Bernhard H F Weber

List of Publications by Year in descending order

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76 papers 4,886 citations

31 h-index

147566

63 g-index

76 all docs 76
docs citations

76 times ranked 7853 citing authors

#	Article	IF	CITATIONS
1	Incidence, progression and risk factors of age-related macular degeneration in 35–95-year-old individuals from three jointly designed German cohort studies. BMJ Open Ophthalmology, 2022, 7, e000912.	0.8	7
2	Vitronectin and Its Interaction with PAI-1 Suggests a Functional Link to Vascular Changes in AMD Pathobiology. Cells, 2022, 11, 1766.	1.8	2
3	Epistatic interactions of genetic loci associated with age-related macular degeneration. Scientific Reports, 2021, 11, 13114.	1.6	4
4	Assigning Co-Regulated Human Genes and Regulatory Gene Clusters. Cells, 2021, 10, 2395.	1.8	4
5	OUP accepted manuscript. Human Molecular Genetics, 2021, , .	1.4	1
6	Performance of Breast Cancer Polygenic Risk Scores in 760 Female <i>CHEK2</i> Germline Mutation Carriers. Journal of the National Cancer Institute, 2021, 113, 893-899.	3.0	21
7	Insights into the loss of the Y chromosome with age in control individuals and in patients with age-related macular degeneration using genotyping microarray data. Human Genetics, 2020, 139, 401-407.	1.8	18
8	Complement C3 Inhibitor Pegcetacoplan for Geographic Atrophy Secondary to Age-Related Macular Degeneration. Ophthalmology, 2020, 127, 186-195.	2.5	306
9	Altered Protein Function Caused by AMD-associated Variant rs704 Links Vitronectin to Disease Pathology. , 2020, 61, 2.		14
10	Pleiotropic Locus 15q24.1 Reveals a Gender-Specific Association with Neovascular but Not Atrophic Age-Related Macular Degeneration (AMD). Cells, 2020, 9, 2257.	1.8	5
11	A mega-analysis of expression quantitative trait loci in retinal tissue. PLoS Genetics, 2020, 16, e1008934.	1.5	22
11	A mega-analysis of expression quantitative trait loci in retinal tissue. PLoS Genetics, 2020, 16, e1008934. Genome-wide association meta-analysis for early age-related macular degeneration highlights novel loci and insights for advanced disease. BMC Medical Genomics, 2020, 13, 120.	0.7	56
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12	Genome-wide association meta-analysis for early age-related macular degeneration highlights novel loci and insights for advanced disease. BMC Medical Genomics, 2020, 13, 120. Clinical Heterogeneity in Autosomal Recessive Bestrophinopathy with Biallelic Mutations in the BEST1	0.7	56
12	Genome-wide association meta-analysis for early age-related macular degeneration highlights novel loci and insights for advanced disease. BMC Medical Genomics, 2020, 13, 120. Clinical Heterogeneity in Autosomal Recessive Bestrophinopathy with Biallelic Mutations in the BEST1 Gene. International Journal of Molecular Sciences, 2020, 21, 9353. Learning from Fifteen Years of Genome-Wide Association Studies in Age-Related Macular Degeneration.	0.7	56 8
12 13 14	Genome-wide association meta-analysis for early age-related macular degeneration highlights novel loci and insights for advanced disease. BMC Medical Genomics, 2020, 13, 120. Clinical Heterogeneity in Autosomal Recessive Bestrophinopathy with Biallelic Mutations in the BEST1 Gene. International Journal of Molecular Sciences, 2020, 21, 9353. Learning from Fifteen Years of Genome-Wide Association Studies in Age-Related Macular Degeneration. Cells, 2020, 9, 2267. Resolving the dark matter of ABCA4 for 1054 Stargardt disease probands through integrated genomics	0.7 1.8 1.8	56 8 18
12 13 14 15	Genome-wide association meta-analysis for early age-related macular degeneration highlights novel loci and insights for advanced disease. BMC Medical Genomics, 2020, 13, 120. Clinical Heterogeneity in Autosomal Recessive Bestrophinopathy with Biallelic Mutations in the BEST1 Gene. International Journal of Molecular Sciences, 2020, 21, 9353. Learning from Fifteen Years of Genome-Wide Association Studies in Age-Related Macular Degeneration. Cells, 2020, 9, 2267. Resolving the dark matter of ABCA4 for 1054 Stargardt disease probands through integrated genomics and transcriptomics. Genetics in Medicine, 2020, 22, 1235-1246. Mutation-Dependent Pathomechanisms Determine the Phenotype in the Bestrophinopathies.	0.7 1.8 1.8	56 8 18

