List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A Protein Corona Modulates Interactions of $\hat{I}\pm$ -Synuclein with Nanoparticles and Alters the Rates of the Microscopic Steps of Amyloid Formation. ACS Nano, 2022, 16, 1102-1118.	14.6	9
2	The low-density lipoprotein receptor-related protein 1 (LRP1) interactome in the human cornea. Experimental Eye Research, 2022, 219, 109081.	2.6	5
3	Structural Remodelling of the Carbonâ€Phosphorus Enzymatic Machinery by a Dual ATPâ€Binding Cassette Module. FASEB Journal, 2022, 36, .	0.5	0
4	Cryo-EM structures of human A2ML1 elucidate the protease-inhibitory mechanism of the A2M family. Nature Communications, 2022, 13, .	12.8	4
5	The conformational change of the protease inhibitor α2-macroglobulin is triggered by the retraction of the cleaved bait region from a central channel. Journal of Biological Chemistry, 2022, 298, 102230.	3.4	1
6	Latency, thermal stability, and identification of an inhibitory compound of mirolysin, a secretory protease of the human periodontopathogen Tannerella forsythia. Journal of Enzyme Inhibition and Medicinal Chemistry, 2021, 36, 1267-1281.	5.2	3
7	ITIH4 acts as a protease inhibitor by a novel inhibitory mechanism. Science Advances, 2021, 7, .	10.3	22
8	PorZ, an Essential Component of the Type IX Secretion System of <i>Porphyromonas gingivalis</i> , Delivers Anionic Lipopolysaccharide to the PorU Sortase for Transpeptidase Processing of T9SS Cargo Proteins. MBio, 2021, 12, .	4.1	17
9	Superoxide dismutase 3 is expressed in bone tissue and required for normal bone homeostasis and mineralization. Free Radical Biology and Medicine, 2021, 164, 399-409.	2.9	8
10	An Integrative Structural Biology Analysis of Von Willebrand Factor Binding and Processing by ADAMTS-13 in Solution. Journal of Molecular Biology, 2021, 433, 166954.	4.2	3
11	The last meal of Tollund Man: new analyses of his gut content. Antiquity, 2021, 95, 1195-1212.	1.0	10
12	Development of selective protease inhibitors via engineering of the bait region of human α2-macroglobulin. Journal of Biological Chemistry, 2021, 297, 100879.	3.4	9
13	Mutation-induced dimerization of transforming growth factor-β–induced protein may drive protein aggregation in granular corneal dystrophy. Journal of Biological Chemistry, 2021, 297, 100858.	3.4	3
14	Identification of polyphenol oxidases in potato tuber (Solanum tuberosum) and purification and characterization of the major polyphenol oxidases. Food Chemistry, 2021, 365, 130454.	8.2	6
15	Structural Investigations of Human A2M Identify a Hollow Native Conformation That Underlies Its Distinctive Protease-Trapping Mechanism. Molecular and Cellular Proteomics, 2021, 20, 100090.	3.8	21
16	A novel approach for production of an active N-terminally truncated Ulp1 (SUMO protease 1) catalytic domain from Escherichia coli inclusion bodies. Protein Expression and Purification, 2020, 166, 105507.	1.3	10
17	Unfolding and partial refolding of a cellulase from the SDS-denatured state: From β-sheet to α-helix and back. Biochimica Et Biophysica Acta - General Subjects, 2020, 1864, 129434.	2.4	18
18	The interactome of stabilized αâ€synuclein oligomers and neuronal proteins. FEBS Journal, 2020, 287, 2037-2054.	4.7	9

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19	Plasmin inhibition by bacterial serpin: Implications in gum disease. FASEB Journal, 2020, 34, 619-630.	0.5	12
20	Transport of a Peptide from Bovine αs1-Casein across Models of the Intestinal and Blood–Brain Barriers. Nutrients, 2020, 12, 3157.	4.1	8
21	α2-Macroglobulin-like protein 1 can conjugate and inhibit proteases through their hydroxyl groups, because of an enhanced reactivity of its thiol ester. Journal of Biological Chemistry, 2020, 295, 16732-16742.	3.4	6
22	STEEP mediates STING ER exit and activation of signaling. Nature Immunology, 2020, 21, 868-879.	14.5	82
