

# Nicole M Thielens

## List of Publications by Year in descending order

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167  
papers

6,456  
citations

44069

48  
h-index

79698

73  
g-index

182  
all docs

182  
docs citations

182  
times ranked

6226  
citing authors

#	ARTICLE	IF	CITATIONS
1	Structure and activation of the C1 complex of complement: unraveling the puzzle. <i>Trends in Immunology</i> , 2004, 25, 368-373.	6.8	223
2	C1q: A fresh look upon an old molecule. <i>Molecular Immunology</i> , 2017, 89, 73-83.	2.2	188
3	Synergy between Ficolin-2 and Pentraxin 3 Boosts Innate Immune Recognition and Complement Deposition. <i>Journal of Biological Chemistry</i> , 2009, 284, 28263-28275.	3.4	184
4	Structural insights into the innate immune recognition specificities of L- and H-ficolins. <i>EMBO Journal</i> , 2007, 26, 623-633.	7.8	170
5	Nanobodies Targeting Mouse/Human VCAM1 for the Nuclear Imaging of Atherosclerotic Lesions. <i>Circulation Research</i> , 2012, 110, 927-937.	4.5	167
6	Structural insights into the Slit-Robo complex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 14923-14928.	7.1	159
7	Substrate Specificities of Recombinant Mannan-binding Lectin-associated Serine Proteases-1 and -2. <i>Journal of Biological Chemistry</i> , 2001, 276, 40880-40887.	3.4	154
8	Levels of mannan-binding lectin-associated serine protease-2 in healthy individuals. <i>Journal of Immunological Methods</i> , 2003, 282, 159-167.	1.4	141
9	C1q restrains autoimmunity and viral infection by regulating CD8 <sup>+</sup> T cell metabolism. <i>Science</i> , 2018, 360, 558-563.	12.6	133
10	DC/L-SIGN recognition of spike glycoprotein promotes SARS-CoV-2 trans-infection and can be inhibited by a glycomimetic antagonist. <i>PLoS Pathogens</i> , 2021, 17, e1009576.	4.7	133
11	The Two Major Oligomeric Forms of Human Mannan-Binding Lectin: Chemical Characterization, Carbohydrate-Binding Properties, and Interaction with MBL-Associated Serine Proteases. <i>Journal of Immunology</i> , 2005, 174, 2870-2877.	0.8	128
12	Interaction Properties of Human Mannan-Binding Lectin (MBL)-Associated Serine Proteases-1 and -2, MBL-Associated Protein 19, and MBL. <i>Journal of Immunology</i> , 2001, 166, 5068-5077.	0.8	124
13	Heteromeric Complexes of Native Collectin Kidney 1 and Collectin Liver 1 Are Found in the Circulation with MASPs and Activate the Complement System. <i>Journal of Immunology</i> , 2013, 191, 6117-6127.	0.8	113
14	Carbohydrate Recognition Properties of Human Ficolins. <i>Journal of Biological Chemistry</i> , 2010, 285, 6612-6622.	3.4	106
15	Studies on the interactions between C-reactive protein and complement proteins. <i>Immunology</i> , 2007, 121, 40-50.	4.4	104
16	The crystal structure of the zymogen catalytic domain of complement protease C1r reveals that a disruptive mechanical stress is required to trigger activation of the C1 complex. <i>EMBO Journal</i> , 2002, 21, 231-239.	7.8	101
17	Interaction of C1q and Mannan-Binding Lectin with Viruses. <i>Immunobiology</i> , 2002, 205, 563-574.	1.9	100
18	Periodontal Ehlers-Danlos Syndrome Is Caused by Mutations in C1R and C1S, which Encode Subcomponents C1r and C1s of Complement. <i>American Journal of Human Genetics</i> , 2016, 99, 1005-1014.	6.2	100

