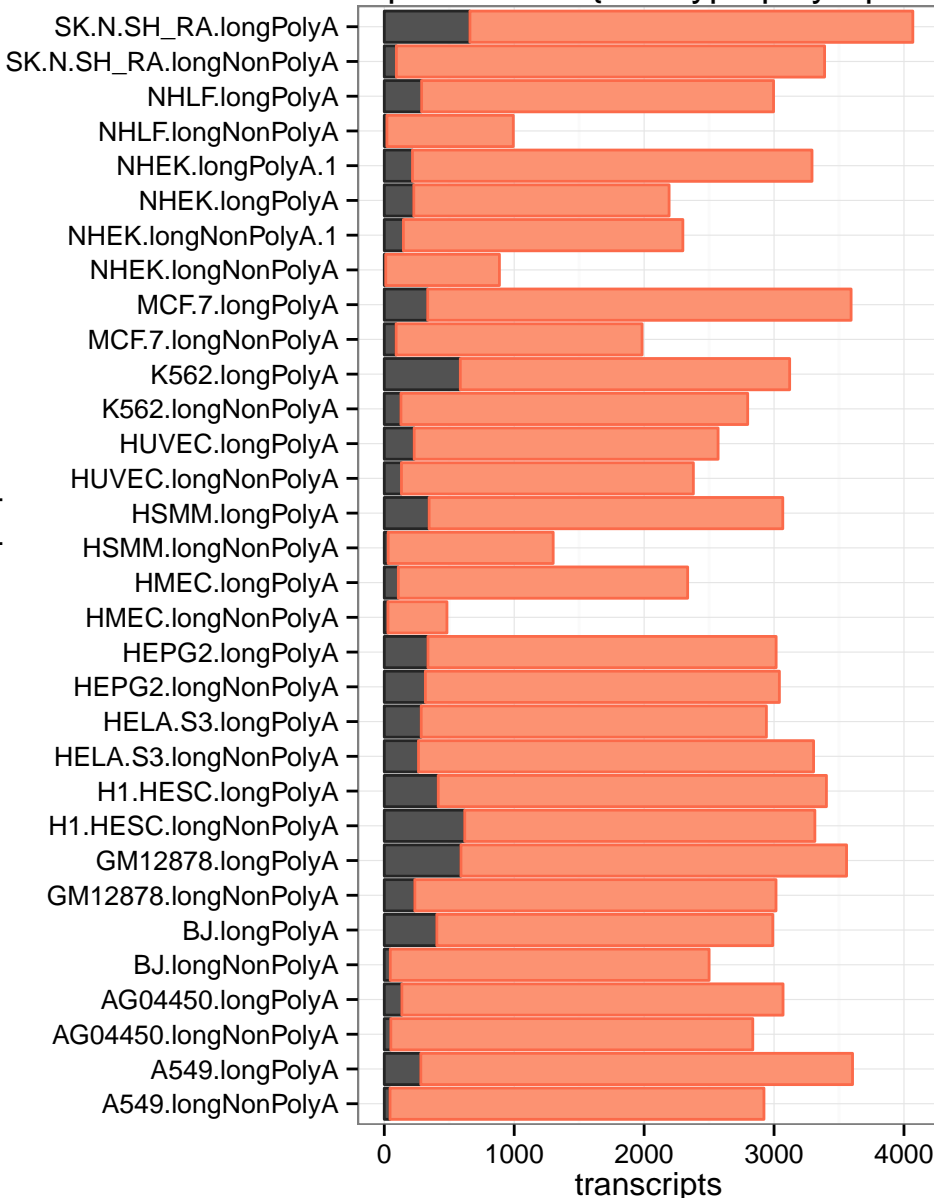


Cell-type Max Expr

For transcripts found in {cell-type/polyA pulldown} pair

RnaSeqExpr



ctsCat



cell-type-this
cell-type-other