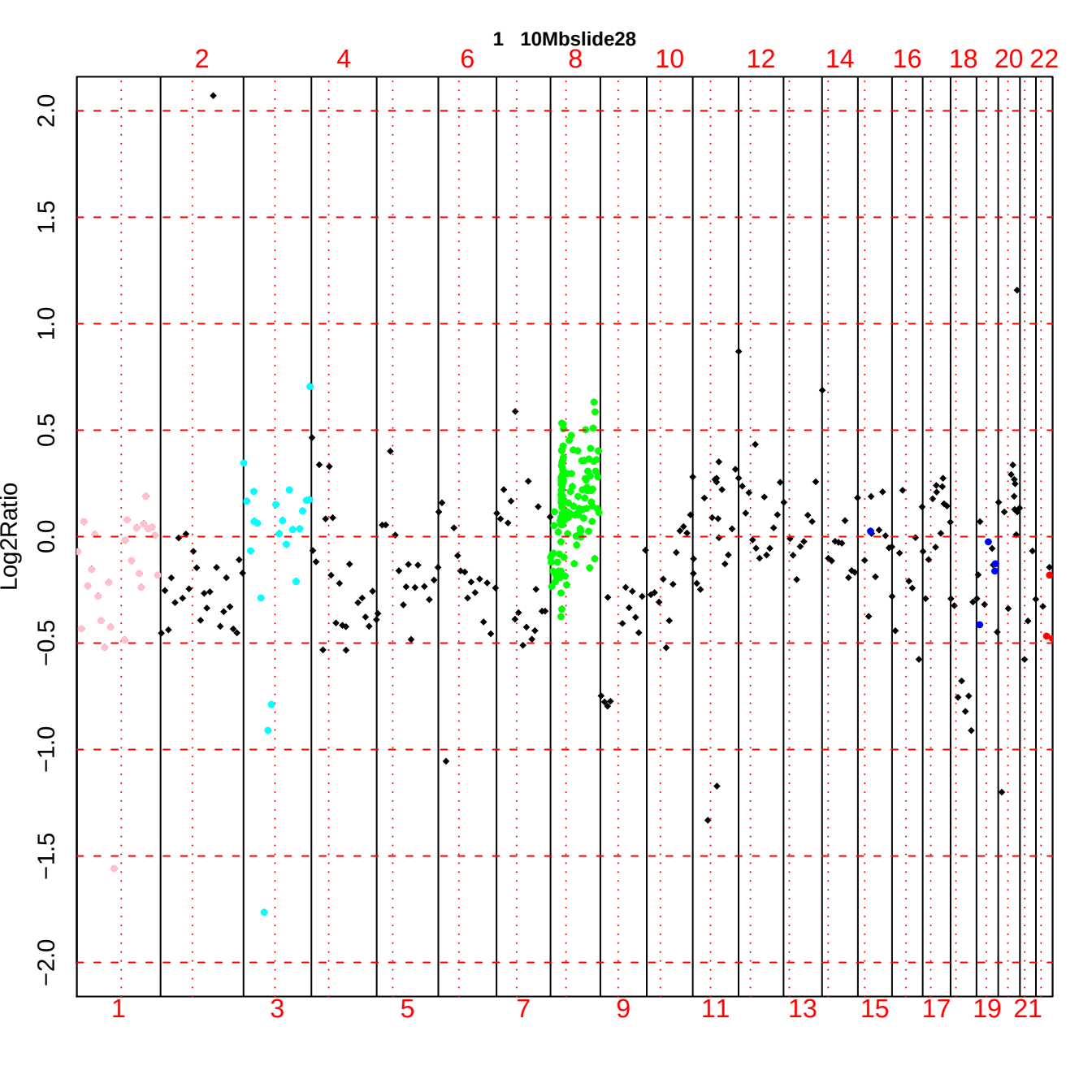
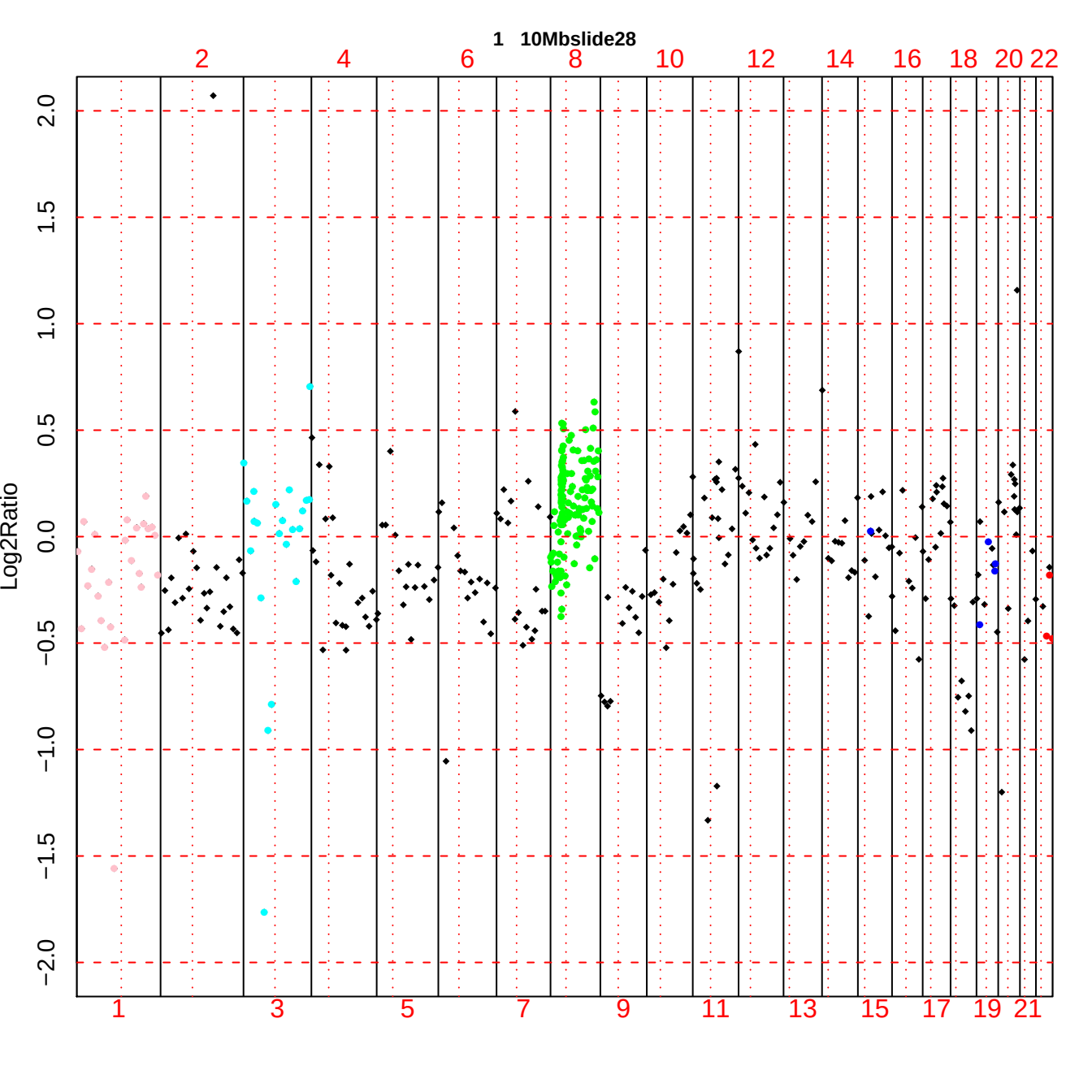

1 10Mbslide28
2
4
6
8
10
12
14
16
18
20
22
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
Log2Ratio
1
3
5
7
9
11
13
15
17
19
21


1 10Mbslide28
