

Alex W. Hewitt

List of Publications by Citations

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322
papers

11,961
citations

58
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95
g-index

358
ext. papers

15,250
ext. citations

7.7
avg, IF

5.85
L-index

#	Paper	IF	Citations
322	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. <i>Nature Genetics</i> , 2016 , 48, 134-43	36.3	769
321	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017 , 542, 186-190	50.4	412
320	Genome-wide association study identifies susceptibility loci for open angle glaucoma at TMCO1 and CDKN2B-AS1. <i>Nature Genetics</i> , 2011 , 43, 574-8	36.3	329
319	Genome-wide meta-analyses of multi-ancestry cohorts identify multiple new susceptibility loci for refractive error and myopia. <i>Nature Genetics</i> , 2013 , 45, 314-8	36.3	314
318	Common variants near CAV1 and CAV2 are associated with primary open-angle glaucoma. <i>Nature Genetics</i> , 2010 , 42, 906-9	36.3	303
317	Genome-wide association analyses identify multiple loci associated with central corneal thickness and keratoconus. <i>Nature Genetics</i> , 2013 , 45, 155-63	36.3	222
316	Genetic influences on handedness: data from 25,732 Australian and Dutch twin families. <i>Neuropsychologia</i> , 2009 , 47, 330-7	3.2	205
315	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018 , 50, 26-41	36.3	186
314	A genome-wide association study for myopia and refractive error identifies a susceptibility locus at 15q25. <i>Nature Genetics</i> , 2010 , 42, 902-5	36.3	179
313	Genome-wide analysis of multi-ancestry cohorts identifies new loci influencing intraocular pressure and susceptibility to glaucoma. <i>Nature Genetics</i> , 2014 , 46, 1126-1130	36.3	171
312	Deep sequencing of uveal melanoma identifies a recurrent mutation in PLCB4. <i>Oncotarget</i> , 2016 , 7, 4624-31	3.1	168
311	A systematic meta-analysis of genetic association studies for diabetic retinopathy. <i>Diabetes</i> , 2009 , 58, 2137-47	0.9	160
310	Genome-wide association analysis identifies TXNRD2, ATXN2 and FOXC1 as susceptibility loci for primary open-angle glaucoma. <i>Nature Genetics</i> , 2016 , 48, 189-94	36.3	159
309	Common variants near ABCA1, AFAP1 and GMDS confer risk of primary open-angle glaucoma. <i>Nature Genetics</i> , 2014 , 46, 1120-1125	36.3	141
308	Genome-wide association meta-analysis highlights light-induced signaling as a driver for refractive error. <i>Nature Genetics</i> , 2018 , 50, 834-848	36.3	135
307	Common genetic determinants of intraocular pressure and primary open-angle glaucoma. <i>PLoS Genetics</i> , 2012 , 8, e1002611	6	131
306	Ancestral LOXL1 variants are associated with pseudoexfoliation in Caucasian Australians but with markedly lower penetrance than in Nordic people. <i>Human Molecular Genetics</i> , 2008 , 17, 710-6	5.6	129

