

Travis Beddoe

List of Publications by Year in descending order

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116
papers

6,841
citations

81839

39
h-index

62565

80
g-index

126
all docs

126
docs citations

126
times ranked

8820
citing authors

#	ARTICLE	IF	CITATIONS
1	AB5 subtilase cytotoxin inactivates the endoplasmic reticulum chaperone BiP. <i>Nature</i> , 2006, 443, 548-552.	13.7	351
2	Butyrophilin 3A1 binds phosphorylated antigens and stimulates human $\gamma\delta$ T cells. <i>Nature Immunology</i> , 2013, 14, 908-916.	7.0	351
3	T cell receptor recognition of a 'super-bulged' major histocompatibility complex class I bound peptide. <i>Nature Immunology</i> , 2005, 6, 1114-1122.	7.0	280
4	Recognition of vitamin B metabolites by mucosal-associated invariant T cells. <i>Nature Communications</i> , 2013, 4, 2142.	5.8	261
5	CD1d-lipid antigen recognition by the $\gamma\delta$ TCR. <i>Nature Immunology</i> , 2013, 14, 1137-1145.	7.0	256
6	Structural basis of a unique interferon- γ signaling axis mediated via the receptor IFNAR1. <i>Nature Immunology</i> , 2013, 14, 901-907.	7.0	255
7	A molecular basis underpinning the T cell receptor heterogeneity of mucosal-associated invariant T cells. <i>Journal of Experimental Medicine</i> , 2014, 211, 1585-1600.	4.2	245
8	Current Threat of Triclabendazole Resistance in <i>Fasciola hepatica</i> . <i>Trends in Parasitology</i> , 2016, 32, 458-469.	1.5	233
9	Incorporation of a non-human glycan mediates human susceptibility to a bacterial toxin. <i>Nature</i> , 2008, 456, 648-652.	13.7	217
10	A T cell receptor flattens a bulged antigenic peptide presented by a major histocompatibility complex class I molecule. <i>Nature Immunology</i> , 2007, 8, 268-276.	7.0	206
11	Structure, biological functions and applications of the AB5 toxins. <i>Trends in Biochemical Sciences</i> , 2010, 35, 411-418.	3.7	204
12	CD94-NKG2A recognition of human leukocyte antigen (HLA)-E bound to an HLA class I leader sequence. <i>Journal of Experimental Medicine</i> , 2008, 205, 725-735.	4.2	198
13	The CDR3 regions of an immunodominant T cell receptor dictate the 'energetic landscape' of peptide-MHC recognition. <i>Nature Immunology</i> , 2005, 6, 171-180.	7.0	187
14	Killer cell immunoglobulin-like receptor 3DL1-mediated recognition of human leukocyte antigen B. <i>Nature</i> , 2011, 479, 401-405.	13.7	174
15	High Resolution Structures of Highly Bulged Viral Epitopes Bound to Major Histocompatibility Complex Class I. <i>Journal of Biological Chemistry</i> , 2005, 280, 23900-23909.	1.6	162
16	Natural HLA Class I Polymorphism Controls the Pathway of Antigen Presentation and Susceptibility to Viral Evasion. <i>Journal of Experimental Medicine</i> , 2004, 200, 13-24.	4.2	159
17	The 2.0-Å... Crystal Structure of eqFP611, a Far Red Fluorescent Protein from the Sea Anemone <i>Entacmaea quadricolor</i> . <i>Journal of Biological Chemistry</i> , 2003, 278, 44626-44631.	1.6	158
18	Crystal structure of the human T cell receptor CD3 heterodimer complexed to the therapeutic mAb OKT3. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 7675-7680.	3.3	148