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19	A Circulating MicroRNA Profile in a Laser-Induced Mouse Model of Choroidal Neovascularization. International Journal of Molecular Sciences, 2020, 21, 2689.	1.8	8
20	Investigating the effects of additional truncating variants in DNA-repair genes on breast cancer risk in BRCA1-positive women. BMC Cancer, 2019, 19, 787.	1.1	10
21	Assessment of Novel Genome-Wide Significant Gene Loci and Lesion Growth in Geographic Atrophy Secondary to Age-Related Macular Degeneration. JAMA Ophthalmology, 2019, 137, 867.	1.4	28
22	The Y227N mutation affects bestrophin-1 protein stability and impairs sperm function in a mouse model of Best vitelliform macular dystrophy. Biology Open, 2019, 8, .	0.6	5
23	Costâ€effective molecular inversion probeâ€based <i>ABCA4</i> sequencing reveals deepâ€intronic variants in Stargardt disease. Human Mutation, 2019, 40, 1749-1759.	1.1	39
24	Retinal Layer Thicknesses in Early Age-Related Macular Degeneration: Results From the German AugUR Study. , 2019, 60, 1581.		34
25	Identification of the retinoschisin-binding site on the retinal Na/K-ATPase. PLoS ONE, 2019, 14, e0216320.	1.1	13
26	On the differences between mega―and meta―mputation and analysis exemplified on the genetics of age―elated macular degeneration. Genetic Epidemiology, 2019, 43, 559-576.	0.6	5
27	The agonistic TSPO ligand XBD173 attenuates the glial response thereby protecting inner retinal neurons in a murine model of retinal ischemia. Journal of Neuroinflammation, 2019, 16, 43.	3.1	35
28	Cell-Type-Specific Complement Expression in the Healthy and Diseased Retina. Cell Reports, 2019, 29, 2835-2848.e4.	2.9	81
29	Y chromosome mosaicism is associated with age-related macular degeneration. European Journal of Human Genetics, 2019, 27, 36-41.	1.4	49
30	Gene panel testing of 5589 <i><scp>BRCA</scp>1/2</i> â€negative index patients with breast cancer in a routine diagnostic setting: results of the German Consortium for Hereditary Breast and Ovarian Cancer. Cancer Medicine, 2018, 7, 1349-1358.	1.3	126
31	BEST1 protein stability and degradation pathways differ between autosomal dominant Best disease and autosomal recessive bestrophinopathy accounting for the distinct retinal phenotypes. Human Molecular Genetics, 2018, 27, 1630-1641.	1.4	32
32	Recurrent structural variation, clustered sites of selection, and disease risk for the complement factor H ($<$ i>CFH) gene family. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E4433-E4442.	3.3	43
33	A mega-analysis of expression quantitative trait loci (eQTL) provides insight into the regulatory architecture of gene expression variation in liver. Scientific Reports, 2018, 8, 5865.	1.6	52
34	A Deep Learning Algorithm for Prediction of Age-Related Eye Disease Study Severity Scale for Age-Related Macular Degeneration from Color Fundus Photography. Ophthalmology, 2018, 125, 1410-1420.	2.5	365
35	Mutational spectrum in a worldwide study of 29,700 families with <i>BRCA1</i> or <i>BRCA2</i> mutations. Human Mutation, 2018, 39, 593-620.	1.1	224
36	Pleiotropic Effects of Risk Factors in Age-Related Macular Degeneration and Seemingly Unrelated Complex Diseases. Advances in Experimental Medicine and Biology, 2018, 1074, 247-255.	0.8	1

