

Stanford



Esther M. John

Professor (Research) of Medicine (Oncology) and, by courtesy, of Health Research and Policy (Epidemiology)

Medicine - Oncology

CONTACT INFORMATION

• Administrative Contact

Sanavy Lou

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Bio

BIO

Esther M. John, PhD, MSPH is Professor in the Department of Medicine (Oncology) and co-leader of the Population Sciences Program of the Stanford Cancer Institute.

Dr. John is a cancer epidemiologist and her research focuses on the etiology and prognosis of breast and prostate cancer, to better understand modifiable lifestyle,

hormonal and genetic causes and outcomes of these common cancers. She has a special interest in understanding cancer health disparities and cancer in Hispanics and

African Americans, two understudied populations.

ACADEMIC APPOINTMENTS

- Professor (Research), Medicine - Oncology
- Professor (Research) (By courtesy), Health Research & Policy
- Member, Stanford Cancer Institute

ADMINISTRATIVE APPOINTMENTS

- Co-leader, Population Sciences Program, Stanford Cancer Institute, (2014- present)
- Co-leader, Cancer Epidemiology Program, Stanford Cancer Institute, (2005-2014)
- Lecturer & Consulting Assistant/Associate/Full Professor, Dept. Health Research & Policy, Stanford University School of Medicine, (1994-2017)
- Director of Research, Cancer Prevention Institute of California, (2015-2018)
- Director of Epidemiology, Cancer Prevention Institute of California, (2000-2002)

BOARDS, ADVISORY COMMITTEES, PROFESSIONAL ORGANIZATIONS

- Member, Editorial Board, Cancer Epidemiology, Biomarkers & Prevention (2013 - present)
- Member, American Association of Cancer Research (AACR) (2007 - present)
- Member, American Public Health Association (APHA) (1991 - present)
- Member, Society of Epidemiologic Research (SER) (1988 - present)

PROFESSIONAL EDUCATION

- D.E.S., Université de Fribourg, Fribourg, Switzerland , Secondary Education
- M.A., University of North Carolina at Chapel Hill, Chapel Hill, NC , Geography

- M.S.P.H., University of North Carolina at Chapel Hill, Chapel Hill, NC , Epidemiology
- Ph.D., University of North Carolina at Chapel Hill, Chapel Hill, NC , Epidemiology

LINKS

- Breast Cancer Family Registry: <http://www.bcfamilyregistry.org>
- Northern California Breast Cancer Family Registry: <http://www.med.stanford.edu/ca-bcfr>
- LEGACY Girls Study: <http://www.legacygirlsstudy.org>

Research & Scholarship

CURRENT RESEARCH AND SCHOLARLY INTERESTS

Dr. John has extensive expertise in conducting population-based epidemiologic studies and has led as Principal Investigator multiple large-scale studies, including multi-center studies with a study site in the San Francisco Bay Area with its diverse population. Many of her studies and collaborations investigated cancer health disparities. Her research has focused on the role of modifiable lifestyle factors (e.g., body size, physical activity, diet), hormonal factors, early-life exposures, genetic variants, and gene-environment interactions; differences in risk factors by race/ethnicity, breast cancer subtypes, and prostate cancer subtypes; risk factors for familial breast cancer and second primary breast cancer, as well as prognostic factors related to survival disparities.

As Principal Investigator, Dr. John has led a number of studies conducted in the San Francisco Bay Area, including:

- the Northern California site of the Breast Cancer Family Registry, an on-going prospective multi-generational cohort of over 13,000 families established in 1995 at six international sites;
- the Northern California site of the WECARE Study that investigates risk factors for second primary breast cancer;
- the California site of the Breast Cancer Health Disparities Study that investigated genetic variability and breast cancer risk and survival in Hispanic and non-Hispanic white populations in the context of genetic admixture;
- the Breast Cancer Etiology in Minorities (BEM) Study, a pooled analysis of risk factors for breast cancer subtypes in racial/ethnic minorities;
- the San Francisco Bay Area Breast Cancer Study, a population-based case-control study in nearly 5,000 Hispanic, African American and non-Hispanic white women that investigated the role of modifiable lifestyle factors and other risk factors;
- the San Francisco Bay Area Prostate Cancer Study, a population-based case-control study of lifestyle and genetic risk factors for advanced and localized disease.

