

relative transcript abundance

0.2  
0.1  
0.0

1

5

9

0.2  
0.1  
0.0

0.3  
0.4  
0.1  
1  
2  
1  
3  
5  
6  
1  
1  
8  
2  
5  
2  
8  
3  
5  
3  
8  
4  
7  
4  
3  
4  
5  
4  
1  
5  
3  
5  
9  
6  
5  
6  
1  
1  
3  
1  
1  
8  
9  
10  
1  
12  
13  
14  
15  
16  
1  
1  
5  
18  
5  
18  
1  
9  
5

genes