

Stacey J Winham

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

120
papers

2,668
citations

28
h-index

47
g-index

132
ext. papers

3,626
ext. citations

6.7
avg, IF

4.93
L-index

| # | Paper | IF | Citations |
|-----|---|------|-----------|
| 120 | Automated quantification of levels of breast terminal duct lobular (TDLU) involution using deep learning.. <i>Npj Breast Cancer</i> , 2022 , 8, 13 | 7.8 | 0 |
| 119 | A Genome-Wide Gene-Based Gene-Environment Interaction Study of Breast Cancer in More than 90,000 Women. <i>Cancer Research Communications</i> , 2022 , 2, 211-219 | | 0 |
| 118 | Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci.. <i>Breast Cancer Research</i> , 2022 , 24, 27 | 8.3 | 1 |
| 117 | Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women.. <i>Scientific Reports</i> , 2022 , 12, 6199 | 4.9 | |
| 116 | Quantification of diet quality utilizing the rapid eating assessment for participants-shortened version in bipolar disorder: Implications for prospective depression and cardiometabolic studies.. <i>Journal of Affective Disorders</i> , 2022 , 310, 150-155 | 6.6 | 0 |
| 115 | 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 | 4 | 7 |
| 114 | Understanding Verbosity: Funding Source and the Length of Consent Forms for Cancer Clinical Trials. <i>Journal of Cancer Education</i> , 2021 , 36, 1248-1252 | 1.8 | 1 |
| 113 | Simplified Breast Risk Tool Integrating Questionnaire Risk Factors, Mammographic Density, and Polygenic Risk Score: Development and Validation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 600-607 | 4 | 4 |
| 112 | MCM3 is a novel proliferation marker associated with longer survival for patients with tubo-ovarian high-grade serous carcinoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , | 5.1 | 2 |
| 111 | What about sex?. <i>Nature Metabolism</i> , 2021 , | 14.6 | 2 |
| 110 | DNA Methylation Profiles of Ovarian Clear Cell Carcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , | 4 | 2 |
| 109 | Incorporating Robustness to Imaging Physics into Radiomic Feature Selection for Breast Cancer Risk Estimation. <i>Cancers</i> , 2021 , 13, | 6.6 | 2 |
| 108 | Body mass index and blood pressure in bipolar patients: Target cardiometabolic markers for clinical practice. <i>Journal of Affective Disorders</i> , 2021 , 282, 637-643 | 6.6 | 2 |
| 107 | Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. <i>Human Genetics and Genomics Advances</i> , 2021 , 2, 100042-100042 | 0.8 | 2 |
| 106 | Testing and estimation of X-chromosome SNP effects: Impact of model assumptions. <i>Genetic Epidemiology</i> , 2021 , 45, 577-592 | 2.6 | 3 |
| 105 | Randomized, Double-Blind Trial on the Impact of Word Count in Cancer Clinical Trial Consent Forms. <i>JCO Oncology Practice</i> , 2021 , 17, e1460-e1472 | 2.3 | 0 |
| 104 | Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. <i>Biological Psychiatry</i> , 2021 , 89, 1127-1137 | 7.9 | 12 |

