

Massimo Zollo

List of Publications by Year in descending order

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136
papers

8,276
citations

81900

39
h-index

48315

88
g-index

139
all docs

139
docs citations

139
times ranked

12443
citing authors

#	ARTICLE	IF	CITATIONS
1	A New Butyrate Releaser Exerts a Protective Action against SARS-CoV-2 Infection in Human Intestine. <i>Molecules</i> , 2022, 27, 862.	3.8	18
2	Loss of Detection of sgN Precedes Viral Abridged Replication in COVID-19-Affected Patientsâ€™ A Target for SARS-CoV-2 Propagation. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1941.	4.1	4
3	SARS-CoV-2 Pandemic Tracing in Italy Highlights Lineages with Mutational Burden in Growing Subsets. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4155.	4.1	3
4	Molecular detection of SARS-CoV-2 eta VOI in Northern Italy: a case report. <i>Clinical Chemistry and Laboratory Medicine</i> , 2022, 60, 61-63.	2.3	0
5	Germline rare variants of lectin pathway genes predispose to asymptomatic SARS-CoV-2 infection in elderly individuals. <i>Genetics in Medicine</i> , 2022, , .	2.4	7
6	A Novel Human Neutralizing mAb Recognizes Delta, Gamma and Omicron Variants of SARS-CoV-2 and Can Be Used in Combination with Sotrovimab. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5556.	4.1	3
7	SARS-CoV-2 Subgenomic N (sgN) Transcripts in Oro-Nasopharyngeal Swabs Correlate with the Highest Viral Load, as Evaluated by Five Different Molecular Methods. <i>Diagnostics</i> , 2021, 11, 288.	2.6	25
8	The metastasis suppressor protein NM23-H1 modulates the PI3K-AKT axis through interaction with the p110 α catalytic subunit. <i>Oncogenesis</i> , 2021, 10, 34.	4.9	10
9	Common variants at 21q22.3 locus influence MX1 and TMPRSS2 gene expression and susceptibility to severe COVID-19. <i>IScience</i> , 2021, 24, 102322.	4.1	60
10	Seroprevalence of SARS-CoV-2-specific antibodies in the town of Ariano Irpino (Avellino, Campania,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	4.9	12
11	Regulatory Noncoding and Predicted Pathogenic Coding Variants of CCR5 Predispose to Severe COVID-19. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5372.	4.1	16
12	Novel human neutralizing mAbs specific for Spike-RBD of SARS-CoV-2. <i>Scientific Reports</i> , 2021, 11, 11046.	3.3	13
13	The TNFRSF13C H159Y Variant Is Associated with Severe COVID-19: A Retrospective Study of 500 Patients from Southern Italy. <i>Genes</i> , 2021, 12, 881.	2.4	12
14	Long-chain polyphosphates impair SARS-CoV-2 infection and replication. <i>Science Signaling</i> , 2021, 14, .	3.6	27
15	Prune-1 drives polarization of tumor-associated macrophages (TAMs) within the lung metastatic niche in triple-negative breast cancer. <i>IScience</i> , 2021, 24, 101938.	4.1	11
16	Functional Genomics of PRUNE1 in Neurodevelopmental Disorders (NDDs) Tied to Medulloblastoma (MB) and Other Tumors. <i>Frontiers in Oncology</i> , 2021, 11, 758146.	2.8	7
17	Protective effects elicited by cow milk fermented with <i>L. Paracasei</i> CBAL74 against SARS-CoV-2 infection in human enterocytes. <i>Journal of Functional Foods</i> , 2021, 87, 104787.	3.4	9
18	Identification of SARS-CoV-2 Proteins from Nasopharyngeal Swabs Probed by Multiple Reaction Monitoring Tandem Mass Spectrometry. <i>ACS Omega</i> , 2021, 6, 34945-34953.	3.5	10

