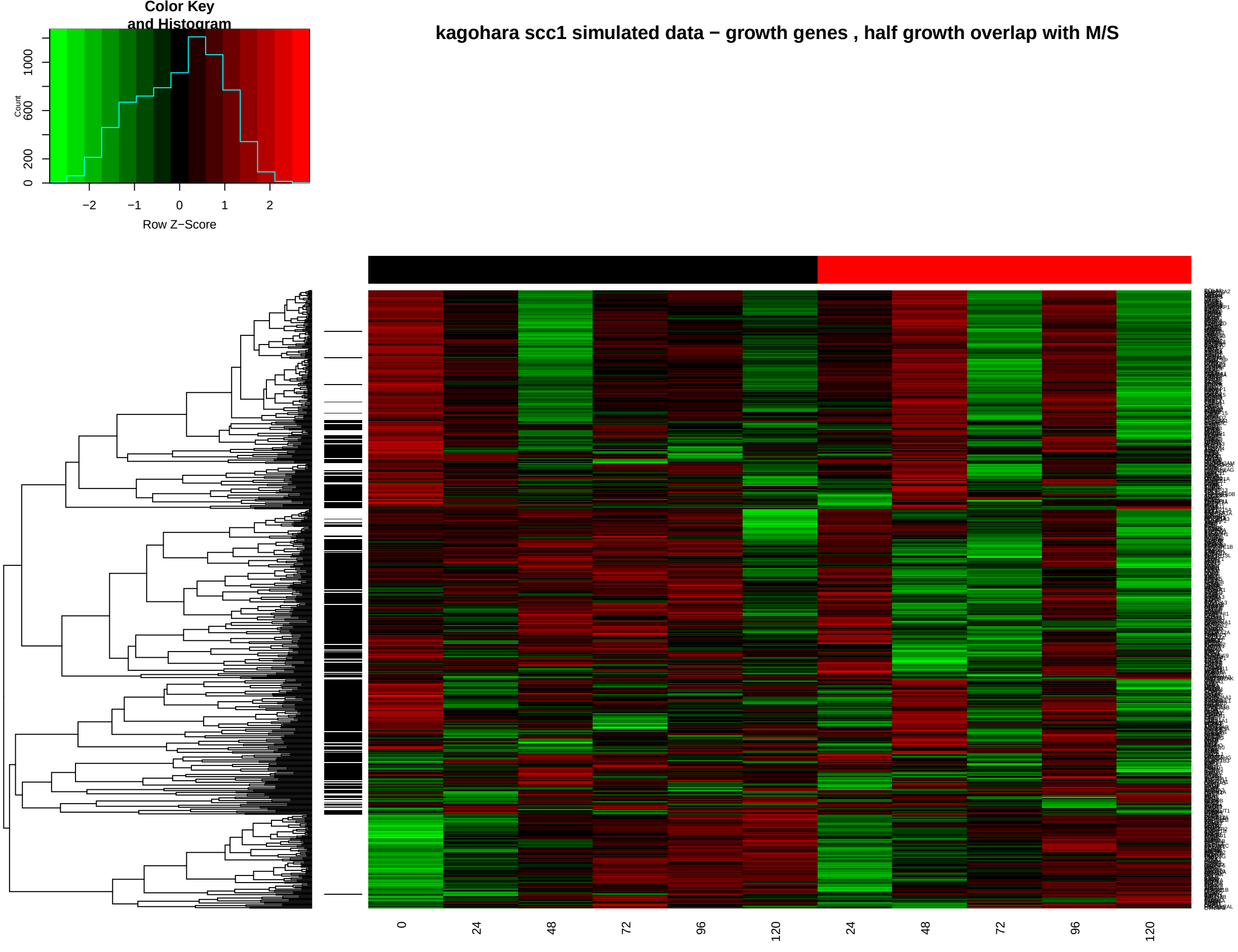
Color Key
and Histogram
Count
0
200
600
1000
−2
−1
0
1
2
Row Z−Score
kagohara scc1 simulated data − growth genes , half growth overlap with M/S
0
24
48
72
96
120
24
48
72
96
120