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| 103 | Identification of a Locus Near Associated With Progression-Free Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 1669-1680 | 4 | 2 |
| 102 | Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021 , 108, 1190-1203 | 11 | 1 |
| 101 | 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 | 9.7 | 14 |
| 100 | Population-based targeted sequencing of 54 candidate genes identifies as a susceptibility gene for high-grade serous ovarian cancer. <i>Journal of Medical Genetics</i> , 2021 , 58, 305-313 | 5.8 | 12 |
| 99 | Circulating CD14 HLA-DR monocytic cells as a biomarker for epithelial ovarian cancer progression. <i>American Journal of Reproductive Immunology</i> , 2021 , 85, e13343 | 3.8 | 1 |
| 98 | Association of mammographic density measures and breast cancer "intrinsic" molecular subtypes. <i>Breast Cancer Research and Treatment</i> , 2021 , 187, 215-224 | 4.4 | 3 |
| 97 | Association of Daily Alcohol Intake, Volumetric Breast Density, and Breast Cancer Risk. <i>JNCI Cancer Spectrum</i> , 2021 , 5, pkaa124 | 4.6 | 0 |
| 96 | Somatic mutations in benign breast disease tissues and association with breast cancer risk. <i>BMC Medical Genomics</i> , 2021 , 14, 185 | 3.7 | 0 |
| 95 | 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 | 6.3 | 5 |
| 94 | Mammographic Variation Measures, Breast Density, and Breast Cancer Risk. <i>American Journal of Roentgenology</i> , 2021 , 217, 326-335 | 5.4 | 3 |
| 93 | Deep Learning Predicts Interval and Screening-detected Cancer from Screening Mammograms: A Case-Case-Control Study in 6369 Women. <i>Radiology</i> , 2021 , 301, 550-558 | 20.5 | 2 |
| 92 | Fully Automated Volumetric Breast Density Estimation from Digital Breast Tomosynthesis. <i>Radiology</i> , 2021 , 301, 561-568 | 20.5 | 3 |
| 91 | Deep-LIBRA: An artificial-intelligence method for robust quantification of breast density with independent validation in breast cancer risk assessment. <i>Medical Image Analysis</i> , 2021 , 73, 102138 | 15.4 | 7 |
| 90 | Evaluation of LIBRA Software for Fully Automated Mammographic Density Assessment in Breast Cancer Risk Prediction. <i>Radiology</i> , 2020 , 296, 24-31 | 20.5 | 5 |
| 89 | Development and Validation of the Gene Expression Predictor of High-grade Serous Ovarian Carcinoma Molecular SubTYPE (PrOTYPE). <i>Clinical Cancer Research</i> , 2020 , 26, 5411-5423 | 12.9 | 21 |
| 88 | Clinical and pathological associations of PTEN expression in ovarian cancer: a multicentre study from the Ovarian Tumour Tissue Analysis Consortium. <i>British Journal of Cancer</i> , 2020 , 123, 793-802 | 8.7 | 16 |
| 87 | Potential pharmacogenomic targets in bipolar disorder: considerations for current testing and the development of decision support tools to individualize treatment selection. <i>International Journal of Bipolar Disorders</i> , 2020 , 8, 23 | 5.4 | 4 |
| 86 | fiddle: a tool to combat publication bias by getting research out of the file drawer and into the scientific community. <i>Clinical Science</i> , 2020 , 134, 2729-2739 | 6.5 | 2 |

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| 85 | Cytotoxic T cell depletion with increasing epithelial abnormality in women with benign breast disease. <i>Breast Cancer Research and Treatment</i> , 2020 , 180, 55-61 | 4.4 | 2 |
| 84 | Breast Cancer Risk and Use of Nonsteroidal Anti-inflammatory Agents After a Benign Breast Biopsy. <i>Cancer Prevention Research</i> , 2020 , 13, 967-976 | 3.2 | 3 |
| 83 | Evening chronotype as a discrete clinical subphenotype in bipolar disorder. <i>Journal of Affective Disorders</i> , 2020 , 266, 556-562 | 6.6 | 17 |
| 82 | The impact of binge eating behavior on lithium- and quetiapine-associated changes in body weight, body mass index, and waist circumference during 6 months of treatment: Findings from the bipolar CHOICE study. <i>Journal of Affective Disorders</i> , 2020 , 266, 772-781 | 6.6 | 9 |
| 81 | Bioinformatics and DNA-extraction strategies to reliably detect genetic variants from FFPE breast tissue samples. <i>BMC Genomics</i> , 2019 , 20, 689 | 4.5 | 17 |
| 80 | Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. <i>Breast Cancer Research</i> , 2019 , 21, 68 | 8.3 | 18 |
| 79 | Hyaline fibrous involution of breast lobules: a histologic finding associated with germline BRCA mutation. <i>Modern Pathology</i> , 2019 , 32, 1263-1270 | 9.8 | 1 |
| 78 | Correlations between sex-related hormones, alcohol dependence and alcohol craving. <i>Drug and Alcohol Dependence</i> , 2019 , 197, 183-190 | 4.9 | 5 |
| 77 | Reveal, Don't Conceal: Transforming Data Visualization to Improve Transparency. <i>Circulation</i> , 2019 , 140, 1506-1518 | 16.7 | 35 |
| 76 | Automated volumetric breast density measures: differential change between breasts in women with and without breast cancer. <i>Breast Cancer Research</i> , 2019 , 21, 118 | 8.3 | 7 |
| 75 | Statistical methods for testing X chromosome variant associations: application to sex-specific characteristics of bipolar disorder. <i>Biology of Sex Differences</i> , 2019 , 10, 57 | 9.3 | 3 |
| 74 | Molecular signatures of X chromosome inactivation and associations with clinical outcomes in epithelial ovarian cancer. <i>Human Molecular Genetics</i> , 2019 , 28, 1331-1342 | 5.6 | 11 |
| 73 | Radiomic Phenotypes of Mammographic Parenchymal Complexity: Toward Augmenting Breast Density in Breast Cancer Risk Assessment. <i>Radiology</i> , 2019 , 290, 41-49 | 20.5 | 36 |
| 72 | A comprehensive gene-environment interaction analysis in Ovarian Cancer using genome-wide significant common variants. <i>International Journal of Cancer</i> , 2019 , 144, 2192-2205 | 7.5 | 11 |
| 71 | Sex-specific effects of gain-of-function P2RX7 variation on bipolar disorder. <i>Journal of Affective Disorders</i> , 2019 , 245, 597-601 | 6.6 | 4 |
| 70 | Sex hormones in alcohol consumption: a systematic review of evidence. <i>Addiction Biology</i> , 2019 , 24, 157-169 | 4.69 | 38 |
| 69 | MyD88 and TLR4 Expression in Epithelial Ovarian Cancer. <i>Mayo Clinic Proceedings</i> , 2018 , 93, 307-320 | 6.4 | 14 |
| 68 | Genomic Analysis Using Regularized Regression in High-Grade Serous Ovarian Cancer. <i>Cancer Informatics</i> , 2018 , 17, 1176935118755341 | 2.4 | 3 |

