

# Isabelle Landrieu

## List of Publications by Year in descending order

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123  
papers

5,349  
citations

76326

40  
h-index

106344

65  
g-index

128  
all docs

128  
docs citations

128  
times ranked

6322  
citing authors

#	ARTICLE	IF	CITATIONS
1	Role of Tau as a Microtubule-Associated Protein: Structural and Functional Aspects. <i>Frontiers in Aging Neuroscience</i> , 2019, 11, 204.	3.4	294
2	Modulators of 14-3-3 Protein-Protein Interactions. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 3755-3778.	6.4	202
3	Identification of the Tau phosphorylation pattern that drives its aggregation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 9080-9085.	7.1	168
4	Cell signaling, post-translational protein modifications and NMR spectroscopy. <i>Journal of Biomolecular NMR</i> , 2012, 54, 217-236.	2.8	153
5	Hepatitis C Virus NS5A Protein Is a Substrate for the Peptidyl-prolyl cis/trans Isomerase Activity of Cyclophilins A and B. <i>Journal of Biological Chemistry</i> , 2009, 284, 13589-13601.	3.4	149
6	Structural Impact of Heparin Binding to Full-Length Tau As Studied by NMR Spectroscopy. <i>Biochemistry</i> , 2006, 45, 12560-12572.	2.5	142
7	<sup>1</sup> H NMR Study on the Binding of Pin1 Trp-Trp Domain with Phosphothreonine Peptides. <i>Journal of Biological Chemistry</i> , 2001, 276, 25150-25156.	3.4	115
8	Prevention of tau seeding and propagation by immunotherapy with a central tau epitope antibody. <i>Brain</i> , 2019, 142, 1736-1750.	7.6	113
9	Structural basis for oxygen degradation domain selectivity of the HIF prolyl hydroxylases. <i>Nature Communications</i> , 2016, 7, 12673.	12.8	109
10	Identification of O-GlcNAc sites within peptides of the Tau protein and their impact on phosphorylation. <i>Molecular BioSystems</i> , 2011, 7, 1420.	2.9	108
11	NMR Analysis of a Tau Phosphorylation Pattern. <i>Journal of the American Chemical Society</i> , 2006, 128, 3575-3583.	13.7	107
12	A small CDC25 dual-specificity tyrosine-phosphatase isoform in <i>Arabidopsis thaliana</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 13380-13385.	7.1	105
13	Mice lacking phosphatase PP2A subunit PR61/B $\alpha$ (Ppp2r5d) develop spatially restricted tauopathy by deregulation of CDK5 and GSK3 $\beta$ . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 6957-6962.	7.1	105
14	NMR observation of Tau in <i>Xenopus</i> oocytes. <i>Journal of Magnetic Resonance</i> , 2008, 192, 252-257.	2.1	100
15	Domain 3 of NS5A Protein from the Hepatitis C Virus Has Intrinsic $\alpha$ -Helical Propensity and Is a Substrate of Cyclophilin A. <i>Journal of Biological Chemistry</i> , 2011, 286, 20441-20454.	3.4	98
16	NMR Investigation of the Interaction between the Neuronal Protein Tau and the Microtubules. <i>Biochemistry</i> , 2007, 46, 3055-3064.	2.5	86
17	Alzheimer disease specific phosphoepitopes of Tau interfere with assembly of tubulin but not binding to microtubules. <i>FASEB Journal</i> , 2009, 23, 1146-1152.	0.5	80
18	Structural characterization by nuclear magnetic resonance of the impact of phosphorylation in the proline-rich region of the disordered Tau protein. <i>Proteins: Structure, Function and Bioinformatics</i> , 2012, 80, 454-462.	2.6	79

