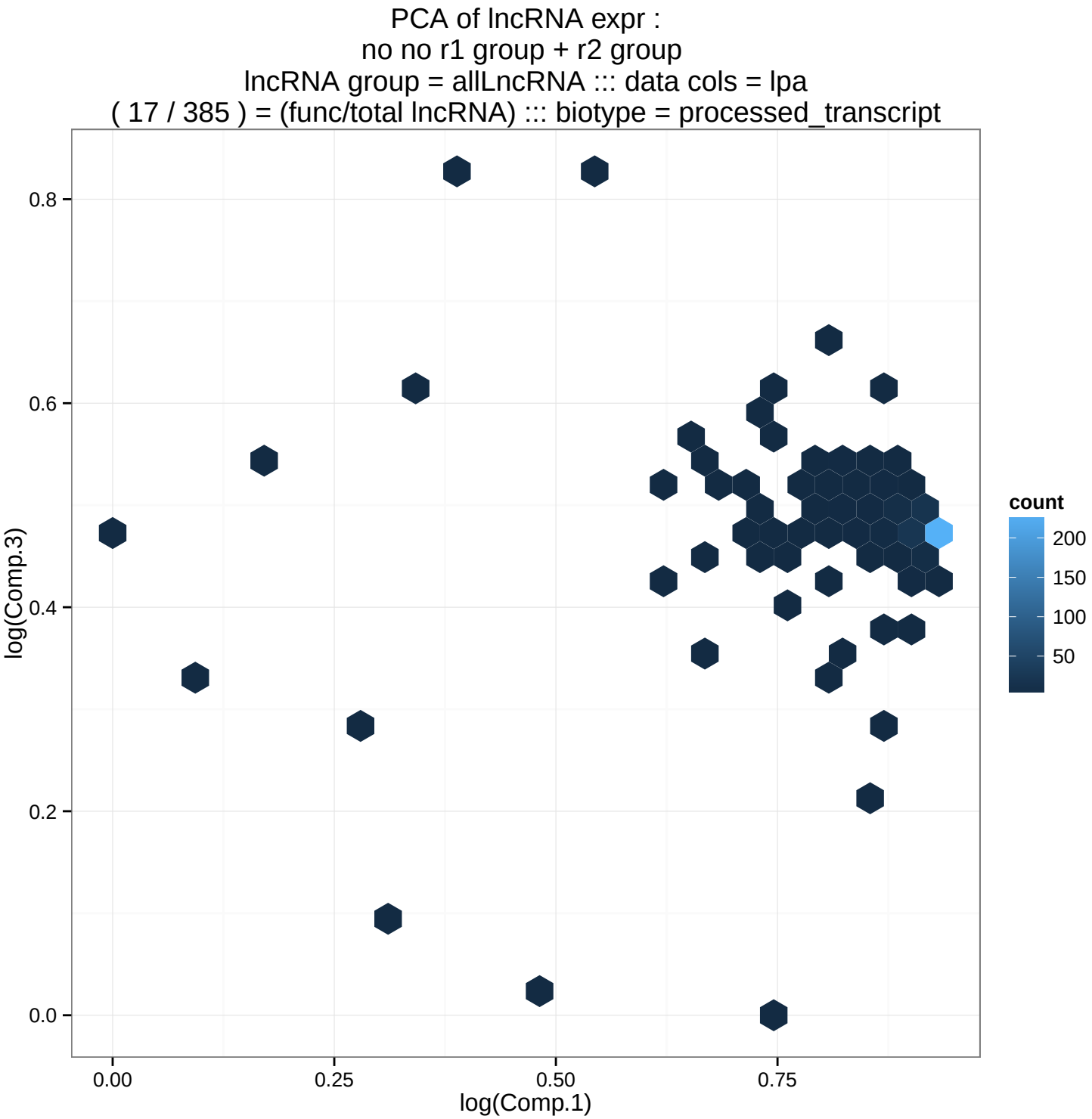
PCA of lncRNA expr :
no no r1 group + r2 group
lncRNA group = allLncRNA ::: data cols = lpa
(17 / 385) = (func/total lncRNA) ::: biotype = processed_transcript
log(Comp.3)
0.0
0.2
0.4
0.6
0.8
log(Comp.1)
0.00
0.25
0.50
0.75
count
200
150
100
50