

Open Access

# The Potential of Rapamycin as a Selected Drug for Apoptosis Induction in Triple Negative Breast Cancer Subtype Using High-Throughput Transcriptome Analysis

## Maryam Peymani\* and Mehdi Roshanian Bakhsh

Shahrekord Branch, Islamic Azad University, Shahrekord, Iran

#### Abstract

Triple-negative breast cancers (TNBC), which lack estrogen and progesterone receptors demonstrate HER2/ neu overexpression. The poor survival outcomes coupled with TNBC patients are partially, due to a lack of therapeutic targets. The aim of this study was to introduce a new drug among the conventional chemotherapy drugs based on the trans cripts of the patients. For this purpose, transcript of MDA-MB-468 cell line as a TNBC model was analyzed in the presence of 15 common drugs in fifteen GEO datasets. Then, reduced gene lists in the presence of each drug in three breast cancer subtypes were compared based on TCGA data. The effect of the selected drug on survival and apoptosis rates of MDA-MB-468 and MCF-7 cells (as triple positive model) were compared. The data showed that the expression of the reduced gene list in the presence of 3000 µg/mL Rapamycin there was a sharp decrease in MDA-MB-468 cell viability and a significant increase in apoptosis compared to MCF-7 cells treated with the same concentration of the drug. The expression of PSAT1 as a gene with highest score among 44 common increased genes between all three subtypes, showed a reduction in presence of Rapamycin based on microarray data and was verified *in vitro* using RT-qPCR. In summery here, we have developed a novel computational method that allows the imputation of drug response in very large clinical cancer genomics data sets, such as TCGA.

### **Recent Publications**

1. Nazempour N (2021)The role of cell surface proteins gene expression in diagnosis, prognosis, and drug resistance of colorectal cancer: In silico analysis and validation. Exp Mol Pathol 123:104688.

2. Mahdevar M (2021) Steroid receptor RNA activator gene footprint in the progression and drug resistance of colorectal cancer through oxidative phosphorylation pathway. Life Sci 285:119950.

3. Miralaei N (2021) Integrated pan-cancer of AURKA expression and drug sensitivity analysis reveals increased expression of AURKA is responsible for drug resistance. Cancer Med 10: 6428-6441.

4. Ghanizade P (2021) Differential expression analysis of CCDC107 and RMRP lncRNA as potential biomarkers in colorectal cancer diagnosis. Nucleosides Nucleotides Nucleic Acids 40: 1144-1158.

#### Biography

Maryam Peymani has her expertise in cancer and genomics. She has recently published several articles in this field. She is currently an Associate Professor of Biology at University of Islamic Azad University Shahrekord branch.

\*Corresponding author: Kevin O. Ochwedo, University of Nairobi/Sub-Sahara International Centre of Excellence for Malaria Research, Nairobi, Kenya

**Received:** 01-Sep-2022, Manuscript No: jpgb-22-133, **Editor assigned:** 05-Sep-2022, Pre QC No: jpgb-22-133(PQ), **Reviewed:** 19-Sep-2022, QC No: jpgb-22-133, **Revised:** 26-Sep-2022, Manuscript No: jpgb-22-133 (R), **Published:** 30-Sep-2022, DOI: 10.4172/jpgb.1000133

Citation: Peymani M, Bakhsh MR (2022) The Potential of Rapamycin as a Selected Drug for Apoptosis Induction in Triple Negative Breast Cancer Subtype Using High-Throughput Transcriptome Analysis. J Plant Genet Breed 6: 133.

**Copyright:** © 2022 Ochwedo KO. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.