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19	Structural Bases for the Affinity-Driven Selection of a Public TCR against a Dominant Human Cytomegalovirus Epitope. <i>Journal of Immunology</i> , 2009, 183, 430-437.	0.8	93
20	Identification of the C1q-binding Sites of Human C1r and C1s. <i>Journal of Biological Chemistry</i> , 2009, 284, 19340-19348.	3.4	84
21	X-ray Structure of the Ca <sup>2+</sup> -binding Interaction Domain of C1s. <i>Journal of Biological Chemistry</i> , 2003, 278, 32157-32164.	3.4	82
22	Characterization of Recombinant Mannan-Binding Lectin-Associated Serine Protease (MASP)-3 Suggests an Activation Mechanism Different from That of MASP-1 and MASP-2. <i>Journal of Immunology</i> , 2004, 172, 4342-4350.	0.8	79
23	Structural biology of the C1 complex of complement unveils the mechanisms of its activation and proteolytic activity. <i>Molecular Immunology</i> , 2002, 39, 383-394.	2.2	78
24	Residue Lys57 in the Collagen-Like Region of Human L-Ficolin and Its Counterpart Lys47 in H-Ficolin Play a Key Role in the Interaction with the Mannan-Binding Lectin-Associated Serine Proteases and the Collectin Receptor Calreticulin. <i>Journal of Immunology</i> , 2009, 182, 456-465.	0.8	77
25	Crystal Structure of the CUB1-EGF-CUB2 Domain of Human MASP-1/3 and Identification of Its Interaction Sites with Mannan-binding Lectin and Ficolins. <i>Journal of Biological Chemistry</i> , 2008, 283, 25715-25724.	3.4	75
26	Baculovirus-mediated Expression of Truncated Modular Fragments from the Catalytic Region of Human Complement Serine Protease C1s. <i>Journal of Biological Chemistry</i> , 1998, 273, 1232-1239.	3.4	73
27	Characterization of the Interaction Between L-Ficolin/P35 and Mannan-Binding Lectin-Associated Serine Proteases-1 and -2. <i>Journal of Immunology</i> , 2002, 169, 5735-5743.	0.8	72
28	Molecular organization of human Ficolin-2. <i>Molecular Immunology</i> , 2007, 44, 401-411.	2.2	72
29	M-Ficolin Interacts with the Long Pentraxin PTX3: A Novel Case of Cross-Talk between Soluble Pattern-Recognition Molecules. <i>Journal of Immunology</i> , 2011, 186, 5815-5822.	0.8	72
30	The Human C1q Globular Domain: Structure and Recognition of Non-Immune Self Ligands. <i>Frontiers in Immunology</i> , 2011, 2, 92.	4.8	72
31	Structural biology of C1: dissection of a complex molecular machinery. <i>Immunological Reviews</i> , 2001, 180, 136-145.	6.0	69
32	Procollagen C-proteinase enhancer grasps the stalk of the C-propeptide trimer to boost collagen precursor maturation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6394-6399.	7.1	66
33	The X-ray Structure of Human Mannan-binding Lectin-associated Protein 19 (MAp19) and Its Interaction Site with Mannan-binding Lectin and L-ficolin. <i>Journal of Biological Chemistry</i> , 2004, 279, 29391-29397.	3.4	65
34	Elucidation of the substrate specificity of the MASP-2 protease of the lectin complement pathway and identification of the enzyme as a major physiological target of the serpin, C1-inhibitor. <i>Molecular Immunology</i> , 2008, 45, 670-677.	2.2	64
35	The N-terminal CUB-Epidermal Growth Factor Module Pair of Human Complement Protease C1r Binds Ca <sup>2+</sup> with High Affinity and Mediates Ca <sup>2+</sup> -dependent Interaction with C1s. <i>Journal of Biological Chemistry</i> , 1999, 274, 9149-9159.	3.4	62
36	Monomeric Structures of the Zymogen and Active Catalytic Domain of Complement Protease C1r. <i>Structure</i> , 2002, 10, 1509-1519.	3.3	59