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19	The Peptidyl Prolyl cis/trans-Isomerase Pin1 Recognizes the Phospho-Thr212-Pro213 Site on Tau. <i>Biochemistry</i> , 2004, 43, 2032-2040.	2.5	77
20	Accepting its Random Coil Nature Allows a Partial NMR Assignment of the Neuronal Tau Protein. <i>ChemBioChem</i> , 2004, 5, 1639-1646.	2.6	74
21	Spectroscopic Studies of GSK3 $\beta$ Phosphorylation of the Neuronal Tau Protein and Its Interaction with the N-terminal Domain of Apolipoprotein E. <i>Journal of Biological Chemistry</i> , 2010, 285, 33435-33444.	3.4	71
22	Exacerbation of C1q dysregulation, synaptic loss and memory deficits in tau pathology linked to neuronal adenosine A2A receptor. <i>Brain</i> , 2019, 142, 3636-3654.	7.6	71
23	Nuclear Magnetic Resonance Spectroscopy Characterization of Interaction of Tau with DNA and Its Regulation by Phosphorylation. <i>Biochemistry</i> , 2015, 54, 1525-1533.	2.5	70
24	Involvement of 14-3-3 in tubulin instability and impaired axon development is mediated by Tau. <i>FASEB Journal</i> , 2015, 29, 4133-4144.	0.5	69
25	Tau phosphorylation regulates the interaction between BIN1's SH3 domain and Tau's proline-rich domain. <i>Acta Neuropathologica Communications</i> , 2015, 3, 58.	5.2	66
26	Molecular Implication of PP2A and Pin1 in the Alzheimer's Disease Specific Hyperphosphorylation of Tau. <i>PLoS ONE</i> , 2011, 6, e21521.	2.5	61
27	Nuclear Magnetic Resonance Analysis of the Acetylation Pattern of the Neuronal Tau Protein. <i>Biochemistry</i> , 2014, 53, 3020-3032.	2.5	60
28	Tau Aggregation in Alzheimer's Disease. <i>Prion</i> , 2007, 1, 21-25.	1.8	58
29	Systematic Identification of Tubulin-interacting Fragments of the Microtubule-associated Protein Tau Leads to a Highly Efficient Promoter of Microtubule Assembly. <i>Journal of Biological Chemistry</i> , 2011, 286, 33358-33368.	3.4	56
30	Stabilizer-Guided Inhibition of Protein-Protein Interactions. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 15720-15724.	13.8	56
31	Molecular mechanisms of the phospho-dependent prolyl cis/trans isomerase Pin1. <i>FEBS Journal</i> , 2007, 274, 5211-5222.	4.7	55
32	Immunophilin FKBP52 induces Tau-P301L filamentous assembly in vitro and modulates its activity in a model of tauopathy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 4584-4589.	7.1	55
33	Characterization of Neuronal Tau Protein as a Target of Extracellular Signal-regulated Kinase. <i>Journal of Biological Chemistry</i> , 2016, 291, 7742-7753.	3.4	54
34	Proline-Directed Random-Coil Chemical Shift Values as a Tool for the NMR Assignment of the Tau Phosphorylation Sites. <i>ChemBioChem</i> , 2004, 5, 73-78.	2.6	53
35	Regulation of Pin1 peptidyl-prolylcis/transisomerase activity by its WW binding module on a multi-phosphorylated peptide of Tau protein. <i>FEBS Letters</i> , 2005, 579, 4159-4164.	2.8	53
36	The Arabidopsis thaliana PIN1At Gene Encodes a Single-domain Phosphorylation-dependent Peptidyl Prolylcis/trans Isomerase. <i>Journal of Biological Chemistry</i> , 2000, 275, 10577-10581.	3.4	49

