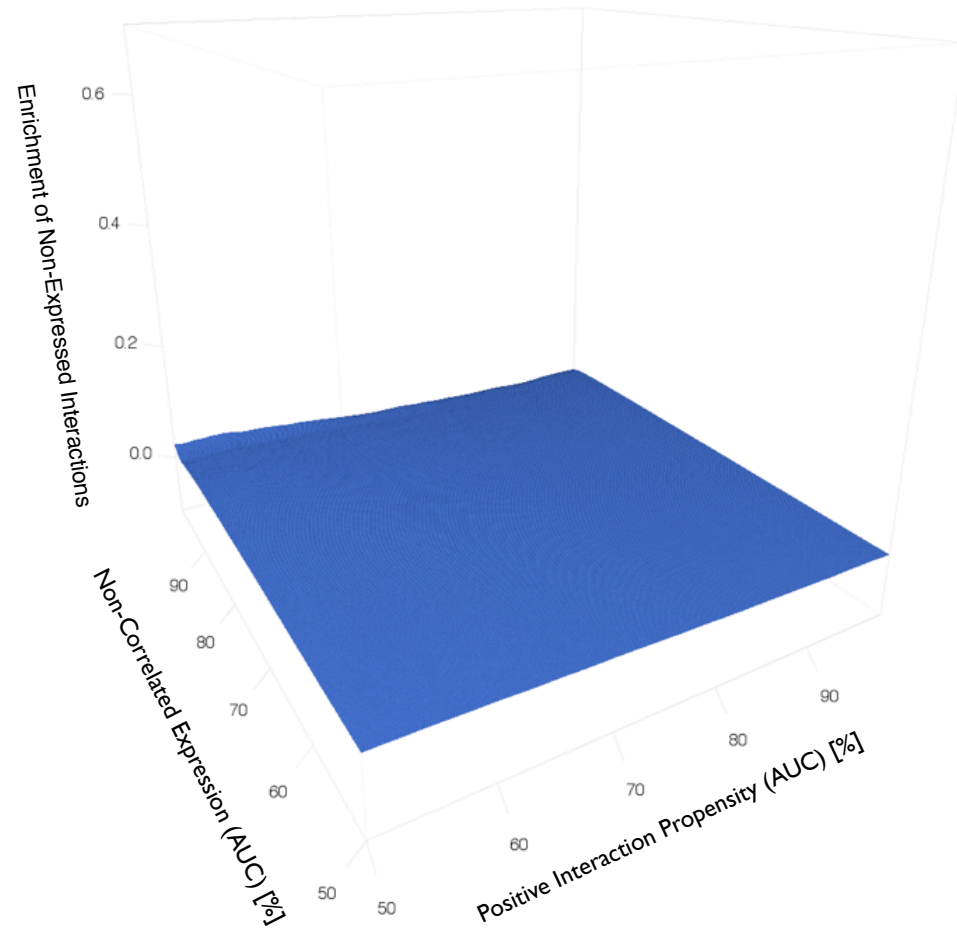
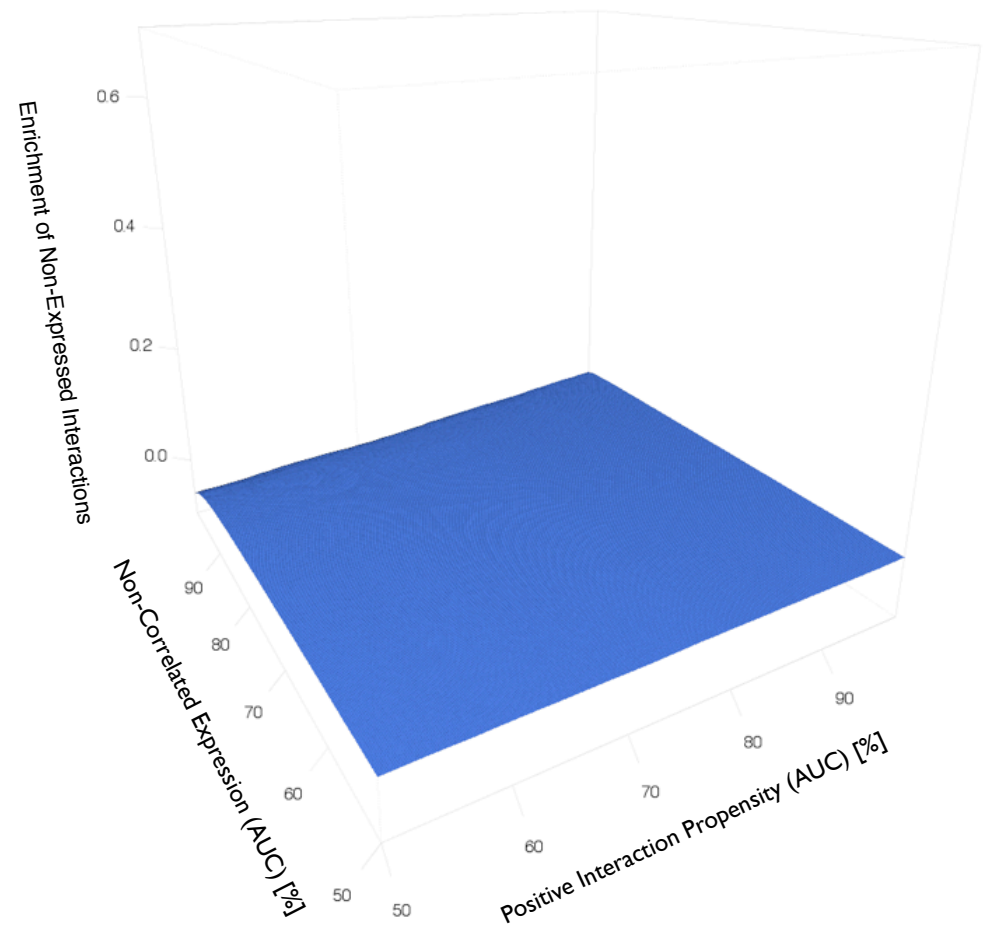
