

Yu-Hui Rogers

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

22,132
citations

858243

12
h-index

1255698

13
g-index

16
all docs

16
docs citations

16
times ranked

26975
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic and functional adaptation in surface ocean planktonic prokaryotes. <i>Nature</i> , 2010, 468, 60-66.	13.7	280
2	DNA Sequencing for the Detection of Human Genome Variation. , 2010, , 27-37.		0
3	DNA Sequencing for the Detection of Human Genome Variation and Polymorphism. , 2009, , 88-100.		0
4	Emerging DNA sequencing technologies for human genomic medicine. <i>Drug Discovery Today</i> , 2008, 13, 569-577.	3.2	57
5	The Diploid Genome Sequence of an Individual Human. <i>PLoS Biology</i> , 2007, 5, e254.	2.6	1,491
6	Evolutionary and Biomedical Insights from the Rhesus Macaque Genome. <i>Science</i> , 2007, 316, 222-234.	6.0	1,283
7	The Sorcerer II Global Ocean Sampling Expedition: Northwest Atlantic through Eastern Tropical Pacific. <i>PLoS Biology</i> , 2007, 5, e77.	2.6	1,757
8	A Sanger/pyrosequencing hybrid approach for the generation of high-quality draft assemblies of marine microbial genomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 11240-11245.	3.3	243
9	Massively parallel sequencing. <i>Nature</i> , 2005, 437, 326-327.	13.7	136
10	Environmental Genome Shotgun Sequencing of the Sargasso Sea. <i>Science</i> , 2004, 304, 66-74.	6.0	3,776
11	The Sequence of the Human Genome. <i>Science</i> , 2001, 291, 1304-1351.	6.0	12,623
12	PBXL Fluorescent Dyes for Ultrasensitive Direct Detection. <i>Journal of Fluorescence</i> , 1999, 9, 197-208.	1.3	4
13	Immobilization of Oligonucleotides onto a Glass Support via Disulfide Bonds: A Method for Preparation of DNA Microarrays. <i>Analytical Biochemistry</i> , 1999, 266, 23-30.	1.1	243
14	Nested genetic bit analysis (N-GBA) for mutation detection in the p53 tumor suppressor gene. <i>Nucleic Acids Research</i> , 1997, 25, 5065-5071.	6.5	37
15	Genetic Bit Analysis: a solid phase method for typing single nucleotide polymorphisms. <i>Nucleic Acids Research</i> , 1994, 22, 4167-4175.	6.5	202