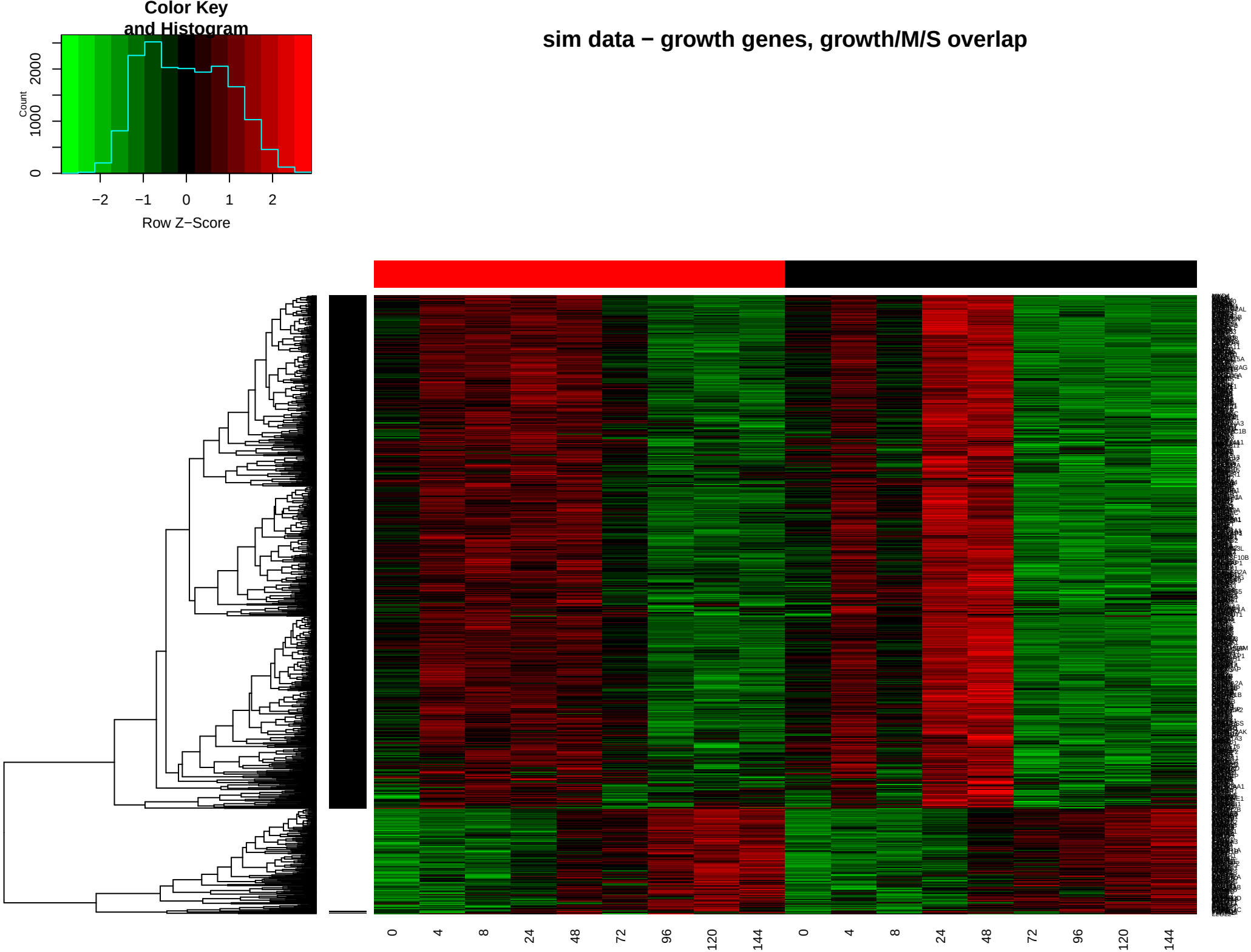
Color Key
and Histogram
Count
0
1000
2000
−2
−1
0
1
2
Row Z−Score
sim data − growth genes, growth/M/S overlap
0
4
8
24
48
72
96
120
144
0
4
8
24
48
72
96
120
144