

Maynard V Olson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8869743/publications.pdf>

Version: 2024-02-01

23
papers

22,591
citations

471061

17
h-index

610482

24
g-index

25
all docs

25
docs citations

25
times ranked

25478
citing authors

#	ARTICLE	IF	CITATIONS
1	Initial sequencing and analysis of the human genome. <i>Nature</i> , 2001, 409, 860-921.	13.7	21,074
2	When Less Is More: Gene Loss as an Engine of Evolutionary Change. <i>American Journal of Human Genetics</i> , 1999, 64, 18-23.	2.6	381
3	Sequencing the chimpanzee genome: insights into human evolution and disease. <i>Nature Reviews Genetics</i> , 2003, 4, 20-28.	7.7	219
4	Transcription and processing of cloned yeast tyrosine tRNA genes microinjected into frog oocytes. <i>Nature</i> , 1979, 278, 137-143.	13.7	154
5	Identification of a Genomic Island Present in the Majority of Pathogenic Isolates of <i>Pseudomonas aeruginosa</i> . <i>Journal of Bacteriology</i> , 2001, 183, 843-853.	1.0	114
6	Genome mosaicism is conserved but not unique in <i>Pseudomonas aeruginosa</i> isolates from the airways of young children with cystic fibrosis. <i>Environmental Microbiology</i> , 2003, 5, 1341-1349.	1.8	102
7	Redefining Genomic Privacy: Trust and Empowerment. <i>PLoS Biology</i> , 2014, 12, e1001983.	2.6	87
8	A genomic sequence analysis of the mouse and human microtubule-associated protein tau. <i>Mammalian Genome</i> , 2001, 12, 700-712.	1.0	59
9	Molecular characterisation of the tyrosine tRNA genes of yeast. <i>Nature</i> , 1977, 267, 639-641.	13.7	57
10	Ancient haplotypes of the HLA Class II region. <i>Genome Research</i> , 2005, 15, 1250-1257.	2.4	54
11	Closing gaps in the human genome with fosmid resources generated from multiple individuals. <i>Nature Genetics</i> , 2008, 40, 96-101.	9.4	50
12	Human Genetic Individuality. <i>Annual Review of Genomics and Human Genetics</i> , 2012, 13, 1-27.	2.5	43
13	Enrichment of super-sized resequencing targets from the human genome. <i>Nature Methods</i> , 2007, 4, 891-892.	9.0	40
14	The Human Genome Project: A Player's Perspective. <i>Journal of Molecular Biology</i> , 2002, 319, 931-942.	2.0	28
15	Targeted, haplotype-resolved resequencing of long segments of the human genome. <i>Genomics</i> , 2005, 86, 759-766.	1.3	23
16	Large-insert genome analysis technology detects structural variation in <i>Pseudomonas aeruginosa</i> clinical strains from cystic fibrosis patients. <i>Genomics</i> , 2008, 91, 530-537.	1.3	22
17	Dr Watson's base pairs. <i>Nature</i> , 2008, 452, 819-820.	13.7	20
18	GENOMICS: The Chimpanzee Genome—A Bittersweet Celebration. <i>Science</i> , 2004, 305, 191-192.	6.0	18

#	ARTICLE	IF	CITATIONS
19	Clone by clone by clone. Nature, 2001, 409, 816-818.	13.7	15
20	A Behind-the-Scenes Story of Precision Medicine *. Genomics, Proteomics and Bioinformatics, 2017, 15, 3-10.	3.0	11
21	Precision medicine at the crossroads. Human Genomics, 2017, 11, 23.	1.4	8
22	What Does a "Normal" Human Genome Look Like?. Science, 2011, 331, 872-872.	6.0	7
23	Detecting Disease-Causing Mutations in the Human Genome by Haplotype Matching. American Journal of Human Genetics, 2006, 79, 958-964.	2.6	4