Table S1. Comparison of chromatin immunoprecipitation data with differentially regulatedgenes in transfected WM-266-4 cells

Experiment	cDNA	Sample-specific genes	Common genes with CT16 upregulated genes	Common genes with CT16 downregulated genes
Histone acetylation	CT16	202	6	1
Histone acetylation	Control	329	0	5
DNA methylation	CT16	778	6	6
DNA methylation	Control	946	2	6