Supplementary Figure 2: (A) Heat map showing the differentially regulated miRNA in the comparison of aACC and naACC carcinoma types as identified by SAM. The heat map was clustered using Euclidean distance and average linkage. (B) All discriminatory miRNA are overexpressed in aACC as compared to naACC. Data analysis was carried out as described in Supplementary Figure 1. The underlined miRNA were selected to be validated by real-time quantitative RT-PCR.