## Patrick England

List of Publications by Year in descending order

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89 papers 6,989 citations

70961 41 h-index 80 g-index

96 all docs 96
docs citations

96 times ranked 11670 citing authors

#	Article	IF	CITATIONS
1	Specificity and affinity of human $Fc\hat{l}^3$ receptors and their polymorphic variants for human $IgG$ subclasses. Blood, 2009, 113, 3716-3725.	0.6	1,218
2	Structural basis of potent Zika–dengue virus antibody cross-neutralization. Nature, 2016, 536, 48-53.	13.7	465
3	Efficient protection and isolation of ubiquitylated proteins using tandem ubiquitinâ€binding entities. EMBO Reports, 2009, 10, 1250-1258.	2.0	407
4	ESAT-6 from Mycobacterium tuberculosis Dissociates from Its Putative Chaperone CFP-10 under Acidic Conditions and Exhibits Membrane-Lysing Activity. Journal of Bacteriology, 2007, 189, 6028-6034.	1.0	272
5	Clostridium difficile toxin expression is inhibited by the novel regulator TcdC. Molecular Microbiology, 2007, 64, 1274-1288.	1.2	213
6	The Disulfide Bonds in Glycoprotein E2 of Hepatitis C Virus Reveal the Tertiary Organization of the Molecule. PLoS Pathogens, 2010, 6, e1000762.	2.1	210
7	AIF promotes chromatinolysis and caspase-independent programmed necrosis by interacting with histone H2AX. EMBO Journal, 2010, 29, 1585-1599.	3.5	197
8	Single-cell deep phenotyping of IgG-secreting cells for high-resolution immune monitoring. Nature Biotechnology, 2017, 35, 977-982.	9.4	193
9	High-throughput single-cell activity-based screening and sequencing of antibodies using droplet microfluidics. Nature Biotechnology, 2020, 38, 715-721.	9.4	163
10	Transglutaminase is essential for IgA nephropathy development acting through IgA receptors. Journal of Experimental Medicine, 2012, 209, 793-806.	4.2	145
11	Full-length extracellular region of the var2CSA variant of PfEMP1 is required for specific, high-affinity binding to CSA. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 4884-4889.	3.3	137
12	Tethering of HP1 proteins to chromatin is relieved by phosphoacetylation of histone H3. EMBO Reports, 2004, 5, 490-496.	2.0	132
13	Fc $\hat{I}^3$ RIV is a mouse IgE receptor that resembles macrophage Fc $\hat{I}$ µRI in humans and promotes IgE-induced lung inflammation. Journal of Clinical Investigation, 2008, $118$ , $3738$ - $3750$ .	3.9	132
14	The high-affinity human IgG receptor Fc $\hat{l}^3$ RI (CD64) promotes IgG-mediated inflammation, anaphylaxis, and antitumor immunotherapy. Blood, 2013, 121, 1563-1573.	0.6	120
15	A neutralizing monoclonal antibody (mAb A24) directed against the transferrin receptor induces apoptosis of tumor T lymphocytes from ATL patients. Blood, 2004, 103, 1838-1845.	0.6	101
16	Attenuation of Rabies Virulence: Takeover by the Cytoplasmic Domain of Its Envelope Protein. Science Signaling, 2010, 3, ra5.	1.6	100
17	Structural Flexibility of a Conserved Antigenic Region in Hepatitis C Virus Glycoprotein E2 Recognized by Broadly Neutralizing Antibodies. Journal of Virology, 2015, 89, 2170-2181.	1.5	96
18	Cadherin-23, myosin VIIa and harmonin, encoded by Usher syndrome type I genes, form a ternary complex and interact with membrane phospholipids. Human Molecular Genetics, 2010, 19, 3557-3565.	1.4	94

