

Mia M Gaudet

List of Publications by Year in descending order

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Version: 2024-02-01

121
papers

12,564
citations

66250

44
h-index

32181

105
g-index

122
all docs

122
docs citations

122
times ranked

18956
citing authors

#	ARTICLE	IF	CITATIONS
1	Prospective evaluation of a breast-cancer risk model integrating classical risk factors and polygenic risk in 15 cohorts from six countries. <i>International Journal of Epidemiology</i> , 2022, 50, 1897-1911.	0.9	43
2	Excess Body Fatness during Early to Mid-Adulthood and Survival from Colorectal and Breast Cancer: A Pooled Analysis of Five International Cohort Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 325-333.	1.1	4
3	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
4	Developing an algorithm across integrated healthcare systems to identify a history of cancer using electronic medical records. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2022, , .	2.2	0
5	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 217-228.	1.1	12
6	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	3.0	45
7	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. <i>International Journal of Cancer</i> , 2021, 148, 307-319.	2.3	35
8	Joint associations of physical activity and body mass index with the risk of established excess body fatness-related cancers among postmenopausal women. <i>Cancer Causes and Control</i> , 2021, 32, 127-138.	0.8	6
9	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	2.9	5
10	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , 2021, 384, 440-451.	13.9	414
11	Association of the Age at Menarche with Site-Specific Cancer Risks in Pooled Data from Nine Cohorts. <i>Cancer Research</i> , 2021, 81, 2246-2255.	0.4	30
12	Risk of Late-Onset Breast Cancer in Genetically Predisposed Women. <i>Journal of Clinical Oncology</i> , 2021, 39, 3430-3440.	0.8	21
13	Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. <i>Human Genetics</i> , 2021, 140, 1353-1365.	1.8	18
14	Risk of Breast Cancer Among Carriers of Pathogenic Variants in Breast Cancer Predisposition Genes Varies by Polygenic Risk Score. <i>Journal of Clinical Oncology</i> , 2021, 39, 2564-2573.	0.8	47
15	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145.	2.9	9
16	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	1.1	19
17	Breast cancer risk factors by mode of detection among screened women in the Cancer Prevention Study-II. <i>Breast Cancer Research and Treatment</i> , 2021, 186, 791-805.	1.1	8
18	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120

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19	Sustained Weight Loss and Risk of Breast Cancer in Women 50 Years and Older: A Pooled Analysis of Prospective Data. <i>Journal of the National Cancer Institute</i> , 2020, 112, 929-937.	3.0	58
20	Epidemiologic risk factors for in situ and invasive ductal breast cancer among regularly screened postmenopausal women by grade in the Cancer Prevention Study-II Nutrition Cohort. <i>Cancer Causes and Control</i> , 2020, 31, 95-103.	0.8	4
21	Ovarian Cancer Risk Factor Associations by Primary Anatomic Site: The Ovarian Cancer Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2010-2018.	1.1	6
22	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581.	9.4	265
23	Contribution of Germline Predisposition Gene Mutations to Breast Cancer Risk in African American Women. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1213-1221.	3.0	51
24	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	1.6	2
25	Erythrocyte levels of cadmium and lead and risk of <sc>B</sc> cell non-Hodgkin lymphoma and multiple myeloma. <i>International Journal of Cancer</i> , 2020, 147, 3110-3118.	2.3	6
26	The Risk of Ovarian Cancer Increases with an Increase in the Lifetime Number of Ovulatory Cycles: An Analysis from the Ovarian Cancer Cohort Consortium (OC3). <i>Cancer Research</i> , 2020, 80, 1210-1218.	0.4	35
27	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
28	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
29	Analgesic Use and Ovarian Cancer Risk: An Analysis in the Ovarian Cancer Cohort Consortium. <i>Journal of the National Cancer Institute</i> , 2019, 111, 137-145.	3.0	43
30	Blood levels of cadmium and lead in relation to breast cancer risk in three prospective cohorts. <i>International Journal of Cancer</i> , 2019, 144, 1010-1016.	2.3	43
31	Mode of detection and breast cancer mortality by follow-up time and tumor characteristics among screened women in Cancer Prevention Study-II. <i>Breast Cancer Research and Treatment</i> , 2019, 177, 679-689.	1.1	12
32	Breast cancer statistics, 2019. <i>Ca-A Cancer Journal for Clinicians</i> , 2019, 69, 438-451.	157.7	2,068
33	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
34	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
35	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
36	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	2.9	52

