

David Valle

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5535175/publications.pdf>

Version: 2024-02-01

79
papers

7,145
citations

117625

34
h-index

76900

74
g-index

82
all docs

82
docs citations

82
times ranked

12522
citing authors

#	ARTICLE	IF	CITATIONS
1	Centers for Mendelian Genomics: A decade of facilitating gene discovery. <i>Genetics in Medicine</i> , 2022, 24, 784-797.	2.4	44
2	Variant-level matching for diagnosis and discovery: Challenges and opportunities. <i>Human Mutation</i> , 2022, , .	2.5	11
3	The impact of GeneMatcher on international data sharing and collaboration. <i>Human Mutation</i> , 2022, , .	2.5	7
4	A promoter variant in the <i>OTC</i> gene associated with late and variable age of onset hyperammonemia. <i>Journal of Inherited Metabolic Disease</i> , 2022, 45, 710-718.	3.6	5
5	IFIH1 loss-of-function variants contribute to very early-onset inflammatory bowel disease. <i>Human Genetics</i> , 2021, 140, 1299-1312.	3.8	17
6	Victor McKusick and his short course. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 3242-3252.	1.2	1
7	PhenoDB, GeneMatcher and VariantMatcher, tools for analysis and sharing of sequence data. <i>Orphanet Journal of Rare Diseases</i> , 2021, 16, 365.	2.7	24
8	Response to Biesecker et al.. <i>American Journal of Human Genetics</i> , 2021, 108, 1807-1808.	6.2	3
9	The utility of exome sequencing for fetal pleural effusions. <i>Prenatal Diagnosis</i> , 2020, 40, 590-595.	2.3	9
10	The Deep Genome Project. <i>Genome Biology</i> , 2020, 21, 18.	8.8	30
11	Genetic diagnosis in first or second trimester pregnancy loss using exome sequencing: a systematic review of human essential genes. <i>Journal of Assisted Reproduction and Genetics</i> , 2019, 36, 1539-1548.	2.5	25
12	Bioinformatics-Based Identification of Expanded Repeats: A Non-reference Intronic Pentamer Expansion in RFC1 Causes CANVAS. <i>American Journal of Human Genetics</i> , 2019, 105, 151-165.	6.2	170
13	2018 Victor A. McKusick Leadership Award Introduction: James R. Lupski. <i>American Journal of Human Genetics</i> , 2019, 104, 389-390.	6.2	0
14	Insights into genetics, human biology and disease gleaned from family based genomic studies. <i>Genetics in Medicine</i> , 2019, 21, 798-812.	2.4	161
15	TBX6-associated congenital scoliosis (TACS) as a clinically distinguishable subtype of congenital scoliosis: further evidence supporting the compound inheritance and TBX6 gene dosage model. <i>Genetics in Medicine</i> , 2019, 21, 1548-1558.	2.4	60
16	Familial monophasic acute transverse myelitis due to the pathogenic variant in <i>VPS37A</i> . <i>Neurology: Genetics</i> , 2018, 4, e213.	1.9	4
17	Defect in phosphoinositide signalling through a homozygous variant in <i>PLCB3</i> causes a new form of spondylometaphyseal dysplasia with corneal dystrophy. <i>Journal of Medical Genetics</i> , 2018, 55, 122-130.	3.2	9
18	Identification of <i>STAC3</i> variants in non-Native American families with overlapping features of Carey-Fineman-Ziter syndrome and Moebius syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2763-2771.	1.2	28