23	Tracing the <i>In Vivo</i> Fate of Nanoparticles with a "Non-Self―Biological Identity. ACS Nano, 2020, 14, 10666-10679.	14.6	12
24	Biochemical mechanisms of aggregation in TGFBI-linked corneal dystrophies. Progress in Retinal and Eye Research, 2020, 77, 100843.	15.5	48
25	Protein Composition of the Subretinal Fluid Suggests Selective Diffusion of Vitreous Proteins in Retinal Detachment. Translational Vision Science and Technology, 2020, 9, 16.	2.2	9
26	Modulation of Small RNA Signatures in Schwann-Cell-Derived Extracellular Vesicles by the p75 Neurotrophin Receptor and Sortilin. Biomedicines, 2020, 8, 450.	3.2	14
27	Mapping and identification of soft corona proteins at nanoparticles and their impact on cellular association. Nature Communications, 2020, 11, 4535.	12.8	122
28	Structural and functional insights into oligopeptide acquisition by the RagAB transporter from Porphyromonas gingivalis. Nature Microbiology, 2020, 5, 1016-1025.	13.3	46
29	Inter-α-inhibitor heavy chain-1 has an integrin-like 3D structure mediating immune regulatory activities and matrix stabilization during ovulation. Journal of Biological Chemistry, 2020, 295, 5278-5291.	3.4	18
30	Kallikrein-Related Peptidase 14 Activates Zymogens of Membrane Type Matrix Metalloproteinases (MT-MMPs)—A CleavEx Based Analysis. International Journal of Molecular Sciences, 2020, 21, 4383.	4.1	5
31	Protein Analysis of the TGFBI ^{R124H} Mouse Model Gives Insight into Phenotype Development of Granular Corneal Dystrophy. Proteomics - Clinical Applications, 2020, 14, e1900072.	1.6	2
32	FAM20C phosphorylation of the RGDSVVYGLR motif in osteopontin inhibits interaction with the αvβ3 integrin. Journal of Cellular Biochemistry, 2020, 121, 4809-4818.	2.6	12
33	Apolipoprotein E Triggers Complement Activation in Joint Synovial Fluid of Rheumatoid Arthritis Patients by Binding C1q. Journal of Immunology, 2020, 204, 2779-2790.	0.8	20
34	Substituting the Thiol Ester of Human A2M or C3 with a Disulfide Produces Native Proteins with Altered Proteolysis-Induced Conformational Changes. Biochemistry, 2020, 59, 4799-4809.	2.5	6
35	Matrix-degrading protease ADAMTS-5 cleaves inter-α-inhibitor and releases active heavy chain 2 in synovial fluids from arthritic patients. Journal of Biological Chemistry, 2019, 294, 15495-15504.	3.4	10
36	Proteolytic processing and activation of gingipain zymogens secreted by T9SS of Porphyromonas gingivalis. Biochimie, 2019, 166, 161-172.	2.6	14

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37	Sortilin gates neurotensin and BDNF signaling to control peripheral neuropathic pain. Science Advances, 2019, 5, eaav9946.	10.3	35
38	The serine protease HtrA1 cleaves misfolded transforming growth factor β–induced protein (TGFBlp) and induces amyloid formation. Journal of Biological Chemistry, 2019, 294, 11817-11828.	3.4	11
39	Staphylococcus saccharolyticus Isolated From Blood Cultures and Prosthetic Joint Infections Exhibits Excessive Genome Decay. Frontiers in Microbiology, 2019, 10, 478.	3.5	12
40	Conservation of the Amyloid Interactome Across Diverse Fibrillar Structures. Scientific Reports, 2019, 9, 3863.	3.3	13
41	Imperfect repeats in the functional amyloid protein FapC reduce the tendency to fragment during fibrillation. Protein Science, 2019, 28, 633-642.	7.6	36
42	Frequently used bioinformatics tools overestimate the damaging effect of allelic variants. Genes and Immunity, 2019, 20, 10-22.	4.1	12
43	A Novel Biological Role for Peptidyl-Arginine Deiminases: Citrullination of Cathelicidin LL-37 Controls the Immunostimulatory Potential of Cell-Free DNA. Journal of Immunology, 2018, 200, 2327-2340.	0.8	27
44	Proteomic profiling of <i><scp>TGFBI</scp></i> â€null mouse corneas reveals only minor changes in matrix composition supportive of <i><scp>TGFBI</scp></i> knockdown as therapy against <i><scp>TGFBI</scp></i> â€linked corneal dystrophies. FEBS Journal, 2018, 285, 101-114.	4.7	24
