

## Methylation Sequence Submission

The sequence file must start with a unique identifier (gene\_CpG ID). On the next line or lines should be the sequences. The sequence can be all on one line or broken up on several lines. We recommend at least 500bp flanking sequence on both sides of the region of interest. The region of interest can be identified by highlighting the sequence (bold, color, italics). See the example below.

P57\_CpG1 (blue), \_CpG2 (gray)

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CTGCAGGGCCTTTCAAGTATGTCCGGAAGTCCTGGTTCCTCCAGGGCGGGGGG
AGGGGGCCAGCTCCTGGCACATGCCTGAGTCACCTTCTCGACTCCCTGCCCTA
ATAATGAACTCCTTTTGGCGGCGGTAGCGGAGGCAGCAGGGGGCGGGGGG
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GCCCCACCAAGAGGCACTGCTGTGGAGGAGGGGTCATTCTCTCTTTTCTAGCA
TAACTACTTCCAGCCAACTAAGGTAGGCAGAAGTGGGGGTCAATATGGCCT
GACCCAAACCTAAGGTCCTAGGCATCTATCAGTGATGTAGGTACAGGCTAGG
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GCTACAGATCTTTTAGATCTCTACAGTATTGTTCCCTCAGGAAAAGAATGGATG
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AGATGCCTCGAAGGCAGATACTTAGGCCTGGGTGTAGAGGGTGGATCCAGCC
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CACCGACGTCGGCATCGCCGAGGGGCTGTGCCAGCTCCATGGTCCGAGGGG
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TTACCGCCCGCAGAGCTCTTTAACTCGAGCCCCGCCCTTGTCGCGCAGCC
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ATGGAGGTGGACAGCGAGTCTGTGCCCGCCTTCTACCGCGAGACGGTGCAGG
TGGGGCGCTGTCGCCTGCAGCTGGGGCCCCGGCCACCCCCGGTGGCCGTGGC
TGTCATCCC CGCTTCTGGGGCCGCGGCTGGCGAGGCCCCCCGACGGCCTAGAG
GAGGCGCCTGAGCAGCCGCCAGCGCCCCAGCCTCGGCCGTGGTCGCGGAGC
CCACCCACCCGCGACCCCGGCCCGGCTTCAGATCTGACCTCAGACCCAAT
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TCCGGAGGTGACCCTGGTCGCGACCTCCGACCCGACTCCGGACCCGATCCCG  
GACGCGAACCCGGACGTGGCGACTCGGGACGGCGAGGAACAGGTCCCTGAG  
CAGGTCTCTGAGCAGGGCGAGGAGTCGGGTGCTGAGCCGGGTGATGAGCTGG  
GAACTGAGCCGGTCTCTGAGCAGGGCGAGGAGCAGGGCGCAGAGCCGGTCCG  
AGGAGAAGGACGAGGAGCCGGAGGAGGAGCAGGGCGCGGAGCCGGTCCGAG  
GAGCAGGGTGC GGAGCCGGTTCGAGGAGCAGAATGGGGAGCCGGTTCGAGGAG  
CAGGACGAGAATCAAGAGCAGCGCGGCCAGGAGCTGAAGGACCAGCCTCTC  
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GCATGGAGGGGGTTAAGTGCGGCGGCAGCCCCGGGGGGCTTGGCCGCGGGA  
CAAGGGGAAATGCTTACACAGCACATTGCGCGGCGACGTAAACAAAGCTGA  
CCCGCCGCGGACCTCGGCGCGGGCGGGGATGGCGCCCCACCCCGGCCCCGC  
GCCCTGCGCACTCCGCGGGTCTCCTCGCGGGTCTCTGGGCCGGCCCCGCCCTG  
ACCGACGCGCGTGTGTCGCCCGCAGACTTCTTCGCCAAGCGCAAGAG

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