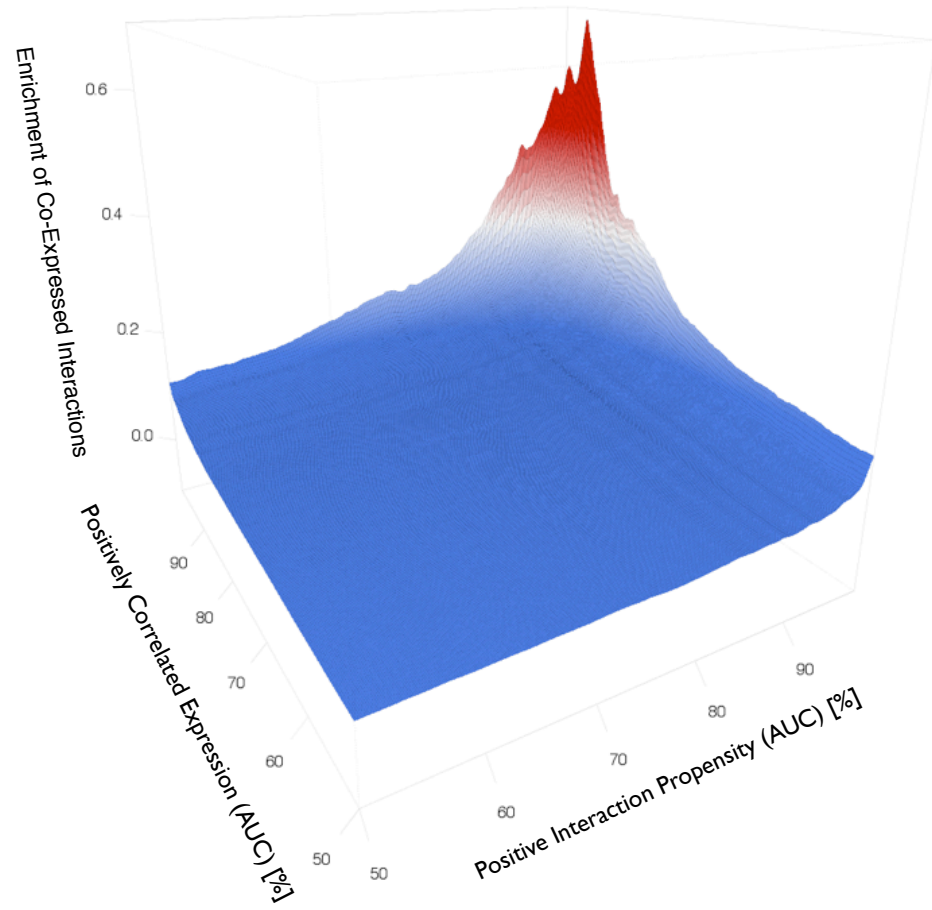