These studies collected and pooled extensive data and biospecimens and continue to support numerous ancillary studies, collaborations and international consortia and have contributed to a better understanding of cancer risk and survival in racial/ethnic minority populations.

Dr. John is also a founding PI of the LEGACY Girls Study, an on-going prospective cohort established in 2011 that investigates early life exposures in relation to pubertal development outcomes, breast tissue characteristics, and behavioral and psychosocial outcomes in the context of having a family history or breast cancer.

Publications

PUBLICATIONS

- **Alcohol consumption, cigarette smoking, and risk of breast cancer for BRCA1 and BRCA2 mutation carriers: results from The BRCA1 and BRCA2 Cohort Consortium.** *Cancer epidemiology, biomarkers & prevention : a publication of the American Association for Cancer Research, cosponsored by the American Society of Preventive Oncology*
Li, H., Terry, M. B., Antoniou, A. C., Phillips, K., Kast, K., Mooij, T. M., Engel, C., Nogues, C., Stoppa-Lyonnet, D., Lasset, C., Berthet, P., Mari, V., Caron, et al 2019

- **Recreational physical activity is associated with reduced breast cancer risk in adult women at high risk for breast cancer: a cohort study of women selected for familial and genetic risk.** *Cancer research*
Kehm, R. D., Genkinger, J. M., MacInnis, R. J., John, E. M., Phillips, K., Dite, G. S., Milne, R. L., Zeinomar, N., Liao, Y., Knight, J. A., Southey, M. C., Chung, W. K., Giles, et al
2019
- **A polygenic risk score for breast cancer in U.S. Latinas and Latin-American women.** *Journal of the National Cancer Institute*
Shieh, Y., Fejerman, L., Lott, P. C., Marker, K., Sawyer, S. D., Hu, D., Huntsman, S., Torres, J., Echeverry, M., Bohorquez, M. E., Martinez-Chequer, J. C., Polanco-Echeverry, G., Estrada-Florez, et al
2019
- **Benign breast disease increases breast cancer risk independent of underlying familial risk profile: Findings from a Prospective Family Study Cohort** *INTERNATIONAL JOURNAL OF CANCER*
Zeinomar, N., Phillips, K., Daly, M. B., Milne, R. L., Dite, G. S., MacInnis, R. J., Liao, Y., Kehm, R. D., Knight, J. A., Southey, M. C., Chung, W. K., Giles, G. G., McLachlan, et al
2019; 145 (2): 370–79
- **A polygenic risk score predicts breast cancer risk in Latinas**
Shieh, Y., Fejerman, L., Sawyer, S. D., Hu, D., Huntsman, S., John, E. M., Kushi, L. H., Torres-Mejia, G., Weitzel, J. N., Haiman, C. A., Ziv, E., Neuhausen, S. L.
