

# Partho Ghosh

## List of Publications by Year in descending order

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50  
papers

3,096  
citations

186265

28  
h-index

223800

46  
g-index

58  
all docs

58  
docs citations

58  
times ranked

3257  
citing authors

#	ARTICLE	IF	CITATIONS
1	Contribution of <i>Streptococcus pyogenes</i> M87 protein to innate immune resistance and virulence. <i>Microbial Pathogenesis</i> , 2022, 169, 105636.	2.9	4
2	Nonimmune antibody interactions of Group A <i>Streptococcus</i> M and M-like proteins. <i>PLoS Pathogens</i> , 2021, 17, e1009248.	4.7	13
3	Determinants of adenine-mutagenesis in diversity-generating retroelements. <i>Nucleic Acids Research</i> , 2021, 49, 1033-1045.	14.5	5
4	Crystal structure of a <i>Thermus aquaticus</i> diversity-generating retroelement variable protein. <i>PLoS ONE</i> , 2019, 14, e0205618.	2.5	5
5	Diversity-generating retroelements: natural variation, classification and evolution inferred from a large-scale genomic survey. <i>Nucleic Acids Research</i> , 2018, 46, 11-24.	14.5	102
6	Variation, Indispensability, and Masking in the M protein. <i>Trends in Microbiology</i> , 2018, 26, 132-144.	7.7	33
7	Template-assisted synthesis of adenine-mutagenized cDNA by a retroelement protein complex. <i>Nucleic Acids Research</i> , 2018, 46, 9711-9725.	14.5	21
8	Group A Streptococcal M1 Protein Provides Resistance against the Antimicrobial Activity of Histones. <i>Scientific Reports</i> , 2017, 7, 43039.	3.3	29
9	Retroelement-guided protein diversification abounds in vast lineages of Bacteria and Archaea. <i>Nature Microbiology</i> , 2017, 2, 17045.	13.3	62
10	Group A streptococcal M protein activates the NLRP3 inflammasome. <i>Nature Microbiology</i> , 2017, 2, 1425-1434.	13.3	73
11	Conservation of the C-type lectin fold for accommodating massive sequence variation in archaeal diversity-generating retroelements. <i>BMC Structural Biology</i> , 2016, 16, 13.	2.3	15
12	Coiled-coil destabilizing residues in the group A <i>Streptococcus</i> M1 protein are required for functional interaction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 9515-9520.	7.1	37
13	Conserved patterns hidden within group A <i>Streptococcus</i> M protein hypervariability recognize human C4b-binding protein. <i>Nature Microbiology</i> , 2016, 1, 16155.	13.3	47
14	Targeted diversity generation by intraterrestrial archaea and archaeal viruses. <i>Nature Communications</i> , 2015, 6, 6585.	12.8	63
15	Group A Streptococcal M1 Protein Sequesters Cathelicidin to Evade Innate Immune Killing. <i>Cell Host and Microbe</i> , 2015, 18, 471-477.	11.0	51
16	M1 Protein from GAS Protects against Histone Killing. <i>FASEB Journal</i> , 2015, 29, 718.18.	0.5	0
17	Mutual Exclusivity of Hyaluronan and Hyaluronidase in Invasive Group A <i>Streptococcus</i> . <i>Journal of Biological Chemistry</i> , 2014, 289, 32303-32315.	3.4	30
18	The Fibrinogen-binding M1 Protein Reduces Pharyngeal Cell Adherence and Colonization Phenotypes of M1T1 Group A <i>Streptococcus</i> . <i>Journal of Biological Chemistry</i> , 2014, 289, 3539-3546.	3.4	22

