

# Patrick M Schaeffer

## List of Publications by Year in descending order

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45  
papers

1,242  
citations

394421

19  
h-index

377865

34  
g-index

48  
all docs

48  
docs citations

48  
times ranked

1200  
citing authors

#	ARTICLE	IF	CITATIONS
1	Rise of the terminator protein tus: A versatile tool in the biotechnologist's toolbox. <i>Analytica Chimica Acta</i> , 2022, 1213, 339946.	5.4	2
2	Delineation of the Ancestral Tus-Dependent Replication Fork Trap. <i>International Journal of Molecular Sciences</i> , 2021, 22, 13533.	4.1	4
3	A new bivalent fluorescent fusion protein for differential Cu(II) and Zn(II) ion detection in aqueous solution. <i>Analytica Chimica Acta</i> , 2020, 1101, 120-128.	5.4	13
4	High-Throughput Differential Scanning Fluorimetry of GFP-Tagged Proteins. <i>Methods in Molecular Biology</i> , 2020, 2089, 69-85.	0.9	10
5	Electrophoretic Mobility Shift Assays with GFP-Tagged Proteins (GFP-EMSA). <i>Methods in Molecular Biology</i> , 2020, 2089, 159-166.	0.9	1
6	Defining specific allergens for improved component-resolved diagnosis of shrimp allergy in adults. <i>Molecular Immunology</i> , 2019, 112, 330-337.	2.2	12
7	Selective protein unfolding: a universal mechanism of action for the development of irreversible inhibitors. <i>Chemical Communications</i> , 2018, 54, 1738-1741.	4.1	11
8	Negative regulators of cell death pathways in cancer: perspective on biomarkers and targeted therapies. <i>Apoptosis: an International Journal on Programmed Cell Death</i> , 2018, 23, 93-112.	4.9	44
9	Functional characterisation of <i>Burkholderia pseudomallei</i> biotin protein ligase: A toolkit for anti-melioidosis drug development. <i>Microbiological Research</i> , 2017, 199, 40-48.	5.3	7
10	A green fluorescent protein-based assay for high-throughput ligand-binding studies of a mycobacterial biotin protein ligase. <i>Microbiological Research</i> , 2017, 205, 35-39.	5.3	6
11	IgE reactivity to shrimp allergens in infants and their cross-reactivity to house dust mite. <i>Pediatric Allergy and Immunology</i> , 2017, 28, 703-707.	2.6	25
12	Green fluorescent protein-based assays for high-throughput functional characterization and ligand-binding studies of biotin protein ligase. <i>Analytical Methods</i> , 2016, 8, 418-424.	2.7	9
13	In-gel detection of biotin-protein conjugates with a green fluorescent streptavidin probe. <i>Analytical Methods</i> , 2015, 7, 2087-2092.	2.7	20
14	Tus-Ter-lock immuno-PCR assays for the sensitive detection of tropomyosin-specific IgE antibodies. <i>Bioanalysis</i> , 2014, 6, 465-476.	1.5	25
15	Dissecting the salt dependence of the Tus-Ter protein-DNA complexes by high-throughput differential scanning fluorimetry of a GFP-tagged Tus. <i>Molecular BioSystems</i> , 2013, 9, 3146.	2.9	17
16	A GFP-tagged nucleoprotein-based aggregation assay for anti-influenza drug discovery and antibody development. <i>Analyst</i> , 2013, 138, 6073.	3.5	4
17	ELISA and immuno-polymerase chain reaction assays for the sensitive detection of melioidosis. <i>Diagnostic Microbiology and Infectious Disease</i> , 2013, 75, 135-138.	1.8	11
18	Improved diagnosis of melioidosis using a 2-dimensional immunoarray. <i>Diagnostic Microbiology and Infectious Disease</i> , 2013, 77, 209-215.	1.8	9