AMER ASSOC CANCER RESEARCH.2019
- **Runs of homozygosity and testicular cancer risk** *ANDROLOGY*
Loveday, C., Sud, A., Litchfield, K., Levy, M., Holroyd, A., Broderick, P., Kote-Jarai, Z., Dunning, A. M., Muir, K., Peto, J., Eeles, R., Easton, D. F., Dudakia, et al
2019; 7 (4): 555–64
- **The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis.** *International journal of epidemiology*
Ooi, B. N., Loh, H., Ho, P. J., Milne, R. L., Giles, G., Gao, C., Kraft, P., John, E. M., Swerdlow, A., Brenner, H., Wu, A. H., Haiman, C., Evans, et al
2019
- **A Genome-wide Association Study of Prostate Cancer in Latinos.** *International journal of cancer*
Du, Z., Hopp, H., Ingles, S. A., Huff, C., Sheng, X., Weaver, B., Stern, M., Hoffmann, T. J., John, E. M., Van Den Eeden, S. K., Strom, S., Leach, R. J., Thompson, et al
2019
- **Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers.** *British journal of cancer*
Qian, F., Rookus, M. A., Leslie, G., Risch, H. A., Greene, M. H., Aalfs, C. M., Adank, M. A., Adlard, J., Agnarsson, B. A., Ahmed, M., Aittomaki, K., Andrulis, I. L., Arnold, et al
2019
- **The functional ALDH2 polymorphism is associated with breast cancer risk: A pooled analysis from the Breast Cancer Association Consortium** *MOLECULAR GENETICS & GENOMIC MEDICINE*
Ugai, T., Milne, R. L., Ito, H., Aronson, K. J., Bolla, M. K., Chan, T., Chan, C. W., Choi, J., Conroy, D. M., Dennis, J., Dunning, A. M., Easton, D. F., Gaborieau, et al
2019; 7 (6)
- **Genetic predisposition to breast cancer among African American women.**
Palmer, J. R., Hu, C., Hart, S., Gnanaolivu, R., Gao, C., Anton-Culver, H., Trentham-Dietz, A., Bernstein, L., Weitzel, J. N., Domchek, S. M., Goldgar, D., Nathanson, K., Pal, et al
AMER SOC CLINICAL ONCOLOGY.2019
- **Association analyses identify 31 new risk loci for colorectal cancer susceptibility** *NATURE COMMUNICATIONS*
Law, P. J., Timofeeva, M., Fernandez-Rozadilla, C., Timofeeva, A., Broderick, P., Studd, J., Fernandez-Tajes, J., Farrington, S., Svinti, V., Palles, C., Orlando, G., Sud, A., Holroyd, et al
2019; 10: 2154
- **Regular use of aspirin and other non-steroidal anti-inflammatory drugs and breast cancer risk for women at familial or genetic risk: a cohort study** *BREAST CANCER RESEARCH*
Kehm, R. D., Hopper, J. L., John, E. M., Phillips, K., MacInnis, R. J., Dite, G. S., Milne, R. L., Liao, Y., Zeinomar, N., Knight, J. A., Southey, M. C., Vahdat, L., Kornhauser, et al

