

S2 Table. Methylation fate in Jurkat T-cell leukemia of sites that are highly methylated in normal T-cells.

Non-promoter sites that are methylated (>70%) in T-cells		Hypomethylation in T-cell leukemia			
Class	#	Fraction of class (%)	Fold-ratio (versus all)	P-value (versus all)	Average cancer-normal difference
All	130,794	18.2			-10.2%
ESSEs	35,392	22.6	1.24	<1e-20	-12.3%
ESSEs unmethylated in ES	3,451	26.5	1.46	<1e-20	-12.5%
Non-ESSE sites unmethylated in ES	2,825	19.0	1.04	0.14	-8.1%
Constitutive enhancers unmethylated in ES	1,266	13.1	0.72	1.9e-3	-4.1%