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37	Structural Basis for Innate Immune Sensing by M-ficolin and Its Control by a pH-dependent Conformational Switch. <i>Journal of Biological Chemistry</i> , 2007, 282, 35814-35820.	3.4	59
38	Modified low density lipoproteins differentially bind and activate the C1 complex of complement. <i>Molecular Immunology</i> , 2007, 44, 1169-1177.	2.2	57
39	Deciphering the Fine Details of C1 Assembly and Activation Mechanisms: "Mission Impossible". <i>Frontiers in Immunology</i> , 2014, 5, 565.	4.8	57
40	Characterization of the interaction between collectin 11 (CL-11, CL-K1) and nucleic acids. <i>Molecular Immunology</i> , 2013, 56, 757-767.	2.2	56
41	Identification of the Site of Human Mannan-Binding Lectin Involved in the Interaction with Its Partner Serine Proteases: The Essential Role of Lys55. <i>Journal of Immunology</i> , 2007, 178, 5710-5716.	0.8	55
42	Human astrovirus coat protein binds C1q and MBL and inhibits the classical and lectin pathways of complement activation. <i>Molecular Immunology</i> , 2010, 47, 792-798.	2.2	55
43	Structure and properties of the Ca <sup>2+</sup> -binding CUB domain, a widespread ligand-recognition unit involved in major biological functions. <i>Biochemical Journal</i> , 2011, 439, 185-193.	3.7	55
44	Oxidative Stress Sensitizes Retinal Pigmented Epithelial (RPE) Cells to Complement-mediated Injury in a Natural Antibody-, Lectin Pathway-, and Phospholipid Epitope-dependent Manner. <i>Journal of Biological Chemistry</i> , 2013, 288, 12753-12765.	3.4	55
45	Expression of recombinant human complement C1q allows identification of the C1r/C1s-binding sites. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 8650-8655.	7.1	55
46	<sc>M</sc> and <sc>B</sc> and <sc>M</sc>ur<sc>G</sc> as scaffolds for the cytoplasmic steps of peptidoglycan biosynthesis. <i>Environmental Microbiology</i> , 2013, 15, 3218-3228.	3.8	54
47	Cutting Edge: C1q Binds Deoxyribose and Heparan Sulfate through Neighboring Sites of Its Recognition Domain. <i>Journal of Immunology</i> , 2010, 185, 808-812.	0.8	52
48	Chemical and functional characterization of a fragment of C1s containing the epidermal growth factor homology region. <i>Biochemistry</i> , 1990, 29, 3570-3578.	2.5	51
49	Structure and Assembly of the Catalytic Region of Human Complement Protease C1 <sub>r</sub> : A Three-Dimensional Model Based on Chemical Cross-Linking and Homology Modeling. <i>Biochemistry</i> , 1997, 36, 6270-6282.	2.5	51
50	Human and Pneumococcal Cell Surface Glyceraldehyde-3-phosphate Dehydrogenase (GAPDH) Proteins Are Both Ligands of Human C1q Protein. <i>Journal of Biological Chemistry</i> , 2012, 287, 42620-42633.	3.4	51
51	Complement Protein C1q Recognizes a Conformationally Modified Form of the Prion Protein. <i>Biochemistry</i> , 2005, 44, 4349-4356.	2.5	49
52	Deciphering Complement Receptor Type 1 Interactions with Recognition Proteins of the Lectin Complement Pathway. <i>Journal of Immunology</i> , 2013, 190, 3721-3731.	0.8	49
53	The Atypical Serine Proteases of the Complement System**Received for publication on October 7, 1997. <i>Advances in Immunology</i> , 1998, , 249-307.	2.2	48
54	The Immunopathology of Complement Proteins and Innate Immunity in Autoimmune Disease. <i>Clinical Reviews in Allergy and Immunology</i> , 2020, 58, 229-251.	6.5	47

