

Cristina Sisu

List of Publications by Year in descending order

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31
papers

43,037
citations

489802

18
h-index

488211

31
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36
all docs

36
docs citations

36
times ranked

79458
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential Expression of RAD51AP1 in Ovarian Cancer: Effects of siRNA In Vitro. <i>Journal of Personalized Medicine</i> , 2022, 12, 201.	1.1	1
2	In Silico Study to Predict the Structural and Functional Consequences of SNPs on Biomarkers of Ovarian Cancer (OC) and BPA Exposure-Associated OC. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1725.	1.8	4
3	Impact of Environmentally Relevant Concentrations of Bisphenol A (BPA) on the Gene Expression Profile in an In Vitro Model of the Normal Human Ovary. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5334.	1.8	9
4	GENCODE 2021. <i>Nucleic Acids Research</i> , 2021, 49, D916-D923.	6.5	633
5	Identification of Potential Bisphenol A (BPA) Exposure Biomarkers in Ovarian Cancer. <i>Journal of Clinical Medicine</i> , 2021, 10, 1979.	1.0	11
6	Non-redundant functions of H2A.Z.1 and H2A.Z.2 in chromosome segregation and cell cycle progression. <i>EMBO Reports</i> , 2021, 22, e52061.	2.0	23
7	GENCODE Pseudogenes. <i>Methods in Molecular Biology</i> , 2021, 2324, 67-82.	0.4	1
8	Pseudogenes as Biomarkers and Therapeutic Targets in Human Cancers. <i>Methods in Molecular Biology</i> , 2021, 2324, 319-337.	0.4	13
9	Is There a Link between Bisphenol A (BPA), a Key Endocrine Disruptor, and the Risk for SARS-CoV-2 Infection and Severe COVID-19?. <i>Journal of Clinical Medicine</i> , 2020, 9, 3296.	1.0	16
10	In Silico and In Vitro Analysis of lncRNA XIST Reveals a Panel of Possible Lung Cancer Regulators and a Five-Gene Diagnostic Signature. <i>Cancers</i> , 2020, 12, 3499.	1.7	9
11	Transcriptional activity and strain-specific history of mouse pseudogenes. <i>Nature Communications</i> , 2020, 11, 3695.	5.8	17
12	Perspectives on ENCODE. <i>Nature</i> , 2020, 583, 693-698.	13.7	123
13	Expanded encyclopaedias of DNA elements in the human and mouse genomes. <i>Nature</i> , 2020, 583, 699-710.	13.7	1,252
14	Differential expression of mTOR components in endometriosis and ovarian cancer: Effects of rapalogues and dual kinase inhibitors on mTORC1 and mTORC2 stoichiometry. <i>International Journal of Molecular Medicine</i> , 2019, 43, 47-56.	1.8	24
15	Deletions of Chromosome 7q Affect Nuclear Organization and HLXB9 Gene Expression in Hematological Disorders. <i>Cancers</i> , 2019, 11, 585.	1.7	21
16	Liquid Biopsies in Lung Cancer: Four Emerging Technologies and Potential Clinical Applications. <i>Cancers</i> , 2019, 11, 331.	1.7	13
17	GENCODE reference annotation for the human and mouse genomes. <i>Nucleic Acids Research</i> , 2019, 47, D766-D773.	6.5	2,350
18	Repeat associated mechanisms of genome evolution and function revealed by the <i>Mus caroli</i> and <i>Mus pahari</i> genomes. <i>Genome Research</i> , 2018, 28, 448-459.	2.4	99

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19	Sixteen diverse laboratory mouse reference genomes define strain-specific haplotypes and novel functional loci. <i>Nature Genetics</i> , 2018, 50, 1574-1583.	9.4	169
20	Loregic: A Method to Characterize the Cooperative Logic of Regulatory Factors. <i>PLoS Computational Biology</i> , 2015, 11, e1004132.	1.5	18
21	A global reference for human genetic variation. <i>Nature</i> , 2015, 526, 68-74.	13.7	13,998
22	Comparative analysis of pseudogenes across three phyla. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 13361-13366.	3.3	72
23	Comparative analysis of the transcriptome across distant species. <i>Nature</i> , 2014, 512, 445-448.	13.7	289
24	Integrating sequence and array data to create an improved 1000 Genomes Project haplotype reference panel. <i>Nature Communications</i> , 2014, 5, 3934.	5.8	364
25	Integrative Annotation of Variants from 1092 Humans: Application to Cancer Genomics. <i>Science</i> , 2013, 342, 1235587.	6.0	341
26	The GENCODE pseudogene resource. <i>Genome Biology</i> , 2012, 13, R51.	13.9	273
27	An integrated encyclopedia of DNA elements in the human genome. <i>Nature</i> , 2012, 489, 57-74.	13.7	15,516
28	An integrated map of genetic variation from 1,092 human genomes. <i>Nature</i> , 2012, 491, 56-65.	13.7	7,199
29	The Influence of Ligand Valency on Aggregation Mechanisms for Inhibiting Bacterial Toxins. <i>ChemBioChem</i> , 2009, 10, 329-337.	1.3	59
30	Classification of proteins based on similarity of two-dimensional protein maps. <i>Biophysical Chemistry</i> , 2008, 138, 11-22.	1.5	4
31	Strong Inhibition of Cholera Toxin by Multivalent GM1 Derivatives. <i>ChemBioChem</i> , 2007, 8, 1500-1503.	1.3	101