

Title page

Idiopathic male infertility and polymorphisms in the DNA methyltransferase genes involved in epigenetic marking

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Figure legends

Figure S1. A predicted effect of the variant rs2424909 on transcription factor-binding site. Comparison between C allele (A) and T allele (B) of rs2424909 for the putative transcriptional factor-binding site.

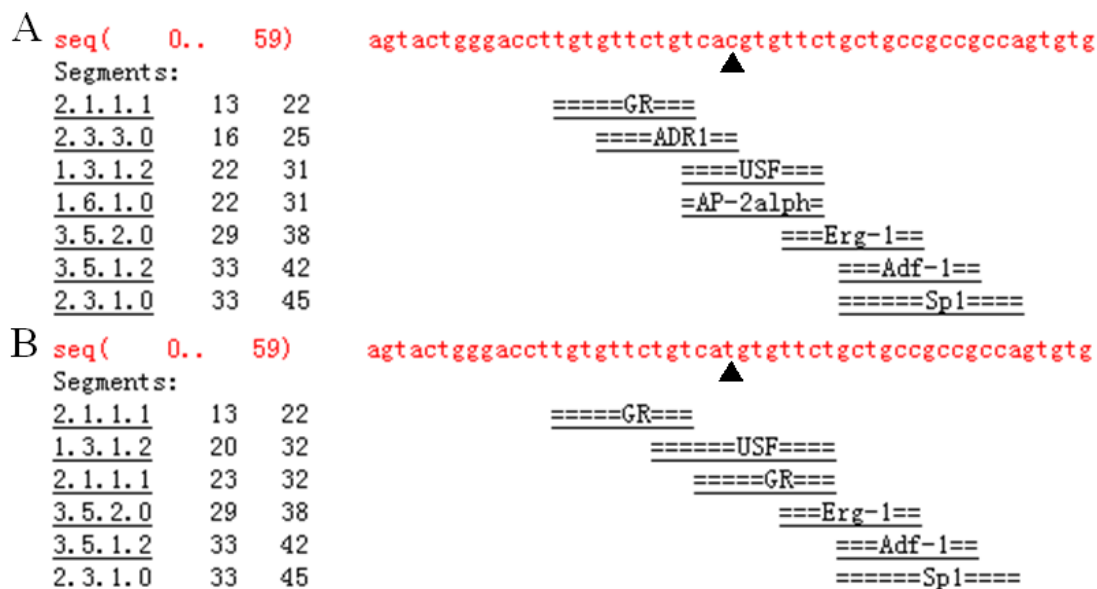


Figure S2. A predicted effect of the variant rs4804490 on transcription factor-binding site. Comparison between C allele (A) and A allele (B) of rs4804490 for the putative transcriptional factor-binding site.



Figure S3. A predicted effect of the variant rs1550117 on transcription factor-binding site. Comparison between A allele (A) and G allele (B) of rs1550117 for the putative transcriptional factor-binding site.



Figure S4. A predicted effect of the variant rs7354779 on transcription factor-binding site. Comparison between C allele (A) and T allele (B) of rs7354779 for the putative transcriptional factor-binding site.