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| 67 | Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. <i>British Journal of Cancer</i> , 2018 , 118, 1123-1129 | 8.7 | 10 |
| 66 | Genetic overlap between endometriosis and endometrial cancer: evidence from cross-disease genetic correlation and GWAS meta-analyses. <i>Cancer Medicine</i> , 2018 , 7, 1978-1987 | 4.8 | 40 |
| 65 | Bipolar disorder with binge eating behavior: a genome-wide association study implicates PRR5-ARHGAP8. <i>Translational Psychiatry</i> , 2018 , 8, 40 | 8.6 | 14 |
| 64 | Mediation analysis of alcohol consumption, DNA methylation, and epithelial ovarian cancer. <i>Journal of Human Genetics</i> , 2018 , 63, 339-348 | 4.3 | 12 |
| 63 | Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study. <i>Journal of Pathology: Clinical Research</i> , 2018 , 4, 250-261 | 5.3 | 38 |
| 62 | Automated and Clinical Breast Imaging Reporting and Data System Density Measures Predict Risk for Screen-Detected and Interval Cancers: A Case-Control Study. <i>Annals of Internal Medicine</i> , 2018 , 168, 757-765 | 8 | 42 |
| 61 | Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. <i>PLoS ONE</i> , 2018 , 13, e0197561 | 3.7 | 9 |
| 60 | Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018 , 9, 3166 | 17.4 | 70 |
| 59 | Macrophagic "Crown-like Structures" Are Associated with an Increased Risk of Breast Cancer in Benign Breast Disease. <i>Cancer Prevention Research</i> , 2018 , 11, 113-119 | 3.2 | 39 |
| 58 | CD56+ immune cell infiltration and MICA are decreased in breast lobules with fibrocystic changes. <i>Breast Cancer Research and Treatment</i> , 2018 , 167, 649-658 | 4.4 | 4 |
| 57 | Model for Predicting Breast Cancer Risk in Women With Atypical Hyperplasia. <i>Journal of Clinical Oncology</i> , 2018 , 36, 1840-1846 | 2.2 | 13 |
| 56 | Why we need to report more than ρ Data were Analyzed by t-tests or ANOVA. <i>ELife</i> , 2018 , 7, | 8.9 | 16 |
| 55 | Transcriptomic Characterization of Endometrioid, Clear Cell, and High-Grade Serous Epithelial Ovarian Carcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 1101-1109 | 4 | 9 |
| 54 | Evaluation of 2 breast cancer risk models in a benign breast disease cohort. <i>Cancer</i> , 2018 , 124, 3319-3328 | 3.4 | 3 |
| 53 | Cigarette smoking is associated with adverse survival among women with ovarian cancer: Results from a pooled analysis of 19 studies. <i>International Journal of Cancer</i> , 2017 , 140, 2422-2435 | 7.5 | 18 |
| 52 | Alterations in the Immune Cell Composition in Premalignant Breast Tissue that Precede Breast Cancer Development. <i>Clinical Cancer Research</i> , 2017 , 23, 3945-3952 | 12.9 | 36 |
| 51 | Mammographic breast density and risk of breast cancer in women with atypical hyperplasia: an observational cohort study from the Mayo Clinic Benign Breast Disease (BBD) cohort. <i>BMC Cancer</i> , 2017 , 17, 84 | 4.8 | 17 |
| 50 | Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017 , 49, 680-691 | 36.3 | 190 |