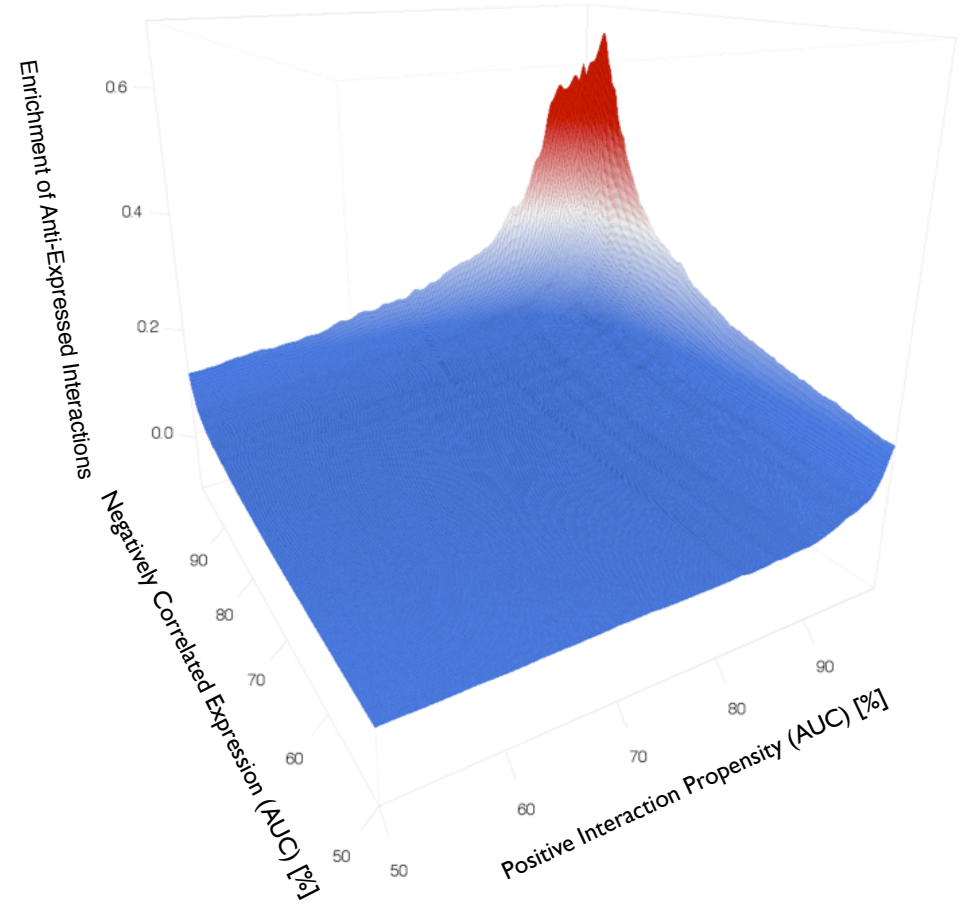
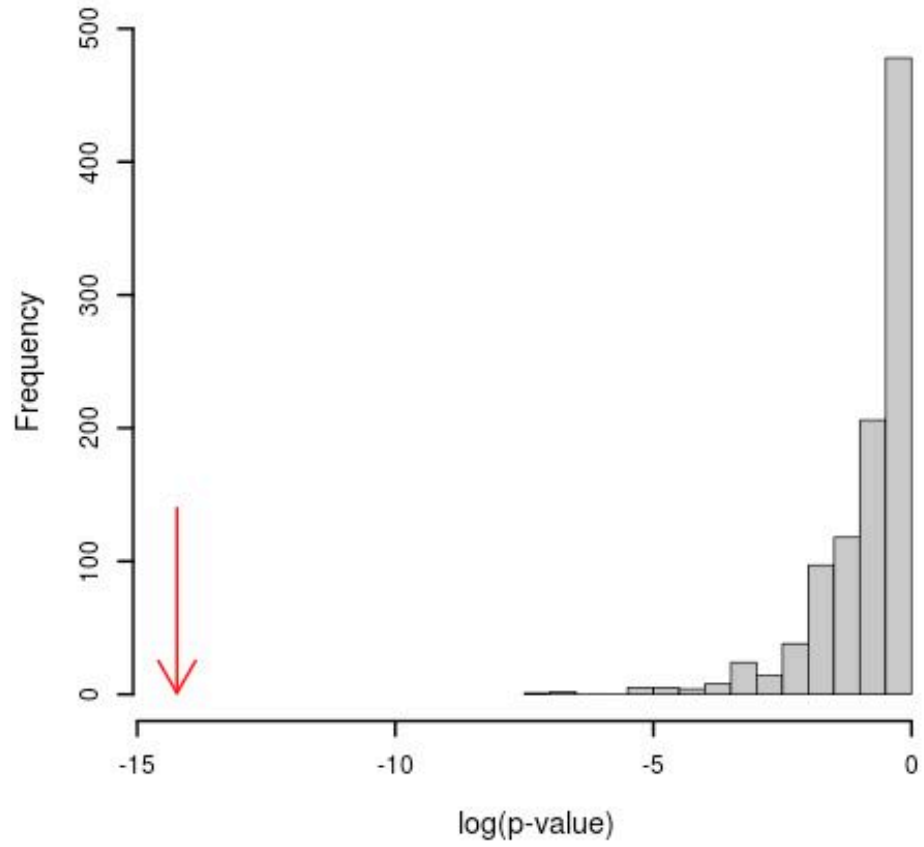


