

# Siddharth Banka

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

99  
papers

2,972  
citations

31  
h-index

53  
g-index

114  
ext. papers

4,394  
ext. citations

7.9  
avg, IF

4.29  
L-index

#	Paper	IF	Citations
99	Refining the clinical phenotype associated with missense variants in exons 38 and 39 of KMT2D.. <i>American Journal of Medical Genetics, Part A</i> , <b>2022</b> ,	2.5	
98	Novel diagnostic DNA methylation epesignatures expand and refine the epigenetic landscapes of Mendelian disorders.. <i>Human Genetics and Genomics Advances</i> , <b>2022</b> , 3, 100075	0.8	1
97	Novel pathogenic variants and quantitative phenotypic analyses of Robinow syndrome: WNT signaling perturbation and phenotypic variability.. <i>Human Genetics and Genomics Advances</i> , <b>2022</b> , 3, 100074	0.8	3
96	Biallelic TMEM260 variants cause truncus arteriosus, with or without renal defects. <i>Clinical Genetics</i> , <b>2022</b> , 101, 127-133	4	1
95	A basement membrane discovery pipeline uncovers network complexity, regulators, and human disease associations.. <i>Science Advances</i> , <b>2022</b> , 8, eabn2265	14.3	4
94	Deficiency of TET3 leads to a genome-wide DNA hypermethylation epesignature in human whole blood. <i>Npj Genomic Medicine</i> , <b>2021</b> , 6, 92	6.2	0
93	100,000 Genomes Pilot on Rare-Disease Diagnosis in Health Care - Preliminary Report. <i>New England Journal of Medicine</i> , <b>2021</b> , 385, 1868-1880	59.2	34
92	A homozygous GRIN1 null variant causes a more severe phenotype of early infantile epileptic encephalopathy. <i>American Journal of Medical Genetics, Part A</i> , <b>2021</b> ,	2.5	1
91	SPEN haploinsufficiency causes a neurodevelopmental disorder overlapping proximal 1p36 deletion syndrome with an epesignature of X chromosomes in females. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 502-516	11	12
90	Personalised virtual gene panels reduce interpretation workload and maintain diagnostic rates of proband-only clinical exome sequencing for rare disorders. <i>Journal of Medical Genetics</i> , <b>2021</b> ,	5.8	2
89	Solving unsolved rare neurological diseases-a Solve-RD viewpoint. <i>European Journal of Human Genetics</i> , <b>2021</b> , 29, 1332-1336	5.3	0
88	Genotype-phenotype correlations and novel molecular insights into the DHX30-associated neurodevelopmental disorders. <i>Genome Medicine</i> , <b>2021</b> , 13, 90	14.4	2
87	Recurrent KCNT2 missense variants affecting p.Arg190 result in a recognizable phenotype. <i>American Journal of Medical Genetics, Part A</i> , <b>2021</b> , 185, 3083-3091	2.5	2
86	Non-coding region variants upstream of MEF2C cause severe developmental disorder through three distinct loss-of-function mechanisms. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 1083-1094	11	8
85	Bi-allelic loss-of-function variants in BCAS3 cause a syndromic neurodevelopmental disorder. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 1069-1082	11	4
84	Haploinsufficiency of ARFGEF1 is associated with developmental delay, intellectual disability, and epilepsy with variable expressivity. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 1901-1911	8.1	1
83	Solving patients with rare diseases through programmatic reanalysis of genome-phenome data. <i>European Journal of Human Genetics</i> , <b>2021</b> , 29, 1337-1347	5.3	4