#	ARTICLE	IF	CITATIONS
19	Thorase variants are associated with defects in glutamatergic neurotransmission that can be rescued by Perampanel. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	20
20	Patients with a Kabuki syndrome phenotype demonstrate DNA methylation abnormalities. <i>European Journal of Human Genetics</i> , 2017, 25, 1335-1344.	2.8	52
21	LMNA Sequences of 60,706 Unrelated Individuals Reveal 132 Novel Missense Variants in A-Type Lamins and Suggest a Link between Variant p.G602S and Type 2 Diabetes. <i>Frontiers in Genetics</i> , 2017, 8, 79.	2.3	17
22	A Syndromic Intellectual Disability Disorder Caused by Variants in <i>TELO2</i> , a Gene Encoding a Component of the TTT Complex. <i>American Journal of Human Genetics</i> , 2016, 98, 909-918.	6.2	35
23	Gonadal mosaicism in <i>ARID1B</i> gene causes intellectual disability and dysmorphic features in three siblings. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 156-161.	1.2	16
24	Disorders of Ornithine and Proline Metabolism. , 2016, , 321-331.		4
25	Lessons learned from the search for genes responsible for rare Mendelian disorders. <i>Molecular Genetics & Genomic Medicine</i> , 2016, 4, 371-375.	1.2	7
26	Neuregulin 3 Knockout Mice Exhibit Behaviors Consistent with Psychotic Disorders. <i>Molecular Neuropsychiatry</i> , 2016, 2, 79-87.	2.9	27
27	The management of pregnancy and delivery in 3-hydroxy-3-methylglutaryl-CoA lyase deficiency. , 2016, 170, 1600-1602.		7
28	GeneMatcher Aids in the Identification of a New Malformation Syndrome with Intellectual Disability, Unique Facial Dysmorphisms, and Skeletal and Connective Tissue Abnormalities Caused by De Novo Variants in <i>HNRNPK</i> . <i>Human Mutation</i> , 2015, 36, 1009-1014.	2.5	56
29	GeneMatcher: A Matching Tool for Connecting Investigators with an Interest in the Same Gene. <i>Human Mutation</i> , 2015, 36, 928-930.	2.5	1,153
30	Identification and Functional Studies of Regulatory Variants Responsible for the Association of <i>NRG3</i> with a Delusion Phenotype in Schizophrenia. <i>Molecular Neuropsychiatry</i> , 2015, 1, 36-46.	2.9	14
31	New Arab family with cerebral dysgenesis, neuropathy, ichthyosis and keratoderma syndrome suggests a possible founder effect for the c.223delG mutation. <i>Journal of Dermatology</i> , 2015, 42, 821-822.	1.2	8
32	Functional Variants in <i>DPYSL2</i> Sequence Increase Risk of Schizophrenia and Suggest a Link to mTOR Signaling. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 61-72.	1.8	39
33	Assessment of incidental findings in 232 whole-exome sequences from the Baylorâ€“Hopkins Center for Mendelian Genomics. <i>Genetics in Medicine</i> , 2015, 17, 782-788.	2.4	41
34	New Tools for Mendelian Disease Gene Identification: PhenoDB Variant Analysis Module; and GeneMatcher, a Web-Based Tool for Linking Investigators with an Interest in the Same Gene. <i>Human Mutation</i> , 2015, 36, 425-431.	2.5	141
35	The Genetic Basis of Mendelian Phenotypes: Discoveries, Challenges, and Opportunities. <i>American Journal of Human Genetics</i> , 2015, 97, 199-215.	6.2	574
36	2014 Victor A. McKusick Leadership Award1. <i>American Journal of Human Genetics</i> , 2015, 96, 374-376.	6.2	0

