Silvia Russo

List of Publications by Year in descending order

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Version: 2024-02-01

136940 118840 4,436 105 32 62 h-index citations g-index papers 106 106 106 5536 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Genes for RNA-binding proteins involved in neural-specific functions and diseases are downregulated in Rubinstein-Taybi iNeurons. Neural Regeneration Research, 2022, 17, 5.	3.0	5
2	Germline variants in genes of the subcortical maternal complex and Multilocus Imprinting Disturbance are associated with miscarriage/infertility or Beckwith–Wiedemann progeny. Clinical Epigenetics, 2022, 14, 43.	4.1	6
3	Trans-acting genetic variants causing multilocus imprinting disturbance (MLID): common mechanisms and consequences. Clinical Epigenetics, 2022, 14, 41.	4.1	14
4	SCN2A Pathogenic Variants and Epilepsy: Heterogeneous Clinical, Genetic and Diagnostic Features. Brain Sciences, 2022, 12, 18.	2.3	5
5	Expanding the Molecular Spectrum of ANKRD11 Gene Defects in 33 Patients with a Clinical Presentation of KBG Syndrome. International Journal of Molecular Sciences, 2022, 23, 5912.	4.1	6
6	Clinical and molecular characterization of patients affected by ⟨scp⟩Beckwithâ€Wiedemann⟨/scp⟩ spectrum conceived through assisted reproduction techniques. Clinical Genetics, 2022, 102, 314-323.	2.0	7
7	Maternal Uniparental Disomy of Chromosome 20 (UPD(20)mat) as Differential Diagnosis of Silver Russell Syndrome: Identification of Three New Cases. Genes, 2021, 12, 588.	2.4	6
8	Histone Deacetylase Inhibitors Ameliorate Morphological Defects and Hypoexcitability of iPSC-Neurons from Rubinstein-Taybi Patients. International Journal of Molecular Sciences, 2021, 22, 5777.	4.1	7
9	Editorial: Overlapping Phenotypes and Genetic Heterogeneity of Rare Neurodevelopmental Disorders. Frontiers in Neurology, 2021, 12, 711288.	2.4	2
10	Interconnected Gene Networks Underpin the Clinical Overlap of HNRNPH1-Related and Rubinstein–Taybi Intellectual Disability Syndromes. Frontiers in Neuroscience, 2021, 15, 745684.	2.8	0
11	Guided Growth in Leg Length Discrepancy in Beckwith-Wiedemann Syndrome: A Consecutive Case Series. Children, 2021, 8, 1152.	1.5	1
12	A novel RAD21 mutation in a boy with mild Cornelia de Lange presentation: Further delineation of the phenotype. European Journal of Medical Genetics, 2020, 63, 103620.	1.3	6
13	Cognitive Profiles and Brain Volume Are Affected in Patients with Silver–Russell Syndrome. Journal of Clinical Endocrinology and Metabolism, 2020, 105, e1478-e1488.	3.6	4
14	Recombinant Chromosome 7 Driven by Maternal Chromosome 7 Pericentric Inversion in a Girl with Features of Silver-Russell Syndrome. International Journal of Molecular Sciences, 2020, 21, 8487.	4.1	1
15	Sleep disordered breathing and daytime hypoventilation in a male with MECP2 mutation. American Journal of Medical Genetics, Part A, 2020, 182, 2982-2987.	1.2	3
16	Loss-of-function maternal-effect mutations of PADI6 are associated with familial and sporadic Beckwith-Wiedemann syndrome with multi-locus imprinting disturbance. Clinical Epigenetics, 2020, 12, 139.	4.1	40
17	Transcriptome Analysis of iPSC-Derived Neurons from Rubinstein-Taybi Patients Reveals Deficits in Neuronal Differentiation. Molecular Neurobiology, 2020, 57, 3685-3701.	4.0	15
18	DNA Methylation in the Diagnosis of Monogenic Diseases. Genes, 2020, 11, 355.	2.4	28

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19	Comparison of Quantitative Analysis of Methylated Alleles Real-Time PCR and Methylation-Specific MLPA for Molecular Diagnosis of Beckwith-Wiedemann Syndrome. Pathobiology, 2019, 86, 217-224.	3.8	1
