Vincenzo De Filippis

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	NOG-Derived Peptides Can Restore Neuritogenesis on a CRASH Syndrome Cell Model. Biomedicines, 2022, 10, 102.	3.2	3
2	Exogenous human α-Synuclein acts in vitro as a mild platelet antiaggregant inhibiting α-thrombin-induced platelet activation. Scientific Reports, 2022, 12, .	3.3	4
3	Mapping specificity, cleavage entropy, allosteric changes and substrates of blood proteases in a high-throughput screen. Nature Communications, 2021, 12, 1693.	12.8	17
4	A conserved Neurite Outgrowth and Guidance motif with biomimetic potential in neuronal Cell Adhesion Molecules. Computational and Structural Biotechnology Journal, 2021, 19, 5622-5636.	4.1	3
5	A serine protease secreted from Bacillus subtilis cleaves human plasma transthyretin to generate an amyloidogenic fragment. Communications Biology, 2020, 3, 764.	4.4	12
6	Effects of point mutations in the binding pocket of the mouse major urinary protein MUP20 on ligand affinity and specificity. Scientific Reports, 2019, 9, 300.	3.3	18
7	Fibronectin-binding protein B (FnBPB) from Staphylococcus aureus protects against the antimicrobial activity of histones. Journal of Biological Chemistry, 2019, 294, 3588-3602.	3.4	39
8	Noncoded amino acids in protein engineering: Structure–activity relationship studies of hirudin–thrombin interaction. Biotechnology and Applied Biochemistry, 2018, 65, 69-80.	3.1	11
9	Protein engineering by chemical methods: Incorporation of nonnatural amino acids as a tool for studying protein folding, stability, and function. Peptide Science, 2018, 110, e24090.	1.8	4
10	Single-Cell and Single-Molecule Analysis Unravels the Multifunctionality of the <i>Staphylococcus aureus</i> Collagen-Binding Protein Cna. ACS Nano, 2017, 11, 2160-2170.	14.6	47
11	Impaired Release of Vitamin D in Dysfunctional Adipose Tissue: New Cues on Vitamin D Supplementation in Obesity. Journal of Clinical Endocrinology and Metabolism, 2017, 102, 2564-2574.	3.6	40
12	Non-canonical proteolytic activation of human prothrombin by subtilisin from Bacillus subtilis may shift the procoagulant–anticoagulant equilibrium toward thrombosis. Journal of Biological Chemistry, 2017, 292, 15161-15179.	3.4	28
13	Molecular mapping of α-thrombin (αT)/β2-glycoprotein I (β2GpI) interaction reveals how β2GpI affects αT functions. Biochemical Journal, 2016, 473, 4629-4650.	3.7	16
14	Osteocalcin and Sex Hormone Binding Globulin Compete on a Specific Binding Site of GPRC6A. Endocrinology, 2016, 157, 4473-4486.	2.8	43
15	A MICU1 Splice Variant Confers High Sensitivity to the Mitochondrial Ca2+ Uptake Machinery of Skeletal Muscle. Molecular Cell, 2016, 64, 760-773.	9.7	97
16	Loop Electrostatics Asymmetry Modulates the Preexisting Conformational Equilibrium in Thrombin. Biochemistry, 2016, 55, 3984-3994.	2.5	17
17	Enhanced neuronal cell differentiation combining biomimetic peptides and a carbon nanotube-polymer scaffold. Nanomedicine: Nanotechnology, Biology, and Medicine, 2015, 11, 621-632.	3.3	39
18	Thrombin inhibits the anti-myeloperoxidase and ferroxidase functions of ceruloplasmin: relevance in rheumatoid arthritis. Free Radical Biology and Medicine, 2015, 86, 279-294.	2.9	36

