Distribution of summation groups in transcript-type/pulldown RPKM distros color by summation group longNonPolyA, IncRNA longNonPolyA, mRNA 1.00 -0.75 -0.50 factor(sumGroups) (1e-05,0.000364] 0.25 -(0.000364, 0.00075](0.00075, 0.00136]count 0.000 (0.00136, 0.00235] (0.00235,0.0042] longPolyA, IncRNA longPolyA, mRNA (0.0042, 0.0075]1.00 -(0.0075,0.0141] (0.0141, 0.0298]0.75 -(0.0298, 0.0847] (0.0847,258] 0.50 -0.25 -0.00 -5 -5 -15 -5 -15 -10 0 -10 0 log(RPKM of transcript)