## Eugenia Galeota

List of Publications by Year in descending order

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840776 1058476 3,297 20 11 14 citations h-index g-index papers 21 21 21 6155 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Identification, semantic annotation and comparison of combinations of functional elements in multiple biological conditions. Bioinformatics, 2022, 38, 1183-1190.	4.1	1
2	Integrated longitudinal immunophenotypic, transcriptional, and repertoire analyses delineate immune responses in patients with COVID-19. Science Immunology, 2021, 6, .	11.9	108
3	A dual role of dLsd1 in oogenesis: regulating developmental genes and repressing transposons. Nucleic Acids Research, 2020, 48, 1206-1224.	14.5	5
4	Genome-wide dynamics of RNA synthesis, processing, and degradation without RNA metabolic labeling. Genome Research, 2020, 30, 1492-1507.	5.5	26
5	Ontology-driven integrative analysis of omics data through Onassis. Scientific Reports, 2020, 10, 703.	3.3	9
6	m6A-Dependent RNA Dynamics in T Cell Differentiation. Genes, 2019, 10, 28.	2.4	32
7	Stability and Evolution of Scientific Networks. , 2018, , 2972-2980.		O
8	Ontology-based annotations and semantic relations in large-scale (epi)genomics data. Briefings in Bioinformatics, 2017, 18, bbw036.	6.5	21
9	Stability and Evolution of Scientific Networks. , 2017, , 1-9.		0
10	Integrated Systems for NGS Data Management and Analysis: Open Issues and Available Solutions. Frontiers in Genetics, 2016, 7, 75.	2.3	45
11	The hierarchical organization of natural protein interaction networks confers self-organization properties on pseudocells. BMC Systems Biology, 2015, 9, S3.	3.0	5
12	Immune modulation of pathologic complete response after neoadjuvant HER2-directed therapies in the NeoSphere trial. Annals of Oncology, 2015, 26, 2429-2436.	1.2	106
13	Abstract P2-03-05: The heterogeneous clinical behavior of luminal breast cancers is associated with different mutational landscapes. , $2015,  ,  .$		O
14	The MIntAct projectâ€"IntAct as a common curation platform for 11 molecular interaction databases. Nucleic Acids Research, 2014, 42, D358-D363.	14.5	1,634
15	Stability and Evolution of Scientific Networks. , 2014, , 2048-2056.		O
16	An immune-related signature for prediction of risk of late recurrences beyond proliferation and ER-related genes in ER-positive breast cancer Journal of Clinical Oncology, 2014, 32, 530-530.	1.6	0
17	A new reference implementation of the PSICQUIC web service. Nucleic Acids Research, 2013, 41, W601-W606.	14.5	91
18	MINT, the molecular interaction database: 2012 update. Nucleic Acids Research, 2012, 40, D857-D861.	14.5	917

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#	Article	IF	CITATIONS
19	Selection in scientific networks. Social Network Analysis and Mining, 2012, 2, 229-237.	2.8	17
20	PSICQUIC and PSISCORE: accessing and scoring molecular interactions. Nature Methods, 2011, 8, 528-529.	19.0	274