

Bruno Reversade

List of Publications by Year in descending order

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Version: 2024-02-01

108
papers

7,347
citations

57719

44
h-index

62565

80
g-index

138
all docs

138
docs citations

138
times ranked

11309
citing authors

#	ARTICLE	IF	CITATIONS
1	Biallelic <i>ZNFX1</i> variants are associated with a spectrum of immunohematological abnormalities. <i>Clinical Genetics</i> , 2022, 101, 247-254.	1.0	12
2	Generation of three human induced pluripotent stem cell lines with <i>IRX5</i> knockout and knockin genetic editions using CRISPR-Cas9 system. <i>Stem Cell Research</i> , 2022, 58, 102627.	0.3	4
3	Huriez syndrome: Additional pathogenic variants supporting allelism to <i>SMARCAD1</i> syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 1752-1760.	0.7	2
4	Discovery of a genetic module essential for assigning left-right asymmetry in humans and ancestral vertebrates. <i>Nature Genetics</i> , 2022, 54, 62-72.	9.4	16
5	Human NLRP1 is a sensor of pathogenic coronavirus 3CL proteases in lung epithelial cells. <i>Molecular Cell</i> , 2022, 82, 2385-2400.e9.	4.5	61
6	HOX epimutations driven by maternal <i>SMCHD1/LRIF1</i> haploinsufficiency trigger homeotic transformations in genetically wildtype offspring. <i>Nature Communications</i> , 2022, 13, .	5.8	5
7	Direct identification of A-to-I editing sites with nanopore native RNA sequencing. <i>Nature Methods</i> , 2022, 19, 833-844.	9.0	35
8	R-SPONDIN2 mesenchymal cells form the bud tip progenitor niche during human lung development. <i>Developmental Cell</i> , 2022, 57, 1598-1614.e8.	3.1	19
9	Human model of <i>IRX5</i> mutations reveals key role for this transcription factor in ventricular conduction. <i>Cardiovascular Research</i> , 2021, 117, 2092-2107.	1.8	17
10	A Micropatterned Human-specific Neuroepithelial Tissue for Modeling Gene and Drug-induced Neurodevelopmental Defects. <i>Advanced Science</i> , 2021, 8, 2001100.	5.6	13
11	Structural basis for distinct inflammasome complex assembly by human NLRP1 and CARD8. <i>Nature Communications</i> , 2021, 12, 188.	5.8	54
12	IL11 is elevated in systemic sclerosis and IL11-dependent ERK signalling underlies TGF β -mediated activation of dermal fibroblasts. <i>Rheumatology</i> , 2021, 60, 5820-5826.	0.9	36
13	Huriez syndrome caused by a large deletion that abrogates the skin-specific isoform of <i>SMARCAD1</i> . <i>British Journal of Dermatology</i> , 2021, 184, 1205-1207.	1.4	3
14	Inherited deficiency of stress granule <i>ZNFX1</i> in patients with monocytosis and mycobacterial disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	47
15	AKT Signaling Modifies the Balance between Cell Proliferation and Migration in Neural Crest Cells from Patients Affected with Bosma Arhinia and Microphthalmia Syndrome. <i>Biomedicine</i> , 2021, 9, 751.	1.4	5
16	Loss of <i>C2orf69</i> defines a fatal autoinflammatory syndrome in humans and zebrafish that evokes a glycogen-storage-associated mitochondriopathy. <i>American Journal of Human Genetics</i> , 2021, 108, 1301-1317.	2.6	11
17	Somatic genetic rescue of a germline ribosome assembly defect. <i>Nature Communications</i> , 2021, 12, 5044.	5.8	44
18	Identical twins carry a persistent epigenetic signature of early genome programming. <i>Nature Communications</i> , 2021, 12, 5618.	5.8	26

