Martin Zenker

List of Publications by Year in descending order

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71102 74163 6,379 108 41 75 citations h-index g-index papers 116 116 116 10105 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	A Single-Gene Cause in 29.5% of Cases of Steroid-Resistant Nephrotic Syndrome. Journal of the American Society of Nephrology: JASN, 2015, 26, 1279-1289.	6.1	499
2	Identifying facial phenotypes of genetic disorders using deep learning. Nature Medicine, 2019, 25, 60-64.	30.7	449
3	Mutation of SHOC2 promotes aberrant protein N-myristoylation and causes Noonan-like syndrome with loose anagen hair. Nature Genetics, 2009, 41, 1022-1026.	21.4	358
4	Noonan syndrome and clinically related disorders. Best Practice and Research in Clinical Endocrinology and Metabolism, 2011, 25, 161-179.	4.7	303
5	Deficiency of UBR1, a ubiquitin ligase of the N-end rule pathway, causes pancreatic dysfunction, malformations and mental retardation (Johanson-Blizzard syndrome). Nature Genetics, 2005, 37, 1345-1350.	21.4	252
6	Childhood cancer predisposition syndromes—A concise review and recommendations by the Cancer Predisposition Working Group of the Society for Pediatric Oncology and Hematology. American Journal of Medical Genetics, Part A, 2017, 173, 1017-1037.	1.2	200
7	Human pluripotent stem cell-derived acinar/ductal organoids generate human pancreas upon orthotopic transplantation and allow disease modelling. Gut, 2017, 66, 473-486.	12.1	174
8	Expansion of the genotypic and phenotypic spectrum in patients with KRAS germline mutations. Journal of Medical Genetics, 2006, 44, 131-135.	3.2	170
9	Cancer spectrum and frequency among children with Noonan, Costello, and cardio-facio-cutaneous syndromes. British Journal of Cancer, 2015, 112, 1392-1397.	6.4	167
10	Mutations in sphingosine-1-phosphate lyase cause nephrosis with ichthyosis and adrenal insufficiency. Journal of Clinical Investigation, 2017, 127, 912-928.	8.2	160
11	Functional Dysregulation of CDC42 Causes Diverse Developmental Phenotypes. American Journal of Human Genetics, 2018, 102, 309-320.	6.2	138
12	The landscape of epilepsy-related GATOR1 variants. Genetics in Medicine, 2019, 21, 398-408.	2.4	137
13	Autosomal recessive Noonan syndrome associated with biallelic LZTR1 variants. Genetics in Medicine, 2018, 20, 1175-1185.	2.4	133
14	ClinGen's RASopathy Expert Panel consensus methods for variant interpretation. Genetics in Medicine, 2018, 20, 1334-1345.	2.4	126
15	SOS1 is the second most common Noonan gene but plays no major role in cardio-facio-cutaneous syndrome. Journal of Medical Genetics, 2007, 44, 651-656.	3.2	114
16	Juvenile myelomonocytic leukaemia and Noonan syndrome. Journal of Medical Genetics, 2014, 51, 689-697.	3.2	112
17	Neu-Laxova Syndrome Is a Heterogeneous Metabolic Disorder Caused by Defects in Enzymes of the L-Serine Biosynthesis Pathway. American Journal of Human Genetics, 2014, 95, 285-293.	6.2	110
18	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. Genetics in Medicine, 2018, 20, 630-638.	2.4	101

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19	Gain-of-Function Mutations of ARHGAP31, a Cdc42/Rac1 GTPase Regulator, Cause Syndromic Cutis Aplasia and Limb Anomalies. American Journal of Human Genetics, 2011, 88, 574-585.	6.2	100
20	Hypertrophic Cardiomyopathy in Noonan Syndrome TreatedÂbyÂMEK-Inhibition. Journal of the American College of Cardiology, 2019, 73, 2237-2239.	2.8	96
21	The spectra of clinical phenotypes in aplasia cutis congenita and terminal transverse limb defects. American Journal of Medical Genetics, Part A, 2009, 149A, 1860-1881.	1.2	92
