

Stuart MacGregor

List of Publications by Year in descending order

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Version: 2024-02-01

283
papers

23,587
citations

13068

68
h-index

9839

141
g-index

317
all docs

317
docs citations

317
times ranked

27762
citing authors

#	ARTICLE	IF	CITATIONS
1	High Polygenic Risk Is Associated with Earlier Trabeculectomy in Patients with Primary Open-Angle Glaucoma. <i>Ophthalmology Glaucoma</i> , 2023, 6, 54-57.	0.9	1
2	Methotrexate-related central neurotoxicity: clinical characteristics, risk factors and genome-wide association study in children treated for acute lymphoblastic leukemia. <i>Haematologica</i> , 2022, 107, 635-643.	1.7	16
3	Normal-tension glaucoma is associated with cognitive impairment. <i>British Journal of Ophthalmology</i> , 2022, 106, 952-956.	2.1	14
4	Multitrait genetic association analysis identifies 50 new risk loci for gastro-oesophageal reflux, seven new loci for Barrett's oesophagus and provides insights into clinical heterogeneity in reflux diagnosis. <i>Gut</i> , 2022, 71, 1053-1061.	6.1	74
5	Examining Evidence for a Causal Association between Telomere Length and Nevus Count. <i>Journal of Investigative Dermatology</i> , 2022, 142, 1502-1505.e6.	0.3	0
6	Genetic overlap analysis of endometriosis and asthma identifies shared loci implicating sex hormones and thyroid signalling pathways. <i>Human Reproduction</i> , 2022, 37, 366-383.	0.4	19
7	Attitudes Towards Polygenic Risk Testing in Individuals with Glaucoma. <i>Ophthalmology Glaucoma</i> , 2022, 5, 436-446.	0.9	10
8	Multi-Trait Genetic Analysis Identifies Autoimmune Loci Associated with Cutaneous Melanoma. <i>Journal of Investigative Dermatology</i> , 2022, 142, 1607-1616.	0.3	11
9	Is Genetic Risk for Sleep Apnea Causally Linked With Glaucoma Susceptibility?. , 2022, 63, 25.		3
10	Genetic Risk of Cardiovascular Disease Is Associated with Macular Ganglion Cell's Inner Plexiform Layer Thinning in an Early Glaucoma Cohort. <i>Ophthalmology Science</i> , 2022, 2, 100108.	1.0	1
11	Acute central nervous system toxicity during treatment of pediatric acute lymphoblastic leukemia: phenotypes, risk factors and genotypes. <i>Haematologica</i> , 2022, 107, 2318-2328.	1.7	3
12	Evaluating a Causal Relationship between Complement Factor I Protein Level and Advanced Age-Related Macular Degeneration Using Mendelian Randomization. <i>Ophthalmology Science</i> , 2022, 2, 100146.	1.0	6
13	The APOE E4 Allele Is Associated with Faster Rates of Neuroretinal Thinning in a Prospective Cohort Study of Suspect and Early Glaucoma. <i>Ophthalmology Science</i> , 2022, 2, 100159.	1.0	4
14	Association of Novel Loci With Keratoconus Susceptibility in a Multitrait Genome-Wide Association Study of the UK Biobank Database and Canadian Longitudinal Study on Aging. <i>JAMA Ophthalmology</i> , 2022, 140, 568.	1.4	5
15	The effect of screening on melanoma incidence and biopsy rates. <i>British Journal of Dermatology</i> , 2022, 187, 515-522.	1.4	22
16	Retinal ganglion cell-specific genetic regulation in primary open-angle glaucoma. <i>Cell Genomics</i> , 2022, 2, 100142.	3.0	9
17	A Polygenic Risk Score Predicts Intraocular Pressure Readings Outside Office Hours and Early Morning Spikes as Measured by Home Tonometry. <i>Ophthalmology Glaucoma</i> , 2021, 4, 411-420.	0.9	11
18	Polygenic Risk Scores Allow Risk Stratification for Keratinocyte Cancer in Organ-Transplant Recipients. <i>Journal of Investigative Dermatology</i> , 2021, 141, 325-333.e6.	0.3	8

