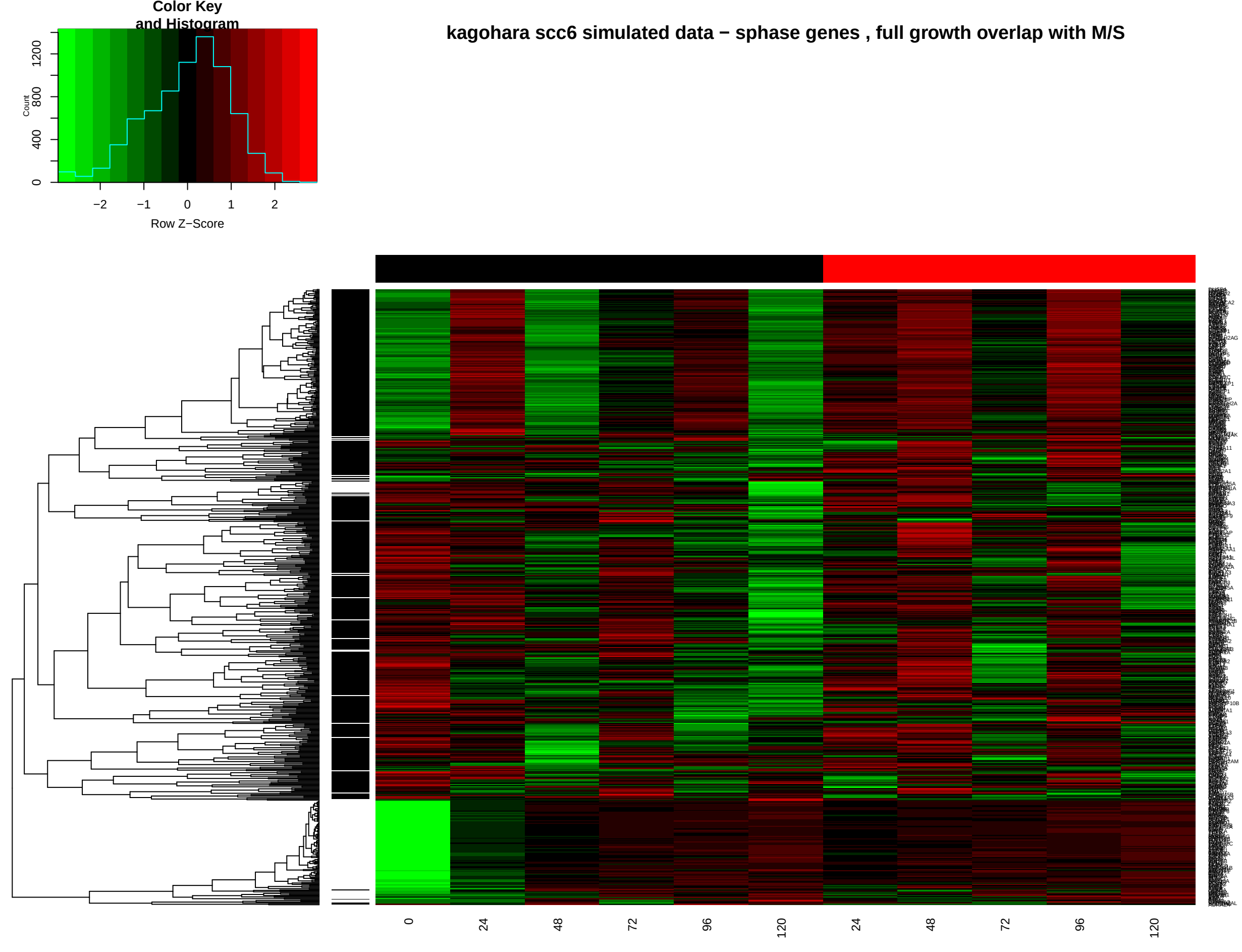
Color Key
and Histogram
Count
0
400
800
1200
−2
−1
0
1
2
Row Z−Score
kagohara scc6 simulated data − sphase genes , full growth overlap with M/S
0
24
48
72
96
120
24
48
72
96
120