305	Genome-wide association identifies ATOH7 as a major gene determining human optic disc size. <i>Human Molecular Genetics</i> , 2010 , 19, 2716-24	5.6	118
304	Angiopoietin receptor TEK mutations underlie primary congenital glaucoma with variable expressivity. <i>Journal of Clinical Investigation</i> , 2016 , 126, 2575-87	15.9	117
303	Nine loci for ocular axial length identified through genome-wide association studies, including shared loci with refractive error. <i>American Journal of Human Genetics</i> , 2013 , 93, 264-77	11	116
302	Central corneal thickness is highly heritable: the twin eye studies. <i>Investigative Ophthalmology and Visual Science</i> , 2005 , 46, 3718-22		115
301	Four novel Loci (19q13, 6q24, 12q24, and 5q14) influence the microcirculation in vivo. <i>PLoS Genetics</i> , 2010 , 6, e1001184	6	111
300	The heritability of ocular traits. <i>Survey of Ophthalmology</i> , 2010 , 55, 561-83	6.1	111
299	Genome-wide association study identifies five new susceptibility loci for primary angle closure glaucoma. <i>Nature Genetics</i> , 2016 , 48, 556-62	36.3	109
298	Common genetic variants near the Brittle Cornea Syndrome locus ZNF469 influence the blinding disease risk factor central corneal thickness. <i>PLoS Genetics</i> , 2010 , 6, e1000947	6	106
297	Association of polymorphisms in the hepatocyte growth factor gene promoter with keratoconus 2011 , 52, 8514-9		101
296	Autosomal dominant optic atrophy: penetrance and expressivity in patients with OPA1 mutations. <i>American Journal of Ophthalmology</i> , 2007 , 143, 656-62	4.9	101
295	AAV-Mediated CRISPR/Cas Gene Editing of Retinal Cells In Vivo 2016 , 57, 3470-6		97
294	Genome-wide association study of intraocular pressure uncovers new pathways to glaucoma. <i>Nature Genetics</i> , 2018 , 50, 1067-1071	36.3	86
293	A common variant near TGFBR3 is associated with primary open angle glaucoma. <i>Human Molecular Genetics</i> , 2015 , 24, 3880-92	5.6	84
292	Myocilin allele-specific glaucoma phenotype database. <i>Human Mutation</i> , 2008 , 29, 207-11	4.7	84
291	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, 438-453	5.6	80
290	Meta-analysis of gene-environment-wide association scans accounting for education level identifies additional loci for refractive error. <i>Nature Communications</i> , 2016 , 7, 11008	17.4	79
289	Insights into the genetic architecture of early stage age-related macular degeneration: a genome-wide association study meta-analysis. <i>PLoS ONE</i> , 2013 , 8, e53830	3.7	79
288	The path to open-angle glaucoma gene discovery: endophenotypic status of intraocular pressure, cup-to-disc ratio, and central corneal thickness 2010 , 51, 3509-14		79

287	Genetic dissection of acute anterior uveitis reveals similarities and differences in associations observed with ankylosing spondylitis. <i>Arthritis and Rheumatology</i> , 2015 , 67, 140-51	9.5	78
286	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	36.3	78
285	Genetic association study of exfoliation syndrome identifies a protective rare variant at LOXL1 and five new susceptibility loci. <i>Nature Genetics</i> , 2017 , 49, 993-1004	36.3	72
284	Genome-wide association study identifies seven novel susceptibility loci for primary open-angle glaucoma. <i>Human Molecular Genetics</i> , 2018 , 27, 1486-1496	5.6	72
283	Current state and future prospects of artificial intelligence in ophthalmology: a review. <i>Clinical and Experimental Ophthalmology</i> , 2019 , 47, 128-139	2.4	72
282	Meta-analysis of genome-wide association studies identifies novel loci that influence cupping and the glaucomatous process. <i>Nature Communications</i> , 2014 , 5, 4883	17.4	71
281	A common variant mapping to CACNA1A is associated with susceptibility to exfoliation syndrome. <i>Nature Genetics</i> , 2015 , 47, 387-92	36.3	70
280	Functional and structural implications of the complement factor H Y402H polymorphism associated with age-related macular degeneration. <i>Investigative Ophthalmology and Visual Science</i> , 2008 , 49, 1763-70		70
279	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	36.3	68
278	The association between time spent outdoors and myopia using a novel biomarker of outdoor light exposure 2012 , 53, 4363-70		66
277	Myopia is associated with lower vitamin D status in young adults 2014 , 55, 4552-9		65
276	Meta-analysis of human methylation data for evidence of sex-specific autosomal patterns. <i>BMC Genomics</i> , 2014 , 15, 981	4.5	65
275	Identification of LOXL1 protein and Apolipoprotein E as components of surgically isolated pseudoexfoliation material by direct mass spectrometry. <i>Experimental Eye Research</i> , 2009 , 89, 479-85	3.7	65
274	PSEN1E9, APPswe, and APOE4 Confer Disparate Phenotypes in Human iPSC-Derived Microglia. <i>Stem Cell Reports</i> , 2019 , 13, 669-683	8	64
273	Glaucoma risk alleles at CDKN2B-AS1 are associated with lower intraocular pressure, normal-tension glaucoma, and advanced glaucoma. <i>Ophthalmology</i> , 2012 , 119, 1539-45	7.3	64
272	Apparent autosomal dominant keratoconus in a large Australian pedigree accounted for by digenic inheritance of two novel loci. <i>Human Genetics</i> , 2008 , 124, 379-86	6.3	63
271	Myopia in young adults is inversely related to an objective marker of ocular sun exposure: the Western Australian Raine cohort study. <i>American Journal of Ophthalmology</i> , 2014 , 158, 1079-85	4.9	62
270	ARHGEF12 influences the risk of glaucoma by increasing intraocular pressure. <i>Human Molecular Genetics</i> , 2015 , 24, 2689-99	5.6	62

