

Alejandra Cervera

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

2,516
citations

840776

11
h-index

677142

22
g-index

29
all docs

29
docs citations

29
times ranked

6038
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic Profile in a Non-Seminoma Testicular Germ-Cell Tumor Cohort Reveals a Potential Biomarker of Sensitivity to Platinum-Based Therapy. <i>Cancers</i> , 2022, 14, 2065.	3.7	5
2	FUNGI: FUSioN Gene Integration toolset. <i>Bioinformatics</i> , 2021, 37, 3353-3355.	4.1	1
3	Human cell transformation by combined lineage conversion and oncogene expression. <i>Oncogene</i> , 2021, 40, 5533-5547.	5.9	12
4	Functional Profiling of FSH and Estradiol in Ovarian Granulosa Cell Tumors. <i>Journal of the Endocrine Society</i> , 2020, 4, bvaa034.	0.2	13
5	Distinct subtypes of diffuse large B-cell lymphoma defined by hypermutated genes. <i>Leukemia</i> , 2019, 33, 2662-2672.	7.2	24
6	Anduril 2: upgraded large-scale data integration framework. <i>Bioinformatics</i> , 2019, 35, 3815-3817.	4.1	31
7	Anagrelide for Gastrointestinal Stromal Tumor. <i>Clinical Cancer Research</i> , 2019, 25, 1676-1687.	7.0	14
8	Modified penetrance of coding variants by cis-regulatory variation contributes to disease risk. <i>Nature Genetics</i> , 2018, 50, 1327-1334.	21.4	167
9	Deltex-1 mutations predict poor survival in diffuse large B-cell lymphoma. <i>Haematologica</i> , 2017, 102, e195-e198.	3.5	23
10	Alternative splicing discriminates molecular subtypes and has prognostic impact in diffuse large B-cell lymphoma. <i>Blood Cancer Journal</i> , 2017, 7, e596-e596.	6.2	14
11	MicroRNAs regulate key cell survival pathways and mediate chemosensitivity during progression of diffuse large B-cell lymphoma. <i>Blood Cancer Journal</i> , 2017, 7, 654.	6.2	26
12	SePIA: RNA and small RNA sequence processing, integration, and analysis. <i>BioData Mining</i> , 2016, 9, 20.	4.0	25
13	A survey of best practices for RNA-seq data analysis. <i>Genome Biology</i> , 2016, 17, 13.	8.8	1,898
14	Low Expression and Somatic Mutations of the KLHL6 Gene Predict Poor Survival in Patients with Activated B-Cell like Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2016, 128, 2926-2926.	1.4	3
15	Low Expression of the CIITA Gene Predicts Poor Outcome in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2016, 128, 2948-2948.	1.4	2
16	Gene expression profiling of pre-eclamptic placentae by RNA sequencing. <i>Scientific Reports</i> , 2015, 5, 14107.	3.3	89
17	Somatic Mutations in E3 Ubiquitin Ligase Deltex 1 Are Associated with Survival in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2014, 124, 1688-1688.	1.4	0
18	Integrative Analysis of Deep Sequencing Data Identifies Estrogen Receptor Early Response Genes and Links ATAD3B to Poor Survival in Breast Cancer. <i>PLoS Computational Biology</i> , 2013, 9, e1003100.	3.2	11

#	ARTICLE	IF	CITATIONS
19	Global Profiling Of Outcome Associated Alternative Splicing Events and Gene Expression In Diffuse Large B-Cell Lymphoma. Blood, 2013, 122, 75-75.	1.4	0
20	Alternative Splicing and Expression of Class II Tubulin Beta (TUBB2B) Are Associated with Outcome in Diffuse Large B-Cell Lymphoma. Blood, 2012, 120, 1557-1557.	1.4	1
21	Authorship attribution as a case of anomaly detection: A neural network model. International Journal of Hybrid Intelligent Systems, 2011, 8, 225-235.	1.2	7
22	Detection of Different Authorship of Text Sequences through Self-organizing Maps and Mutual Information Function. Lecture Notes in Computer Science, 2010, , 186-195.	1.3	2
23	A Neural Network May Show the Best Way to Learn How to Count for Students in Elementary Math Courses. , 2008, , .		0
24	Decreasing Neighborhood Revisited in Self-Organizing Maps. Lecture Notes in Computer Science, 2008, , 671-679.	1.3	1