## Chloé B Steen

List of Publications by Year in descending order

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<u> ΓΗΙ ΟÃ Ο Β ΣΤΕΕΝ</u>

#	Article	IF	CITATIONS
1	Determining cell type abundance and expression from bulk tissues with digital cytometry. Nature Biotechnology, 2019, 37, 773-782.	17.5	2,396
2	Profiling Cell Type Abundance and Expression in Bulk Tissues with CIBERSORTx. Methods in Molecular Biology, 2020, 2117, 135-157.	0.9	249
3	Noninvasive Early Identification of Therapeutic Benefit from Immune Checkpoint Inhibition. Cell, 2020, 183, 363-376.e13.	28.9	206
4	T cell characteristics associated with toxicity to immune checkpoint blockade in patients with melanoma. Nature Medicine, 2022, 28, 353-362.	30.7	132
5	Atlas of clinically distinct cell states and ecosystems across human solid tumors. Cell, 2021, 184, 5482-5496.e28.	28.9	116
6	The landscape of tumor cell states and ecosystems in diffuse large B cell lymphoma. Cancer Cell, 2021, 39, 1422-1437.e10.	16.8	102
7	Inferring gene expression from cell-free DNA fragmentation profiles. Nature Biotechnology, 2022, 40, 585-597.	17.5	63
8	ALIX and ESCRT-III Coordinately Control Cytokinetic Abscission during Germline Stem Cell Division In Vivo. PLoS Genetics, 2015, 11, e1004904.	3.5	54
9	Computational approaches for characterizing the tumor immune microenvironment. Immunology, 2019, 158, 70-84.	4.4	30
10	Towards Non-Invasive Classification of DLBCL Genetic Subtypes By Ctdna Profiling. Blood, 2019, 134, 551-551.	1.4	9
11	Mutational dynamics and immune evasion in diffuse large B-cell lymphoma explored in a relapse-enriched patient series. Blood Advances, 2020, 4, 1859-1866.	5.2	7
12	Heterogeneity of Regulatory T Cells in B-Cell Non-Hodgkin Lymphoma. Blood, 2020, 136, 27-28.	1.4	1
13	Mutational Dynamics and Evolutionary Divergence in DLBCL: A Call for Relapse Sampling. Blood, 2019, 134, 1497-1497.	1.4	0
14	Noninvasive Cell-of-Origin Classification of Diffuse Large B-Cell Lymphoma Using Inferred Gene Expression from Cell-Free DNA Sequencing. Blood, 2021, 138, 37-37.	1.4	0