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19	Metabolic pathway analyses identify proline biosynthesis pathway as a promoter of liver tumorigenesis. <i>Journal of Hepatology</i> , 2020, 72, 725-735.	1.8	71
20	Congenital insensitivity to pain with anhidrosis syndrome: A series from Jordan. <i>Clinical Neurology and Neurosurgery</i> , 2020, 189, 105636.	0.6	12
21	Enteroviral 3C protease activates the human NLRP1 inflammasome in airway epithelia. <i>Science</i> , 2020, 370, .	6.0	151
22	Heterozygous missense variant in EIF6 gene: A novel form of Shwachmanâ€“Diamond syndrome?. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2010-2020.	0.7	11
23	Mitchell-Riley syndrome iPSC exhibit reduced pancreatic endoderm differentiation due to an <i>RFX6</i> mutation. <i>Development (Cambridge)</i> , 2020, 147, .	1.2	10
24	A loss-of-function NUA2 mutation in humans causes anencephaly due to impaired Hippo-YAP signaling. <i>Journal of Experimental Medicine</i> , 2020, 217, .	4.2	25
25	Loss of MTX2 causes mandibuloacral dysplasia and links mitochondrial dysfunction to altered nuclear morphology. <i>Nature Communications</i> , 2020, 11, 4589.	5.8	30
26	Nextâ€“generation sequencing in a series of 80 fetuses with complex cardiac malformations and/or heterotaxy. <i>Human Mutation</i> , 2020, 41, 2167-2178.	1.1	21
27	ELABELA antagonizes intrarenal renin-angiotensin system to lower blood pressure and protects against renal injury. <i>American Journal of Physiology - Renal Physiology</i> , 2020, 318, F1122-F1135.	1.3	35
28	Congenital posterior cervical spine malformation due to biallelic c.240â€“T>G <i>RIPPLY2</i> variant: A discrete entity. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 1466-1472.	0.7	5
29	Mitochondrial peptide BRAWNIN is essential for vertebrate respiratory complex III assembly. <i>Nature Communications</i> , 2020, 11, 1312.	5.8	87
30	A <i>GLI3</i> variant leading to polydactyly in heterozygotes and Pallisterâ€“Hallâ€“like syndrome in a homozygote. <i>Clinical Genetics</i> , 2020, 97, 915-919.	1.0	3
31	Loss of PYCR2 Causes Neurodegeneration by Increasing Cerebral Glycine Levels via SHMT2. <i>Neuron</i> , 2020, 107, 82-94.e6.	3.8	30
32	Loss-of-function mutations in UDP-Glucose 6-Dehydrogenase cause recessive developmental epileptic encephalopathy. <i>Nature Communications</i> , 2020, 11, 595.	5.8	35
33	Dominant-negative NFKBIA mutation promotes IL-1 ^{Î²} production causing hepatic disease with severe immunodeficiency. <i>Journal of Clinical Investigation</i> , 2020, 130, 5817-5832.	3.9	17
34	R-spondin signalling is essential for the maintenance and differentiation of mouse nephron progenitors. <i>ELife</i> , 2020, 9, .	2.8	20
35	Generation of four H1 hESC sublines carrying a hemizygous knock-out/mutant MECP2. <i>Stem Cell Research</i> , 2019, 40, 101533.	0.3	1
36	Editorial overview: Cilia in development and disease. <i>Current Opinion in Genetics and Development</i> , 2019, 56, iii-iv.	1.5	0