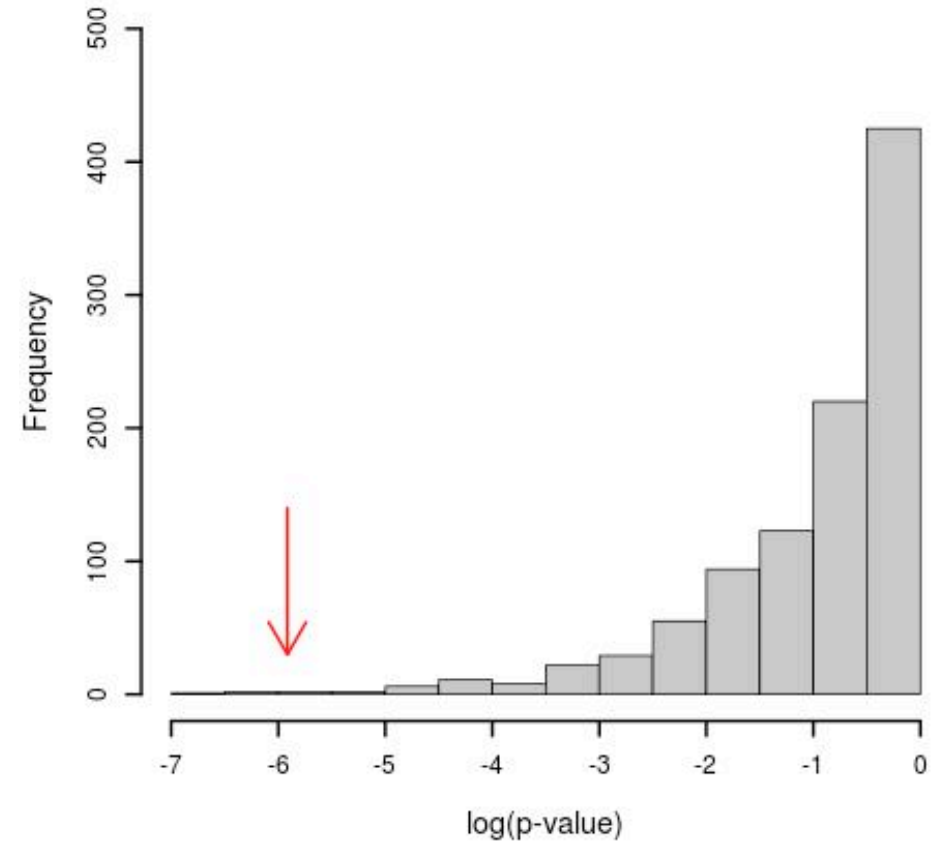
**A****Transcript-Transcript correlations****B****Protein-Transcript correlations****Figure S1**

