

Supplementary Figure 1. MiRNA expression profiles in pPCL patients. Patients are in columns, miRNAs in rows. The color scale bar represents the relative miRNA expression changes normalized by the standard deviation. The patients' molecular characteristics are shown above the matrix; n indicates unavailable information. (A) Hierarchical clustering of the samples using the 46 most variable miRNAs. (B) Heatmap of the differentially expressed miRNAs between pPCL patients carrying t(4;14), t(11;14), or t(14;16)/t(14;20). Differentially expressed miRNAs were identified by means of a SAM three-class analysis. MiRNA expression of the pPCL case (PCL-020) not carrying any IGH chromosomal translocation (and therefore not included in the analysis) is also shown.

