

John I Glass

List of Publications by Year in descending order

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63
papers

11,604
citations

109321

35
h-index

114465

63
g-index

63
all docs

63
docs citations

63
times ranked

10531
citing authors

#	ARTICLE	IF	CITATIONS
1	Creation of a Bacterial Cell Controlled by a Chemically Synthesized Genome. <i>Science</i> , 2010, 329, 52-56.	12.6	2,177
2	A Whole-Cell Computational Model Predicts Phenotype from Genotype. <i>Cell</i> , 2012, 150, 389-401.	28.9	1,177
3	Complete Chemical Synthesis, Assembly, and Cloning of a <i>Mycoplasma genitalium</i> Genome. <i>Science</i> , 2008, 319, 1215-1220.	12.6	1,122
4	Design and synthesis of a minimal bacterial genome. <i>Science</i> , 2016, 351, aad6253.	12.6	1,077
5	Essential genes of a minimal bacterium. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 425-430.	7.1	806
6	Genome of the Bacterium <i>Streptococcus pneumoniae</i> Strain R6. <i>Journal of Bacteriology</i> , 2001, 183, 5709-5717.	2.2	717
7	Genome Sequence of Avery's Virulent Serotype 2 Strain D39 of <i>Streptococcus pneumoniae</i> and Comparison with That of Unencapsulated Laboratory Strain R6. <i>Journal of Bacteriology</i> , 2007, 189, 38-51.	2.2	429
8	Genome Transplantation in Bacteria: Changing One Species to Another. <i>Science</i> , 2007, 317, 632-638.	12.6	422
9	The complete sequence of the mucosal pathogen <i>Ureaplasma urealyticum</i> . <i>Nature</i> , 2000, 407, 757-762.	27.8	383
10	Creating Bacterial Strains from Genomes That Have Been Cloned and Engineered in Yeast. <i>Science</i> , 2009, 325, 1693-1696.	12.6	289
11	The Sorcerer II Global Ocean Sampling Expedition: Metagenomic Characterization of Viruses within Aquatic Microbial Samples. <i>PLoS ONE</i> , 2008, 3, e1456.	2.5	276
12	Essence of life: essential genes of minimal genomes. <i>Trends in Cell Biology</i> , 2011, 21, 562-568.	7.9	167
13	Synthetic Generation of Influenza Vaccine Viruses for Rapid Response to Pandemics. <i>Science Translational Medicine</i> , 2013, 5, 185ra68.	12.4	164
14	Cloning whole bacterial genomes in yeast. <i>Nucleic Acids Research</i> , 2010, 38, 2558-2569.	14.5	156
15	Sequencing Multimegabase-Template DNA with BigDye Terminator Chemistry. <i>Genome Research</i> , 1998, 8, 557-561.	5.5	153
16	A Metagenomic Framework for the Study of Airborne Microbial Communities. <i>PLoS ONE</i> , 2013, 8, e81862.	2.5	127
17	Molecular Methods for the Detection of <i>Mycoplasma</i> and <i>Ureaplasma</i> Infections in Humans. <i>Journal of Molecular Diagnostics</i> , 2012, 14, 437-450.	2.8	124
18	A Genome-Scale Metabolic Reconstruction of <i>Mycoplasma genitalium</i> , iPS189. <i>PLoS Computational Biology</i> , 2009, 5, e1000285.	3.2	119

