

# Christof Angermueller

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7423542/publications.pdf>

Version: 2024-02-01

5  
papers

3,085  
citations

1478505

6  
h-index

1872680

6  
g-index

8  
all docs

8  
docs citations

8  
times ranked

5049  
citing authors

#	ARTICLE	IF	CITATIONS
1	Deep learning for computational biology. <i>Molecular Systems Biology</i> , 2016, 12, 878.	7.2	1,059
2	Single-cell genome-wide bisulfite sequencing for assessing epigenetic heterogeneity. <i>Nature Methods</i> , 2014, 11, 817-820.	19.0	954
3	Parallel single-cell sequencing links transcriptional and epigenetic heterogeneity. <i>Nature Methods</i> , 2016, 13, 229-232.	19.0	602
4	DeepCpG: accurate prediction of single-cell DNA methylation states using deep learning. <i>Genome Biology</i> , 2017, 18, 67.	8.8	361
5	Genome-Scale Oscillations in DNA Methylation during Exit from Pluripotency. <i>Cell Systems</i> , 2018, 7, 63-76.e12.	6.2	70