## Gildas Bertho

List of Publications by Year in descending order

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		257450	276875
75	1,930	24	41
papers	citations	h-index	g-index
70	70	70	2755
79	79	79	2755
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Structural analysis of unstable norbixin isomers guided by pure shift nuclear magnetic resonance. Magnetic Resonance in Chemistry, 2022, 60, 504-514.	1.9	1
2	Ultrahigh-Resolution NMR with Water Signal Suppression for a Deeper Understanding of the Action of Antimetabolic Drugs on Diffuse Large B-Cell Lymphoma. Journal of Proteome Research, 2022, 21, 1041-1051.	3.7	9
3	Distinction between 2′- and 3′-Phosphate Isomers of a Fluorescent NADPH Analogue Led to Strong Inhibition of Cancer Cells Migration. Antioxidants, 2021, 10, 723.	5.1	1
4	Adenomyosis is associated with specific proton nuclear magnetic resonance (1H-NMR) serum metabolic profiles. Fertility and Sterility, 2021, 116, 243-254.	1.0	7
5	The complex metabolism of poststerone in male rats. Journal of Steroid Biochemistry and Molecular Biology, 2021, 212, 105897.	2.5	2
6	Loss of prion protein control of glucose metabolism promotes neurodegeneration in model of prion diseases. PLoS Pathogens, 2021, 17, e1009991.	4.7	11
7	On the Supra‣UMO Interaction: Case Study of a Sudden Change of Electronic Structure as a Functional Emergence. Chemistry - A European Journal, 2021, 27, 17889-17899.	3.3	3
8	Endometriosis phenotypes are associated with specific serum metabolic profiles determined by proton-nuclear magnetic resonance. Reproductive BioMedicine Online, 2020, 41, 640-652.	2.4	14
9	The follicular fluid metabolome differs according to the endometriosis phenotype. Reproductive BioMedicine Online, 2020, 41, 1023-1037.	2.4	20
10	Real-Time and Non-invasive Monitoring of the Activation of the IRE1 $\hat{l}_{\pm}$ -XBP1 Pathway in Individuals with Hemodynamic Impairment. EBioMedicine, 2018, 27, 284-292.	6.1	12
11	Urinary metabolic profiling of asymptomatic acute intermittent porphyria using a rule-mining-based algorithm. Metabolomics, 2018, 14, 10.	3.0	7
12	Insights into the interaction of high potency inhibitor IRCâ€083864 with phosphatase CDC25. Proteins: Structure, Function and Bioinformatics, 2017, 85, 593-601.	2.6	7
13	Model of the Interaction between the NF- $\hat{l}^{\circ}$ B Inhibitory Protein p100 and the E3 Ubiquitin Ligase $\hat{l}^{2}$ -TrCP based on NMR and Docking Experiments. Journal of Chemical Information and Modeling, 2017, 57, 223-233.	5.4	7
14	Specific Physical Exercise Improves Energetic Metabolism in the Skeletal Muscle of Amyotrophic-Lateral- Sclerosis Mice. Frontiers in Molecular Neuroscience, 2017, 10, 332.	2.9	37
15	Rule-Mining for the Early Prediction of Chronic Kidney Disease Based on Metabolomics and Multi-Source Data. PLoS ONE, 2016, 11, e0166905.	2.5	19
16	Interaction of a small molecule Natura- $\hat{l}_{\pm}$ and STAT3-SH2 domain to block Y705 phosphorylation and inhibit lupus nephritis. Biochemical Pharmacology, 2016, 99, 123-131.	4.4	6
17	Expression in yeast, new substrates, and construction of a first 3D model of human orphan cytochrome P450 2U1: Interpretation of substrate hydroxylation regioselectivity from docking studies. Biochimica Et Biophysica Acta - General Subjects, 2015, 1850, 1426-1437.	2.4	16
18	The urinary metabolome of chronic kidney disease. Kidney International, 2014, 85, 1239-1240.	5.2	5

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19	Novel RANK Antagonists for the Treatment of Bone-Resorptive Disease: Theoretical Predictions and Experimental Validation. Journal of Bone and Mineral Research, 2014, 29, 1466-1477.	2.8	12
