

Resource Summary Report

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Susan G. Komen Tissue Bank

RRID:SCR_004708

Type: Tool

Proper Citation

Susan G. Komen Tissue Bank (RRID:SCR_004708)

Resource Information

URL: <https://komentissuebank.iu.edu/>

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Description: The goals of the Susan G. Komen for the Cure Tissue Bank at the IU Simon Cancer Center are to acquire biomolecule and tissue specimens from the entire continuum of breast development: puberty to menopause and to make these specimens or the digital data derived from them available and accessible to researchers across the globe.

Abbreviations: KTB

Synonyms: Tissue Bank at the IU Simon Cancer Center, Susan G. Komen for the Cure Tissue Bank, Komen Tissue Bank

Resource Type: biomaterial supply resource, material resource, tissue bank

Related Condition: Breast cancer

Resource Name: Susan G. Komen Tissue Bank

Resource ID: SCR_004708

Alternate IDs: nlx_70500

Record Creation Time: 20220129T080226+0000

Record Last Update: 20260727T040355+0000

Ratings and Alerts

No rating or validation information has been found for Susan G. Komen Tissue Bank.

No alerts have been found for Susan G. Komen Tissue Bank.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Zhao T, et al. (2024) A proteome-wide association study identifies putative causal proteins for breast cancer risk. British journal of cancer, 131(11), 1796.

Bhat-Nakshatri P, et al. (2024) Single-nucleus chromatin accessibility and transcriptomic map of breast tissues of women of diverse genetic ancestry. Nature medicine, 30(12), 3482.

Guo Q, et al. (2022) Mammary collagen is under reproductive control with implications for breast cancer. Matrix biology : journal of the International Society for Matrix Biology, 105, 104.

Niehoff NM, et al. (2020) Outdoor air pollution and terminal duct lobular involution of the normal breast. Breast cancer research : BCR, 22(1), 100.

Bodelon C, et al. (2020) Polygenic risk score for the prediction of breast cancer is related to lesser terminal duct lobular unit involution of the breast. NPJ breast cancer, 6, 41.

Hastings ML, et al. (2012) Sensitive PCR-based quantitation of cell-free circulating microRNAs. Methods (San Diego, Calif.), 58(2), 144.