

Josep Vendrell

List of Publications by Year in descending order

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67
papers

3,535
citations

201575

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133188

59
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docs citations

70
times ranked

3931
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization, Recombinant Production and Structure-Function Analysis of NvCI, A Picomolar Metallocoxyypeptidase Inhibitor from the Marine Snail <i>Nerita versicolor</i> . <i>Marine Drugs</i> , 2019, 17, 511.	2.2	4
2	Synthesis and Structural/Functional Characterization of Selective M14 Metallocoxyypeptidase Inhibitors Based on Phosphinic Pseudopeptide Scaffold: Implications on the Design of Specific Optical Probes. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 1917-1931.	2.9	8
3	Discovery of Mechanism-Based Inactivators for Human Pancreatic Carboxypeptidase A from a Focused Synthetic Library. <i>ACS Medicinal Chemistry Letters</i> , 2017, 8, 1122-1127.	1.3	8
4	A functional and structural study of the major metalloprotease secreted by the pathogenic fungus <i>Aspergillus fumigatus</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 1946-1957.	2.5	22
5	Carboxypeptidase B. , 2013, , 1324-1329.		4
6	Insect Gut Carboxypeptidase 3. , 2013, , 1370-1375.		0
7	Linking amyloid protein aggregation and yeast survival. <i>Molecular BioSystems</i> , 2011, 7, 1121.	2.9	26
8	Structural and Functional Analysis of the Complex between Citrate and the Zinc Peptidase Carboxypeptidase A. <i>Enzyme Research</i> , 2011, 2011, 1-8.	1.8	11
9	Analysis of a new crystal form of procarboxypeptidase B: Further insights into the catalytic mechanism. <i>Biopolymers</i> , 2010, 93, 178-185.	1.2	11
10	Deciphering the role of the thermodynamic and kinetic stabilities of SH3 domains on their aggregation inside bacteria. <i>Proteomics</i> , 2010, 10, 4172-4185.	1.3	23
11	The X-Ray Structure of Carboxypeptidase A Inhibited by a Thiirane Mechanism-Based Inhibitor. <i>Chemical Biology and Drug Design</i> , 2010, 75, 29-34.	1.5	10
12	Progress in metallocoxyypeptidases and their small molecular weight inhibitors. <i>Biochimie</i> , 2010, 92, 1484-1500.	1.3	41
13	Aromatic Organic Compounds as Scaffolds for Metallocoxyypeptidase Inhibitor Design. <i>Chemical Biology and Drug Design</i> , 2009, 73, 75-82.	1.5	4
14	A new type of five-membered heterocyclic inhibitors of basic metallocoxyypeptidases. <i>European Journal of Medicinal Chemistry</i> , 2009, 44, 3266-3271.	2.6	7
15	Cyclobutane-containing peptides: Evaluation as novel metallocoxyypeptidase inhibitors and modelling of their mode of action. <i>Bioorganic and Medicinal Chemistry</i> , 2009, 17, 3824-3828.	1.4	42
16	Design, Selection, and Characterization of Thioflavin-Based Intercalation Compounds with Metal Chelating Properties for Application in Alzheimer's Disease. <i>Journal of the American Chemical Society</i> , 2009, 131, 1436-1451.	6.6	196
17	Direct interaction between a human digestive protease and the mucoadhesive poly(acrylic acid). <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2008, 64, 784-791.	2.5	14
18	Thioxophosphoranyl aryl- and heteroaryloxiranes as the representants of a new class of metallocoxyypeptidase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2008, 16, 4823-4828.	1.4	8

