

# Christian Munk

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/118925/publications.pdf>

Version: 2024-02-01

15  
papers

2,130  
citations

759233

12  
h-index

888059

17  
g-index

17  
all docs

17  
docs citations

17  
times ranked

3147  
citing authors

#	ARTICLE	IF	CITATIONS
1	Long-term Absolute Risk of Cervical Intraepithelial Neoplasia Grade 3 or Worse Following Human Papillomavirus Infection: Role of Persistence. <i>Journal of the National Cancer Institute</i> , 2010, 102, 1478-1488.	6.3	485
2	GPCRdb: an information system for G protein-coupled receptors. <i>Nucleic Acids Research</i> , 2016, 44, D356-D364.	14.5	472
3	GPCRdb in 2018: adding GPCR structure models and ligands. <i>Nucleic Acids Research</i> , 2018, 46, D440-D446.	14.5	421
4	GPCRdb in 2021: integrating GPCR sequence, structure and function. <i>Nucleic Acids Research</i> , 2021, 49, D335-D343.	14.5	254
5	An online resource for GPCR structure determination and analysis. <i>Nature Methods</i> , 2019, 16, 151-162.	19.0	108
6	GPCR activation mechanisms across classes and macro/microscales. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 879-888.	8.2	98
7	Combinatorial expression of GPCR isoforms affects signalling and drug responses. <i>Nature</i> , 2020, 587, 650-656.	27.8	87
8	Integrating structural and mutagenesis data to elucidate GPCR ligand binding. <i>Current Opinion in Pharmacology</i> , 2016, 30, 51-58.	3.5	52
9	The G protein database, GproteinDb. <i>Nucleic Acids Research</i> , 2022, 50, D518-D525.	14.5	49
10	An online GPCR structure analysis platform. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 875-878.	8.2	16
11	TMEM45A, SERPINB5 and p16INK4A transcript levels are predictive for development of high-grade cervical lesions. <i>American Journal of Cancer Research</i> , 2016, 6, 1524-36.	1.4	13
12	Structural insight to mutation effects uncover a common allosteric site in class C GPCRs. <i>Bioinformatics</i> , 2017, 33, 1116-1120.	4.1	9
13	Physical state and viral load as predictive biomarkers for persistence and progression of HPV16-positive cervical lesions: results from a population based long-term prospective cohort study. <i>American Journal of Cancer Research</i> , 2012, 2, 192-203.	1.4	8
14	HPV16 viral load and physical state measurement as a potential immediate triage strategy for HR-HPV-infected women: a study in 644 women with single HPV16 infections. <i>American Journal of Cancer Research</i> , 2018, 8, 715-722.	1.4	8
15	Methylation of CpG 5962 in L1 of the human papillomavirus 16 genome as a potential predictive marker for viral persistence: A prospective large cohort study using cervical swab samples. <i>Cancer Medicine</i> , 2020, 9, 1058-1068.	2.8	7