Distribution of summation groups in transcript-type/pulldown RPKM distros color by summation group longNonPolyA, IncRNA longNonPolyA, mRNA 20000 factor(sumGroups) 10000 -(1e-05,0.000364] (0.000364, 0.00075](0.00075,0.00136] (0.00136, 0.00235]0 count (0.00235,0.0042] longPolyA, IncRNA longPolyA, mRNA (0.0042, 0.0075] (0.0075,0.0141] (0.0141, 0.0298]20000 -(0.0298, 0.0847] (0.0847,258] 10000 -0 --5 -10 -5 -15 -10 0 -15 0 5 log(RPKM of transcript)