22	Haploinsufficiency of the NOTCH1 Receptor as a Cause of Adams–Oliver Syndrome With Variable Cardiac Anomalies. Circulation: Cardiovascular Genetics, 2015, 8, 572-581.	5.1	84
23	Fucosyltransferase 2 (FUT2) non-secretor status and blood group B are associated with elevated serum lipase activity in asymptomatic subjects, and an increased risk for chronic pancreatitis: a genetic association study. Gut, 2015, 64, 646-656.	12.1	82
24	Clinical manifestations of mutations in RAS and related intracellular signal transduction factors. Current Opinion in Pediatrics, 2011, 23, 443-451.	2.0	81
25	Mutations in <i>GRIP1</i> cause Fraser syndrome. Journal of Medical Genetics, 2012, 49, 303-306.	3.2	79
26	Genotype and phenotype in patients with Noonan syndrome and a RIT1 mutation. Genetics in Medicine, 2016, 18, 1226-1234.	2.4	77
27	Heterozygous Loss-of-Function Mutations in DLL4 Cause Adams-Oliver Syndrome. American Journal of Human Genetics, 2015, 97, 475-482.	6.2	73
28	Cardioâ€facioâ€cutaneous syndrome: Does genotype predict phenotype?. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2011, 157, 129-135.	1.6	72
29	Activating Mutations Affecting the Dbl Homology Domain of SOS2 Cause Noonan Syndrome. Human Mutation, 2015, 36, 1080-1087.	2.5	67
30	Assessing the gene–disease association of 19 genes with the RASopathies using the ClinGen gene curation framework. Human Mutation, 2018, 39, 1485-1493.	2.5	66
31	Manitoba-oculo-tricho-anal (MOTA) syndrome is caused by mutations in FREM1. Journal of Medical Genetics, 2011, 48, 375-382.	3.2	60
32	A Dual Phenotype of Periventricular Nodular Heterotopia and Frontometaphyseal Dysplasia in One Patient Caused by a Single FLNA Mutation Leading to Two Functionally Different Aberrant Transcripts. American Journal of Human Genetics, 2004, 74, 731-737.	6.2	55
33	Specific mosaic <i><scp>KRAS</scp></i> mutations affecting codon 146 cause oculoectodermal syndrome and encephalocraniocutaneous lipomatosis. Clinical Genetics, 2016, 90, 334-342.	2.0	55
34	Rapidly progressive hypertrophic cardiomyopathy in an infant with Noonan syndrome with multiple lentigines: Palliative treatment with a rapamycin analog. American Journal of Medical Genetics, Part A, 2015, 167, 744-751.	1.2	53
35	Genetic screening confirms heterozygous mutations in ACAN as a major cause of idiopathic short stature. Scientific Reports, 2017, 7, 12225.	3.3	53
36	Genetics of nephrotic syndrome: new insights into molecules acting at the glomerular filtration barrier. Journal of Molecular Medicine, 2009, 87, 849-857.	3.9	52

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37	Genetic and Pathogenetic Aspects of Noonan Syndrome and Related Disorders. Hormone Research, 2009, 72, 57-63.	1.8	51
38	Synaptic activity controls localization and function of Ct <scp>BP</scp> 1 via binding to <scp>B</scp> assoon and <scp>P</scp> iccolo. EMBO Journal, 2015, 34, 1056-1077.	7.8	51
39	Enhanced MAPK1 Function Causes a Neurodevelopmental Disorder within the RASopathy Clinical Spectrum. American Journal of Human Genetics, 2020, 107, 499-513.	6.2	48
40	NEMF mutations that impair ribosome-associated quality control are associated with neuromuscular disease. Nature Communications, 2020, 11, 4625.	12.8	47
41	A variable combination of features of Noonan syndrome and neurofibromatosis type I are caused by mutations in the <i>NF1</i> gene. American Journal of Medical Genetics, Part A, 2006, 140A, 2749-2756.	1.2	46
42	Genetic Basis and Pancreatic Biology of Johanson-Blizzard Syndrome. Endocrinology and Metabolism Clinics of North America, 2006, 35, 243-253.	3.2	44
43	Phenotype and natural history in Marshall–Smith syndrome. American Journal of Medical Genetics, Part A, 2010, 152A, 2714-2726.	1.2	44