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| 49 | Breast Cancer Risk and Progressive Histology in Serial Benign Biopsies. <i>Journal of the National Cancer Institute</i> , 2017 , 109, | 9.7 | 9 |
| 48 | Data visualization, bar naked: A free tool for creating interactive graphics. <i>Journal of Biological Chemistry</i> , 2017 , 292, 20592-20598 | 5.4 | 48 |
| 47 | Dose-Response Association of CD8+ Tumor-Infiltrating Lymphocytes and Survival Time in High-Grade Serous Ovarian Cancer. <i>JAMA Oncology</i> , 2017 , 3, e173290 | 13.4 | 152 |
| 46 | Association between mammographic breast density and histologic features of benign breast disease. <i>Breast Cancer Research</i> , 2017 , 19, 134 | 8.3 | 15 |
| 45 | Postlactational involution biomarkers plasminogen and phospho-STAT3 are linked with active age-related lobular involution. <i>Breast Cancer Research and Treatment</i> , 2017 , 166, 133-143 | 4.4 | |
| 44 | NanoString-based breast cancer risk prediction for women with sclerosing adenosis. <i>Breast Cancer Research and Treatment</i> , 2017 , 166, 641-650 | 4.4 | 6 |
| 43 | An integrative approach to assess X-chromosome inactivation using allele-specific expression with applications to epithelial ovarian cancer. <i>Genetic Epidemiology</i> , 2017 , 41, 898-914 | 2.6 | 7 |
| 42 | Characterization of fusion genes in common and rare epithelial ovarian cancer histologic subtypes. <i>Oncotarget</i> , 2017 , 8, 46891-46899 | 3.3 | 17 |
| 41 | Analyses of germline variants associated with ovarian cancer survival identify functional candidates at the 1q22 and 19p12 outcome loci. <i>Oncotarget</i> , 2017 , 8, 64670-64684 | 3.3 | 5 |
| 40 | Breast cancer risk by the extent and type of atypical hyperplasia. <i>Cancer</i> , 2016 , 122, 3087-8 | 6.4 | 5 |
| 39 | Extent of atypical hyperplasia stratifies breast cancer risk in 2 independent cohorts of women. <i>Cancer</i> , 2016 , 122, 2971-8 | 6.4 | 39 |
| 38 | Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 1503-1510 | 4 | 42 |
| 37 | Clinicopathologic features of breast cancers that develop in women with previous benign breast disease. <i>Cancer</i> , 2016 , 122, 378-85 | 6.4 | 25 |
| 36 | Clinical features of bipolar spectrum with binge eating behaviour. <i>Journal of Affective Disorders</i> , 2016 , 201, 95-8 | 6.6 | 26 |
| 35 | Recreational physical inactivity and mortality in women with invasive epithelial ovarian cancer: evidence from the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , 2016 , 115, 95-101 | 8.7 | 28 |
| 34 | Accumulating evidence for a role of TCF7L2 variants in bipolar disorder with elevated body mass index. <i>Bipolar Disorders</i> , 2016 , 18, 124-35 | 3.8 | 20 |
| 33 | Natural history of age-related lobular involution and impact on breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2016 , 155, 423-30 | 4.4 | 23 |
| 32 | Prevalence and correlates of DSM-5 eating disorders in patients with bipolar disorder. <i>Journal of Affective Disorders</i> , 2016 , 191, 216-21 | 6.6 | 49 |

