



Epigenetic reprogramming induced by lipids fosters mammary cell plasticity in non-transformed breast epithelial cells

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Introduction

- Understanding the genesis of sporadic estrogen receptor negative breast cancer (ERnegBC) is a significantly unmet clinical need.
- Genes involved in **lipid metabolism** are overexpressed in the contralateral unaffected breast of women with ERnegBC (1).
- Exposure of non-transformed breast epithelial cells to lipids results in significant changes in histone PTMs and gene expression. The upregulated genes are involved in neural pathways and stemness (2).
- Neural genes are highly expressed in a specific type of ERnegBC: Triple Negative Breast Cancer (3).
- Our aim is to identify potential mechanisms linking lipids and epigenetic reprogramming to the genesis of ERnegBC**

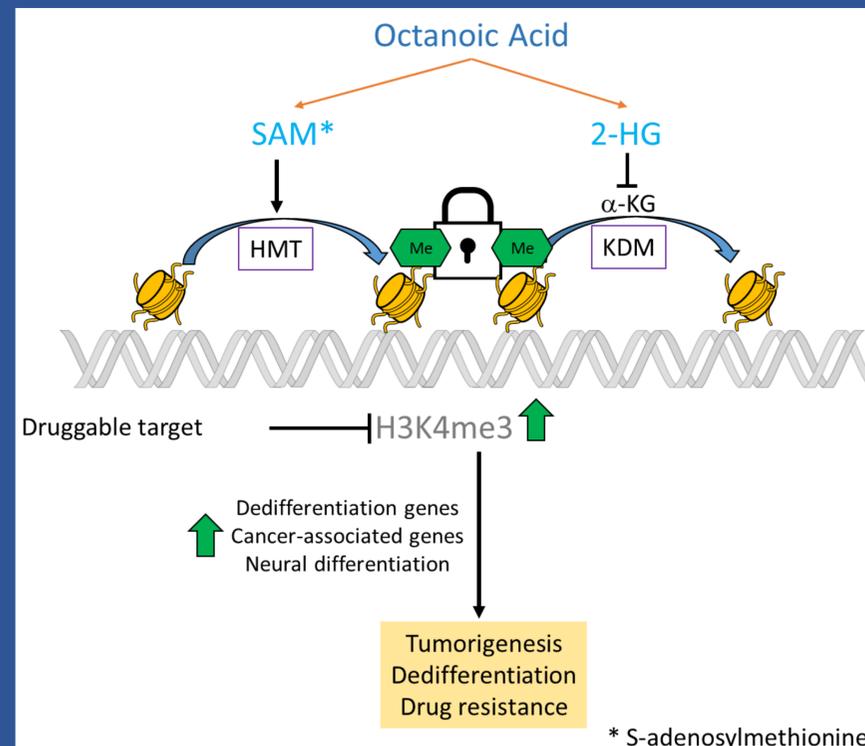
Methods

- ¹³C-glucose tracing was performed in MCF-10A cells exposed to octanoic acid (OA) ± PHGDH inhibitor.
- CUT&RUN for H3K4me3 was performed in MCF-10A exposed to OA. MACS2, DiffBind and ChIPseeker were used to call and annotate peaks. HOMER was used for Transcription factor (TF) binding motif enrichment analysis.
- Gene expression was measured by qPCR in mammary microstructures derived from breast tissue exposed to OA.
- Single-cell RNA-Seq (scRNA-seq) was performed on primary human breast epithelial cells exposed to OA. The digital expression matrix file containing UMIs will be analyzed with the Seurat package version 2.3.4 R version 3.5.3 The Aldefluor assay was used to identify stem-like (ALDH+) cells upon OA.
- To determine if lipid-exposed cells adopt a neural-like phenotype, cells were grown on Poly-D-Lysine/Laminin (PDL/LM) coated plates.