269	Genome-wide association study for sight-threatening diabetic retinopathy reveals association with genetic variation near the GRB2 gene. <i>Diabetologia</i> , 2015 , 58, 2288-97	10.3	60
268	Complex genetics of complex traits: the case of primary open-angle glaucoma. <i>Clinical and Experimental Ophthalmology</i> , 2006 , 34, 472-84	2.4	60
267	Large-scale cis- and trans-eQTL analyses identify thousands of genetic loci and polygenic scores that regulate blood gene expression. <i>Nature Genetics</i> , 2021 , 53, 1300-1310	36.3	60
266	Genome-wide association meta-analysis of individuals of European ancestry identifies new loci explaining a substantial fraction of hair color variation and heritability. <i>Nature Genetics</i> , 2018 , 50, 652-658	36.3	59
265	Enriched retinal ganglion cells derived from human embryonic stem cells. <i>Scientific Reports</i> , 2016 , 6, 30552	4.2	59
264	Meta-analysis of Genome-Wide Association Studies Identifies Novel Loci Associated With Optic Disc Morphology. <i>Genetic Epidemiology</i> , 2015 , 39, 207-16	2.6	58
263	A Glaucoma Case-control Study of the WDR36 Gene D658G sequence variant. <i>American Journal of Ophthalmology</i> , 2006 , 142, 324-5	4.9	58
262	Drusen in patient-derived hiPSC-RPE models of macular dystrophies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, E8214-E8223	11.5	57
261	Large scale international replication and meta-analysis study confirms association of the 15q14 locus with myopia. The CREAM consortium. <i>Human Genetics</i> , 2012 , 131, 1467-80	6.3	57
260	Angiopoietin-1 is required for Schlemm's canal development in mice and humans. <i>Journal of Clinical Investigation</i> , 2017 , 127, 4421-4436	15.9	57
259	Electrical Stimulation Promotes Cardiac Differentiation of Human Induced Pluripotent Stem Cells. <i>Stem Cells International</i> , 2016 , 2016, 1718041	5	57
258	Identification of myopia-associated WNT7B polymorphisms provides insights into the mechanism underlying the development of myopia. <i>Nature Communications</i> , 2015 , 6, 6689	17.4	56
257	Recessive mutations in SLC38A8 cause foveal hypoplasia and optic nerve misrouting without albinism. <i>American Journal of Human Genetics</i> , 2013 , 93, 1143-50	11	56
256	Projected worldwide disease burden from giant cell arteritis by 2050. <i>Journal of Rheumatology</i> , 2015 , 42, 119-25	4.1	56
255	A Global Social Media Survey of Attitudes to Human Genome Editing. <i>Cell Stem Cell</i> , 2016 , 18, 569-72	18	56
254	Copy number variations of TBK1 in Australian patients with primary open-angle glaucoma. <i>American Journal of Ophthalmology</i> , 2015 , 159, 124-30.e1	4.9	54
253	How significant is a family history of glaucoma? Experience from the Glaucoma Inheritance Study in Tasmania. <i>Clinical and Experimental Ophthalmology</i> , 2007 , 35, 793-9	2.4	54
252	Association of TCF4 and CLU polymorphisms with Fuchs' endothelial dystrophy and implication of CLU and TGFBI proteins in the disease process. <i>European Journal of Human Genetics</i> , 2012 , 20, 632-8	5.3	51

