

Different DNA methylome, transcriptome and histological features in uterine fibroids with and without MED12 mutations

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Supplemental Table S10. Increased DNA methylation and mRNA expression levels of SATB2 or NRG1 in uterine fibroids with and without MED12 mutations.

	MED12m-positive uterine fibroids (n = 9)	MED12m-negative uterine fibroids (n = 12)	Total
DNA methylation (15% or more DNA methylation level than myometrium)			
SATB2	88.9% (8/9)	75% (9/12)	81% (17/21)
NRG1	100% (9/9)	75% (9/12)	85.7% (18/21)
mRNA expression (2-fold or more mRNA expression level than myometrium)			
SATB2	100% (9/9)	75% (9/12)	85.7% (18/21)
NRG1	100% (9/9)	66.7% (8/12)	81.0% (17/21)