82	The diagnostic utility of clinical exome sequencing in 60 patients with hearing loss disorders: A single-institution experience. <i>Clinical Otolaryngology</i> , <b>2021</b> , 46, 1257-1262	1.8	0
81	Biallelic variants in PCDHGC4 cause a novel neurodevelopmental syndrome with progressive microcephaly, seizures, and joint anomalies. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 2138-2149	8.1	1
80	New insights into the clinical and molecular spectrum of the novel CYFIP2-related neurodevelopmental disorder and impairment of the WRC-mediated actin dynamics. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 543-554	8.1	9
79	Neutrophil dysfunction triggers inflammatory bowel disease in G6PC3 deficiency. <i>Journal of Leukocyte Biology</i> , <b>2021</b> , 109, 1147-1154	6.5	2
78	Mutation-specific pathophysiological mechanisms define different neurodevelopmental disorders associated with SATB1 dysfunction. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 346-356	11	7
77	ERBB4 exonic deletions on chromosome 2q34 in patients with intellectual disability or epilepsy. <i>European Journal of Human Genetics</i> , <b>2021</b> , 29, 1377-1383	5.3	2
76	Impaired eIF5A function causes a Mendelian disorder that is partially rescued in model systems by spermidine. <i>Nature Communications</i> , <b>2021</b> , 12, 833	17.4	13
75	Clinical delineation, sex differences, and genotype-phenotype correlation in pathogenic KDM6A variants causing X-linked Kabuki syndrome type 2. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 1202-1210	8.1	3
74	Post-translational formation of hypusine in eIF5A: implications in human neurodevelopment. <i>Amino Acids</i> , <b>2021</b> , 1	3.5	4
73	Delineating the molecular and phenotypic spectrum of the SETD1B-related syndrome. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 2122-2137	8.1	3
72	Presence of pathogenic copy number variants (CNVs) is correlated with socioeconomic status. <i>Journal of Medical Genetics</i> , <b>2020</b> , 57, 70-72	5.8	2
71	Whole-genome sequencing of patients with rare diseases in a national health system. <i>Nature</i> , <b>2020</b> , 583, 96-102	50.4	139
70	Opposite Modulation of RAC1 by Mutations in TRIO Is Associated with Distinct, Domain-Specific Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , <b>2020</b> , 106, 338-355	11	21
69	Expanding Clinical Presentations Due to Variations in THOC2 mRNA Nuclear Export Factor. <i>Frontiers in Molecular Neuroscience</i> , <b>2020</b> , 13, 12	6.1	4
68	A restricted spectrum of missense KMT2D variants cause a multiple malformations disorder distinct from Kabuki syndrome. <i>Genetics in Medicine</i> , <b>2020</b> , 22, 867-877	8.1	17
67	Telemedicine strategy of the European Reference Network ITHACA for the diagnosis and management of patients with rare developmental disorders. <i>Orphanet Journal of Rare Diseases</i> , <b>2020</b> , 15, 103	4.2	14
66	Delineation of a Human Mendelian Disorder of the DNA Demethylation Machinery: TET3 Deficiency. <i>American Journal of Human Genetics</i> , <b>2020</b> , 106, 234-245	11	22
65	Evidence for 28 genetic disorders discovered by combining healthcare and research data. <i>Nature</i> , <b>2020</b> , 586, 757-762	50.4	103