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19	Durability of Humoral Immune Responses to SARS-CoV-2 in Citizens of Ariano Irpino (Campania, Italy): A Longitudinal Observational Study With an 11.5-Month Follow-Up. <i>Frontiers in Public Health</i> , 2021, 9, 801609.	2.7	5
20	A Structurally Simple Vaccine Candidate Reduces Progression and Dissemination of Triple-Negative Breast Cancer. <i>IScience</i> , 2020, 23, 101250.	4.1	14
21	Pattern of Relapse and Treatment Response in WNT-Activated Medulloblastoma. <i>Cell Reports Medicine</i> , 2020, 1, 100038.	6.5	24
22	Biallelic mutations in neurofascin cause neurodevelopmental impairment and peripheral demyelination. <i>Brain</i> , 2019, 142, 2948-2964.	7.6	43
23	Metastatic group 3 medulloblastoma is driven by PRUNE1 targeting NME1-OTX2-SNAIL via PTEN inhibition. <i>Brain</i> , 2018, 141, 1300-1319.	7.6	22
24	A competitive cell-permeable peptide impairs Nme-1 (NDPK-A) and Prune-1 interaction: therapeutic applications in cancer. <i>Laboratory Investigation</i> , 2018, 98, 571-581.	3.7	5
25	RIP1-HAT1-SIRT Complex Identification and Targeting in Treatment and Prevention of Cancer. <i>Clinical Cancer Research</i> , 2018, 24, 2886-2900.	7.0	40
26	MBRS-52. TARGETING PRUNE-1 IN A GEMM OF METASTATIC MEDULLOBLASTOMA: A POTENTIAL ROUTE OF INHIBITION FOR NEW FUTURE THERAPIES. <i>Neuro-Oncology</i> , 2018, 20, i139-i139.	1.2	0
27	Heterogeneity within the PF-EPN-B ependymoma subgroup. <i>Acta Neuropathologica</i> , 2018, 136, 227-237.	7.7	86
28	Immunotherapy Bridge 2017 and Melanoma Bridge 2017: meeting abstracts. <i>Journal of Translational Medicine</i> , 2018, 16, .	4.4	2
29	Intertumoral Heterogeneity within Medulloblastoma Subgroups. <i>Cancer Cell</i> , 2017, 31, 737-754.e6.	16.8	836
30	The phenotypic and molecular spectrum of PEHO syndrome and PEHO-like disorders. <i>Brain</i> , 2017, 140, e49-e49.	7.6	33
31	PRUNE is crucial for normal brain development and mutated in microcephaly with neurodevelopmental impairment. <i>Brain</i> , 2017, 140, 940-952.	7.6	62
32	Reply: PRUNE1: a disease-causing gene for secondary microcephaly. <i>Brain</i> , 2017, 140, e62-e62.	7.6	0
33	In vivo bioluminescence imaging using orthotopic xenografts towards patient's derived-xenograft Medulloblastoma models. <i>Quarterly Journal of Nuclear Medicine and Molecular Imaging</i> , 2017, 61, 95-101.	0.7	5
34	Fenretinide (4-HPR) Targets Caspase-9, ERK 1/2 and the Wnt3a/ β 2-Catenin Pathway in Medulloblastoma Cells and Medulloblastoma Cell Spheroids. <i>PLoS ONE</i> , 2016, 11, e0154111.	2.5	24
35	Therapeutic Impact of Cytoreductive Surgery and Irradiation of Posterior Fossa Ependymoma in the Molecular Era: A Retrospective Multicohort Analysis. <i>Journal of Clinical Oncology</i> , 2016, 34, 2468-2477.	1.6	160
36	Natural compounds for pediatric cancer treatment. <i>Naunyn-Schmiedeberg's Archives of Pharmacology</i> , 2016, 389, 131-149.	3.0	26