2019; 21

- **Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer** *NATURE COMMUNICATIONS*
Ferreira, M. A., Gamazon, E. R., Al-Ejeh, F., Aittomaki, K., Andrulis, I. L., Anton-Culver, H., Arason, A., Arndt, V., Aronson, K. J., Arun, B. K., Asseryanis, E., Azzollini, J., Balmana, et al
2019; 10
- **Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer.** *Nature communications*
Ferreira, M. A., Gamazon, E. R., Al-Ejeh, F., Aittomaki, K., Andrulis, I. L., Anton-Culver, H., Arason, A., Arndt, V., Aronson, K. J., Arun, B. K., Asseryanis, E., Azzollini, J., Balmana, et al
2019; 10 (1): 1741
- **10-year performance of four models of breast cancer risk:a validation study** *LANCET ONCOLOGY*
Terry, M., Liao, Y., Whittemore, A. S., Leoce, N., Buchsbaum, R., Zeinotar, N., Dite, G. S., Chung, W. K., Knight, J. A., Southey, M. C., Milne, R. L., Goldgar, D., Giles, et al
2019; 20 (4): 504–17
- **Enrollment and biospecimen collection in a multiethnic family cohort: the Northern California site of the Breast Cancer Family Registry** *CANCER CAUSES & CONTROL*
John, E. M., Sangaramoorthy, M., Koo, J., Whittemore, A. S., West, D. W.
2019; 30 (4): 395–408
- **Genome-wide association study of germline variants and breast cancer-specific mortality** *BRITISH JOURNAL OF CANCER*
Escala-Garcia, M., Guo, Q., Doerk, T., Canisius, S., Keeman, R., Dennis, J., Beesley, J., Lecarpentier, J., Bolla, M. K., Wang, Q., Abraham, J., Andrulis, I. L., Anton-Culver, et al
2019; 120 (6): 647–57
- **Enrollment and biospecimen collection in a multiethnic family cohort: the Northern California site of the Breast Cancer Family Registry.** *Cancer causes & control : CCC*
John, E. M., Sangaramoorthy, M., Koo, J., Whittemore, A. S., West, D. W.
2019
- **Risk-Reducing Oophorectomy and Breast Cancer Risk Across the Spectrum of Familial Risk** *JNCI-JOURNAL OF THE NATIONAL CANCER INSTITUTE*
Terry, M., Daly, M. B., Phillips, K., Ma, X., Zeinomar, N., Leoce, N., Dite, G. S., MacInnis, R. J., Chung, W. K., Knight, J. A., Southey, M. C., Milne, R. L., Goldgar, et al
2019; 111 (3): 331–34
- **Genome-wide association study of germline variants and breast cancer-specific mortality.** *British journal of cancer*
Escala-Garcia, M., Guo, Q., Dork, T., Canisius, S., Keeman, R., Dennis, J., Beesley, J., Lecarpentier, J., Bolla, M. K., Wang, Q., Abraham, J., Andrulis, I. L., Anton-Culver, et al
2019
- **10-year performance of four models of breast cancer risk: a validation study.** *The Lancet. Oncology*
Terry, M. B., Liao, Y., Whittemore, A. S., Leoce, N., Buchsbaum, R., Zeinomar, N., Dite, G. S., Chung, W. K., Knight, J. A., Southey, M. C., Milne, R. L., Goldgar, D., Giles, et al
2019
- **A pooled analysis of breast-feeding and breast cancer risk by hormone receptor status in parous Hispanic women.** *Epidemiology (Cambridge, Mass.)*
Sangaramoorthy, M., Hines, L. M., Torres-Mejia, G., Phipps, A. I., Baumgartner, K. B., Wu, A. H., Koo, J., Ingles, S. A., Slattery, M. L., John, E. M.
2019
- **Benign Breast Disease Increases Breast Cancer Risk Independent of Underlying Familial Risk Profile: Findings from a Prospective Family Study Cohort (ProF-SC).** *International journal of cancer*
Zeinomar, N., Phillips, K., Daly, M. B., Milne, R. L., Dite, G. S., MacInnis, R. J., Liao, Y., Kehm, R. D., Knight, J. A., Southey, M. C., Chung, W. K., Giles, G. G., McLachlan, et al
2019
- **Association of Prepubertal and Adolescent Androgen Concentrations With Timing of Breast Development and Family History of Breast Cancer.** *JAMA network open*
Houghton, L. C., Knight, J. A., Wei, Y., Romeo, R. D., Goldberg, M., Andrulis, I. L., Bradbury, A. R., Buys, S. S., Daly, M. B., John, E. M., Chung, W. K., Santella, R. M., Stanczyk, et al
2019; 2 (2): e190083

