



Supplementary Materials

DNA methylation of *FKBP5* as predictor of overall survival in Malignant Pleural Mesothelioma

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<u>cg03546163</u>

Figure S1. Locations of cg03546163 (CpG2, in bold) and a second CpG site very close (CpG1, in red), investigated by Sequenom MassARRAY. Flanking primers are underlined. The CpG sites could not be tested individually due to MassARRAY technology constraints, but had to be tested jointly as a single unit: the methylation level is the cumulative value of all the sites within the CpG unit.



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