20	Urinary Metabolic Fingerprint of Acute Intermittent Porphyria Analyzed by <sup>1</sup> H NMR Spectroscopy. Analytical Chemistry, 2014, 86, 2166-2174.	6.5	21
21	Toward Stable Electron Paramagnetic Resonance Oximetry Probes: Synthesis, Characterization, and Metabolic Evaluation of New Ester Derivatives of a Tris-( <i>para</i> -carboxyltetrathiaaryl)methyl (TAM) Radical. Chemical Research in Toxicology, 2013, 26, 1561-1569.	3.3	10
22	A new derivative detected in accelerated ageing of artesunate-amodiaquine fixed dose combination tablets. Journal of Pharmaceutical and Biomedical Analysis, 2013, 81-82, 20-26.	2.8	6
23	Thiolactone Sulfoxides as New Reactive Metabolites Acting as Bis-Electrophiles: Implication in Clopidogrel and Prasugrel Bioactivation. Chemical Research in Toxicology, 2013, 26, 794-802.	3.3	14
24	Tryptophan Depletion and the Kinase GCN2 Mediate IFN-γ–Induced Autophagy. Journal of Immunology, 2012, 189, 2954-2964.	0.8	38
25	Structural and Functional Characterization of Nrf2 Degradation by the Glycogen Synthase Kinase $3\hat{l}^2$ -TrCP Axis. Molecular and Cellular Biology, 2012, 32, 3486-3499.	2.3	338
26	Contact-based ligand-clustering approach for the identification of active compounds in virtual screening. Advances and Applications in Bioinformatics and Chemistry, 2012, 5, 61.	2.6	20
27	Cytochromes P450 Catalyze Both Steps of the Major Pathway of Clopidogrel Bioactivation, whereas Paraoxonase Catalyzes the Formation of a Minor Thiol Metabolite Isomer. Chemical Research in Toxicology, 2012, 25, 348-356.	3.3	108
28	Naphthalene-dioxygenase catalysed cis-dihydroxylation of bicyclic azaarenes. RSC Advances, 2012, 2, 605-615.	3.6	8
29	Metabolic Activation of Prasugrel: Nature of the Two Competitive Pathways Resulting in the Opening of Its Thiophene Ring. Chemical Research in Toxicology, 2012, 25, 1058-1065.	3.3	21
30	Paraoxonase-1 and clopidogrel efficacy. Nature Medicine, 2011, 17, 1040-1041.	30.7	50
31	NMR Applications for Identifying & Samp;#946;-TrCP Protein-Ligand Interactions. Mini-Reviews in Medicinal Chemistry, 2011, 11, 283-297.	2.4	4
32	Formation and Fate of a Sulfenic Acid Intermediate in the Metabolic Activation of the Antithrombotic Prodrug Prasugrel. Chemical Research in Toxicology, 2010, 23, 1268-1274.	3.3	30
33	Automatic clustering of docking poses in virtual screening process using self-organizing map. Bioinformatics, 2010, 26, 53-60.	4.1	63
34	Synthetic studies towards diazepanone scaffolds. Tetrahedron: Asymmetry, 2009, 20, 2320-2330.	1.8	20
35	Oxidative and Reductive Metabolism of Tris( <i>p</i> -carboxyltetrathiaaryl)methyl Radicals by Liver Microsomes. Chemical Research in Toxicology, 2009, 22, 1342-1350.	3.3	34
36	Possible role of region 152-156 in the structural duality of a peptide fragment from sheep prion protein. Protein Science, 2009, 13, 3151-3160.	7.6	25

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37	Metabolic Oxidative Cleavage of Thioesters: Evidence for the Formation of Sulfenic Acid Intermediates in the Bioactivation of the Antithrombotic Prodrugs Ticlopidine and Clopidogrel. Chemical Research in Toxicology, 2009, 22, 369-373.	3.3	88
38	Oxidation of tris-(p-carboxyltetrathiaaryl)methyl radical EPR probes: evidence for their oxidative decarboxylation and molecular origin of their specific ability to react with O2誉~'. Chemical Communications, 2009, , 1416.	4.1	27
39	Diastereoselective Additions to (3 <i>S</i> )â€3â€Aminodehydrocaprolactams: Development of a Versatile Synthesis of New Substituted Cyclic <scp>L</scp> â€Lysines. European Journal of Organic Chemistry, 2008, 2008, 1901-1909.	2.4	10