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19	Single-domain antibodies recognize selectively small oligomeric forms of amyloid $\hat{l}^2$ , prevent $\hat{Al}^2$ -induced neurotoxicity and inhibit fibril formation. Molecular Immunology, 2009, 46, 695-704.	1.0	87
20	A global benchmark study using affinity-based biosensors. Analytical Biochemistry, 2009, 386, 194-216.	1.1	85
21	A short synthetic peptide inhibits signal transduction, migration and angiogenesis mediated by Tie2 receptor. EMBO Reports, 2004, 5, 262-267.	2.0	81
22	The CymR Regulator in Complex with the Enzyme CysK Controls Cysteine Metabolism in Bacillus subtilis. Journal of Biological Chemistry, 2008, 283, 35551-35560.	1.6	81
23	Quality assessment and optimization of purified protein samples: why and how?. Microbial Cell Factories, 2014, 13, 180.	1.9	79
24	IgG subclasses determine pathways of anaphylaxis in mice. Journal of Allergy and Clinical Immunology, 2017, 139, 269-280.e7.	1.5	78
25	Structural Basis for the ABO Blood-Group Dependence of Plasmodium falciparum Rosetting. PLoS Pathogens, 2012, 8, e1002781.	2.1	74
26	Knowledge-based Design of Reagentless Fluorescent Biosensors from Recombinant Antibodies. Journal of Molecular Biology, 2002, 318, 429-442.	2.0	71
27	A Multilaboratory Comparison of Calibration Accuracy and the Performance of External References in Analytical Ultracentrifugation. PLoS ONE, 2015, 10, e0126420.	1.1	71
28	Structural basis of myosin V Rab GTPase-dependent cargo recognition. Proceedings of the National Academy of Sciences of the United States of America, 2013, $110$ , $20443-20448$ .	3.3	70
29	The structural basis of Arf effector specificity: the crystal structure of ARF6 in a complex with JIP4. EMBO Journal, 2009, 28, 2835-2845.	3.5	68
30	Structural and Thermodynamic Bases for the Design of Pure Prolactin Receptor Antagonists. Journal of Biological Chemistry, 2007, 282, 33118-33131.	1.6	66
31	Mechanistic Insight into Bunyavirus-Induced Membrane Fusion from Structure-Function Analyses of the Hantavirus Envelope Glycoprotein Gc. PLoS Pathogens, 2016, 12, e1005813.	2.1	66
32	Calcium-Induced Folding and Stabilization of the Intrinsically Disordered RTX Domain of the CyaA Toxin. Biophysical Journal, 2010, 99, 3744-3753.	0.2	64
33	Rabies virus matrix protein interplay with elF3, new insights into rabies virus pathogenesis. Nucleic Acids Research, 2007, 35, 1522-1532.	<b>6.</b> 5	62
34	Identification and Characterization of the Binding Site of the Respiratory Syncytial Virus Phosphoprotein to RNA-Free Nucleoprotein. Journal of Virology, 2015, 89, 3484-3496.	1.5	60
35	Crystal Structure of an Affinity-matured Prolactin Complexed to Its Dimerized Receptor Reveals the Topology of Hormone Binding Site 2. Journal of Biological Chemistry, 2010, 285, 8422-8433.	1.6	59
36	Energetic and Kinetic Contributions of Contact Residues of Antibody D1.3 in the Interaction with Lysozyme. Biochemistry, 1997, 36, 164-172.	1.2	58

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37	A Druggable Pocket at the Nucleocapsid/Phosphoprotein Interaction Site of Human Respiratory Syncytial Virus. Journal of Virology, 2015, 89, 11129-11143.	1.5	56
38	Stable trimerization of recombinant rabies virus glycoprotein ectodomain is required for interaction with the p75NTR receptor. Journal of General Virology, 2005, 86, 2543-2552.	1.3	53
39	Coordinated recruitment of Spir actin nucleators and myosin V motors to Rab11 vesicle membranes. ELife, 2016, $5$ , .	2.8	53
40	Structure of a <i>Plasmodium falciparum</i> PfEMP1 rosetting domain reveals a role for the N-terminal segment in heparin-mediated rosette inhibition. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 5243-5248.	3.3	51
41	The FHAâ€containing protein GarA acts as a phosphorylationâ€dependent molecular switch in mycobacterial signaling. FEBS Letters, 2009, 583, 301-307.	1.3	46
42	Structural Characterization of the Stem–Stem Dimerization Interface between Prolactin Receptor Chains Complexed with the Natural Hormone. Journal of Molecular Biology, 2010, 404, 112-126.	2.0	45
43	Insights into the Rrf2 repressor familyâ€∫–â€∫the structure of CymR, the global cysteine regulator of <i>Bacillusâ€∫subtilis⟨/i⟩. FEBS Journal, 2011, 278, 2689-2701.</i>	2.2	45
44	SARS-CoV-2 Nsp3 unique domain SUD interacts with guanine quadruplexes and G4-ligands inhibit this interaction. Nucleic Acids Research, 2021, 49, 7695-7712.	6.5	43
45	Peroxynitrite transforms nerve growth factor into an apoptotic factor for motor neurons. Free Radical Biology and Medicine, 2006, 41, 1632-1644.	1.3	41
46	Several regions of the repeat domain of the Staphylococcus capraeautolysin, AtlC, are involved in fibronectin binding. FEMS Microbiology Letters, 2002, 213, 193-197.	0.7	40
47	The epitope arrangement on flavivirus particles contributes to Mab C10's extraordinary neutralization breadth across Zika and dengue viruses. Cell, 2021, 184, 6052-6066.e18.	13.5	38
48	Pilotin–secretin recognition in the type II secretion system of <i>Klebsiella oxytoca</i> Molecular Microbiology, 2011, 82, 1422-1432.	1.2	37
49	Outer Membrane Targeting of Secretin PulD Protein Relies on Disordered Domain Recognition by a Dedicated Chaperone. Journal of Biological Chemistry, 2011, 286, 38833-38843.	1.6	36
50	SepF is the FtsZ anchor in archaea, with features of an ancestral cell division system. Nature Communications, 2021, 12, 3214.	5.8	35
51	Characterization of the elongasome core PBP2 : MreC complex of <i>Helicobacter pylori</i> Molecular Microbiology, 2011, 82, 68-86.	1.2	34
52	SUN Proteins Belong to a Novel Family of $\hat{l}^2$ -(1,3)-Glucan-modifying Enzymes Involved in Fungal Morphogenesis. Journal of Biological Chemistry, 2013, 288, 13387-13396.	1.6	34
53	Structural basis for the increased processivity of D-family DNA polymerases in complex with PCNA. Nature Communications, 2020, 11, 1591.	5.8	34
54	Analysis of SUMOylated proteins using SUMO-traps. Scientific Reports, 2013, 3, 1690.	1.6	32