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37	Ovarian cancer risk factors by tumor aggressiveness: An analysis from the Ovarian Cancer Cohort Consortium. <i>International Journal of Cancer</i> , 2019, 145, 58-69.	2.3	28
38	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	2.6	711
39	Reply to "Mosaic loss of chromosome Y in leukocytes matters". <i>Nature Genetics</i> , 2019, 51, 7-9.	9.4	7
40	Dietary Energy Density, Glycemic Load, Glycemic Index, and Risk for Endometrial Cancer in the CPS-II Nutrition Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 113-115.	1.1	10
41	Obesity, physical activity, and breast cancer survival among older breast cancer survivors in the Cancer Prevention Study-II Nutrition Cohort. <i>Breast Cancer Research and Treatment</i> , 2018, 167, 133-145.	1.1	36
42	A comprehensive analysis of polymorphic variants in steroid hormone and insulin-like growth factor metabolism and risk of <i>in situ</i> breast cancer: Results from the Breast and Prostate Cancer Cohort Consortium. <i>International Journal of Cancer</i> , 2018, 142, 1182-1188.	2.3	0
43	Prevalence of pathogenic/likely pathogenic variants in the 24 cancer genes of the ACMG Secondary Findings v2.0 list in a large cancer cohort and ethnicity-matched controls. <i>Genome Medicine</i> , 2018, 10, 99.	3.6	15
44	Reproducibility of non-fasting plasma metabolomics measurements across processing delays. <i>Metabolomics</i> , 2018, 14, 129.	1.4	16
45	Pooled Analysis of Nine Cohorts Reveals Breast Cancer Risk Factors by Tumor Molecular Subtype. <i>Cancer Research</i> , 2018, 78, 6011-6021.	0.4	67
46	Untargeted Metabolomics Identifies Novel Potential Biomarkers of Habitual Food Intake in a Cross-Sectional Study of Postmenopausal Women. <i>Journal of Nutrition</i> , 2018, 148, 932-943.	1.3	57
47	Serum metabolomic profiles associated with postmenopausal hormone use. <i>Metabolomics</i> , 2018, 14, 97.	1.4	24
48	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
49	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	9.4	184
50	Pooled analysis of active cigarette smoking and invasive breast cancer risk in 14 cohort studies. <i>International Journal of Epidemiology</i> , 2017, 46, dyw288.	0.9	56
51	Associations of parity and age at first pregnancy with overall and cause-specific mortality in the Cancer Prevention Study II. <i>Fertility and Sterility</i> , 2017, 107, 179-188.e6.	0.5	14
52	The American Cancer Society's Cancer Prevention Study 3 (CPS-3): Recruitment, study design, and baseline characteristics. <i>Cancer</i> , 2017, 123, 2014-2024.	2.0	42
53	The relationship between physical activity, obesity, and lung cancer risk by smoking status in a large prospective cohort of US adults. <i>Cancer Causes and Control</i> , 2017, 28, 1357-1368.	0.8	23
54	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099

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55	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
56	Higher-than-expected population prevalence of potentially pathogenic germline <i>TP53</i> variants in individuals unselected for cancer history. <i>Human Mutation</i> , 2017, 38, 1723-1730.	1.1	40
57	Gene-environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. <i>International Journal of Cancer</i> , 2017, 141, 1830-1840.	2.3	20
58	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	45
59	Influence of Fasting Status and Sample Preparation on Metabolic Biomarker Measurements in Postmenopausal Women. <i>PLoS ONE</i> , 2016, 11, e0167832.	1.1	10
60	Evaluation of a Novel Difficulty of Smoking Cessation Phenotype Based on Number of Quit Attempts. <i>Nicotine and Tobacco Research</i> , 2016, 19, ntw234.	1.4	5
61	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. <i>JAMA Oncology</i> , 2016, 2, 1295.	3.4	285
62	Mosaic loss of chromosome Y is associated with common variation near <i>TCL1A</i> . <i>Nature Genetics</i> , 2016, 48, 563-568.	9.4	134
63	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016, 76, 5103-5114.	0.4	100
64	Pre- and postdiagnostic diet in relation to mortality among breast cancer survivors in the CPS-II Nutrition Cohort. <i>Cancer Causes and Control</i> , 2016, 27, 1303-1314.	0.8	40
65	Dietary Energy Density and Postmenopausal Breast Cancer Incidence in the Cancer Prevention Study II Nutrition Cohort. <i>Journal of Nutrition</i> , 2016, 146, 2045-2050.	1.3	16
66	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	5.8	86
67	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
68	Body Mass Index, Waist Circumference, Diabetes, and Risk of Liver Cancer for U.S. Adults. <i>Cancer Research</i> , 2016, 76, 6076-6083.	0.4	119
69	Interactions between breast cancer susceptibility loci and menopausal hormone therapy in relationship to breast cancer in the Breast and Prostate Cancer Cohort Consortium. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 531-540.	1.1	2
70	Ovarian Cancer Risk Factors by Histologic Subtype: An Analysis From the Ovarian Cancer Cohort Consortium. <i>Journal of Clinical Oncology</i> , 2016, 34, 2888-2898.	0.8	349
71	CWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. <i>Human Molecular Genetics</i> , 2016, 25, ddw092.	1.4	19
72	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	2.2	26