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19	The 2.2 Å... Crystal Structure of a Pocilloporin Pigment Reveals a Nonplanar Chromophore Conformation. <i>Structure</i> , 2003, 11, 275-284.	1.6	127
20	The crystal structure of myelin oligodendrocyte glycoprotein, a key autoantigen in multiple sclerosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 11059-11064.	3.3	121
21	Biased T Cell Receptor Usage Directed against Human Leukocyte Antigen DQ8-Restricted Gliadin Peptides Is Associated with Celiac Disease. <i>Immunity</i> , 2012, 37, 611-621.	6.6	121
22	Antigen Ligation Triggers a Conformational Change within the Constant Domain of the $\hat{I}\hat{I}^2$ T Cell Receptor. <i>Immunity</i> , 2009, 30, 777-788.	6.6	111
23	Recognition of \hat{I}^2 -linked self glycolipids mediated by natural killer T cell antigen receptors. <i>Nature Immunology</i> , 2011, 12, 827-833.	7.0	111
24	Structural basis for a major histocompatibility complex class Ibâ€restricted T cell response. <i>Nature Immunology</i> , 2006, 7, 256-264.	7.0	109
25	A structural basis for selection and cross-species reactivity of the semi-invariant NKT cell receptor in CD1d/glycolipid recognition. <i>Journal of Experimental Medicine</i> , 2006, 203, 661-673.	4.2	105
26	Hard wiring of T cell receptor specificity for the major histocompatibility complex is underpinned by TCR adaptability. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 10608-10613.	3.3	101
27	Natural micropolymorphism in human leukocyte antigens provides a basis for genetic control of antigen recognition. <i>Journal of Experimental Medicine</i> , 2009, 206, 209-219.	4.2	93
28	Subtle Changes in Peptide Conformation Profoundly Affect Recognition of the Non-Classical MHC Class I Molecule HLA-E by the CD94â€NKG2 Natural Killer Cell Receptors. <i>Journal of Molecular Biology</i> , 2008, 377, 1297-1303.	2.0	88
29	The nascent polypeptide-associated complex (NAC) promotes interaction of ribosomes with the mitochondrial surface in vivo. <i>FEBS Letters</i> , 2002, 516, 213-216.	1.3	87
30	The Heterodimeric Assembly of the CD94-NKG2 Receptor Family and Implications for Human Leukocyte Antigen-E Recognition. <i>Immunity</i> , 2007, 27, 900-911.	6.6	87
31	A minimal binding footprint on CD1d-glycolipid is a basis for selection of the unique human NKT TCR. <i>Journal of Experimental Medicine</i> , 2008, 205, 939-949.	4.2	83
32	The High Resolution Crystal Structure of the Human Tumor Suppressor Maspin Reveals a Novel Conformational Switch in the G-helix. <i>Journal of Biological Chemistry</i> , 2005, 280, 22356-22364.	1.6	69
33	The structural basis for autonomous dimerization of the pre-T-cell antigen receptor. <i>Nature</i> , 2010, 467, 844-848.	13.7	68
34	NKT TCR Recognition of CD1d- \hat{I}^2 -Galactosylceramide. <i>Journal of Immunology</i> , 2011, 187, 4705-4713.	0.4	62
35	Functional and Structural Characteristics of NY-ESO-1-related HLA A2-restricted Epitopes and the Design of a Novel Immunogenic Analogue. <i>Journal of Biological Chemistry</i> , 2004, 279, 23438-23446.	1.6	61
36	Delivery of nascent polypeptides to the mitochondrial surface. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2002, 1592, 35-39.	1.9	58

