```
        5'
- 120   TCCCACACCACTCGGCTCAGCCAATGGGAGGGCGCGCCGCGGGGCTTGGAGGTCCGCGCC
                                  - 39    - 33          - 23       - 15     - 9
                              - 41  - 37      - 30              - 17    - 11          - 1
 - 60   AACGGTGCTGAAAACCCTGCGCGCGAACGGCGGTCTCCGGGAGCGCGCTCGCGGCAGCCC
        GACTCT
            3'
```

Figure S5. Sequence of the 5' regulatory region of rat *Fads1*. CpG dinucleotides are indicated by bold text. Numbers above the text indicate the locations relative to the transcription start site (bases) of individual CpG dinucleotides which were measured by pyrosequencing. Arrow marks the transcription start site.