#	ARTICLE	IF	CITATIONS
37	A novel bile acid biosynthesis defect due to a deficiency of peroxisomal ABCD3. <i>Human Molecular Genetics</i> , 2015, 24, 361-370.	2.9	115
38	Reduced PLP2 expression increases ER-stress-induced neuronal apoptosis and risk for adverse neurological outcomes after hypoxia ischemia injury. <i>Human Molecular Genetics</i> , 2015, 24, 7221-7226.	2.9	16
39	Enhanced conversion of induced neuronal cells (iN cells) from human fibroblasts: Utility in uncovering cellular deficits in mental illness-associated chromosomal abnormalities. <i>Neuroscience Research</i> , 2015, 101, 57-61.	1.9	14
40	Novel Deletion of <i>SERPINF1</i> Causes Autosomal Recessive Osteogenesis Imperfecta Type VI in Two Brazilian Families. <i>Molecular Syndromology</i> , 2014, 5, 268-275.	0.8	18
41	Mutations in <i>AlstrÅm</i> protein impair terminal differentiation of cardiomyocytes. <i>Nature Communications</i> , 2014, 5, 3416.	12.8	66
42	Mutations in <i>PCYT1A</i> , Encoding a Key Regulator of Phosphatidylcholine Metabolism, Cause Spondylometaphyseal Dysplasia with Cone-Rod Dystrophy. <i>American Journal of Human Genetics</i> , 2014, 94, 105-112.	6.2	53
43	<i>henoDB</i> : A New Web-Based Tool for the Collection, Storage, and Analysis of Phenotypic Features. <i>Human Mutation</i> , 2013, 34, 566-571.	2.5	64
44	Linkage and association on 8p21.2-p21.1 in schizophrenia. , 2011, 156, 188-197.		26
45	<i>PHR1</i> is a vesicle-bound protein abundantly expressed in mature olfactory neurons. <i>Laryngoscope</i> , 2010, 120, 1002-1010.	2.0	1
46	Familiality of Novel Factorial Dimensions of Schizophrenia. <i>Archives of General Psychiatry</i> , 2009, 66, 591.	12.3	71
47	Fine Mapping on Chromosome 10q22-q23 Implicates Neuregulin 3 in Schizophrenia. <i>American Journal of Human Genetics</i> , 2009, 84, 21-34.	6.2	81
48	Functional genomics and SNP analysis of human genes encoding proline metabolic enzymes. <i>Amino Acids</i> , 2008, 35, 655-664.	2.7	46
49	<i>PRODH</i> variants and risk for schizophrenia. <i>Amino Acids</i> , 2008, 35, 673-679.	2.7	54
50	X chromosome cDNA microarray screening identifies a functional PLP2 promoter polymorphism enriched in patients with X-linked mental retardation. <i>Genome Research</i> , 2007, 17, 641-648.	5.5	16
51	Recurrent 10q22-q23 Deletions: A Genomic Disorder on 10q Associated with Cognitive and Behavioral Abnormalities. <i>American Journal of Human Genetics</i> , 2007, 80, 938-947.	6.2	101
52	Overexpression of proline oxidase induces proline-dependent and mitochondria-mediated apoptosis. <i>Molecular and Cellular Biochemistry</i> , 2007, 295, 85-92.	3.1	98
53	A microdeletion in Xp11.3 accounts for co-segregation of retinitis pigmentosa and mental retardation in a large kindred. <i>American Journal of Medical Genetics, Part A</i> , 2006, 140A, 349-357.	1.2	23
54	Functional Consequences of <i>PRODH</i> Missense Mutations. <i>American Journal of Human Genetics</i> , 2005, 76, 409-420.	6.2	134

