

Supplementary Table 5 Overview of inter assay variability per region

	High methylated sample (n=5)			Low methylated sample (n=5)		
	Mean	SD	CV	Mean	SD	CV
<i>DMR0</i>	86.1	2.30	0.027	7.83	2.14	0.273
<i>DMR2</i>	90.2	3.13	0.035	6.26	2.71	0.433
<i>CTCF3</i>	90.7	1.82	0.020	8.06	2.88	0.357
<i>CTCF6</i>	84.6	2.11	0.025	7.90	3.15	0.399
<i>H19</i>	94.0	2.59	0.028	5.67	2.19	0.386

Mean represents mean of all CpGs in the 5 high or low methylated samples. SD represents mean of all SDs per CpG per region.

CV=coefficient of variation.