

Expected TS that interact with HK (threshold: 50 %)

STRING	94 %	75.8 %	88 %	92.6 %	91.8 %
IMEx	90 %	55.7 %	70.7 %	83.8 %	87.6 %
HI-2012	77.5 %	31.6 %	42.7 %	73.9 %	76.2 %
Havugimana	90.7 %	70.1 %	71.7 %	86.9 %	93.7 %
Bossi	89.6 %	58 %	72.9 %	85.4 %	87.7 %
	Body Map	Gene Atlas	HPA	HPA All	RNAseq Atlas

Percent

100
75
50
25
0