

# Scott Schwartz

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10421299/publications.pdf>

Version: 2024-02-01

13  
papers

14,520  
citations

687363

13  
h-index

1125743

13  
g-index

13  
all docs

13  
docs citations

13  
times ranked

19465  
citing authors

#	ARTICLE	IF	CITATIONS
1	Initial sequencing and comparative analysis of the mouse genome. <i>Nature</i> , 2002, 420, 520-562.	27.8	6,319
2	A Greedy Algorithm for Aligning DNA Sequences. <i>Journal of Computational Biology</i> , 2000, 7, 203-214.	1.6	4,325
3	Genome sequence of the Brown Norway rat yields insights into mammalian evolution. <i>Nature</i> , 2004, 428, 493-521.	27.8	1,943
4	Human-Mouse Alignments with BLASTZ. <i>Genome Research</i> , 2003, 13, 103-107.	5.5	1,071
5	Covariation in Frequencies of Substitution, Deletion, Transposition, and Recombination During Eutherian Evolution. <i>Genome Research</i> , 2003, 13, 13-26.	5.5	263
6	MultiPipMaker and supporting tools: alignments and analysis of multiple genomic DNA sequences. <i>Nucleic Acids Research</i> , 2003, 31, 3518-3524.	14.5	196
7	Distinguishing Regulatory DNA From Neutral Sites. <i>Genome Research</i> , 2003, 13, 64-72.	5.5	118
8	Patterns of Insertions and Their Covariation With Substitutions in the Rat, Mouse, and Human Genomes. <i>Genome Research</i> , 2004, 14, 517-527.	5.5	73
9	Generation and Comparative Analysis of a 3.3 Mb of Mouse Genomic Sequence Orthologous to the Region of Human Chromosome 7q11.23 Implicated in Williams Syndrome. <i>Genome Research</i> , 2002, 12, 3-15.	5.5	72
10	GALA, a Database for Genomic Sequence Alignments and Annotations. <i>Genome Research</i> , 2003, 13, 732-741.	5.5	45
11	Sequence and comparative analysis of the rabbit $\beta$ -like globin gene cluster reveals a rapid mode of evolution in a G + C-rich region of mammalian genomes. <i>Journal of Molecular Biology</i> , 1991, 222, 233-249.	4.2	40
12	PipTools: A Computational Toolkit to Annotate and Analyze Pairwise Comparisons of Genomic Sequences. <i>Genomics</i> , 2002, 80, 681-690.	2.9	32
13	EnteriX 2003: visualization tools for genome alignments of Enterobacteriaceae. <i>Nucleic Acids Research</i> , 2003, 31, 3527-3532.	14.5	23