251	WNT10A exonic variant increases the risk of keratoconus by decreasing corneal thickness. <i>Human Molecular Genetics</i> , 2015 , 24, 5060-8	5.6	50
250	Measurement of Systemic Mitochondrial Function in Advanced Primary Open-Angle Glaucoma and Leber Hereditary Optic Neuropathy. <i>PLoS ONE</i> , 2015 , 10, e0140919	3.7	47
249	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. <i>Nature Communications</i> , 2021 , 12, 1258	17.4	47
248	Novel pleiotropic risk loci for melanoma and nevus density implicate multiple biological pathways. <i>Nature Communications</i> , 2018 , 9, 4774	17.4	47
247	Gene therapy for visual loss: Opportunities and concerns. <i>Progress in Retinal and Eye Research</i> , 2019 , 68, 31-53	20.5	46
246	The association between pterygium and conjunctival ultraviolet autofluorescence: the Norfolk Island Eye Study. <i>Acta Ophthalmologica</i> , 2013 , 91, 363-70	3.7	45
245	Australian and New Zealand Registry of Advanced Glaucoma: methodology and recruitment. <i>Clinical and Experimental Ophthalmology</i> , 2012 , 40, 569-75	2.4	45
244	PAX6 mutations may be associated with high myopia. <i>Ophthalmic Genetics</i> , 2007 , 28, 179-82	1.2	45
243	Vitreous biomarkers in diabetic retinopathy: a systematic review and meta-analysis. <i>Journal of Diabetes and Its Complications</i> , 2014 , 28, 419-25	3.2	44
242	A comparative analysis of high-throughput platforms for validation of a circulating microRNA signature in diabetic retinopathy. <i>Scientific Reports</i> , 2015 , 5, 10375	4.9	43
241	What is the appropriate age cut-off for cycloplegia in refraction?. <i>Acta Ophthalmologica</i> , 2014 , 92, e458-62	3.2	43
240	Raine eye health study: design, methodology and baseline prevalence of ophthalmic disease in a birth-cohort study of young adults. <i>Ophthalmic Genetics</i> , 2013 , 34, 199-208	1.2	41
239	Mutation in TMEM98 in a large white kindred with autosomal dominant nanophthalmos linked to 17p12-q12. <i>JAMA Ophthalmology</i> , 2014 , 132, 970-7	3.9	40
238	Heritability of central corneal thickness in nuclear families 2009 , 50, 4087-90		40
237	Cardiac Repair With a Novel Population of Mesenchymal Stem Cells Resident in the Human Heart. <i>Stem Cells</i> , 2015 , 33, 3100-13	5.8	39
236	Optic disc morphology--rethinking shape. <i>Progress in Retinal and Eye Research</i> , 2009 , 28, 227-48	20.5	39
235	Reliability and validity of conjunctival ultraviolet autofluorescence measurement. <i>British Journal of Ophthalmology</i> , 2012 , 96, 801-5	5.5	39
234	Genetic dissection of myopia: evidence for linkage of ocular axial length to chromosome 5q. <i>Ophthalmology</i> , 2008 , 115, 1053-1057.e2	7.3	39