References

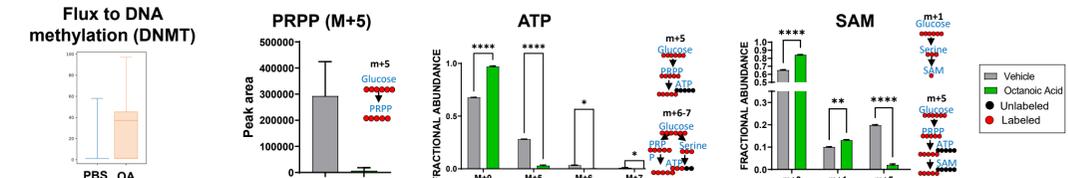
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Lipid exposure increases the production of **SAM** and **2-HG** resulting in **epigenetic fostered plasticity**, which selects cells with a multi-potential **embryonic or stem-like state** and reprograms differentiation to a **neural/neural crest-like state**

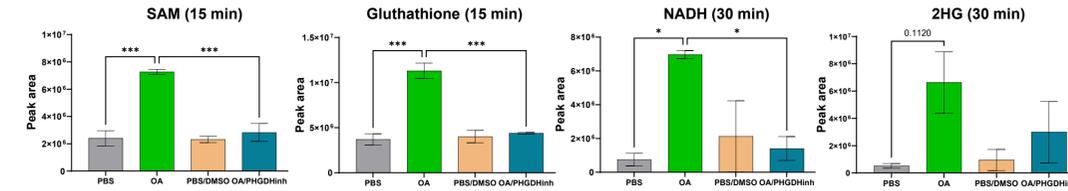


Results

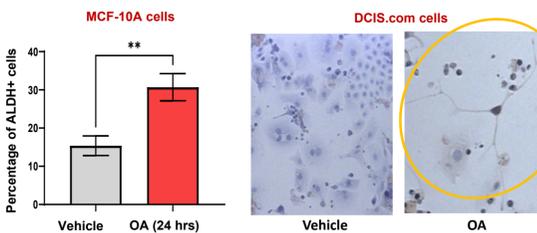
A OA decreased flux through the pentose phosphate pathway (PPP) & increased fractional abundance of SAM M+1 through the serine pathway



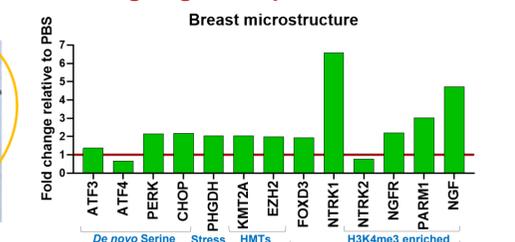
B Metabolites levels increased by OA; prevented by inhibiting PHGDH



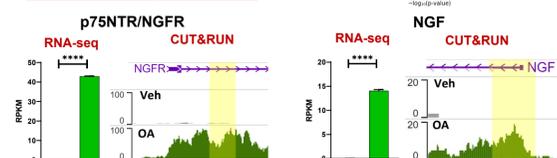
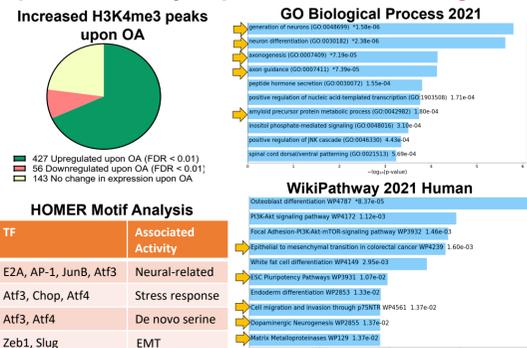
C OA enriched ALDH+ cells and cells develop a neural-like phenotype



D Exposure of breast microstructure to OA changes gene expression



E 661 differentially enriched H3K4me3 peaks upon OA mainly in promoters of neural genes



F OA changes the distribution of cell subpopulations (e.g. LP3) and modulates gene expression within each subcluster (e.g. BSL1)

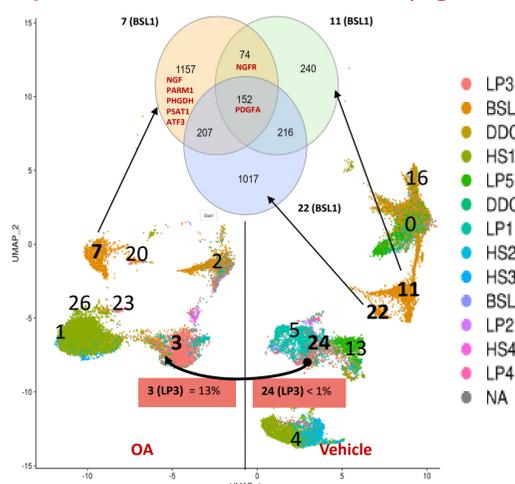


Figure 1. Metabolomics (A,B). Aldefluor test and development of neural-like phenotype (C). qPCR (D). CUT&RUN for H3K4me3 (E). scRNAseq (F).