Expected TS that interact with HK (threshold: 20 %)

STRING	92.8 %	66.9 %	84 %	85.2 %	89.6 %
IMEx	87.9 %	43.8 %	63.2 %	71.6 %	83.6 %
HI-2012	74.5 %	21.9 %	34.1 %	58.5 %	69.8 %
Havugimana	90.1 %	58.7 %	65.2 %	76.9 %	91.4 %
Bossi	87.6 %	46.8 %	65.4 %	74.2 %	84.2 %
	Body Map	Gene Atlas	HPA	HPA AII	RNAseq Atlas