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37	Prune. , 2016, , 3840-3843.		0
38	Glioblastoma stem cells differentiation through epigenetic modulation is driven by miR-296-5p/HMGA1/Sox2 axis. Translational Cancer Research, 2016, 5, S782-S788.	1.0	2
39	<i>M/R</i> β4a deficiency accelerates medulloblastoma formation <i>in vivo</i> . International Journal of Cancer, 2015, 136, 2293-2303.	5.1	40
40	A functional connectome: regulation of Wnt/TCF-dependent transcription by pairs of pathway activators. Molecular Cancer, 2015, 14, 206.	19.2	15
41	Molecular Biology and Genetics of Medulloblastoma. , 2015, , 265-286.		0
42	Progress on Nme (NDP kinase/Nm23/Awd) gene family-related functions derived from animal model systems: studies on development, cardiovascular disease, and cancer metastasis exemplified. Naunyn-Schmiedeberg's Archives of Pharmacology, 2015, 388, 109-117.	3.0	13
43	A therapeutic approach to treat prostate cancer by targeting Nm23-H1/h-Prune interaction. Naunyn-Schmiedeberg's Archives of Pharmacology, 2015, 388, 257-269.	3.0	20
44	Tissue invasion and metastasis: Molecular, biological and clinical perspectives. Seminars in Cancer Biology, 2015, 35, S244-S275.	9.6	408
45	Designing a broad-spectrum integrative approach for cancer prevention and treatment. Seminars in Cancer Biology, 2015, 35, S276-S304.	9.6	220
46	Early Targets of miR-34a in Neuroblastoma. Molecular and Cellular Proteomics, 2014, 13, 2114-2131.	3.8	29
47	Cytogenetic Prognostication Within Medulloblastoma Subgroups. Journal of Clinical Oncology, 2014, 32, 886-896.	1.6	263
48	H-Prune through GSK-3β interaction sustains canonical WNT/β-catenin signaling enhancing cancer progression in NSCLC. Oncotarget, 2014, 5, 5736-5749.	1.8	42
49	Immune Cells Within the Tumor Microenvironment. , 2014, , 1-23.		2
50	MicroRNA 199b-5p delivery through stable nucleic acid lipid particles (SNALPs) in tumorigenic cell lines. Naunyn-Schmiedeberg's Archives of Pharmacology, 2013, 386, 287-302.	3.0	30
51	Genetics of recurrent medulloblastoma. Lancet Oncology, The, 2013, 14, 1147-1148.	10.7	7
52	Dipyridamole prevents triple-negative breast-cancer progression. Clinical and Experimental Metastasis, 2013, 30, 47-68.	3.3	48
53	TERT promoter mutations are highly recurrent in SHH subgroup medulloblastoma. Acta Neuropathologica, 2013, 126, 917-929.	7.7	146
54	Epigenetic Silencing of DKK3 in Medulloblastoma. International Journal of Molecular Sciences, 2013, 14, 7492-7505.	4.1	18

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55	Mapping Functional Interaction Sites of Human Prune C-terminal Domain by NMR Spectroscopy in Human Cell Lysates. <i>Chemistry - A European Journal</i> , 2013, 19, 12217-12220.	3.3	12
56	Neuroblastoma tumorigenesis is regulated through the Nm23-H1/h-Prune C-terminal interaction. <i>Scientific Reports</i> , 2013, 3, 1351.	3.3	34
57	gH625 is a viral derived peptide for effective delivery of intrinsically disordered proteins. <i>International Journal of Nanomedicine</i> , 2013, 8, 2555.	6.7	20
58	GSK3 β regulates physiological migration of stem/progenitor cells via cytoskeletal rearrangement. <i>Journal of Clinical Investigation</i> , 2013, 123, 1705-1717.	8.2	32
59	GSK3 β regulates physiological migration of stem/progenitor cells via cytoskeletal rearrangement. <i>Journal of Clinical Investigation</i> , 2013, 123, 3183-3183.	8.2	0
60	The micro-RNA 199b-5p regulatory circuit involves Hes1, CD15, and epigenetic modifications in medulloblastoma. <i>Neuro-Oncology</i> , 2012, 14, 596-612.	1.2	48
61	The metallophosphodiesterase Mpped2 impairs tumorigenesis in neuroblastoma. <i>Cell Cycle</i> , 2012, 11, 569-581.	2.6	30
62	Subgroup-specific structural variation across 1,000 medulloblastoma genomes. <i>Nature</i> , 2012, 488, 49-56.	27.8	761
63	Novel pyrimidopyrimidine derivatives for inhibition of cellular proliferation and motility induced by h-prune in breast cancer. <i>European Journal of Medicinal Chemistry</i> , 2012, 57, 41-50.	5.5	22
64	Spatial and temporal expressions of prune reveal a role in Müller gliogenesis during <i>Xenopus</i> retinal development. <i>Gene</i> , 2012, 509, 93-103.	2.2	8
65	Targeting monocyte chemotactic protein-1 synthesis with bindarit induces tumor regression in prostate and breast cancer animal models. <i>Clinical and Experimental Metastasis</i> , 2012, 29, 585-601.	3.3	84
66	Molecular networks that regulate cancer metastasis. <i>Seminars in Cancer Biology</i> , 2012, 22, 234-249.	9.6	296
67	Tumor microenvironment: a main actor in the metastasis process. <i>Clinical and Experimental Metastasis</i> , 2012, 29, 381-395.	3.3	155
68	Norcantharidin impairs medulloblastoma growth by inhibition of Wnt/ β -catenin signaling. <i>Journal of Neuro-Oncology</i> , 2012, 106, 59-70.	2.9	36
69	Loss of 10q26.1-q26.3 in association with 7q34-q36.3 gain or 17q24.3-q25.3 gain predict poor outcome in pediatric medulloblastoma. <i>Cancer Letters</i> , 2011, 308, 215-224.	7.2	3
70	Upregulation of miR-21 by Ras in vivo and its role in tumor growth. <i>Oncogene</i> , 2011, 30, 275-286.	5.9	130
71	Correlation of NM23-H1 cytoplasmic expression with metastatic stage in human prostate cancer tissue. <i>Naunyn-Schmiedeberg's Archives of Pharmacology</i> , 2011, 384, 489-498.	3.0	25
72	A critical evaluation of biochemical activities reported for the nucleoside diphosphate kinase/Nm23/Awd family proteins: opportunities and missteps in understanding their biological functions. <i>Naunyn-Schmiedeberg's Archives of Pharmacology</i> , 2011, 384, 331-339.	3.0	37