44	Activating Mutations of RRAS2 Are a Rare Cause of Noonan Syndrome. American Journal of Human Genetics, 2019, 104, 1223-1232.	6.2	43
45	<i>WDR73</i> Mutations Cause Infantile Neurodegeneration and Variable Glomerular Kidney Disease. Human Mutation, 2015, 36, 1021-1028.	2.5	42
46	Further delineation of Malan syndrome. Human Mutation, 2018, 39, 1226-1237.	2.5	42
47	NFIB Haploinsufficiency Is Associated with Intellectual Disability and Macrocephaly. American Journal of Human Genetics, 2018, 103, 752-768.	6.2	40
48	Foodâ€Derived Xenoâ€microRNAs: Influence of Diet and Detectability in Gastrointestinal Tractâ€"Proofâ€ofâ€Principle Study. Molecular Nutrition and Food Research, 2019, 63, e1800076.	3.3	40
49	Expansion of the phenotypic spectrum and description of molecular findings in a cohort of patients with oculocutaneous mosaic RASopathies. Molecular Genetics & Expansion of the phenotypic spectrum and description of molecular findings in a cohort of patients	1.2	39
50	bFGF-mediated pluripotency maintenance in human induced pluripotent stem cells is associated with NRAS-MAPK signaling. Cell Communication and Signaling, 2018, 16, 96.	6.5	38
51	Genotype and phenotype spectrum of NRAS germline variants. European Journal of Human Genetics, 2017, 25, 823-831.	2.8	36
52	CTLA-4-mediated posttranslational modifications direct cytotoxic T-lymphocyte differentiation. Cell Death and Differentiation, 2017, 24, 1739-1749.	11.2	36
53	Variants in nuclear factor I genes influence growth and development. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2019, 181, 611-626.	1.6	32
54	Elucidating the genetic architecture of Adams-Oliver syndrome in a large European cohort. Human Mutation, 2018, 39, 1246-1261.	2.5	31

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55	Activating MRAS mutations cause Noonan syndrome associated with hypertrophic cardiomyopathy. Human Molecular Genetics, 2020, 29, 1772-1783.	2.9	30
56	Noonan syndrome: improving recognition and diagnosis. Archives of Disease in Childhood, 2022, 107, 1073-1078.	1.9	28
57	Deletions in the 3′ Part of the <i>NFIX</i> Gene Including a Recurrent Alu-Mediated Deletion of Exon 6 and 7 Account for Previously Unexplained Cases of Marshall-Smith Syndrome. Human Mutation, 2014, 35, 1092-1100.	2.5	26
58	Clinical and molecular analysis of RASopathies in a group of Turkish patients. Clinical Genetics, 2013, 83, 181-186.	2.0	24
59	SPRED2 loss-of-function causes a recessive Noonan syndrome-like phenotype. American Journal of Human Genetics, 2021, 108, 2112-2129.	6.2	23
60	Genetic variation of the RASGRF1 regulatory region affects human hippocampus-dependent memory. Frontiers in Human Neuroscience, 2014, 8, 260.	2.0	22
61	Valenced action/inhibition learning in humans is modulated by a genetic variant linked to dopamine D2 receptor expression. Frontiers in Systems Neuroscience, 2014, 8, 140.	2.5	22
62	Genetic basis of hypertrophic cardiomyopathy in children. Clinical Research in Cardiology, 2019, 108, 282-289.	3.3	22
63	Responsiveness of sphingosine phosphate lyase insufficiency syndrome to vitamin <scp>B6</scp> cofactor supplementation. Journal of Inherited Metabolic Disease, 2020, 43, 1131-1142.	3.6	21
64	Surgery in Focal Congenital Hyperinsulinism (CHI) - The "Hyperinsulinism Germany International" Experience in 30 Children. Pediatric Endocrinology Reviews, 2016, 14, 129-137.	1.2	20
65	Dealing with the incidental finding of secondary variants by the example of SRNS patients undergoing targeted next-generation sequencing. Pediatric Nephrology, 2016, 31, 73-81.	1.7	19
66	MRI Spectrum of Brain Involvement in Sphingosine-1-Phosphate Lyase Insufficiency Syndrome. American Journal of Neuroradiology, 2020, 41, 1943-1948.	2.4	19