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37	SMCHD1 is involved in <i>de novo</i> methylation of the DUX4-encoding D4Z4 macrosatellite. <i>Nucleic Acids Research</i> , 2019, 47, 2822-2839.	6.5	39
38	Novel mutation in HTRA1 in a family with diffuse white matter lesions and inflammatory features. <i>Neurology: Genetics</i> , 2019, 5, e345.	0.9	9
39	Homozygous <i>NLRP1</i> gain-of-function mutation in siblings with a syndromic form of recurrent respiratory papillomatosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 19055-19063.	3.3	92
40	Homozygous Null TBX4 Mutations Lead to Posterior Amelia with Pelvic and Pulmonary Hypoplasia. <i>American Journal of Human Genetics</i> , 2019, 105, 1294-1301.	2.6	17
41	Bone matrix hypermineralization associated with low bone turnover in a case of Nasu-Hakola disease. <i>Bone</i> , 2019, 123, 48-55.	1.4	7
42	Genome-wide CRISPR-Cas9 Screen Identifies Leukemia-Specific Dependence on a Pre-mRNA Metabolic Pathway Regulated by DCPS. <i>Cancer Cell</i> , 2018, 33, 386-400.e5.	7.7	99
43	KIAA1109 Variants Are Associated with a Severe Disorder of Brain Development and Arthrogryposis. <i>American Journal of Human Genetics</i> , 2018, 102, 116-132.	2.6	46
44	Novel mutations in the ciliopathy-associated gene CPLANE1 (C5orf42) cause OFD syndrome type VI rather than Joubert syndrome. <i>European Journal of Medical Genetics</i> , 2018, 61, 585-595.	0.7	22
45	ENPP1 Mutation Causes Recessive Cole Disease by Altering Melanogenesis. <i>Journal of Investigative Dermatology</i> , 2018, 138, 291-300.	0.3	23
46	Structural basis of RIP2 activation and signaling. <i>Nature Communications</i> , 2018, 9, 4993.	5.8	65
47	Human DPP9 represses NLRP1 inflammasome and protects against autoinflammatory diseases via both peptidase activity and FIIND domain binding. <i>Journal of Biological Chemistry</i> , 2018, 293, 18864-18878.	1.6	172
48	RSPO2 inhibition of RNF43 and ZNRF3 governs limb development independently of LGR4/5/6. <i>Nature</i> , 2018, 557, 564-569.	13.7	141
49	A homozygous loss-of-function CAMK2A mutation causes growth delay, frequent seizures and severe intellectual disability. <i>ELife</i> , 2018, 7, .	2.8	53
50	FSHD2- and BAMS-associated mutations confer opposing effects on SMCHD1 function. <i>Journal of Biological Chemistry</i> , 2018, 293, 9841-9853.	1.6	33
51	Cenani-Lenz syndactyly syndrome - a case report of a family with isolated syndactyly. <i>BMC Medical Genetics</i> , 2018, 19, 125.	2.1	9
52	De novo mutations in SMCHD1 cause Bosma arhinia microphthalmia syndrome and abrogate nasal development. <i>Nature Genetics</i> , 2017, 49, 249-255.	9.4	88
53	ELABELA-APJ axis protects from pressure overload heart failure and angiotensin II-induced cardiac damage. <i>Cardiovascular Research</i> , 2017, 113, 760-769.	1.8	111
54	Katanin p80, NuMA and cytoplasmic dynein cooperate to control microtubule dynamics. <i>Scientific Reports</i> , 2017, 7, 39902.	1.6	25

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55	Fifteen years of research on oral“facial”digital syndromes: from 1 to 16 causal genes. <i>Journal of Medical Genetics</i> , 2017, 54, 371-380.	1.5	85
56	Loss-of-Function Mutations in LGI4, a Secreted Ligand Involved in Schwann Cell Myelination, Are Responsible for Arthrogyriposis Multiplex Congenita. <i>American Journal of Human Genetics</i> , 2017, 100, 659-665.	2.6	19
57	CDK10 Mutations in Humans and Mice Cause Severe Growth Retardation, Spine Malformations, and Developmental Delays. <i>American Journal of Human Genetics</i> , 2017, 101, 391-403.	2.6	35
58	Alternative Progenitor Cells Compensate to Rebuild the Coronary Vasculature in Elabela- and Apj-Deficient Hearts. <i>Developmental Cell</i> , 2017, 42, 655-666.e3.	3.1	88
59	Discriminative Features in Three Autosomal Recessive Cutis Laxa Syndromes: Cutis Laxa IIA, Cutis Laxa IIB, and Geroderma Osteoplastica. <i>International Journal of Molecular Sciences</i> , 2017, 18, 635.	1.8	18
60	Long-Term Culture of Self-renewing Pancreatic Progenitors Derived from Human Pluripotent Stem Cells. <i>Stem Cell Reports</i> , 2017, 8, 1675-1688.	2.3	55
61	ELABELA deficiency promotes preeclampsia and cardiovascular malformations in mice. <i>Science</i> , 2017, 357, 707-713.	6.0	181
62	Identification of Common Genetic Variants Influencing Spontaneous Dizygotic Twinning and Female Fertility. <i>American Journal of Human Genetics</i> , 2016, 98, 898-908.	2.6	89
63	Loss-of-Function Mutations in ELMO2 Cause Intraosseous Vascular Malformation by Impeding RAC1 Signaling. <i>American Journal of Human Genetics</i> , 2016, 99, 299-317.	2.6	23
64	Germline NLRP1 Mutations Cause Skin Inflammatory and Cancer Susceptibility Syndromes via Inflammasome Activation. <i>Cell</i> , 2016, 167, 187-202.e17.	13.5	317
65	Loss of Iroquois homeobox transcription factors 3 and 5 in osteoblasts disrupts cranial mineralization. <i>Bone Reports</i> , 2016, 5, 86-95.	0.2	21
66	Incidentalome from Genomic Sequencing: A Barrier to Personalized Medicine?. <i>EBioMedicine</i> , 2016, 5, 211-216.	2.7	23
67	A novel ICK mutation causes ciliary disruption and lethal endocrine-cerebro-osteodysplasia syndrome. <i>Cilia</i> , 2016, 5, 8.	1.8	37
68	Discovery and Structure“Activity Relationship of a Bioactive Fragment of ELABELA that Modulates Vascular and Cardiac Functions. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 2962-2972.	2.9	100
69	De Novo Loss-of-Function Mutations in USP9X Cause a Female-Specific Recognizable Syndrome with Developmental Delay and Congenital Malformations. <i>American Journal of Human Genetics</i> , 2016, 98, 373-381.	2.6	95
70	Leveraging ancestry to improve causal variant identification in exome sequencing for monogenic disorders. <i>European Journal of Human Genetics</i> , 2016, 24, 113-119.	1.4	3
71	Quantitative imaging reveals real-time Pou5f3“Nanog complexes driving dorsoventral mesendoderm patterning in zebrafish. <i>ELife</i> , 2016, 5, .	2.8	34
72	The Apelin receptor enhances Nodal/TGF β 2 signaling to ensure proper cardiac development. <i>ELife</i> , 2016, 5, .	2.8	34