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37	The Talin Rod IBS2 $\beta$ -Helix Interacts with the $\beta$ 3 Integrin Cytoplasmic Tail Membrane-proximal Helix by Establishing Charge Complementary Salt Bridges. <i>Journal of Biological Chemistry</i> , 2008, 283, 24212-24223.	3.4	49
38	The elusive tau molecular structures: can we translate the recent breakthroughs into new targets for intervention?. <i>Acta Neuropathologica Communications</i> , 2019, 7, 31.	5.2	49
39	Selective intracellular accumulation of the major metabolite issued from the activation of the prodrug ethionamide in mycobacteria. <i>Journal of Antimicrobial Chemotherapy</i> , 2006, 58, 768-772.	3.0	47
40	Neurofibrillary degeneration of the Alzheimer-type: an alternate pathway to neuronal apoptosis?. <i>Biochemical Pharmacology</i> , 2003, 66, 1619-1625.	4.4	45
41	BIN1 recovers tauopathy-induced long-term memory deficits in mice and interacts with Tau through Thr348 phosphorylation. <i>Acta Neuropathologica</i> , 2019, 138, 631-652.	7.7	44
42	A Phosphorylation-Induced Turn Defines the Alzheimer's Disease AT8 Antibody Epitope on the Tau Protein. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 6819-6823.	13.8	41
43	Selectivity via Cooperativity: Preferential Stabilization of the p65/14-3-3 Interaction with Semisynthetic Natural Products. <i>Journal of the American Chemical Society</i> , 2020, 142, 11772-11783.	13.7	41
44	Arabidopsis PASTICCINO2 Is an Antiphosphatase Involved in Regulation of Cyclin-Dependent Kinase A. <i>Plant Cell</i> , 2006, 18, 1426-1437.	6.6	40
45	Characterization of the AT180 epitope of phosphorylated Tau protein by a combined nuclear magnetic resonance and fluorescence spectroscopy approach. <i>Biochemical and Biophysical Research Communications</i> , 2011, 412, 743-746.	2.1	40
46	Mechanism of Tau-Promoted Microtubule Assembly As Probed by NMR Spectroscopy. <i>Journal of the American Chemical Society</i> , 2014, 136, 12615-12623.	13.7	40
47	Structural Basis for the Non-Immunosuppressive Character of the Cyclosporin A Analogue Debio 025. <i>Biochemistry</i> , 2010, 49, 4679-4686.	2.5	39
48	Pin1 : A Therapeutic Target in Alzheimer Neurodegeneration. <i>Journal of Molecular Neuroscience</i> , 2002, 19, 275-288.	2.3	38
49	Characterization of the Arabidopsis thaliana Arath;CDC25 dual-specificity tyrosine phosphatase. <i>Biochemical and Biophysical Research Communications</i> , 2004, 322, 734-739.	2.1	38
50	Monitoring of the ethionamide pro-drug activation in mycobacteria by 1H high resolution magic angle spinning NMR. <i>Biochemical and Biophysical Research Communications</i> , 2005, 331, 452-458.	2.1	38
51	Microinjection of recombinant O-GlcNAc transferase potentiates Xenopus oocytes M-phase entry. <i>Biochemical and Biophysical Research Communications</i> , 2008, 369, 539-546.	2.1	38
52	Comparative analysis of Erk phosphorylation suggests a mixed strategy for measuring phosphoform distributions. <i>Molecular Systems Biology</i> , 2011, 7, 482.	7.2	38
53	Solution Structure of the Single-domain Prolyl Cis/Trans Isomerase PIN1At from Arabidopsis thaliana. <i>Journal of Molecular Biology</i> , 2002, 320, 321-332.	4.2	36
54	Regulation of the interaction between the neuronal BIN1 isoform 1 and Tau proteins: role of the SH3 domain. <i>FEBS Journal</i> , 2017, 284, 3218-3229.	4.7	35