40	The key-role of tyrosine 155 in the mechanism of prion transconformation as highlighted by a study of sheep mutant peptides. Peptides, 2008, 29, 1073-1084.	2.4	4
41	Transfer-NMR and Docking Studies Identify the Binding of the Peptide Derived from Activating Transcription Factor 4 to Protein Ubiquitin Ligase $\hat{l}^2$ -TrCP. Competition STD-NMR with $\hat{l}^2$ -Catenin. Biochemistry, 2008, 47, 14-29.	2.5	28
42	Structure of the Complex between Phosphorylated Substrates and the SCF $\hat{l}^2$ -TrCP Ubiquitin Ligase Receptor: A Combined NMR, Molecular Modeling, and Docking Approach. Journal of Chemical Information and Modeling, 2008, 48, 2350-2361.	5.4	9
43	Phosphorylation-dependent structure of ATF4 peptides derived from a human ATF4 protein, a member of the family of transcription factors. Peptides, 2007, 28, 2253-2267.	2.4	17
44	Structural Studies on 24P-lîºBî± Peptide Derived from a Human lκB-α Protein Related to the Inhibition of the Activity of the Transcription Factor NF-κBâ€. Biochemistry, 2007, 46, 2958-2972.	2.5	16
45	Allosteric Tuning of the Intra-Cavity Binding Properties of a Calix[6]arene through External Binding to a ZnII Center Coordinated to Amino Side Chains. Chemistry - A European Journal, 2007, 13, 2078-2088.	3.3	29
46	Efficient synthesis of polyfunctionalised enantiopure diazepanone scaffolds. Tetrahedron Letters, 2007, 48, 8149-8152.	1.4	26
47	Ecdysteroids from the medicinal fernMicrosorum scolopendria (Burm. f.). Phytochemical Analysis, 2007, 18, 441-450.	2.4	28
48	STD and TRNOESY NMR studies for the epitope mapping of the phosphorylation motif of the oncogenic protein $\hat{I}^2$ -catenin recognized by a selective monoclonal antibody. FEBS Letters, 2006, 580, 5411-5422.	2.8	11
49	NMR studies for identifying phosphopeptide ligands of the HIV-1 protein Vpu binding to the F-box protein $\hat{I}^2$ -TrCP. Peptides, 2006, 27, 194-210.	2.4	17
50	Fixing the conformations of diamineplatinum(II)-GpG chelates: NMR and CD signatures of individual rotamers. Journal of Biological Inorganic Chemistry, 2006, 11, 139-152.	2.6	10
51	Synthesis and glycosidase inhibitory activity of new hexa-substituted C8-glycomimetics. Beilstein Journal of Organic Chemistry, 2005, 1, 12.	2.2	15
52	STD and TRNOESY NMR Studies on the Conformation of the Oncogenic Protein $\hat{l}^2$ -Catenin Containing the Phosphorylated Motif DpSGXXpS Bound to the $\hat{l}^2$ -TrCP Protein. Journal of Biological Chemistry, 2005, 280, 29107-29116.	3.4	23
53	First evidence that cytochrome P450 may catalyze both S-oxidation and epoxidation of thiophene derivatives. Biochemical and Biophysical Research Communications, 2005, 338, 450-455.	2.1	86
54	Solution structure of a peptide derived from the oncogenic protein $\hat{l}^2$ -Catenin in its phosphorylated and nonphosphorylated states. Peptides, 2005, 26, 227-241.	2.4	20

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55	Epitope Mapping of the Phosphorylation Motif of the HIV-1 Protein Vpu Bound to the Selective Monoclonal Antibody Using TRNOESY and STD NMR Spectroscopy. Biochemistry, 2004, 43, 14555-14565.	2.5	10
56	3-Carboxy-4-phosphonocyclopentane amino acids: New metabotropic glutamate receptor ligands. Amino Acids, 2003, 24, 303-310.	2.7	7
57	Synthesis and glycosidase inhibitory activity of enantiopure polyhydroxylated octahydroindoles and decahydroquinolines, analogs to castanospermine. Tetrahedron, 2003, 59, 8721-8730.	1.9	24
58	NMR Studies of the Phosphorylation Motif of the HIV-1 Protein Vpu Bound to the F-Box Protein $\hat{l}^2$ -TrCP. Biochemistry, 2003, 42, 14741-14751.	2.5	42