233	Assessing the Genetic Predisposition of Education on Myopia: A Mendelian Randomization Study. <i>Genetic Epidemiology</i> , 2016 , 40, 66-72	2.6	39
232	Hypomethylation of the IL17RC promoter in peripheral blood leukocytes is not a hallmark of age-related macular degeneration. <i>Cell Reports</i> , 2013 , 5, 1527-35	10.6	37
231	Mitochondrial oxidative phosphorylation compensation may preserve vision in patients with OPA1-linked autosomal dominant optic atrophy. <i>PLoS ONE</i> , 2011 , 6, e21347	3.7	37
230	Cross-ancestry genome-wide association analysis of corneal thickness strengthens link between complex and Mendelian eye diseases. <i>Nature Communications</i> , 2018 , 9, 1864	17.4	37
229	Higher prevalence of myocilin mutations in advanced glaucoma in comparison with less advanced disease in an Australasian disease registry. <i>Ophthalmology</i> , 2013 , 120, 1135-43	7.3	36
228	Effect of birth parameters on retinal vascular caliber: the Twins Eye Study in Tasmania. <i>Hypertension</i> , 2009 , 53, 487-93	8.5	36
227	Methods of Retinal Ganglion Cell Differentiation From Pluripotent Stem Cells. <i>Translational Vision Science and Technology</i> , 2014 , 3, 7	3.3	36
226	Genome engineering in ophthalmology: Application of CRISPR/Cas to the treatment of eye disease. <i>Progress in Retinal and Eye Research</i> , 2016 , 53, 1-20	20.5	36
225	Rapid inexpensive genome-wide association using pooled whole blood. <i>Genome Research</i> , 2009 , 19, 2075-80	5.9	35
224	Mitochondrial replacement in an iPSC model of Leber's hereditary optic neuropathy. <i>Aging</i> , 2017 , 9, 1341-63	15.35	35
223	Genetically low vitamin D concentrations and myopic refractive error: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2017 , 46, 1882-1890	7.8	34
222	Quantitative genetic analysis of the retinal vascular caliber: the Australian Twins Eye Study. <i>Hypertension</i> , 2009 , 54, 788-95	8.5	34
221	Disease severity of familial glaucoma compared with sporadic glaucoma. <i>JAMA Ophthalmology</i> , 2006 , 124, 950-4		34
220	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	3.6	33
219	Association of genetic variants in the TMCO1 gene with clinical parameters related to glaucoma and characterization of the protein in the eye 2012 , 53, 4917-25		33
218	Twins eye study in Tasmania (TEST): rationale and methodology to recruit and examine twins. <i>Twin Research and Human Genetics</i> , 2009 , 12, 441-54	2.2	33
217	Elevation of serum asymmetrical and symmetrical dimethylarginine in patients with advanced glaucoma 2012 , 53, 1923-7		33
216	Study of mitochondrial respiratory defects on reprogramming to human induced pluripotent stem cells. <i>Aging</i> , 2016 , 8, 945-57	5.6	33