67	Germline PTPN11 and somatic PIK3CA variant in a boy with megalencephaly-capillary malformation syndrome (MCAP) - pure coincidence?. European Journal of Human Genetics, 2015, 23, 409-412.	2.8	17
68	Adams–Oliver syndrome caused by mutations of the <i>EOGT</i> gene. American Journal of Medical Genetics, Part A, 2019, 179, 2246-2251.	1.2	17
69	Variants of SOS2 are a rare cause of Noonan syndrome with particular predisposition for lymphatic complications. European Journal of Human Genetics, 2021, 29, 51-60.	2.8	17
70	Long-term outcomes of childhood onset Noonan compared to sarcomere hypertrophic cardiomyopathy. Cardiovascular Diagnosis and Therapy, 2019, 9, S299-S309.	1.7	16
71	Genetic testing in inherited endocrine disorders: joint position paper of the European reference network on rare endocrine conditions (Endo-ERN). Orphanet Journal of Rare Diseases, 2020, 15, 144.	2.7	15
72	Neurologic and neurodevelopmental complications in cardiofaciocutaneous syndrome are associated with genotype: A multinational cohort study. Genetics in Medicine, 2022, 24, 1556-1566.	2.4	15

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73	Genotype–epigenotype–phenotype correlations in females with frontometaphyseal dysplasia. American Journal of Medical Genetics, Part A, 2006, 140A, 1069-1073.	1.2	14
74	Expanding the genotypic and phenotypic spectrum of severe serine biosynthesis disorders. Human Mutation, 2020, 41, 1615-1628.	2.5	14
75	ABO blood type B and fucosyltransferase 2 non-secretor status as genetic risk factors for chronic pancreatitis. Gut, 2016, 65, 353-354.	12.1	13
76	A Postzygotic SMO Mutation Caused the Original Case of Happle–Tinschert Syndrome. Acta Dermato-Venereologica, 2018, 98, 534-535.	1.3	11
77	Evolutionary conserved networks of human height identify multiple Mendelian causes of short stature. European Journal of Human Genetics, 2019, 27, 1061-1071.	2.8	11
78	Mosaic <scp>RASopathy</scp> due to <scp><i>KRAS</i></scp> variant <scp>G12D</scp> with segmental overgrowth and associated peripheral vascular malformations. American Journal of Medical Genetics, Part A, 2021, 185, 3122-3128.	1.2	11
79	Cubilin Single Nucleotide Polymorphism Variants are Associated with Macroangiopathy While a Matrix Metalloproteinase-9 Single Nucleotide Polymorphism Flip-Flop may Indicate Susceptibility of Diabetic Nephropathy in Type-2 Diabetic Patients. Nephron, 2019, 141, 156-165.	1.8	10
80	The seventh international <scp>RASopathies</scp> symposium: Pathways to a cureâ€"expanding knowledge, enhancing research, and therapeutic discovery. American Journal of Medical Genetics, Part A, 2022, 188, 1915-1927.	1.2	10
81	Long-term Culture of EBV-induced Human Lymphoblastoid Cell Lines Reveals Chromosomal Instability. Journal of Histochemistry and Cytochemistry, 2020, 68, 239-251.	2.5	9
82	Ablepharon macrostomia syndrome: A distinct genetic entity clinically related to the group of FRAS–FREM complex disorders. American Journal of Medical Genetics, Part A, 2013, 161, 3012-3017.	1.2	7
83	Characteristic dental pattern with hypodontia and short roots in Fraser syndrome. American Journal of Medical Genetics, Part A, 2020, 182, 1681-1689.	1.2	7
84	The clinical significance of A2ML1 variants in Noonan syndrome has to be reconsidered. European Journal of Human Genetics, 2021, 29, 524-527.	2.8	7
85	Gut microbial similarity in twins is driven by shared environment and aging. EBioMedicine, 2022, 79, 104011.	6.1	7
86	Profiling of the Bacterial Microbiota along the Murine Alimentary Tract. International Journal of Molecular Sciences, 2022, 23, 1783.	4.1	6
