

# Florian Battke

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/488840/publications.pdf>

Version: 2024-02-01

28  
papers

773  
citations

840776

11  
h-index

610901

24  
g-index

28  
all docs

28  
docs citations

28  
times ranked

1731  
citing authors

#	ARTICLE	IF	CITATIONS
1	Biomarkers Associated with Immune-Related Adverse Events under Checkpoint Inhibitors in Metastatic Melanoma. <i>Cancers</i> , 2022, 14, 302.	3.7	15
2	Trio exome sequencing is highly relevant in prenatal diagnostics. <i>Prenatal Diagnosis</i> , 2022, 42, 845-851.	2.3	18
3	The question of WGS's clinical utility remains unanswered. <i>European Journal of Human Genetics</i> , 2021, 29, 722-723.	2.8	7
4	Controlling my genome with my smartphone: first clinical experiences of the PROMISE system. <i>Clinical Research in Cardiology</i> , 2021, , 1.	3.3	3
5	Copy number variation profiling in pharmacogenes using panel-based exome resequencing and correlation to human liver expression. <i>Human Genetics</i> , 2020, 139, 137-149.	3.8	9
6	Are Pathogenic Germline Variants in Metastatic Melanoma Associated with Resistance to Combined Immunotherapy?. <i>Cancers</i> , 2020, 12, 1101.	3.7	9
7	&lt;p&gt;Circulating Tumor DNA Correlates with Outcome in Metastatic Melanoma Treated by BRAF and MEK Inhibitors â€ Results of a Prospective Biomarker Study&lt;/p&gt;. <i>OncoTargets and Therapy</i> , 2020, Volume 13, 5017-5032.	2.0	9
8	Interference With Complex IV as a Model of Age-Related Decline in Synaptic Connectivity. <i>Frontiers in Molecular Neuroscience</i> , 2020, 13, 43.	2.9	11
9	Sequencing for an interdisciplinary molecular tumor board in patients with advanced breast cancer: experiences from a case series. <i>Oncotarget</i> , 2020, 11, 3279-3285.	1.8	2
10	Tumor mutation burden and circulating tumor DNA in combined CTLA-4 and PD-1 antibody therapy in metastatic melanoma â€ results of a prospective biomarker study. , 2019, 7, 180.		137
11	Targeted gene sequencing in 6994 individuals with neurodevelopmental disorder with epilepsy. <i>Genetics in Medicine</i> , 2019, 21, 2496-2503.	2.4	45
12	Molecular tumor analysis and liquid biopsy: a feasibility investigation analyzing circulating tumor DNA in patients with central nervous system lymphomas. <i>BMC Cancer</i> , 2019, 19, 192.	2.6	32
13	A New Panel-Based Next-Generation Sequencing Method for ADME Genes Reveals Novel Associations of Common and Rare Variants With Expression in a Human Liver Cohort. <i>Frontiers in Genetics</i> , 2019, 10, 7.	2.3	37
14	My Genome Belongs to Me: Controlling Third Party Computation on Genomic Data. <i>Proceedings on Privacy Enhancing Technologies</i> , 2019, 2019, 108-132.	2.8	6
15	Rare Variants in Neurodegeneration Associated Genes Revealed by Targeted Panel Sequencing in a German ALS Cohort. <i>Frontiers in Molecular Neuroscience</i> , 2016, 9, 92.	2.9	41
16	The Maternal Transcriptome of the Crustacean <i>Parhyale hawaiiensis</i> Is Inherited Asymmetrically to Invariant Cell Lineages of the Ectoderm and Mesoderm. <i>PLoS ONE</i> , 2013, 8, e56049.	2.5	23
17	REVEALâ€visual eQTL analytics. <i>Bioinformatics</i> , 2012, 28, i542-i548.	4.1	0
18	iHAT: interactive Hierarchical Aggregation Table for Genetic Association Data. <i>BMC Bioinformatics</i> , 2012, 13, S2.	2.6	16

#	ARTICLE	IF	CITATIONS
19	An eQTL biological data visualization challenge and approaches from the visualization community. BMC Bioinformatics, 2012, 13, S8.	2.6	9
20	iHAT: Interactive hierarchical aggregation table. , 2011, , .		2
21	TIALA &#x2014; Time series alignment analysis. , 2011, , .		4
22	Mayday SeaSight: Combined Analysis of Deep Sequencing and Microarray Data. PLoS ONE, 2011, 6, e16345.	2.5	17
23	The PII protein GlnK is a pleiotropic regulator for morphological differentiation and secondary metabolism in <i>Streptomyces coelicolor</i> . Applied Microbiology and Biotechnology, 2011, 92, 1219-1236.	3.6	34
24	<i>GaggleBridge</i>: collaborative data analysis. Bioinformatics, 2011, 27, 2612-2613.	4.1	4
25	Mayday - integrative analytics for expression data. BMC Bioinformatics, 2010, 11, 121.	2.6	102
26	The dynamic architecture of the metabolic switch in <i>Streptomyces coelicolor</i> . BMC Genomics, 2010, 11, 10.	2.8	171
27	Integrative systems biology visualization with MAYDAY. Journal of Integrative Bioinformatics, 2010, 7, .	1.5	9
28	Post-Hybridization Quality Measures for Oligos in Genome-Wide Microarray Experiments. Lecture Notes in Computer Science, 2008, , 64-75.	1.3	1