64	Reply: Expanding the clinical and genetic spectrum of PCYT2-related disorders. <i>Brain</i> , <b>2020</b> , 143, e77	11.2	0
63	Haploinsufficiency of ATP6V0C possibly underlies 16p13.3 deletions that cause microcephaly, seizures, and neurodevelopmental disorder. <i>American Journal of Medical Genetics, Part A</i> , <b>2020</b> ,	2.5	2
62	Profound vitamin D deficiency in four siblings with Imlerslund-Grasbeck syndrome with homozygous CUBN mutation. <i>JIMD Reports</i> , <b>2019</b> , 49, 43-47	1.9	5
61	Mutations in PCYT2 disrupt etherlipid biosynthesis and cause a complex hereditary spastic paraplegia. <i>Brain</i> , <b>2019</b> , 142, 3382-3397	11.2	40
60	Spatially clustering de novo variants in CYFIP2, encoding the cytoplasmic FMRP interacting protein 2, cause intellectual disability and seizures. <i>European Journal of Human Genetics</i> , <b>2019</b> , 27, 747-759	5.3	25
59	Expanding the genetic and phenotypic spectrum of branched-chain amino acid transferase 2 deficiency. <i>Journal of Inherited Metabolic Disease</i> , <b>2019</b> , 42, 809-817	5.4	8
58	The clinical spectrum of the congenital myasthenic syndrome resulting from COL13A1 mutations. <i>Brain</i> , <b>2019</b> , 142, 1547-1560	11.2	20
57	Heterozygous Variants in KMT2E Cause a Spectrum of Neurodevelopmental Disorders and Epilepsy. <i>American Journal of Human Genetics</i> , <b>2019</b> , 104, 1210-1222	11	31
56	The clinical presentation caused by truncating CHD8 variants. <i>Clinical Genetics</i> , <b>2019</b> , 96, 72-84	4	20
55	Genotype-phenotype specificity in Menke-Hennekam syndrome caused by missense variants in exon 30 or 31 of CREBBP. <i>American Journal of Medical Genetics, Part A</i> , <b>2019</b> , 179, 1058-1062	2.5	14
54	Bi-allelic Loss-of-Function CACNA1B Mutations in Progressive Epilepsy-Dyskinesia. <i>American Journal of Human Genetics</i> , <b>2019</b> , 104, 948-956	11	17
53	HIST1H1E heterozygous protein-truncating variants cause a recognizable syndrome with intellectual disability and distinctive facial gestalt: A study to clarify the HIST1H1E syndrome phenotype in 30 individuals. <i>American Journal of Medical Genetics, Part A</i> , <b>2019</b> , 179, 2049-2055	2.5	10
52	A comparative analysis of KMT2D missense variants in Kabuki syndrome, cancers and the general population. <i>Journal of Human Genetics</i> , <b>2019</b> , 64, 161-170	4.3	16
51	A maternally inherited frameshift CDKL5 variant in a male with global developmental delay and late-onset generalized epilepsy. <i>American Journal of Medical Genetics, Part A</i> , <b>2019</b> , 179, 507-511	2.5	1
50	Kabuki syndrome: international consensus diagnostic criteria. <i>Journal of Medical Genetics</i> , <b>2019</b> , 56, 89-95.8	5.8	80
49	A new microdeletion syndrome involving TBC1D24, ATP6V0C, and PDPK1 causes epilepsy, microcephaly, and developmental delay. <i>Genetics in Medicine</i> , <b>2019</b> , 21, 1058-1064	8.1	12
48	Histone Lysine Methylases and Demethylases in the Landscape of Human Developmental Disorders. <i>American Journal of Human Genetics</i> , <b>2018</b> , 102, 175-187	11	108
47	Growth hormone deficiency as a cause for short stature in Wiedemann-Steiner Syndrome. <i>Endocrinology, Diabetes and Metabolism Case Reports</i> , <b>2018</b> , 2018,	1.4	7

