

# Licochalcone A is a candidate for breast cancer prevention through its reprogramming of metabolic and antioxidant pathways

Atieh Hajirahimkhan<sup>1</sup>, Elizabeth T. Bartom<sup>2</sup>, Sriram Chandrasekaran<sup>3</sup>, Xiaoling Xuei<sup>4</sup>, Susan E. Clare<sup>1</sup>, Seema A. Khan<sup>1</sup>

<sup>1</sup>Division of Breast Surgery, Robert H. Lurie Comprehensive Cancer Center, Northwestern University, <sup>2</sup>The Louis A. Simpson and Kimberly K. Querrey Biomedical Research Center, Northwestern University, Chicago, IL. <sup>3</sup>Department of Biomedical Engineering, University of Michigan, Ann Arbor, MI, <sup>4</sup>School of Medicine, Indiana University, Indianapolis, IN.

## BACKGROUND

- Less than 5% of women who could benefit from breast cancer risk reduction drugs report taking them, mainly due to the adverse effects of these medications.<sup>1</sup> There is no drug for preventing ER- cancer.
- Prevention strategies with optimal efficacy, less toxicity, and greater acceptance are needed.**
- Natural products with significantly lower toxicity and sufficient efficacy to shift the breast microenvironment to a tumor preventive milieu are ideal candidates.<sup>2</sup>
- Previously, we have shown that licochalconeA (LicA) from licorice inhibits aromatase activity and has antioxidant potential.<sup>3,4,5</sup>
- We now report on the response of high-risk postmenopausal human breast tissue, breast pre-malignant and malignant cells to LicA treatment *in vitro* and *in vivo*.

## OBJECTIVE

LicA modulates metabolic and antioxidant pathways in the breast leading to a tumor preventive environment.

## METHODS

- Contralateral unaffected breast tissue of 6 postmenopausal women, who had bilateral mastectomy due to unilateral breast cancer were obtained and processed to microstructures.
- Microstructures were treated with DMSO and LicA (5  $\mu$ M) for 24 h, prior to RNA extraction and total RNA sequencing.
- Differential gene expression was determined. Gene ontology (GO) pathway analysis was performed. The enriched pathways with combined enrichment scores > 4 and FDR < 0.05 were considered statistically significant. The differential gene expression results were further analyzed with computational metabolic flux analysis and NanoString metabolism panel. Modulated pathways with P < 0.05 were considered significant.
- Live cell imaging/proliferation was analyzed in DCIS.COM/ER+ PR+, DCIS.COM, MCF-7, and MDA-MB-231 cells treated with single and repeated doses of LicA.
- Female athymic nude mice inoculated with luminal or triple negative breast cancer cells, received LicA or vehicle for 28 days at the dose of 80 mg/kg/day and tumor volume was evaluated.



Supported by the Postdoctoral Fellowships: American Cancer Society 131667-PF-18-049-01NEC, RHLCCC Translational Bridge, NCI-CONSORT-T32, and by Bramsen-Hamill Foundation.