**A****B****Figure S2**

**HUR: p-value distribution**



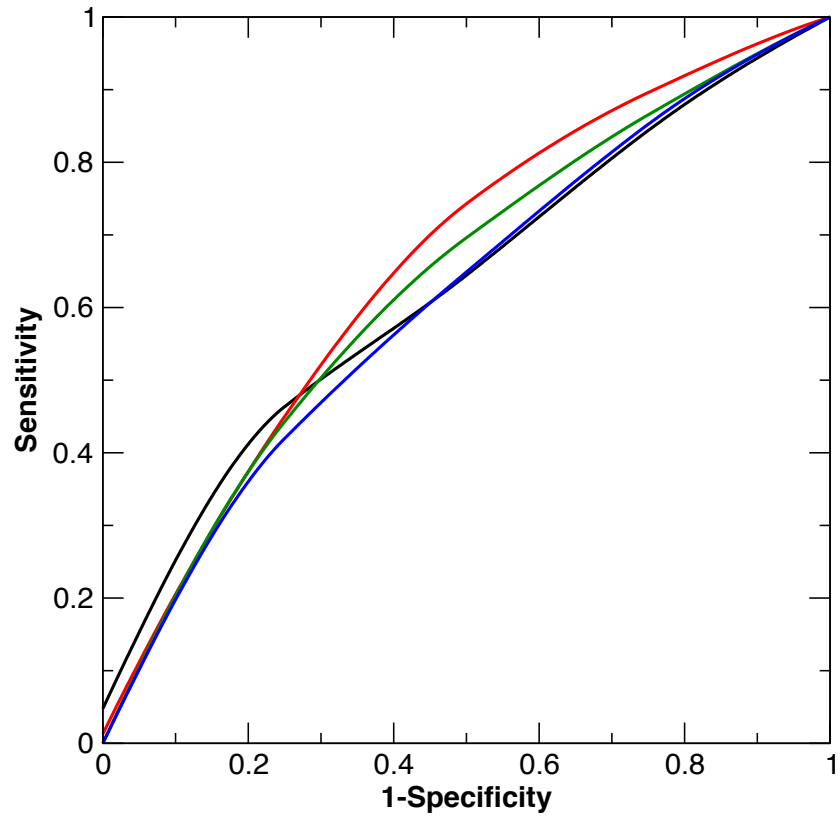
**LIN28B: p-value distribution**



**Figure S3**

A

ROC Comparison (HUR)



ROC Comparison (LIN28B)

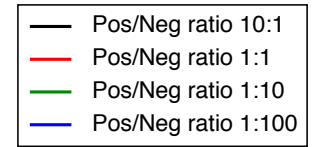
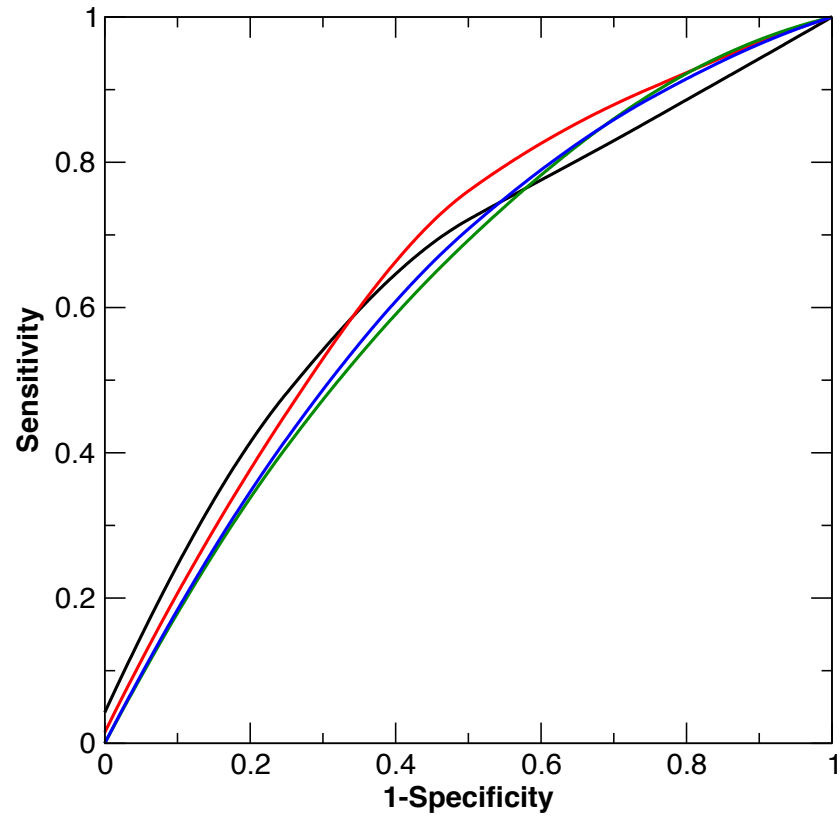
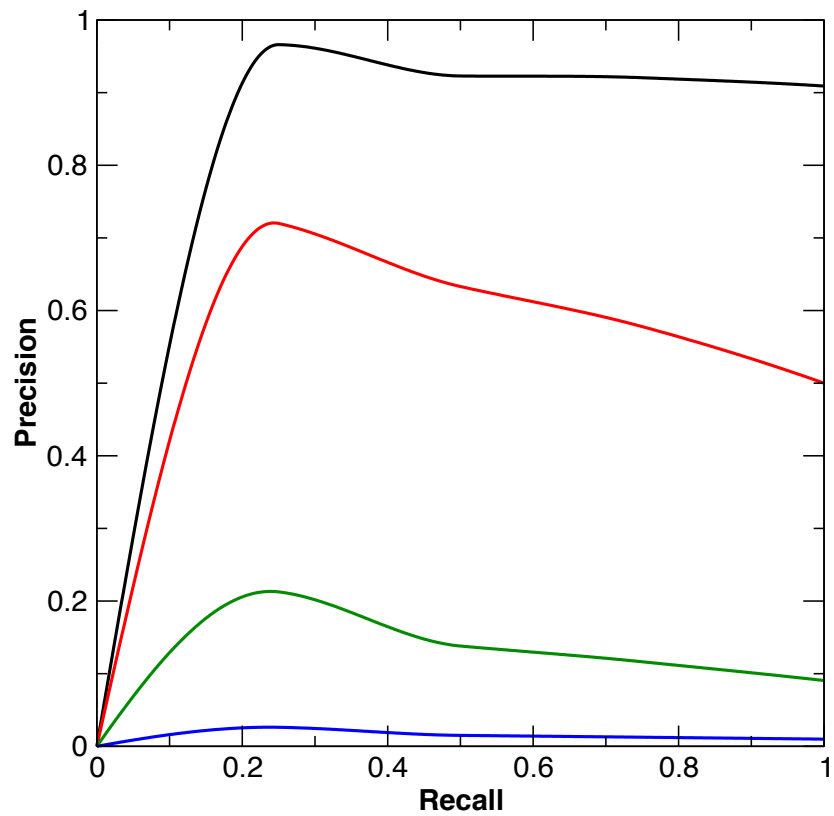


