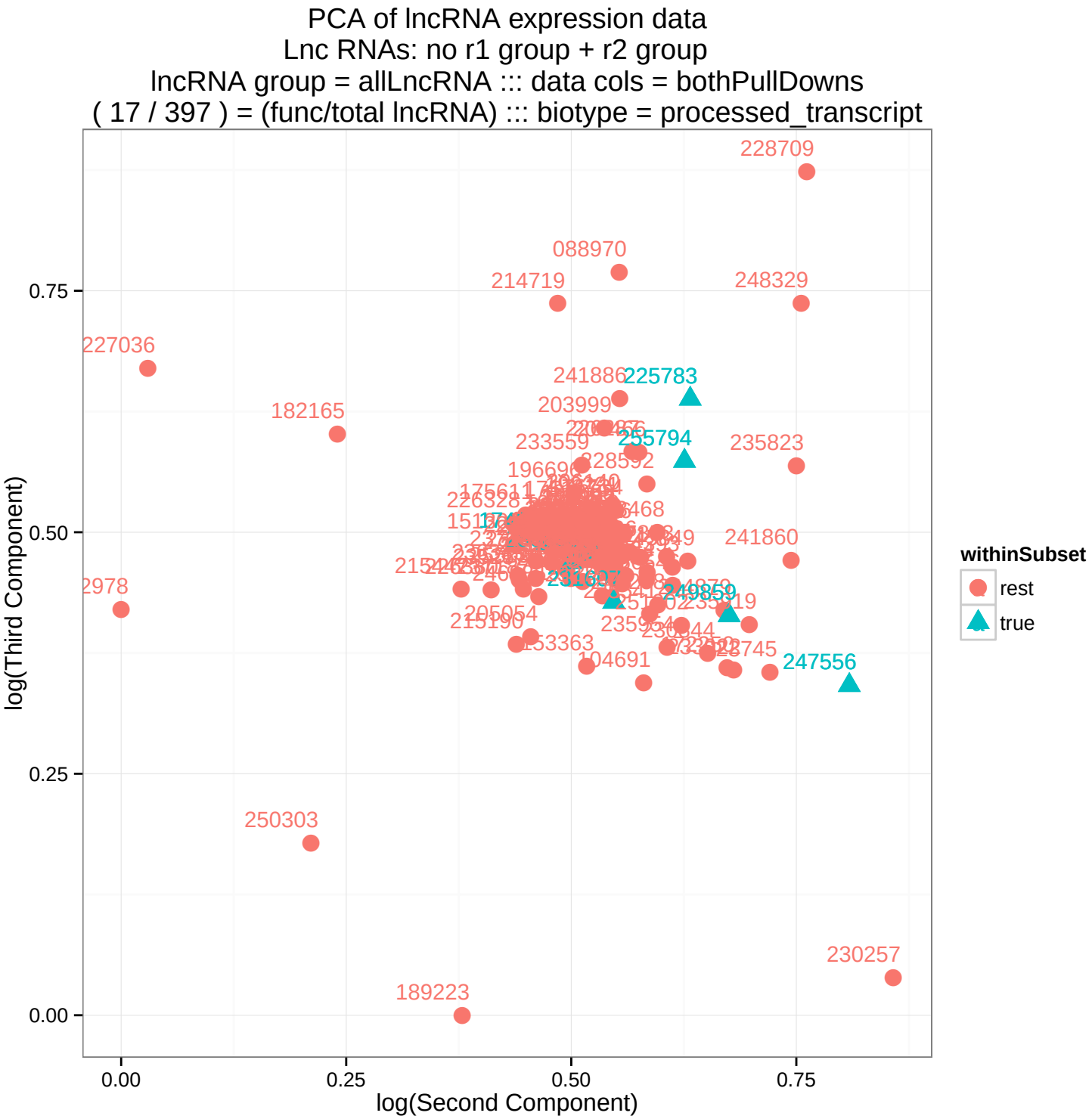
PCA of lncRNA expression data
Lnc RNAs: no r1 group + r2 group
lncRNA group = allLncRNA ::: data cols = bothPullDowns
(17 / 397) = (func/total lncRNA) ::: biotype = processed_transcript
log(Third Component)
log(Second Component)
0.00
0.25
0.50
0.75
228709
088970
214719
248329
227036
241886
225783
203999
182165
233559
255794
235823
228592
241860
2978
249859
205054
215190
153363
104691
247556
250303
230257
189223
withinSubset
rest
true