45	A Screening Method for the Isolation of Bacteria Capable of Degrading Toxic Steroidal Glycoalkaloids Present in Potato. Frontiers in Microbiology, 2018, 9, 2648.	3.5	21
46	α-Synucleins from Animal Species Show Low Fibrillation Propensities and Weak Oligomer Membrane Disruption. Biochemistry, 2018, 57, 5145-5158.	2.5	15
47	Small-Molecule Probes for Affinity-Guided Introduction of Biocompatible Handles on Metal-Binding Proteins. Bioconjugate Chemistry, 2018, 29, 3016-3025.	3.6	16
48	Serum Amyloid P Component (SAP) Interactome in Human Plasma Containing Physiological Calcium Levels. Biochemistry, 2017, 56, 896-902.	2.5	14
49	Female versus male biological identities of nanoparticles determine the interaction with immune cells in fish. Environmental Science: Nano, 2017, 4, 895-906.	4.3	31
50	Reactive Center Loop Insertion in $\hat{I}\pm -1$ -Antitrypsin Captured by Accelerated Molecular Dynamics Simulation. Biochemistry, 2017, 56, 634-646.	2.5	20
51	Mirolysin, a LysargiNase from <i>Tannerella forsythia</i> , proteolytically inactivates the human cathelicidin, LL-37. Biological Chemistry, 2017, 398, 395-409.	2.5	18
52	Human Lysozyme Peptidase Resistance Is Perturbed by the Anionic Glycolipid Biosurfactant Rhamnolipid Produced by the Opportunistic PathogenPseudomonas aeruginosa. Biochemistry, 2017, 56, 260-270.	2.5	6
53	Structural and Functional Implications of Human Transforming Growth Factor Î ² -Induced Protein, TGFBIp, in Corneal Dystrophies. Structure, 2017, 25, 1740-1750.e2.	3.3	24
54	Transcriptome analysis of the response of Burmese python to digestion. GigaScience, 2017, 6, 1-18.	6.4	17

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55	Characterisation of protein families in spider digestive fluids and their role in extra-oral digestion. BMC Genomics, 2017, 18, 600.	2.8	39
56	Activation of Complement by Pigment Epithelium–Derived Factor in Rheumatoid Arthritis. Journal of Immunology, 2017, 199, 1113-1121.	0.8	4
57	Analysis of Factor D Isoforms in Malpuech–Michels–Mingarelli–Carnevale Patients Highlights the Role of MASP-3 as a Maturase in the Alternative Pathway of Complement. Journal of Immunology, 2017, 199, 2158-2170.	0.8	43
58	Critical Influence of Cosolutes and Surfaces on the Assembly of Serpin-Derived Amyloid Fibrils. Biophysical Journal, 2017, 113, 580-596.	0.5	20
59	Mutation-Induced Deamidation of Corneal Dystrophy-Related Transforming Growth Factor β-Induced Protein. Biochemistry, 2017, 56, 6470-6480.	2.5	4
60	"Ant-egg―cataract revisited. Graefe's Archive for Clinical and Experimental Ophthalmology, 2017, 255, 119-125.	1.9	1
61	An Aberrant Phosphorylation of Amyloid Precursor Protein Tyrosine Regulates Its Trafficking and the Binding to the Clathrin Endocytic Complex in Neural Stem Cells of Alzheimer's Disease Patients. Frontiers in Molecular Neuroscience, 2017, 10, 59.	2.9	28
62	Extracellular superoxide dismutase is present in secretory vesicles of human neutrophils and released upon stimulation. Free Radical Biology and Medicine, 2016, 97, 478-488.	2.9	29
63	Detection of proteolytic signatures for Parkinson's disease. Future Neurology, 2016, 11, 15-32.	0.5	0
64	LASIK surgery of granular corneal dystrophy type 2 patients leads to accumulation and differential proteolytic processing of transforming growth factor betaâ€induced protein (TGFBIp). Proteomics, 2016, 16, 539-543.	2.2	20
65	Combinatorial Biomolecular Nanopatterning for Highâ€Throughput Screening of Stemâ€Cell Behavior. Advanced Materials, 2016, 28, 1472-1476.	21.0	17
66	The outer-membrane export signal of Porphyromonas gingivalis type IX secretion system (T9SS) is a conserved C-terminal β-sandwich domain. Scientific Reports, 2016, 6, 23123.	3.3	52
67	Genomic and exoproteomic analyses of cold―and alkalineâ€adapted bacteria reveal an abundance of secreted subtilisinâ€like proteases. Microbial Biotechnology, 2016, 9, 245-256.	4.2	9
