

# Gary Saunders

## List of Publications by Citations

**Source:** <https://exaly.com/author-pdf/5969167/gary-saunders-publications-by-citations.pdf>

**Version:** 2024-04-24

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

9

papers

3,487

citations

7

h-index

9

g-index

9

ext. papers

4,357

ext. citations

15.4

avg, IF

3.36

L-index

#	Paper	IF	Citations
9	GENCODE: the reference human genome annotation for The ENCODE Project. <i>Genome Research</i> , <b>2012</b> , 22, 1760-74	9.7	3142
8	The European Genome-phenome Archive of human data consented for biomedical research. <i>Nature Genetics</i> , <b>2015</b> , 47, 692-5	36.3	224
7	Federated discovery and sharing of genomic data using Beacons. <i>Nature Biotechnology</i> , <b>2019</b> , 37, 220-224	44.5	42
6	Genetics of mating and sex determination in the parasitic nematode <i>Haemonchus contortus</i> . <i>Genetics</i> , <b>2008</b> , 180, 1877-87	4	33
5	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> , <b>2011</b> , 6, e23216	3.7	21
4	The Data Use Ontology to streamline responsible access to human biomedical datasets. <i>Cell Genomics</i> , <b>2021</b> , 1, None		10
3	The European Variation Archive: a FAIR resource of genomic variation for all species. <i>Nucleic Acids Research</i> , <b>2021</b> ,	20.1	10
2	The ELIXIR Human Copy Number Variations Community: building bioinformatics infrastructure for research. <i>F1000Research</i> , <b>2020</b> , 9,	3.6	4
1	Beacon v2 and Beacon Networks: a "lingua franca" for federated data discovery in biomedical genomics, and beyond.. <i>Human Mutation</i> , <b>2022</b> ,	4.7	1