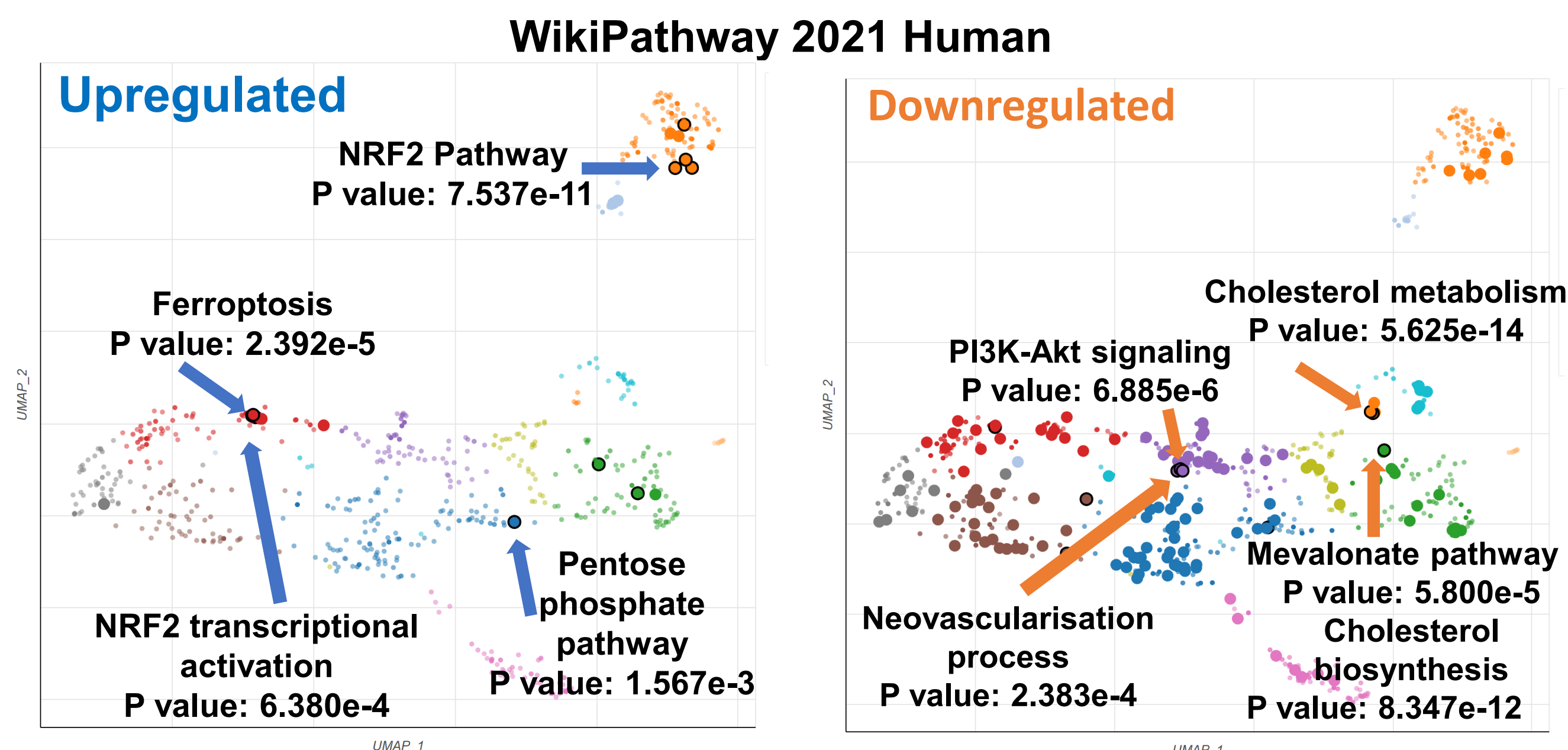
Copies of this poster may not be reproduced without permission from the author of this poster. Contact them at [atieh.hajirahimkhan@northwestern.edu](mailto:atieh.hajirahimkhan@northwestern.edu) for permission to reprint and/or distribute.

*Can a natural product protect high risk women from breast cancer?*

**Licochalcone A is a good candidate**

- ✓ In high-risk women's breast microstructures
- ✓ In ER+ and ER- pre-malignant and malignant breast cells
- ✓ In xenograft mouse models of luminal BC and TNBC

