

# Julian Weldon Adlard

## List of Publications by Citations

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54  
papers

4,330  
citations

25  
h-index

56  
g-index

56  
ext. papers

5,677  
ext. citations

8  
avg, IF

3.74  
L-index

#	Paper	IF	Citations
54	Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. <i>JAMA - Journal of the American Medical Association</i> , <b>2017</b> , 317, 2402-2416	27.4	1140
53	Cancer risks for BRCA1 and BRCA2 mutation carriers: results from prospective analysis of EMBRACE. <i>Journal of the National Cancer Institute</i> , <b>2013</b> , 105, 812-22	9.7	616
52	Association of type and location of BRCA1 and BRCA2 mutations with risk of breast and ovarian cancer. <i>JAMA - Journal of the American Medical Association</i> , <b>2015</b> , 313, 1347-61	27.4	286
51	Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003212	6	209
50	Effect of BRCA Mutations on Metastatic Relapse and Cause-specific Survival After Radical Treatment for Localised Prostate Cancer. <i>European Urology</i> , <b>2015</b> , 68, 186-93	10.2	192
49	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 680-691	36.3	190
48	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 1767-1778	36.3	186
47	Mutational spectrum in a worldwide study of 29,700 families with BRCA1 or BRCA2 mutations. <i>Human Mutation</i> , <b>2018</b> , 39, 593-620	4.7	138
46	Cancer Risks Associated With Germline Pathogenic Variants: An International Study of 524 Families. <i>Journal of Clinical Oncology</i> , <b>2020</b> , 38, 674-685	2.2	133
45	Prediction of Breast and Prostate Cancer Risks in Male BRCA1 and BRCA2 Mutation Carriers Using Polygenic Risk Scores. <i>Journal of Clinical Oncology</i> , <b>2017</b> , 35, 2240-2250	2.2	101
44	Tumour risks and genotype-phenotype correlations associated with germline variants in succinate dehydrogenase subunit genes , and. <i>Journal of Medical Genetics</i> , <b>2018</b> , 55, 384-394	5.8	97
43	Comprehensive Study of the Clinical Phenotype of Germline BAP1 Variant-Carrying Families Worldwide. <i>Journal of the National Cancer Institute</i> , <b>2018</b> , 110, 1328-1341	9.7	97
42	Identification of a BRCA2-specific modifier locus at 6p24 related to breast cancer risk. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003173	6	90
41	Refined histopathological predictors of BRCA1 and BRCA2 mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 3419	8.3	82
40	Common breast cancer susceptibility alleles are associated with tumour subtypes in BRCA1 and BRCA2 mutation carriers: results from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , <b>2011</b> , 13, R110	8.3	62
39	Common alleles at 6q25.1 and 1p11.2 are associated with breast cancer risk for BRCA1 and BRCA2 mutation carriers. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 3304-21	5.6	62
38	Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 15	8.3	58

37	Prostate Cancer Risks for Male BRCA1 and BRCA2 Mutation Carriers: A Prospective Cohort Study. <i>European Urology</i> , <b>2020</b> , 77, 24-35	10.2	53
36	Associations of common breast cancer susceptibility alleles with risk of breast cancer subtypes in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 3416	8.3	46
35	MLH1 -93G>A promoter polymorphism and risk of mismatch repair deficient colorectal cancer. <i>International Journal of Cancer</i> , <b>2008</b> , 123, 2456-9	7.5	43
34	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , <b>2020</b> , 22, 1653-1666	8.1	34
33	DNA glycosylases involved in base excision repair may be associated with cancer risk in BRCA1 and BRCA2 mutation carriers. <i>PLoS Genetics</i> , <b>2014</b> , 10, e1004256	6	33
32	Robust diagnostic genetic testing using solution capture enrichment and a novel variant-filtering interface. <i>Human Mutation</i> , <b>2014</b> , 35, 434-41	4.7	32
31	Comprehensive Cancer-Predisposition Gene Testing in an Adult Multiple Primary Tumor Series Shows a Broad Range of Deleterious Variants and Atypical Tumor Phenotypes. <i>American Journal of Human Genetics</i> , <b>2018</b> , 103, 3-18	11	27
30	Thalidomide in the treatment of cancer. <i>Anti-Cancer Drugs</i> , <b>2000</b> , 11, 787-91	2.4	27
29	Risk-reducing salpingo-oophorectomy, natural menopause, and breast cancer risk: an international prospective cohort of BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2020</b> , 22, 8	8.3	22
28	Association of Genomic Domains in and with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , <b>2020</b> , 80, 624-638	10.1	22
27	Height and Body Mass Index as Modifiers of Breast Cancer Risk in BRCA1/2 Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 350-364	9.7	22
26	Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 4732-47	5.6	21
25	A nonsynonymous polymorphism in IRS1 modifies risk of developing breast and ovarian cancers in BRCA1 and ovarian cancer in BRCA2 mutation carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 1362-70	4	20
24	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 61	8.3	16
23	Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. <i>Breast Cancer Research and Treatment</i> , <b>2017</b> , 161, 117-134	4.4	15
22	Effects of BRCA2 cis-regulation in normal breast and cancer risk amongst BRCA2 mutation carriers. <i>Breast Cancer Research</i> , <b>2012</b> , 14, R63	8.3	15
21	Cas9-based enrichment and single-molecule sequencing for precise characterization of genomic duplications. <i>Laboratory Investigation</i> , <b>2020</b> , 100, 135-146	5.9	15
20	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , <b>2019</b> , 121, 180-192	8.7	13