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55	Structural Insights into the Recognition Properties of Human Ficolins. <i>Journal of Innate Immunity</i> , 2010, 2, 17-23.	3.8	46
56	Direct interaction between CD91 and C1q. <i>FEBS Journal</i> , 2010, 277, 3526-3537.	4.7	45
57	<i>Trypanosoma cruzi</i> calreticulin inhibits the complement lectin pathway activation by direct interaction with L-Ficolin. <i>Molecular Immunology</i> , 2014, 60, 80-85.	2.2	45
58	The Role of the Individual Domains in the Structure and Function of the Catalytic Region of a Modular Serine Protease, C1r. <i>Journal of Immunology</i> , 2001, 167, 5202-5208.	0.8	43
59	The lectin-like activity of human C1q and its implication in DNA and apoptotic cell recognition. <i>FEBS Letters</i> , 2008, 582, 3111-3116.	2.8	43
60	Recombinant human complement subcomponent C1s lacking .beta.-hydroxyasparagine, sialic acid, and one of its two carbohydrate chains still reassembles with C1q and C1r to form a functional C1 complex. <i>Biochemistry</i> , 1992, 31, 4254-4262.	2.5	40
61	Assembly and Enzymatic Properties of the Catalytic Domain of Human Complement Protease C1r. <i>Journal of Biological Chemistry</i> , 2001, 276, 36233-36240.	3.4	40
62	Two parallel routes of the complement-mediated antibody-dependent enhancement of HIV-1 infection. <i>Aids</i> , 1997, 11, 949-958.	2.2	39
63	The Role of Nanometer-Scaled Ligand Patterns in Polyvalent Binding by Large Mannan-Binding Lectin Oligomers. <i>Journal of Immunology</i> , 2012, 188, 1292-1306.	0.8	39
64	Contribution of rare and predicted pathogenic gene variants to childhood-onset lupus: a large, genetic panel analysis of British and French cohorts. <i>Lancet Rheumatology</i> , The, 2020, 2, e99-e109.	3.9	38
65	[4] Human complement serine proteases and and their proenzymes. <i>Methods in Enzymology</i> , 1993, 223, 61-82.	1.0	37
66	Functional Characterization of Complement Proteases C1s/Mannan-binding Lectin-associated Serine Protease-2 (MASP-2) Chimeras Reveals the Higher C4 Recognition Efficacy of the MASP-2 Complement Control Protein Modules. <i>Journal of Biological Chemistry</i> , 2005, 280, 41811-41818.	3.4	36
67	The chaperone and potential mannan-binding lectin (MBL) co-receptor calreticulin interacts with MBL through the binding site for MBL-associated serine proteases. <i>FEBS Journal</i> , 2008, 275, 515-526.	4.7	35
68	Functional Characterization of the Recombinant Human C1 Inhibitor Serpin Domain: Insights into Heparin Binding. <i>Journal of Immunology</i> , 2010, 184, 4982-4989.	0.8	34
69	Active Human Complement Reduces the Zika Virus Load via Formation of the Membrane-Attack Complex. <i>Frontiers in Immunology</i> , 2018, 9, 2177.	4.8	33
70	Interaction of C1q With Pentraxin 3 and IgM Revisited: Mutational Studies With Recombinant C1q Variants. <i>Frontiers in Immunology</i> , 2019, 10, 461.	4.8	32
71	Antibodies targeting circulating protective molecules in lupus nephritis: Interest as serological biomarkers. <i>Autoimmunity Reviews</i> , 2018, 17, 890-899.	5.8	30
72	Editorial: The Role of Complement in Health and Disease. <i>Frontiers in Immunology</i> , 2019, 10, 1869.	4.8	30

