## Rachael E Workman

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10777120/publications.pdf

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1040056 1199594 1,725 11 9 12 citations g-index h-index papers 14 14 14 3101 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Metagenomic next-generation sequencing of rectal swabs for the surveillance of antimicrobial-resistant organisms on the Illumina Miseq and Oxford MinION platforms. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 95-102.	2.9	16
2	High-quality chromosome-scale assembly of the walnut (Juglans regia L.) reference genome. GigaScience, 2020, 9, .	6.4	83
3	Characterization of human telomerase reverse transcriptase promoter methylation and transcription factor binding in differentiated thyroid cancer cell lines. Genes Chromosomes and Cancer, 2019, 58, 530-540.	2.8	21
4	Nanopore native RNA sequencing of a human poly(A) transcriptome. Nature Methods, 2019, 16, 1297-1305.	19.0	411
5	Single-molecule, full-length transcript sequencing provides insight into the extreme metabolism of the ruby-throated hummingbird Archilochus colubris. GigaScience, 2018, 7, 1-12.	6.4	67
6	Protein engineering strategies for improving the selective methylation of target CpG sites by a dCas9-directed cytosine methyltransferase in bacteria. PLoS ONE, 2018, 13, e0209408.	2.5	9
7	Detecting DNA cytosine methylation using nanopore sequencing. Nature Methods, 2017, 14, 407-410.	19.0	820
8	Multiplexed analysis of fixed tissue RNA using Ligation in situ Hybridization. Nucleic Acids Research, 2017, 45, e128-e128.	14.5	7
9	Targeted DNA methylation in human cells using engineered dCas9-methyltransferases. Scientific Reports, 2017, 7, 6732.	3.3	73
10	Common mycelial networks impact competition in an invasive grass. American Journal of Botany, 2016, 103, 1041-1049.	1.7	12
11	Nanopore sequencing detects structural variants in cancer. Cancer Biology and Therapy, 2016, 17, 246-253.	3.4	130