## Angela J Cree

List of Publications by Year in descending order

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		471371	454834
32	2,555	17	30
papers	citations	h-index	g-index
32	32	32	4266
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. Nature Genetics, 2016, 48, 134-143.	9.4	1,167
2	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. Nature Communications, 2021, 12, 1258.	5.8	196
3	Multitrait analysis of glaucoma identifies new risk loci and enables polygenic prediction of disease susceptibility and progression. Nature Genetics, 2020, 52, 160-166.	9.4	192
4	Association between the SERPING1 gene and age-related macular degeneration: a two-stage case–control study. Lancet, The, 2008, 372, 1828-1834.	6.3	156
5	Eplerenone for chronic central serous chorioretinopathy in patients with active, previously untreated disease for more than 4 months (VICI): a randomised, double-blind, placebo-controlled trial. Lancet, The, 2020, 395, 294-303.	6.3	134
6	New insights into the genetics of primary open-angle glaucoma based on meta-analyses of intraocular pressure and optic disc characteristics Human Molecular Genetics, 2017, 26, ddw399.	1.4	120
7	Meta-analysis of genome-wide association studies identifies novel loci that influence cupping and the glaucomatous process. Nature Communications, 2014, 5, 4883.	5.8	89
8	Pharmacogenetic Associations with Vascular Endothelial Growth Factor Inhibition in Participants with Neovascular Age-related Macular Degeneration in the IVAN Study. Ophthalmology, 2013, 120, 2637-2643.	2.5	59
9	Support for the involvement of complement factor I in age-related macular degeneration. European Journal of Human Genetics, 2010, 18, 15-16.	1.4	54
10	Sorsby fundus dystrophy – A review of pathology and disease mechanisms. Experimental Eye Research, 2017, 165, 35-46.	1.2	45
11	Age-related Macular Degeneration and Modification of Systemic Complement Factor H Production Through Liver Transplantation. Ophthalmology, 2013, 120, 1612-1618.	2.5	39
12	An Intraocular Pressure Polygenic Risk Score Stratifies Multiple Primary Open-Angle Glaucoma Parameters Including Treatment Intensity. Ophthalmology, 2020, 127, 901-907.	2.5	37
13	Rare Genetic Variants in Complement Factor I Lead to Low FI Plasma Levels Resulting in Increased Risk of Age-Related Macular Degeneration. , 2020, 61, 18.		36
14	Association of Genetic Variants With Response to Anti–Vascular Endothelial Growth Factor Therapy in Age-Related Macular Degeneration. JAMA Ophthalmology, 2018, 136, 875.	1.4	30
15	Characteristics of p.Gln368Ter Myocilin Variant and Influence of Polygenic Risk on Glaucoma Penetrance in the UK Biobank. Ophthalmology, 2021, 128, 1300-1311.	2.5	27
16	Ex-vivo models of the Retinal Pigment Epithelium (RPE) in long-term culture faithfully recapitulate key structural and physiological features of native RPE. Tissue and Cell, 2017, 49, 447-460.	1.0	22
17	Multi-trait genome-wide association study identifies new loci associated with optic disc parameters. Communications Biology, 2019, 2, 435.	2.0	22
18	Complement factor I and age-related macular degeneration. Molecular Vision, 2014, 20, 1253-7.	1.1	18

#	Article	IF	CITATIONS
19	The effect of systemic levels of TNF-alpha and complement pathway activity on outcomes of VEGF inhibition in neovascular AMD. Eye, 2022, 36, 2192-2199.	1.1	16
20	A convenient protocol for establishing a human cell culture model of the outer retina F1000Research, 2018, 7, 1107.	0.8	13
21	Clinical efficacy of eplerenone versus placebo for central serous chorioretinopathy: study protocol for the VICI randomised controlled trial. Eye, 2019, 33, 295-303.	1.1	13
22	In vitro stem cell modelling demonstrates a proofâ€ofâ€concept for excess functional mutant TIMP3 as the cause of S orsby f undus d ystrophy. Journal of Pathology, 2020, 252, 138-150.	2.1	10
23	A small gene sequencing panel realises a high diagnostic rate in patients with congenital nystagmus following basic phenotyping. Scientific Reports, 2019, 9, 13229.	1.6	9
24	Developing and validating a multivariable prediction model which predicts progression of intermediate to late age-related macular degeneration—the PINNACLE trial protocol. Eye, 2023, 37, 1275-1283.	1,1	9
25	Comprehensive sequencing of the myocilin gene in a selected cohort of severe primary open-angle glaucoma patients. Scientific Reports, 2019, 9, 3100.	1.6	8
26	Oligomeric AÎ $^2$ 1-42 Induces an AMD-Like Phenotype and Accumulates in Lysosomes to Impair RPE Function. Cells, 2021, 10, 413.	1.8	8
27	Prevalence and phenotype associations of complement factor I mutations in geographic atrophy. Human Mutation, 2021, 42, 1139-1152.	1.1	8
28	Epigenetic Age Acceleration Is Not Associated with Age-Related Macular Degeneration. International Journal of Molecular Sciences, 2021, 22, 13457.	1.8	8
29	A novel, wearable, electronic visual aid to assist those with reduced peripheral vision. PLoS ONE, 2019, 14, e0223755.	1.1	5
30	Eplerenone versus placebo for chronic central serous chorioretinopathy: the VICI RCT. Efficacy and Mechanism Evaluation, 2021, 8, 1-82.	0.9	4
31	A Genome-Wide Complement for Central Serous Chorioretinopathy. JAMA Ophthalmology, 2018, 136, 1136.	1.4	1
32	AMD Risk Alleles Are Not Implicated in Age-Related Macular Degeneration in Patients with Liver Transplantation. Ophthalmology Retina, 2018, 2, 872-874.	1,2	0