

Supplementary materials

Table 1S. Expression levels of *miR-205*, *BCL2* and *YAPI* determined in control and patient groups using the 2- Δ Ct method.

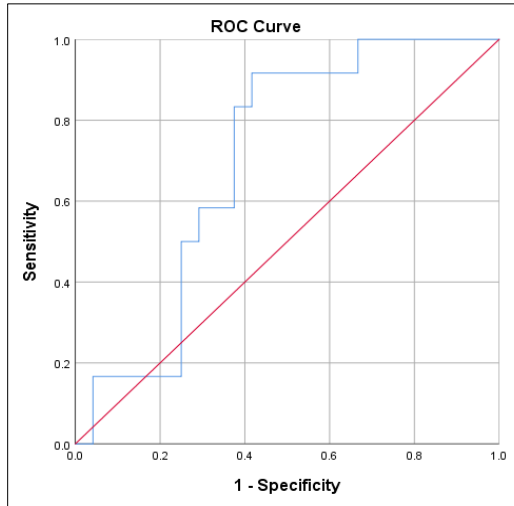
groups	Means Ct of Target gene	Means Ct of HKG	Δ Ct	2 ^{-ΔCt}	experimental group/ Control group	Fold of gene expression
<i>MiR-205</i>						
Patients	36.19	12.35	23.85	0.0000001	0.0000001/0.0000002	0.33
Control	34.68	12.39	22.29	0.0000002	0.0000002/0.0000002	1.00
<i>BCL2</i>						
Patients	21.67	14.59	7.08	0.00736	0.00736/0.00383	1.92
Control	22.29	14.26	8.03	0.00383	0.00383/0.00383	1.00
<i>YAPI</i>						
Patients	21.85	14.59	7.26	0.00651	0.00651/0.00451	1.44
Control	22.05	14.26	7.79	0.00451	0.00451/0.00451	1.00

*Target Gene: *miR-205*, *BCL2* and *YAPI*. HKG: House- Keeping genes (*GAPDH* and *snRNAU6*)

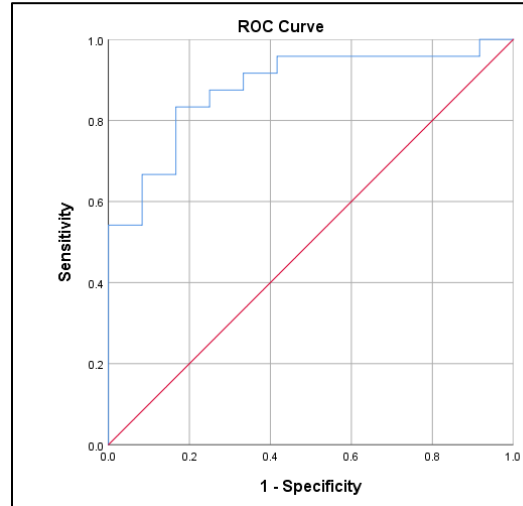
Table 2S. Pearson correlation coefficient factor.

		<i>miR-205</i>	<i>Bcl2</i>	<i>YAPI</i>
<i>miR-205</i>	Pearson Correlation	1	0.365*	0.236
	p value		0.029	0.166
<i>Bcl2</i>	Pearson Correlation		1	0.556**
	p value			0.000
<i>YAPI</i>	Pearson Correlation			1
	p value			

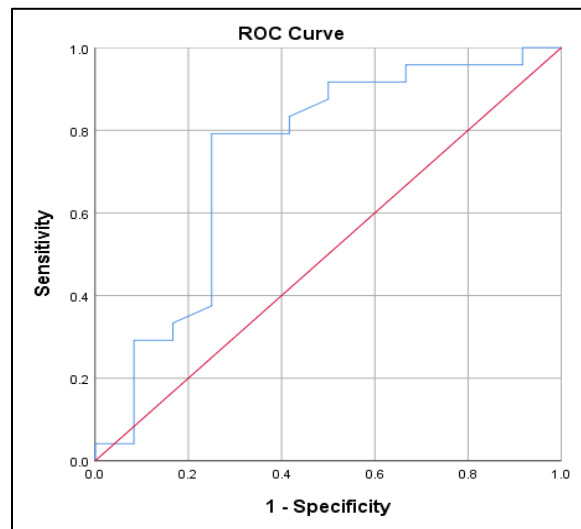
*: Significant p value; **: Highly Significant p value



(a) *miR-205*



(b) *BCL2*



(c) *YAP1*

Figure 1S. Receiver Operating Characteristic (ROC) curve of (a) *MicroRNA-205*, (b) *BCL2*, and (c) *YAP1*.