20	Pathogenic Variants in STXBP1 and in Genes for GABAa Receptor Subunities Cause Atypical Rett/Rett-like Phenotypes. International Journal of Molecular Sciences, 2019, 20, 3621.	4.1	29
21	Generation of three iPSC lines (IAli002, IAli004, IAli003) from Rubinstein-Taybi syndrome 1 patients carrying CREBBP non sense c.4435G>T, p.(Gly1479*) and c.3474G>A, p.(Trp1158*) and missense c.4627G>T, p.(Asp1543Tyr) mutations. Stem Cell Research, 2019, 40, 101553.	0.7	6
22	Discrepant molecular and clinical diagnoses in Beckwith-Wiedemann and Silver-Russell syndromes. Genetical Research, 2019, 101, e3.	0.9	17
23	Analysis of the Phenotypes in the Rett Networked Database. International Journal of Genomics, 2019, 2019, 1-9.	1.6	23
24	Molecular Etiology Disclosed by Array CGH in Patients With Silver–Russell Syndrome or Similar Phenotypes. Frontiers in Genetics, 2019, 10, 955.	2.3	11
25	A KHDC3L mutation resulting in recurrent hydatidiform mole causes genome-wide DNA methylation loss in oocytes and persistent imprinting defects post-fertilisation. Genome Medicine, 2019, 11, 84.	8.2	45
26	The phenotypic variations of multi-locus imprinting disturbances associated with maternal-effect variants of NLRP5 range from overt imprinting disorder to apparently healthy phenotype. Clinical Epigenetics, 2019, 11, 190.	4.1	22
27	Clinical and molecular diagnosis, screening and management of Beckwith–Wiedemann syndrome: an international consensus statement. Nature Reviews Endocrinology, 2018, 14, 229-249.	9.6	388
28	Recommendations for a nomenclature system for reporting methylation aberrations in imprinted domains. Epigenetics, 2018, 13, 117-121.	2.7	70
29	Segmental Maternal UPD of Chromosome 7q in a Patient With Pendred and Silver Russell Syndromes-Like Features. Frontiers in Genetics, 2018, 9, 600.	2.3	4
30	Workload measurement for molecular genetics laboratory: A survey study. PLoS ONE, 2018, 13, e0206855.	2.5	6
31	Recurrence and Familial Inheritance of Intronic NIPBL Pathogenic Variant Associated With Mild CdLS. Frontiers in Neurology, 2018, 9, 967.	2.4	6
32	Use of nutritional devices in Cornelia de Lange syndrome: Data from a large Italian cohort. American Journal of Medical Genetics, Part A, 2018, 176, 1865-1871.	1.2	2
33	The phenotype of <i>SCN8A</i> developmental and epileptic encephalopathy. Neurology, 2018, 91, e1112-e1124.	1.1	114
34	Generation of the Rubinstein-Taybi syndrome type 2 patient-derived induced pluripotent stem cell line (IAli001-A) carrying the EP300 exon 23 stop mutation c.3829A†>†T, p.(Lys1277*). Stem Cell Research, 20 30, 175-179.)18).7	4
35	Taurine Administration Recovers Motor and Learning Deficits in an Angelman Syndrome Mouse Model. International Journal of Molecular Sciences, 2018, 19, 1088.	4.1	14
36	iPSC-derived neurons of CREBBP - and EP300 -mutated Rubinstein-Taybi syndrome patients show morphological alterations and hypoexcitability. Stem Cell Research, 2018, 30, 130-140.	0.7	19

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37	Clinical Manifestations and Metabolic Outcomes of Seven Adults with Silver-Russell Syndrome. Journal of Clinical Endocrinology and Metabolism, 2018, 103, 2225-2233.	3.6	10
38	Nomenclature and definition in asymmetric regional body overgrowth. American Journal of Medical Genetics, Part A, 2017, 173, 1735-1738.	1.2	36
39	Phenotypes and genotypes in individuals with <i>SMC1A</i> variants. American Journal of Medical Genetics, Part A, 2017, 173, 2108-2125.	1.2	69
40	Diagnosis and management of Silver–Russell syndrome: first international consensus statement. Nature Reviews Endocrinology, 2017, 13, 105-124.	9.6	336