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73	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv279.	3.0	152
74	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
75	Anthropometry and head and neck cancer: a pooled analysis of cohort data. <i>International Journal of Epidemiology</i> , 2015, 44, 673-681.	0.9	32
76	Moderate-to-vigorous physical activity and leisure-time sitting in relation to ovarian cancer risk in a large prospective US cohort. <i>Cancer Causes and Control</i> , 2015, 26, 1691-1697.	0.8	33
77	Association of breast cancer risk loci with breast cancer survival. <i>International Journal of Cancer</i> , 2015, 137, 2837-2845.	2.3	33
78	Vitamin D-Associated Genetic Variation and Risk of Breast Cancer in the Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 627-630.	1.1	20
79	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	9.4	513
80	Genetic risk variants associated with in situ breast cancer. <i>Breast Cancer Research</i> , 2015, 17, 82.	2.2	25
81	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	56
82	Plasma carotenoids and breast cancer risk in the Cancer Prevention Study II Nutrition Cohort. <i>Cancer Causes and Control</i> , 2015, 26, 1233-1244.	0.8	24
83	The Authors Reply. <i>American Journal of Epidemiology</i> , 2015, 182, 974-975.	1.6	0
84	Tobacco and Alcohol in Relation to Male Breast Cancer: An Analysis of the Male Breast Cancer Pooling Project Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 520-531.	1.1	19
85	A Genome-Wide Pleiotropy Scan Does Not Identify New Susceptibility Loci for Estrogen Receptor Negative Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e85955.	1.1	8
86	Evidence for an Association of Dietary Flavonoid Intake with Breast Cancer Risk by Estrogen Receptor Status Is Limited. <i>Journal of Nutrition</i> , 2014, 144, 1603-1611.	1.3	29
87	Additive Interactions Between Susceptibility Single-Nucleotide Polymorphisms Identified in Genome-Wide Association Studies and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , 2014, 180, 1018-1027.	1.6	36
88	Post-GWAS gene-environment interplay in breast cancer: results from the Breast and Prostate Cancer Cohort Consortium and a meta-analysis on 79 000 women. <i>Human Molecular Genetics</i> , 2014, 23, 5260-5270.	1.4	37
89	Artificially and Sugar-Sweetened Carbonated Beverage Consumption Is Not Associated with Risk of Lymphoid Neoplasms in Older Men and Women. <i>Journal of Nutrition</i> , 2014, 144, 2041-2049.	1.3	25
90	Establishment of the Cancer Prevention Study II Nutrition Cohort Colorectal Tissue Repository. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2694-2702.	1.1	23