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73	Mass Spectrometry Analysis of the Oligomeric C1q Protein Reveals the B Chain as the Target of Trypsin Cleavage and Interaction with Fucoidan. <i>Biochemistry</i> , 2005, 44, 2602-2609.	2.5	29
74	Assembly of C1 and the MBL and ficolin-MASP complexes: Structural insights. <i>Immunobiology</i> , 2007, 212, 279-288.	1.9	29
75	CD91 interacts with mannan-binding lectin (MBL) through the MBL-associated serine protease-binding site. <i>FEBS Journal</i> , 2010, 277, 4956-4964.	4.7	29
76	Complement Protein C1q Forms a Complex with Cytotoxic Prion Protein Oligomers. <i>Journal of Biological Chemistry</i> , 2010, 285, 19267-19276.	3.4	29
77	Mannan-binding lectin and C1q bind to distinct structures and exert differential effects on macrophages. <i>European Journal of Immunology</i> , 2000, 30, 1706-1713.	2.9	27
78	Human L-Ficolin Recognizes Phosphocholine Moieties of Pneumococcal Teichoic Acid. <i>Journal of Immunology</i> , 2014, 193, 5699-5708.	0.8	27
79	Structural and Functional Characterization of a Single-Chain Form of the Recognition Domain of Complement Protein C1q. <i>Frontiers in Immunology</i> , 2016, 7, 79.	4.8	27
80	Peptide Inhibitor of Complement C1 (PIC1) Rapidly Inhibits Complement Activation after Intravascular Injection in Rats. <i>PLoS ONE</i> , 2015, 10, e0132446.	2.5	27
81	Analysis of the N-linked oligosaccharides of human C1s using electrospray ionisation mass spectrometry. <i>FEBS Letters</i> , 1995, 358, 323-328.	2.8	26
82	C1R Mutations Trigger Constitutive Complement 1 Activation in Periodontal Ehlers-Danlos Syndrome. <i>Frontiers in Immunology</i> , 2019, 10, 2537.	4.8	26
83	Human complement subcomponent C2: purification and proteolytic cleavage in fluid phase by C1 <sub>1,s</sub> , C1 <sub>1,r</sub> -C1 <sub>1,s</sub> AND C1 <sub>1,r</sub> . <i>FEBS Letters</i> , 1982, 141, 19-24.	2.8	25
84	Purification and characterization of C4-binding protein from human serum. <i>FEBS Letters</i> , 1981, 132, 49-54.	2.8	24
85	Calcium-Dependent Complex Formation Between PBP2 and Lytic Transglycosylase SltB1 of <i>Pseudomonas aeruginosa</i> . <i>Microbial Drug Resistance</i> , 2012, 18, 298-305.	2.0	24
86	A novel peptide inhibitor of classical and lectin complement activation including ABO incompatibility. <i>Molecular Immunology</i> , 2013, 53, 132-139.	2.2	24
87	Enhancement of Ebola Virus Infection via Ficolin-1 Interaction with the Mucin Domain of GP Glycoprotein. <i>Journal of Virology</i> , 2016, 90, 5256-5269.	3.4	24
88	Association between the Presence of Autoantibodies Targeting Ficolin-3 and Active Nephritis in Patients with Systemic Lupus Erythematosus. <i>PLoS ONE</i> , 2016, 11, e0160879.	2.5	24
89	Structure and functions of the interaction domains of C1r and C1s: keystones of the architecture of the C1 complex. <i>Immunopharmacology</i> , 1999, 42, 3-13.	2.0	22
90	The Serine Protease Domain of MASP-3: Enzymatic Properties and Crystal Structure in Complex with Ecotin. <i>PLoS ONE</i> , 2013, 8, e67962.	2.5	22

