

miR-451 expression impacts key biological processes through regulation of PI3K/AKT pathway and sensitizes human head and neck cancer cell lines to Allitinib - a EGFR inhibitor.

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Introduction: MicroRNAs (miRNAs) are a class of small non-coding RNAs involved in gene expression regulation. They have important roles in several biological processes such as cell development, differentiation, cell cycle regulation, proliferation, migration, invasion, apoptosis and chemoresistance. MicroRNA-451 has emerged as an important tumor suppressor that is downregulated in several tumor types. Moreover, activation of the PI3K/AKT signaling pathway has been associated with the loss of miR-451 expression and poor prognosis in head and neck squamous cell (HNSCC). Activation of PI3K/AKT pathway has been described as an important mechanism implicated in tumor progression and resistance to epidermal growth factor receptor (EGFR) inhibitors. Up to 90% HNSCC present higher expression levels of EGFR, this is associated with poor prognosis. Allitinib is a new irreversible and promising selective EGFR inhibitor for the treatment of solid tumors. **Objectives:** We evaluated the biological role of miR-451 overexpression in head and neck cancer cell lines, and its impact in the response to Allitinib treatment in these cell lines. **Methods:** We determined miR-451 expression in six human HNSCC cell lines by RT-qPCR. Overexpression of miR-451 was performed in two cell lines (JHU-12 and JHU-28) and functional assays were conducted to assess its biological impact regarding invasion, migration, proliferation, apoptosis, cytotoxicity and colony formation. We also evaluated the protein levels of PI3K/AKT pathway members by western blotting. The role of miR-451 overexpression in Allitinib treatment was evaluated monitoring the cell viability using MTS assay. **Results:** miR-451 overexpression inhibited migration and clonogenic potential of JHU-12, and invasion capacity of JHU-28. Furthermore, we observed a reduction of proliferation rate in JHU-12 and JHU-28. Also, miR-451 overexpression was associated with lower protein levels of PI3K and phosphorylated AKT. Interestingly, we observed that miR-451 transfected JHU-28 cell line was more sensitive to Allitinib treatment. **Conclusion:** Our results demonstrated that miR-451 should be involved in the control of progression and metastatic potential in HNSCC, potentially through the regulation of PI3K/AKT pathway. Also, our data demonstrated that miR-451 overexpression might sensitize HNSCC cell line to Allitinib treatment, and its role as a potential new strategy for the treatment of chemo resistant HNSCC need to be explored.

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