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73	Mutation in TWINKLE in a Large Iranian Family with Progressive External Ophthalmoplegia, Myopathy, Dysphagia and Dysphonia, and Behavior Change. <i>Archives of Iranian Medicine</i> , 2016, 19, 87-91.	0.2	4
74	Intellectual disability, muscle weakness and characteristic face in three siblings: A newly described recessive syndrome mapping to 3p24.3â€“p25.3. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2508-2515.	0.7	9
75	Loss of the scavenger mRNA decapping enzyme DCPS causes syndromic intellectual disability with neuromuscular defects. <i>Human Molecular Genetics</i> , 2015, 24, 3163-3171.	1.4	31
76	ELABELA Is an Endogenous Growth Factor that Sustains hESC Self-Renewal via the PI3K/AKT Pathway. <i>Cell Stem Cell</i> , 2015, 17, 635.	5.2	5
77	ELABELA Is an Endogenous Growth Factor that Sustains hESC Self-Renewal via the PI3K/AKT Pathway. <i>Cell Stem Cell</i> , 2015, 17, 435-447.	5.2	139
78	Recurrent De Novo Mutations Affecting Residue Arg138 of Pyrroline-5-Carboxylate Synthase Cause a Progeroid Form of Autosomal-Dominant Cutis Laxa. <i>American Journal of Human Genetics</i> , 2015, 97, 483-492.	2.6	70
79	Gmnc Is a Master Regulator of the Multiciliated Cell Differentiation Program. <i>Current Biology</i> , 2015, 25, 3267-3273.	1.8	83
80	Multiple Self-Healing Palmoplantar Carcinoma: A Familial Predisposition to Skin Cancer with Primary Palmoplantar and Conjunctival Lesions. <i>Journal of Investigative Dermatology</i> , 2015, 135, 304-308.	0.3	20
81	The hormonal peptide Elabela guides angioblasts to the midline during vasculogenesis. <i>ELife</i> , 2015, 4, .	2.8	86
82	Katanin p80 Regulates Human Cortical Development by Limiting Centriole and Cilia Number. <i>Neuron</i> , 2014, 84, 1240-1257.	3.8	89
83	C5orf42 is the major gene responsible for OFD syndrome type VI. <i>Human Genetics</i> , 2014, 133, 367-377.	1.8	71
84	ELABELA: A Hormone Essential for Heart Development Signals via the Apelin Receptor. <i>Developmental Cell</i> , 2013, 27, 672-680.	3.1	384
85	Optimal histone H3 to linker histone H1 chromatin ratio is vital for mesodermal competence in <i>Xenopus</i> . <i>Development (Cambridge)</i> , 2013, 140, 853-860.	1.2	13
86	The snRNA-processing complex, Integrator, is required for ciliogenesis and dynein recruitment to the nuclear envelope via distinct mechanisms. <i>Biology Open</i> , 2013, 2, 1390-1396.	0.6	18
87	Nuclear-localized Asunder regulates cytoplasmic dynein localization via its role in the Integrator complex. <i>Molecular Biology of the Cell</i> , 2013, 24, 2954-2965.	0.9	23
88	Definition of the phenotypic spectrum of Temtamy preaxial brachydactyly syndrome associated with autosomal recessive CHYS1 mutations. <i>Middle East Journal of Medical Genetics</i> , 2012, 1, 64-70.	0.0	6
89	Haploinsufficiency for AAGAB causes clinically heterogeneous forms of punctate palmoplantar keratoderma. <i>Nature Genetics</i> , 2012, 44, 1272-1276.	9.4	78
90	Human Asunder promotes dynein recruitment and centrosomal tethering to the nucleus at mitotic entry. <i>Molecular Biology of the Cell</i> , 2012, 23, 4713-4724.	0.9	19

