IncRNA group = IDRlessthan0_1 ::: data cols = Ipa (26 / 1131) = (func/total IncRNA) ::: biotype = lincRNA star plot of total expression over all samples SK.N.SH_RA.longPolyA A549.longPolyA NHLF.longPolyA AG04450.longPolyA 10 -NHEK.longPolyA BJ.longPolyA MCF.7.longPolyA GM12878.longPolyA factor(withinSubset) false — true K562.longPolyA H1.HESC.longPolyA HUVEC.longPolyA HELA.S3.longPolyA HSMM.longPolyA HEPG2.longPolyA HMEC.longPolyA

variable