







Zhang et al Fig. 9

Fig.9 Network Statistics: 278 nodes,783 edges

Top 25 transcription factors from differential network

	Name	Total Edges	Edges In	Edges Out
1	SP1/SP3	63	14	49
2	AP-1	52	27	25
3	p53	41	20	21
4	c-Myc	35	23	12
5	STAT3	24	12	12
6	NF-kB	20	4	16
7	c-Jun	20	10	10
8	STAT1	19	11	8
9	YY1	19	3	16
10	LXR-alpha/RXR-alpha	17	5	12
11	EGR1	16	9	7
12	PU.1	16	9	7
13	SMAD3	16	10	6
14	Oct-1	14	6	8
15	RUNX2	14	7	7
16	STAT5	14	5	9
17	c-Fos	14	8	6
18	SP1	14	4	10
19	DEC1	13	7	6
20	ETS1	13	7	6
21	GATA-1	13	4	9
22	RelA	13	7	6
23	AML1/ETO	12	1	11
24	c-Myb	12	8	4
25	C/EBPbeta	12	6	6

