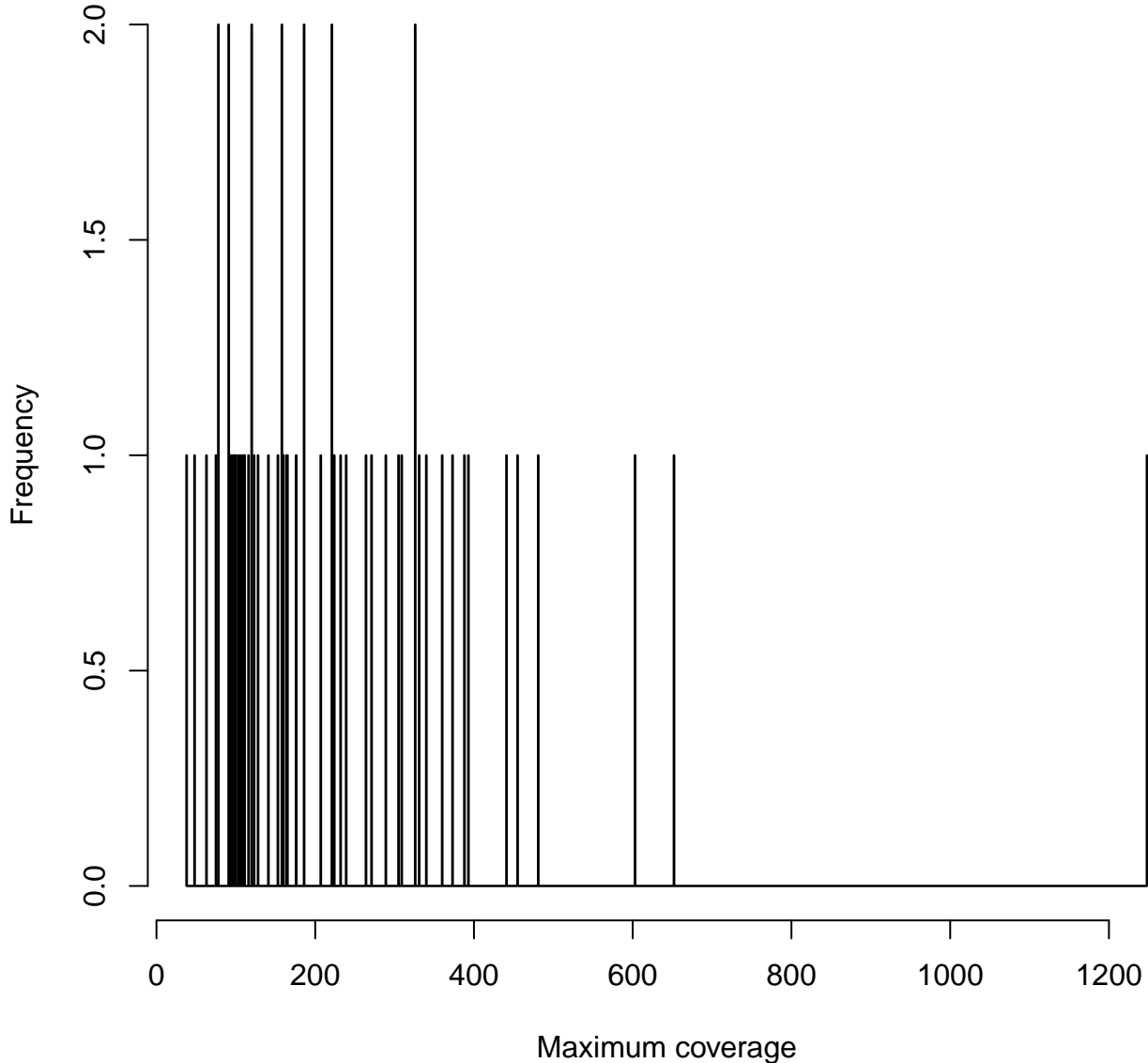
**jx in Intropolis but not "annotated" by knownGene:
255 samples; n jx: 58**



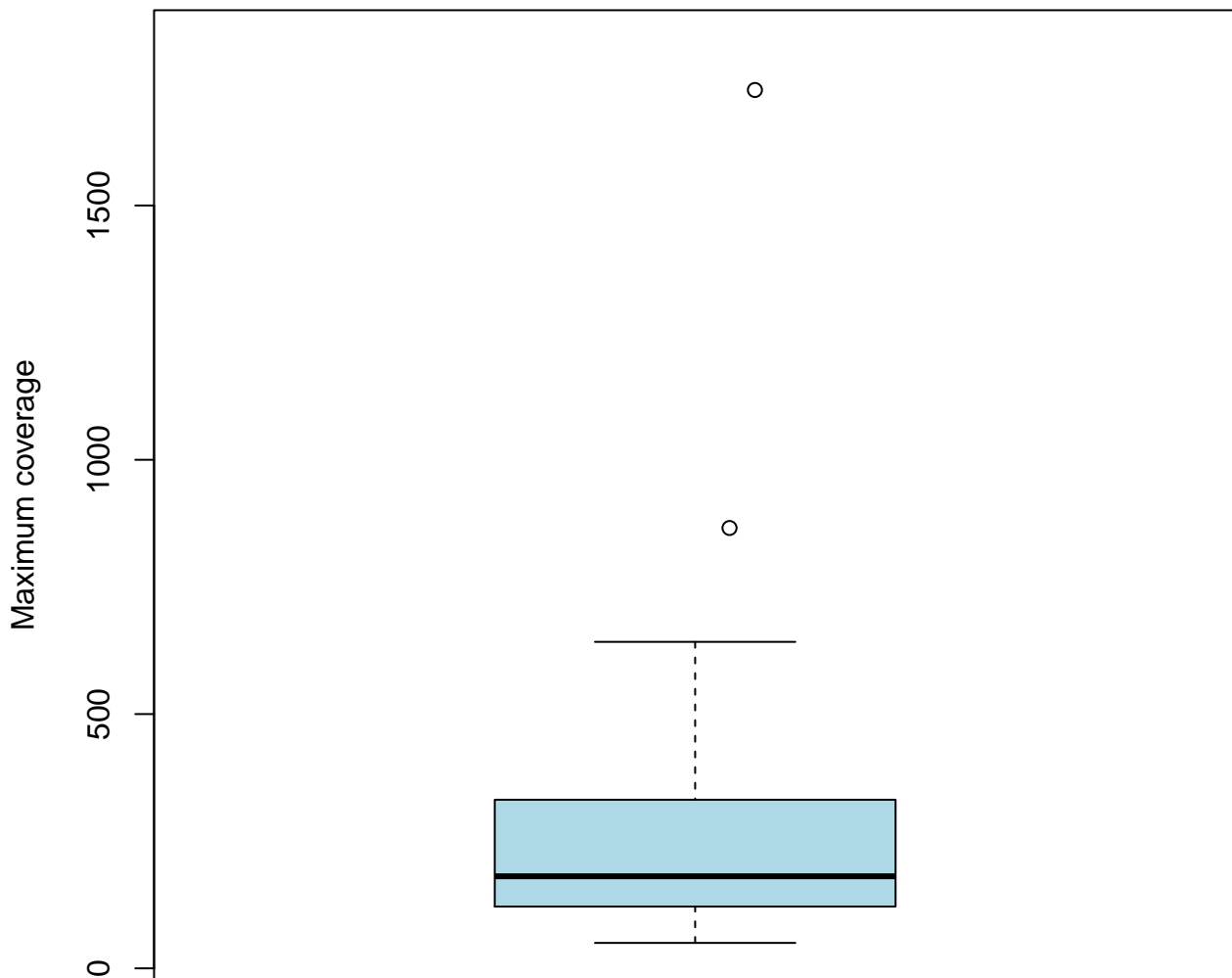