- **Association of Prepubertal and Adolescent Androgen Concentrations With Timing of Breast Development and Family History of Breast Cancer** *JAMA NETWORK OPEN*
Houghton, L. C., Knight, J. A., Wei, Y., Romeo, R. D., Goldberg, M., Andrulis, I. L., Bradbury, A. R., Buys, S. S., Daly, M. B., John, E. M., Chung, W. K., Santella, R. M., Stanczyk, et al
2019; 2 (2)
- **Shared heritability and functional enrichment across six solid cancers.** *Nature communications*
Jiang, X., Finucane, H. K., Schumacher, F. R., Schmit, S. L., Tyrer, J. P., Han, Y., Michailidou, K., Lesueur, C., Kuchenbaecker, K. B., Dennis, J., Conti, D. V., Casey, G., Gaudet, et al
2019; 10 (1): 431
- **Genome-wide association study identifies susceptibility loci for B-cell childhood acute lymphoblastic leukemia (vol 9, 1340, 2018)** *NATURE COMMUNICATIONS*
Vijayakrishnan, J., Studd, J., Broderick, P., Kinnersley, B., Holroyd, A., Law, P. J., Kumar, R., Allan, J. M., Harrison, C. J., Moorman, A. V., Vora, A., Roman, E., Rachakonda, et al
2019; 10: 419
- **Germline variation at 8q24 and prostate cancer risk in men of European ancestry (vol 9, 4616, 2018)** *NATURE COMMUNICATIONS*
Matejic, M., Saunders, E. J., Dadaev, T., Brook, M. N., Wang, K., Sheng, X., Al Olama, A., Schumacher, F. R., Ingles, S. A., Govindasami, K., Benlloch, S., Berndt, S. I., Albanes, et al
2019; 10: 382
- **Identification of novel common breast cancer risk variants at the 6q25 locus among Latinas.** *Breast cancer research : BCR*
Hoffman, J., Fejerman, L., Hu, D., Huntsman, S., Li, M., John, E. M., Torres-Mejia, G., Kushi, L., Ding, Y. C., Weitzel, J., Neuhausen, S. L., Lott, P., COLUMBUS Consortium, et al
2019; 21 (1): 3
- **Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma (vol 9, 3707, 2018)** *NATURE COMMUNICATIONS*
Went, M., Sud, A., Foersti, A., Halvarsson, B., Weinhold, N., Kimber, S., van Duin, M., Thorleifsson, G., Holroyd, A., Johnson, D. C., Li, N., Orlando, G., Law, et al
2019; 10: 213
- **Large-scale transcriptome-wide association study identifies new prostate cancer risk regions (vol 9, 4079, 2018)** *NATURE COMMUNICATIONS*
Mancuso, N., Gayther, S., Gusev, A., Zheng, W., Penney, K. L., Kote-Jarai, Z., Eeles, R., Freedman, M., Haiman, C., Pasaniuc, B., Henderson, B. E., Benlloch, S., Schumacher, et al
2019; 10: 171
- **Author Correction: Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci.** *Nature genetics*
Schumacher, F. R., Olama, A. A., Berndt, S. I., Benlloch, S., Ahmed, M., Saunders, E. J., Dadaev, T., Leongamornlert, D., Anokian, E., Cieza-Borrella, C., Goh, C., Brook, M. N., Sheng, et al
2019
- **Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility (vol 8, 1892, 2017)** *NATURE COMMUNICATIONS*
Sud, A., Thomsen, H., Law, P. J., Foersti, A., Filho, M., Holroyd, A., Broderick, P., Orlando, G., Lenive, O., Wright, L., Cooke, R., Easton, D., Pharoah, et al
2019; 10: 157
- **Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes** *AMERICAN JOURNAL OF HUMAN GENETICS*
Mavaddat, N., Michailidou, K., Dennis, J., Lush, M., Fachal, L., Lee, A., Tyrer, J. P., Chen, T., Wang, Q., Bolla, M. K., Yang, X., Adank, M. A., Ahearn, et al
2019; 104 (1): 21–34
- **Identification of novel susceptibility loci and genes for prostate cancer risk: A transcriptome-wide association study in over 140,000 European descendants.** *Cancer research*
Wu, L., Wang, J., Cai, Q., Cavazos, T. B., Emami, N. C., Long, J., Shu, X. O., Lu, Y., Guo, X., Bauer, J. A., Pasaniuc, B., Penney, K. L., Freedman, et al
2019
- **Two truncating variants in FANCC and breast cancer risk.** *Scientific reports*
Dörk, T., Peterlongo, P., Mannermaa, A., Bolla, M. K., Wang, Q., Dennis, J., Ahearn, T., Andrulis, I. L., Anton-Culver, H., Arndt, V., Aronson, K. J., Augustinsson, A., Freeman, et al
2019; 9 (1): 12524