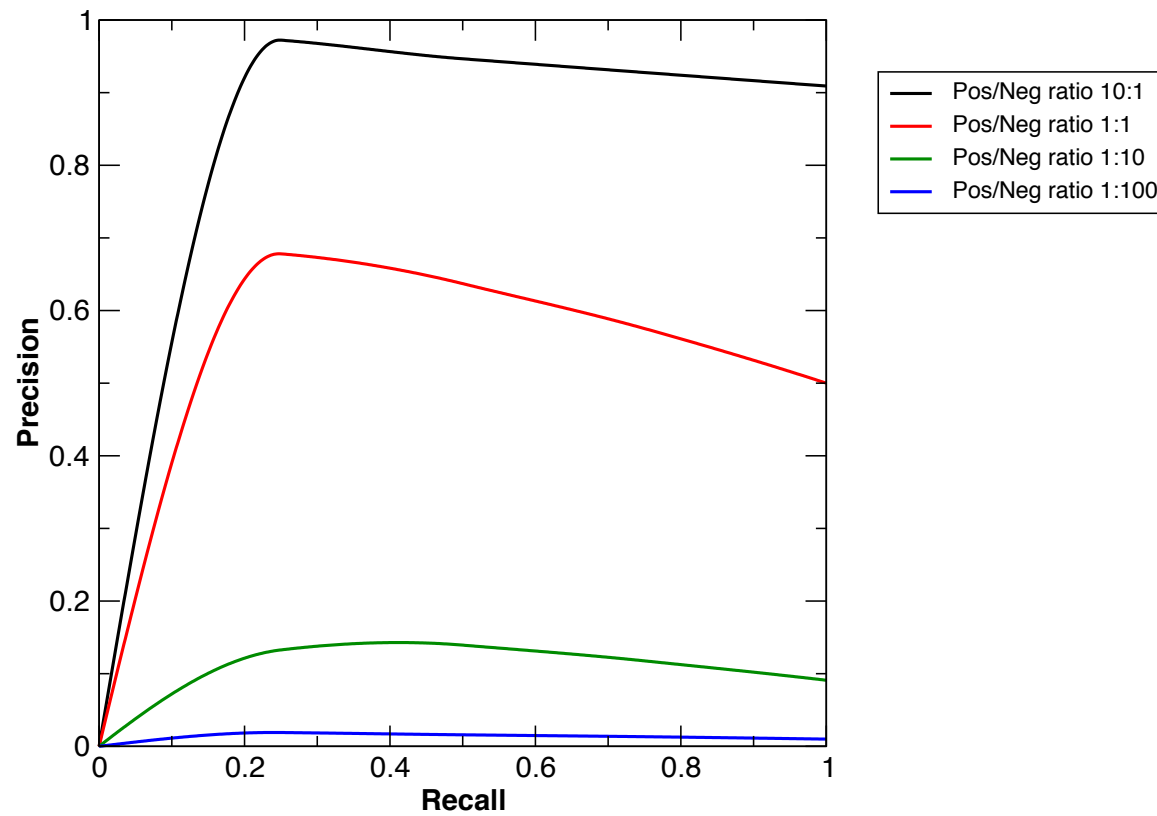
Figure S4

**B**

**Precision-Recall Comparison (HUR)**



**Precision-Recall Comparison (LIN28B)**



— Pos/Neg ratio 10:1  
— Pos/Neg ratio 1:1  
— Pos/Neg ratio 1:10  
— Pos/Neg ratio 1:100