68	Disulfide Bond Pattern of Transforming Growth Factor Î ² -Induced Protein. Biochemistry, 2016, 55, 5610-5621.	2.5	10
69	How Glycosaminoglycans Promote Fibrillation of Salmon Calcitonin. Journal of Biological Chemistry, 2016, 291, 16849-16862.	3.4	15
70	Dynamic protein coronas revealed as a modulator of silver nanoparticle sulphidation in vitro. Nature Communications, 2016, 7, 11770.	12.8	136
71	Structural and functional probing of PorZ, an essential bacterial surface component of the type-IX secretion system of human oral-microbiomic Porphyromonas gingivalis Scientific Reports, 2016, 6, 37708.	3.3	58
72	Keratin 12 missense mutation induces the unfolded protein response and apoptosis in Meesmann epithelial corneal dystrophy. Human Molecular Genetics, 2016, 25, 1176-1191.	2.9	22

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73	Enzymatic and Structural Characterization of the Major Endopeptidase in the Venus Flytrap Digestion Fluid. Journal of Biological Chemistry, 2016, 291, 2271-2287.	3.4	16
74	Carbamylated LL-37 as a modulator of the immune response. Innate Immunity, 2016, 22, 218-229.	2.4	32
75	Antagonism between Staphylococcus epidermidis and Propionibacterium acnes and its genomic basis. BMC Genomics, 2016, 17, 152.	2.8	131
76	Transglutaminase 2-Catalyzed Intramolecular Cross-Linking of Osteopontin. Biochemistry, 2016, 55, 294-303.	2.5	14
77	The Compact and Biologically Relevant Structure of Inter-α-inhibitor Is Maintained by the Chondroitin Sulfate Chain and Divalent Cations. Journal of Biological Chemistry, 2016, 291, 4658-4670.	3.4	7
78	The spider hemolymph clot proteome reveals high concentrations of hemocyanin and von Willebrand factor-like proteins. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2016, 1864, 233-241.	2.3	24
79	Near-complete 1H, 13C, 15N resonance assignments of dimethylsulfoxide-denatured TGFBIp FAS1-4 A546T. Biomolecular NMR Assignments, 2016, 10, 25-29.	0.8	2
80	SILAC-MS Based Characterization of LPS and Resveratrol Induced Changes in Adipocyte Proteomics – Resveratrol as Ameliorating Factor on LPS Induced Changes. PLoS ONE, 2016, 11, e0159747.	2.5	17
81	Hepatocytes respond differently to major dietary trans fatty acid isomers, elaidic acid and trans-vaccenic acid. Proteome Science, 2015, 13, 31.	1.7	16
82	Protein Composition of TGFBI-R124C- and TGFBI-R555W- Associated Aggregates Suggests Multiple Mechanisms Leading to Lattice and Granular Corneal Dystrophy. , 2015, 56, 4653.		28
83	New Insights to Clathrin and Adaptor Protein 2 for the Design and Development of Therapeutic Strategies. International Journal of Molecular Sciences, 2015, 16, 29446-29453.	4.1	21
84	KLIKK proteases of Tannerella forsythia: putative virulence factors with a unique domain structure. Frontiers in Microbiology, 2015, 6, 312.	3.5	40
85	Mirolase, a novel subtilisin-like serine protease from the periodontopathogen Tannerella forsythia. Biological Chemistry, 2015, 396, 261-275.	2.5	29
86	Metal Ion-dependent Heavy Chain Transfer Activity of TSG-6 Mediates Assembly of the Cumulus-Oocyte Matrix. Journal of Biological Chemistry, 2015, 290, 28708-28723.	3.4	46
87	Characterization of the gila monster (Heloderma suspectum suspectum) venom proteome. Journal of Proteomics, 2015, 117, 1-11.	2.4	25
88	The effects of hypochlorous acid and neutrophil proteases on the structure and function of extracellular superoxide dismutase. Free Radical Biology and Medicine, 2015, 81, 38-46.	2.9	10
89	Miropin, a Novel Bacterial Serpin from the Periodontopathogen Tannerella forsythia, Inhibits a Broad Range of Proteases by Using Different Peptide Bonds within the Reactive Center Loop. Journal of Biological Chemistry, 2015, 290, 658-670.	3.4	42
90	The amido-pentadienoate-functionality of the rakicidins is a thiol reactive electrophile – development of a general synthetic strategy. Chemical Communications, 2015, 51, 12427-12430.	4.1	22

