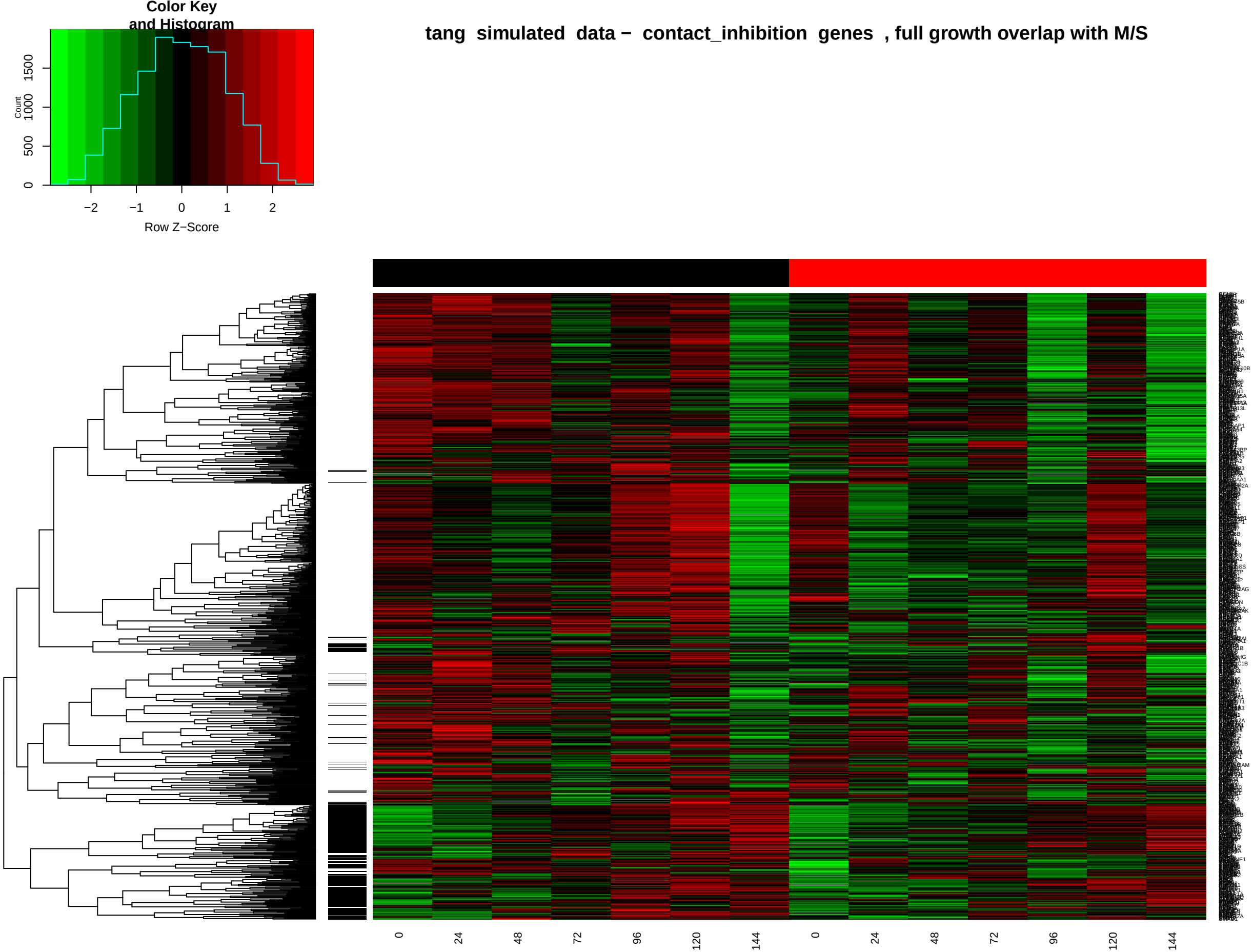
Color Key
and Histogram
Count
0
500
1000
1500
−2
−1
0
1
2
Row Z−Score
tang simulated data − contact_inhibition genes , full growth overlap with M/S
0
24
48
72
96
120
144
0
24
48
72
96
120
144