

Resource Summary Report

Generated by [NIF](#) on Aug 13, 2026

Susan G. Komen for the Cure

RRID:SCR_004842

Type: Tool

Proper Citation

Susan G. Komen for the Cure (RRID:SCR_004842)

Resource Information

URL: <http://ww5.komen.org/>

Proper Citation: Susan G. Komen for the Cure (RRID:SCR_004842)

Description: Susan G. Komen for the Cure is the global leader of the breast cancer movement, having invested more than \$1.9 billion since inception in 1982. As the world's largest grassroots network of breast cancer survivors and activists, we're working together to save lives, empower people, ensure quality care for all and energize science to find the cures. Thanks to events like the Susan G. Komen Race for the Cure and the Susan G. Komen 3-Day for the Cure, and generous contributions from our partners, sponsors and fellow supporters, we have become the largest source of nonprofit funds dedicated to the fight against breast cancer in the world.

Abbreviations: Susan G. Komen

Synonyms: Susan G. Komen

Resource Type: funding resource

Resource Name: Susan G. Komen for the Cure

Resource ID: SCR_004842

Alternate IDs: Wikidata: Q246898, grid.428834.1, nlx_82488, Crossref funder ID: 100000869, ISNI: 0000 0001 0241 5320

Alternate URLs: <https://ror.org/02nadbe75>

Record Creation Time: 20220129T080226+0000

Record Last Update: 20260808T185828+0000

Ratings and Alerts

No rating or validation information has been found for Susan G. Komen for the Cure.

No alerts have been found for Susan G. Komen for the Cure.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 24 mentions in open access literature.

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

Franks JL, et al. (2020) In silico APC/C substrate discovery reveals cell cycle-dependent degradation of UHRF1 and other chromatin regulators. PLoS biology, 18(12), e3000975.

Funnell T, et al. (2019) Integrated structural variation and point mutation signatures in cancer genomes using correlated topic models. PLoS computational biology, 15(2), e1006799.

Lukong KE, et al. (2017) Understanding breast cancer - The long and winding road. BBA clinical, 7, 64.

Zhao SG, et al. (2015) Development and validation of a novel platform-independent metastasis signature in human breast cancer. PloS one, 10(5), e0126631.

Callaway JB, et al. (2015) Source and Purity of Dengue-Viral Preparations Impact Requirement for Enhancing Antibody to Induce Elevated IL-1 β Secretion: A Primary Human Monocyte Model. PloS one, 10(8), e0136708.

Ambert KH, et al. (2013) Virk: an active learning-based system for bootstrapping knowledge base development in the neurosciences. Frontiers in neuroinformatics, 7, 38.

Eckman DM, et al. (2013) Weekly doxorubicin increases coronary arteriolar wall and adventitial thickness. PloS one, 8(2), e57554.

Tabchy A, et al. (2013) Systematic identification of combinatorial drivers and targets in cancer cell lines. PloS one, 8(4), e60339.

Wallace JA, et al. (2013) Ets2 in tumor fibroblasts promotes angiogenesis in breast cancer. PloS one, 8(8), e71533.

Kumar S, et al. (2012) Tissue transglutaminase constitutively activates HIF-1 β promoter and nuclear factor- κ B via a non-canonical pathway. PloS one, 7(11), e49321.

D'Amato NC, et al. (2012) Evidence for phenotypic plasticity in aggressive triple-negative breast cancer: human biology is recapitulated by a novel model system. PloS one, 7(9), e45684.

Wang L, et al. (2012) Using Internet search engines to obtain medical information: a comparative study. *Journal of medical Internet research*, 14(3), e74.

Kumar A, et al. (2012) Evidence that GTP-binding domain but not catalytic domain of transglutaminase 2 is essential for epithelial-to-mesenchymal transition in mammary epithelial cells. *Breast cancer research : BCR*, 14(1), R4.

Li N, et al. (2011) Inhibition of cell proliferation by an anti-EGFR aptamer. *PloS one*, 6(6), e20299.

Stojadinovic A, et al. (2011) Consensus recommendations for advancing breast cancer: risk identification and screening in ethnically diverse younger women. *Journal of Cancer*, 2, 210.

Kumar A, et al. (2011) Evidence that aberrant expression of tissue transglutaminase promotes stem cell characteristics in mammary epithelial cells. *PloS one*, 6(6), e20701.

Figueiro G, et al. (2011) Control region variability of haplogroup C1d and the tempo of the peopling of the Americas. *PloS one*, 6(6), e20978.

Dong J, et al. (2010) Shorter exposures to harder X-rays trigger early apoptotic events in *Xenopus laevis* embryos. *PloS one*, 5(1), e8970.

Liang Y, et al. (2010) BRIT1/MCPH1 is essential for mitotic and meiotic recombination DNA repair and maintaining genomic stability in mice. *PLoS genetics*, 6(1), e1000826.

Sun Y, et al. (2010) Adiponectin deficiency promotes tumor growth in mice by reducing macrophage infiltration. *PloS one*, 5(8), e11987.