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1.5
1.0
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0.0
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Log2Ratio
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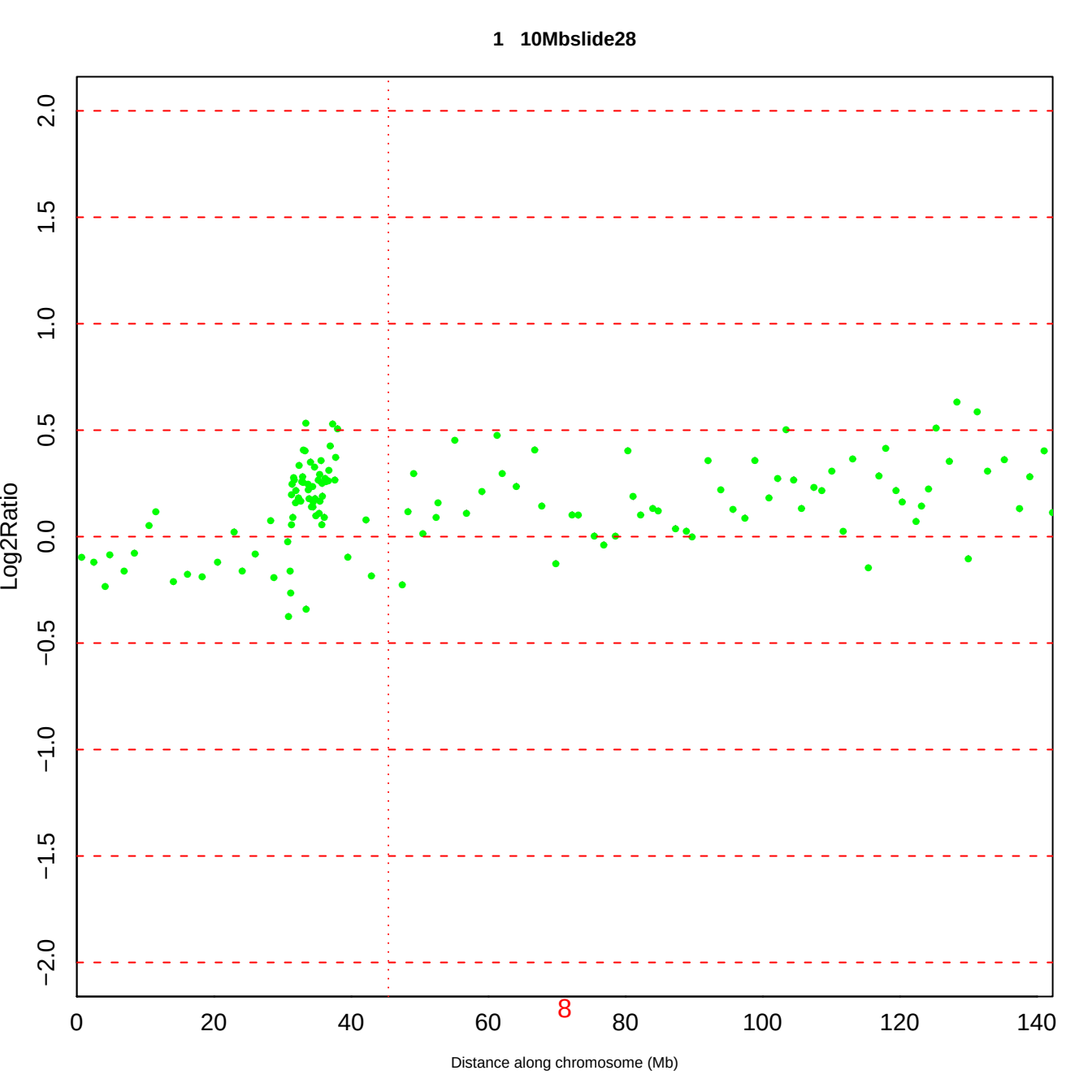
19
21


1 10Mbslide28
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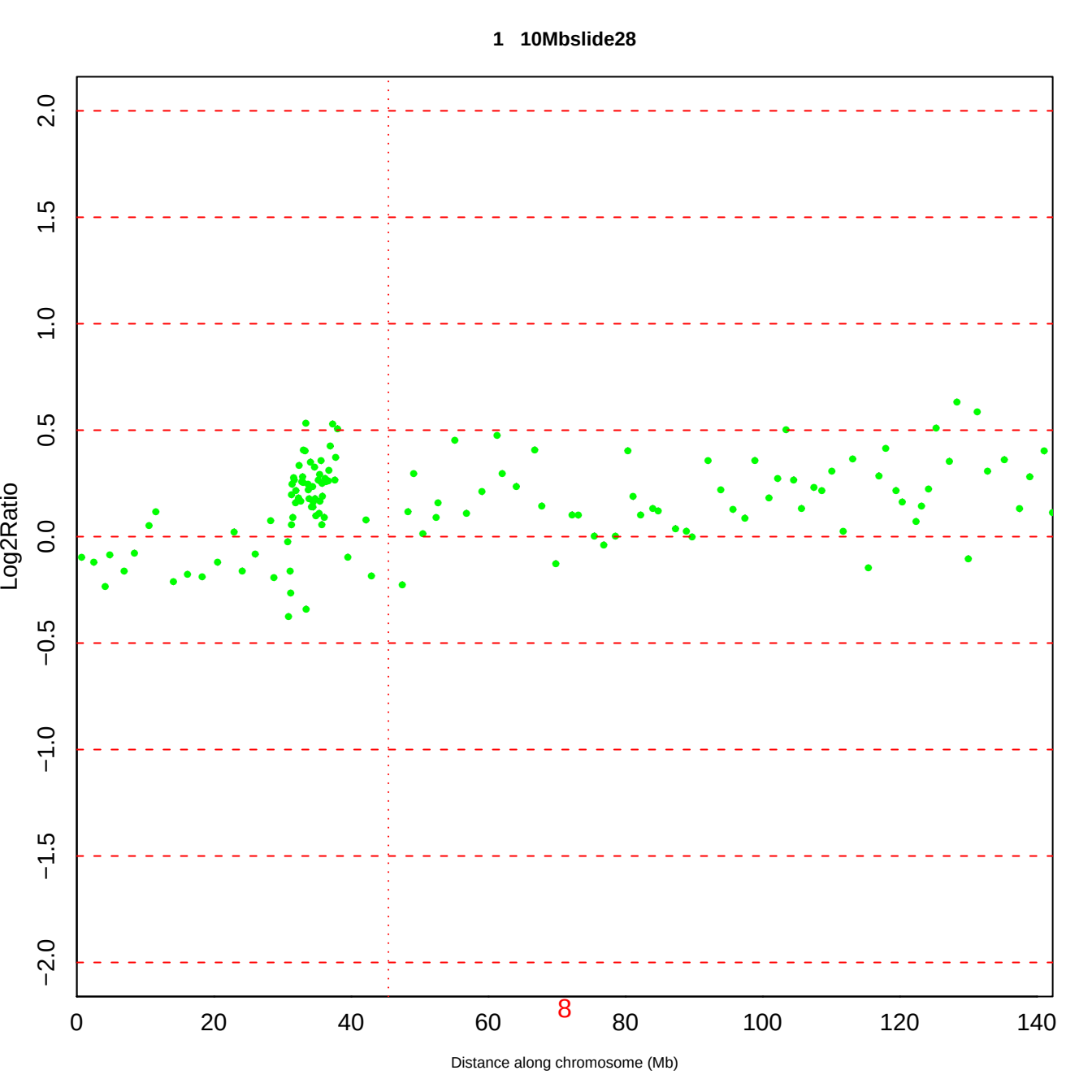
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
0
20
40
60