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55	Conformational Flexibility in the Immunoglobulin-Like Domain of the Hepatitis C Virus Glycoprotein E2. MBio, 2017, 8, .	1.8	31
56	Calcium-dependent disorder-to-order transitions are central to the secretion and folding of the CyaA toxin of Bordetella pertussis, the causative agent of whooping cough. Toxicon, 2018, 149, 37-44.	0.8	29
57	Essential dynamic interdependence of FtsZ and SepF for Z-ring and septum formation in Corynebacterium glutamicum. Nature Communications, 2020, 11, 1641.	5.8	29
58	Binding of the Unorthodox Transcription Activator, Crl, to the Components of the Transcription Machinery. Journal of Biological Chemistry, 2008, 283, 33455-33464.	1.6	28
59	Structural activation of the transcriptional repressor EthR from Mycobacterium tuberculosis by single amino acid change mimicking natural and synthetic ligands. Nucleic Acids Research, 2012, 40, 3018-3030.	6.5	28
60	Structure of the prefusion-locking broadly neutralizing antibody RVC20 bound to the rabies virus glycoprotein. Nature Communications, 2020, 11, 596.	5.8	28
61	The Scc Spirochetal Coiled-Coil Protein Forms Helix-Like Filaments and Binds to Nucleic Acids Generating Nucleoprotein Structures. Journal of Bacteriology, 2006, 188, 469-476.	1.0	24
62	An extracellular <scp> <i>Leptospira interrogans</i> </scp> leucineâ€rich repeat protein binds human E― and VEâ€cadherins. Cellular Microbiology, 2019, 21, e12949.	1.1	23
63	The N-terminal Domain of Hepatocyte Growth Factor Inhibits the Angiogenic Behavior of Endothelial Cells Independently from Binding to the c-met Receptor. Journal of Biological Chemistry, 2003, 278, 37400-37408.	1.6	20
64	Diversity and junction residues as hotspots of binding energy in an antibody neutralizing the dengue virus. FEBS Journal, 2006, 273, 34-46.	2,2	19
65	Apparent cooperativity for carbamoylphosphate in Escherichia coli aspartate transcarbamoylase only reflects cooperativity for aspartate. FEBS Journal, 1994, 222, 775-780.	0.2	18
66	Crl Binds to Domain 2 of $ f  < \sup S < \sup and Confers a Competitive Advantage on a Natural  f  < \sup S < \min and Confers a Competitive Advantage on a Natural  f  < \sup A < \min and Confers a Competitive Advantage on a Natural  f  < \sup A < \sup A$	1.0	17
67	Evolution and activation mechanism of the flavivirus class II membrane-fusion machinery. Nature Communications, 2022, $13$ , .	5.8	17
68	Hydrodynamic characterization of recombinant human fibrinogen species. Thrombosis Research, 2013, 132, e48-e53.	0.8	16
69	Development of a highly specific and sensitive VHH-based sandwich immunoassay for the detection of the SARS-CoV-2 nucleoprotein. Journal of Biological Chemistry, 2022, 298, 101290.	1.6	16
70	The Tip of the Four N-Terminal $\hat{l}_{\pm}$ -Helices of Clostridium sordellii Lethal Toxin Contains the Interaction Site with Membrane Phosphatidylserine Facilitating Small GTPases Glucosylation. Toxins, 2016, 8, 90.	1.5	15
71	Analysis of the Escherichia coli glucosamine-6-phosphate synthase activity by isothermal titration calorimetry and differential scanning calorimetry. Archives of Biochemistry and Biophysics, 2010, 498, 95-104.	1.4	14
72	Residue 146 regulates prolactin receptor folding, basal activity and ligand-responsiveness: Potential implications in breast tumorigenesis. Molecular and Cellular Endocrinology, 2015, 401, 173-188.	1.6	14