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19	Oxidation of Met1606 in von Willebrand factor is a risk factor for thrombotic and septic complications in chronic renal failure. Biochemical Journal, 2012, 442, 423-432.	3.7	18
20	Modeling ADAMTS13-von Willebrand Factor interaction: Implications for oxidative stress-related cardiovascular diseases and type 2A von Willebrand Disease. Biophysical Chemistry, 2012, 160, 1-11.	2.8	7
21	A novel protein from the serum of Python sebae, structurally homologous with type-γ phospholipase A2 inhibitor, displays antitumour activity. Biochemical Journal, 2011, 440, 251-262.	3.7	13
22	Formation of methionine sulfoxide by peroxynitrite at position 1606 of von Willebrand factor inhibits its cleavage by ADAMTS-13: A new prothrombotic mechanism in diseases associated with oxidative stress. Free Radical Biology and Medicine, 2010, 48, 446-456.	2.9	56
23	Chemical synthesis and characterization of wildâ€type and biotinylated Nâ€terminal domain 1–64 of β2â€glycoprotein I. Protein Science, 2010, 19, 1065-1078.	7.6	23
24	Thrombin Inhibition by Serpins Disrupts Exosite II. Journal of Biological Chemistry, 2010, 285, 38621-38629.	3.4	21
25	Conformational and biochemical characterization of a biologically active rat recombinant Protease Nexin-1 expressed in E. coli. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2009, 1794, 602-614.	2.3	11
26	Fibrinogen-elongated Î ³ Chain Inhibits Thrombin-induced Platelet Response, Hindering the Interaction with Different Receptors. Journal of Biological Chemistry, 2008, 283, 30193-30204.	3.4	34
27	o-Nitrotyrosine andp-iodophenylalanine as spectroscopic probes for structural characterization of SH3 complexes. Protein Science, 2007, 16, 1257-1265.	7.6	7
28	3-Nitrotyrosine as a spectroscopic probe for investigating protein-protein interactions. Protein Science, 2006, 15, 976-986.	7.6	65
29	Effect of Na+ binding on the conformation, stability and molecular recognition properties of thrombin. Biochemical Journal, 2005, 390, 485-492.	3.7	44
30	Incorporation of the fluorescent amino acid 7-azatryptophan into the core domain 1-47 of hirudin as a probe of hirudin folding and thrombin recognition. Protein Science, 2004, 13, 1489-1502.	7.6	50
31	Probing the Hirudinâ^'Thrombin Interaction by Incorporation of Noncoded Amino Acids and Molecular Dynamics Simulation,. Biochemistry, 2002, 41, 13556-13569.	2.5	28
32	Minimal structural requirements for agonist activity of PAR-2 activating peptides. Bioorganic and Medicinal Chemistry Letters, 2002, 12, 21-24.	2.2	7
33	Incorporation of noncoded amino acids into the Nâ€ŧerminal domain 1â€47 of hirudin yields a highly potent and selective thrombin inhibitor. Protein Science, 1999, 8, 2213-2217.	7.6	21
34	Limited proteolysis of bovine αâ€ŀactalbumin: Isolation and characterization of protein domains. Protein Science, 1999, 8, 2290-2303.	7.6	50
35	Enhanced Protein Thermostability by Ala → Aib Replacementâ€. Biochemistry, 1998, 37, 1686-1696.	2.5	49
36	Synthesis and Characterization of More Potent Analogues of Hirudin Fragment 1â^'47 Containing Non-Natural Amino Acids,. Biochemistry, 1998, 37, 13507-13515.	2.5	35

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37	The complete mature bovine prion protein highly expressed inEscherichia coli: biochemical and structural studies. FEBS Letters, 1997, 412, 359-364.	2.8	24
38	Probing the partly folded states of proteins by limited proteolysis. Folding & Design, 1997, 2, R17-R26.	4.5	279
39	Limited proteolysis of ribonuclease A with thermolysin in trifluoroethanol. Protein Science, 1997, 6, 860-872.	7.6	28
40	Acid-Induced Molten Globule State of a Fully Active Mutant of Human Interleukin-6. Biochemistry, 1996, 35, 11503-11511.	2.5	33
41	Limited Proteolysis of Lysozyme in Trifluoroethanol. Isolation and Characterization of a Partially Active Enzyme Derivative. FEBS Journal, 1995, 230, 779-787.	0.2	25
42	Structure, Stability and Biological Properties of a N-terminally Truncated form of Recombinant Human Interleukin-6 Containing a Single Disulfide Bond. FEBS Journal, 1995, 227, 573-581.	0.2	24
43	Core Domain of Hirudin from the Leech Hirudinaria manillensis: Chemical Synthesis, Purification, and Characterization of a Trp3 Analog of Fragment 1-47. Biochemistry, 1995, 34, 9552-9564.	2.5	41
44	Limited proteolysis of cytochromecin trifluoroethanol. FEBS Letters, 1995, 362, 266-270.	2.8	35
45	Probing the Structure of Hirudin from Hirudinaria manillensis by Limited Proteolysis. Isolation, Characterization and Thrombin-Inhibitory Properties of N-Terminal Fragments. FEBS Journal, 1994, 226, 323-333.	0.2	37
46	The thermodynamics of the unfolding of an isolated protein subdomain The 255-316 C-terminal fragment of thermolysin. FEBS Letters, 1994, 344, 154-156.	2.8	14
47	NMR Solution Structure of the C-Terminal Fragment 255-316 of Thermolysin: A Dimer Formed by Subunits Having the Native Structure. Biochemistry, 1994, 33, 14834-14847.	2.5	29
48	Cumulative stabilizing effects of glycine to alanine substitutions in Bacillus subtilis neutral protease. Protein Engineering, Design and Selection, 1992, 5, 543-550.	2.1	32
49	Synthesis and conformational studies of peptides encompassing the carboxyâ€ŧerminal helix of thermolysin. International Journal of Peptide and Protein Research, 1990, 35, 396-405.	0.1	3
50	Semisynthesis of carboxyâ€ŧerminal fragments of thermolysin. International Journal of Peptide and Protein Research, 1990, 35, 219-221.	0.1	19