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| 31 | Investigation of Exomic Variants Associated with Overall Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 446-54 | 4 | 6 |
| 30 | Mucocele-like lesions of the breast: a clinical outcome and histologic analysis of 102 cases. <i>Human Pathology</i> , 2016 , 49, 33-8 | 3.7 | 22 |
| 29 | Reinventing Biostatistics Education for Basic Scientists. <i>PLoS Biology</i> , 2016 , 14, e1002430 | 9.7 | 27 |
| 28 | From Static to Interactive: Transforming Data Visualization to Improve Transparency. <i>PLoS Biology</i> , 2016 , 14, e1002484 | 9.7 | 35 |
| 27 | Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. <i>Oncotarget</i> , 2016 , 7, 72381-72394 | 3.3 | 11 |
| 26 | A targeted genetic association study of epithelial ovarian cancer susceptibility. <i>Oncotarget</i> , 2016 , 7, 7381-9 | 3.9 | 7 |
| 25 | Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016 , 45, 884-95 | 7.8 | 45 |
| 24 | Modeling X Chromosome Data Using Random Forests: Conquering Sex Bias. <i>Genetic Epidemiology</i> , 2016 , 40, 123-32 | 2.6 | 4 |
| 23 | Exome genotyping arrays to identify rare and low frequency variants associated with epithelial ovarian cancer risk. <i>Human Molecular Genetics</i> , 2016 , 25, 3600-3612 | 5.6 | 9 |
| 22 | Chronic Recreational Physical Inactivity and Epithelial Ovarian Cancer Risk: Evidence from the Ovarian Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 1114-24 | 4.4 | 27 |
| 21 | Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. <i>Human Genetics</i> , 2016 , 135, 741-56 | 6.3 | 18 |
| 20 | Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016 , 48, 667-674 | 36.3 | 56 |
| 19 | Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016 , 45, 1619-1630 | 7.8 | 77 |
| 18 | Beyond bar and line graphs: time for a new data presentation paradigm. <i>PLoS Biology</i> , 2015 , 13, e1002128 | 9.7 | 367 |
| 17 | Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. <i>Clinical Cancer Research</i> , 2015 , 21, 5264-76 | 12.9 | 24 |
| 16 | Genetics of cardiovascular disease: Importance of sex and ethnicity. <i>Atherosclerosis</i> , 2015 , 241, 219-28 | 3.1 | 68 |
| 15 | Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015 , 5, 17369 | 4.9 | 27 |
| 14 | Associations of prodynorphin sequence variation with alcohol dependence and related traits are phenotype-specific and sex-dependent. <i>Scientific Reports</i> , 2015 , 5, 15670 | 4.9 | 4 |

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| 13 | Sex differences in the risk of rapid cycling and other indicators of adverse illness course in patients with bipolar I and II disorder. <i>Bipolar Disorders</i> , 2015 , 17, 670-6 | 3.8 | 28 |
| 12 | Methylation of leukocyte DNA and ovarian cancer: relationships with disease status and outcome. <i>BMC Medical Genomics</i> , 2014 , 7, 21 | 3.7 | 17 |
| 11 | Genome-wide investigation of regional blood-based DNA methylation adjusted for complete blood counts implicates BNC2 in ovarian cancer. <i>Genetic Epidemiology</i> , 2014 , 38, 457-66 | 2.6 | 16 |
| 10 | Bipolar disorder with comorbid binge eating history: a genome-wide association study implicates APOB. <i>Journal of Affective Disorders</i> , 2014 , 165, 151-8 | 6.6 | 28 |
| 9 | Applications of multifactor dimensionality reduction to genome-wide data using the R package RMDRR. <i>Methods in Molecular Biology</i> , 2013 , 1019, 479-98 | 1.4 | 3 |
| 8 | Epigenome-wide ovarian cancer analysis identifies a methylation profile differentiating clear-cell histology with epigenetic silencing of the HERG K ⁺ channel. <i>Human Molecular Genetics</i> , 2013 , 22, 3038-47 | 5.6 | 49 |
| 7 | Gene-environment interactions in genome-wide association studies: current approaches and new directions. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2013 , 54, 1120-34 | 7.9 | 46 |
| 6 | A Weighted Random Forests Approach to Improve Predictive Performance. <i>Statistical Analysis and Data Mining</i> , 2013 , 6, 496-505 | 1.4 | 49 |
| 5 | SNP interaction detection with Random Forests in high-dimensional genetic data. <i>BMC Bioinformatics</i> , 2012 , 13, 164 | 3.6 | 71 |
| 4 | The effect of retrospective sampling on estimates of prediction error for multifactor dimensionality reduction. <i>Annals of Human Genetics</i> , 2011 , 75, 46-61 | 2.2 | 3 |
| 3 | An R package implementation of multifactor dimensionality reduction. <i>BioData Mining</i> , 2011 , 4, 24 | 4.3 | 18 |
| 2 | A comparison of internal validation techniques for multifactor dimensionality reduction. <i>BMC Bioinformatics</i> , 2010 , 11, 394 | 3.6 | 18 |
| 1 | Polygenic Risk Modelling for Prediction of Epithelial Ovarian Cancer Risk | | 1 |