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55	Fragment-based Differential Targeting of PPI Stabilizer Interfaces. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 6694-6707.	6.4	35
56	Major Differences between the Self-Assembly and Seeding Behavior of Heparin-Induced and in Vitro Phosphorylated Tau and Their Modulation by Potential Inhibitors. <i>ACS Chemical Biology</i> , 2019, 14, 1363-1379.	3.4	34
57	NMR spectroscopy of the neuronal tau protein: normal function and implication in Alzheimer's disease. <i>Biochemical Society Transactions</i> , 2010, 38, 1006-1011.	3.4	33
58	The FK506-binding protein FKBP52 in vitro induces aggregation of truncated Tau forms with prion-like behavior. <i>FASEB Journal</i> , 2015, 29, 3171-3181.	0.5	33
59	Tamoxifen Inhibits CDK5 Kinase Activity by Interacting with p35/p25 and Modulates the Pattern of Tau Phosphorylation. <i>Chemistry and Biology</i> , 2015, 22, 472-482.	6.0	33
60	Structural Analysis of Escherichia coli OpgG, a Protein Required for the Biosynthesis of Osmoregulated Periplasmic Glucans. <i>Journal of Molecular Biology</i> , 2004, 342, 195-205.	4.2	32
61	Regions of Tau Implicated in the Paired Helical Fragment Core as Defined by NMR. <i>ChemBioChem</i> , 2005, 6, 1849-1856.	2.6	32
62	Structural Basis of Tau Interaction With BIN1 and Regulation by Tau Phosphorylation. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 421.	2.9	32
63	Direct Crosstalk Between O-GlcNAcylation and Phosphorylation of Tau Protein Investigated by NMR Spectroscopy. <i>Frontiers in Endocrinology</i> , 2018, 9, 595.	3.5	32
64	Proline Conformation in a Functional Tau Fragment. <i>Journal of Molecular Biology</i> , 2016, 428, 79-91.	4.2	31
65	Inhibition of Tau seeding by targeting Tau nucleation core within neurons with a single domain antibody fragment. <i>Molecular Therapy</i> , 2022, 30, 1484-1499.	8.2	31
66	Phosphorylation and O-GlcNAcylation of the PHF-1 Epitope of Tau Protein Induce Local Conformational Changes of the C-Terminus and Modulate Tau Self-Assembly Into Fibrillar Aggregates. <i>Frontiers in Molecular Neuroscience</i> , 2021, 14, 661368.	2.9	30
67	Plasmodium falciparum Inhibitor-3 Homolog Increases Protein Phosphatase Type 1 Activity and Is Essential for Parasitic Survival. <i>Journal of Biological Chemistry</i> , 2012, 287, 1306-1321.	3.4	29
68	Inhibition of 14-3-3/Tau by Hybrid Small-Molecule Peptides Operating via Two Different Binding Modes. <i>ACS Chemical Neuroscience</i> , 2018, 9, 2639-2654.	3.5	29
69	Studying the Natively Unfolded Neuronal Tau Protein by Solution NMR Spectroscopy. <i>Protein and Peptide Letters</i> , 2006, 13, 235-246.	0.9	28
70	Characterization of ERM transactivation domain binding to the ACID/PTOV domain of the Mediator subunit MED25. <i>Nucleic Acids Research</i> , 2015, 43, 7110-7121.	14.5	28
71	Control of Protein-Protein Interactions: A Structure-Based Discovery of Low Molecular Weight Inhibitors of the Interactions between Pin1 WW Domain and Phosphopeptides. <i>Journal of Medicinal Chemistry</i> , 2005, 48, 4815-4823.	6.4	26
72	Unraveling a phosphorylation event in a folded protein by NMR spectroscopy: phosphorylation of the Pin1 WW domain by PKA. <i>Journal of Biomolecular NMR</i> , 2013, 55, 323-337.	2.8	26

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73	Isomerization and Oligomerization of Truncated and Mutated Tau Forms by FKBP52 are Independent Processes. <i>Journal of Molecular Biology</i> , 2016, 428, 1080-1090.	4.2	26
74	The Peptidyl-Prolyl Isomerase and Chaperone Par27 of <i>Bordetella pertussis</i> as the Prototype for a New Group of Parvulins. <i>Journal of Molecular Biology</i> , 2008, 376, 414-426.	4.2	25
75	Identification of a <i>Plasmodium falciparum</i> inhibitor motif involved in the binding and regulation activity of protein phosphatase type 1. <i>FEBS Journal</i> , 2014, 281, 4519-4534.	4.7	25
76	NMR Meets Tau: Insights into Its Function and Pathology. <i>Biomolecules</i> , 2016, 6, 28.	4.0	25
77	Zinc Binding to Tau Influences Aggregation Kinetics and Oligomer Distribution. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5979.	4.1	25
78	p13 and the WW Domain of PIN1 Bind to the Same Phosphothreonine-Proline Epitope. <i>Journal of Biological Chemistry</i> , 2001, 276, 1434-1438.	3.4	24
79	Structural and Functional Characterization of the Interaction between Cyclophilin B and a Heparin-derived Oligosaccharide. <i>Journal of Biological Chemistry</i> , 2007, 282, 34148-34158.	3.4	24
80	High-Resolution Magic Angle Spinning NMR of the Neuronal Tau Protein Integrated in Alzheimer's-Like Paired Helical Fragments. <i>Journal of the American Chemical Society</i> , 2005, 127, 10138-10139.	13.7	23
81	A functional fragment of Tau forms fibers without the need for an intermolecular cysteine bridge. <i>Biochemical and Biophysical Research Communications</i> , 2014, 445, 299-303.	2.1	23
82	Single Domain Antibody Fragments as New Tools for the Detection of Neuronal Tau Protein in Cells and in Mice Studies. <i>ACS Chemical Neuroscience</i> , 2019, 10, 3997-4006.	3.5	23
83	NMR Spectroscopy of the Main Protease of SARS-CoV-2 and Fragment-Based Screening Identify Three Protein Hotspots and an Antiviral Fragment. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 25428-25435.	13.8	22
84	Exploring the Molecular Function of PIN1 by Nuclear Magnetic Resonance. <i>Current Protein and Peptide Science</i> , 2006, 7, 179-194.	1.4	21
85	Towards understanding the phosphorylation code of tau. <i>Biochemical Society Transactions</i> , 2012, 40, 698-703.	3.4	20
86	Studying Posttranslational Modifications by In-Cell NMR. <i>Chemistry and Biology</i> , 2008, 15, 311-312.	6.0	17
87	Phosphorylation in intrinsically disordered regions regulates the activity of Neurogenin2. <i>BMC Biochemistry</i> , 2014, 15, 24.	4.4	17
88	Nuclear Magnetic Resonance Spectroscopy for the Identification of Multiple Phosphorylations of Intrinsically Disordered Proteins. <i>Journal of Visualized Experiments</i> , 2016, , .	0.3	17
89	Set-up and screening of a fragment library targeting the 14-3-3 protein interface. <i>MedChemComm</i> , 2019, 10, 1796-1802.	3.4	17
90	Design of Drug-Like Protein-Protein Interaction Stabilizers Guided By Chelation-Controlled Bioactive Conformation Stabilization. <i>Chemistry - A European Journal</i> , 2020, 26, 7131-7139.	3.3	17