**Figure S4**

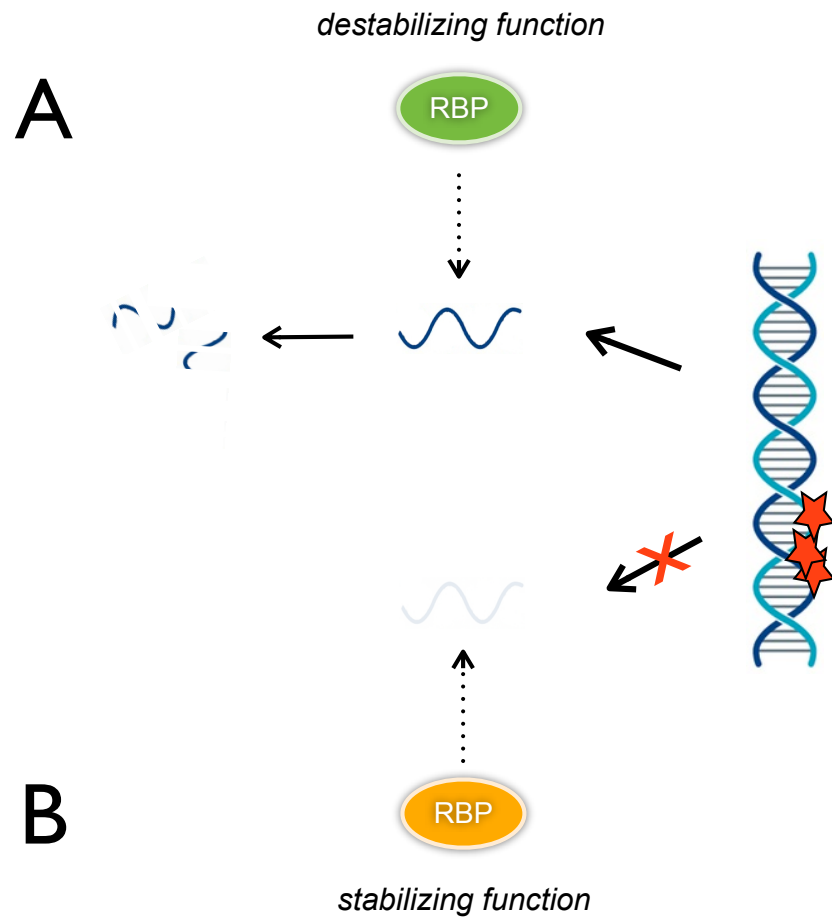


Figure S5

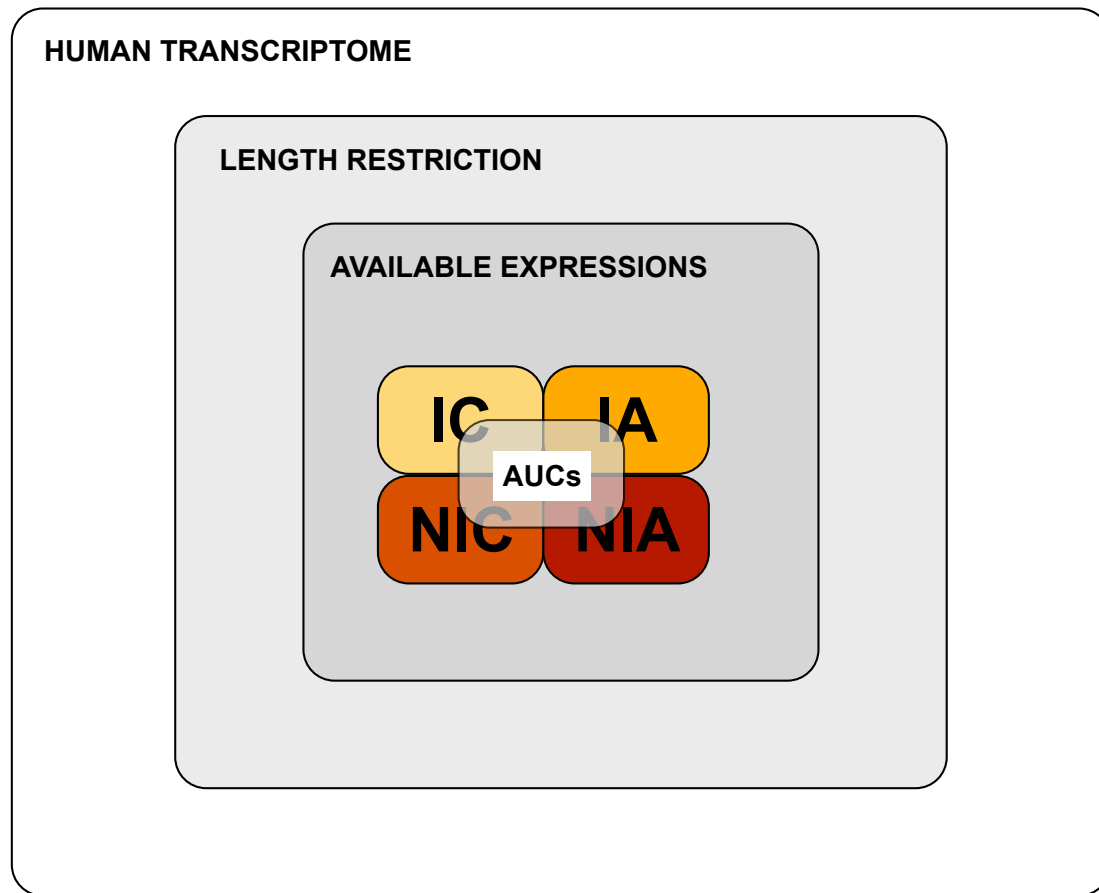