215	Giant cell arteritis: ophthalmic manifestations of a systemic disease. <i>Graefes Archive for Clinical and Experimental Ophthalmology</i> , 2016 , 254, 2291-2306	3.8	33
214	Genome-wide association studies for diabetic macular edema and proliferative diabetic retinopathy. <i>BMC Medical Genetics</i> , 2018 , 19, 71	2.1	32
213	Genetic variants near PDGFRA are associated with corneal curvature in Australians 2012 , 53, 7131-6		31
212	Drug discovery using induced pluripotent stem cell models of neurodegenerative and ocular diseases. <i>Pharmacology & Therapeutics</i> , 2017 , 177, 32-43	13.9	30
211	Genome-Wide Association Shows that Pigmentation Genes Play a Role in Skin Aging. <i>Journal of Investigative Dermatology</i> , 2017 , 137, 1887-1894	4.3	30
210	A genetic variant regulating miR-126 is associated with sight threatening diabetic retinopathy. <i>Diabetes and Vascular Disease Research</i> , 2015 , 12, 133-8	3.3	30
209	Novel quantitative trait loci for central corneal thickness identified by candidate gene analysis of osteogenesis imperfecta genes. <i>Human Genetics</i> , 2010 , 127, 33-44	6.3	30
208	Single cell RNA sequencing of stem cell-derived retinal ganglion cells. <i>Scientific Data</i> , 2018 , 5, 180013	8.2	29
207	Association of genetic variants with primary angle closure glaucoma in two different populations. <i>PLoS ONE</i> , 2013 , 8, e67903	3.7	29
206	Optic disc evaluation in optic neuropathies: the optic disc assessment project. <i>Ophthalmology</i> , 2011 , 118, 964-70	7.3	29
205	Classification of iris colour: review and refinement of a classification schema. <i>Clinical and Experimental Ophthalmology</i> , 2011 , 39, 462-71	2.4	29
204	An Interactive Multimedia Approach to Improving Informed Consent for Induced Pluripotent Stem Cell Research. <i>Cell Stem Cell</i> , 2016 , 18, 307-8	18	27
203	Pterygium and conjunctival ultraviolet autofluorescence in young Australian adults: the Raine study. <i>Clinical and Experimental Ophthalmology</i> , 2015 , 43, 300-7	2.4	26
202	Automated Cell Culture Systems and Their Applications to Human Pluripotent Stem Cell Studies. <i>SLAS Technology</i> , 2018 , 23, 315-325	3	26
201	Genotypic and phenotypic spectrum of X-linked retinoschisis in Australia. <i>Clinical and Experimental Ophthalmology</i> , 2005 , 33, 233-9	2.4	26
200	Smartphone use in ophthalmology: What is their place in clinical practice?. <i>Survey of Ophthalmology</i> , 2020 , 65, 250-262	6.1	26
199	Glaucoma spectrum and age-related prevalence of individuals with FOXC1 and PITX2 variants. <i>European Journal of Human Genetics</i> , 2017 , 25, 839-847	5.3	25
198	Analysis combining correlated glaucoma traits identifies five new risk loci for open-angle glaucoma. <i>Scientific Reports</i> , 2018 , 8, 3124	4.9	25

197	Genetic investigation into the endophenotypic status of central corneal thickness and optic disc parameters in relation to open-angle glaucoma. <i>American Journal of Ophthalmology</i> , 2012 , 154, 833-842.	4.9	25
196	Risk Alleles Associated with Neovascularization in a Pachychoroid Phenotype. <i>Ophthalmology</i> , 2016 , 123, 2628-2630	7.3	25
195	Prevalence of Keratoconus Based on Scheimpflug Imaging: The Raine Study. <i>Ophthalmology</i> , 2021 , 128, 515-521	7.3	25
194	Distribution of astigmatism as a function of age in an Australian population. <i>Acta Ophthalmologica</i> , 2015 , 93, e377-85	3.7	24
193	Establishment and evolution of the Australian Inherited Retinal Disease Register and DNA Bank. <i>Clinical and Experimental Ophthalmology</i> , 2013 , 41, 476-83	2.4	24
192	Association of eNOS polymorphisms with primary angle-closure glaucoma 2013 , 54, 2108-14		24
191	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021 ,	50.4	24
190	TIMP1, TIMP2, and TIMP4 are increased in aqueous humor from primary open angle glaucoma patients. <i>Molecular Vision</i> , 2015 , 21, 1162-72	2.3	24
189	Association of Genetic Variation With Keratoconus. <i>JAMA Ophthalmology</i> , 2020 , 138, 174-181	3.9	24
188	A Need for Better Understanding Is the Major Determinant for Public Perceptions of Human Gene Editing. <i>Human Gene Therapy</i> , 2019 , 30, 36-43	4.8	23
187	Identification of a candidate gene for astigmatism 2013 , 54, 1260-7		23
186	Genetic Association at the 9p21 Glaucoma Locus Contributes to Sex Bias in Normal-Tension Glaucoma 2016 , 57, 3416-21		23
185	Development of a Modular Automated System for Maintenance and Differentiation of Adherent Human Pluripotent Stem Cells. <i>SLAS Discovery</i> , 2017 , 22, 1016-1025	3.4	22
184	Replication of genetic loci implicated in diabetic retinopathy 2014 , 55, 1666-71		22
183	Heritable features of the optic disc: a novel twin method for determining genetic significance. <i>Investigative Ophthalmology and Visual Science</i> , 2007 , 48, 2469-75		22
182	Macular Ganglion Cell-Inner Plexiform Layer Loss Precedes Peripapillary Retinal Nerve Fiber Layer Loss in Glaucoma with Lower Intraocular Pressure. <i>Ophthalmology</i> , 2019 , 126, 1119-1130	7.3	21
181	Genome-wide meta-analysis identifies novel loci associated with age-related macular degeneration. <i>Journal of Human Genetics</i> , 2020 , 65, 657-665	4.3	21
180	A COL17A1 Splice-Altering Mutation Is Prevalent in Inherited Recurrent Corneal Erosions. <i>Ophthalmology</i> , 2016 , 123, 709-22	7.3	21

