## Jessica I Hoell

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5044691/publications.pdf

Version: 2024-02-01

43 papers

2,859 citations

331538
21
h-index

42 g-index

44 all docs

44 docs citations

44 times ranked 6400 citing authors

#	Article	IF	CITATIONS
1	YBX1 mediates translation of oncogenic transcripts to control cell competition in AML. Leukemia, 2022, 36, 426-437.	3.3	18
2	Mutational mechanisms shaping the coding and noncoding genome of germinal center derived B-cell lymphomas. Leukemia, 2021, 35, 2002-2016.	3.3	34
3	Fragile X mental retardation protein protects against tumour necrosis factor-mediated cell death and liver injury. Gut, 2020, 69, 133-145.	6.1	14
4	RNA-Binding Proteins in Acute Leukemias. International Journal of Molecular Sciences, 2020, 21, 3409.	1.8	36
5	YBX1 Indirectly Targets Heterochromatin-Repressed Inflammatory Response-Related Apoptosis Genes through Regulating CBX5 mRNA. International Journal of Molecular Sciences, 2020, 21, 4453.	1.8	11
6	Facing the large variety of life-limiting conditions in children. European Journal of Pediatrics, 2019, 178, 1893-1902.	1.3	28
7	Mutated SON putatively causes a cancer syndrome comprising high-risk medulloblastoma combined with café-au-lait spots. Familial Cancer, 2019, 18, 353-358.	0.9	4
8	An alternative approach to establishing unbiased colorectal cancer risk estimation in Lynch syndrome. Genetics in Medicine, 2019, 21, 2706-2712.	1.1	11
9	Genomic and transcriptomic changes complement each other in the pathogenesis of sporadic Burkitt lymphoma. Nature Communications, 2019, 10, 1459.	5.8	99
10	Pediatric ALL relapses after allo-SCT show high individuality, clonal dynamics, selective pressure, and druggable targets. Blood Advances, 2019, 3, 3143-3156.	2.5	4
11	Deciphering human ribonucleoprotein regulatory networks. Nucleic Acids Research, 2019, 47, 570-581.	<b>6.</b> 5	54
12	Diagnostic challenges in a child with early onset desmoplastic medulloblastoma and homozygous variants in MSH2 and MSH6. European Journal of Human Genetics, 2018, 26, 440-444.	1.4	18
13	Outcome of relapse after allogeneic <scp>HSCT</scp> in children with <scp>ALL</scp> enrolled in the <scp>ALL</scp> â€ <scp>SCT</scp> 2003/2007 trial. British Journal of Haematology, 2018, 180, 82-89.	1.2	50
14	Tumor Necrosis Factor-Mediated Survival of CD169 <sup>+</sup> Cells Promotes Immune Activation during Vesicular Stomatitis Virus Infection. Journal of Virology, 2018, 92, .	1.5	16
15	Challenges of palliative care in children with inborn metabolic diseases. Orphanet Journal of Rare Diseases, 2018, 13, 112.	1.2	8
16	Transcriptome-wide analysis uncovers the targets of the RNA-binding protein MSI2 and effects of MSI2's RNA-binding activity on IL-6 signaling. Journal of Biological Chemistry, 2018, 293, 15359-15369.	1.6	31
17	Advance care planning and outcome in pediatric palliative home care. Oncotarget, 2018, 9, 17867-17875.	0.8	15
18	Tâ $\in$ ell acute lymphoblastic leukemia in infants has distinct genetic and epigenetic features compared to childhood cases. Genes Chromosomes and Cancer, 2017, 56, 159-167.	1.5	15