80
100
120
140
Distance along chromosome (Mb)


1 10Mbslide28
Log2Ratio
2.0
1.5
1.0
0.5
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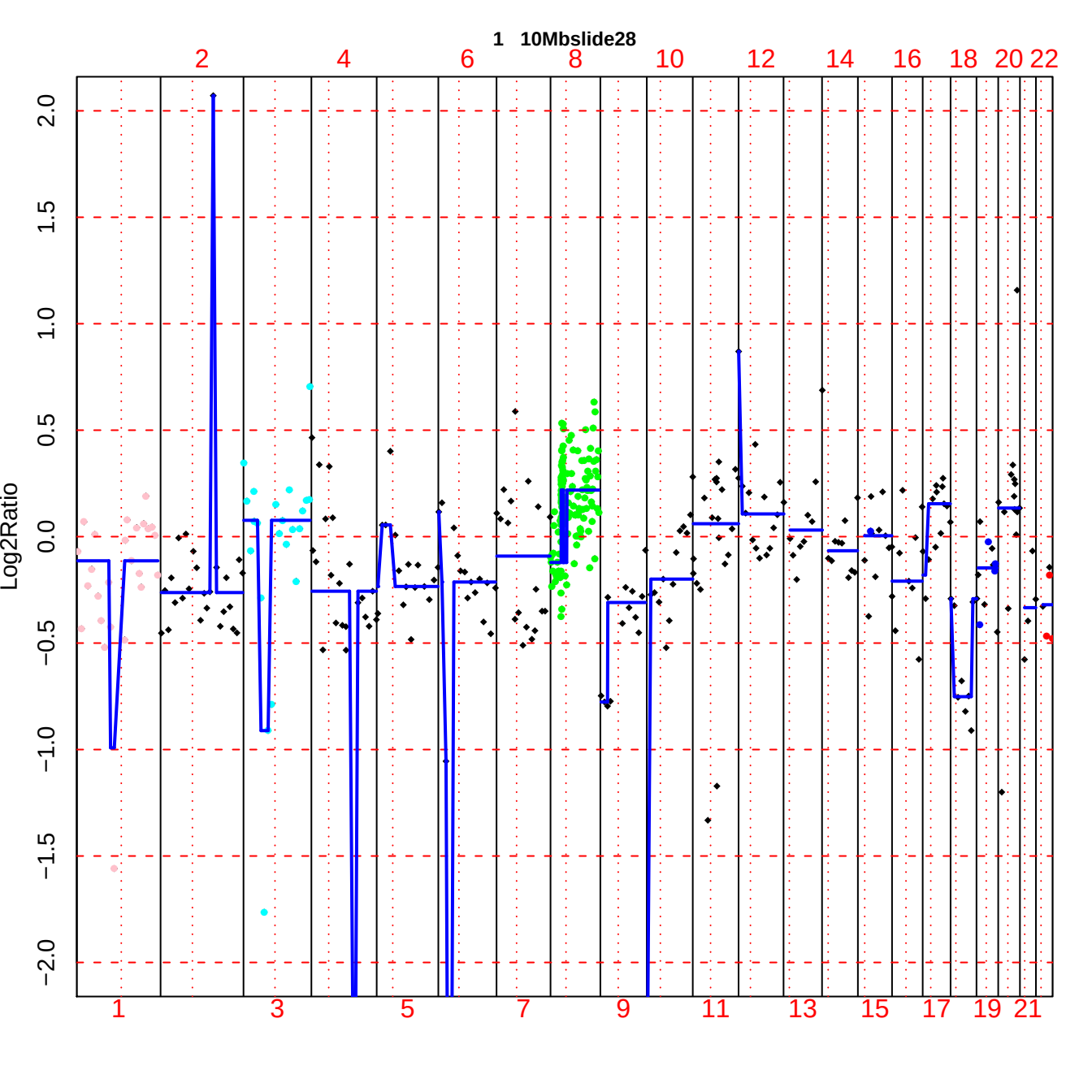
−0.5
−1.0
−1.5
−2.0
0
20
40
60
80
100
120
140