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91	Tau pathology modulates Pin1 post-translational modifications and may be relevant as biomarker. <i>Neurobiology of Aging</i> , 2013, 34, 757-769.	3.1	16
92	The Study of Posttranslational Modifications of Tau Protein by Nuclear Magnetic Resonance Spectroscopy: Phosphorylation of Tau Protein by ERK2 Recombinant Kinase and Rat Brain Extract, and Acetylation by Recombinant Creb-Binding Protein. <i>Methods in Molecular Biology</i> , 2017, 1523, 179-213.	0.9	15
93	Two Tau binding sites on tubulin revealed by thiol-disulfide exchanges. <i>Scientific Reports</i> , 2018, 8, 13846.	3.3	15
94	Phosphorylated full-length Tau interacts with 14-3-3 proteins via two short phosphorylated sequences, each occupying a binding groove of 14-3-3 dimer. <i>FEBS Journal</i> , 2021, 288, 1918-1934.	4.7	13
95	Solution NMR Study of the Monomeric Form of p13 Protein Sheds Light on the Hinge Region Determining the Affinity for a Phosphorylated Substrate. <i>Journal of Biological Chemistry</i> , 2002, 277, 12375-12381.	3.4	12
96	Microtubule and MAPs. <i>Methods in Cell Biology</i> , 2010, 95, 449-480.	1.1	12
97	Molecular Basis for an Ancient Partnership between Prolyl Isomerase Pin1 and Phosphatase Inhibitor-2. <i>Biochemistry</i> , 2011, 50, 6567-6578.	2.5	12
98	Solution Structure of the N-Terminal Domain of Mediator Subunit MED26 and Molecular Characterization of Its Interaction with EAF1 and TAF7. <i>Journal of Molecular Biology</i> , 2017, 429, 3043-3055.	4.2	12
99	Backbone chemical shift assignments of human 14-3-3 $\beta$ . <i>Biomolecular NMR Assignments</i> , 2019, 13, 103-107.	0.8	11
100	ELISE NMR: Experimental liquid sealing of NMR samples. <i>Journal of Magnetic Resonance</i> , 2006, 181, 199-202.	2.1	10
101	A $\beta$ -Turn Motif in the Steroid Hormone Receptor's Ligand-Binding Domains Interacts with the Peptidyl-prolyl Isomerase (PPIase) Catalytic Site of the Immunophilin FKBP52. <i>Biochemistry</i> , 2016, 55, 5366-5376.	2.5	10
102	Adoption of a Turn Conformation Drives the Binding Affinity of p53 C-Terminal Domain Peptides to 14-3-3 $\beta$ . <i>ACS Chemical Biology</i> , 2020, 15, 262-271.	3.4	10
103	Dynamic interactions and Ca <sup>2+</sup> -binding modulate the holdase-type chaperone activity of S100B preventing tau aggregation and seeding. <i>Nature Communications</i> , 2021, 12, 6292.	12.8	10
104	Mitochondrial Asparaginyl-tRNA Synthetase Encoded by the Yeast Nuclear Gene YCR24c. <i>FEBS Journal</i> , 1997, 243, 268-273.	0.2	8
105	Selective backbone labelling of ILV methyl labelled proteins. <i>Journal of Biomolecular NMR</i> , 2009, 43, 219-227.	2.8	8
106	Nuclear Magnetic Resonance Spectroscopy Insights into Tau Structure in Solution: Impact of Post-translational Modifications. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1184, 35-45.	1.6	8
107	Ranking High Affinity Ligands of Low Solubility by NMR Spectroscopy. <i>ACS Medicinal Chemistry Letters</i> , 2011, 2, 485-487.	2.8	7
108	NMR reveals the intrinsically disordered domain 2 of NS5A protein as an allosteric regulator of the hepatitis C virus RNA polymerase NS5B. <i>Journal of Biological Chemistry</i> , 2017, 292, 18024-18043.	3.4	7