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19	Bacterial antisense RNAs are mainly the product of transcriptional noise. <i>Science Advances</i> , 2016, 2, e1501363.	10.3	118
20	Small repeating units within the <i>Ureaplasma urealyticum</i> MB antigen gene encode serovar specificity and are associated with antigen size variation. <i>Infection and Immunity</i> , 1995, 63, 891-898.	2.2	109
21	<i>Ureaplasma urealyticum</i> biovar specificity and diversity are encoded in multiple-banded antigen gene. <i>Journal of Clinical Microbiology</i> , 1994, 32, 1464-1469.	3.9	103
22	Detection and Characterization of Human <i>Ureaplasma</i> Species and Serovars by Real-Time PCR. <i>Journal of Clinical Microbiology</i> , 2010, 48, 2715-2723.	3.9	89
23	A Structurally Distinct Human Mycoplasma Protein that Generically Blocks Antigen-Antibody Union. <i>Science</i> , 2014, 343, 656-661.	12.6	85
24	Extensive Horizontal Gene Transfer in <i>Ureaplasmas</i> from Humans Questions the Utility of Serotyping for Diagnostic Purposes. <i>Journal of Clinical Microbiology</i> , 2011, 49, 2818-2826.	3.9	68
25	Assembly of Large, High G+C Bacterial DNA Fragments in Yeast. <i>ACS Synthetic Biology</i> , 2012, 1, 267-273.	3.8	65
26	Direct transfer of whole genomes from bacteria to yeast. <i>Nature Methods</i> , 2013, 10, 410-412.	19.0	64
27	Discovery of a novel bicycloproline P2 bearing peptidyl β -ketoamide LY514962 as HCV protease inhibitor. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2004, 14, 251-256.	2.2	61
28	Synergistic Antiviral Activity of Human Interferon Combinations in the Hepatitis C Virus Replicon System. <i>Journal of Interferon and Cytokine Research</i> , 2003, 23, 247-257.	1.2	60
29	Sequence analysis of a complete 1.66 Mb <i>Prochlorococcus marinus</i> MED4 genome cloned in yeast. <i>Nucleic Acids Research</i> , 2012, 40, 10375-10383.	14.5	56
30	A systems biology <i>tour de force</i> for a near-minimal bacterium. <i>Molecular Systems Biology</i> , 2009, 5, 330.	7.2	53
31	Sequence analysis of the chromosomal region around and within the V-1-encoding gene of <i>Mycoplasma pulmonis</i> : evidence for DNA inversion as a mechanism for V-1 variation. <i>Infection and Immunity</i> , 1996, 64, 472-479.	2.2	51
32	Mutations in ribosomal proteins and ribosomal RNA confer macrolide resistance in human <i>Ureaplasma</i> spp.. <i>International Journal of Antimicrobial Agents</i> , 2011, 37, 377-379.	2.5	41
33	Chromosomal Mutations Responsible for Fluoroquinolone Resistance in <i>Ureaplasma</i> Species in the United States. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 2780-2783.	3.2	39
34	Transfer RNA Misidentification Scrambles Sense Codon Recoding. <i>ChemBioChem</i> , 2013, 14, 1967-1972.	2.6	39
35	P4 and P1 ² optimization of bicycloproline P2 bearing tetrapeptidyl β -ketoamides as HCV protease inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2004, 14, 5007-5011.	2.2	37
36	Transferring whole genomes from bacteria to yeast spheroplasts using entire bacterial cells to reduce DNA shearing. <i>Nature Protocols</i> , 2014, 9, 743-750.	12.0	37

