

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

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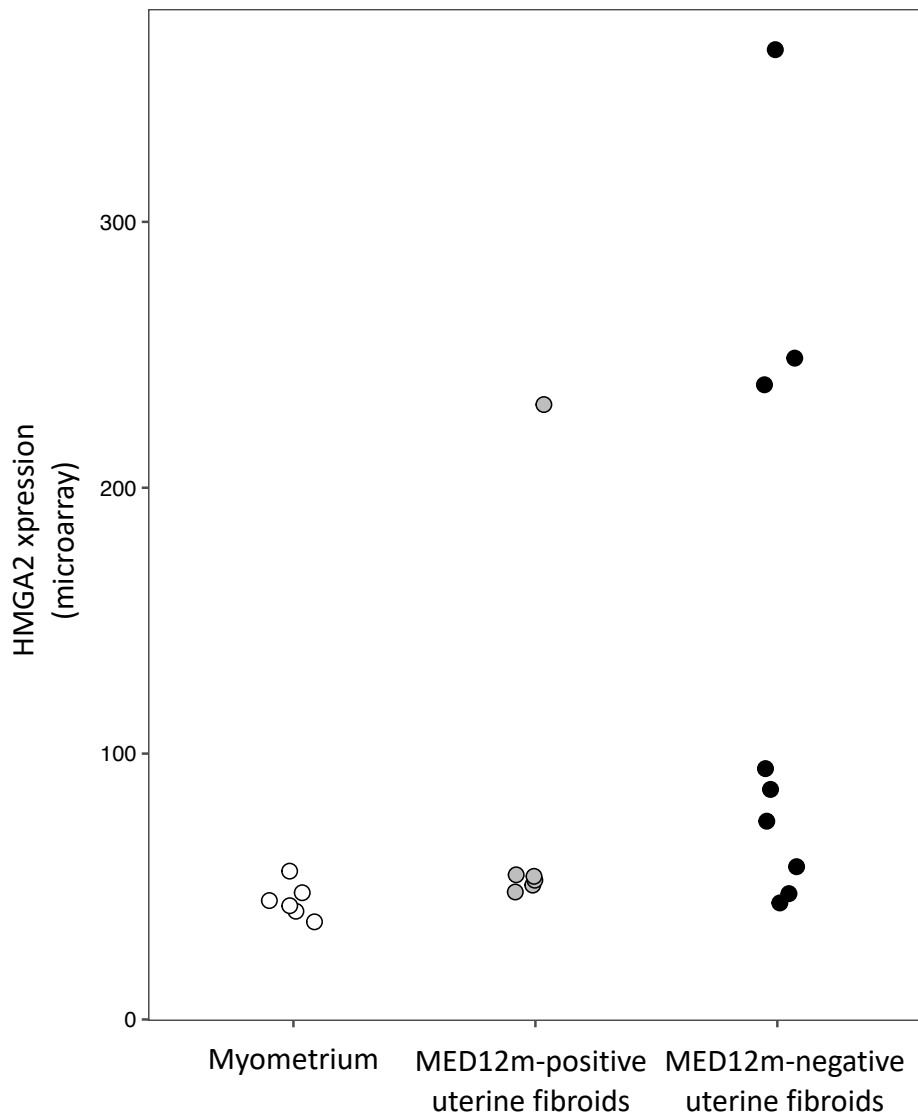
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Supplemental Figure S2. Expression levels of HMGA2 in the MED12m-positive and -negative uterine fibroids.

The expression levels in the transcriptome analysis are shown in the dot plot. Vertical axis indicates the signal intensity of HMGA2 in the myometrium (n = 6), MED12m-positive uterine fibroids (n = 6), and -negative uterine fibroids (n = 9).