



## Aberrant expression of circ-0072309 in breast carcinoma as a promising biomarker for diagnosis

Hajar Zeidi<sup>1</sup>, Rezvan Esmaeili<sup>2</sup>, Sadegh Babashah<sup>1, \*</sup>

<sup>1</sup> Department of Molecular Genetics, Faculty of Biological Sciences, Tarbiat Modares University, P.O. Box: 14115-154, Tehran, Iran

<sup>2</sup> Genetics Department, Breast Cancer Research Center, Motamed Cancer Institute, ACECR, Tehran, Iran

Corresponding author: babashah@modares.ac.ir

### Background

Breast cancer (BC) is one of the most common cancers and the second highest mortality in women worldwide. Different studies have shown that there is a close correlation between circRNAs and initiation and progression of different cancers and diseases; however, knowledge about the correlation between non-coding RNAs, especially the circRNAs and BC, is not fully explored. In this study, the expression level of circ-0072309 (LIFR) in BC was investigated experimentally and miRNA target genes for it was proposed.

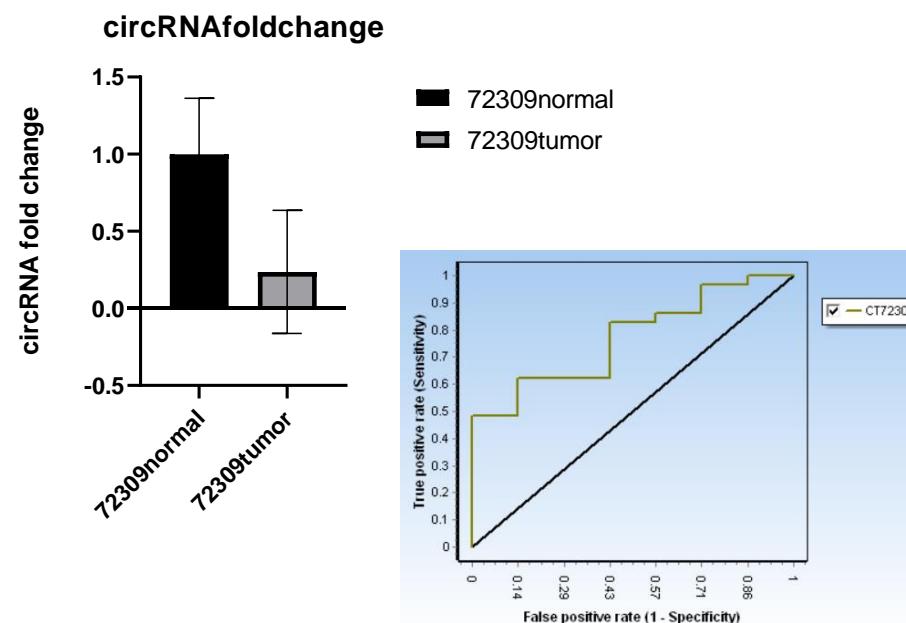
### Materials and Methods

BC tissue specimens (twenty-nine cases) and their corresponding adjacent normal tissue were collected during the surgery and has-circ-0072309 expression in breast cancer tissues was analyzed using qRT-PCR. A series of functional experiments were carried out to investigate has-circ-0072309 function in breast cancer development. Additionally, miRNAs that potentially target has-circ-0072309 were predicted by bioinformatic method. miRTargetlink database (<https://ccbweb.cs.uni-saarland.de/mirtargetlink>) and DIANA database (<http://diana.imis.athena-innovation.gr/DianaTools/index.php?r=mirpath/index/>) were used to predict interactions of miRNAs with LIFR.

### Results

The lower expression levels of circ-0072309 in BC tissues was showed compared to paired adjacent normal tissues with P value ( $P < 0.002$ ). The area under the receiver operating characteristic (ROC) curve was 0.768. In addition, a total 10 miRNAs that can be targeted by candidate circRNA was predicted base on bioinformatics databases. Moreover, miRNAs selection tools predicted that miR-125a-5p and miR-203a-3p can be targeted by circ-0072309.

Circ_0072309	Mean	SD	N	P-value
-Δct (T)	-9.7751	2.3489	29	0.002
-Δct (N)	-8.1425	1.3201	29	



### Conclusion

Our findings revealed that the has-circ-0072309 plays an essential role in BC progression and circ-0072309 could serve as a prognostic biomarker of BC.

### References

1. Yan L, Zheng M, Wang H. Circular RNA hsa\_circ\_0072309 inhibits proliferation and invasion of breast cancer cells via targeting miR-492. *Cancer Management and Research* 2019;11:1033–1041.
2. Zhao Y, Li J, Li J, Xu L, Lian W. The decreased circular RNA hsa\_circ\_0072309 promotes cell apoptosis of ischemic stroke by sponging miR-100. *Eur Rev Med Pharmacol Sci.* 2020;24:4420-4429.
3. Pang W, Huang F, Zhang X, Ye M, Huang Y, Huang X, Pang J, Cai C, Wang Z. Circular RNA hsa circ 0072309 inhibits non-small cell lung cancer progression by sponging miR-580-3p. *Bioscience Reports* 2020;40:1-10.