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19	Diversity-generating Retroelements in Phage and Bacterial Genomes. <i>Microbiology Spectrum</i> , 2014, 2, .	3.0	47
20	Structure of the Essential Plasmodium Host Cell Traversal Protein SPECT1. <i>PLoS ONE</i> , 2014, 9, e114685.	2.5	12
21	Coiled-coil irregularities of the M1 protein structure promote M1-fibrinogen interaction and influence group A <i>Streptococcus</i> host cell interactions and virulence. <i>Journal of Molecular Medicine</i> , 2013, 91, 861-869.	3.9	14
22	Structure of the Essential Diversity-Generating Retroelement Protein bAvd and Its Functionally Important Interaction with Reverse Transcriptase. <i>Structure</i> , 2013, 21, 266-276.	3.3	26
23	Surface display of a massively variable lipoprotein by a <i>Legionella</i> diversity-generating retroelement. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8212-8217.	7.1	48
24	Structure and Interactions of the Cytoplasmic Domain of the Yersinia Type III Secretion Protein YscD. <i>Journal of Bacteriology</i> , 2012, 194, 5949-5958.	2.2	18
25	Two Translation Products of <i>Yersinia yscQ</i> Assemble To Form a Complex Essential to Type III Secretion. <i>Biochemistry</i> , 2012, 51, 1669-1677.	2.5	49
26	The Nonideal Coiled Coil of M Protein and Its Multifarious Functions in Pathogenesis. <i>Advances in Experimental Medicine and Biology</i> , 2011, 715, 197-211.	1.6	31
27	Streptococcal M1 protein constructs a pathological host fibrinogen network. <i>Nature</i> , 2011, 472, 64-68.	27.8	100
28	Conservation of the C-type lectin fold for massive sequence variation in a <i>Treponema</i> diversity-generating retroelement. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 14649-14653.	7.1	39
29	A Solvent-Exposed Patch in Chaperone-Bound YopE Is Required for Translocation by the Type III Secretion System. <i>Journal of Bacteriology</i> , 2010, 192, 3114-3122.	2.2	14
30	<i>Listeria monocytogenes</i> Internalin B Activates Junctional Endocytosis to Accelerate Intestinal Invasion. <i>PLoS Pathogens</i> , 2010, 6, e1000900.	4.7	86
31	M1 Protein Allows Group A Streptococcal Survival in Phagocyte Extracellular Traps through Cathelicidin Inhibition. <i>Journal of Innate Immunity</i> , 2009, 1, 202-214.	3.8	157
32	Coiled-Coil Irregularities and Instabilities in Group A <i>Streptococcus</i> M1 Are Required for Virulence. <i>Science</i> , 2008, 319, 1405-1408.	12.6	137
33	The Type III Secretion Chaperone SycE Promotes a Localized Disorder-to-Order Transition in the Natively Unfolded Effector YopE. <i>Journal of Biological Chemistry</i> , 2008, 283, 20857-20863.	3.4	39
34	Selective Ligand Recognition by a Diversity-Generating Retroelement Variable Protein. <i>PLoS Biology</i> , 2008, 6, e131.	5.6	47
35	Investigation of the Mechanism of Binding between Internalin B and Heparin Using Surface Plasmon Resonance. <i>Biochemistry</i> , 2007, 46, 2697-2706.	2.5	9
36	The C-type lectin fold as an evolutionary solution for massive sequence variation. <i>Nature Structural and Molecular Biology</i> , 2005, 12, 886-892.	8.2	92

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37	Process of Protein Transport by the Type III Secretion System. <i>Microbiology and Molecular Biology Reviews</i> , 2004, 68, 771-795.	6.6	351
38	GW domains of the <i>Listeria monocytogenes</i> invasion protein InlB are required for potentiation of Met activation. <i>Molecular Microbiology</i> , 2004, 52, 257-271.	2.5	67
39	Characterization of the calcium-binding sites of <i>Listeria monocytogenes</i> InlB. <i>Biochemical and Biophysical Research Communications</i> , 2004, 316, 379-386.	2.1	7
40	Multiple Regions of Internalin B Contribute to Its Ability to Turn on the Ras-Mitogen-activated Protein Kinase Pathway. <i>Journal of Biological Chemistry</i> , 2003, 278, 7783-7789.	3.4	43
41	In Vivo Phospholipase Activity of the <i>Pseudomonas aeruginosa</i> Cytotoxin ExoU and Protection of Mammalian Cells with Phospholipase A2 Inhibitors. <i>Journal of Biological Chemistry</i> , 2003, 278, 41326-41332.	3.4	172
42	Three-Dimensional Secretion Signals in Chaperone-Effector Complexes of Bacterial Pathogens. <i>Molecular Cell</i> , 2002, 9, 971-980.	9.7	162
43	GW domains of the <i>Listeria monocytogenes</i> invasion protein InlB are SH3-like and mediate binding to host ligands. <i>EMBO Journal</i> , 2002, 21, 5623-5634.	7.8	107
44	Structure of the Rho-activating domain of <i>Escherichia coli</i> cytotoxic necrotizing factor 1. <i>Nature Structural Biology</i> , 2001, 8, 584-588.	9.7	95
45	Structure of the <i>Yersinia</i> type III secretory system chaperone SycE. , 2001, 8, 974-978.		68
46	Structure of the InlB Leucine-Rich Repeats, a Domain that Triggers Host Cell Invasion by the Bacterial Pathogen <i>L. monocytogenes</i> . <i>Molecular Cell</i> , 1999, 4, 1063-1072.	9.7	161
47	Crystal structure of colicin Ia. <i>Nature</i> , 1997, 385, 461-464.	27.8	250
48	A carboxy-terminal fragment of colicin Ia forms ion channels. <i>Journal of Membrane Biology</i> , 1993, 134, 85-92.	2.1	19
49	Diversity-generating Retroelements in Phage and Bacterial Genomes. , 0, , 1237-1252.		8
50	An M protein coiled coil unfurls and exposes its hydrophobic core to capture LL-37. <i>ELife</i> , 0, 11, .	6.0	7