

Histone Deacetylase Inhibitors Reprogram Triple Negative Breast Cancer to be Less Aggressive by Targeting Cell Proliferation, Metastasis, and Cancer Stemness

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Triple negative breast cancer (TNBC) is highly aggressive and the most difficult type of breast cancer to treat due to its lack of receptors to target. Novel studies have explored the epigenetic mechanisms to inhibit or reverse aggressive characteristics of this subtype of breast cancer. Histone deacetylases (HDACs) control gene expression by changing the chromatin conformation to a closed state, which leads to transcriptional repression of certain genes. Recent studies have suggested that specific inhibitors targeting HDAC may cause tumor suppressor genes to be re-expressed in cancer cells, reversing cancer progression. Epigenetic mechanisms may also play a role in reprogramming cancer stem cells (CSCs) in TNBC. CSCs are primarily drivers of recurrence and metastasis, and therefore associated with poorer prognoses. In this study, we have examined the effects of HDAC inhibitors on cell proliferation, metastasis, and cancer stemness in TNBC cells. Two TNBC cell lines, 4T1 and SUM-159, were treated with the HDAC inhibitors, Vorinostat (SAHA) and Panobinostat, and their effects on cell proliferation, tumor suppressor genes and metastatic markers were determined by MTT assay and qPCR analysis. HDAC inhibitors on the formation of mammospheres and the stemness properties were further determined. We found that HDAC inhibitors decrease cell proliferation and downregulate cancer genes such as MYC, NFKb, and MMP9 while upregulating tumor suppressor genes such as CDKN1a. HDAC inhibitors have a modest effect on mammosphere forming efficiency and the CD24^{low}/CD44^{high} subpopulation. In conclusion, HDAC inhibitors target the cell proliferation pathway and to a lesser extent cancer stemness in TNBC. Supported by NIH R25ES020721.

