

# Anja FÃ¼llgrabe

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6546269/publications.pdf>

Version: 2024-02-01

15  
papers

2,438  
citations

687363

13  
h-index

996975

15  
g-index

20  
all docs

20  
docs citations

20  
times ranked

5949  
citing authors

#	ARTICLE	IF	CITATIONS
1	Expression Atlas updateâ€”an integrated database of gene and protein expression in humans, animals and plants. <i>Nucleic Acids Research</i> , 2016, 44, D746-D752.	14.5	526
2	ArrayExpress update â€” from bulk to single-cell expression data. <i>Nucleic Acids Research</i> , 2019, 47, D711-D715.	14.5	497
3	Expression Atlas: gene and protein expression across multiple studies and organisms. <i>Nucleic Acids Research</i> , 2018, 46, D246-D251.	14.5	365
4	Expression Atlas update: from tissues to single cells. <i>Nucleic Acids Research</i> , 2020, 48, D77-D83.	14.5	363
5	A high-resolution mRNA expression time course of embryonic development in zebrafish. <i>ELife</i> , 2017, 6, .	6.0	257
6	Dynamics of Lgr6 + Progenitor Cells in the Hair Follicle, Sebaceous Gland, and Interfollicular Epidermis. <i>Stem Cell Reports</i> , 2015, 5, 843-855.	4.8	87
7	Expression Atlas update: gene and protein expression in multiple species. <i>Nucleic Acids Research</i> , 2022, 50, D129-D140.	14.5	78
8	From ArrayExpress to BioStudies. <i>Nucleic Acids Research</i> , 2021, 49, D1502-D1506.	14.5	53
9	A proteomics sample metadata representation for multiomics integration and big data analysis. <i>Nature Communications</i> , 2021, 12, 5854.	12.8	45
10	The RNASeq-er APIâ€”a gateway to systematically updated analysis of public RNA-seq data. <i>Bioinformatics</i> , 2017, 33, 2218-2220.	4.1	35
11	Guidelines for reporting single-cell RNA-seq experiments. <i>Nature Biotechnology</i> , 2020, 38, 1384-1386.	17.5	27
12	Coordinated hedgehog signaling induces new hair follicles in adult skin. <i>ELife</i> , 2020, 9, .	6.0	27
13	A user guide for the online exploration and visualization of PCAWG data. <i>Nature Communications</i> , 2020, 11, 3400.	12.8	23
14	Ten quick tips for biocuration. <i>PLoS Computational Biology</i> , 2019, 15, e1006906.	3.2	21
15	Common and distinct transcriptional signatures of mammalian embryonic lethality. <i>Nature Communications</i> , 2019, 10, 2792.	12.8	16