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19	Cardiovascular Disease Predicts Structural and Functional Progression in Early Glaucoma. <i>Ophthalmology</i> , 2021, 128, 58-69.	2.5	24
20	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using population-based cohort studies and two-sample Mendelian randomization analyses. <i>International Journal of Cancer</i> , 2021, 148, 1338-1350.	2.3	9
21	The effects of eight serum lipid biomarkers on age-related macular degeneration risk: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2021, 50, 325-336.	0.9	25
22	Germline variation in the insulin-like growth factor pathway and risk of Barrett's esophagus and esophageal adenocarcinoma. <i>Carcinogenesis</i> , 2021, 42, 369-377.	1.3	11
23	Genetic analysis of endometriosis and depression identifies shared loci and implicates causal links with gastric mucosa abnormality. <i>Human Genetics</i> , 2021, 140, 529-552.	1.8	36
24	A comprehensive re-assessment of the association between vitamin D and cancer susceptibility using Mendelian randomization. <i>Nature Communications</i> , 2021, 12, 246.	5.8	39
25	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. <i>Nature Communications</i> , 2021, 12, 1258.	5.8	196
26	Predicting the Future of Genetic Risk Profiling of Glaucoma. <i>JAMA Ophthalmology</i> , 2021, 139, 224.	1.4	15
27	A multi-ethnic genome-wide association study implicates collagen matrix integrity and cell differentiation pathways in keratoconus. <i>Communications Biology</i> , 2021, 4, 266.	2.0	36
28	Time spent outdoors in childhood is associated with reduced risk of myopia as an adult. <i>Scientific Reports</i> , 2021, 11, 6337.	1.6	34
29	IMI 2021 Yearly Digest. , 2021, 62, 7.		36
30	Symptom-level modelling unravels the shared genetic architecture of anxiety and depression. <i>Nature Human Behaviour</i> , 2021, 5, 1432-1442.	6.2	45
31	Genetic variation affects morphological retinal phenotypes extracted from UK Biobank optical coherence tomography images. <i>PLoS Genetics</i> , 2021, 17, e1009497.	1.5	50
32	Associations of sleep apnoea with glaucoma and age-related macular degeneration: an analysis in the United Kingdom Biobank and the Canadian Longitudinal Study on Aging. <i>BMC Medicine</i> , 2021, 19, 104.	2.3	19
33	Polyunsaturated Fatty Acid Levels and the Risk of Keratinocyte Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 1591-1598.	1.1	10
34	Polygenic Risk Scores Stratify Keratinocyte Cancer Risk among Solid Organ Transplant Recipients with Chronic Immunosuppression in a High Ultraviolet Radiation Environment. <i>Journal of Investigative Dermatology</i> , 2021, 141, 2866-2875.e2.	0.3	4
35	Identification of a Locus Near <i>ULK1</i> Associated With Progression-Free Survival in Ovarian Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 1669-1680.	1.1	5
36	Evaluation of Shared Genetic Susceptibility to High and Low Myopia and Hyperopia. <i>JAMA Ophthalmology</i> , 2021, 139, 601.	1.4	22