- **The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer.** *NPJ breast cancer*
Figlioli, G., Bogliolo, M., Catucci, I., Caleca, L., Lasheras, S. V., Pujol, R., Kiiski, J. I., Muranen, T. A., Barnes, D. R., Dennis, J., Michailidou, K., Bolla, M. K., Leslie, et al
2019; 5: 38
- **Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer.** *JAMA network open*
Watt, G. P., Reiner, A. S., Smith, S. A., Stram, D. O., Capanu, M., Malone, K. E., Lynch, C. F., John, E. M., Knight, J. A., Mellekjær, L., Bernstein, L., Brooks, J. D., Woods, et al
2019; 2 (9): e1912259
- **Publisher Correction: Shared heritability and functional enrichment across six solid cancers.** *Nature communications*
Jiang, X., Finucane, H. K., Schumacher, F. R., Schmit, S. L., Tyrer, J. P., Han, Y., Michailidou, K., Lesueur, C., Kuchenbaecker, K. B., Dennis, J., Conti, D. V., Casey, G., Gaudet, et al
2019; 10 (1): 4386
- **Association of Genomic Domains in BRCA1 and BRCA2 with Prostate Cancer Risk and Aggressiveness.** *Cancer research*
Patel, V. L., Busch, E. L., Friebel, T. M., Cronin, A., Leslie, G., McGuffog, L., Adlard, J., Agata, S., Agnarsson, B. A., Ahmed, M., Aittomäki, K., Alducci, E., Andrulis, et al
2019
- **Estrogenic activity, race/ethnicity, and Indigenous American ancestry among San Francisco Bay Area women.** *PLoS one*
Sanchez, S. S., Tachachartvanich, P., Stanczyk, F. Z., Gomez, S. L., John, E. M., Smith, M. T., Fejerman, L.
2019; 14 (3): e0213809
- **Considerations when using breast cancer risk models for women with negative BRCA1/BRCA2 mutation results.** *Journal of the National Cancer Institute*
MacInnis, R. J., Liao, Y., Knight, J. A., Milne, R. L., Whittemore, A. S., Chung, W. K., Leco, N., Buchsbaum, R., Zeinomar, N., Dite, G. S., Southey, M. C., Goldgar, D., Giles, et al
2019
- **Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study** *CANCER EPIDEMIOLOGY BIOMARKERS & PREVENTION*
Adams, C. D., Richmond, R., Ferreira, D., Spiller, W., Tan, V., Zheng, J., Wurtz, P., Donovan, J., Hamdy, F., Neal, D., Lane, J., Smith, G., Relton, et al
2019; 28 (1): 208–16
- **Assessing patient readiness for personalized genomic medicine.** *Journal of community genetics*
Frost, C. J., Andrulis, I. L., Buys, S. S., Hopper, J. L., John, E. M., Terry, M. B., Bradbury, A., Chung, W. K., Colbath, K., Quintana, N., Gamarra, E., Egleston, B., Galpern, et al
2019; 10 (1): 109–20
- **The functional ALDH2 polymorphism is associated with breast cancer risk: A pooled analysis from the Breast Cancer Association Consortium.** *Molecular genetics & genomic medicine*
Ugai, T., Milne, R. L., Ito, H., Aronson, K. J., Bolla, M. K., Chan, T., Chan, C. W., Choi, J. Y., Conroy, D. M., Dennis, J., Dunning, A. M., Easton, D. F., Gaborieau, et al
2019: e707
- **A Pooled Analysis of Breastfeeding and Breast Cancer Risk by Hormone Receptor Status in Parous Hispanic Women.** *Epidemiology (Cambridge, Mass.)*
Sangaramoorthy, M., Hines, L. M., Torres-Mejía, G., Phipps, A. I., Baumgartner, K. B., Wu, A. H., Koo, J., Ingles, S. A., Slattery, M. L., John, E. M.
2019; 30 (3): 449–57
- **Regular use of aspirin and other non-steroidal anti-inflammatory drugs and breast cancer risk for women at familial or genetic risk: a cohort study.** *Breast cancer research : BCR*
Kehm, R. D., Hopper, J. L., John, E. M., Phillips, K. A., MacInnis, R. J., Dite, G. S., Milne, R. L., Liao, Y., Zeinomar, N., Knight, J. A., Southey, M. C., Vahdat, L., Kornhauser, et al
2019; 21 (1): 52
- **Alcohol consumption, cigarette smoking, and familial breast cancer risk: findings from the Prospective Family Study Cohort (ProF-SC).** *Breast cancer research : BCR*
Zeinomar, N., Knight, J. A., Genkinger, J. M., Phillips, K. A., Daly, M. B., Milne, R. L., Dite, G. S., Kehm, R. D., Liao, Y., Southey, M. C., Chung, W. K., Giles, G. G., McLachlan, et al
2019; 21 (1): 128
- **Re-evaluating genetic variants identified in candidate gene studies of breast cancer risk using data from nearly 280,000 women of Asian and European ancestry.** *EBioMedicine*