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91	Structures of parasite calreticulins provide insights into their flexibility and dual carbohydrate/peptide-binding properties. <i>IUCr</i> , 2016, 3, 408-419.	2.2	21
92	M-ficolin and leukosialin (CD43): new partners in neutrophil adhesion. <i>Journal of Leukocyte Biology</i> , 2012, 91, 469-474.	3.3	20
93	C1q and Mannose-Binding Lectin Interact with CR1 in the Same Region on CCP24-25 Modules. <i>Frontiers in Immunology</i> , 2018, 9, 453.	4.8	19
94	Chemical characterization and location of ionic interactions involved in the assembly of the C1 complex of human complement. <i>The Protein Journal</i> , 1993, 12, 771-781.	1.1	17
95	Functional Role of the Linker between the Complement Control Protein Modules of Complement Protease C1s. <i>Journal of Immunology</i> , 2005, 175, 4536-4542.	0.8	17
96	Deciphering Key Residues Involved in the Virulence-promoting Interactions between <i>Streptococcus pneumoniae</i> and Human Plasminogen. <i>Journal of Biological Chemistry</i> , 2017, 292, 2217-2225.	3.4	17
97	Molecular and Cellular Interactions of Scavenger Receptor SR-F1 With Complement C1q Provide Insights Into Its Role in the Clearance of Apoptotic Cells. <i>Frontiers in Immunology</i> , 2020, 11, 544.	4.8	17
98	Classical Complement Pathway Components C1r and C1s: Purification from Human Serum and in Recombinant Form and Functional Characterization. <i>Methods in Molecular Biology</i> , 2014, 1100, 43-60.	0.9	17
99	Complement C1r serine protease contributes to kidney fibrosis. <i>American Journal of Physiology - Renal Physiology</i> , 2019, 317, F1293-F1304.	2.7	16
100	Structural and Functional Studies on C1r and C1s: New Insights into the Mechanisms Involved in C1 Activity and Assembly. <i>Immunobiology</i> , 1998, 199, 303-316.	1.9	15
101	Activation of classical pathway of complement cascade by soluble oligomers of prion. <i>Cellular Microbiology</i> , 2007, 9, 2870-2879.	2.1	15
102	Autoantibodies Targeting Ficolin-2 in Systemic Lupus Erythematosus Patients With Active Nephritis. <i>Arthritis Care and Research</i> , 2018, 70, 1263-1268.	3.4	14
103	Impact of the surface charge of polydiacetylene micelles on their interaction with human innate immune protein C1q and the complement system. <i>International Journal of Pharmaceutics</i> , 2018, 536, 434-439.	5.2	14
104	Effect of lactoperoxidase-catalyzed iodination on the calcium-dependent interactions of human C.hivin.1s. Location of the iodination sites. <i>Biochemistry</i> , 1991, 30, 7135-7141.	2.5	13
105	HIV-1 rsgp41 depends on calcium for binding of human C1q but not for binding of gp120. <i>Molecular Immunology</i> , 1995, 32, 371-374.	2.2	13
106	Structure, Function and Molecular Genetics of Human and Murine C1r. <i>Immunobiology</i> , 2002, 205, 365-382.	1.9	13
107	Studies on the mechanisms of allergen-induced activation of the classical and lectin pathways of complement. <i>Molecular Immunology</i> , 2003, 39, 839-846.	2.2	13
108	Recombinant C1q variants modulate macrophage responses but do not activate the classical complement pathway. <i>Molecular Immunology</i> , 2020, 117, 65-72.	2.2	12

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109	Molecular Basis of Complement C1q Collagen-Like Region Interaction with the Immunoglobulin-Like Receptor LAIR-1. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5125.	4.1	12
110	Protective Effect of Surfactant Protein D in Pulmonary Vaccinia Virus Infection: Implication of A27 Viral Protein. <i>Viruses</i> , 2013, 5, 928-953.	3.3	11
111	Two Different Missense C1S Mutations, Associated to Periodontal Ehlers-Danlos Syndrome, Lead to Identical Molecular Outcomes. <i>Frontiers in Immunology</i> , 2019, 10, 2962.	4.8	10
112	Involvement of Surfactant Protein D in Ebola Virus Infection Enhancement via Glycoprotein Interaction. <i>Viruses</i> , 2019, 11, 15.	3.3	10
113	Complement System and Alarmin HMGB1 Crosstalk: For Better or Worse. <i>Frontiers in Immunology</i> , 2022, 13, 869720.	4.8	10
114	Editorial: State-of-the-Art Research on C1q and the Classical Complement Pathway. <i>Frontiers in Immunology</i> , 2016, 7, 398.	4.8	9
115	Catalytically inactive Gla-domainless factor Xa binds to TFPI and restores <i>ex vivo</i> coagulation in hemophilia plasma. <i>Haematologica</i> , 2017, 102, e483-e485.	3.5	9
116	Interaction of Complement Defence Collagens C1q and Mannose-Binding Lectin with BMP-1/Tolloid-like Proteinases. <i>Scientific Reports</i> , 2017, 7, 16958.	3.3	9
117	Headless C1q: a new molecular tool to decipher its collagen-like functions. <i>FEBS Journal</i> , 2021, 288, 2030-2041.	4.7	8
118	Soluble FAS Ligand Enhances Suboptimal CD40L/IL-21-Mediated Human Memory B Cell Differentiation into Antibody-Secreting Cells. <i>Journal of Immunology</i> , 2021, 207, 449-458.	0.8	8
119	Neutron scattering study of the ( $^3$ -B) catalytic domains of complement proteases Cl <sub>1,r</sub> and Cl <sub>1,s</sub> . <i>FEBS Letters</i> , 1990, 269, 19-22.	2.8	7
120	Association of Terminal Complement Proteins in Solution and Modulation by Suramin,. <i>Biochemistry</i> , 1999, 38, 6807-6816.	2.5	7
121	Human ficolin-2 recognition versatility extended: An update on the binding of ficolin-2 to sulfated/phosphated carbohydrates. <i>FEBS Letters</i> , 2014, 588, 4694-4700.	2.8	7
122	Mode of PEG Coverage on Carbon Nanotubes Affects Binding of Innate Immune Protein C1q. <i>Journal of Physical Chemistry B</i> , 2018, 122, 757-763.	2.6	7
123	A study of a covalent-like interaction between soluble nascent C4b and C4-binding protein. <i>BBA - Proteins and Proteomics</i> , 1982, 704, 197-203.	2.1	6
124	A model system for the study of the assembly and regulation of human complement C3 convertase (classical pathway). <i>European Journal of Immunology</i> , 1986, 16, 617-622.	2.9	6
125	A Recombinant Chimeric Epidermal Growth Factor-like Module with High Binding Affinity for Integrins. <i>Journal of Biological Chemistry</i> , 2003, 278, 19834-19843.	3.4	6
126	Immunization with synthetic SARS-CoV-2 S glycoprotein virus-like particles protects macaques from infection. <i>Cell Reports Medicine</i> , 2022, 3, 100528.	6.5	6



