Doruk Beyter

List of Publications by Year in descending order

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1163117 1474206 1,981 9 8 9 citations h-index g-index papers 15 15 15 5125 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Ratatosk: hybrid error correction of long reads enables accurate variant calling and assembly. Genome Biology, 2021, 22, 28.	8.8	35
2	PopDel identifies medium-size deletions simultaneously in tens of thousands of genomes. Nature Communications, 2021, 12, 730.	12.8	9
3	Long-read sequencing of 3,622 Icelanders provides insight into the role of structural variants in human diseases and other traits. Nature Genetics, 2021, 53, 779-786.	21.4	156
4	Differences between germline genomes of monozygotic twins. Nature Genetics, 2021, 53, 27-34.	21.4	83
5	GraphTyper2 enables population-scale genotyping of structural variation using pangenome graphs. Nature Communications, 2019, 10, 5402.	12.8	96
6	ProteoStorm: An Ultrafast Metaproteomics Database Search Framework. Cell Systems, 2018, 7, 463-467.e6.	6.2	27
7	Strelka2: fast and accurate calling of germline and somatic variants. Nature Methods, 2018, 15, 591-594.	19.0	939
8	Extrachromosomal oncogene amplification drives tumour evolution and genetic heterogeneity. Nature, 2017, 543, 122-125.	27.8	530
9	Diversity, Productivity, and Stability of an Industrial Microbial Ecosystem. Applied and Environmental Microbiology, 2016, 82, 2494-2505.	3.1	46