**jx in Intropolis but not "annotated" by knownGene:
256 samples; n jx: 58**



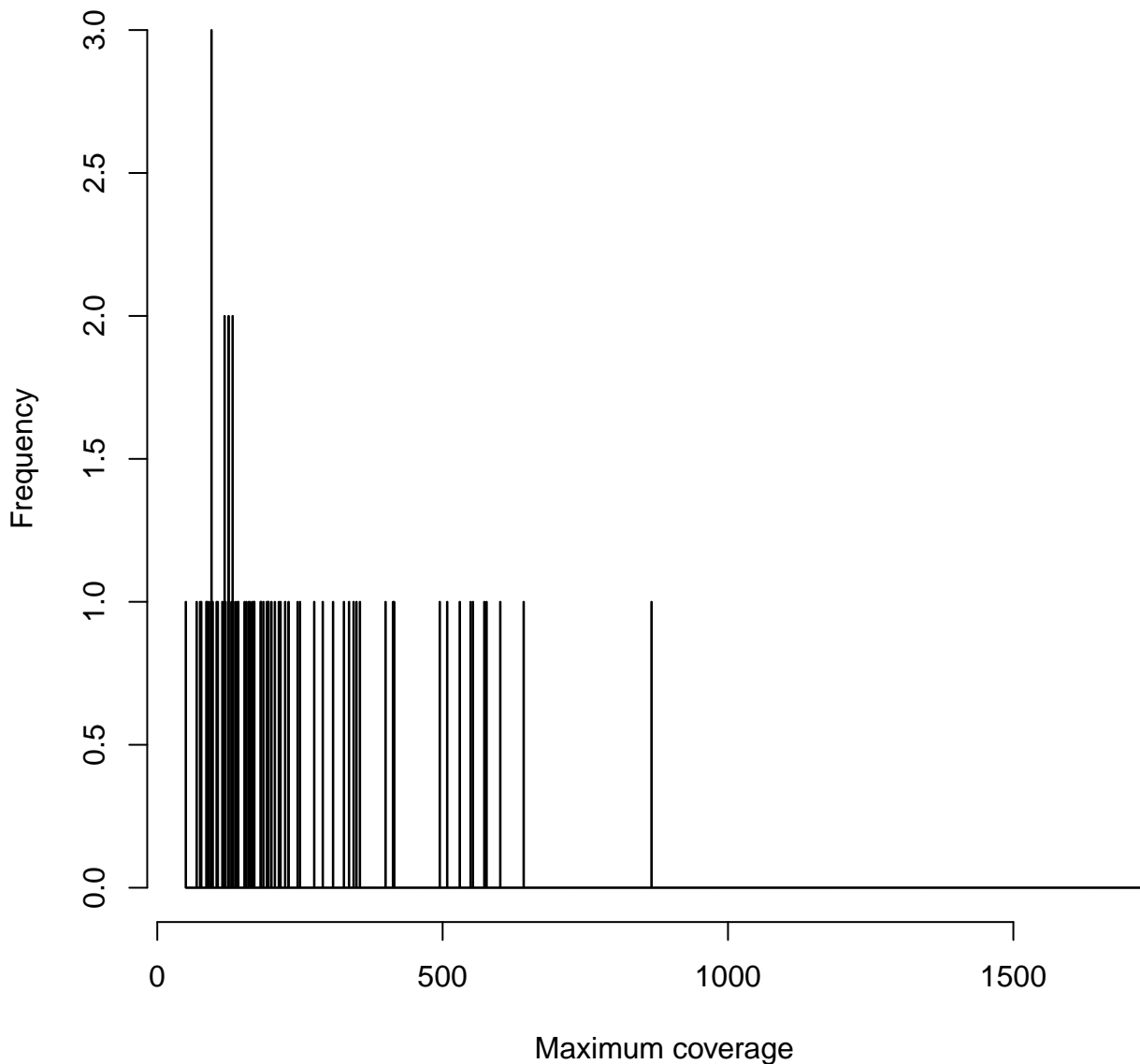
**jx in Intropolis but not "annotated" by knownGene:
256 samples; n jx: 58**