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37	Enzymatic Properties of an Ecto-nucleoside Triphosphate Diphosphohydrolase from <i>Legionella pneumophila</i> . <i>Journal of Biological Chemistry</i> , 2008, 283, 12909-12918.	1.6	54
38	Crystal Structure of LipL32, the Most Abundant Surface Protein of Pathogenic <i>Leptospira</i> spp.. <i>Journal of Molecular Biology</i> , 2009, 387, 1229-1238.	2.0	53
39	Disparate thermodynamics governing T cell receptor-MHC-I interactions implicate extrinsic factors in guiding MHC restriction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 6641-6646.	3.3	52
40	A mortise-tenon joint in the transmembrane domain modulates autotransporter assembly into bacterial outer membranes. <i>Nature Communications</i> , 2014, 5, 4239.	5.8	46
41	Targeting of a natural killer cell receptor family by a viral immunoevasin. <i>Nature Immunology</i> , 2013, 14, 699-705.	7.0	41
42	Crystal Structure of a <i>Legionella pneumophila</i> Ecto-Triphosphate Diphosphohydrolase, A Structural and Functional Homolog of the Eukaryotic NTPDases. <i>Structure</i> , 2010, 18, 228-238.	1.6	39
43	The Endoplasmic Reticulum-Mitochondrion Tether ERMES Orchestrates Fungal Immune Evasion, Illuminating Inflammasome Responses to Hyphal Signals. <i>MSphere</i> , 2016, 1, .	1.3	39
44	Structure and Function of the Oxidoreductase DsbA1 from <i>Neisseria meningitidis</i> . <i>Journal of Molecular Biology</i> , 2009, 394, 931-943.	2.0	36
45	The Structure of H-2Kb and Kb*08 Complexed to a Herpes Simplex Virus Determinant: Evidence for a Conformational Switch That Governs T Cell Repertoire Selection and Viral Resistance. <i>Journal of Immunology</i> , 2004, 173, 402-409.	0.4	31
46	Antagonism of Antiviral and Allogeneic Activity of a Human Public CTL Clonotype by a Single Altered Peptide Ligand: Implications for Allograft Rejection. <i>Journal of Immunology</i> , 2005, 174, 5593-5601.	0.4	30
47	Tetrahydrolipstatin Inhibition, Functional Analyses, and Three-dimensional Structure of a Lipase Essential for Mycobacterial Viability. <i>Journal of Biological Chemistry</i> , 2010, 285, 30050-30060.	1.6	30
48	The Interaction of KIR3DL1*001 with HLA Class I Molecules Is Dependent upon Molecular Microarchitecture within the Bw4 Epitope. <i>Journal of Immunology</i> , 2015, 194, 781-789.	0.4	25
49	A Biophysical Analysis of the Tetratricopeptide Repeat-rich Mitochondrial Import Receptor, Tom70, Reveals an Elongated Monomer That Is Inherently Flexible, Unstable, and Unfolds via a Multistate Pathway. <i>Journal of Biological Chemistry</i> , 2004, 279, 46448-46454.	1.6	24
50	Hijacking of a Substrate-binding Protein Scaffold for use in Mycobacterial Cell Wall Biosynthesis. <i>Journal of Molecular Biology</i> , 2006, 359, 983-997.	2.0	23
51	Crystal Structure of a UDP-glucose-specific Glycosyltransferase from a Mycobacterium Species. <i>Journal of Biological Chemistry</i> , 2008, 283, 27881-27890.	1.6	23
52	Structural and Biochemical Characterization of the Oxidoreductase NmDsbA3 from <i>Neisseria meningitidis</i> . <i>Journal of Biological Chemistry</i> , 2008, 283, 32452-32461.	1.6	23
53	Structure-function analyses of a pertussis-like toxin from pathogenic <i>Escherichia coli</i> reveal a distinct mechanism of inhibition of trimeric G-proteins. <i>Journal of Biological Chemistry</i> , 2017, 292, 15143-15158.	1.6	23
54	Crocodilepox Virus Evolutionary Genomics Supports Observed Poxvirus Infection Dynamics on Saltwater Crocodile (<i>Crocodylus porosus</i>). <i>Viruses</i> , 2019, 11, 1116.	1.5	23

