<u>The Expression of hsa-miR-155-5p in Plasma Samples Of Breast Cancer Before And After Chemotherapy</u>

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Abstract	Breast cancer has emerged as the most common cancer-related mortality among women worldwide. Therefore, early cancer detection using biomarkers such as microRNA is needed. One of microRNAs that has an important role in breast cancer development is miR-155. HsamiR-155-5p is an oncomir that is commonly dysregulated in breast cancer. This study aims to determine the expression of hsa-miR-155-5p in breast cancer patient \tilde{A} \hat{C} \hat{A} \hat{C} \hat{A} ms plasma before and after chemotherapy. We collected 64 samples from breast cancer patients admitted to Dr. Sardjito Hospital in Yogyakarta. RNA from plasma was extracted using RNA Isolation Kit miRCURY-Biofluid. cDNA synthesis was performed using cDNA Synthesis kit II and quantification of miR-155-5p using ExiLent SYBR Green master mix (Exiqon). qRT-PCR results were then analyzed with Livak's method and compared (before and after chemotherapy) with t-test. Expression of miR-155-5p in the breast cancer patients \tilde{A} \tilde{C} \tilde{C} \tilde{C} method to before chemotherapy was significantly increased (10.59 times) when compared to before chemotherapy (p = 0.001). We concluded that there was upregulated expression of miR-155-5p after chemotherapy than before chemotherapy. There has not been a known, relevant pathway between hsa-miR-155-5p and chemotherapy regimens nor resistance to chemotherapy. \tilde{C} \tilde{C} has a mirror resistance to chemotherapy.
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Author	DWI NUR INDAH SARI, S.Si, M.Sc., M.Sc.