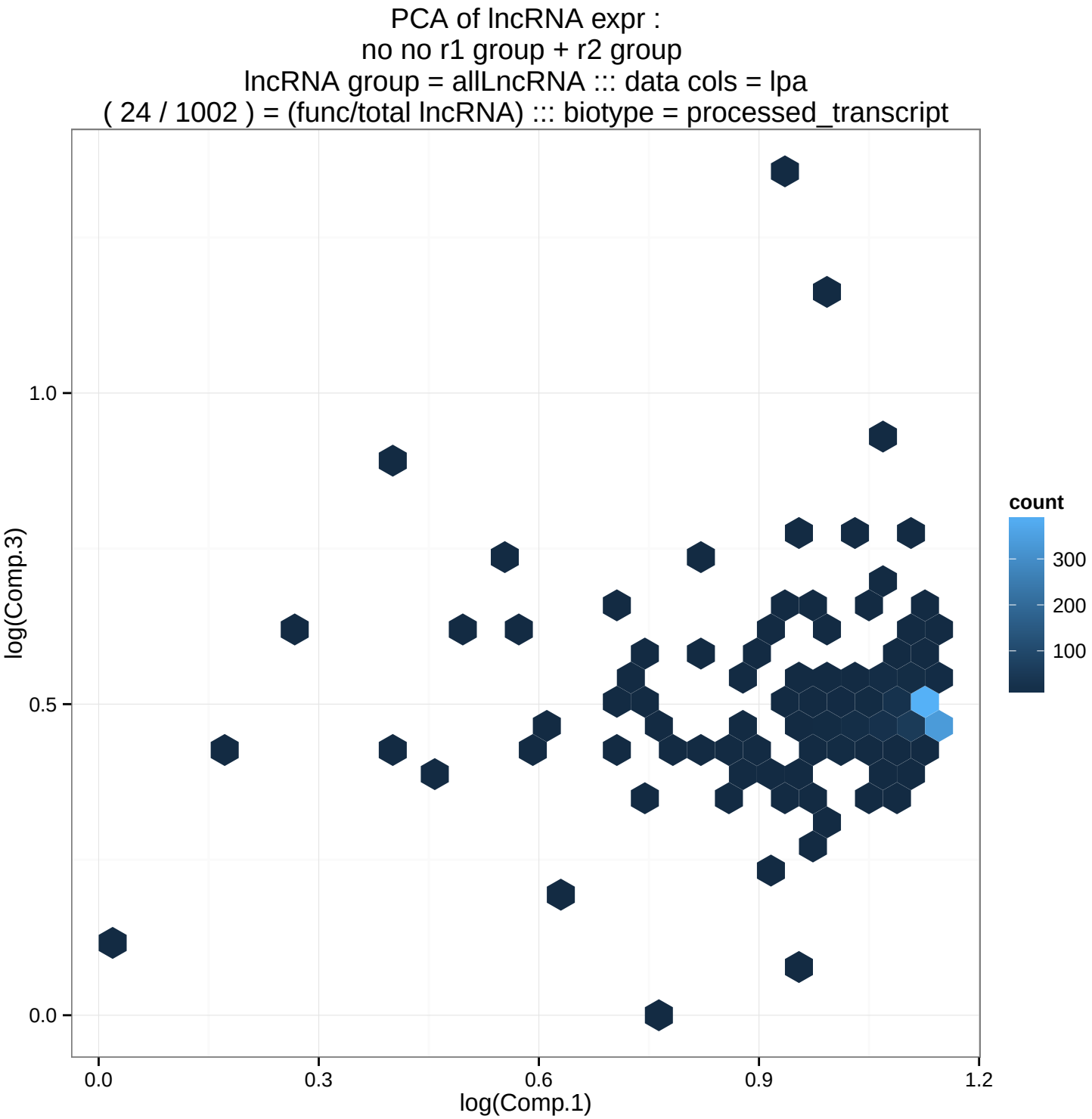
PCA of lncRNA expr :
no no r1 group + r2 group
lncRNA group = allLncRNA ::: data cols = lpa
( 24 / 1002 ) = (func/total lncRNA) ::: biotype = processed_transcript
log(Comp.3)
1.0
0.5
0.0
0.0
0.3
0.6
0.9
1.2
log(Comp.1)
count
300
200
100