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109	H/D exchange of a 15 N labelled Tau fragment as measured by a simple Relax-EXSY experiment. Journal of Magnetic Resonance, 2014, 249, 32-37.	2.1	6
110	The O- <sup>12</sup> -linked N-acetylglucosaminylation of the Lamin B receptor and its impact on DNA binding and phosphorylation. Biochimica Et Biophysica Acta - General Subjects, 2018, 1862, 825-835.	2.4	6
111	Deciphering the Structure and Formation of Amyloids in Neurodegenerative Diseases With Chemical Biology Tools. Frontiers in Chemistry, 2022, 10, .	3.6	6
112	Identification of YHR019 in Saccharomyces cerevisiae chromosome VIII as the gene for the cytosolic asparaginyl-tRNA synthetase. , 1998, 14, 527-533.		5
113	Recombinant Production of the p10CKS1At Protein from Arabidopsis thaliana and 13C and 15N Double-Isotopic Enrichment for NMR Studies. Protein Expression and Purification, 1999, 16, 144-151.	1.3	5
114	Letter to the editor: sequence-specific 1H, 13C and 15N chemical shift backbone NMR assignment and secondary structure of the Arabidopsis thaliana PIN1At protein. Journal of Biomolecular NMR, 2000, 17, 271-272.	2.8	3
115	1H, 15N and 13C assignments of the N-terminal domain of the Mediator complex subunit MED26. Biomolecular NMR Assignments, 2016, 10, 233-236.	0.8	3
116	NMR spectroscopy of the main protease of SARS-CoV-2 and fragment-based screening identify three protein hotspots and an antiviral fragment. Angewandte Chemie, 2021, 133, 25632.	2.0	2
117	Assignment of the 1H, 13C and 15N resonances and secondary structure of the monomeric p13suc1 protein of Saccharomyces pombe. Journal of Biomolecular NMR, 2002, 23, 155-156.	2.8	1
118	1H, 13C, and 15N chemical shift assignment of human PACSIN1/syndapin I SH3 domain in solution. Biomolecular NMR Assignments, 2020, 14, 175-178.	0.8	1
119	Proline-Directed Random-Coil Chemical Shift Values as a Tool for the NMR Assignment of the Tau Phosphorylation Sites. ChemBioChem, 2004, 5, 256-256.	2.6	0
120	P4-032: MOLECULAR CHARACTERISATION OF BRIDGING INTEGRATOR 1 (BIN1) INTERACTION WITH TAU. , 2014, 10, P794-P794.		0
121	Nanobodies (VHHs) for targeting tau in Alzheimer's disease and tauopathies. IBRO Reports, 2019, 6, S100.	0.3	0
122	F20601: MAJOR DIFFERENCES BETWEEN THE SELF-ASSEMBLY, SEEDING BEHAVIOR, AND INTERACTION WITH MODULATORS OF HEPARIN-INDUCED VERSUS IN-VITRO PHOSPHORYLATED TAU. Alzheimer's and Dementia, 2019, 15, P524.	0.8	0
123	9. Post-translational Modifications of the Proteome: The Example of Tau in the Neuron and the Brain. , 2017, , 198-223.		0