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91	Mutations in IRX5 impair craniofacial development and germ cell migration via SDF1. <i>Nature Genetics</i> , 2012, 44, 709-713.	9.4	68
92	Multiple self-healing squamous epithelioma is caused by a disease-specific spectrum of mutations in TGFBR1. <i>Nature Genetics</i> , 2011, 43, 365-369.	9.4	147
93	Loss of CHSY1, a Secreted FRINGE Enzyme, Causes Syndromic Brachydactyly in Humans via Increased NOTCH Signaling. <i>American Journal of Human Genetics</i> , 2010, 87, 768-778.	2.6	82
94	Mutations in PYCR1 cause cutis laxa with progeroid features. <i>Nature Genetics</i> , 2009, 41, 1016-1021.	9.4	211
95	05-P010 Histone variant H3.3 is required for <i>Xenopus</i> embryonic development. <i>Mechanisms of Development</i> , 2009, 126, S115.	1.7	0
96	06-P037 Dubowitz syndrome maps to chromosome 12 within an interval of 70 genes. <i>Mechanisms of Development</i> , 2009, 126, S130-S131.	1.7	0
97	The opposing homeobox genes Goosecoid and Vent1/2 self-regulate <i>Xenopus</i> patterning. <i>EMBO Journal</i> , 2007, 26, 2955-2965.	3.5	73
98	Embryonic Dorsal-Ventral Signaling: Secreted Frizzled-Related Proteins as Inhibitors of Tolloid Proteinases. <i>Cell</i> , 2006, 124, 147-159.	13.5	199
99	Default neural induction: neuralization of dissociated <i>Xenopus</i> cells is mediated by Ras/MAPK activation. <i>Genes and Development</i> , 2005, 19, 1022-1027.	2.7	128
100	Sirenomelia in Bmp7 and Tsg compound mutant mice: requirement for Bmp signaling in the development of ventral posterior mesoderm. <i>Development (Cambridge)</i> , 2005, 132, 2489-2499.	1.2	75
101	Depletion of Bmp2, Bmp4, Bmp7 and Spemann organizer signals induces massive brain formation in <i>Xenopus</i> embryos. <i>Development (Cambridge)</i> , 2005, 132, 3381-3392.	1.2	131
102	Regulation of ADMP and BMP2/4/7 at Opposite Embryonic Poles Generates a Self-Regulating Morphogenetic Field. <i>Cell</i> , 2005, 123, 1147-1160.	13.5	232
103	Chordin Is Required for the Spemann Organizer Transplantation Phenomenon in <i>Xenopus</i> Embryos. <i>Developmental Cell</i> , 2003, 4, 219-230.	3.1	130
104	The pro-BMP activity of Twisted gastrulation is independent of BMP binding. <i>Development (Cambridge)</i> , 2003, 130, 4047-4056.	1.2	65
105	Connective-tissue growth factor (CTGF) modulates cell signalling by BMP and TGF- β 2. <i>Nature Cell Biology</i> , 2002, 4, 599-604.	4.6	817
106	Proteolytic cleavage of Chordin as a switch for the dual activities of Twisted gastrulation in BMP signaling. <i>Development (Cambridge)</i> , 2001, 128, 4439-4447.	1.2	134
107	Proteolytic cleavage of Chordin as a switch for the dual activities of Twisted gastrulation in BMP signaling. <i>Development (Cambridge)</i> , 2001, 128, 4439-47.	1.2	65
108	Gene expression profiles in normal and Otx2 ^{-/-} early gastrulating mouse embryos. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 14388-14393.	3.3	51