

# Lina Zelinger

## List of Publications by Year in descending order

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Version: 2024-02-01

20  
papers

1,029  
citations

623574

14  
h-index

887953

17  
g-index

20  
all docs

20  
docs citations

20  
times ranked

1783  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Missense Mutation in DHDDS, Encoding Dehydrodolichyl Diphosphate Synthase, Is Associated with Autosomal-Recessive Retinitis Pigmentosa in Ashkenazi Jews. <i>American Journal of Human Genetics</i> , 2011, 88, 207-215.	2.6	120
2	NRL-Regulated Transcriptome Dynamics of Developing Rod Photoreceptors. <i>Cell Reports</i> , 2016, 17, 2460-2473.	2.9	104
3	Mutations in IMPG2, Encoding Interphotoreceptor Matrix Proteoglycan 2, Cause Autosomal-Recessive Retinitis Pigmentosa. <i>American Journal of Human Genetics</i> , 2010, 87, 199-208.	2.6	98
4	Gene Augmentation Therapy Restores Retinal Function and Visual Behavior in a Sheep Model of CNGA3 Achromatopsia. <i>Molecular Therapy</i> , 2015, 23, 1423-1433.	3.7	93
5	Mutations in RAB28, Encoding a Farnesylated Small GTPase, Are Associated with Autosomal-Recessive Cone-Rod Dystrophy. <i>American Journal of Human Genetics</i> , 2013, 93, 110-117.	2.6	85
6	A homozygous nonsense CEP250 mutation combined with a heterozygous nonsense C2orf71 mutation is associated with atypical Usher syndrome. <i>Journal of Medical Genetics</i> , 2014, 51, 460-469.	1.5	78
7	Loss of the Metalloprotease ADAM9 Leads to Cone-Rod Dystrophy in Humans and Retinal Degeneration in Mice. <i>American Journal of Human Genetics</i> , 2009, 84, 683-691.	2.6	76
8	Mutations in ARL2BP, Encoding ADP-Ribosylation-Factor-Like 2 Binding Protein, Cause Autosomal-Recessive Retinitis Pigmentosa. <i>American Journal of Human Genetics</i> , 2013, 93, 321-329.	2.6	67
9	Genetics and Disease Expression in the CNGA3 Form of Achromatopsia. <i>Ophthalmology</i> , 2015, 122, 997-1007.	2.5	61
10	Identification of Mutations Causing Inherited Retinal Degenerations in the Israeli and Palestinian Populations Using Homozygosity Mapping. , 2014, 55, 1149.		46
11	Whole Exome Sequencing Reveals GUCY2D as a Major Gene Associated With Cone and Cone-Rod Dystrophy in Israel. <i>Investigative Ophthalmology and Visual Science</i> , 2015, 56, 420-430.	3.3	32
12	RNA Biology in Retinal Development and Disease. <i>Trends in Genetics</i> , 2018, 34, 341-351.	2.9	29
13	Exome Sequencing Identifies a Founder Frameshift Mutation in an Alternative Exon of USH1C as the Cause of Autosomal Recessive Retinitis Pigmentosa with Late-Onset Hearing Loss. <i>PLoS ONE</i> , 2012, 7, e51566.	1.1	27
14	Mutations in <i>CRB1</i> are a Relatively Common Cause of Autosomal Recessive Early-Onset Retinal Degeneration in the Israeli and Palestinian Populations. , 2013, 54, 2068.		25
15	Cone Dystrophy with Supernormal Rod Response. <i>Ophthalmology</i> , 2013, 120, 2338-2343.	2.5	23
16	Regulation of Noncoding Transcriptome in Developing Photoreceptors by Rod Differentiation Factor NRL. , 2017, 58, 4422.		19
17	Nonsyndromic Early-Onset Cone-Rod Dystrophy and Limb-Girdle Muscular Dystrophy in a Consanguineous Israeli Family are Caused by Two Independent yet Linked Mutations in <i>ALMS1</i> and <i>DYSF</i> . <i>Human Mutation</i> , 2015, 36, 836-841.	1.1	17
18	TRPM1 Mutations are the Most Common Cause of Autosomal Recessive Congenital Stationary Night Blindness (CSNB) in the Palestinian and Israeli Populations. <i>Scientific Reports</i> , 2019, 9, 12047.	1.6	14

#	ARTICLE	IF	CITATIONS
19	An ancient autosomal haplotype bearing a rare achromatopsia-causing founder mutation is shared among Arab Muslims and Oriental Jews. <i>Human Genetics</i> , 2010, 128, 261-267.	1.8	12
20	SSBP1 faux pas in mitonuclear tango causes optic neuropathy. <i>Journal of Clinical Investigation</i> , 2019, 130, 62-64.	3.9	3