

viSNE enables visualization of high dimensional single-heterogeneity of leukemia

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Single-cell mass cytometry for analysis of immune system functional states. <i>Current Opinion in Immunology</i> , 2013, 25, 484-494.	2.4	196
3	High-Dimensional Analysis of Human CD8+ T Cell Phenotype, Function, and Antigen Specificity. <i>Current Topics in Microbiology and Immunology</i> , 2013, 377, 61-84.	0.7	11
4	A multidimensional overview. <i>Nature Reviews Cancer</i> , 2013, 13, 439-439.	12.8	3
5	LA2SNE: A novel stochastic neighbor embedding approach for microbiome data visualization. , 2014, , .		0
6	Single-cell sphingosine kinase activity measurements in primary leukemia. <i>Analytical and Bioanalytical Chemistry</i> , 2014, 406, 7027-7036.	1.9	16
7	Characterizing heterogeneity in leukemic cells using single-cell gene expression analysis. <i>Genome Biology</i> , 2014, 15, 525.	3.8	54
8	OpenCyto: An Open Source Infrastructure for Scalable, Robust, Reproducible, and Automated, End-to-End Flow Cytometry Data Analysis. <i>PLoS Computational Biology</i> , 2014, 10, e1003806.	1.5	185
9	Defining minimal residual disease in acute myeloid leukemia: which platforms are ready for "prime time"? <i>Hematology American Society of Hematology Education Program</i> , 2014, 2014, 222-233.	0.9	48
11	Transient partial permeabilization with saponin enables cellular barcoding prior to surface marker staining. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2014, 85, 1011-1019.	1.1	108
12	Transcriptional mechanisms of cell fate decisions revealed by single cell expression profiling. <i>BioEssays</i> , 2014, 36, 419-426.	1.2	24
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22	Single-Cell Trajectory Detection Uncovers Progression and Regulatory Coordination in Human B Cell Development. <i>Cell</i> , 2014, 157, 714-725.	13.5	838
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24	Single-cell technologies for monitoring immune systems. <i>Nature Immunology</i> , 2014, 15, 128-135.	7.0	337
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