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55	EcxAB Is a Founding Member of a New Family of Metalloprotease AB5 Toxins with a Hybrid Cholera-like B Subunit. <i>Structure</i> , 2013, 21, 2003-2013.	1.6	22
56	Determination of the prevalence and intensity of <i>Fasciola hepatica</i> infection in dairy cattle from six irrigation regions of Victoria, South-eastern Australia, further identifying significant triclabendazole resistance on three properties. <i>Veterinary Parasitology</i> , 2020, 277, 109019.	0.7	22
57	Structural Basis of Subtilase Cytotoxin SubAB Assembly. <i>Journal of Biological Chemistry</i> , 2013, 288, 27505-27516.	1.6	21
58	Galectin-11: A novel host mediator targeting specific stages of the gastrointestinal nematode parasite, <i>Haemonchus contortus</i> . <i>International Journal for Parasitology</i> , 2015, 45, 791-796.	1.3	20
59	A novel ex vivo immunoproteomic approach characterising <i>Fasciola hepatica</i> tegumental antigens identified using immune antibody from resistant sheep. <i>International Journal for Parasitology</i> , 2017, 47, 555-567.	1.3	20
60	A conserved energetic footprint underpins recognition of human leukocyte antigen-E by two distinct $\hat{I}\hat{I}^2$ T cell receptors. <i>Journal of Biological Chemistry</i> , 2017, 292, 21149-21158.	1.6	20
61	Development of a multiplex quantitative PCR assay for detection and quantification of DNA from <i>Fasciola hepatica</i> and the intermediate snail host, <i>Austropeplea tomentosa</i> , in water samples. <i>Veterinary Parasitology</i> , 2018, 259, 17-24.	0.7	20
62	The production, purification and crystallization of a pocilloporin pigment from a reef-forming coral. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2003, 59, 597-599.	2.5	19
63	The B Subunit of an AB5 Toxin Produced by <i>Salmonella enterica</i> Serovar Typhi Up-Regulates Chemokines, Cytokines, and Adhesion Molecules in Human Macrophage, Colonic Epithelial, and Brain Microvascular Endothelial Cell Lines. <i>Infection and Immunity</i> , 2013, 81, 673-683.	1.0	19
64	Disrupting the Allosteric Interaction between the <i>Plasmodium falciparum</i> cAMP-dependent Kinase and Its Regulatory Subunit. <i>Journal of Biological Chemistry</i> , 2016, 291, 25375-25386.	1.6	14
65	The development and deployment of a field-based loop mediated isothermal amplification assay for virulent <i>Dichelobacter nodosus</i> detection on Australian sheep. <i>PLoS ONE</i> , 2018, 13, e0204310.	1.1	14
66	The RNA-Dependent-RNA Polymerase, an Emerging Antiviral Drug Target for the Hendra Virus. <i>Current Drug Targets</i> , 2014, 15, 103-113.	1.0	13
67	Crystal Structure and Comparative Functional Analyses of a <i>Mycobacterium</i> Aldo-Keto Reductase. <i>Journal of Molecular Biology</i> , 2010, 398, 26-39.	2.0	12
68	First insight into CD59-like molecules of adult <i>Fasciola hepatica</i> . <i>Experimental Parasitology</i> , 2014, 144, 57-64.	0.5	12
69	Assessment of a rtPCR for the detection of virulent and benign <i>Dichelobacter nodosus</i> , the causative agent of ovine footrot, in Australia. <i>BMC Veterinary Research</i> , 2018, 14, 252.	0.7	12
70	Towards understanding the liver fluke transmission dynamics on farms: Detection of liver fluke transmitting snail and liver fluke-specific environmental DNA in water samples from an irrigated dairy farm in Southeast Australia. <i>Veterinary Parasitology</i> , 2021, 291, 109373.	0.7	12
71	Efficient production of recombinant IL-21 proteins for pre-clinical studies by a two-step dilution refolding method. <i>International Immunopharmacology</i> , 2013, 16, 376-381.	1.7	11
72	Multiple ecto-nucleoside triphosphate diphosphohydrolases facilitate intracellular replication of <i>Legionella pneumophila</i> . <i>Biochemical Journal</i> , 2014, 462, 279-289.	1.7	11