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37	Targeted Chromosomal Knockouts in <i>Mycoplasma pneumoniae</i> . Applied and Environmental Microbiology, 2010, 76, 5297-5299.	3.1	32
38	Fluoroquinolone Resistance in <i>Ureaplasma parvum</i> in the United States. Journal of Clinical Microbiology, 2006, 44, 1590-1591.	3.9	31
39	Cytokine-Activated Natural Killer Cells Exert Direct Killing of Hepatoma Cells Harboring Hepatitis C Virus Replicons. Journal of Interferon and Cytokine Research, 2006, 26, 854-865.	1.2	30
40	RecA mediates MgpB and MgpC phase and antigenic variation in <i>Mycoplasma genitalium</i> , but plays a minor role in DNA repair. Molecular Microbiology, 2012, 85, 669-683.	2.5	30
41	Tuning Gene Activity by Inducible and Targeted Regulation of Gene Expression in Minimal Bacterial Cells. ACS Synthetic Biology, 2018, 7, 1538-1552.	3.8	30
42	P1 and P3 optimization of novel bicycloproline P2 bearing tetrapeptidyl $\hat{\pm}$ -ketoamide based HCV protease inhibitors. Bioorganic and Medicinal Chemistry Letters, 2004, 14, 257-261.	2.2	29
43	Complete Genome Sequences of <i>Mycoplasma leachii</i> Strain PG50 ^T and the Pathogenic <i>Mycoplasma mycoides</i> subsp. <i>mycoides</i> Small Colony Biotype Strain Gladysdale. Journal of Bacteriology, 2012, 194, 4448-4449.	2.2	29
44	Bacterial genome reduction using the progressive clustering of deletions via yeast sexual cycling. Genome Research, 2015, 25, 435-444.	5.5	27
45	New Selectable Marker for Manipulating the Simple Genomes of <i>Mycoplasma</i> Species. Antimicrobial Agents and Chemotherapy, 2009, 53, 4429-4432.	3.2	26
46	Simultaneous non-contiguous deletions using large synthetic DNA and site-specific recombinases. Nucleic Acids Research, 2014, 42, e111-e111.	14.5	24
47	<i>Streptococcus pneumoniae</i> as a genomics platform for broad-spectrum antibiotic discovery. Current Opinion in Microbiology, 2002, 5, 338-342.	5.1	23
48	The costimulatory immunogen LPS induces the B-Cell clones that infiltrate transplanted human kidneys. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 6036-6041.	7.1	23
49	Pyruvate Oxidase Is a Determinant of Avery's Rough Morphology. Journal of Bacteriology, 2004, 186, 8164-8171.	2.2	20
50	Random insertion and gene disruption via transposon mutagenesis of <i>Ureaplasma parvum</i> using a mini-transposon plasmid. International Journal of Medical Microbiology, 2014, 304, 1218-1225.	3.6	20
51	Novel P4 truncated tripeptidyl $\hat{\pm}$ -ketoamides as HCV protease inhibitors. Bioorganic and Medicinal Chemistry Letters, 2004, 14, 263-266.	2.2	19
52	Efficient size-independent chromosome delivery from yeast to cultured cell lines. Nucleic Acids Research, 2017, 45, gkw1252.	14.5	18
53	P1 and P1; Optimization of [3,4]-Bicycloproline P2 Incorporated Tetrapeptidyl $\hat{\pm}$ -Ketoamide Based HCV Protease Inhibitors. Letters in Drug Design and Discovery, 2005, 2, 118-123.	0.7	17
54	Enhancement of Targeted Homologous Recombination in <i>Mycoplasma mycoides</i> subsp. <i>capri</i> by Inclusion of Heterologous <i>recA</i> . Applied and Environmental Microbiology, 2010, 76, 6951-6954.	3.1	17

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55	Different base per unit length ratios exist in single-stranded RNA and single-stranded DNA. <i>Nucleic Acids Research</i> , 1980, 8, 5739-5751.	14.5	16
56	Phylogenetic Analysis of the 16S-23S rRNA Intergenic Spacer Regions of the Genus <i>Ureaplasma</i> .. <i>Journal of Veterinary Medical Science</i> , 1996, 58, 191-195.	0.9	15
57	Genome Sequences of <i>Mycoplasma alligatoris</i> A21JP2 ^T and <i>Mycoplasma crocodyli</i> MP145 ^T. <i>Journal of Bacteriology</i> , 2011, 193, 2892-2893.	2.2	13
58	Genotypic Characterization of <i>Ureaplasma</i> Serovars from Clinical Isolates by Pulsed-Field Gel Electrophoresis. <i>Journal of Clinical Microbiology</i> , 2011, 49, 3325-3328.	3.9	13
59	Synthetic Genomics and the Construction of a Synthetic Bacterial Cell. <i>Perspectives in Biology and Medicine</i> , 2012, 55, 473-489.	0.5	13
60	Rescue of mutant fitness defects using in vitro reconstituted designer transposons in <i>Mycoplasma mycoides</i> . <i>Frontiers in Microbiology</i> , 2014, 5, 369.	3.5	12
61	Size Variation of a Major Serotype-Specific Antigen of <i>Ureaplasma urealyticum</i> . <i>Annals of the New York Academy of Sciences</i> , 1994, 730, 299-301.	3.8	10
62	Rescue of Infectious Sindbis Virus by Yeast Spheroplast-Mammalian Cell Fusion. <i>Viruses</i> , 2021, 13, 603.	3.3	4
63	<i>Ureaplasma urealyticum</i> : an opportunity for combinatorial genomics. <i>Trends in Microbiology</i> , 2001, 9, 163.	7.7	3