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37	Views of ophthalmologists on the genetics of age-related macular degeneration: Results of a qualitative study. PLoS ONE, 2018, 13, e0209328.	1.1	9
38	Pathomechanism of mutated and secreted retinoschisin in X-linked juvenile retinoschisis. Experimental Eye Research, 2018, 177, 23-34.	1.2	14
39	Evaluation of serum sphingolipids and the influence of genetic risk factors in age-related macular degeneration. PLoS ONE, 2018, 13, e0200739.	1.1	19
40	Effects of genetic variants in the TSPO gene on protein structure and stability. PLoS ONE, 2018, 13, e0195627.	1.1	19
41	Prevalence of pathogenic BRCA1/2 germline mutations among 802 women with unilateral triple-negative breast cancer without family cancer history. BMC Cancer, 2018, 18, 265.	1.1	84
42	Choroidal Flow Signal in Late-Onset Stargardt Disease and Age-Related Macular Degeneration: An OCT-Angiography Study., 2018, 59, AMD122.		38
43	Investigating the modulation of genetic effects on late AMD by age and sex: Lessons learned and two additional loci. PLoS ONE, 2018, 13, e0194321.	1.1	19
44	On the impact of different approaches to classify age-related macular degeneration: Results from the German AugUR study. Scientific Reports, 2018, 8, 8675.	1.6	31
45	Twenty-year follow-up of a familial case of PTH1R-associated primary failure of tooth eruption. American Journal of Orthodontics and Dentofacial Orthopedics, 2017, 151, 598-606.	0.8	6
46	Recombinant Haplotypes Narrow the ARMS2/HTRA1 Association Signal for Age-Related Macular Degeneration. Genetics, 2017, 205, 919-924.	1.2	65
47	Retinoschisin is linked to retinal Na/K-ATPase signaling and localization. Molecular Biology of the Cell, 2017, 28, 2178-2189.	0.9	26
48	The Xâ€linked juvenile retinoschisis protein retinoschisin is a novel regulator of mitogenâ€activated protein kinase signalling and apoptosis in the retina. Journal of Cellular and Molecular Medicine, 2017, 21, 768-780.	1.6	18
49	Genetic pleiotropy between age-related macular degeneration and 16 complex diseases and traits. Genome Medicine, 2017, 9, 29.	3.6	52
50	An Eye on Age-Related Macular Degeneration: The Role of MicroRNAs in Disease Pathology. Molecular Diagnosis and Therapy, 2017, 21, 31-43.	1.6	78
51	Mutations in the Genes for Interphotoreceptor Matrix Proteoglycans, IMPG1 and IMPG2, in Patients with Vitelliform Macular Lesions. Genes, 2017, 8, 170.	1.0	24
52	Mutation Spectrum of the <i>ABCA4</i> Gene in 335 Stargardt Disease Patients From a Multicenter German Cohortâ€"Impact of Selected Deep Intronic Variants and Common SNPs., 2017, 58, 394.		104
53	Progression of Late-Onset Stargardt Disease. , 2016, 57, 5186.		57
54	Distinct Genetic Risk Profile of the Rapidly Progressing Diffuse-Trickling Subtype of Geographic Atrophy in Age-Related Macular Degeneration (AMD). , 2016, 57, 2463.		22