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73	Phosphorylcholineâ^'Carbohydrateâ^' Protein Conjugates Efficiently Induce Hapten-Specific Antibodies Which Recognize BothStreptococcuspneumoniaeandNeisseriameningitidis:Â A Potential Multitarget Vaccine against Respiratory Infections. Journal of Medicinal Chemistry, 2004, 47, 3916-3919.	2.9	13
74	Energetics of 5-bromo-4-chloro-3-indolyl- $\hat{l}$ ±-D-mannose binding to the Parkia platyce phalaseed lectin and its use for MAD phasing. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 326-331.	0.7	13
75	Interaction of a Partially Disordered Antisigma Factor with Its Partner, the Signaling Domain of the TonB-Dependent Transporter HasR. PLoS ONE, 2014, 9, e89502.	1.1	13
76	The catalytic domains of <i>Clostridium sordellii</i> lethal toxin and related large clostridial glucosylating toxins specifically recognize the negatively charged phospholipids phosphatidylserine and phosphatidic acid. Cellular Microbiology, 2015, 17, 1477-1493.	1.1	13
77	The Translocation Domain of Botulinum Neurotoxin A Moderates the Propensity of the Catalytic Domain to Interact with Membranes at Acidic pH. PLoS ONE, 2016, 11, e0153401.	1.1	13
78	High-affinity autoreactive plasma cells disseminate through multiple organs in patients with immune thrombocytopenic purpura. Journal of Clinical Investigation, 2022, 132, .	3.9	13
79	Dengue virus NS1 protein conveys proâ€inflammatory signals by docking onto highâ€density lipoproteins. EMBO Reports, 2022, 23, .	2.0	13
80	A multi-laboratory benchmark study of isothermal titration calorimetry (ITC) using Ca2+ and Mg2+ binding to EDTA. European Biophysics Journal, 2021, 50, 429-451.	1.2	12
81	Structural and functional features of Crl proteins and identification of conserved surface residues required for interaction with the RpoS/ $l$ fS subunit of RNA polymerase. Biochemical Journal, 2014, 463, 215-224.	1.7	11
82	Conformational switch of harmonin, a submembrane scaffold protein of the hair cell mechanoelectrical transduction machinery. FEBS Letters, 2017, 591, 2299-2310.	1.3	9
83	The stress sigma factor of RNA polymerase RpoS/ $\hat{l}fS$ is a solvent-exposed open molecule in solution. Biochemical Journal, 2018, 475, 341-354.	1.7	7
84	The Stable Interaction Between Signal Peptidase LepB of Escherichia coli and Nuclease Bacteriocins Promotes Toxin Entry into the Cytoplasm. Journal of Biological Chemistry, 2015, 290, 30783-30796.	1.6	5
85	Macromolecular interactions in vitro, comparing classical and novel approaches. European Biophysics Journal, 2021, 50, 313-330.	1.2	5
86	A Residue Quartet in the Extracellular Domain of the Prolactin Receptor Selectively Controls Mitogen-activated Protein Kinase Signaling. Journal of Biological Chemistry, 2015, 290, 11890-11904.	1.6	4
87	Efficient monitoring of protein ubiquitylation levels using <scp>TUBE</scp> sâ€based microarrays. FEBS Letters, 2016, 590, 2748-2756.	1.3	4
88	Real-Time Surface Plasmon Resonance (SPR) for the Analysis of Interactions Between SUMO Traps and Mono- or PolySUMO Moieties. Methods in Molecular Biology, 2016, 1475, 99-107.	0.4	2
89	Community-building and promotion of technological excellence in molecular biophysics: the ARBRE–MOBIEU network. European Biophysics Journal, 2021, 50, 307-311.	1.2	1