Timothy Butler

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/384313/publications.pdf

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20 papers 2,183 citations

471371 17 h-index ⁷⁵²⁵⁷³
20
g-index

25 all docs

25 docs citations

25 times ranked

3571 citing authors

#	Article	IF	CITATIONS
1	Somatic mutation rates scale with lifespan across mammals. Nature, 2022, 604, 517-524.	13.7	211
2	Reliable detection of somatic mutations in solid tissues by laser-capture microdissection and low-input DNA sequencing. Nature Protocols, 2021, 16, 841-871.	5.5	82
3	Lineage tracing of human development through somatic mutations. Nature, 2021, 595, 85-90.	13.7	79
4	The mutational landscape of human somatic and germline cells. Nature, 2021, 597, 381-386.	13.7	180
5	Convergent somatic mutations in metabolism genes in chronic liver disease. Nature, 2021, 598, 473-478.	13.7	87
6	Using social network analysis to complete literature reviews: a new systematic approach for independent researchers to detect and interpret prominent research programs within large collections of relevant literature. International Journal of Social Research Methodology: Theory and Practice, 2020, 23, 483-496.	2.3	13
7	Somatic Evolution in Non-neoplastic IBD-Affected Colon. Cell, 2020, 182, 672-684.e11.	13.5	122
8	Transient commensal clonal interactions can drive tumor metastasis. Nature Communications, 2020, 11, 5799.	5.8	30
9	The mutational landscape of normal human endometrial epithelium. Nature, 2020, 580, 640-646.	13.7	338
10	Extensive heterogeneity in somatic mutation and selection in the human bladder. Science, 2020, 370, 75-82.	6.0	195
11	Two cryptic species of California mustard within Caulanthus lasiophyllus. American Journal of Botany, 2020, 107, 1815-1830.	0.8	2
12	Circulating tumor DNA dynamics using patient-customized assays are associated with outcome in neoadjuvantly treated breast cancer. Journal of Physical Education and Sports Management, 2019, 5, a003772.	0.5	24
13	Circulating-tumor DNA as an early detection and diagnostic tool. Current Opinion in Genetics and Development, 2017, 42, 14-21.	1.5	84
14	Non-random fragmentation patterns in circulating cell-free DNA reflect epigenetic regulation. BMC Genomics, 2015, 16, S1.	1.2	120
15	Exome Sequencing of Cell-Free DNA from Metastatic Cancer Patients Identifies Clinically Actionable Mutations Distinct from Primary Disease. PLoS ONE, 2015, 10, e0136407.	1.1	97
16	Circulating tumor cells in hepatocellular carcinoma: a pilot study of detection, enumeration, and next-generation sequencing in cases and controls. BMC Cancer, 2015, 15, 206.	1.1	103
17	Phylogenetic analyses of melanoma reveal complex patterns of metastatic dissemination. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 10995-11000.	3.3	146
18	Assessing Copy Number Alterations in Targeted, Amplicon-Based Next-Generation Sequencing Data. Journal of Molecular Diagnostics, 2015, 17, 53-63.	1.2	129

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#	Article	IF	CITATIONS
19	Transcriptome Analysis of a Petal Anthocyanin Polymorphism in the Arctic Mustard, Parrya nudicaulis. PLoS ONE, 2014, 9, e101338.	1.1	18
20	Arctic Mustard Flower Color Polymorphism Controlled by Petal-Specific Downregulation at the Threshold of the Anthocyanin Biosynthetic Pathway. PLoS ONE, 2011, 6, e18230.	1.1	80