Biostatics Statement

We hereby declare that we have used online statistical tools for our analysis

1. For analyzing microarray data we have used Affymatrix TAC software for differential miRNA analysis. One Way ANOVA paired method was applied for assessing statistical significance. p-value of 0.05 was considered significant.

2. For analyzing qRTPCR data SPSS version 16.0 software (SPSS Inc., Chicago, Illinois, USA) and Graph Pad Prism version 5 (San Diego, California: Graph Pad Software Inc., 2007) were used to perform statistical analysis. Two-tailed p values ≤ 0.05 were considered significant.

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