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19	The Crystal Structure of Thrombin-activable Fibrinolysis Inhibitor (TAFI) Provides the Structural Basis for Its Intrinsic Activity and the Short Half-life of TAFI. <i>Journal of Biological Chemistry</i> , 2008, 283, 29416-29423.	1.6	31
20	Metallo-carboxypeptidases: Emerging Drug Targets in Biomedicine. <i>Current Pharmaceutical Design</i> , 2007, 13, 347-364.	0.9	12
21	Metallo-carboxypeptidases: Emerging Drug Targets in Biomedicine. <i>Current Pharmaceutical Design</i> , 2007, 13, 349-366.	0.9	95
22	Caught after the Act: A Human A-Type Metallo-carboxypeptidase in a Product Complex with a Cleaved Hexapeptide. <i>Biochemistry</i> , 2007, 46, 6921-6930.	1.2	20
23	Ile-Phe Dipeptide Self-Assembly: Clues to Amyloid Formation. <i>Biophysical Journal</i> , 2007, 92, 1732-1741.	0.2	129
24	Self-assembly of human latexin into amyloid-like oligomers. <i>BMC Structural Biology</i> , 2007, 7, 75.	2.3	6
25	Structural and functional characterization of binding sites in metallo-carboxypeptidases based on Optimal Docking Area analysis. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 68, 131-144.	1.5	10
26	AGGRESCAN: a server for the prediction and evaluation of "hot spots" of aggregation in polypeptides. <i>BMC Bioinformatics</i> , 2007, 8, 65.	1.2	845
27	Response of the digestive system of <i>Helicoverpa zea</i> to ingestion of potato carboxypeptidase inhibitor and characterization of an uninhibited carboxypeptidase B. <i>Insect Biochemistry and Molecular Biology</i> , 2006, 36, 654-664.	1.2	34
28	Mutagenesis of the central hydrophobic cluster in A β 42 Alzheimer's peptide. Side-chain properties correlate with aggregation propensities. <i>FEBS Journal</i> , 2006, 273, 658-668.	2.2	164
29	Detailed molecular comparison between the inhibition mode of A/B-type carboxypeptidases in the zymogen state and by the endogenous inhibitor latexin. <i>Cellular and Molecular Life Sciences</i> , 2005, 62, 1996-2014.	2.4	19
30	Prediction of "hot spots" of aggregation in disease-linked polypeptides. <i>BMC Structural Biology</i> , 2005, 5, 18.	2.3	173
31	Structural basis of the resistance of an insect carboxypeptidase to plant protease inhibitors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 16602-16607.	3.3	64
32	Structure of human carboxypeptidase A4 with its endogenous protein inhibitor, latexin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 3978-3983.	3.3	89
33	Human kallikrein 6 activity is regulated via an autoproteolytic mechanism of activation/inactivation. <i>Biological Chemistry</i> , 2004, 385, 517-24.	1.2	62
34	Secondary Binding Site of the Potato Carboxypeptidase Inhibitor. Contribution to Its Structure, Folding, and Biological Properties. <i>Biochemistry</i> , 2004, 43, 7973-7982.	1.2	18
35	Amyloid Fibril Formation by a Partially Structured Intermediate State of β -Chymotrypsin. <i>Journal of Molecular Biology</i> , 2004, 342, 321-331.	2.0	206
36	Carboxypeptidase B. , 2004, , 831-833.		1

