Tim Stuart

List of Publications by Year in descending order

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TIM STUADT

#	Article	IF	CITATIONS
1	Characterizing cellular heterogeneity in chromatin state with scCUT&Tag-pro. Nature Biotechnology, 2022, 40, 1220-1230.	17.5	46
2	Single nucleus multi-omics identifies human cortical cell regulatory genome diversity. Cell Genomics, 2022, 2, 100107.	6.5	58
3	Integrated analysis of multimodal single-cell data. Cell, 2021, 184, 3573-3587.e29.	28.9	5,912
4	Single-cell chromatin state analysis with Signac. Nature Methods, 2021, 18, 1333-1341.	19.0	595
5	geneBasis: an iterative approach for unsupervised selection of targeted gene panels from scRNA-seq. Genome Biology, 2021, 22, 333.	8.8	15
6	A Genomeâ€Wide Association Study of Nonâ€Photochemical Quenching in response to local seasonal climates in <i>Arabidopsis thaliana</i> . Plant Direct, 2019, 3, e00138.	1.9	25
7	Integrative single-cell analysis. Nature Reviews Genetics, 2019, 20, 257-272.	16.3	932
8	Comprehensive Integration of Single-Cell Data. Cell, 2019, 177, 1888-1902.e21.	28.9	9,755
9	A single-cell transcriptional roadmap for cardiopharyngeal fate diversification. Nature Cell Biology, 2019, 21, 674-686.	10.3	78
10	Toward a Common Coordinate Framework for the Human Body. Cell, 2019, 179, 1455-1467.	28.9	81
11	Approaches for theÂAnalysis and Interpretation of Whole Genome Bisulfite Sequencing Data. Methods in Molecular Biology, 2018, 1767, 299-310.	0.9	6
12	DNA methylation profiles of diverse <i>Brachypodium distachyon</i> align with underlying genetic diversity. Genome Research, 2016, 26, 1520-1531.	5.5	34
13	Unique cell-type-specific patterns of DNA methylation in the root meristem. Nature Plants, 2016, 2, 16058.	9.3	159
14	Population scale mapping of transposable element diversity reveals links to gene regulation and epigenomic variation. ELife, 2016, 5, .	6.0	181