Distance along chromosome (Mb)


1 10Mbslide28
2
4
6
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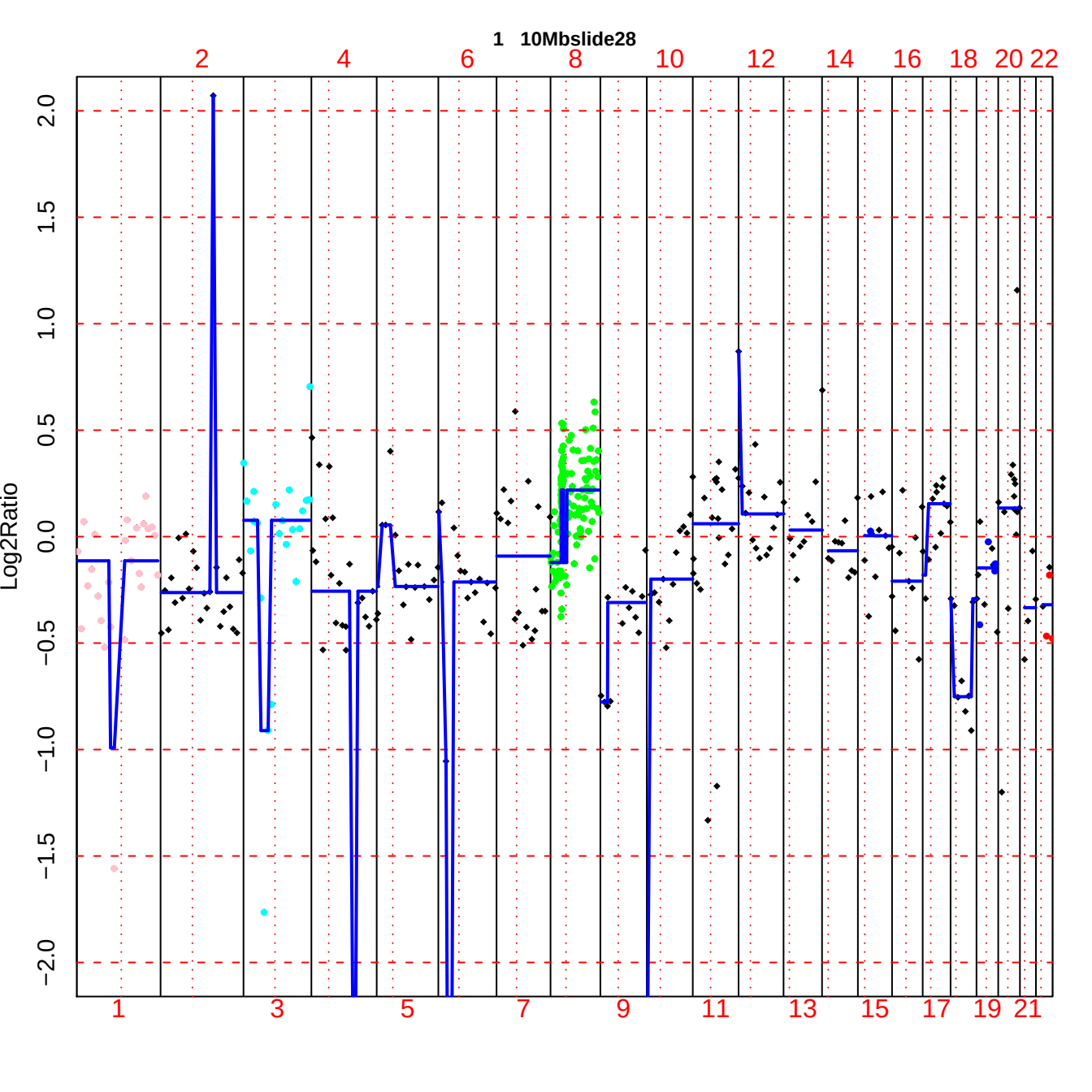
10
12
14
16
18
20
22
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
Log2Ratio