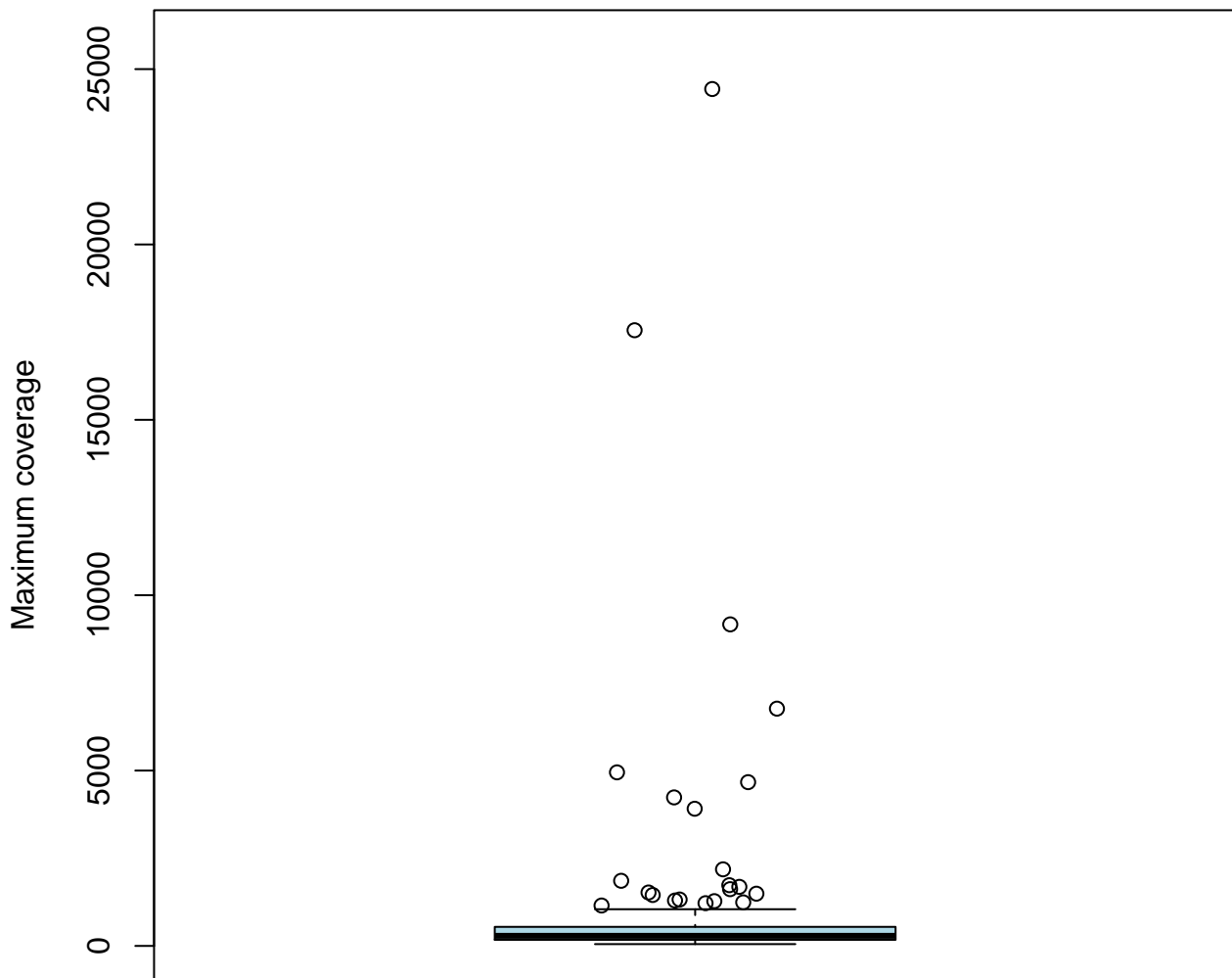
**jx in Intropolis but not "annotated" by knownGene:
257 samples; n jx: 71**



**jx in Intropolis but not "annotated" by knownGene:
257 samples; n jx: 71**



**jx in Intropolis but not "annotated" by knownGene:
258 samples; n jx: 213**



**jx in Intropolis but not "annotated" by knownGene:
258 samples; n jx: 213**