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37	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
38	Automated AI labeling of optic nerve head enables insights into cross-ancestry glaucoma risk and genetic discovery in >280,000 images from UKB and CLSA. <i>American Journal of Human Genetics</i> , 2021, 108, 1204-1216.	2.6	39
39	Large-scale cross-cancer fine-mapping of the 5p15.33 region reveals multiple independent signals. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100041.	1.0	6
40	Genetic Relationship Between Endometriosis and Melanoma. <i>Frontiers in Reproductive Health</i> , 2021, 3, .	0.6	5
41	649Personal history of keratinocyte carcinoma is a marker of inherited cancer risk: Mendelian randomization analyses. <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	0
42	Association of Monogenic and Polygenic Risk With the Prevalence of Open-Angle Glaucoma. <i>JAMA Ophthalmology</i> , 2021, 139, 1023.	1.4	15
43	Assessing the genetic relationship between gastro-esophageal reflux disease and risk of COVID-19 infection. <i>Human Molecular Genetics</i> , 2021, , .	1.4	7
44	Characteristics of p.Gln368Ter Myocilin Variant and Influence of Polygenic Risk on Glaucoma Penetrance in the UK Biobank. <i>Ophthalmology</i> , 2021, 128, 1300-1311.	2.5	27
45	Coffee consumption and risk of breast cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2021, 16, e0236904.	1.1	9
46	Genetically determined risk of keratinocyte carcinoma and risk of other cancers. <i>International Journal of Epidemiology</i> , 2021, 50, 1316-1324.	0.9	1
47	Germline variants are associated with increased primary melanoma tumor thickness at diagnosis. <i>Human Molecular Genetics</i> , 2021, 29, 3578-3587.	1.4	3
48	Genetically determined cutaneous nevi and risk of cancer. <i>International Journal of Cancer</i> , 2021, , .	2.3	1
49	Is there a causal relationship between vitamin D and melanoma risk? A Mendelian randomization study. <i>British Journal of Dermatology</i> , 2020, 182, 97-103.	1.4	18
50	Potential influence of socioeconomic status on genetic correlations between alcohol consumption measures and mental health. <i>Psychological Medicine</i> , 2020, 50, 484-498.	2.7	44
51	Genetic heterogeneity in self-reported depressive symptoms identified through genetic analyses of the PHQ-9. <i>Psychological Medicine</i> , 2020, 50, 2385-2396.	2.7	46
52	An Intraocular Pressure Polygenic Risk Score Stratifies Multiple Primary Open-Angle Glaucoma Parameters Including Treatment Intensity. <i>Ophthalmology</i> , 2020, 127, 901-907.	2.5	37
53	Using Mendelian randomization to evaluate the causal relationship between serum C-reactive protein levels and age-related macular degeneration. <i>European Journal of Epidemiology</i> , 2020, 35, 139-146.	2.5	66
54	Association of Genetic Variation With Keratoconus. <i>JAMA Ophthalmology</i> , 2020, 138, 174.	1.4	34

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55	Does polygenic risk influence associations between sun exposure and melanoma? A prospective cohort analysis. <i>British Journal of Dermatology</i> , 2020, 183, 303-310.	1.4	13
56	Overlapping genetic architecture between Parkinson disease and melanoma. <i>Acta Neuropathologica</i> , 2020, 139, 347-364.	3.9	23
57	Can vitamin D levels affect your risk of melanoma?. <i>British Journal of Dermatology</i> , 2020, 182, e19-e19.	1.4	0
58	Multiplex melanoma families are enriched for polygenic risk. <i>Human Molecular Genetics</i> , 2020, 29, 2976-2985.	1.4	9
59	Gene Discovery Using Twins. <i>Twin Research and Human Genetics</i> , 2020, 23, 90-93.	0.3	0
60	Association of Myopia and Intraocular Pressure With Retinal Detachment in European Descent Participants of the UK Biobank Cohort. <i>JAMA Ophthalmology</i> , 2020, 138, 671.	1.4	23
61	Genome-wide association meta-analysis of corneal curvature identifies novel loci and shared genetic influences across axial length and refractive error. <i>Communications Biology</i> , 2020, 3, 133.	2.0	22
62	Meta-analysis of 542,934 subjects of European ancestry identifies new genes and mechanisms predisposing to refractive error and myopia. <i>Nature Genetics</i> , 2020, 52, 401-407.	9.4	180
63	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020, 11, 3353.	5.8	75
64	Body mass index and height and risk of cutaneous melanoma: Mendelian randomization analyses. <i>International Journal of Epidemiology</i> , 2020, 49, 1236-1245.	0.9	21
65	Multitrait analysis of glaucoma identifies new risk loci and enables polygenic prediction of disease susceptibility and progression. <i>Nature Genetics</i> , 2020, 52, 160-166.	9.4	192
66	Investigating the genetic and causal relationship between initiation or use of alcohol, caffeine, cannabis and nicotine. <i>Drug and Alcohol Dependence</i> , 2020, 210, 107966.	1.6	12
67	Genome-wide meta-analysis identifies novel loci associated with age-related macular degeneration. <i>Journal of Human Genetics</i> , 2020, 65, 657-665.	1.1	59
68	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , 2020, 52, 494-504.	9.4	138
69	Rationale and protocol for the 7- and 8-year longitudinal assessments of eye health in a cohort of young adults in the Raine Study. <i>BMJ Open</i> , 2020, 10, e033440.	0.8	5
70	The Genetics of Myopia. , 2020, , 95-132.		10
71	Genome-Wide Association Meta-Analysis of Single-Nucleotide Polymorphisms and Symptomatic Venous Thromboembolism during Therapy for Acute Lymphoblastic Leukemia and Lymphoma in Caucasian Children. <i>Cancers</i> , 2020, 12, 1285.	1.7	5
72	Abstract 30: Cross-cancer cross-tissue transcriptome-wide association study (TWAS) of 11 cancers identifies 56 novel genes. , 2020, , .		0

