Expected TS that interact with HK (threshold: 12.5 %)

	Body Map	Gene Atlas	HPA	HPA All	RNAseq Atlas
STRING	92.2 %	63.5 %	81.8 %	79.8 %	88.3 %
IMEx	87.1 %	39.9 %	59.1 %	63.7 %	81.2 %
HI–2012	73.3 %	19.4 %	31 %	49.5 %	66.6 %
Havugimana	89.8 %	54.5 %	61.3 %	69.8 %	90.1 %
Bossi	86.8 %	43.1 %	61.5 %	66.6 %	82.2 %

Percent: 100, 75, 50, 25, 0