87	Cutis marmorata telangiectatica congenita being caused by postzygotic GNA11 mutations. European Journal of Medical Genetics, 2022, 65, 104472.	1.3	6
88	Generation of functional cardiomyocytes from rat embryonic and induced pluripotent stem cells using feeder-free expansion and differentiation in suspension culture. PLoS ONE, 2018, 13, e0192652.	2.5	5
89	ADA2 deficiency in a patient with Noonan syndromeâ€like disorder with loose anagen hair: The coâ€occurrence of two rare syndromes. American Journal of Medical Genetics, Part A, 2019, 179, 2474-2480.	1.2	5
90	Two unrelated families with variable expression of Fraser syndrome due to the same pathogenic variant in the <scp><i>FRAS1</i></scp> gene. American Journal of Medical Genetics, Part A, 2020, 182, 773-779.	1.2	5

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91	Refinement of the critical genomic region for congenital hyperinsulinismÂin the Chromosome 9p deletion syndrome. Wellcome Open Research, 2019, 4, 149.	1.8	5
92	<i>WARS1</i> and <i>SARS1</i> : Two tRNA synthetases implicated in autosomal recessive microcephaly. Human Mutation, 2022, 43, 1454-1471.	2.5	5
93	Expanding the clinical phenotype of <scp>RASopathies</scp> in 38 Turkish patients, including the rare <scp><i>LZTR1</i></scp> , <scp><i>RAF1</i></scp> , <scp><i>RIT1</i></scp> variants, and large deletion in <scp><i>NF1</i></scp> . American Journal of Medical Genetics, Part A, 2021, 185, 3623-3633.	1.2	4
94	A 2q24.2 microdeletion containing <i>TANK</i> as novel candidate gene for intellectual disability. American Journal of Medical Genetics, Part A, 2019, 179, 832-836.	1.2	3
95	Mosaic Neurofibromatosis Type 1 With Multiple Cutaneous Diffuse and Plexiform Neurofibromas of the Lower Leg. Anticancer Research, 2020, 40, 3423-3427.	1.1	3
96	Aplasia cutis congenita in a CDC42 ―related developmental phenotype. American Journal of Medical Genetics, Part A, 2021, 185, 850-855.	1.2	3
97	Neurofibromatosis Type 1 With Cherubism-like Phenotype, Multiple Osteolytic Bone Lesions of Lower Extremities, and Alagille-syndrome: Case Report With Literature Survey. In Vivo, 2021, 35, 1711-1736.	1.3	3
98	Refinement of the critical genomic region for hypoglycaemia in the Chromosome 9p deletion syndrome. Wellcome Open Research, 2019, 4, 149.	1.8	3
99	Oral HRAS Mutation in Orofacial Nevus Sebaceous Syndrome (Schimmelpenning-Feuerstein-Mims-Syndrome): A Case Report With a Literature Survey. In Vivo, 2022, 36, 274-293.	1.3	3
100	Unrecognized juvenile nephropathic cystinosis. Kidney International, 2018, 94, 1027.	5.2	2
101	Outcomes in growth hormone-treated Noonan syndrome children: impact of PTPN11 mutation status. Endocrine Connections, 2022, 11, .	1.9	2
102	Correlation of PET-MRI, pathology, LOH and surgical success in a case of CHI with atypical large pancreatic focus. Journal of the Endocrine Society, 2022, 6, bvac056.	0.2	2
103	<i>KRAS</i> Mutation in an Implant-associated Peripheral Giant Cell Granuloma of the Jaw: Implications of Genetic Analysis of the Lesion for Treatment Concept and Surveillance. In Vivo, 2021, 35, 947-953.	1.3	1
104	In Vitro Modeling of Congenital Hypertrophic Cardiomyopathy using Induced Pluripotent Stem Cell-Derived Cardiomyocytes. Thoracic and Cardiovascular Surgeon, 2017, 65, S1-S110.	1.0	1
105	Congenital and Inherited Anomalies. , 0, , 58-68.		1
106	The mosaic hedgehog spectrum: another lesson on the polymorphy of mosaicism. British Journal of Dermatology, 2019, 182, 22-23.	1.5	0
107	Novel Germ Line Mutations in the KRAS2 Gene Cause Noonan Syndrome and Deregulate Hematopoietic Cell Growth Blood, 2005, 106, 1602-1602.	1.4	0
108	BRAF Mutations in Juvenile Myelomonocytic Leukemia Blood, 2007, 110, 4602-4602.	1.4	0