179	Associations of birth weight with ocular biometry, refraction, and glaucomatous endophenotypes: the Australian Twins Eye Study. <i>American Journal of Ophthalmology</i> , 2010 , 150, 909-16	4.9	21
178	Genetic isolates in ophthalmic diseases. <i>Ophthalmic Genetics</i> , 2008 , 29, 149-61	1.2	21
177	Genome-wide association study for refractive astigmatism reveals genetic co-determination with spherical equivalent refractive error: the CREAM consortium. <i>Human Genetics</i> , 2015 , 134, 131-46	6.3	20
176	Genome-wide association study success in ophthalmology. <i>Current Opinion in Ophthalmology</i> , 2014 , 25, 386-93	5.1	20
175	Heritability of strabismus: genetic influence is specific to eso-deviation and independent of refractive error. <i>Twin Research and Human Genetics</i> , 2012 , 15, 624-30	2.2	20
174	Heritability of the iridotrabecular angle width measured by optical coherence tomography in Chinese children: the Guangzhou twin eye study. <i>Investigative Ophthalmology and Visual Science</i> , 2008 , 49, 1356-61		20
173	Friedreich's ataxia induced pluripotent stem cell-derived cardiomyocytes display electrophysiological abnormalities and calcium handling deficiency. <i>Aging</i> , 2017 , 9, 1440-1452	5.6	20
172	Comparative performance of the BGI and Illumina sequencing technology for single-cell RNA-sequencing. <i>NAR Genomics and Bioinformatics</i> , 2020 , 2, lqaa034	3.7	19
171	Multiallelic copy number variation in the complement component 4A (C4A) gene is associated with late-stage age-related macular degeneration (AMD). <i>Journal of Neuroinflammation</i> , 2016 , 13, 81	10.1	19
170	Participant understanding and recall of informed consent for induced pluripotent stem cell biobanking. <i>Cell and Tissue Banking</i> , 2016 , 17, 449-56	2.2	19
169	Defined Medium Conditions for the Induction and Expansion of Human Pluripotent Stem Cell-Derived Retinal Pigment Epithelium. <i>Stem Cell Reviews and Reports</i> , 2016 , 12, 179-88	6.4	19
168	Associations between depression and anxiety symptoms and retinal vessel caliber in adolescents and young adults. <i>Psychosomatic Medicine</i> , 2014 , 76, 732-8	3.7	19
167	Genetic loci for retinal arteriolar microcirculation. <i>PLoS ONE</i> , 2013 , 8, e65804	3.7	19
166	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	12.1	19
165	Genotype-free demultiplexing of pooled single-cell RNA-seq. <i>Genome Biology</i> , 2019 , 20, 290	18.3	19
164	Common Sequence Variation in the VEGFC Gene Is Associated with Diabetic Retinopathy and Diabetic Macular Edema. <i>Ophthalmology</i> , 2015 , 122, 1828-36	7.3	18
163	Biallelic CPAMD8 Variants Are a Frequent Cause of Childhood and Juvenile Open-Angle Glaucoma. <i>Ophthalmology</i> , 2020 , 127, 758-766	7.3	18
162	Myocilin Predictive Genetic Testing for Primary Open-Angle Glaucoma Leads to Early Identification of At-Risk Individuals. <i>Ophthalmology</i> , 2017 , 124, 303-309	7.3	17

