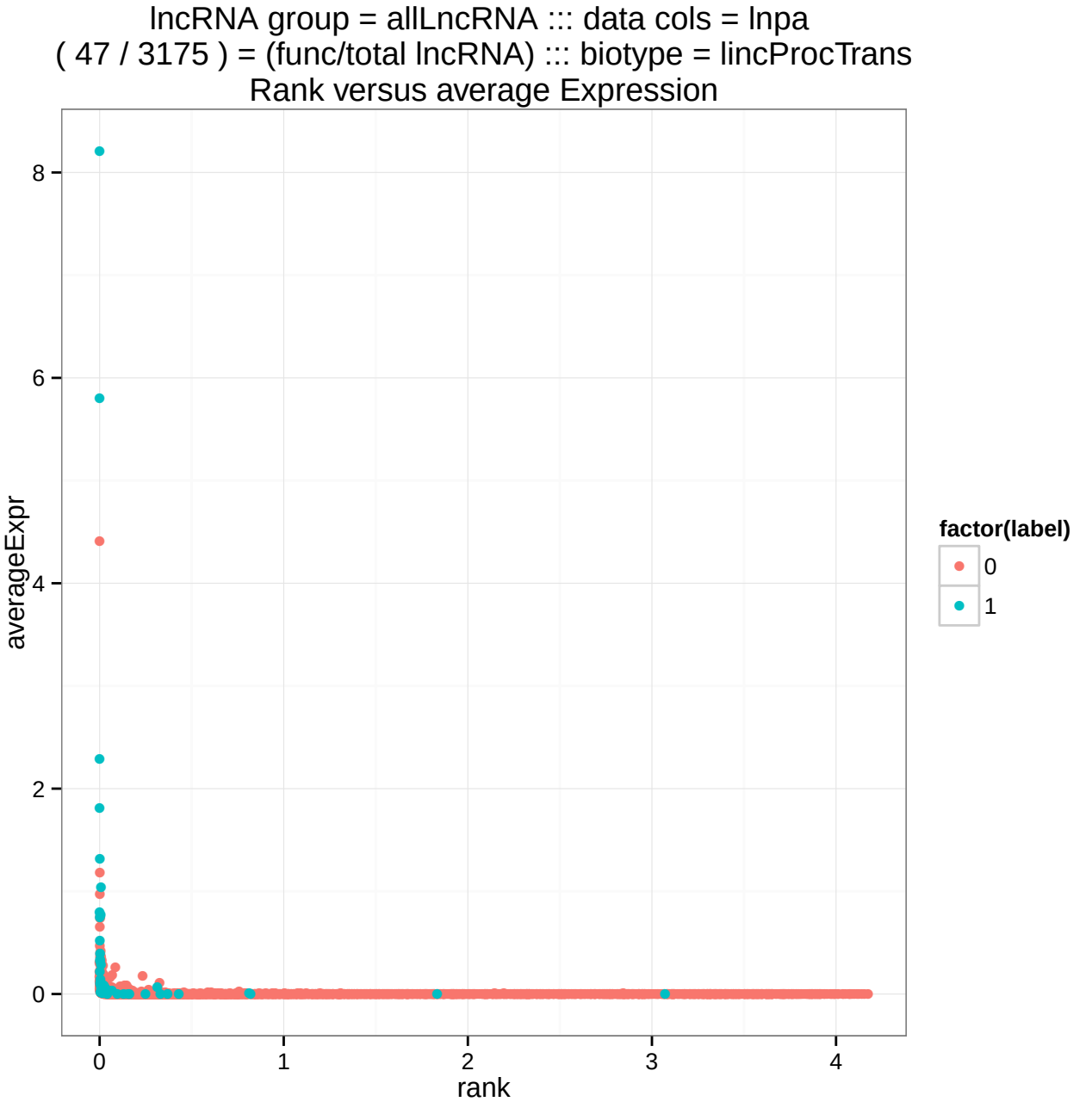
lncRNA group = allLncRNA ::: data cols = lnpa
(47 / 3175) = (func/total lncRNA) ::: biotype = lincProcTrans
Rank versus average Expression
averageExpr
rank
0
2
4
6
8
0
1
2
3
4
factor(label)
0
1