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73	Current Status for Controlling the Overlooked Caprine Fasciolosis. <i>Animals</i> , 2021, 11, 1819.	1.0	11
74	Evaluation of the Role of Galectins in Parasite Immunity. <i>Methods in Molecular Biology</i> , 2015, 1207, 371-395.	0.4	11
75	Cloning, expression, purification and crystallographic studies of galectin-11 from domestic sheep (<i>Ovis aries</i>). <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2015, 71, 993-997.	0.4	10
76	Thermotolerance and molecular chaperone function of an SGT1-like protein from the psychrophilic yeast, <i>Glaciozyma antarctica</i> . <i>Cell Stress and Chaperones</i> , 2016, 21, 707-715.	1.2	10
77	Direct serogrouping of <i>Dichelobacter nodosus</i> from Victorian farms using conventional multiplex polymerase chain reaction. <i>BMC Research Notes</i> , 2018, 11, 108.	0.6	10
78	Optimization of a Loop Mediated Isothermal Amplification (LAMP) Assay for In-Field Detection of <i>Dichelobacter nodosus</i> With aprV2 (VDN LAMP) in Victorian Sheep Flocks. <i>Frontiers in Veterinary Science</i> , 2019, 6, 67.	0.9	10
79	The mitochondrial protein targeting suppressor (<i>mts1</i>) mutation maps to the mRNA-binding domain of Npl3p and affects translation on cytoplasmic polysomes. <i>Molecular Microbiology</i> , 2002, 35, 1277-1285.	1.2	9
80	Tracking the Unfolding Pathway of a Multirepeat Protein via Tryptophan Scanning. <i>Journal of Biological Chemistry</i> , 2006, 281, 24345-24350.	1.6	9
81	Molecular characterisation and vaccine efficacy of two novel developmentally regulated surface tegument proteins of <i>Fasciola hepatica</i> . <i>Veterinary Parasitology</i> , 2020, 286, 109244.	0.7	9
82	The 2.0 Å... Crystal Structure of a Pociilloporin at pH 3.5: The Structural Basis for the Linkage Between Color Transition and Halide Binding. <i>Photochemistry and Photobiology</i> , 2006, 82, 359.	1.3	8
83	Isothermal Nucleic Acid Amplification Technologies for the Detection of Equine Viral Pathogens. <i>Animals</i> , 2021, 11, 2150.	1.0	8
84	Crystal structure of fuculose aldolase from the Antarctic psychrophilic yeast <i>Glaciozyma antarctica</i> PI12. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2016, 72, 831-839.	0.4	7
85	Proteomic identification of galectin-11 and -14 ligands from <i>Fasciola hepatica</i> . <i>International Journal for Parasitology</i> , 2019, 49, 921-932.	1.3	7
86	Proteomic identification of galectin-11 and 14 ligands from <i>Haemonchus contortus</i> . <i>PeerJ</i> , 2018, 6, e4510.	0.9	7
87	Evaluation of Immunogenicity and Efficacy of <i>Fasciola hepatica</i> Tetraspanin 2 (TSP2) Fused to <i>E. coli</i> Heat-Labile Enterotoxin B Subunit LTB Adjuvant Following Intranasal Vaccination of Cattle. <i>Vaccines</i> , 2021, 9, 1213.	2.1	7
88	OZITX, a pertussis toxin-like protein for occluding inhibitory G protein signalling including G α _z . <i>Communications Biology</i> , 2022, 5, 256.	2.0	7
89	Expression, purification, crystallization and preliminary X-ray characterization of a putative glycosyltransferase of the GT-A fold found in mycobacteria. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2008, 64, 428-431.	0.7	6
90	Cloning, expression, purification and preliminary X-ray diffraction studies of a novel AB5toxin. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013, 69, 912-915.	0.7	6

