

Supplementary table

Hypermethylation of 3.3-kb repeats and hypomethylation of Sat2 repeats in cervical tumors

№	sample	3.3-kb repeats	Sat2 repeats
1	8	hyper	hypo
2	12	hyper	hypo
3	13	hyper	hypo
4	14	hyper	hypo
5	16	hyper	hypo
6	19		
7	21		hypo
8	22		hypo
9	26		hypo
10	28		hypo
11	30		
12	31		
13	33		hypo
Frequency of changes		5/13	10/13

Numbers of samples are indicated as in Figure 3.
Hyper – hypermethylation, hypo – hypomethylation,
empty box – the absence of changes.