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91	ADAM10 controls collagen signaling and cell migration on collagen by shedding the ectodomain of discoidin domain receptor 1 (DDR1). Molecular Biology of the Cell, 2015, 26, 659-673.	2.1	41
92	Optimized co-solute paramagnetic relaxation enhancement for the rapid NMR analysis of a highly fibrillogenic peptide. Journal of Biomolecular NMR, 2015, 62, 129-142.	2.8	26
93	Fibril Core of Transforming Growth Factor Beta-Induced Protein (TGFBIp) Facilitates Aggregation of Corneal TGFBIp. Biochemistry, 2015, 54, 2943-2956.	2.5	19
94	Calcium Regulates the Activity and Structural Stability of Tpr, a Bacterial Calpain-like Peptidase. Journal of Biological Chemistry, 2015, 290, 27248-27260.	3.4	11
95	Early Events in the Amyloid Formation of the A546T Mutant of Transforming Growth Factor β-Induced Protein in Corneal Dystrophies Compared to the Nonfibrillating R555W and R555Q Mutants. Biochemistry, 2015, 54, 5546-5556.	2.5	6
96	Characterization of the gila monster (Heloderma suspectum suspectum) venom proteome. Data in Brief, 2015, 3, 137-142.	1.0	12
97	Clearance Kinetics and Matrix Binding Partners of the Receptor for Advanced Glycation End Products. PLoS ONE, 2014, 9, e88259.	2.5	16
98	Insight into the Protein Composition of Immunoglobulin Light Chain Deposits of Eyelid, Orbital and Conjunctival Amyloidosis. Journal of Proteomics and Bioinformatics, 2014, s8, .	0.4	7
99	Comparison of two phenotypically distinct lattice corneal dystrophies caused by mutations in the transforming growth factor beta induced (<i>TGFBI</i>) gene. Proteomics - Clinical Applications, 2014, 8, 168-177.	1.6	24
100	Proteomics and the Eye. Proteomics - Clinical Applications, 2014, 8, 127-129.	1.6	5
101	Peptidyl Arginine Deiminase from Porphyromonas gingivalis Abolishes Anaphylatoxin C5a Activity. Journal of Biological Chemistry, 2014, 289, 32481-32487.	3.4	83
102	Proteome reference maps of the <i>Lotus japonicus</i> nodule and root. Proteomics, 2014, 14, 230-240.	2.2	21
103	Spider genomes provide insight into composition and evolution of venom and silk. Nature Communications, 2014, 5, 3765.	12.8	235
104	Incorporation of Pentraxin 3 into Hyaluronan Matrices Is Tightly Regulated and Promotes Matrix Cross-linking. Journal of Biological Chemistry, 2014, 289, 30481-30498.	3.4	67
105	The Autolysis of Human HtrA1 Is Governed by the Redox State of Its N-Terminal Domain. Biochemistry, 2014, 53, 3851-3857.	2.5	19
106	Proteomics of Fuchs' Endothelial Corneal Dystrophy Support That the Extracellular Matrix of Descemet's Membrane Is Disordered. Journal of Proteome Research, 2014, 13, 4659-4667.	3.7	36
107	Template-directed covalent conjugation of DNA to native antibodies, transferrin and other metal-binding proteins. Nature Chemistry, 2014, 6, 804-809.	13.6	152
108	Carbamylation of immunoglobulin abrogates activation of the classical complement pathway. European Journal of Immunology, 2014, 44, 3403-3412.	2.9	23

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109	A Common Polymorphism in Extracellular Superoxide Dismutase Affects Cardiopulmonary Disease Risk by Altering Protein Distribution. Circulation: Cardiovascular Genetics, 2014, 7, 659-666.	5.1	31
110	Unconditioned commercial embryo culture media contain a large variety of non-declared proteins: a comprehensive proteomics analysis. Human Reproduction, 2014, 29, 2421-2430.	0.9	63
111	Coagulation Factor XIIIa Substrates in Human Plasma. Journal of Biological Chemistry, 2014, 289, 6526-6534.	3.4	55
112	The Role of Stable α-Synuclein Oligomers in the Molecular Events Underlying Amyloid Formation. Journal of the American Chemical Society, 2014, 136, 3859-3868.	13.7	218
113	Preparation of uniformly 13C,15N-labeled recombinant human amylin for solid-state NMR investigation. Protein Expression and Purification, 2014, 99, 119-130.	1.3	7
