

## Expected TS that interact with HK (threshold: 15 %)

	Body Map	Gene Atlas	HPA	HPA All	RNAseq Atlas
STRING	92.8 %	64.8 %	82.3 %	82.4 %	88.3 %
IMEx	87.9 %	41.4 %	60 %	67.3 %	81.2 %
HI-2012	74.5 %	20.4 %	31.6 %	53.7 %	66.6 %
Havugimana	90.1 %	55.9 %	62 %	73.4 %	90.1 %
Bossi	87.6 %	44.5 %	62.2 %	70.1 %	82.2 %

**Percent**