#	Article	lF	Citations
19	Palliative care for children with a yet undiagnosed syndrome. European Journal of Pediatrics, 2017, 176, 1319-1327.	1.3	16
20	End-of-life care in children with hematologic malignancies. Oncotarget, 2017, 8, 89939-89948.	0.8	15
21	Immunohistochemical detection of inhibitor of DNA binding 3 mutational variants in mature aggressive B-cell lymphoma. Haematologica, 2016, 101, e259-e261.	1.7	2
22	Effective treatment of spasticity using dronabinol in pediatric palliative care. European Journal of Paediatric Neurology, 2016, 20, 898-903.	0.7	37
23	Experiences in palliative home care of infants with life-limiting conditions. European Journal of Pediatrics, 2016, 175, 321-327.	1.3	12
24	Symptoms and management of pediatric patients with incurable brain tumors in palliative home care. European Journal of Paediatric Neurology, 2016, 20, 261-269.	0.7	24
25	The PARA-suite: PAR-CLIP specific sequence read simulation and processing. PeerJ, 2016, 4, e2619.	0.9	8
26	Palliative Outpatient Care in Children with Hematologic Malignancies Compared to Solid Tumors. Blood, 2016, 128, 5977-5977.	0.6	0
27	The <i>PCBP1</i> gene encoding poly(rc) binding protein i is recurrently mutated in <scp>B</scp> urkitt lymphoma. Genes Chromosomes and Cancer, 2015, 54, 555-564.	1.5	29
28	Biochemical and bioinformatic methods for elucidating the role of RNA–protein interactions in posttranscriptional regulation. Briefings in Functional Genomics, 2015, 14, 102-114.	1.3	10
29	Genomics and drug profiling of fatal TCF3-HLFâ^positive acute lymphoblastic leukemia identifies recurrent mutation patterns and therapeutic options. Nature Genetics, 2015, 47, 1020-1029.	9.4	190
30	DNA methylome analysis in Burkitt and follicular lymphomas identifies differentially methylated regions linked to somatic mutation and transcriptional control. Nature Genetics, 2015, 47, 1316-1325.	9.4	119
31	MINCR is a MYC-induced IncRNA able to modulate MYC's transcriptional network in Burkitt lymphoma cells. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E5261-70.	3.3	91
32	The Cell Cycle Regulator CCDC6 Is a Key Target of RNA-Binding Protein EWS. PLoS ONE, 2015, 10, e0119066.	1.1	9
33	PAR-CLIP (Photoactivatable Ribonucleoside-Enhanced Crosslinking and Immunoprecipitation). Methods in Enzymology, 2014, 539, 113-161.	0.4	90
34	Recurrent <i>RHOA</i> mutations in pediatric <scp>B</scp> urkitt lymphoma treated according to the NHLâ€BFM protocols. Genes Chromosomes and Cancer, 2014, 53, 911-916.	1.5	51
35	Rapid Creation of Stable Mammalian Cell Lines for Regulated Expression of Proteins Using the Gateway® Recombination Cloning Technology and Flp-In T-REx® Lines. Methods in Enzymology, 2013, 529, 99-124.	0.4	22
36	MicroRNAs in Human Cancer. Advances in Experimental Medicine and Biology, 2013, 774, 1-20.	0.8	606

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37	Wholeâ€genome pairedâ€end analysis confirms remarkable genomic stability of atypical teratoid/rhabdoid tumors. Genes Chromosomes and Cancer, 2013, 52, 983-985.	1.5	4
38	Comprehensive Analysis of Long Non-Coding RNAs in Ovarian Cancer Reveals Global Patterns and Targeted DNA Amplification. PLoS ONE, 2013, 8, e80306.	1.1	90
39	In-Depth miRNA Profiling Of Germinal Center Derived B-Cell Lymphomas By Next Generation Sequencing: A Report From The German Icgc-Mmml-Seq Project. Blood, 2013, 122, 2500-2500.	0.6	4
40	Recurrent mutation of the ID3 gene in Burkitt lymphoma identified by integrated genome, exome and transcriptome sequencing. Nature Genetics, 2012, 44, 1316-1320.	9.4	389
41	RNA targets of wild-type and mutant FET family proteins. Nature Structural and Molecular Biology, 2011, 18, 1428-1431.	3.6	321
42	mRNA and protein levels of FUS, EWSR1, and TAF15 are upregulated in liposarcoma. Genes Chromosomes and Cancer, 2011, 50, 338-347.	1.5	31
43	Targeted Mutation Reveals Essential Functions of the Homeodomain Transcription Factor Shox2 in Sinoatrial and Pacemaking Development. Circulation, 2007, 115, 1830-1838.	1.6	222