59	Antibiotic Resistance Peptides:  Interaction of Peptides Conferring Macrolide and Ketolide Resistance with Staphylococcus aureus Ribosomes. Conformation of Bound Peptides As Determined by Transferred NOE Experiments. Biochemistry, 2002, 41, 4218-4229.	2.5	23
60	The First Water-Soluble Copper(I) Calix[6]arene Complex Presenting a Hydrophobic Ligand Binding Pocket: A Remarkable Model for Active Sites in Metalloenzymes. Angewandte Chemie - International Edition, 2002, 41, 1044-1046.	13.8	71
61	Covalent bonding of bridged pyridinium aldehyde derivatives with guanine N7 is controlled by CpG site conformationâ€. Perkin Transactions II RSC, 2001, , 1771-1780.	1.1	0
62	Purification and structure of the major product obtained by reaction of NADPH and NMNH with the myeloperoxidase/hydrogen peroxide/chloride system. FEBS Journal, 2001, 268, 2889-2895.	0.2	10
63	Solution structure of the sheep prion PrP〚142-166ã€: a possible site for the conformational conversion of prion protein. Comptes Rendus De L'Academie Des Sciences - Series IIc: Chemistry, 2001, 4, 739-743.	0.1	1
64	A novel mechanism of antibiotic resistance: study of the complex state of peptides with bacterial Staphylococcus aureus ribosomes. Comptes Rendus De L'Academie Des Sciences - Series IIc: Chemistry, 2001, 4, 745-750.	0.1	0
65	Identification and quantitative analysis of the phytoecdysteroids in Silene species (Caryophyllaceae) by high-performance liquid chromatography. Journal of Chromatography A, 2001, 935, 309-319.	3.7	42
66	Sheep Prion Protein Synthetic Peptide Spanning Helix 1 and β-Strand 2 (Residues 142–166) Shows β-Hairpin Structure in Solution. Journal of Biological Chemistry, 2001, 276, 46364-46370.	3.4	32
67	Microbial hydroxylation/functionalization of terpenoid synthons derived from communic acids. Phytochemistry, 2000, 54, 23-27.	2.9	4
68	Lincomycin and clindamycin conformations. A fragment shared by macrolides, ketolides and lincosamides determined from TRNOE ribosome-bound conformations. Bioorganic and Medicinal Chemistry, 2000, 8, 1225-1243.	3.0	27
69	Conformations in solution and bound to bacterial ribosomes of ketolides, HMR 3647 (telithromycin) and RU 72366: A new class of highly potent antibacterials. Bioorganic and Medicinal Chemistry, 2000, 8, 1579-1597.	3.0	24
70	Dissociation–equilibrium constant and bound conformation for weak antibiotic binding interaction with different bacterial ribosomes â€. Perkin Transactions II RSC, 2000, , 2363-2371.	1.1	13
71	Conformational analysis of josamycin, a 16-membered macrolide free in solution and bound to bacterial ribosomes. Journal of the Chemical Society Perkin Transactions II, 1999, , 529-544.	0.9	9
72	Conformational Analysis of Ketolide, Conformations of RU 004 in Solution and Bound to Bacterial Ribosomes. Journal of Medicinal Chemistry, 1998, 41, 3373-3386.	6.4	39

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73	Transferred nuclear Overhauser effect study of macrolide–ribosome interactions: correlation between antibiotic activities and bound conformations. Bioorganic and Medicinal Chemistry, 1998, 6, 209-221.	3.0	44
74	Solution conformation of methylated macrolide antibiotics roxithromycin and erythromycin using NMR and molecular modelling. Ribosome-bound conformation determined by TRNOE and formation of cytochrome P450-metabolite complex. International Journal of Biological Macromolecules, 1998, 22, 103-127.	<b>7.</b> 5	29
75	Conformation of macrolides antibiotics bound to ribosomes as determined from transferred nuclear Overhauser effect spectroscopy. Journal De Chimie Physique Et De Physico-Chimie Biologique, 1998, 95, 423-429.	0.2	8