## Balamurugan Periaswamy

List of Publications by Year in descending order

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623734 940533 17 1,839 14 16 citations h-index g-index papers 18 18 18 3583 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Analysis of the human protein interactome and comparison with yeast, worm and fly interaction datasets. Nature Genetics, 2006, 38, 285-293.	21.4	433
2	A curated compendium of phosphorylation motifs. Nature Biotechnology, 2007, 25, 285-286.	17.5	345
3	A macromolecular approach to eradicate multidrug resistant bacterial infections while mitigating drug resistance onset. Nature Communications, 2018, 9, 917.	12.8	261
4	An evaluation of human protein-protein interaction data in the public domain. BMC Bioinformatics, 2006, 7, S19.	2.6	201
5	Microbiota-Derived Hydrogen Fuels Salmonella Typhimurium Invasion of the Gut Ecosystem. Cell Host and Microbe, 2013, 14, 641-651.	11.0	145
6	Enhanced CellClassifier: a multi-class classification tool for microscopy images. BMC Bioinformatics, 2010, 11, 30.	2.6	78
7	<i>Salmonella enterica</i> Serovar Typhimurium Binds to HeLa Cells via Fim-Mediated Reversible Adhesion and Irreversible Type Three Secretion System 1-Mediated Docking. Infection and Immunity, 2011, 79, 330-341.	2.2	78
8	Granulocytes Impose a Tight Bottleneck upon the Gut Luminal Pathogen Population during Salmonella Typhimurium Colitis. PLoS Pathogens, 2014, 10, e1004557.	4.7	73
9	Point mutations in the major outer membrane protein drive hypervirulence of a rapidly expanding clone of $\langle i \rangle$ Campylobacter jejuni $\langle i \rangle$ . Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10690-10695.	7.1	56
10	A Macromolecule Reversing Antibiotic Resistance Phenotype and Repurposing Drugs as Potent Antibiotics. Advanced Science, 2020, 7, 2001374.	11.2	53
11	Identification of Novel Phosphorylation Motifs Through an Integrative Computational and Experimental Analysis of the Human Phosphoproteome. Journal of Proteomics and Bioinformatics, 2011, 04, 22-35.	0.4	31
12	A Novel Phage Element of Salmonella enterica Serovar Enteritidis P125109 Contributes to Accelerated Type III Secretion System 2-Dependent Early Inflammation Kinetics in a Mouse Colitis Model. Infection and Immunity, 2012, 80, 3236-3246.	2.2	26
13	Accelerated Type III Secretion System 2-Dependent Enteropathogenesis by a <i>Salmonella enterica</i> Serovar Enteritidis PT4/6 Strain. Infection and Immunity, 2009, 77, 3569-3577.	2.2	25
14	Live Attenuated S. Typhimurium Vaccine with Improved Safety in Immuno-Compromised Mice. PLoS ONE, 2012, 7, e45433.	2.5	25
15	Putative Integrative Mobile Elements That Exploit the Xer Recombination Machinery Carrying $\langle i \rangle$ bla $\langle i \rangle <$ sub $\langle i \rangle$ (Integral of the Xer Recombination Machinery Carrying $\langle i \rangle$ bla $\langle i \rangle <$ sub $\langle i \rangle$ (Integral of the Xer Recombination Machinery Carrying Putation Points Integral of the Xer Recombination Machinery Carrying Singapore. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	8
16	Comprehensive Identification of Fim-Mediated Inversions in Uropathogenic Escherichia coli with Structural Variation Detection Using Relative Entropy. MSphere, 2019, 4, .	2.9	1
17	Draft Genome Sequence of Singapore <i>Klebsiella pneumoniae</i> subsp. <i>pneumoniae</i> lsolate DS32358_14, Which Contains the Carbapenemase Gene <i>bla</i> <sub>VIM-1</sub> . Genome Announcements, 2018, 6, .	0.8	0