Additional File 4.

Title: MicroRNA selected for this study from miRNAs significantly associated with young women's breast cancer in discovery set and its validation.

Description: Clusters were constructed using Average hierarchical clustering method. Statistics of the discovery set were calculated via t-test. FDR stands for False Discovery rate multitesting corrected p-value. Only miRNAs remarked in bold were validated on an independent set of patients. Fold change of the validation set were calculated using ΔΔCt method and p-values were estimated via ANOVA test comparing the age groups and adjusted for multiple testing using Tukey's method. Bold denotes significant associated results in validation set.

Additional File 4. MicroRNA selected for this study from miRNAs significantly associated with young women's breast cancer in discovery set and its validation.

cluster	miR	Discovery set (n=37)		Validation set (n=56)
		FDR	Fold change	Adjusted p-value
Node 15	hsa-miR-1275	1.40×10^{-4}	5.89	**2.86×10^{-3}**
	hsa-miR-1207-5p	7.60×10^{-4}	4.86	0.047
	hsa-miR-4281	1.37×10^{-3}	4.61	
	hsa-miR-1268	7.2×10^{-4}	5.01	
	hsa-miR-663	1.29×10^{-3}	4.04	
	hsa-miR-149	4.10×10^{-3}	-4.03	0.712
	hsa-miR-762	1.40×10^{-4}	6.65	
	hsa-miR-3196	1.40×10^{-4}	6.74	**1.98×10^{-4}**
	hsa-miR-1908	1.40×10^{-4}	5.41	
	hsa-miR-1228*	1.40×10^{-4}	6.02	**4.73×10^{-5}**
	hsa-miR-3141	1.90×10^{-4}	5.30	
	hsa-miR-92b	7.00×10^{-4}	-4.60	**0.012**
Node 63	hsa-miR-139-5p	0.020	-3.13	**0.018**
	hsa-miR-132	1.40×10^{-4}	-5.57	0.908
	hsa-miR-379	0.092	-2.14	0.052
	hsa-miR-409-3p	0.043	-2.62	0.107
	hsa-miR-134	0.086	-2.20	
	hsa-miR-127-3p	0.050	-2.53	
	hsa-miR-433	0.093	-2.15	0.132
	hsa-miR-181a-2*	0.029	-2.92	

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