41	Cancer Risk in Beckwith-Wiedemann Syndrome: A Systematic Review and Meta-Analysis Outlining a Novel (Epi)Genotype Specific Histotype Targeted Screening Protocol. Journal of Pediatrics, 2016, 176, 142-149.e1.	1.8	119
42	A new prognostic index of severity of intellectual disabilities in Cornelia de Lange syndrome. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2016, 172, 179-189.	1.6	7
43	EMQN best practice guidelines for the molecular genetic testing and reporting of chromosome 11p15 imprinting disorders: Silver–Russell and Beckwith–Wiedemann syndrome. European Journal of Human Genetics, 2016, 24, 1377-1387.	2.8	68
44	Thrombocytopenia and Cornelia de Lange syndrome: Still an enigma?. American Journal of Medical Genetics, Part A, 2016, 170, 130-134.	1.2	4
45	A multi-method approach to the molecular diagnosis of overt and borderline 11p15.5 defects underlying Silver–Russell and Beckwith–Wiedemann syndromes. Clinical Epigenetics, 2016, 8, 23.	4.1	54
46	Sleep disorders in Cornelia de Lange syndrome. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2016, 172, 214-221.	1.6	14
47	Functional characterization of CDK5 and CDK5R1 mutations identified in patients with non-syndromic intellectual disability. Journal of Human Genetics, 2016, 61, 283-293.	2.3	12
48	Recommendations of the Scientific Committee of the Italian Beckwith–Wiedemann Syndrome Association on the diagnosis, management and follow-up of the syndrome. European Journal of Medical Genetics, 2016, 59, 52-64.	1.3	76
49	Prenatal molecular testing for Beckwith–Wiedemann and Silver–Russell syndromes: a challenge for molecular analysis and genetic counseling. European Journal of Human Genetics, 2016, 24, 784-793.	2.8	44
50	Blood oxidative stress and metallothionein expression in Rett syndrome: Probing for markers. World Journal of Biological Psychiatry, 2016, 17, 198-209.	2.6	11
51	(Epi)genotype–phenotype correlations in Beckwith–Wiedemann syndrome. European Journal of Human Genetics, 2016, 24, 183-190.	2.8	113
52	Constitutional de novo deletion of the FBXW7 gene in a patient with focal segmental glomerulosclerosis and multiple primitive tumors. Scientific Reports, 2015, 5, 15454.	3.3	10
53	Antiepileptic drugs in Rett Syndrome. European Journal of Paediatric Neurology, 2015, 19, 446-452.	1.6	13
54	Familial intragenic duplication of ANKRD11 underlying three patients of KBG syndrome. Molecular Cytogenetics, 2015, 8, 20.	0.9	18

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55	Epilepsy in Rett syndromeâ€"Lessons from the Rett networked database. Epilepsia, 2015, 56, 569-576.	5.1	47
56	Complex <i>de novo</i> chromosomal rearrangement at 15q11–q13 involving an intrachromosomal triplication in a patient with a severe neuropsychological phenotype: Clinical report and review of the literature. American Journal of Medical Genetics, Part A, 2015, 167, 221-230.	1.2	14
57	Somatic Mosaicism as Modulator of the Global and Intellectual Phenotype in Epimutated Angelman Syndrome Patients. Journal of Intellectual Disability - Diagnosis and Treatment, 2015, 3, 126-137.	0.3	1
58	Overall and allele-specific expression of the SMC1 Agene in female Cornelia de Lange syndrome patients and healthy controls. Epigenetics, 2014, 9, 973-979.	2.7	10
59	Cervical spine malformation in cornelia de lange syndrome: A report of three patients. American Journal of Medical Genetics, Part A, 2014, 164, 1520-1524.	1.2	6
60	Think About It. Journal of Child Neurology, 2014, 29, NP74-NP77.	1.4	10
61	Recent insights into genotype–phenotype relationships in patients with Rett syndrome using a fine grain scale. Research in Developmental Disabilities, 2014, 35, 2976-2986.	2.2	40