46	De Novo Pathogenic Variants in CACNA1E Cause Developmental and Epileptic Encephalopathy with Contractures, Macrocephaly, and Dyskinesias. <i>American Journal of Human Genetics</i> , <b>2018</b> , 103, 666-678	11	44
45	Diagnosing childhood-onset inborn errors of metabolism by next-generation sequencing. <i>Archives of Disease in Childhood</i> , <b>2017</b> , 102, 1019-1029	2.2	24
44	RAC1 Missense Mutations in Developmental Disorders with Diverse Phenotypes. <i>American Journal of Human Genetics</i> , <b>2017</b> , 101, 466-477	11	73
43	ACTB Loss-of-Function Mutations Result in a Pleiotropic Developmental Disorder. <i>American Journal of Human Genetics</i> , <b>2017</b> , 101, 1021-1033	11	50
42	TBC1D24 genotype-phenotype correlation: Epilepsies and other neurologic features. <i>Neurology</i> , <b>2016</b> , 87, 77-85	6.5	75
41	Role of reverse phenotyping in interpretation of next generation sequencing data and a review of INPP5E related disorders. <i>European Journal of Paediatric Neurology</i> , <b>2016</b> , 20, 286-295	3.8	25
40	Small 6q16.1 Deletions Encompassing POU3F2 Cause Susceptibility to Obesity and Variable Developmental Delay with Intellectual Disability. <i>American Journal of Human Genetics</i> , <b>2016</b> , 98, 363-72	11	26
39	Response to: Mutation in MMP2 gene may result in scleroderma-like skin thickening by Bader-Meunier et al. <i>Annals of the Rheumatic Diseases</i> , <b>2016</b> , 75, e2	2.4	1
38	A Restricted Repertoire of De Novo Mutations in ITPR1 Cause Gillespie Syndrome with Evidence for Dominant-Negative Effect. <i>American Journal of Human Genetics</i> , <b>2016</b> , 98, 981-992	11	53
37	Distinct genetic architectures for syndromic and nonsyndromic congenital heart defects identified by exome sequencing. <i>Nature Genetics</i> , <b>2016</b> , 48, 1060-5	36.3	200
36	Dominant Mutations in the Autoimmune Regulator AIRE Are Associated with Common Organ-Specific Autoimmune Diseases. <i>Immunity</i> , <b>2015</b> , 42, 1185-96	32.3	156
35	Leri's pleonosteosis, a congenital rheumatic disease, results from microduplication at 8q22.1 encompassing GDF6 and SDC2 and provides insight into systemic sclerosis pathogenesis. <i>Annals of the Rheumatic Diseases</i> , <b>2015</b> , 74, 1249-56	2.4	19
34	Exome sequencing identifies ATP4A gene as responsible of an atypical familial type I gastric neuroendocrine tumour. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 2914-22	5.6	49
33	Further delineation of the KAT6B molecular and phenotypic spectrum. <i>European Journal of Human Genetics</i> , <b>2015</b> , 23, 1165-70	5.3	45
32	Novel KDM6A (UTX) mutations and a clinical and molecular review of the X-linked Kabuki syndrome (KS2). <i>Clinical Genetics</i> , <b>2015</b> , 87, 252-8	4	80
31	Kabuki syndrome: expanding the phenotype to include microphthalmia and anophthalmia. <i>Clinical Dysmorphology</i> , <b>2015</b> , 24, 135-9	0.9	18
30	RAP1-mediated MEK/ERK pathway defects in Kabuki syndrome. <i>Journal of Clinical Investigation</i> , <b>2015</b> , 125, 3585-99	15.9	48
29	Exome Sequencing Identifies a Dominant TNNT3 Mutation in a Large Family with Distal Arthrogyposis. <i>Molecular Syndromology</i> , <b>2014</b> , 5, 218-28	1.5	6

