

# Gary Saunders

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5969167/publications.pdf>

Version: 2024-02-01

9  
papers

4,788  
citations

1306789

7  
h-index

1473754

9  
g-index

9  
all docs

9  
docs citations

9  
times ranked

15102  
citing authors

#	ARTICLE	IF	CITATIONS
1	GENCODE: The reference human genome annotation for The ENCODE Project. <i>Genome Research</i> , 2012, 22, 1760-1774.	2.4	4,217
2	The European Genome-phenome Archive of human data consented for biomedical research. <i>Nature Genetics</i> , 2015, 47, 692-695.	9.4	338
3	Federated discovery and sharing of genomic data using Beacons. <i>Nature Biotechnology</i> , 2019, 37, 220-224.	9.4	75
4	The European Variation Archive: a FAIR resource of genomic variation for all species. <i>Nucleic Acids Research</i> , 2022, 50, D1216-D1220.	6.5	50
5	Genetics of Mating and Sex Determination in the Parasitic Nematode <i>Haemonchus contortus</i> . <i>Genetics</i> , 2008, 180, 1877-1887.	1.2	40
6	The Data Use Ontology to streamline responsible access to human biomedical datasets. <i>Cell Genomics</i> , 2021, 1, 100028.	3.0	31
7	Annotation of Two Large Contiguous Regions from the <i>Haemonchus contortus</i> Genome Using RNA-seq and Comparative Analysis with <i>Caenorhabditis elegans</i> . <i>PLoS ONE</i> , 2011, 6, e23216.	1.1	22
8	Beacon v2 and Beacon networks: A <i>lingua franca</i> for federated data discovery in biomedical genomics, and beyond. <i>Human Mutation</i> , 2022, , .	1.1	10
9	The ELIXIR Human Copy Number Variations Community: building bioinformatics infrastructure for research. <i>F1000Research</i> , 2020, 9, 1229.	0.8	5