114	Secreted major Venus flytrap chitinase enables digestion of Arthropod prey. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 374-383.	2.3	33
115	Distal Renal Tubules Are Deficient in Aggresome Formation and Autophagy upon Aldosterone Administration. PLoS ONE, 2014, 9, e101258.	2.5	8
116	Proteome Analysis of Human Sebaceous Follicle Infundibula Extracted from Healthy and Acne-Affected Skin. PLoS ONE, 2014, 9, e107908.	2.5	50
117	Identification of Transglutaminase Reactive Residues in Human Osteopontin and Their Role in Polymerization. PLoS ONE, 2014, 9, e113650.	2.5	14
118	The Human Eye Proteome Project: Perspectives on an emerging proteome. Proteomics, 2013, 13, 2500-2511.	2.2	75
119	Species Differences Take Shape at Nanoparticles: Protein Corona Made of the Native Repertoire Assists Cellular Interaction. Environmental Science & Technology, 2013, 47, 14367-14375.	10.0	75
120	Mutation in transforming growth factor beta induced protein associated with granular corneal dystrophy type 1 reduces the proteolytic susceptibility through local structural stabilization. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 2812-2822.	2.3	33
121	Monodisperse and LPS-free Aggregatibacter actinomycetemcomitans leukotoxin: Interactions with human β2 integrins and erythrocytes. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 546-558.	2.3	30
122	Inhibition of gingipains by their profragments as the mechanism protecting Porphyromonas gingivalis against premature activation of secreted proteases. Biochimica Et Biophysica Acta - General Subjects, 2013, 1830, 4218-4228.	2.4	21
123	Hydrogen peroxide induce modifications of human extracellular superoxide dismutase that results in enzyme inhibition. Redox Biology, 2013, 1, 24-31.	9.0	80
124	Off-pathway aggregation can inhibit fibrillation at high protein concentrations. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 677-687.	2.3	12
125	Murine Extracellular Superoxide Dismutase Is Converted into the Inactive Fold by the Ser195Cys Mutation. Biochemistry, 2013, 52, 3369-3375.	2.5	3
126	The Insoluble TGFBIp Fraction of the Cornea Is Covalently Linked via a Disulfide Bond to Type XII Collagen. Biochemistry, 2013, 52, 2821-2827.	2.5	21

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127	Inactivation of Epidermal Growth Factor by Porphyromonas gingivalis as a Potential Mechanism for Periodontal Tissue Damage. Infection and Immunity, 2013, 81, 55-64.	2.2	46
128	Investigations on Collectin Liver 1. Journal of Biological Chemistry, 2013, 288, 23407-23420.	3.4	69
129	Differential Regulation of Extracellular Tissue Inhibitor of Metalloproteinases-3 Levels by Cell Membrane-bound and Shed Low Density Lipoprotein Receptor-related Protein 1. Journal of Biological Chemistry, 2013, 288, 332-342.	3.4	64
130	Inter-α-inhibitor Impairs TSG-6-induced Hyaluronan Cross-linking. Journal of Biological Chemistry, 2013, 288, 29642-29653.	3.4	60
131	Aldosterone and angiotensin II induce protein aggregation in renal proximal tubules. Physiological Reports, 2013, 1, e00064.	1.7	11
132	Effects of Elaidic Acid on Lipid Metabolism in HepG2 Cells, Investigated by an Integrated Approach of Lipidomics, Transcriptomics and Proteomics. PLoS ONE, 2013, 8, e74283.	2.5	35
133	Ribosomal Protein L22 (RPL22) accumulates as aggregates in distal renal tubules after aldosterone administration. FASEB Journal, 2013, 27, 705.5.	0.5	0
134	Serine protease HtrA1 accumulates in corneal transforming growth factor beta induced protein (TGFBIp) amyloid deposits. Molecular Vision, 2013, 19, 861-76.	1.1	26
135	Vesicular signalling and immune modulation as hedonic fingerprints: proteomic profiling in the chronic mild stress depression model. Journal of Psychopharmacology, 2012, 26, 1569-1583.	4.0	24
