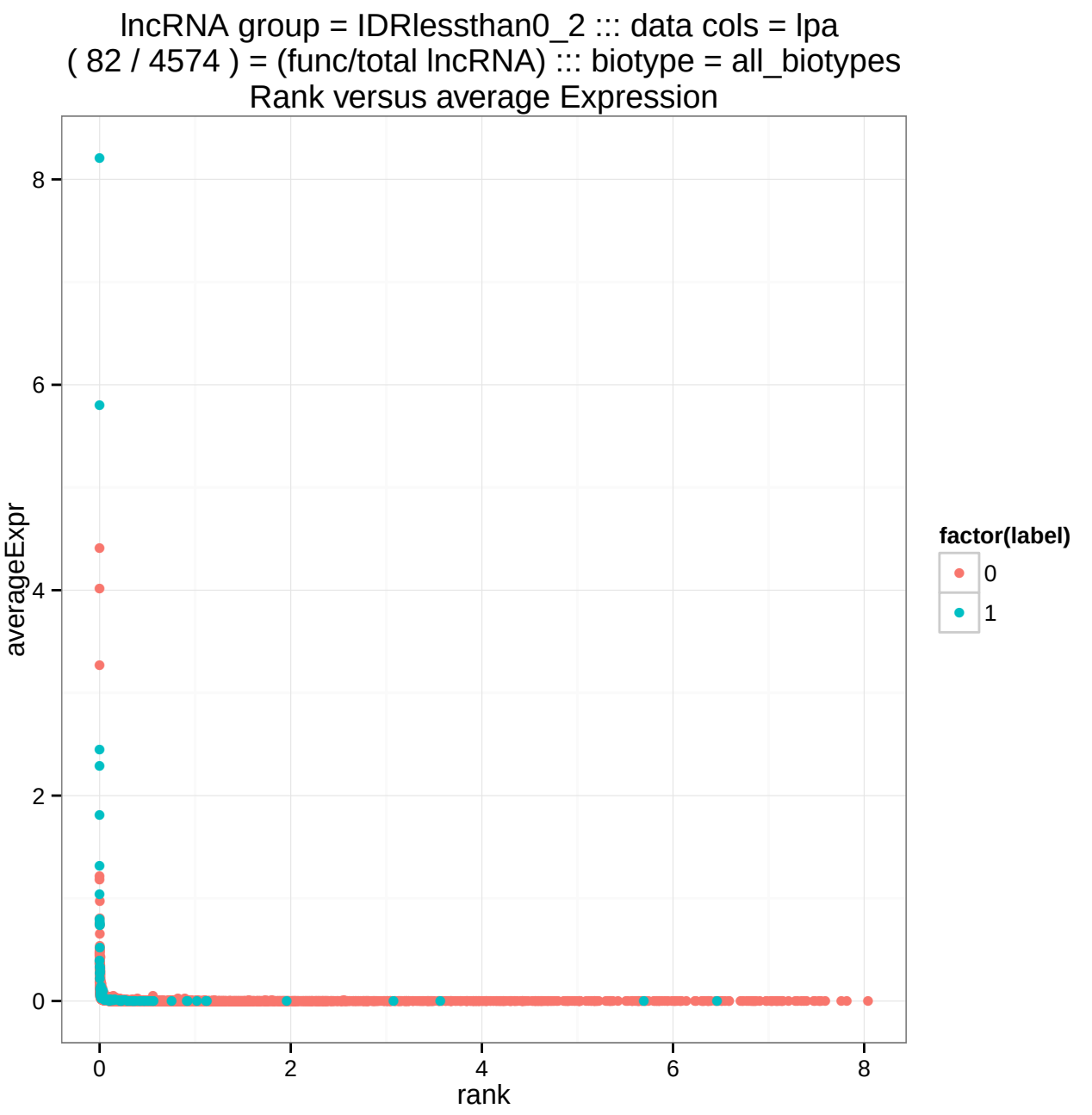
lncRNA group = IDRlessthan0_2 ::: data cols = lpa
(82 / 4574) = (func/total lncRNA) ::: biotype = all_biotypes
Rank versus average Expression
averageExpr
rank
0
2
4
6
8
factor(label)
0
1