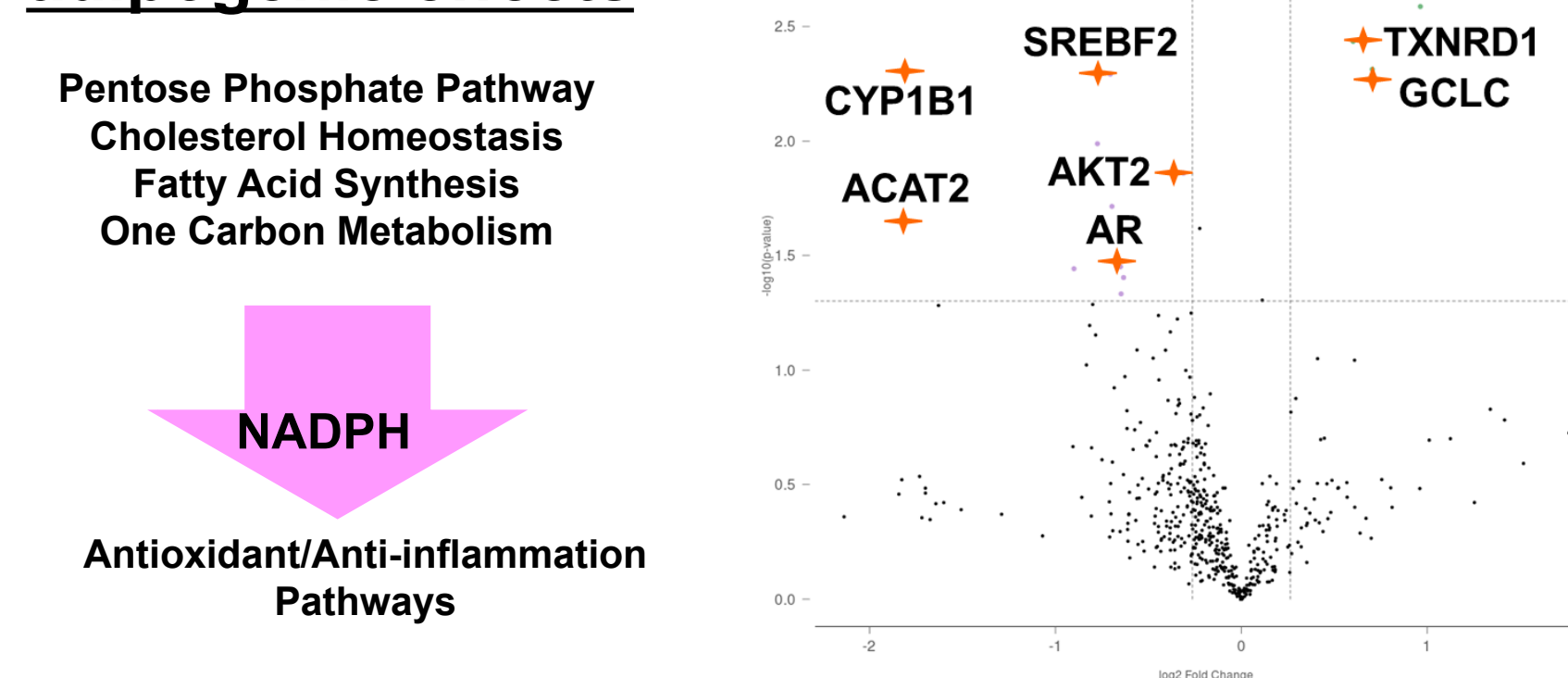
## RESULTS



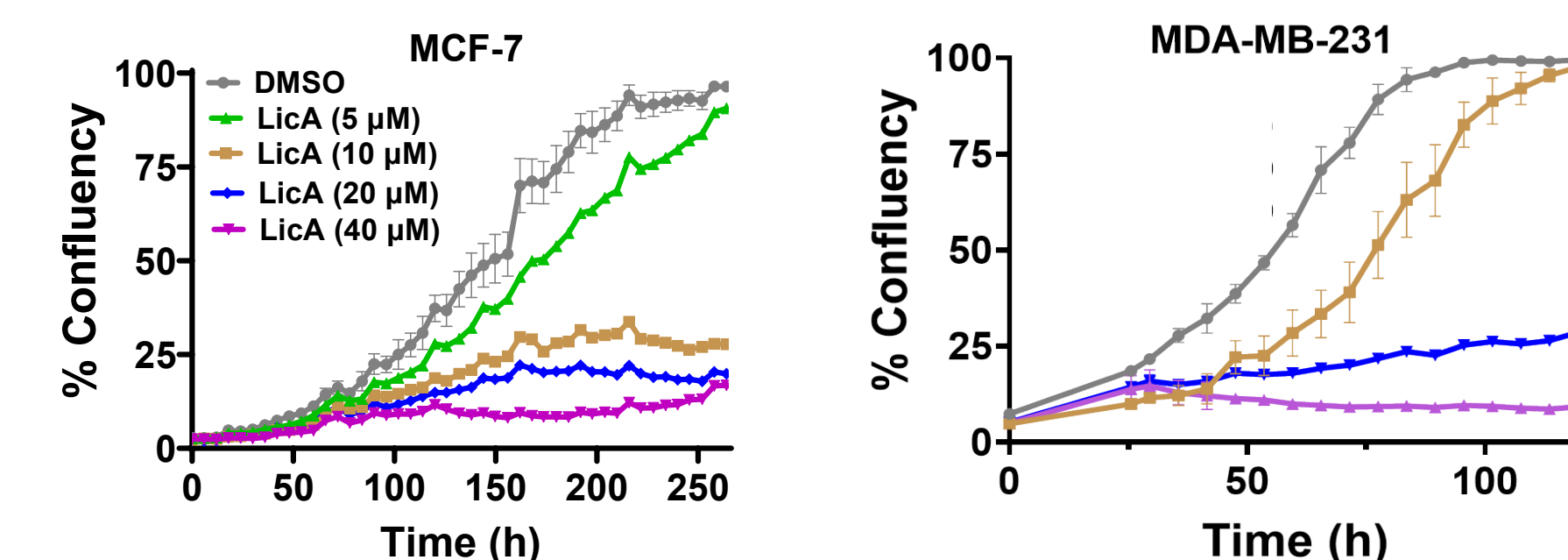
## REFERENCES

- Mol. Cell. Endocrinol. 2021, 530: 111284.
- Pharmacol. Rev. 2016, 68: 1026.
- Chem. Res. Toxicol. 2015, 28: 2130.
- Cancer Prev. Res. 2018, 11: 819.
- bioRxiv*, doi:10.1101/2022.05.06.490985.
- Genome Biol. 2019, 20: 49.