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73	Abstract 1194: Cross-cancer GWAS meta-analysis of more than 370,000 cases and 530,000 controls identifies multiple novel cancer risk regions. , 2020, , .		0
74	Inherited Contributions to Melanoma Risk. , 2019, , 225-248.		0
75	Association between coffee consumption and overall risk of being diagnosed with or dying from cancer among >300 000 UK Biobank participants in a large-scale Mendelian randomization study. International Journal of Epidemiology, 2019, 48, 1447-1456.	0.9	29
76	Determining Possible Shared Genetic Architecture Between Myopia and Primary Open-Angle Glaucoma. , 2019, 60, 3142.		10
77	Genome-wide association analysis of 95%549 individuals identifies novel loci and genes influencing optic disc morphology. Human Molecular Genetics, 2019, 28, 3680-3690.	1.4	19
78	Gastroesophageal reflux GWAS identifies risk loci that also associate with subsequent severe esophageal diseases. Nature Communications, 2019, 10, 4219.	5.8	58
79	Genetic Correlations Between Diabetes and Glaucoma: An Analysis of Continuous and Dichotomous Phenotypes. American Journal of Ophthalmology, 2019, 206, 245-255.	1.7	12
80	Combined analysis of keratinocyte cancers identifies novel genome-wide loci. Human Molecular Genetics, 2019, 28, 3148-3160.	1.4	46
81	Exome-Derived Adiponectin-Associated Variants Implicate Obesity and Lipid Biology. American Journal of Human Genetics, 2019, 105, 15-28.	2.6	21
82	Assessment of melanoma candidate genes in a meta-analysis of 16534 melanoma cases. Journal of the European Academy of Dermatology and Venereology, 2019, 33, e369-e370.	1.3	0
83	New insight into human sweet taste: a genome-wide association study of the perception and intake of sweet substances. American Journal of Clinical Nutrition, 2019, 109, 1724-1737.	2.2	53
84	Implementing MR-PRESSO and GCTA-CGSMR for pleiotropy assessment in Mendelian randomization studies from a practitioner's perspective. Genetic Epidemiology, 2019, 43, 609-616.	0.6	126
85	Risk factors for symptomatic venous thromboembolism during therapy for childhood acute lymphoblastic leukemia. Thrombosis Research, 2019, 178, 132-138.	0.8	16
86	Mendelian Randomization Study for Genetically Predicted Polyunsaturated Fatty Acids Levels on Overall Cancer Risk and Mortality. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 1015-1023.	1.1	19
87	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. Psychological Medicine, 2019, 49, 1218-1226.	2.7	74
88	No Association Between Vitamin D Status and Risk of Barrett's Esophagus or Esophageal Adenocarcinoma: A Mendelian Randomization Study. Clinical Gastroenterology and Hepatology, 2019, 17, 2227-2235.e1.	2.4	16
89	Effect of increased body mass index on risk of diagnosis or death from cancer. British Journal of Cancer, 2019, 120, 565-570.	2.9	20
90	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. Nature Genetics, 2019, 51, 452-469.	9.4	89