19	Oral Contraceptive Use and Breast Cancer Risk: Retrospective and Prospective Analyses From a BRCA1 and BRCA2 Mutation Carrier Cohort Study. <i>JNCI Cancer Spectrum</i> , <b>2018</b> , 2, pky023	4.6	13
18	Risks of breast or ovarian cancer in BRCA1 or BRCA2 predictive test negatives: findings from the EMBRACE study. <i>Genetics in Medicine</i> , <b>2018</b> , 20, 1575-1582	8.1	12
17	The :p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , <b>2019</b> , 5, 38	7.8	12
16	Lymphocyte telomere length is long in BRCA1 and BRCA2 mutation carriers regardless of cancer-affected status. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 1018-24	4	12
15	Increasing Evidence for the Association of Breast Implant-Associated Anaplastic Large Cell Lymphoma and Li Fraumeni Syndrome. <i>Case Reports in Genetics</i> , <b>2019</b> , 2019, 5647940	0.7	10
14	Additional loss of MSH2 and MSH6 expression in sporadic deficient mismatch repair colorectal cancer due to MLH1 promoter hypermethylation. <i>Journal of Clinical Pathology</i> , <b>2019</b> , 72, 443-447	3.9	9
13	Oral contraceptive use and ovarian cancer risk for BRCA1/2 mutation carriers: an international cohort study. <i>American Journal of Obstetrics and Gynecology</i> , <b>2021</b> , 225, 51.e1-51.e17	6.4	9
12	Histopathology of melanocytic lesions in a family with an inherited BAP1 mutation. <i>Journal of Cutaneous Pathology</i> , <b>2016</b> , 43, 287-9	1.7	8
11	Fine-Scale Mapping at 9p22.2 Identifies Candidate Causal Variants That Modify Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>PLoS ONE</i> , <b>2016</b> , 11, e0158801	3.7	7
10	Characterization and Genomic Localization of a SMAD4 Processed Pseudogene. <i>Journal of Molecular Diagnostics</i> , <b>2017</b> , 19, 933-940	5.1	4
9	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , <b>2021</b> , 12, 1078	17.4	4
8	Increased Sensitivity of Diagnostic Mutation Detection by Re-analysis Incorporating Local Reassembly of Sequence Reads. <i>Molecular Diagnosis and Therapy</i> , <b>2017</b> , 21, 685-692	4.5	3
7	Breast and Prostate Cancer Risks for Male BRCA1 and BRCA2 Pathogenic Variant Carriers Using Polygenic Risk Scores. <i>Journal of the National Cancer Institute</i> , <b>2021</b> ,	9.7	3
6	Prostate Cancer Risk by BRCA2 Genomic Regions. <i>European Urology</i> , <b>2020</b> , 78, 494-497	10.2	2
5	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous BRCA1 or BRCA2 pathogenic variant. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 1726-1737	8.1	2
4	SDHC pheochromocytoma and paraganglioma: A UK-wide case series. <i>Clinical Endocrinology</i> , <b>2021</b> ,	3.4	2
3	Multiple primary cancers (renal papillary, lymphoma and teratoma) and hepatic cysts in association with a pathogenic germline mutation in the MET gene. <i>Familial Cancer</i> , <b>2021</b> , 20, 81-83	3	1
2	Long-read nanopore sequencing enables accurate confirmation of a recurrent PMS2 insertion-deletion variant located in a region of complex genomic architecture. <i>Cancer Genetics</i> , <b>2021</b> , 256-257, 122-126	2.3	1

- 1 Homozygosity for the pathogenic RET hotspot variant p.Cys634Trp: A consanguineous family with MEN2A. *European Journal of Medical Genetics*, **2021**, 64, 104141 2.6