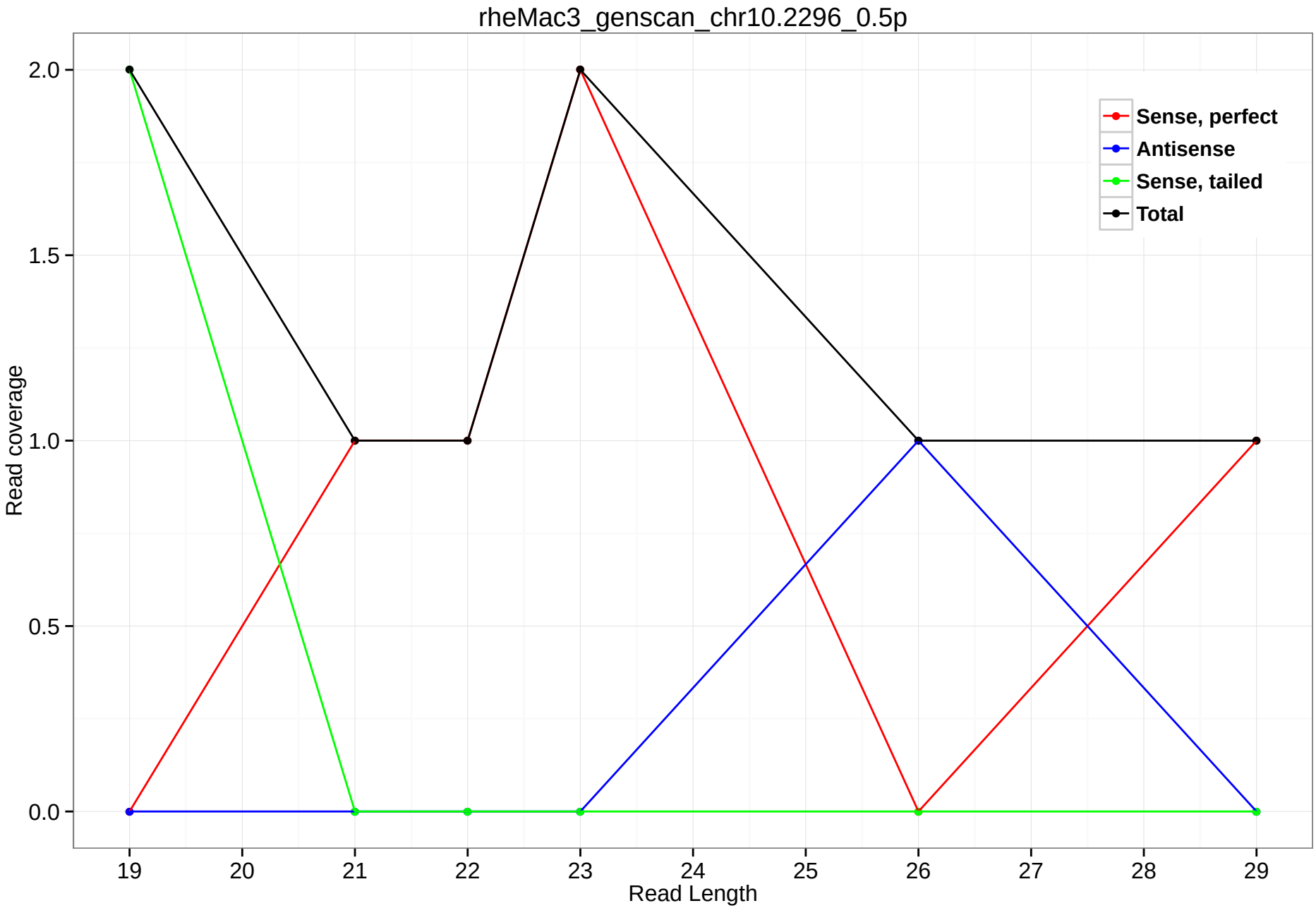
rheMac3_genscan_chr10.2296_0.5p
Read coverage
2.0
1.5
1.0
0.5
0.0
19
20
21
22
23
24
25
26
27
28
29
Read Length
Sense, perfect
Antisense
Sense, tailed
Total