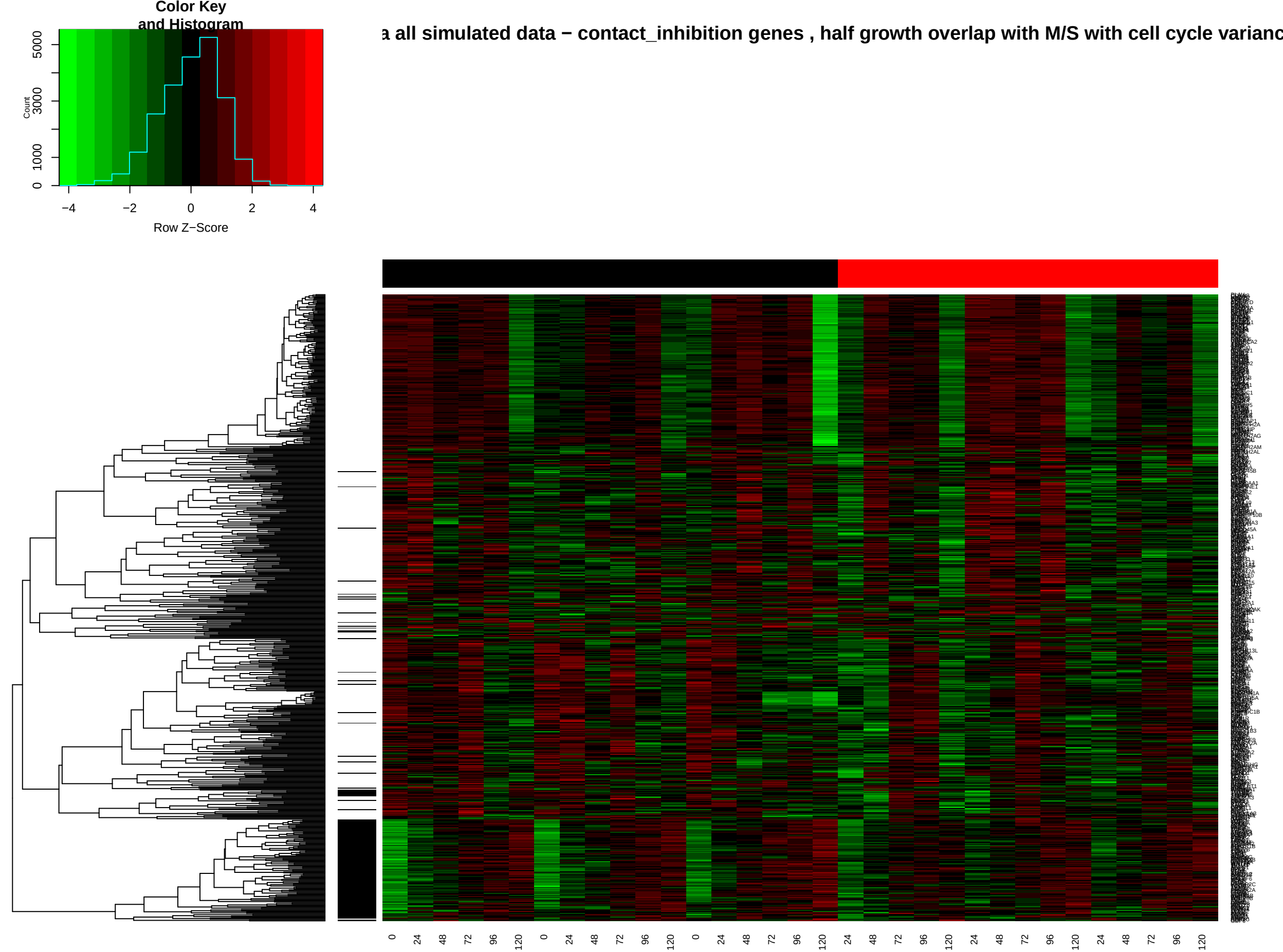
Color Key
and Histogram
Count
0
1000
3000
5000
−4
−2
0
2
4
Row Z−Score
a all simulated data − contact_inhibition genes , half growth overlap with M/S with cell cycle varianc
0
24
48
72
96
120