

# Laurent Gil

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6929319/publications.pdf>

Version: 2024-02-01

18  
papers

19,142  
citations

430754

18  
h-index

839398

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g-index

18  
all docs

18  
docs citations

18  
times ranked

46692  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Ensembl Variant Effect Predictor. <i>Genome Biology</i> , 2016, 17, 122.	3.8	5,181
2	Ensembl 2018. <i>Nucleic Acids Research</i> , 2018, 46, D754-D761.	6.5	2,710
3	The new NHGRI-EBI Catalog of published genome-wide association studies (GWAS Catalog). <i>Nucleic Acids Research</i> , 2017, 45, D896-D901.	6.5	1,932
4	Ensembl 2016. <i>Nucleic Acids Research</i> , 2016, 44, D710-D716.	6.5	1,372
5	Ensembl 2014. <i>Nucleic Acids Research</i> , 2014, 42, D749-D755.	6.5	1,211
6	Ensembl 2015. <i>Nucleic Acids Research</i> , 2015, 43, D662-D669.	6.5	1,145
7	Ensembl 2020. <i>Nucleic Acids Research</i> , 2020, 48, D682-D688.	6.5	1,076
8	Ensembl 2019. <i>Nucleic Acids Research</i> , 2019, 47, D745-D751.	6.5	879
9	Ensembl 2013. <i>Nucleic Acids Research</i> , 2012, 41, D48-D55.	6.5	856
10	Ensembl 2012. <i>Nucleic Acids Research</i> , 2012, 40, D84-D90.	6.5	840
11	Ensembl 2017. <i>Nucleic Acids Research</i> , 2017, 45, D635-D642.	6.5	535
12	Ensembl Genomes 2020 "enabling non-vertebrate genomic research. <i>Nucleic Acids Research</i> , 2020, 48, D689-D695.	6.5	416
13	Ensembl variation resources. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	1.4	377
14	The Polygenic Score Catalog as an open database for reproducibility and systematic evaluation. <i>Nature Genetics</i> , 2021, 53, 420-425.	9.4	293
15	A joint NCBI and EMBL-EBI transcript set for clinical genomics and research. <i>Nature</i> , 2022, 604, 310-315.	13.7	162
16	Locus Reference Genomic: reference sequences for the reporting of clinically relevant sequence variants. <i>Nucleic Acids Research</i> , 2014, 42, D873-D878.	6.5	73
17	MeRy-B: a web knowledgebase for the storage, visualization, analysis and annotation of plant NMR metabolomic profiles. <i>BMC Plant Biology</i> , 2011, 11, 104.	1.6	54
18	An integrative genomics approach for deciphering the complex interactions between ascorbate metabolism and fruit growth and composition in tomato. <i>Comptes Rendus - Biologies</i> , 2009, 332, 1007-1021.	0.1	30