62	Genomic imbalances in patients with a clinical presentation in the spectrum of Cornelia de Lange syndrome. BMC Medical Genetics, 2013, 14, 41.	2.1	15
63	Prevalence of beckwith–wiedemann syndrome in North West of Italy. American Journal of Medical Genetics, Part A, 2013, 161, 2481-2486.	1.2	93
64	Cornelia de Lange individuals with new and recurrent <i>SMC1A</i> mutations enhance delineation of mutation repertoire and phenotypic spectrum. American Journal of Medical Genetics, Part A, 2013, 161, 2909-2919.	1.2	31
65	Molecular characterization of a mosaic NIPBL deletion in a Cornelia de Lange patient with severe phenotype. European Journal of Medical Genetics, 2013, 56, 138-143.	1.3	24
66	Germline mosaicism in cornelia de lange syndrome: Dilemmas and risk figures. American Journal of Medical Genetics, Part A, 2013, 161, 1825-1826.	1.2	2
67	Quantitative DNA methylation analysis improves epigenotype-phenotype correlations in Beckwith-Wiedemann syndrome. Epigenetics, 2013, 8, 1053-1060.	2.7	33
68	Intragenic and large NIPBL rearrangements revealed by MLPA in Cornelia de Lange patients. European Journal of Human Genetics, 2012, 20, 734-741.	2.8	23
69	The KCNQ1OT1 imprinting control region and non-coding RNA: new properties derived from the study of Beckwith–Wiedemann syndrome and Silver–Russell syndrome cases. Human Molecular Genetics, 2012, 21, 10-25.	2.9	135
70	Deletion of the AP1S2 gene in a child with psychomotor delay and hypotonia. European Journal of Medical Genetics, 2012, 55, 124-127.	1.3	8
71	Complex rearrangement involving 9p deletion and duplication in a syndromic patient: Genotype/phenotype correlation and review of the literature. Gene, 2012, 502, 40-45.	2.2	16
72	Juxtaposition of heterochromatic and euchromatic regions by chromosomal translocation mediates a heterochromatic long-range position effect associated with a severe neurological phenotype. Molecular Cytogenetics, 2012, 5, 16.	0.9	22

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73	Rett networked database: An integrated clinical and genetic network of rett syndrome databases. Human Mutation, 2012, 33, 1031-1036.	2.5	19
74	Electroclinical pattern in <i>MECP2</i> duplication syndrome: Eight new reported cases and review of literature. Epilepsia, 2012, 53, 1146-1155.	5.1	34
75	Medical care of adolescents and women with Rett syndrome: An Italian study. American Journal of Medical Genetics, Part A, 2012, 158A, 13-18.	1.2	26
76	Nephrological findings and genotype–phenotype correlation in Beckwith–Wiedemann syndrome. Pediatric Nephrology, 2012, 27, 397-406.	1.7	55
77	Role of UBE3A and ATP10A genes in autism susceptibility region $15q11-q13$ in an Italian population: A positive replication for UBE3A. Psychiatry Research, 2011, 185, 33-38.	3.3	29
78	Mutations in the Small GTPase Gene RAB39B Are Responsible for X-linked Mental Retardation Associated with Autism, Epilepsy, and Macrocephaly. American Journal of Human Genetics, 2010, 86, 185-195.	6.2	220
79	Hypertrophic cardiomyopathy in a girl with Cornelia de Lange syndrome due to mutation in <i>SMC1A</i> . American Journal of Medical Genetics, Part A, 2010, 152A, 2127-2129.	1.2	23
80	Noonan syndrome associated with both a new Jnkâ€activating familial <i>SOS1</i> and a de novo <i>RAF1</i> mutations. American Journal of Medical Genetics, Part A, 2010, 152A, 2176-2184.	1.2	20
81	Efficacy of levetiracetam in the treatment of drug-resistant Rett syndrome. Epilepsy Research, 2010, 88, 112-117.	1.6	18
82	Genetic investigations on 8 patients affected by ring 20 chromosome syndrome. BMC Medical Genetics, 2010, 11, 146.	2.1	86
83	Epilepsy in Rett syndrome: Clinical and genetic features. Epilepsy and Behavior, 2010, 19, 296-300.	1.7	68