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19	Differential Tus-Ter binding and lock formation: implications for DNA replication termination in Escherichia coli. <i>Molecular BioSystems</i> , 2012, 8, 2783.	2.9	20
20	A universal immuno-PCR platform for comparative and ultrasensitive quantification of dual affinity-tagged proteins in complex matrices. <i>Analyst, The</i> , 2012, 137, 5193.	3.5	10
21	Rapid determination of protein stability and ligand binding by differential scanning fluorimetry of GFP-tagged proteins. <i>RSC Advances</i> , 2012, 2, 11892.	3.6	30
22	A polyplex qPCR-based binding assay for protein-DNA interactions. <i>Analyst, The</i> , 2012, 137, 4111.	3.5	11
23	Combining RNA-DNA swapping and quantitative polymerase chain reaction for the detection of influenza A nucleoprotein. <i>Analytical Biochemistry</i> , 2012, 420, 121-126.	2.4	15
24	IgG-detection devices for the Tus-Ter-lock immuno-PCR diagnostic platform. <i>Analyst, The</i> , 2011, 136, 4815.	3.5	24
25	Development of a protease activity assay using heat-sensitive Tus-GFP fusion protein substrates. <i>Analytical Biochemistry</i> , 2011, 415, 126-133.	2.4	10
26	Ultrasensitive detection of antibodies using a new Tus-Ter-lock immunoPCR system. <i>Molecular BioSystems</i> , 2010, 6, 1173.	2.9	27
27	Quantitative determination of protein stability and ligand binding using a green fluorescent protein reporter system. <i>Molecular BioSystems</i> , 2010, 6, 1285.	2.9	57
28	Synthesis and Applications of Covalent Protein-DNA Conjugates. <i>Australian Journal of Chemistry</i> , 2009, 62, 1328.	0.9	9
29	Site-specific covalent attachment of DNA to proteins using a photoactivatable Tus-Ter complex. <i>Chemical Communications</i> , 2009, , 3050.	4.1	25
30	Single-molecule studies of fork dynamics in Escherichia coli DNA replication. <i>Nature Structural and Molecular Biology</i> , 2008, 15, 170-176.	8.2	136
31	Multiple oligomeric forms of Escherichia coli DnaB helicase revealed by electrospray ionisation mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2007, 21, 132-140.	1.5	8
32	Proteomic dissection of DNA polymerization. <i>Expert Review of Proteomics</i> , 2006, 3, 197-211.	3.0	11
33	Kinetic and Crystallographic Analysis of Mutant Escherichia coli Aminopeptidase P: Insights into Substrate Recognition and the Mechanism of Catalysis. <i>Biochemistry</i> , 2006, 45, 964-975.	2.5	41
34	A Molecular Mousetrap Determines Polarity of Termination of DNA Replication in E. coli. <i>Cell</i> , 2006, 125, 1309-1319.	28.9	114
35	Monomeric solution structure of the helicase-binding domain of Escherichia coli DnaG primase. <i>FEBS Journal</i> , 2006, 273, 4997-5009.	4.7	25
36	Helicase binding to DnaI exposes a cryptic DNA-binding site during helicase loading in Bacillus subtilis. <i>Nucleic Acids Research</i> , 2006, 34, 5247-5258.	14.5	50

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37	A molecular mousetrap determines polarity of replication fork arrest at Tus•Ter sites in <i>E. coli</i> . <i>FASEB Journal</i> , 2006, 20, A911.	0.5	0
38	Protein-Protein Interactions in the Eubacterial Replisome. <i>IUBMB Life</i> , 2005, 57, 5-12.	3.4	74
39	Crystal and Solution Structures of the Helicase-binding Domain of <i>Escherichia coli</i> Primase. <i>Journal of Biological Chemistry</i> , 2005, 280, 11495-11504.	3.4	62
40	Integron-associated Mobile Gene Cassettes Code for Folded Proteins: The Structure of Bal32a, a New Member of the Adaptable $\beta$ -Barrel Family. <i>Journal of Molecular Biology</i> , 2005, 346, 1229-1241.	4.2	20
41	Optimization of an <i>Escherichia coli</i> system for cell-free synthesis of selectively <sup>15</sup> N-labelled proteins for rapid analysis by NMR spectroscopy. <i>FEBS Journal</i> , 2004, 271, 4084-4093.	0.2	87
42	Expression, purification, crystallization, and NMR studies of the helicase interaction domain of <i>Escherichia coli</i> DnaG primase. <i>Protein Expression and Purification</i> , 2004, 33, 304-310.	1.3	11
43	Molecular tectonics II: Synthesis of molecular sheets by self-assembly of complementary molecular units in the solid state. <i>Tetrahedron Letters</i> , 1996, 37, 1405-1408.	1.4	41
44	Molecular Tectonics: Self-Assembly of Charged Molecular Tectons into One- and Two-Dimensional Solids. , 1996, , 129-142.		3
45	A molecular approach to solid-state synthesis: prediction and synthesis of self-assembled infinite rods. <i>Journal of the Chemical Society Chemical Communications</i> , 1994, , 2135.	2.0	69