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91	Multi-trait genome-wide association study identifies new loci associated with optic disc parameters. <i>Communications Biology</i> , 2019, 2, 435.	2.0	22
92	Myocilin Gene Gln368Ter Variant Penetrance and Association With Glaucoma in Population-Based and Registry-Based Studies. <i>JAMA Ophthalmology</i> , 2019, 137, 28.	1.4	32
93	Abstract 1592: Genome-wide meta-analysis of keratinocytic cancers identifies 26 novel risk loci. , 2019, , .		0
94	Abstract 1588: Germline variation in DNA repair genes and risk of Barrett's esophagus and esophageal adenocarcinoma. , 2019, , .		0
95	Polyunsaturated fatty acids and risk of melanoma: A Mendelian randomisation analysis. <i>International Journal of Cancer</i> , 2018, 143, 508-514.	2.3	18
96	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.	1.3	62
97	Genome-wide association study identifies seven novel susceptibility loci for primary open-angle glaucoma. <i>Human Molecular Genetics</i> , 2018, 27, 1486-1496.	1.4	111
98	Analysis combining correlated glaucoma traits identifies five new risk loci for open-angle glaucoma. <i>Scientific Reports</i> , 2018, 8, 3124.	1.6	33
99	Genome-wide association study of paclitaxel and carboplatin disposition in women with epithelial ovarian cancer. <i>Scientific Reports</i> , 2018, 8, 1508.	1.6	3
100	Assessment of moderate coffee consumption and risk of epithelial ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2018, 47, 450-459.	0.9	15
101	Height and overall cancer risk and mortality: evidence from a Mendelian randomisation study on 310,000 UK Biobank participants. <i>British Journal of Cancer</i> , 2018, 118, 1262-1267.	2.9	46
102	Interactions Between Genetic Variants and Environmental Factors Affect Risk of Esophageal Adenocarcinoma and Barrett's Esophagus. <i>Clinical Gastroenterology and Hepatology</i> , 2018, 16, 1598-1606.e4.	2.4	16
103	Novel pleiotropic risk loci for melanoma and nevus density implicate multiple biological pathways. <i>Nature Communications</i> , 2018, 9, 4774.	5.8	87
104	Understanding the role of bitter taste perception in coffee, tea and alcohol consumption through Mendelian randomization. <i>Scientific Reports</i> , 2018, 8, 16414.	1.6	36
105	Cell-type-specific eQTL of primary melanocytes facilitates identification of melanoma susceptibility genes. <i>Genome Research</i> , 2018, 28, 1621-1635.	2.4	67
106	Genome-wide association meta-analysis highlights light-induced signaling as a driver for refractive error. <i>Nature Genetics</i> , 2018, 50, 834-848.	9.4	239
107	Cross-ancestry genome-wide association analysis of corneal thickness strengthens link between complex and Mendelian eye diseases. <i>Nature Communications</i> , 2018, 9, 1864.	5.8	63
108	Association Between Population Density and Genetic Risk for Schizophrenia. <i>JAMA Psychiatry</i> , 2018, 75, 901.	6.0	67