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91	Vacuolation Activity and Intracellular Trafficking of ArtB, the Binding Subunit of an AB5 Toxin Produced by <i>Salmonella enterica</i> Serovar Typhi. <i>Infection and Immunity</i> , 2017, 85, .	1.0	6
92	Analysis of daily variation in the release of faecal eggs and coproantigen of <i>Fasciola hepatica</i> in naturally infected dairy cattle and the impact on diagnostic test sensitivity. <i>Veterinary Parasitology</i> , 2021, 298, 109504.	0.7	6
93	The Role of Anti-Viral Effector Molecules in Mollusc Hemolymph. <i>Biomolecules</i> , 2022, 12, 345.	1.8	6
94	Sialoglycan-binding patterns of bacterial AB5 toxin B subunits correlate with host range and toxicity, indicating evolution independent of A subunits. <i>Journal of Biological Chemistry</i> , 2022, 298, 101900.	1.6	6
95	The protein encoded by the MFT1 gene is a targeting factor for mitochondrial precursor proteins, and not a core ribosomal protein. <i>FEBS Letters</i> , 1997, 407, 220-224.	1.3	5
96	The Consequences of Stigma for Knowledge Production: Sheep Producers' Attitudes to Footrot Diagnostics and Control in Australia. <i>Frontiers in Veterinary Science</i> , 2020, 7, 354.	0.9	5
97	Crystal structures of pertussis toxin with NAD ⁺ and analogs provide structural insights into the mechanism of its cytosolic ADP-ribosylation activity. <i>Journal of Biological Chemistry</i> , 2022, 298, 101892.	1.6	5
98	Expression, purification, crystallization and preliminary X-ray diffraction analysis of an essential lipoprotein implicated in cell-wall biosynthesis in <i>Mycobacteria</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2005, 61, 1081-1083.	0.7	4
99	The oligomeric assembly of galectin-11 is critical for anti-parasitic activity in sheep (<i>Ovis aries</i>). <i>Communications Biology</i> , 2020, 3, 464.	2.0	4
100	A structural characterization of the isoniazid <i>Mycobacterium tuberculosis</i> drug target, Rv2971, in its unliganded form. <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2014, 70, 572-577.	0.4	3
101	A structural and functional investigation of a novel protein from <i>Mycobacterium smegmatis</i> implicated in mycobacterial macrophage survivability. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2014, 70, 2264-2276.	2.5	3
102	Evaluation of loop-mediated isothermal amplification (LAMP) assay for detection of aprV2 positive <i>Dichelobacter nodosus</i> in-field by secondary users. <i>BMC Research Notes</i> , 2019, 12, 534.	0.6	3
103	Current Status of Loop-Mediated Isothermal Amplification Technologies for the Detection of Honey Bee Pathogens. <i>Frontiers in Veterinary Science</i> , 2021, 8, 659683.	0.9	3
104	<i>Fasciola hepatica</i> Control Practices on a Sample of Dairy Farms in Victoria, Australia. <i>Frontiers in Veterinary Science</i> , 2021, 8, 669117.	0.9	3
105	Bovine Natural Antibody Relationships to Specific Antibodies and <i>Fasciola hepatica</i> Burdens after Experimental Infection and Vaccination with Glutathione S-Transferase. <i>Veterinary Sciences</i> , 2022, 9, 58.	0.6	3
106	Cloning, expression and crystallisation of SGT1 co-chaperone protein from <i>Glaciozyma antarctica</i> . <i>AIP Conference Proceedings</i> , 2013, .	0.3	2
107	Teladorsagia Circumcincta Galectin-Mucosal Interactome in Sheep. <i>Veterinary Sciences</i> , 2021, 8, 216.	0.6	2
108	Cloning, expression, purification and preliminary X-ray diffraction studies of a mycobacterial protein implicated in bacterial survival in macrophages. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013, 69, 566-569.	0.7	1

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109	Evaluation of the Role of Galectins in Parasite Immunity. <i>Methods in Molecular Biology</i> , 2022, 2442, 475-515.	0.4	1
110	Development of molecular detection methods of <i>Bovicola ovis</i> from sheep fleece. <i>Parasitology Research</i> , 2022, 121, 1597.	0.6	1
111	Ribosome binding to the surface of mitochondria: mechanisms and consequences. <i>Biochemical Society Transactions</i> , 2000, 28, A457-A457.	1.6	0
112	How opposites attract. <i>Immunology and Cell Biology</i> , 2011, 89, 163-164.	1.0	0
113	Air sampling for detection of infectious laryngotracheitis (ILT) in commercial poultry flocks. <i>BMC Research Notes</i> , 2020, 13, 556.	0.6	0
114	CD1d lipid-antigen recognition by the α TCR. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2014, 70, C244-C244.	0.0	0
115	Rapid Evolution of Bacterial Exotoxin B Subunits Independent of A subunits: Sialic Acid Binding Preferences Correlate with Host Range and Intrinsic Toxicity. <i>FASEB Journal</i> , 2018, 32, 673.3.	0.2	0
116	The Biomolecules Journal Club: Highlights on Recent Papersâ€™1. <i>Biomolecules</i> , 2022, 12, 86.	1.8	0