1
3
5
7
9
11
13
15
17
19
21


1 10Mbslide28
2
4
6
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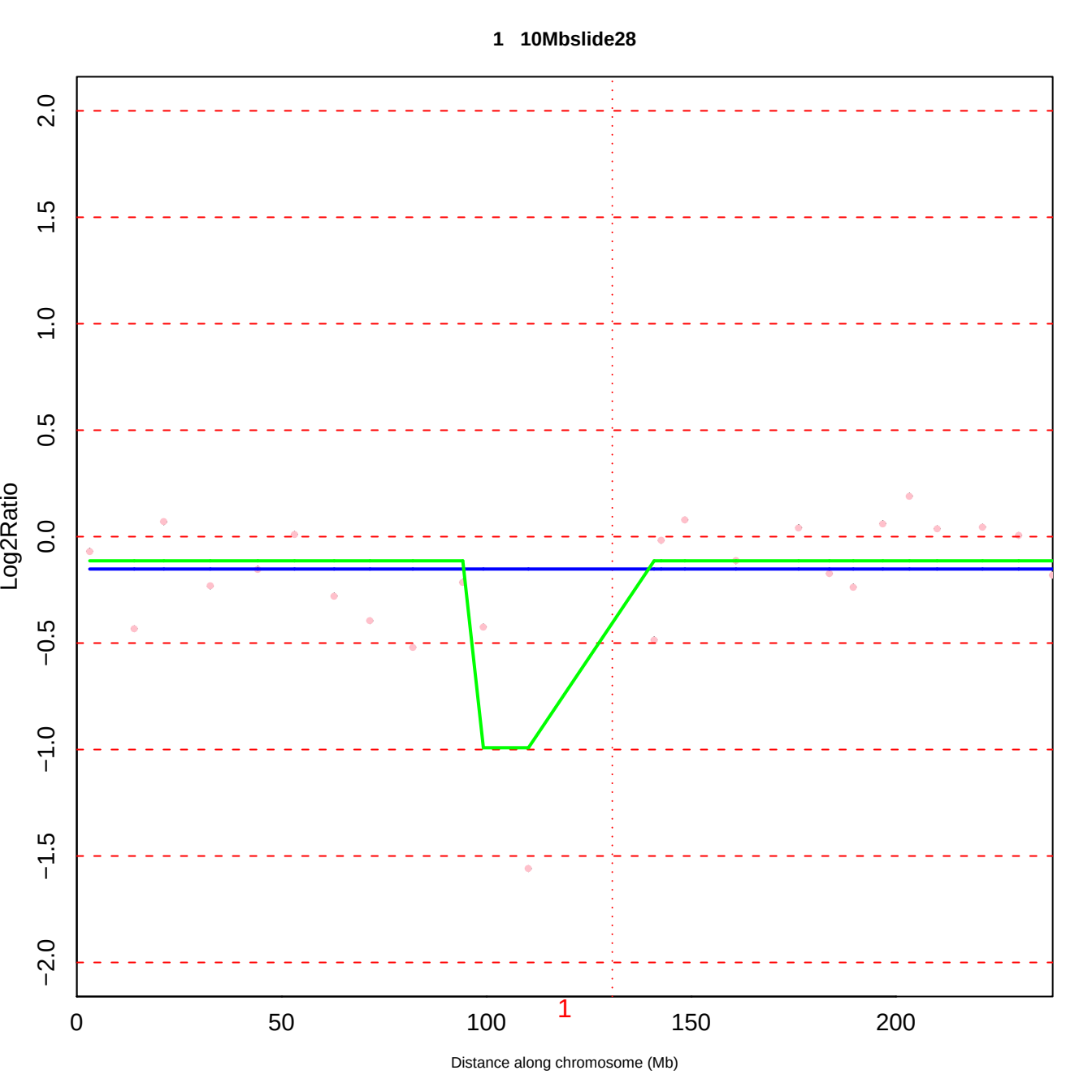
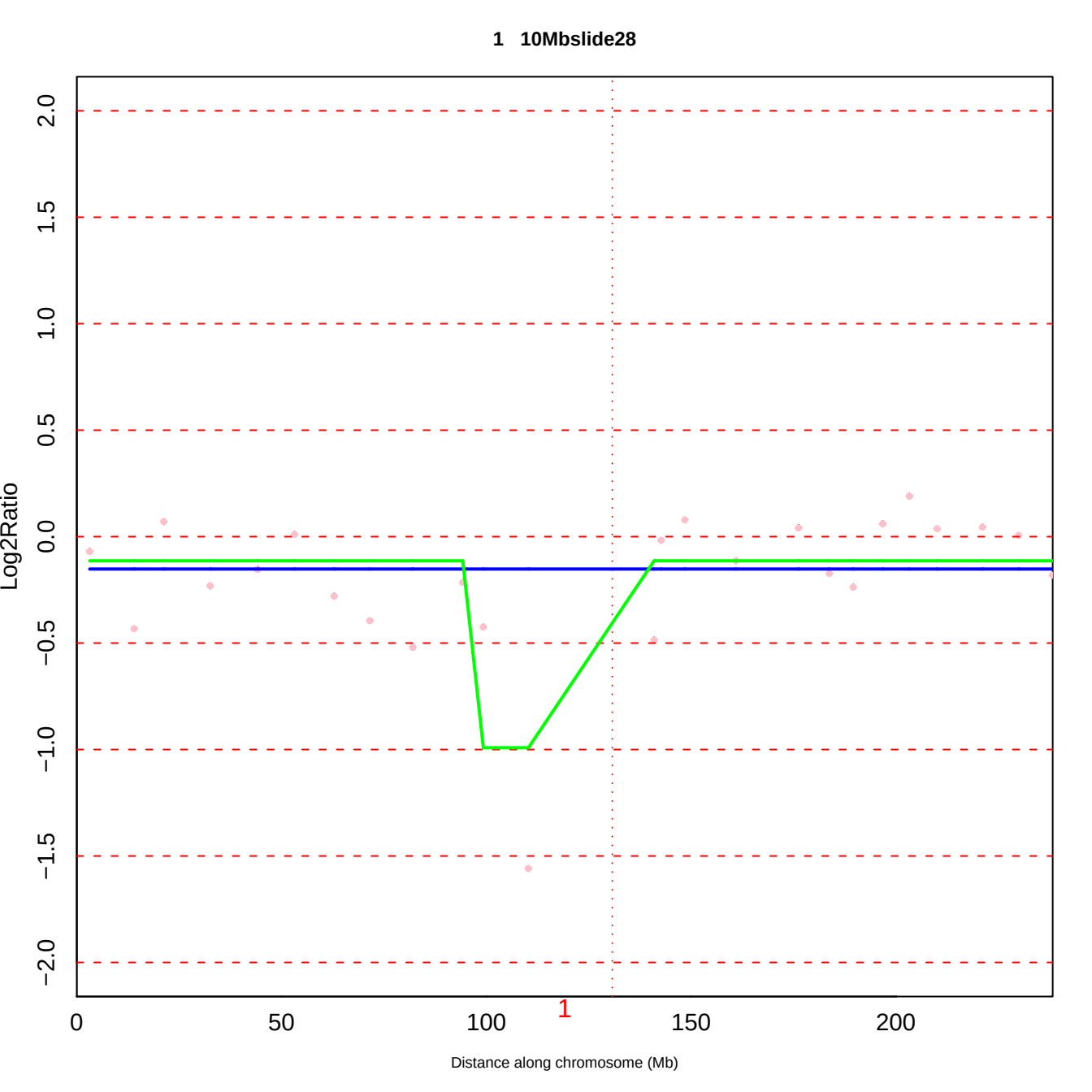
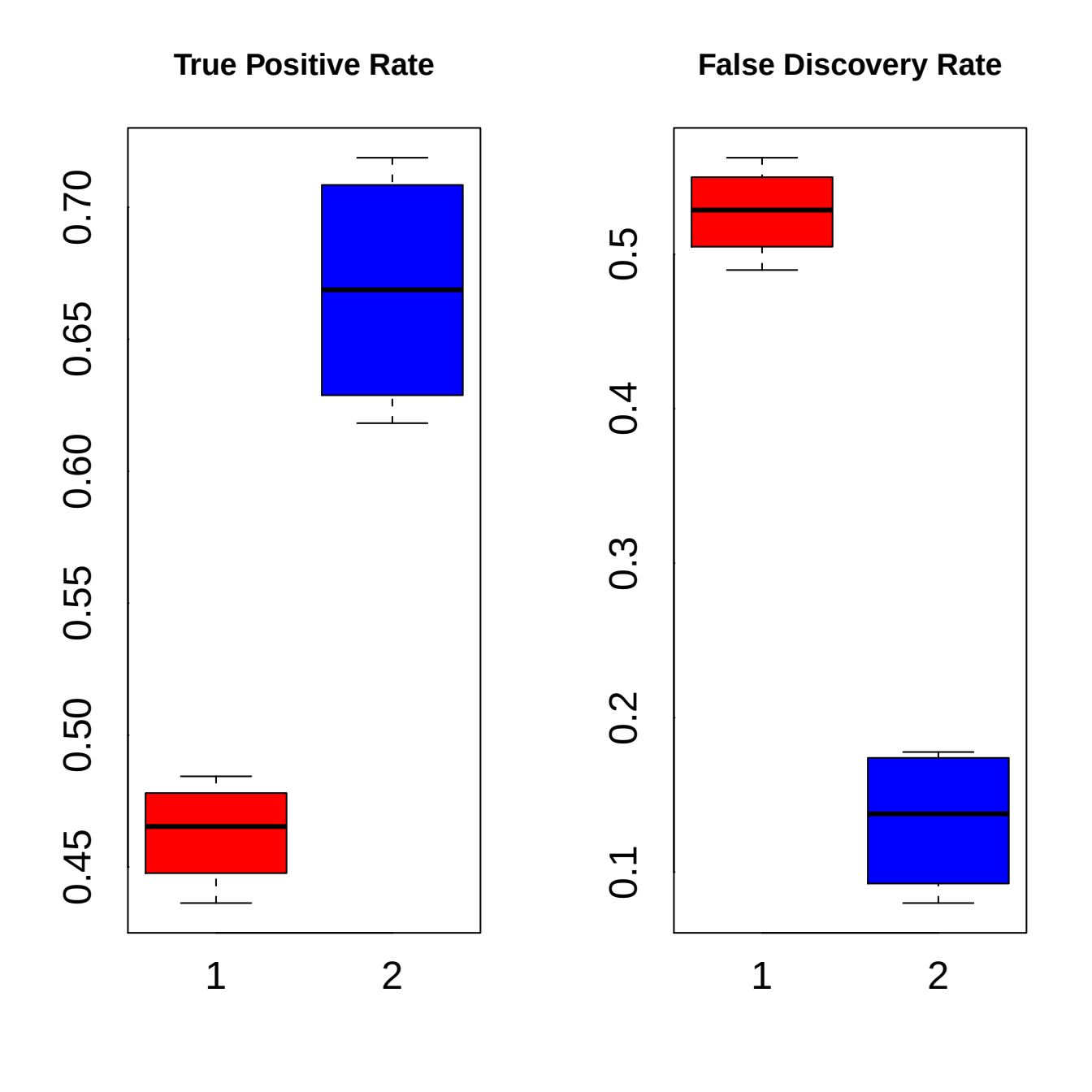
16
18
20
22
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
Log2Ratio
1
3
5
7
9
11
13
15
17
19
21


1 10Mbslide28
Log2Ratio
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
0
50
100
150
200
1
Distance along chromosome (Mb)


1 10Mbslide28
Log2Ratio
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
0
50
100
150
200
1
Distance along chromosome (Mb)


True Positive Rate
0.45
0.50
0.55
0.60
0.65
0.70
1
2
False Discovery Rate
0.1
0.2
0.3
0.4
0.5
1
2