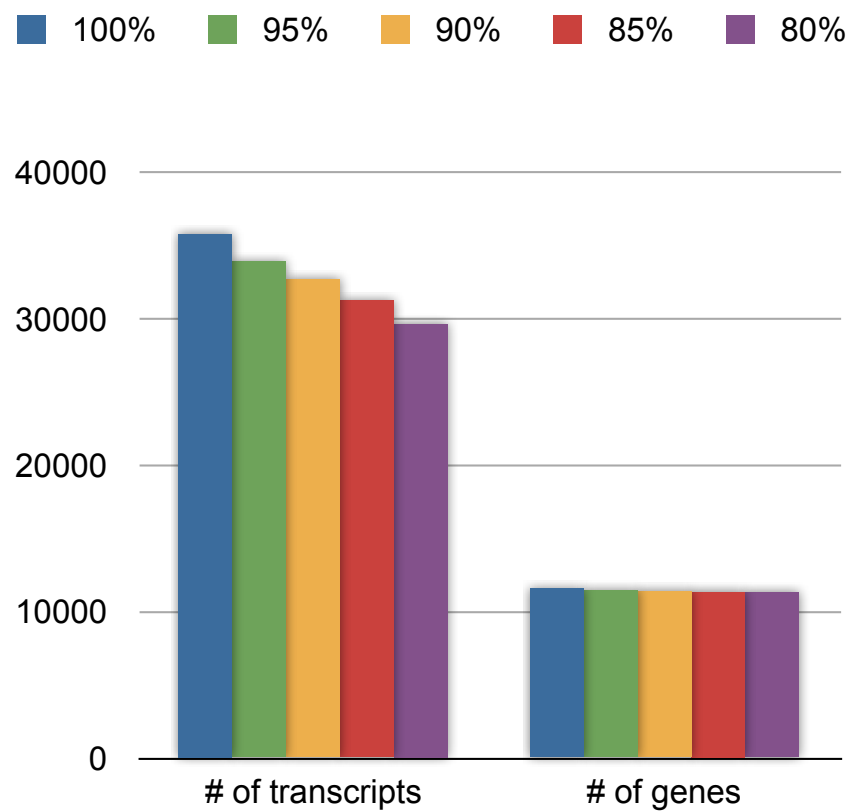


Figure S6

### Sequence redundancy reduction in mRNAs database



**Figure S7**