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73	Detection of erbB2 copy number variations in plasma of patients with esophageal carcinoma. <i>BMC Cancer</i> , 2011, 11, 126.	2.6	22
74	MiR-34a Targeting of Notch Ligand Delta-Like 1 Impairs CD15+/CD133+ Tumor-Propagating Cells and Supports Neural Differentiation in Medulloblastoma. <i>PLoS ONE</i> , 2011, 6, e24584.	2.5	149
75	<i>Prune.</i> , 2011, , 3108-3111.		0
76	ASAP1 promotes tumor cell motility and invasiveness, stimulates metastasis formation in vivo, and correlates with poor survival in colorectal cancer patients. <i>Oncogene</i> , 2010, 29, 2393-2403.	5.9	85
77	The Quassinoid Derivative NBT-272 Targets Both the AKT and ERK Signaling Pathways in Embryonal Tumors. <i>Molecular Cancer Therapeutics</i> , 2010, 9, 3145-3157.	4.1	14
78	Regulation of divalent metal transporter 1 (DMT1) non-IRE isoform by the microRNA Let-7d in erythroid cells. <i>Haematologica</i> , 2010, 95, 1244-1252.	3.5	82
79	The miR-17-92 MicroRNA Cluster Regulates Multiple Components of the TGF- β Pathway in Neuroblastoma. <i>Molecular Cell</i> , 2010, 40, 762-773.	9.7	279
80	GSK3 β Signaling Regulates the Motility of Hematopoietic Progenitors Via Prune.. <i>Blood</i> , 2010, 116, 1553-1553.	1.4	0
81	MicroRNA-199b-5p Impairs Cancer Stem Cells through Negative Regulation of HES1 in Medulloblastoma. <i>PLoS ONE</i> , 2009, 4, e4998.	2.5	233
82	The Nm23-H1-h-Prune complex in cellular physiology: a "tip of the iceberg" protein network perspective. <i>Molecular and Cellular Biochemistry</i> , 2009, 329, 149-159.	3.1	32
83	Xenopus NM23-X4 regulates retinal gliogenesis through interaction with p27Xic1. <i>Neural Development</i> , 2009, 4, 1.	2.4	33
84	Galectin-1 is a major effector of TrkB-mediated neuroblastoma aggressiveness. <i>Oncogene</i> , 2009, 28, 2015-2023.	5.9	61
85	Phosphorylation of nm23-H1 by CKI induces its complex formation with h-prune and promotes cell motility. <i>Oncogene</i> , 2008, 27, 1853-1864.	5.9	48
86	Investigating h-Prune activation of Wnt signalling in breast cancer. <i>Breast Cancer Research</i> , 2008, 10, .	5.0	0
87	Human Metastasis Regulator Protein H-Prune is a Short-Chain Exopolyphosphatase. <i>Biochemistry</i> , 2008, 47, 9707-9713.	2.5	120
88	Regulation of DMT1 (non IRE isoform) by MicroRNA LET-7D. <i>Blood</i> , 2008, 112, 416-416.	1.4	0
89	<i>Prune.</i> , 2008, , 2487-2489.		0
90	Domain mapping on the human metastasis regulator protein h-Prune reveals a C-terminal dimerization domain. <i>Biochemical Journal</i> , 2007, 407, 199-205.	3.7	16