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109	Accuracy of Inferred APOE Genotypes for a Range of Genotyping Arrays and Imputation Reference Panels. <i>Journal of Alzheimer's Disease</i> , 2018, 64, 49-54.	1.2	9
110	Combining common genetic variants and non-genetic risk factors to predict risk of cutaneous melanoma. <i>Human Molecular Genetics</i> , 2018, 27, 4145-4156.	1.4	34
111	Genome-wide association study of intraocular pressure uncovers new pathways to glaucoma. <i>Nature Genetics</i> , 2018, 50, 1067-1071.	9.4	152
112	Family-Based Genome-Wide Association Study of South Indian Pedigrees Supports <i>WNT7B</i> as a Central Corneal Thickness Locus. , 2018, 59, 2495.		11
113	Genome-Wide Association Study Identifies a Susceptibility Locus for Comitant Esotropia and Suggests a Parent-of-Origin Effect. , 2018, 59, 4054.		21
114	GWAS of lifetime cannabis use reveals new risk loci, genetic overlap with psychiatric traits, and a causal effect of schizophrenia liability. <i>Nature Neuroscience</i> , 2018, 21, 1161-1170.	7.1	436
115	Vitamin D and overall cancer risk and cancer mortality: a Mendelian randomization study. <i>Human Molecular Genetics</i> , 2018, 27, 4315-4322.	1.4	49
116	Assessing the Incremental Contribution of Common Genomic Variants to Melanoma Risk Prediction in Two Population-Based Studies. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2617-2624.	0.3	52
117	Testosterone Pathway Genetic Polymorphisms in Relation to Primary Open-Angle Glaucoma: An Analysis in Two Large Datasets. , 2018, 59, 629.		14
118	Genomic locus modulating corneal thickness in the mouse identifies <i>POU6F2</i> as a potential risk of developing glaucoma. <i>PLoS Genetics</i> , 2018, 14, e1007145.	1.5	31
119	Inherited Contributions to Melanoma Risk. , 2018, , 1-23.		1
120	Abstract 234: Understanding melanoma susceptibility through GWAS of risk phenotypes. , 2018, , .		0
121	New insights into the genetics of primary open-angle glaucoma based on meta-analyses of intraocular pressure and optic disc characteristics.. <i>Human Molecular Genetics</i> , 2017, 26, ddw399.	1.4	120
122	A Novel Approach for Pathway Analysis of GWAS Data Highlights Role of BMP Signaling and Muscle Cell Differentiation in Colorectal Cancer Susceptibility. <i>Twin Research and Human Genetics</i> , 2017, 20, 1-9.	0.3	36
123	Haplotype reference consortium panel: Practical implications of imputations with large reference panels. <i>Human Mutation</i> , 2017, 38, 1025-1032.	1.1	43
124	Genome-Wide Association Shows that Pigmentation Genes Play a Role in Skin Aging. <i>Journal of Investigative Dermatology</i> , 2017, 137, 1887-1894.	0.3	48
125	Genetically low vitamin D concentrations and myopic refractive error: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2017, 46, 1882-1890.	0.9	47
126	Meta-analysis identifies five novel loci associated with endometriosis highlighting key genes involved in hormone metabolism. <i>Nature Communications</i> , 2017, 8, 15539.	5.8	230

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127	Germline variation in inflammation-related pathways and risk of Barrett's oesophagus and oesophageal adenocarcinoma. <i>Gut</i> , 2017, 66, 1739-1747.	6.1	38
128	A common intronic variant of PARP1 confers melanoma risk and mediates melanocyte growth via regulation of MITF. <i>Nature Genetics</i> , 2017, 49, 1326-1335.	9.4	51
129	Contribution of Mutations in Known Mendelian Glaucoma Genes to Advanced Early-Onset Primary Open-Angle Glaucoma. , 2017, 58, 1537.		13
130	Whole exome sequencing implicates eye development, the unfolded protein response and plasma membrane homeostasis in primary open-angle glaucoma. <i>PLoS ONE</i> , 2017, 12, e0172427.	1.1	8
131	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.	0.8	7
132	Association of Polymorphisms in MACRO Domain Containing 2 With Thyroid-Associated Orbitopathy. , 2016, 57, 3129.		12
133	Author Response: Stronger Association of CDKN2B-AS1 Variants in Female Normal-Tension Glaucoma Patients in a Japanese Population. , 2016, 57, 6418.		0
134	Genetic Association at the 9p21 Glaucoma Locus Contributes to Sex Bias in Normal-Tension Glaucoma. , 2016, 57, 3416.		26
135	Polymorphisms in genes in the androgen pathway and risk of Barrett's esophagus and esophageal adenocarcinoma. <i>International Journal of Cancer</i> , 2016, 138, 1146-1152.	2.3	10
136	Rare variants in optic disc area gene <i>CARD10</i> enriched in primary open-angle glaucoma. <i>Molecular Genetics & Genomic Medicine</i> , 2016, 4, 624-633.	0.6	14
137	GWAS study using DNA pooling strategy identifies association of variant rs4910623 in OR52B4 gene with anti-VEGF treatment response in age-related macular degeneration. <i>Scientific Reports</i> , 2016, 6, 37924.	1.6	23
138	Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. <i>Human Genetics</i> , 2016, 135, 741-756.	1.8	19
139	When do myopia genes have their effect? Comparison of genetic risks between children and adults. <i>Genetic Epidemiology</i> , 2016, 40, 756-766.	0.6	34
140	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 1619-1630.	0.9	111
141	Sweet Taste Perception is Associated with Body Mass Index at the Phenotypic and Genotypic Level. <i>Twin Research and Human Genetics</i> , 2016, 19, 465-471.	0.3	13
142	Assessing the Genetic Predisposition of Education on Myopia: A Mendelian Randomization Study. <i>Genetic Epidemiology</i> , 2016, 40, 66-72.	0.6	56
143	Genome-wide association studies in oesophageal adenocarcinoma and Barrett's oesophagus: a large-scale meta-analysis. <i>Lancet Oncology</i> , The, 2016, 17, 1363-1373.	5.1	133
144	Assessment of polygenic effects links primary open-angle glaucoma and age-related macular degeneration. <i>Scientific Reports</i> , 2016, 6, 26885.	1.6	21

