

Giuseppe-Antonio Saldi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7178665/publications.pdf>

Version: 2024-02-01

11
papers

376
citations

1684188

5
h-index

2053705

5
g-index

13
all docs

13
docs citations

13
times ranked

356
citing authors

#	ARTICLE	IF	CITATIONS
1	GABA-receptive microglia selectively sculpt developing inhibitory circuits. <i>Cell</i> , 2021, 184, 4048-4063.e32.	28.9	142
2	Gene regulatory network reconstruction using single-cell RNA sequencing of barcoded genotypes in diverse environments. <i>ELife</i> , 2020, 9, .	6.0	116
3	Genetic and epigenetic coordination of cortical interneuron development. <i>Nature</i> , 2021, 597, 693-697.	27.8	56
4	High-performance single-cell gene regulatory network inference at scale: the Inferelator 3.0. <i>Bioinformatics</i> , 2022, 38, 2519-2528.	4.1	32
5	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. <i>PLoS Computational Biology</i> , 2021, 17, e1008569.	3.2	19
6	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. , 2021, 17, e1008569.		0
7	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. , 2021, 17, e1008569.		0
8	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. , 2021, 17, e1008569.		0
9	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. , 2021, 17, e1008569.		0
10	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. , 2021, 17, e1008569.		0
11	Optimal tuning of weighted kNN- and diffusion-based methods for denoising single cell genomics data. , 2021, 17, e1008569.		0