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91	Oophorectomy and Hysterectomy and Cancer Incidence in the Cancer Prevention Study-II Nutrition Cohort. <i>Obstetrics and Gynecology</i> , 2014, 123, 1247-1255.	1.2	41
92	Waist circumference, body mass index, and postmenopausal breast cancer incidence in the Cancer Prevention Study-II Nutrition Cohort. <i>Cancer Causes and Control</i> , 2014, 25, 737-745.	0.8	43
93	Body mass index and risk of head and neck cancer by race: the Carolina Head and Neck Cancer Epidemiology Study. <i>Annals of Epidemiology</i> , 2014, 24, 160-164.e1.	0.9	10
94	Active Smoking and Breast Cancer Risk: Original Cohort Data and Meta-Analysis. <i>Journal of the National Cancer Institute</i> , 2013, 105, 515-525.	3.0	224
95	Type I and II Endometrial Cancers: Have They Different Risk Factors?. <i>Journal of Clinical Oncology</i> , 2013, 31, 2607-2618.	0.8	613
96	Tubal Sterilization and Breast Cancer Incidence: Results From the Cancer Prevention Study II Nutrition Cohort and Meta-Analysis. <i>American Journal of Epidemiology</i> , 2013, 177, 492-499.	1.6	8
97	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
98	Identification of a BRCA2-Specific Modifier Locus at 6p24 Related to Breast Cancer Risk. <i>PLoS Genetics</i> , 2013, 9, e1003173.	1.5	105
99	Evidence of Gene-Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. <i>PLoS Genetics</i> , 2013, 9, e1003284.	1.5	136
100	Recreational Physical Activity and Leisure-Time Sitting in Relation to Postmenopausal Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 1906-1912.	1.1	77
101	A Comparison of the Polytomous Logistic Regression and Joint Cox Proportional Hazards Models for Evaluating Multiple Disease Subtypes in Prospective Cohort Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 275-285.	1.1	61
102	Obesity-related markers and breast cancer in CPS-II Nutrition Cohort. <i>International Journal of Molecular Epidemiology and Genetics</i> , 2013, 4, 156-66.	0.4	21
103	Prospective Studies of Body Mass Index with Head and Neck Cancer Incidence and Mortality. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 497-503.	1.1	51
104	Analysis of Serum Metabolic Profiles in Women with Endometrial Cancer and Controls in a Population-Based Case-Control Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2012, 97, 3216-3223.	1.8	46
105	Learning from disease heterogeneity. <i>Lancet Oncology</i> , The, 2012, 13, 862-863.	5.1	2
106	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012, 21, 5373-5384.	1.4	168
107	Common variants at 12p11, 12q24, 9p21, 9q31.2 and in ZNF365 are associated with breast cancer risk for BRCA1 and/or BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2012, 14, R33.	2.2	78
108	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. <i>Journal of the National Cancer Institute</i> , 2011, 103, 250-263.	3.0	596

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109	Risk factors by molecular subtypes of breast cancer across a population-based study of women 56 years or younger. <i>Breast Cancer Research and Treatment</i> , 2011, 130, 587-597.	1.1	154
110	Associations of common variants at 1p11.2 and 14q24.1 (RAD51L1) with breast cancer risk and heterogeneity by tumor subtype: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 4693-4706.	1.4	71
111	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. <i>Breast Cancer Research</i> , 2010, 12, R110.	2.2	82
112	Do adipokines underlie the association between known risk factors and breast cancer among a cohort of United States women?. <i>Cancer Epidemiology</i> , 2010, 34, 580-586.	0.8	44
113	No Association between <i>FTO</i> or <i>HHEX</i> and Endometrial Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2106-2109.	1.1	24
114	Body mass index and risk of head and neck cancer in a pooled analysis of case-control studies in the International Head and Neck Cancer Epidemiology (INHANCE) Consortium. <i>International Journal of Epidemiology</i> , 2010, 39, 1091-1102.	0.9	89
115	Common Genetic Variants and Modification of Penetrance of BRCA2-Associated Breast Cancer. <i>PLoS Genetics</i> , 2010, 6, e1001183.	1.5	85
116	DNA Hypermethylation of <i>ESR1</i> and <i>PGR</i> in Breast Cancer: Pathologic and Epidemiologic Associations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 3036-3043.	1.1	60
117	Genetic variation in SIPA1 in relation to breast cancer risk and survival after breast cancer diagnosis. <i>International Journal of Cancer</i> , 2009, 124, 1716-1720.	2.3	22
118	<i>HSD17B1</i> Genetic Variants and Hormone Receptor-Defined Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2766-2772.	1.1	11
119	Genetic variation in tumor necrosis factor and lymphotoxin-alpha (TNF-LTA) and breast cancer risk. <i>Human Genetics</i> , 2007, 121, 483-490.	1.8	62
120	Genetic variation of Cytochrome P450 1B1 (CYP1B1) and risk of breast cancer among Polish women. <i>Pharmacogenetics and Genomics</i> , 2006, 16, 547-553.	0.7	23
121	Comprehensive Assessment of Genetic Variation of Catechol-O-Methyltransferase and Breast Cancer Risk. <i>Cancer Research</i> , 2006, 66, 9781-9785.	0.4	21