

		# TFs	# Targets	# Edges
H1ESC differentiation	hESC ChIP	57	17147	279501
	hESC KD	96	10751	103446
	hESC ChIP&KD	25	4891	9301
Reprogramming sequencing data	mESC ChIP	18	14572	60479
	mESC KD	52	11601	76616
	mESC ChIP&KD	17	4711	8020
Hepatocyte dedifferentiation	HepG2 ChIP	40	12238	162659
	mESC ChIP	28	13820	79201
	mESC KD	93	11417	108144
	mESC ChIP&KD	28	4502	8134

Supplemental Table S5. Number of regulators, targets, and edges in different gold standard networks. For each gold standard network, we only count the regulators and targets that are present in the input (gene expression and regulatory features) of the dataset.