

Noah Spies

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5630755/publications.pdf>

Version: 2024-02-01

14
papers

1,217
citations

1163117

8
h-index

1281871

11
g-index

20
all docs

20
docs citations

20
times ranked

2481
citing authors

#	ARTICLE	IF	CITATIONS
1	A robust benchmark for detection of germline large deletions and insertions. Nature Biotechnology, 2020, 38, 1347-1355.	17.5	233
2	A crowdsourced set of curated structural variants for the human genome. PLoS Computational Biology, 2020, 16, e1007933.	3.2	6
3	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
4	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
5	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
6	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
7	Haplotype-resolved and integrated genome analysis of the cancer cell line HepG2. Nucleic Acids Research, 2019, 47, 3846-3861.	14.5	45
8	Comprehensive, integrated, and phased whole-genome analysis of the primary ENCODE cell line K562. Genome Research, 2019, 29, 472-484.	5.5	78
9	Genome-wide reconstruction of complex structural variants using read clouds. Nature Methods, 2017, 14, 915-920.	19.0	96
10	Extensive sequencing of seven human genomes to characterize benchmark reference materials. Scientific Data, 2016, 3, 160025.	5.3	575
11	Cell-lineage heterogeneity and driver mutation recurrence in pre-invasive breast neoplasia. Genome Medicine, 2015, 7, 28.	8.2	17
12	Concepts in solid tumor evolution. Trends in Genetics, 2015, 31, 208-214.	6.7	51
13	svviz: a read viewer for validating structural variants. Bioinformatics, 2015, 31, 3994-3996.	4.1	46
14	Constraint and divergence of global gene expression in the mammalian embryo. ELife, 2015, 4, e05538.	6.0	3