

## Betweenness of TS genes in all subnetworks: z-scores

STRING	[-0.5, -0.1] mean = -0.3	[-1.4, 0.3] mean = -0.7	[-0.6, 0.8] mean = -0.2	[-0.6, 1.2] mean = -0.1	[-1.7, -0.4] mean = -0.7
IMEx	[-0.8, 0] mean = -0.3	[-1.9, 3.6] mean = -0.7	[-1.3, 0.6] mean = -0.5	[-1, 0.1] mean = -0.5	[-1.9, -0.5] mean = -0.9
HI-2012	[-0.2, 0] mean = -0.2	[-1.5, 5] mean = -0.3	[-0.8, 1.2] mean = -0.3	[-1, 0.9] mean = -0.4	[-1, 0.8] mean = -0.3
Havugimana	[-0.6, -0.6] mean = -0.6	[-3.9, -0.1] mean = -1.3	[-1, 2.2] mean = -0.2	[-1.2, 2.2] mean = -0.4	[-1.3, 0] mean = -0.5
Bossi	[-1, -0.3] mean = -0.5	[-3.1, 0] mean = -1.8	[-1.4, 1.1] mean = -0.5	[-1.3, 0.6] mean = -0.5	[-2.8, -0.1] mean = -1.5
	Body Map	Gene Atlas	HPA	HPA All	RNAseq Atlas

