

Supplementary Table 1. MiRNAs whose expression in pPCL is significantly positively associated with the DNA copy number (CN) of the genomic locus harboring the corresponding gene (Kendall's τ correlation, one-sided p-value < .05).

miRNA	Chromosomal location ^a	CN=1,	CN=2,	CN=3,	CN=4,	CN=5,
		n	n	n	n	n
hsa-miR-197	1p13.3	4	11	1	/	/
hsa-miR-186	1p31.1	3	13	/	/	/
hsa-miR-30e	1p34.2	3	12	/	1	/
hsa-miR-30e*	1p34.2	3	12	/	1	/
hsa-miR-30c	1p34.2	3	12	/	1	/
hsa-miR-29c*	1q32.2	/	9	6	/	1
hsa-miR-28-5p	3q28	1	13	2	/	/
hsa-miR-103	5q35.1	1	13	1	1	/
hsa-miR-148a	7p15.2	1	13	1	1	/
hsa-miR-210	11p15.5	/	14	1	1	/
hsa-miR-483-5p	11p15.5	/	14	1	1	/
hsa-miR-331-3p	12q22	2	13	/	1	/
hsa-miR-15a	13q14.3	12	3	1	/	/
hsa-miR-92a	13q31.3	9	6	1	/	/
hsa-miR-19a	13q31.3	9	6	1	/	/
hsa-miR-20a	13q31.3	9	6	1	/	/
hsa-miR-20a*	13q31.3	9	6	1	/	/
hsa-miR-342-3p	14q32.2	3	12	/	1	/
hsa-miR-630	15q24.1	1	14	/	1	/
hsa-miR-1225-5p	16p13.3	2	13	/	1	/
hsa-miR-140-5p	16q22.1	6	8	/	1	1
hsa-miR-22	17p13.3	5	11	/	/	/
hsa-miR-181d	19p13.12	/	13	2	1	/

^aFor miRNAs produced from distinct genomic *loci*, the expression of the mature miRNA was correlated with the DNA CN of each of these genomic locations. The chromosomal location of the genomic *locus* resulted to be associated with mature miRNA expression is indicated, along with the number of pPCL cases having the reported inferred DNA copy number of that *locus*.