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To: Shicheng Guo (shicheng.guo@hotmail.com)

CLEP-D-15-00211

MiR-34a-5p promotes multi-chemoresistance of osteosarcoma through the down-regulation of the DLL1 gene

Youguang Pu; Fangfang Zhao; Haiyan Wang; Wenjing Cai; Jin Gao; Shanbao Cai

Clinical Epigenetics

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I would like to invite you to review the manuscript above which has been submitted to Clinical Epigenetics. Further details including the full abstract can be found at the end of this email.

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Best wishes,

Carmen Jeronimo, PhD

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CLEP-D-15-00211

Research

MiR-34a-5p promotes multi-chemoresistance of osteosarcoma through the down-regulation of the DLL1 gene

Clinical Epigenetics

Abstract: Abstract

Background: MicroRNAs have been identified as key players in the development and progression of osteosarcoma (OS). The molecular mechanisms that lead to OS development and metastasis are poorly understood. MiR-34a-5p has been implicated in the tumorigenesis and progression of several types of cancer, however, its role in OS remains elusive.

Results: We found that miR-34a-5p promotes multi-chemoresistance in OS cells. Using a systematic analysis of two multi-chemosensitive (G-292 and MG63.2) and two resistant (SJSA-1 and MNNG/HOS) OS cell lines, we showed that miR-34a-5p promotes OS multi-drug resistance via its repression of the Delta-like ligand 1 (DLL1) gene, the ligand of the Notch pathway, and thus negatively correlates with OS chemoresistance. siRNA-mediated repression of the DLL1 gene suppressed cell apoptosis and de-sensitized G-292 and MG63.2 cells, while overexpression of DLL1 sensitized SJSA-1 and MNNG/HOS cells to drug-induced cell death. Parallel to the changes in the drug-induced cell death, the activity of the ATF2/ATF3/ATF4 signaling pathway was drastically altered by a forced reversal of miR-34a-5p or DLL1 levels in OS cells.

Conclusions: We demonstrated that DLL1 is a direct target of miR-34a-5p and negatively regulates the multi-chemoresistance of OS through the activation of the ATF2/ATF3/ATF4 signaling pathway. This study also provided a new set of genes in this newly identified miR-34a-5p/DLL1/ATF axis as diagnostic targets for guided anti-OS chemotherapy, including the level of the miR-34a-5p gene, DLL1 gene and the ATF2/ATF3/ATF4 signaling pathway-associated genes.

