Panfeng Wang

List of Publications by Year
in descending order

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38	600	687220 13	713332 21
papers	citations	h-index	g-index

38	38	38	861
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Clinical and genetic features of retinoschisis in 120 families withRS1mutations. British Journal of Ophthalmology, 2023, 107, 367-372.	2.1	6
2	Landscape of pathogenic variants in six preâ€mRNA processing factor genes for retinitis pigmentosa based on large inâ€house data sets and database comparisons. Acta Ophthalmologica, 2022, , .	0.6	4
3	Autosomal Dominant Retinitis Pigmentosaâ€"Associated TOPORS Protein Truncating Variants Are Exclusively Located in the Region of Amino Acid Residues 807 to 867. , 2022, 63, 19.		1
4	Different Phenotypes Represent Advancing Stages of ABCA4-Associated Retinopathy: A Longitudinal Study of 212 Chinese Families From a Tertiary Center. , 2022, 63, 28.		6
5	Spectrum-frequency and genotypeâ€"phenotype analysis of rhodopsin variants. Experimental Eye Research, 2021, 203, 108405.	1.2	14
6	Clinical and Genetic Analysis of 63 Families Demonstrating Early and Advanced Characteristic Fundus as the Signature of CRB1 Mutations. American Journal of Ophthalmology, 2021, 223, 160-168.	1.7	10
7	Dominant RP in the Middle While Recessive in Both the N- and C-Terminals Due to RP1 Truncations: Confirmation, Refinement, and Questions. Frontiers in Cell and Developmental Biology, 2021, 9, 634478.	1.8	6
8	Pathogenicity evaluation and the genotypeâ€"phenotype analysis of OPA1 variants. Molecular Genetics and Genomics, 2021, 296, 845-862.	1.0	6
9	Pathogenic variants and associated phenotypic spectrum of TSPAN12 based on data from a large cohort. Graefe's Archive for Clinical and Experimental Ophthalmology, 2021, 259, 2929-2939.	1.0	7
10	Characterization of PROM1 p.Arg373Cys Variant in a Cohort of Chinese Patients: Macular Dystrophy Plus Peripheral Bone-Spicule Degeneration. , 2021, 62, 19.		6
11	An Early Diagnostic Clue for COL18A1- and LAMA1-Associated Diseases: High Myopia With Alopecia Areata in the Cranial Midline. Frontiers in Cell and Developmental Biology, 2021, 9, 644947.	1.8	1
12	Novel variants in GUCY2D causing retinopathy and the genotype-phenotype correlation. Experimental Eye Research, 2021, 208, 108637.	1.2	7
13	Severe exudative vitreoretinopathy as a common feature for CTNNB1, KIF11 and NDP variants plus sector degeneration for KIF11. American Journal of Ophthalmology, 2021, , .	1.7	2
14	Novel BMP4 Truncations Resulted in Opposite Ocular Anomalies: Pathologic Myopia Rather Than Microphthalmia. Frontiers in Cell and Developmental Biology, 2021, 9, 769636.	1.8	2
15	<scp>RPE</scp>65 mutation frequency and phenotypic variation according to exome sequencing in a tertiary centre for genetic eye diseases in China. Acta Ophthalmologica, 2020, 98, e181-e190.	0.6	21
16	Clinical manifestation and genetic analysis in Chinese early onset Xâ€linked retinoschisis. Molecular Genetics & Genomic Medicine, 2020, 8, e1421.	0.6	14
17	Structural variations in a non-coding region at 1q32.1 are responsible for the NYS7 locus in two large families. Human Genetics, 2020, 139, 1057-1064.	1.8	17
18	Genotypes and phenotypes of genes associated with achromatopsia: A reference for clinical genetic testing. Molecular Vision, 2020, 26, 588-602.	1.1	6

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19	Generation and Characterization of Induced Pluripotent Stem Cells and Retinal Organoids From a Leberâ€™s Congenital Amaurosis Patient With Novel RPE65 Mutations. Frontiers in Molecular Neuroscience, 2019, 12, 212.	1.4	30
20	CPSF1 mutations are associated with early-onset high myopia and involved in retinal ganglion cell axon projection. Human Molecular Genetics, 2019, 28, 1959-1970.	1.4	27
21	An Ophthalmic Targeted Exome Sequencing Panel as a Powerful Tool to Identify Causative Mutations in Patients Suspected of Hereditary Eye Diseases. Translational Vision Science and Technology, 2019, 8, 21.	1.1	54
22	Novel ocular findings in oculodentodigital dysplasia (ODDD): a case report and literature review. Ophthalmic Genetics, 2019, 40, 54-59.	0.5	5
23	Biallelic mutations in USP45, encoding a deubiquitinating enzyme, are associated with Leber congenital amaurosis. Journal of Medical Genetics, 2019, 56, 325-331.	1.5	16
24	Germline Mutations in CTNNB1 Associated With Syndromic FEVR or Norrie Disease. , 2019, 60, 93.		26
25	Mutation profile of glaucoma candidate genes in Mauritanian families with primary congenital glaucoma. Molecular Vision, 2019, 25, 373-381.	1.1	3
26	Phenotypic characterization of patients with early-onset high myopia due to mutations in or : Why not Stickler syndrome?. Molecular Vision, 2018, 24, 560-573.	1.1	15
27	Molecular genetics of cone-rod dystrophy in Chinese patients: New data from 61 probands and mutation overview of 163 probands. Experimental Eye Research, 2016, 146, 252-258.	1.2	60
28	Molecular genetics of Leber congenital amaurosis in Chinese: New data from 66 probands and mutation overview of 159 probands. Experimental Eye Research, 2016, 149, 93-99.	1.2	30
29	Exome sequencing reveals CHM mutations in six families with atypical choroideremia initially diagnosed as retinitis pigmentosa. International Journal of Molecular Medicine, 2014, 34, 573-577.	1.8	28
30	Cone-Rod Dysfunction Is a Sign of Early-Onset High Myopia. Optometry and Vision Science, 2013, 90, 1327-1330.	0.6	13
31	PAX6Mutations Identified in 4 of 35 Families with Microcornea. , 2012, 53, 6338.		14
32	KIF21A novel deletion and recurrent mutation in patients with congenital fibrosis of the extraocular muscles-1. International Journal of Molecular Medicine, 2011, 28, 973-5.	1.8	12
33	High Myopia Is Not Associated with the SNPs in theTGIF, Lumican,TGFB1, andHGFGenes. , 2009, 50, 1546.		43
34	Evaluation of MFRP as a candidate gene for high hyperopia. Molecular Vision, 2009, 15, 181-6.	1.1	11
35	An evaluation of OPTC and EPYC as candidate genes for high myopia. Molecular Vision, 2009, 15, 2045-9.	1.1	8
36	Novel SOX2 Mutation Associated With Ocular Coloboma in a Chinese Family. JAMA Ophthalmology, 2008, 126, 709.	2.6	37

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37	Linkage analysis of two families with X-linked recessive congenital motor nystagmus. Journal of Human Genetics, 2006, 51, 76-80.	1.1	20
38	Novel mutations of the PAX6 gene identified in Chinese patients with aniridia. Molecular Vision, 2006, 12, 644-8.	1.1	12