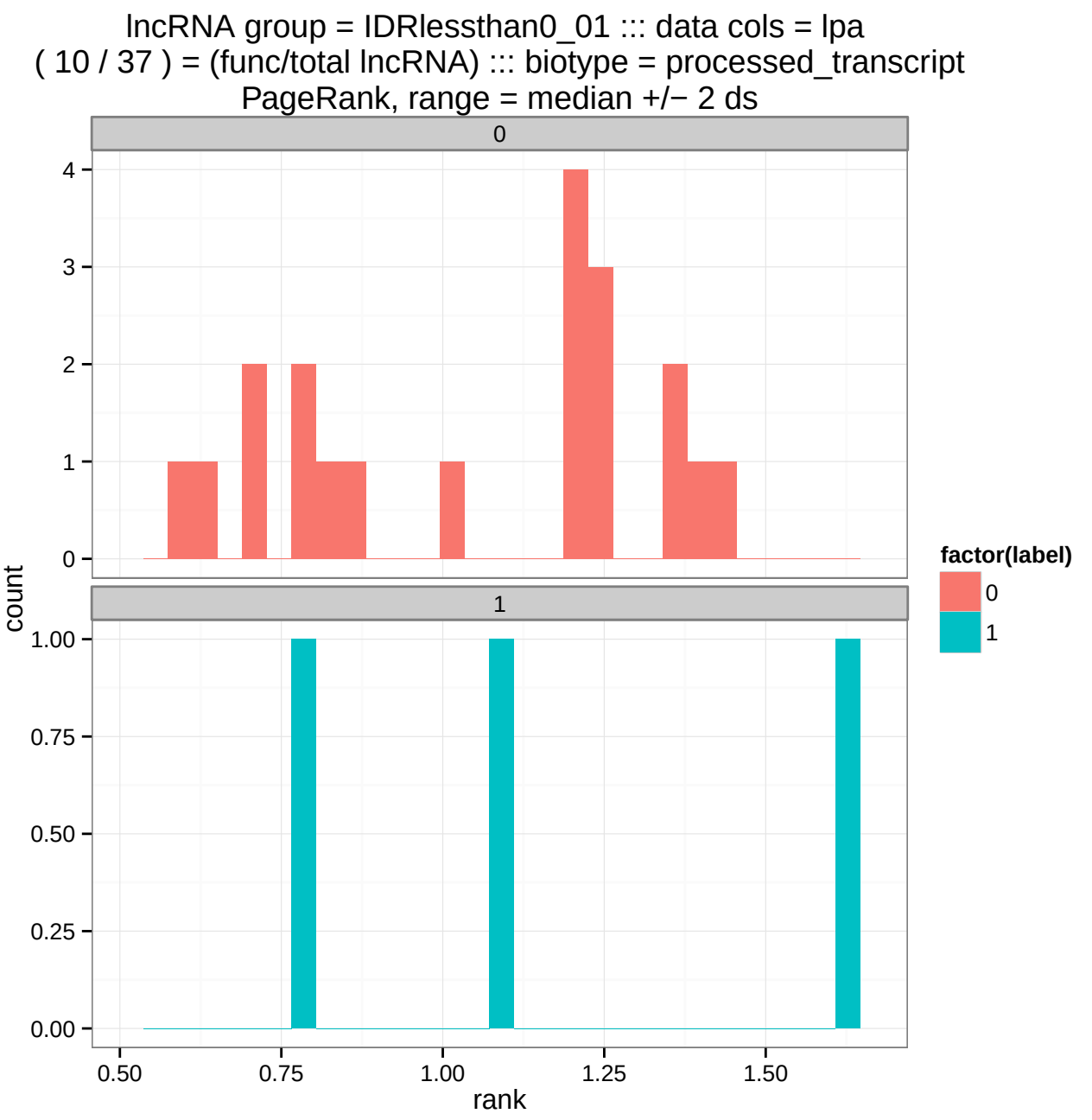
lncRNA group = IDRlessthan0_01 ::: data cols = lpa
(10 / 37) = (func/total lncRNA) ::: biotype = processed_transcript
PageRank, range = median +/− 2 ds
0
1
count
rank
0.50
0.75
1.00
1.25
1.50
0.00
0.25
0.50
0.75
1.00
0
1
2
3
4
factor(label)
0
1