

# Christopher Hunter

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3034527/publications.pdf>

Version: 2024-02-01

19  
papers

6,440  
citations

567281

15  
h-index

794594

19  
g-index

20  
all docs

20  
docs citations

20  
times ranked

11810  
citing authors

#	ARTICLE	IF	CITATIONS
1	Patterns of somatic mutation in human cancer genomes. <i>Nature</i> , 2007, 446, 153-158.	27.8	2,802
2	Intragenic ERBB2 kinase mutations in tumours. <i>Nature</i> , 2004, 431, 525-526.	27.8	757
3	Minimum information about a marker gene sequence (MIMARKS) and minimum information about any (x) sequence (MlxS) specifications. <i>Nature Biotechnology</i> , 2011, 29, 415-420.	17.5	608
4	Somatic Mutations of the Protein Kinase Gene Family in Human Lung Cancer. <i>Cancer Research</i> , 2005, 65, 7591-7595.	0.9	429
5	A Hypermutation Phenotype and Somatic <i>MSH6</i> Mutations in Recurrent Human Malignant Gliomas after Alkylator Chemotherapy. <i>Cancer Research</i> , 2006, 66, 3987-3991.	0.9	383
6	Mutation analysis of 24 known cancer genes in the NCI-60 cell line set. <i>Molecular Cancer Therapeutics</i> , 2006, 5, 2606-2612.	4.1	374
7	A screen of the complete protein kinase gene family identifies diverse patterns of somatic mutations in human breast cancer. <i>Nature Genetics</i> , 2005, 37, 590-592.	21.4	318
8	Recurrent KRAS codon 146 mutations in human colorectal cancer. <i>Cancer Biology and Therapy</i> , 2006, 5, 928-932.	3.4	200
9	Reporting guidelines for human microbiome research: the STORMS checklist. <i>Nature Medicine</i> , 2021, 27, 1885-1892.	30.7	170
10	EBI metagenomics—a new resource for the analysis and archiving of metagenomic data. <i>Nucleic Acids Research</i> , 2014, 42, D600-D606.	14.5	127
11	Sequence analysis of the protein kinase gene family in human testicular germ cell tumors of adolescents and adults. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 42-46.	2.8	96
12	COVID-19 pandemic reveals the peril of ignoring metadata standards. <i>Scientific Data</i> , 2020, 7, 188.	5.3	56
13	<i>Eupolybothrus cavernicolus</i> Komeriki & Stoev sp. n. (Chilopoda: Lithobiomorpha: Lithobiidae): the first eukaryotic species description combining transcriptomic, DNA barcoding and micro-CT imaging data. <i>Biodiversity Data Journal</i> , 2013, 1, e1013.	0.8	46
14	Major submissions tool developments at the European nucleotide archive. <i>Nucleic Acids Research</i> , 2012, 40, D43-D47.	14.5	32
15	GigaDB: promoting data dissemination and reproducibility. <i>Database: the Journal of Biological Databases and Curation</i> , 2014, 2014, bau018-bau018.	3.0	20
16	Experiences in integrated data and research object publishing using GigaDB. <i>International Journal on Digital Libraries</i> , 2017, 18, 99-111.	1.5	14
17	MlxS-SA: a MlxS extension defining the minimum information standard for sequence data from symbiont-associated micro-organisms. <i>ISME Communications</i> , 2022, 2, .	4.2	3
18	Looking back: forward looking. <i>GigaScience</i> , 2017, 6, 1-3.	6.4	2

#	ARTICLE	IF	CITATIONS
19	A Decade of GigaScience: GigaDB and the Open Data Movement. GigaScience, 2022, 11, .	6.4	2