#	ARTICLE	IF	CITATIONS
55	Bipolar I Disorder and Schizophrenia: A Single-Nucleotide Polymorphism Screen of 64 Candidate Genes among Ashkenazi Jewish Case-Parent Trios. <i>American Journal of Human Genetics</i> , 2005, 77, 918-936.	6.2	358
56	A SCIENCE OF THE INDIVIDUAL: Implications for a Medical School Curriculum. <i>Annual Review of Genomics and Human Genetics</i> , 2005, 6, 313-330.	6.2	34
57	PHR1, a PH Domain-Containing Protein Expressed in Primary Sensory Neurons. <i>Molecular and Cellular Biology</i> , 2004, 24, 9137-9151.	2.3	16
58	Genetics, Individuality, and Medicine in the 21st Century**Previously presented at the annual meeting of The American Society of Human Genetics, in Los Angeles, on November 5, 2003.. <i>American Journal of Human Genetics</i> , 2004, 74, 374-381.	6.2	20
59	Genomewide Linkage Scan for Schizophrenia Susceptibility Loci among Ashkenazi Jewish Families Shows Evidence of Linkage on Chromosome 10q22. <i>American Journal of Human Genetics</i> , 2003, 73, 601-611.	6.2	99
60	Introductory Speech for Charles Scriver**Amended from the introduction previously presented at the annual meeting of The American Society of Human Genetics, in San Diego, on October 16, 2001.. <i>American Journal of Human Genetics</i> , 2002, 70, 314-316.	6.2	0
61	Online Mendelian Inheritance In Man (OMIM). <i>Human Mutation</i> , 2000, 15, 57-61.	2.5	503
62	Genetics, Biology and Disease. <i>Annual Review of Genomics and Human Genetics</i> , 2000, 1, 1-19.	6.2	29
63	PHR1 Encodes an Abundant, Pleckstrin Homology Domain-containing Integral Membrane Protein in the Photoreceptor Outer Segments. <i>Journal of Biological Chemistry</i> , 1999, 274, 35676-35685.	3.4	18
64	40 years of the annual Bar Harbor Course™ (1960-1999): a pictorial history. <i>Clinical Genetics</i> , 1999, 55, 398-415.	2.0	6
65	You give me fever. <i>Nature Genetics</i> , 1999, 22, 121-122.	21.4	12
66	Hyperornithinaemia-Hyperammonaemia-Homocitrullinuria syndrome is caused by mutations in a gene encoding a mitochondrial ornithine transporter. <i>Nature Genetics</i> , 1999, 22, 151-158.	21.4	184
67	Gyrate Atrophy of the Choroid and Retina: Lymphocyte Ornithine-Î-Aminotransferase Activity in Different Mutations and Carriers. <i>Pediatric Research</i> , 1998, 44, 381-385.	2.3	18
68	Isolation of the human PEX12 gene, mutated in group 3 of the peroxisome biogenesis disorders. <i>Nature Genetics</i> , 1997, 15, 385-388.	21.4	142
69	Human PEX7 encodes the peroxisomal PTS2 receptor and is responsible for rhizomelic chondrodysplasia punctata. <i>Nature Genetics</i> , 1997, 15, 369-376.	21.4	415
70	Mutations in PEX1 are the most common cause of peroxisome biogenesis disorders. <i>Nature Genetics</i> , 1997, 17, 445-448.	21.4	233
71	A ubiquitin C-terminal hydrolase gene on the proximal short arm of the X chromosome: implications for X-linked retinal disorders. <i>Human Molecular Genetics</i> , 1996, 5, 533-538.	2.9	31
72	Mutations in the PTS1 receptor gene, PXR1, define complementation group 2 of the peroxisome biogenesis disorders. <i>Nature Genetics</i> , 1995, 9, 115-125.	21.4	432

#	ARTICLE	IF	CITATIONS
73	Mice lacking ornithineâ€“â€“ aminoâ€“transferase have paradoxical neonatal hypoorithinaemia and retinal degeneration. <i>Nature Genetics</i> , 1995, 11, 185-190.	21.4	106
74	Transcriptional regulation of the gene for the second component of human complement: Promoter analysis. <i>European Journal of Immunology</i> , 1994, 24, 393-400.	2.9	5
75	Penetrating the peroxisome. <i>Nature</i> , 1993, 361, 682-683.	27.8	73
76	Mutations in the 70K peroxisomal membrane protein gene in Zellweger syndrome. <i>Nature Genetics</i> , 1992, 1, 16-23.	21.4	238
77	Chronic cardiomyopathy and weakness or acute coma in children with a defect in carnitine uptake. <i>Annals of Neurology</i> , 1991, 30, 709-716.	5.3	206
78	Allopurinol-Induced Orotidinuria. <i>New England Journal of Medicine</i> , 1990, 322, 1641-1645.	27.0	124
79	Proline oxidase in cultured mammalian cells. <i>Journal of Cellular Physiology</i> , 1977, 91, 369-376.	4.1	25