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91	New mutations identified in the ocular albinism type 1 gene. <i>Gene</i> , 2007, 402, 20-27.	2.2	5
92	Comparative Proteomic Expression Profile in All-transRetinoic Acid Differentiated Neuroblastoma Cell Line. <i>Journal of Proteome Research</i> , 2007, 6, 2550-2564.	3.7	30
93	Molecular classification of nodal metastasis in primary larynx squamous cell carcinoma. <i>Translational Research</i> , 2007, 150, 233-245.	5.0	25
94	Understanding h-prune biology in the fight against cancer. <i>Clinical and Experimental Metastasis</i> , 2007, 24, 637-645.	3.3	19
95	H-prune-nm23-H1 protein complex and correlation to pathways in cancer metastasis. <i>Journal of Bioenergetics and Biomembranes</i> , 2006, 38, 205-213.	2.3	33
96	PRUNE and NM23-M1 expression in embryonic and adult mouse brain. <i>Journal of Bioenergetics and Biomembranes</i> , 2006, 38, 233-246.	2.3	30
97	Glycogen Synthase Kinase 3 and h-prune Regulate Cell Migration by Modulating Focal Adhesions. <i>Molecular and Cellular Biology</i> , 2006, 26, 898-911.	2.3	111
98	Telencephalic Embryonic Subtractive Sequences: A Unique Collection of Neurodevelopmental Genes. <i>Journal of Neuroscience</i> , 2005, 25, 7586-7600.	3.6	6
99	Tagging genes with cassette-exchange sites. <i>Nucleic Acids Research</i> , 2005, 33, e44-e44.	14.5	18
100	Overexpression of h-prune in breast cancer is correlated with advanced disease status. <i>Clinical Cancer Research</i> , 2005, 11, 199-205.	7.0	32
101	Unraveling Genes and Pathways Influenced by H-Prune PDE Overexpression: A Model to Study Cellular Motility. <i>Cell Cycle</i> , 2004, 3, 756-759.	2.6	33
102	Prune cAMP phosphodiesterase binds nm23-H1 and promotes cancer metastasis. <i>Cancer Cell</i> , 2004, 5, 137-149.	16.8	132
103	Unraveling genes and pathways influenced by H-prune PDE overexpression: a model to study cellular motility. <i>Cell Cycle</i> , 2004, 3, 758-61.	2.6	15
104	Method to express and purify nm23-H2 protein from baculovirus-infected cells. <i>BioTechniques</i> , 2003, 35, 384-391.	1.8	12
105	Allelic polymorphisms in the transcriptional regulatory region of human SEL1L. <i>Mutation Research - Mutation Research Genomics</i> , 2001, 458, 71-76.	1.1	5
106	Complete mutation scanning of the human SEL 1L gene: a candidate gene for type 1 diabetes. <i>Acta Diabetologica</i> , 2001, 38, 191-192.	2.5	8
107	No evidence forSEL1L as a candidate gene forIDDM11-conferred susceptibility. <i>Diabetes/Metabolism Research and Reviews</i> , 2001, 17, 292-295.	4.0	14
108	Amplification and overexpression of PRUNE in human sarcomas and breast carcinomasâ€“a possible mechanism for altering the nm23-H1 activity. <i>Oncogene</i> , 2001, 20, 6881-6890.	5.9	52