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127	Identification of a cryptic protein kinase CK2 phosphorylation site in human complement protease C1r, and its use to probe intramolecular interaction. FEBS Letters, 1996, 386, 15-20.	2.8	5
128	Editorial: The double life of M-ficolin: what functions when circulating in serum and tethered to leukocyte surfaces?. Journal of Leukocyte Biology, 2011, 90, 410-412.	3.3	5
129	Complement C1q Interacts With LRP1 Clusters II and IV Through a Site Close but Different From the Binding Site of Its C1r and C1s-Associated Proteases. Frontiers in Immunology, 2020, 11, 583754.	4.8	5
130	Insights into the ligand binding specificity of SRECâ€™ (scavenger receptor expressed by endothelial) Tj ETQq0 0 0 rBT /Overlock 10 Tf	2.3	5
131	Comparative study of the fluid-phase proteolytic cleavage of human complement subcomponents C4 and C2 by Cs and Cr2 -Cs2. FEBS Letters, 1984, 165, 111-116.	2.8	4
132	MASP interactions with plasma-derived MBL. Molecular Immunology, 2012, 52, 79-87.	2.2	4
133	Role of C1q in Efferocytosis and Self-Tolerance â€” Links With Autoimmunity. , 2015, , .		4
134	Structure of the C1 complex of complement. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E5766-E5767.	7.1	4
135	Recognition protein C1q of innate immunity agglutinates nanodiamonds without activating complement. Nanomedicine: Nanotechnology, Biology, and Medicine, 2019, 18, 292-302.	3.3	4
136	Transient pentameric IgM fulfill biological functionâ€”Effect of expression host and transfection on IgM properties. PLoS ONE, 2020, 15, e0229992.	2.5	4
137	Arrangement of the C1 complex of complement. Biochemical Society Transactions, 1990, 18, 1148-1151.	3.4	3
138	Identification of the C1q binding sites of C1r and C1s: A refined 3D model of the C1 complex. Molecular Immunology, 2008, 45, 4097.	2.2	3
139	Biophysical Characterization of the Oligomeric States of Recombinant Immunoglobulins Type-M and Their C1q-Binding Kinetics by Biolayer Interferometry. Frontiers in Bioengineering and Biotechnology, 0, 10, .	4.1	3
140	C4 binding to artificial systems. FEBS Letters, 1981, 133, 151-156.	2.8	2
141	Ultrastructure of human C4-binding protein: proposition for a new model. European Journal of Immunology, 1985, 15, 941-945.	2.9	2
142	Innate immune sensing: Ligand recognition by M-ficolin is subject to a pH-dependent conformational switch. Molecular Immunology, 2007, 44, 3928-3929.	2.2	2
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