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55	Features of Age-Related Macular Degeneration in the General Adults and Their Dependency on Age, Sex, and Smoking: Results from the German KORA Study. PLoS ONE, 2016, 11, e0167181.	1.1	27
56	Prevalence of <i>BRCA1/2 </i> germline mutations in 21â€401 families with breast and ovarian cancer. Journal of Medical Genetics, 2016, 53, 465-471.	1.5	179
57	Differentiation of murine models of "negative ERG―by single and repetitive light stimuli. Documenta Ophthalmologica, 2016, 132, 101-109.	1.0	17
58	Long-Term Macular Changes in the First Proband of Autosomal Dominant Vitreoretinochoroidopathy (ADVIRC) Due to a Newly Identified Mutation in <i>BEST1</i> . Ophthalmic Genetics, 2016, 37, 102-108.	0.5	16
59	Multiallelic copy number variation in the complement component 4A (C4A) gene is associated with late-stage age-related macular degeneration (AMD). Journal of Neuroinflammation, 2016, 13, 81.	3.1	31
60	Mutations in (i) EXOSC2 (i) are associated with a novel syndrome characterised by retinitis pigmentosa, progressive hearing loss, premature ageing, short stature, mild intellectual disability and distinctive gestalt. Journal of Medical Genetics, 2016, 53, 419-425.	1.5	69
61	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
62	In-Depth Functional Diagnostics of Mouse Models by Single-Flash and Flicker Electroretinograms without Adapting Background Illumination. Advances in Experimental Medicine and Biology, 2016, 854, 619-625.	0.8	5
63	MonoallelicABCA4Mutations Appear Insufficient to Cause Retinopathy: A Quantitative Autofluorescence Study., 2015, 56, 8179.		38
64	Migraine and Vasospasm in Glaucoma: Age-Related Evaluation of 2027 Patients With Glaucoma or Ocular Hypertension., 2015, 56, 7999.		51
65	The genetics of age-related macular degeneration (AMD) – Novel targets for designing treatment options?. European Journal of Pharmaceutics and Biopharmaceutics, 2015, 95, 194-202.	2.0	33
66	Variation of the group 5 grass pollen allergen content of airborne pollen in relation to geographic location and time in season. Journal of Allergy and Clinical Immunology, 2015, 136, 87-95.e6.	1.5	155
67	Reticular Pseudodrusen in Sorsby FundusÂDystrophy. Ophthalmology, 2015, 122, 1555-1562.	2.5	58
68	Common synonymous variants in ABCA4 are protective for chloroquine induced maculopathy (toxic) Tj ETQq0 (O 0 rgBT /C	verlock 10 Tf
69	Sorsby Fundus Dystrophy: Novel Mutations, Novel Phenotypic Characteristics, and Treatment Outcomes., 2015, 56, 2664.		55
70	Synonymous variants in HTRA1 implicated in AMD susceptibility impair its capacity to regulate TGF- \hat{l}^2 signaling. Human Molecular Genetics, 2015, 24, 6361-6373.	1.4	42
71	Clinical and Genetic Factors Associated with Progression of Geographic Atrophy Lesions in Age-Related Macular Degeneration. PLoS ONE, 2015, 10, e0126636.	1.1	61
72	The Role of the Complement System in Age-Related Macular Degeneration. Deutsches Ärzteblatt International, 2014, 111, 133-8.	0.6	47

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73	A Novel Antibody against Human Properdin Inhibits the Alternative Complement System and Specifically Detects Properdin from Blood Samples. PLoS ONE, 2014, 9, e96371.	1.1	44
74	A Circulating MicroRNA Profile Is Associated with Late-Stage Neovascular Age-Related Macular Degeneration. PLoS ONE, 2014, 9, e107461.	1.1	62
75	Modelling the Genetic Risk in Age-Related Macular Degeneration. PLoS ONE, 2012, 7, e37979.	1.1	79
76	Release of Bet v 1 from birch pollen from 5 European countries. Results from the HIALINE study. Atmospheric Environment, 2012, 55, 496-505.	1.9	141