- Yang, Y., Shu, X., Shu, X. O., Bolla, M. K., Kweon, S. S., Cai, Q., Michailidou, K., Wang, Q., Dennis, J., Park, B., Matsuo, K., Kwong, A., Park, et al
2019
- **Race/ethnicity and accuracy of self-reported female first-degree family history of breast and other cancers in the Northern California Breast Cancer Family Registry.** *Cancer epidemiology, biomarkers & prevention : a publication of the American Association for Cancer Research, cosponsored by the American Society of Preventive Oncology*
John, E. M., Canchola, A. J., Sanagaramoorthy, M., Koo, J., Whittemore, A. S., West, D. W.
2019
 - **Surveillance of cancer among sexual and gender minority populations: Where are we and where do we need to go?** *Cancer*
Gomez, S. L., Duffy, C., Griggs, J. J., John, E. M.
2019
 - **CYP2D6 phenotype, tamoxifen, and risk of contralateral breast cancer in the WECARE Study.** *Breast cancer research : BCR*
Brooks, J. D., Comen, E. A., Reiner, A. S., Orlow, I., Leong, S. F., Liang, X., Mellemkjar, L., Knight, J. A., Lynch, C. F., John, E. M., Bernstein, L., Woods, M., Doody, et al
2018; 20 (1): 149
 - **Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes.** *American journal of human genetics*
Mavaddat, N., Michailidou, K., Dennis, J., Lush, M., Fachal, L., Lee, A., Tyrer, J. P., Chen, T., Wang, Q., Bolla, M. K., Yang, X., Adank, M. A., Ahearn, et al
2018
 - **Risk-Reducing Oophorectomy and Breast Cancer Risk Across the Spectrum of Familial Risk.** *Journal of the National Cancer Institute*
Terry, M. B., Daly, M. B., Phillips, K. A., Ma, X., Zeinomar, N., Leoce, N., Dite, G. S., MacInnis, R. J., Chung, W. K., Knight, J. A., Southey, M. C., Milne, R. L., Goldgar, et al
2018
 - **Germline variation at 8q24 and prostate cancer risk in men of European ancestry.** *Nature communications*
Matejic, M., Saunders, E. J., Dadaev, T., Brook, M. N., Wang, K., Sheng, X., Olama, A. A., Schumacher, F. R., Ingles, S. A., Govindasami, K., Benlloch, S., Berndt, S. I., Albanes, et al
2018; 9 (1): 4616
 - **Age-specific breast cancer risk by body mass index and familial risk: prospective family study cohort (ProF-SC).** *Breast cancer research : BCR*
Hopper, J. L., Dite, G. S., MacInnis, R. J., Liao, Y., Zeinomar, N., Knight, J. A., Southey, M. C., Milne, R. L., Chung, W. K., Giles, G. G., Genkinger, J. M., McLachlan, S., Friedlander, et al
2018; 20 (1): 132
 - **Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study.** *Cancer epidemiology, biomarkers & prevention : a publication of the American Association for Cancer Research, cosponsored by the American Society of Preventive Oncology*
Adams, C., Richmond, R. C., Santos Ferreira, D. L., Spiller, W., Tan, V. Y., Zheng, J., Wurtz, P., Donovan, J. L., Hamdy, F. C., Neal, D. E., Lane, J. A., Davey Smith, G., Relton, et al
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