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37	Procarboxypeptidase A from the insect pest <i>Helicoverpa armigera</i> and its derived enzyme. <i>FEBS Journal</i> , 2003, 270, 3026-3035.	0.2	27
38	NMR solution structure of the activation domain of human procarboxypeptidase A2. <i>Protein Science</i> , 2003, 12, 296-305.	3.1	9
39	Major Kinetic Traps for the Oxidative Folding of Leech Carboxypeptidase Inhibitor. <i>Biochemistry</i> , 2003, 42, 6754-6761.	1.2	26
40	The Unfolding Pathway of Leech Carboxypeptidase Inhibitor. <i>Journal of Biological Chemistry</i> , 2002, 277, 17538-17543.	1.6	19
41	Identification and Characterization of Three Members of the Human Metalloprocarboxypeptidase Gene Family. <i>Journal of Biological Chemistry</i> , 2002, 277, 14954-14964.	1.6	69
42	Human Procarboxypeptidase B: Three-dimensional Structure and Implications for Thrombin-activatable Fibrinolysis Inhibitor (TAFI). <i>Journal of Molecular Biology</i> , 2002, 321, 537-547.	2.0	66
43	Crystal structure of a novel Mid-gut procarboxypeptidase from the cotton pest <i>Helicoverpa armigera</i> . <i>Journal of Molecular Biology</i> , 2001, 313, 629-638.	2.0	42
44	The Crystal Structure of the Inhibitor-complexed Carboxypeptidase D Domain II and the Modeling of Regulatory Carboxypeptidases. <i>Journal of Biological Chemistry</i> , 2001, 276, 16177-16184.	1.6	71
45	Structure of a novel leech carboxypeptidase inhibitor determined free in solution and in complex with human carboxypeptidase A2. <i>Nature Structural Biology</i> , 2000, 7, 322-328.	9.7	71
46	Metalloprocarboxypeptidases and their protein inhibitors. <i>BBA - Proteins and Proteomics</i> , 2000, 1477, 284-298.	2.1	129
47	Mapping the Pro-region of Carboxypeptidase B by Protein Engineering. <i>Journal of Biological Chemistry</i> , 1999, 274, 19925-19933.	1.6	45
48	Carboxypeptidases. , 1999, , 13-34.		8
49	Cutting at the right place. The importance of selective limited proteolysis in the activation of proproteinase E. <i>FEBS Journal</i> , 1998, 251, 839-844.	0.2	7
50	Comparative Analysis of the Sequences and Three-Dimensional Models of Human Procarboxypeptidases A1, A2 and B. <i>Biological Chemistry</i> , 1998, 379, 149-156.	1.2	13
51	A Carboxypeptidase Inhibitor from the Medical Leech <i>Hirudo medicinalis</i> . <i>Journal of Biological Chemistry</i> , 1998, 273, 32927-32933.	1.6	78
52	Overexpression of Human Procarboxypeptidase A2 in <i>Pichia pastoris</i> and Detailed Characterization of Its Activation Pathway. <i>Journal of Biological Chemistry</i> , 1998, 273, 3535-3541.	1.6	52
53	Characterisation and preliminary X-ray diffraction analysis of human pancreatic procarboxypeptidase A2. <i>FEBS Letters</i> , 1997, 420, 7-10.	1.3	19
54	The activation pathway of procarboxypeptidase B from porcine pancreas: Participation of the active enzyme in the proteolytic processing. <i>Protein Science</i> , 1995, 4, 1792-1800.	3.1	27

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55	The Sequence and Conformation of Human Pancreatic Procarboxypeptidase A2. <i>Journal of Biological Chemistry</i> , 1995, 270, 6651-6657.	1.6	31
56	Procarboxypeptidase in rat pancreas Overall characterization and comparison of the activation processes. <i>FEBS Journal</i> , 1994, 222, 55-64.	0.2	22
57	Advances in metallo-procarboxypeptidases. , 1994, , 19-27.		0
58	Advances in metallo-procarboxypeptidases. Emerging details on the inhibition mechanism and on the activation process. <i>FEBS Journal</i> , 1993, 211, 381-389.	0.2	77
59	Pancreatic Procarboxypeptidases: Their Activation Processes Related to the Structural Features of the Zymogens and Activation Segments. <i>Biological Chemistry Hoppe-Seyler</i> , 1992, 373, 387-392.	1.4	12
60	Differential scanning calorimetric study of carboxypeptidase B, procarboxypeptidase B and its globular activation domain. <i>FEBS Journal</i> , 1991, 200, 663-670.	0.2	46
61	Sequence-Specific ¹ H NMR Assignments and Determination of the Secondary Structure for the Activation Domain Isolated from Pancreatic Procarboxypeptidase B. <i>Biochemistry</i> , 1990, 29, 7515-7522.	1.2	19
62	Autolysis of proproteinase E in bovine procarboxypeptidase A ternary complex gives rise to subunit III. <i>FEBS Letters</i> , 1990, 277, 37-41.	1.3	19
63	The separation of pancreatic procarboxypeptidases by high-performance liquid chromatography and chromatofocusing. <i>Journal of Chromatography A</i> , 1989, 481, 233-243.	1.8	7
64	Enzymatic and chemical fragmentation of proteins: A simple laboratory visualization of differences in yield, specificity and applicability. <i>Biochemical Education</i> , 1988, 16, 174-176.	0.1	2
65	Complete amino acid analysis of proteins by dansyl derivatization and reversed-phase liquid chromatography. <i>Journal of Chromatography A</i> , 1986, 358, 401-413.	1.8	88
66	Nuclear magnetic resonance studies on the isolated activation segment from porcine pancreatic procarboxypeptidase A. <i>Biochemical Society Transactions</i> , 1985, 13, 344-345.	1.6	0
67	Isolation and re-association of the subunits from the pro-(carboxypeptidase A)â€™pro-(proteinase E) binary complex from pig pancreas. <i>Biochemical Journal</i> , 1982, 205, 449-452.	1.7	17