136	Polymorphic Fibrillation of the Destabilized Fourth Fasciclin-1 Domain Mutant A546T of the Transforming Growth Factor-β-induced Protein (TGFBIp) Occurs through Multiple Pathways with Different Oligomeric Intermediates. Journal of Biological Chemistry, 2012, 287, 34730-34742.	3.4	21
137	Structural insights into triple-helical collagen cleavage by matrix metalloproteinase 1. Proceedings of the United States of America, 2012, 109, 12461-12466.	7.1	185
138	Disruption of gingipain oligomerization into non-covalent cell-surface attached complexes. Biological Chemistry, 2012, 393, 971-977.	2.5	15
139	Unique Structural Features Facilitate Lizard Tail Autotomy. PLoS ONE, 2012, 7, e51803.	2.5	37
140	Human Complement C3 Is a Substrate for Transglutaminases. A Functional Link between Non-Protease-Based Members of the Coagulation and Complement Cascades. Biochemistry, 2012, 51, 4735-4742.	2.5	24
141	The Neuroendocrine Protein 7B2 Is Intrinsically Disordered. Biochemistry, 2012, 51, 7456-7464.	2.5	8
142	Synthesis and Evaluation of Silanediols as Highly Selective Uncompetitive Inhibitors of Human Neutrophil Elastase. Journal of Medicinal Chemistry, 2012, 55, 7900-7908.	6.4	29
143	Human Cornea Proteome: Identification and Quantitation of the Proteins of the Three Main Layers Including Epithelium, Stroma, and Endothelium. Journal of Proteome Research, 2012, 11, 4231-4239.	3.7	92
144	Composition and proteolytic processing of corneal deposits associated with mutations in the TGFBI gene. Experimental Eye Research, 2012, 96, 163-170.	2.6	50

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145	<scp>MS D</scp> ata <scp>M</scp> iner: A webâ€based software tool to analyze, compare, and share mass spectrometry protein identifications. Proteomics, 2012, 12, 2792-2796.	2.2	45
146	The Protein Composition of the Digestive Fluid from the Venus Flytrap Sheds Light on Prey Digestion Mechanisms. Molecular and Cellular Proteomics, 2012, 11, 1306-1319.	3.8	83
147	The C-terminal proteolytic processing of extracellular superoxide dismutase is redox regulated. Free Radical Biology and Medicine, 2012, 52, 191-197.	2.9	15
148	Identification of a potential biomarker panel for the intake of the common dietary trans fat elaidic acid (transâ^†9-C18:1). Journal of Proteomics, 2012, 75, 2685-2696.	2.4	7
149	Human inter-α-inhibitor is a substrate for factor XIIIa and tissue transglutaminase. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 1624-1630.	2.3	10
150	Lack of the Receptor for Advanced Glycation End-Products Attenuates E. coli Pneumonia in Mice. PLoS ONE, 2011, 6, e20132.	2.5	47
151	The role of higher-order protein structure in supporting binding by heteroclitic monoclonal antibodies: The monoclonal antibody KIM185 to CD18 also binds C4-binding protein. Molecular Immunology, 2011, 49, 38-47.	2.2	4
152	Hydrogen exchange mass spectrometry as an analytical tool for the analysis of amyloid fibrillogenesis. International Journal of Mass Spectrometry, 2011, 302, 167-173.	1.5	6
153	Human Phenotypically Distinct TGFBI Corneal Dystrophies Are Linked to the Stability of the Fourth FAS1 Domain of TGFBIp. Journal of Biological Chemistry, 2011, 286, 4951-4958.	3.4	55
154	Inhibition of Staphylococcus aureus cysteine proteases by human serpin potentially limits staphylococcal virulence. Biological Chemistry, 2011, 392, 483-9.	2.5	27
155	Reactive-site mutants of N-TIMP-3 that selectively inhibit ADAMTS-4 and ADAMTS-5: biological and structural implications. Biochemical Journal, 2010, 431, 113-122.	3.7	59
156	The concentration of extracellular superoxide dismutase in plasma is maintained by LRP-mediated endocytosis. Free Radical Biology and Medicine, 2010, 49, 894-899.	2.9	18
157	Divorcing folding from function: How acylation affects the membrane-perturbing properties of an antimicrobial peptide. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2010, 1804, 806-820.	2.3	21
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