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145	Pooled genome wide association detects association upstream of FCRL3 with Graves's disease. BMC Genomics, 2016, 17, 939.	1.2	10
146	Meta-analysis of genome-wide association scans accounting for education level identifies additional loci for refractive error. Nature Communications, 2016, 7, 11008.	5.8	104
147	Childhood gene-environment interactions and age-dependent effects of genetic variants associated with refractive error and myopia: The CREAM Consortium. Scientific Reports, 2016, 6, 25853.	1.6	80
148	Chronic gastroesophageal reflux disease shares genetic background with esophageal adenocarcinoma and Barrett's esophagus. Human Molecular Genetics, 2016, 25, 828-835.	1.4	31
149	Genome-wide association analysis identifies TXNRD2, ATXN2 and FOXC1 as susceptibility loci for primary open-angle glaucoma. Nature Genetics, 2016, 48, 189-194.	9.4	211
150	Germline polymorphisms in an enhancer of <i>PSIP1</i> are associated with progression-free survival in epithelial ovarian cancer. Oncotarget, 2016, 7, 6353-6368.	0.8	29
151	Abstract 4487: An INDEL variant confers melanoma risk through PARP1 expression regulation. , 2016, , .		0
152	<i>PARP1</i> polymorphisms play opposing roles in melanoma occurrence and survival. International Journal of Cancer, 2015, 136, 2488-2489.	2.3	7
153	Polymorphisms in Genes of Relevance for Oestrogen and Oxytocin Pathways and Risk of Barrett's Oesophagus and Oesophageal Adenocarcinoma: A Pooled Analysis from the BEACON Consortium. PLoS ONE, 2015, 10, e0138738.	1.1	9
154	Accurate Imputation-Based Screening of Gln368Ter Myocilin Variant in Primary Open-Angle Glaucoma. , 2015, 56, 5087.		17
155	Genetic burden associated with varying degrees of disease severity in endometriosis. Molecular Human Reproduction, 2015, 21, 594-602.	1.3	30
156	Genetic and Environmental Factors in Conjunctival UV Autofluorescence. JAMA Ophthalmology, 2015, 133, 406.	1.4	30
157	A common variant near TGFBR3 is associated with primary open angle glaucoma. Human Molecular Genetics, 2015, 24, 3880-3892.	1.4	105
158	Association between endometriosis and the interleukin 1A (IL1A) locus. Human Reproduction, 2015, 30, 239-248.	0.4	58
159	ARHGEF12 influences the risk of glaucoma by increasing intraocular pressure. Human Molecular Genetics, 2015, 24, 2689-2699.	1.4	79
160	Meta-analysis of Genome-Wide Association Studies Identifies Novel Loci Associated With Optic Disc Morphology. Genetic Epidemiology, 2015, 39, 207-216.	0.6	72
161	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. Nature Genetics, 2015, 47, 987-995.	9.4	218
162	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	3.2	33

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163	Retinal microvessels reflect familial vulnerability to psychotic symptoms: A comparison of twins discordant for psychotic symptoms and controls. <i>Schizophrenia Research</i> , 2015, 164, 47-52.	1.1	41
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