

Alice E Davidson

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

949
citations

516710

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610901

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29
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docs citations

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times ranked

1564
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Retinal Structure and Function in Achromatopsia. <i>Ophthalmology</i> , 2014, 121, 234-245. | 5.2 | 145 |
| 2 | Clinical and Molecular Analysis of Stargardt Disease With Preserved Foveal Structure and Function. <i>American Journal of Ophthalmology</i> , 2013, 156, 487-501.e1. | 3.3 | 100 |
| 3 | Autosomal-Dominant Corneal Endothelial Dystrophies CHED1 and PPCD1 Are Allelic Disorders Caused by Non-coding Mutations in the Promoter of OVOL2. <i>American Journal of Human Genetics</i> , 2016, 98, 75-89. | 6.2 | 70 |
| 4 | 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. | 6.2 | 67 |
| 5 | Antisense Therapy for a Common Corneal Dystrophy Ameliorates TCF4 Repeat Expansion-Mediated Toxicity. <i>American Journal of Human Genetics</i> , 2018, 102, 528-539. | 6.2 | 59 |
| 6 | CRISPR/Cas9-targeted enrichment and long-read sequencing of the Fuchs endothelial corneal dystrophy-associated TCF4 triplet repeat. <i>Genetics in Medicine</i> , 2019, 21, 2092-2102. | 2.4 | 56 |
| 7 | Mutations in CPAMD8 Cause a Unique Form of Autosomal-Recessive Anterior Segment Dysgenesis. <i>American Journal of Human Genetics</i> , 2016, 99, 1338-1352. | 6.2 | 47 |
| 8 | Ectopic GRHL2 Expression Due to Non-coding Mutations Promotes Cell State Transition and Causes Posterior Polymorphous Corneal Dystrophy 4. <i>American Journal of Human Genetics</i> , 2018, 102, 447-459. | 6.2 | 45 |
| 9 | Genetic Variants Associated With Corneal Biomechanical Properties and Potentially Conferring Susceptibility to Keratoconus in a Genome-Wide Association Study. <i>JAMA Ophthalmology</i> , 2019, 137, 1005. | 2.5 | 45 |
| 10 | TCF4-mediated Fuchs endothelial corneal dystrophy: Insights into a common trinucleotide repeat-associated disease. <i>Progress in Retinal and Eye Research</i> , 2021, 81, 100883. | 15.5 | 40 |
| 11 | A multi-ethnic genome-wide association study implicates collagen matrix integrity and cell differentiation pathways in keratoconus. <i>Communications Biology</i> , 2021, 4, 266. | 4.4 | 36 |
| 12 | Brittle Cornea Syndrome ZNF469 Mutation Carrier Phenotype and Segregation Analysis of Rare ZNF469 Variants in Familial Keratoconus. <i>Investigative Ophthalmology and Visual Science</i> , 2015, 56, 578-586. | 3.3 | 33 |
| 13 | Heterozygous deletions at the ZEB1 locus verify haploinsufficiency as the mechanism of disease for posterior polymorphous corneal dystrophy type 3. <i>European Journal of Human Genetics</i> , 2016, 24, 985-991. | 2.8 | 33 |
| 14 | Screening of a Large Cohort of Leber Congenital Amaurosis and Retinitis Pigmentosa Patients Identifies Novel <i>LCA5</i> Mutations and New Genotype-Phenotype Correlations. <i>Human Mutation</i> , 2013, 34, 1537-1546. | 2.5 | 32 |
| 15 | Identification of Six Novel Mutations in <i>ZEB1</i> and Description of the Associated Phenotypes in Patients with Posterior Polymorphous Corneal Dystrophy 3. <i>Annals of Human Genetics</i> , 2015, 79, 1-9. | 0.8 | 29 |
| 16 | Association of <i>CHRD1</i> Mutations and Variants with X-linked Megalocornea, Neuhäuser Syndrome and Central Corneal Thickness. <i>PLoS ONE</i> , 2014, 9, e104163. | 2.5 | 27 |
| 17 | Genotype-Phenotype Correlation for <i>TGFBI</i> Corneal Dystrophies Identifies p.(G623D) as a Novel Cause of Epithelial Basement Membrane Dystrophy. , 2016, 57, 5407. | | 22 |
| 18 | IPSC-Derived Corneal Endothelial-like Cells Act as an Appropriate Model System to Assess the Impact of <i>SLC4A11</i> Variants on Pre-mRNA Splicing. , 2019, 60, 3084. | | 18 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Machine Learning Algorithms to Detect Subclinical Keratoconus: Systematic Review. <i>JMIR Medical Informatics</i> , 2021, 9, e27363. | 2.6 | 12 |
| 20 | A new paradigm for delivering personalised care: integrating genetics with surgical interventions in BEST1 mutations. <i>Eye</i> , 2020, 34, 577-583. | 2.1 | 11 |
| 21 | The utility of massively parallel sequencing for posterior polymorphous corneal dystrophy type 3 molecular diagnosis. <i>Experimental Eye Research</i> , 2019, 182, 160-166. | 2.6 | 8 |
| 22 | CUGC for posterior polymorphous corneal dystrophy (PPCD). <i>European Journal of Human Genetics</i> , 2020, 28, 126-131. | 2.8 | 4 |
| 23 | Coincidental Occurrence of Schnyder Corneal Dystrophy and Posterior Polymorphous Corneal Dystrophy Type 3. <i>Cornea</i> , 2019, 38, 758-760. | 1.7 | 3 |
| 24 | Non-Penetrance for Ocular Phenotype in Two Individuals Carrying Heterozygous Loss-of-Function ZEB1 Alleles. <i>Genes</i> , 2021, 12, 677. | 2.4 | 3 |
| 25 | Posterior corneal vesicles are not associated with the genetic variants that cause posterior polymorphous corneal dystrophy. <i>Acta Ophthalmologica</i> , 2022, 100, . | 1.1 | 3 |
| 26 | Novel disease-causing variants and phenotypic features of X-linked megalocornea. <i>Acta Ophthalmologica</i> , 2021, , . | 1.1 | 1 |
| 27 | Should Patients with Kearns-Sayre Syndrome and Corneal Endothelial Failure Be Genotyped for a TCF4 Trinucleotide Repeat, Commonly Associated with Fuchs Endothelial Corneal Dystrophy?. <i>Genes</i> , 2021, 12, 1918. | 2.4 | 0 |