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109	SEL1L, the human homolog of <i>C. elegans</i> sel-1: refined physical mapping, gene structure and identification of polymorphic markers. <i>Human Genetics</i> , 2000, 106, 227-235.	3.8	20
110	Identification of two paralogous regions mapping to the short and long arms of human chromosome 2 comprising LIS1 pseudogenes. <i>Cytogenetic and Genome Research</i> , 1999, 86, 225-232.	1.1	6
111	Evidence for interaction between human PRUNE and nm23-H1 NDPKinase. <i>Oncogene</i> , 1999, 18, 7244-7252.	5.9	77
112	Host response to EBV infection in X-linked lymphoproliferative disease results from mutations in an SH2-domain encoding gene. <i>Nature Genetics</i> , 1998, 20, 129-135.	21.4	720
113	A new candidate region for the positional cloning of the XLP gene. <i>European Journal of Human Genetics</i> , 1998, 6, 509-517.	2.8	11
114	FACL4, a New Gene Encoding Long-Chain Acyl-CoA Synthetase 4, Is Deleted in a Family with Alport Syndrome, Elliptocytosis, and Mental Retardation. <i>Genomics</i> , 1998, 47, 350-358.	2.9	114
115	The Gene Encoding a Cationic Amino Acid Transporter (SLC7A4) Maps to the Region Deleted in the Velocardiofacial Syndrome. <i>Genomics</i> , 1998, 49, 230-236.	2.9	52
116	A method for point mutation analysis that links SSCP and dye primer fluorescent sequencing. <i>Molecular and Cellular Probes</i> , 1998, 12, 125-131.	2.1	4
117	Sequencing Analysis of Forty-Eight Human Image cDNA Clones Similar to <i>Drosophila</i> Mutant Protein. <i>DNA Sequence</i> , 1998, 9, 307-315.	0.7	5
118	A Method to Direct Sequence Cosmid LAWRIST16 Clones. <i>DNA Sequence</i> , 1997, 7, 229-233.	0.7	2
119	Identification by Shotgun Sequencing, Genomic Organization, and Functional Analysis of a Fourth Arylsulfatase Gene (ARSF) from the Xp22.3 Region. <i>Genomics</i> , 1997, 42, 192-199.	2.9	37
120	A semi-automated method for preparing high-quality plasmid templates in 96-well format ready for automated DNA sequencing. <i>Technical Tips Online</i> , 1997, 2, 184-187.	0.2	0
121	Rox, a novel bHLHZip protein expressed in quiescent cells that heterodimerizes with Max, binds a non-canonical E box and acts as a transcriptional repressor. <i>EMBO Journal</i> , 1997, 16, 2892-2906.	7.8	126
122	A mammalian homologue of the <i>Drosophila</i> retinal degeneration B gene: implications for the evolution of phototransduction mechanisms. <i>Genes and Function</i> , 1997, 1, 205-213.	2.8	16
123	Molecular Cloning, Expression Pattern, and Chromosomal Localization of the Human Na ⁺ /Cl ⁻ Thiiazide-Sensitive Cotransporter (SLC12A3). <i>Genomics</i> , 1996, 35, 486-493.	2.9	123
124	Identification and mapping of human cDNAs homologous to <i>Drosophila</i> mutant genes through EST database searching. <i>Nature Genetics</i> , 1996, 13, 167-174.	21.4	177
125	Long-range sequence analysis in Xq28: thirteen known and six candidate genes in 219.4 kb of high GC DNA between the RCP/GCP and G6PD loci. <i>Human Molecular Genetics</i> , 1996, 5, 659-668.	2.9	69
126	Sequence and gene content in 52 kb including and centromeric to the G6PD gene in Xq28. <i>DNA Sequence</i> , 1995, 6, 1-11.	0.7	11

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127	A manual high-throughput M13 DNA preparation. <i>BioTechniques</i> , 1994, 16, 370-2.	1.8	5
128	Use of a sensitive fluorescent intercalating dye to detect PCR products of low copy number and high molecular weight.. <i>Genome Research</i> , 1993, 3, 115-119.	5.5	7
129	Sequence of mouse glucose-6-phosphate dehydrogenase cDNA. <i>DNA Sequence</i> , 1993, 3, 319-322.	0.7	26
130	Screening for steroid sulfatase (STS) gene deletions by multiplex DNA amplification. <i>Human Genetics</i> , 1990, 84, 571-3.	3.8	47
131	Molecular heterogeneity of steroid sulfatase deficiency: A multicenter study on 57 unrelated patients, at DNA and protein levels. <i>Genomics</i> , 1989, 4, 36-40.	2.9	102
132	Molecular characterization of a gene of the 'EGF family' expressed in undifferentiated human NTERA2 teratocarcinoma cells. <i>EMBO Journal</i> , 1989, 8, 1987-91.	7.8	68
133	Molecular studies of marsupial X chromosomes reveal limited sequence homology of mammalian X-linked genes. <i>Genomics</i> , 1987, 1, 19-28.	2.9	21
134	MicroRNAs and Cancer Stem Cells in Medulloblastoma. , 0, , .		0
135	Common Variants at 21q22.3 Locus Influence <i>MX1</i> Gene Expression and Susceptibility to Severe COVID-19. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0
136	Pattern of Relapse and Treatment Response in WNT-Activated Medulloblastoma. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0