84	Inherited and Sporadic Epimutations at the <i>IGF2-H19</i> Locus in Beckwith-Wiedemann Syndrome and Wilms’ Tumor. Endocrine Development, 2009, 14, 1-9.	1.3	48
85	A $<$ i>CDKL5 $<$ /i> mutated child with precocious puberty. American Journal of Medical Genetics, Part A, 2009, 149A, 1046-1051.	1.2	21
86	Novel mutations in the CDKL5 gene, predicted effects and associated phenotypes. Neurogenetics, 2009, 10, 241-250.	1.4	62
87	Premature chromatid separation is not a useful diagnostic marker for Cornelia de Lange syndrome. Chromosome Research, 2009, 17, 763-771.	2.2	49
88	Hypomethylation at multiple maternally methylated imprinted regions including PLAGL1 and GNAS loci in Beckwith–Wiedemann syndrome. European Journal of Human Genetics, 2009, 17, 611-619.	2.8	194
89	Clinical and electroencephalographic features in patients with CDKL5 mutations: Two new Italian cases and review of the literature. Epilepsy and Behavior, 2008, 12, 326-331.	1.7	36
90	Mechanisms causing imprinting defects in familial Beckwith–Wiedemann syndrome with Wilms' tumour. Human Molecular Genetics, 2007, 16, 254-264.	2.9	100

#	Article	IF	CITATIONS
91	Prenatal/neonatal pathology in two cases of Cornelia de Lange syndrome harboring novel mutations of NIPBL. Genetics in Medicine, 2007, 9, 188-194.	2.4	18
92	Evaluation of autism traits in Angelman syndrome: a resource to unfold autism genes. Neurogenetics, 2007, 8, 169-178.	1.4	81
93	X-linked Cornelia de Lange syndrome owing to SMC1L1 mutations. Nature Genetics, 2006, 38, 528-530.	21.4	393
94	Mutations and novel polymorphisms in coding regions and UTRs of CDK5R1 and OMG genes in patients with non-syndromic mental retardation. Neurogenetics, 2006, 7, 59-66.	1.4	32
95	A fluorescent method for detecting low-grade 11patUPD mosaicism in Beckwith–Wiedemann syndrome. Molecular and Cellular Probes, 2003, 17, 295-299.	2.1	11
96	Genomic inversions of human chromosome 15q11-q13 in mothers of Angelman syndrome patients with class II (BP2/3) deletions. Human Molecular Genetics, 2003, 12, 849-858.	2.9	131
97	Small familial supernumerary ring chromosome 2: FISH characterization and genotypeâ€phenotype correlation. American Journal of Medical Genetics Part A, 2002, 111, 319-323.	2.4	13
98	Refinement of the NHS locus on chromosome Xp22.13 and analysis of five candidate genes. European Journal of Human Genetics, 2002, 10, 516-520.	2.8	23
99	Refined FISH characterization of a de novo $1p22$ - $p36.2$ paracentric inversion and associated $1p21$ - 22 deletion in a patient with signs of $1p36$ microdeletion syndrome. American Journal of Medical Genetics Part A, 2001, 99, 308-313.	2.4	7
100	Molecular characterization of FRAXE-positive subjects with mental impairment in two unrelated Italian families. American Journal of Medical Genetics Part A, 1998, 75, 304-308.	2.4	13
101	Search for mutations in pancreatic sufficient cystic fibrosis Italian patients: detection of 90% of molecular defects and identification of three novel mutations. Human Genetics, 1995, 96, 312-8.	3.8	23
102	Genetic diagnosis by chorionic villus sampling before 8 gestational weeks: Efficiency, reliability, and risks on 317 completed pregnancies. Prenatal Diagnosis, 1992, 12, 789-799.	2.3	86
103	Frequency of Cystic Fibrosis Mutations Among Italian Patients. Advances in Experimental Medicine and Biology, 1991, 290, 387-390.	1.6	0
104	Frequency of the Î"F508 mutation in a sample of 175 Italian cystic fibrosis patients. Human Genetics, 1990, 85, 400-402.	3.8	9
105	Prenatal diagnosis and linkage disequilibrium with cystic fibrosis for markers surrounding D7S8. Human Genetics, 1990, 85, 275-8.	3.8	3