28	Expanding the clinical and molecular spectrum of thiamine pyrophosphokinase deficiency: a treatable neurological disorder caused by TPK1 mutations. <i>Molecular Genetics and Metabolism</i> , <b>2014</b> , 113, 301-6	3.7	40
27	The genetic basis of DOORS syndrome: an exome-sequencing study. <i>Lancet Neurology</i> , <b>2014</b> , 13, 44-58	24.1	96
26	A clinical and molecular review of ubiquitous glucose-6-phosphatase deficiency caused by G6PC3 mutations. <i>Orphanet Journal of Rare Diseases</i> , <b>2013</b> , 8, 84	4.2	53
25	Trisomy 18 mosaicism: report of two cases. <i>World Journal of Pediatrics</i> , <b>2013</b> , 9, 179-81	4.6	10
24	G6PC3 mutations cause non-syndromic severe congenital neutropenia. <i>Molecular Genetics and Metabolism</i> , <b>2013</b> , 108, 138-41	3.7	16
23	MLL2 mosaic mutations and intragenic deletion-duplications in patients with Kabuki syndrome. <i>Clinical Genetics</i> , <b>2013</b> , 83, 467-71	4	34
22	Weaver syndrome and EZH2 mutations: Clarifying the clinical phenotype. <i>American Journal of Medical Genetics, Part A</i> , <b>2013</b> , 161A, 2972-80	2.5	80
21	Extended spectrum of human glucose-6-phosphatase catalytic subunit 3 deficiency: novel genotypes and phenotypic variability in severe congenital neutropenia. <i>Journal of Pediatrics</i> , <b>2012</b> , 160, 679-683.e2	3.6	60
20	Corneal confocal microscopy detects small-fiber neuropathy in Charcot-Marie-Tooth disease type 1A patients. <i>Muscle and Nerve</i> , <b>2012</b> , 46, 698-704	3.4	67
19	How genetically heterogeneous is Kabuki syndrome?: MLL2 testing in 116 patients, review and analyses of mutation and phenotypic spectrum. <i>European Journal of Human Genetics</i> , <b>2012</b> , 20, 381-8	5.3	113
18	Germline mutations in the oncogene EZH2 cause Weaver syndrome and increased human height. <i>Oncotarget</i> , <b>2011</b> , 2, 1127-33	3.3	110
17	Further delineation of the phenotype of severe congenital neutropenia type 4 due to mutations in G6PC3. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 18-22	5.3	40
16	Pernicious anemia - genetic insights. <i>Autoimmunity Reviews</i> , <b>2011</b> , 10, 455-9	13.6	29
15	Identification and characterization of an inborn error of metabolism caused by dihydrofolate reductase deficiency. <i>American Journal of Human Genetics</i> , <b>2011</b> , 88, 216-25	11	68
14	Mutations in PRDM5 in brittle cornea syndrome identify a pathway regulating extracellular matrix development and maintenance. <i>American Journal of Human Genetics</i> , <b>2011</b> , 88, 767-777	11	84
13	Mutations in PRDM5 in Brittle Cornea Syndrome Identify a Pathway Regulating Extracellular Matrix Development and Maintenance. <i>American Journal of Human Genetics</i> , <b>2011</b> , 89, 346	11	2
12	A novel 800 kb microduplication of chromosome 16q22.1 resulting in learning disability and epilepsy may explain phenotypic variability in a family with 15q13 microdeletion. <i>American Journal of Medical Genetics, Part A</i> , <b>2011</b> , 155A, 1453-7	2.5	3
11	Spectrum of MLL2 (ALR) mutations in 110 cases of Kabuki syndrome. <i>American Journal of Medical Genetics, Part A</i> , <b>2011</b> , 155A, 1511-6	2.5	132

10	Variability of bone marrow morphology in G6PC3 mutations: is there a genotype-phenotype correlation or age-dependent relationship?. <i>American Journal of Hematology</i> , <b>2011</b> , 86, 235-7	7.1	16
9	De-novo duplication of 5(q13.3q21.1) in a child with vitreo-retinal dysplasia and learning disability. <i>Clinical Dysmorphology</i> , <b>2010</b> , 19, 73-75	0.9	
8	Early diagnosis and treatment of cobalamin deficiency of infancy owing to occult maternal pernicious anemia. <i>Journal of Pediatric Hematology/Oncology</i> , <b>2010</b> , 32, 319-22	1.2	13
7	Mutations in the G6PC3 gene cause Dursun syndrome. <i>American Journal of Medical Genetics, Part A</i> , <b>2010</b> , 152A, 2609-11	2.5	31
6	Array comparative genomic hybridisation-based identification of two imbalances of chromosome 1p in a 9-year-old girl with a monosomy 1p36 related phenotype and a family history of learning difficulties: a case report. <i>Journal of Medical Case Reports</i> , <b>2008</b> , 2, 355	1.2	7
5	First report of occurrence of choroid plexus papilloma and medulloblastoma in the same patient. <i>Child's Nervous System</i> , <b>2007</b> , 23, 587-9	1.7	6
4	Liver cyst caused by the peritoneal catheter of a cerebrospinal fluid shunt. <i>Pediatric Neurosurgery</i> , <b>2007</b> , 43, 343-4	0.9	2
3	Liver cyst caused by the peritoneal catheter of a CSF shunt. <i>Pediatric Neurosurgery</i> , <b>2007</b> , 43, 444-5	0.9	3
2	A basement membrane discovery pipeline uncovers network complexity, new regulators, and human disease associations		1
1	Delineation of the First Human Mendelian Disorder of the DNA Demethylation Machinery: TET3 Deficiency		1