

B D C , B

D . C , B

A 19010-2899

Methodology

Statistical Analysis Results

the field (## #4#0#5#6/DMR0 5#0#6) of
4.,),6*),(/ 0 5#)1+*#),(/ ()1S) 8# .## S) ,5 3), ()#4 -#(.#.! In F

Acknowledgements

Support for this research was provided

Total Methylation Values:

hybrid mice, one allele of these genes will be methylated while the other is unmethylated based on genetic strain (BL/6 vs. *castaneus*). We will continue to focus our analysis on methylated alleles to test our hypothesis that the methylation at these non-