

**Table S12**

Xu\_Table 4. Reduction of E2F6 does not affect the transcriptome of Ntera2 cells

	RefSeq ID	Gene Symbol	E2F6 A*	E2F6 B*	Experiment 1#	Experiment 2#
<b>upregulated</b>	<b>NM_016124.3</b>	<b>RHD</b>	<b>1.4</b>	<b>0.82</b>	<b>1.68</b>	<b>2.68</b>
	NM_000088.2	COL1A1	1.31	1.49	1.51	0.14
<b>downregulated</b>	<b>NM_003714.2</b>	<b>STC2</b>	<b>2.41</b>	<b>2.28</b>	<b>1.54</b>	<b>4.13</b>
	<b>NM_032592.1</b>	<b>PHACS</b>	<b>1.61</b>	<b>1.18</b>	<b>1.60</b>	<b>2.49</b>
	<b>NM_016040.3</b>	<b>TMED5</b>	<b>1.06</b>	<b>0.83</b>	<b>1.63</b>	<b>2.32</b>
	<b>NM_004331.2</b>	<b>BNIP3L</b>	<b>1.13</b>	<b>1.21</b>	<b>1.55</b>	<b>1.58</b>
	<b>NM_004048.2</b>	<b>B2M</b>	<b>1.64</b>	<b>1.46</b>	<b>1.51</b>	<b>1.47</b>
	NM_001542.2	IGSF3	1.43	1.27	1.56	1.22
	NM_001038.4	SCNN1A	1.88	1.19	1.54	1.1
	NM_003161.2	RPS6KB1	1.18	0.47	1.50	1.05
	NM_014078.4	MRPL13	1.26	0.65	1.56	0.99
	NM_015608.2	C10orf137	2.03	1.35	1.55	0.87
	NM_002315.1	LMO1	1.01	0.93	1.61	0.84
	NM_022346.3	HCAP-G	2.57	1.97	1.50	0.78
	NM_025061.3	LRRC8E	1.03	1.15	1.52	0.73
	NM_020826.1	SYT13	1.04	0.94	1.53	0.58
	NM_138328.2	RHBDL3	1.27	1.46	1.50	0.57
	NM_002152.2	HRC	1.37	1.58	1.80	0.48
	NM_138813.2	ATP8B3	1.18	0.89	1.56	0.33
	NM_000702.2	ATP1A2	1.28	0.84	1.72	0.18

\*shown are the fold enrichments (log2) of E2F6 binding from two independent ChIP-chip experiments

# shown are the fold change in RNA levels from two independent shE2F6 experiments