**Metabolic flux (left) and NanoString metabolism analysis (right) support antioxidant and anti-adipogenic effects**

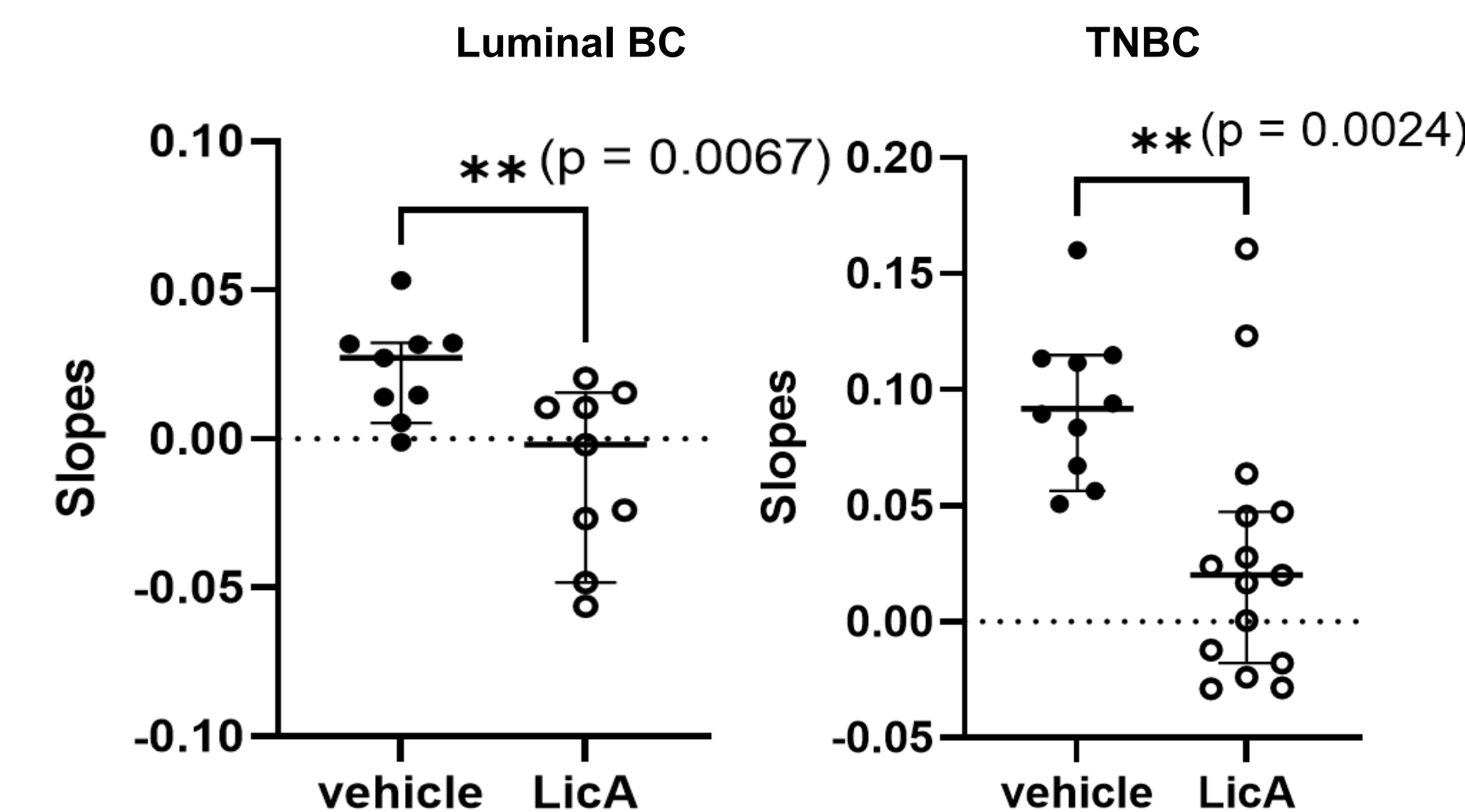


**LicA retards *in vitro* proliferation**



Representative live cell imaging data using IncuCyte showing that LicA can retard proliferation of MCF-7 (ER+), MDA-MB-231 (ER-) breast cancer cells.

**LicA reduces the growth of mammary tumors**



The slopes of tumor growth for every animal was generated using linear regression. The differences in tumor growth was statistically compared using unpaired t-test (two tailed). Data presented as median with 95% CI.