161	Assessment of polygenic effects links primary open-angle glaucoma and age-related macular degeneration. <i>Scientific Reports</i> , 2016 , 6, 26885	4.9	17
160	The p53 codon 72 PRO/PRO genotype may be associated with initial central visual field defects in caucasians with primary open angle glaucoma. <i>PLoS ONE</i> , 2012 , 7, e45613	3.7	17
159	Prevalence of FOXC1 Variants in Individuals With a Suspected Diagnosis of Primary Congenital Glaucoma. <i>JAMA Ophthalmology</i> , 2019 , 137, 348-355	3.9	16
158	Role of lysophosphatidic acid in the retinal pigment epithelium and photoreceptors. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2018 , 1863, 750-761	5	16
157	Genetic and environmental factors in conjunctival UV autofluorescence. <i>JAMA Ophthalmology</i> , 2015 , 133, 406-12	3.9	16
156	The optic nerve head in hereditary optic neuropathies. <i>Nature Reviews Neurology</i> , 2009 , 5, 277-87	15	16
155	A myocilin Gln368STOP homozygote does not exhibit a more severe glaucoma phenotype than heterozygous cases. <i>American Journal of Ophthalmology</i> , 2006 , 141, 402-3	4.9	16
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51	Profile of ocular trauma in the Solomon Islands. <i>Clinical and Experimental Ophthalmology</i> , 2014 , 42, 440-444	6.4	3
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46	Effect of phacoemulsification cataract surgery on intraocular pressure in early glaucoma: A prospective multi-site study. <i>Clinical and Experimental Ophthalmology</i> , 2020 , 48, 442-449	2.4	2
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41	Comparative analysis of loop-mediated isothermal amplification (LAMP)-based assays for rapid detection of SARS-CoV-2 genes. <i>Scientific Reports</i> , 2021 , 11, 22493	4.9	2
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37	The Ark: a customizable web-based data management tool for health and medical research. <i>Bioinformatics</i> , 2017 , 33, 624-626	7.2	2
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23	The Immortal Life of Ethics? The Alienation of Body Tissue, Ethics and the Informed Consent Procedure Within Induced Pluripotent Stem Cell Research 2017 , 61-87		1
22	Mitochondrial replacement in an iPSC model of Leber's hereditary optic neuropathy		1
21	Efficacy and dynamics of self-targeting CRISPR/Cas constructs for gene editing in the retina		1
20	Vision impairment and refractive errors in refugees presenting to community optometry clinics in Victoria, Australia. <i>Australasian journal of optometry, The</i> , 2020 , 103, 668-674	2.7	1
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14	Genome-wide linkage and association analysis of primary open-angle glaucoma endophenotypes in the Norfolk Island isolate. <i>Molecular Vision</i> , 2017 , 23, 660-665	2.3	0
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1	Exposure to Smoking and Alcohol, and Passive Smoking during Childhood: Effect on the Retinal Nerve Fibre Layer in Young Adulthood. <i>Ophthalmic Epidemiology</i> , 2021 , 1-8	1.9	