## Ivano Eberini

List of Publications by Year in descending order

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100 papers 2,722 citations

30 h-index 214800 47 g-index

105 all docs 105 docs citations

105 times ranked 3829 citing authors

#	Article	IF	CITATIONS
1	The oxysterol–CXCR2 axis plays a key role in the recruitment of tumor-promoting neutrophils. Journal of Experimental Medicine, 2013, 210, 1711-1728.	8.5	167
2	The Molecular Basis of Lecithin:Cholesterol Acyltransferase Deficiency Syndromes. Arteriosclerosis, Thrombosis, and Vascular Biology, 2005, 25, 1972-1978.	2.4	158
3	Proteins of rat serum, urine, and cerebrospinal fluid: VI. Further protein identifications and interstrain comparison. Electrophoresis, 2001, 22, 3043-3052.	2.4	96
4	Acute-Phase Proteins Before Cerebral Ischemia in Stroke-Prone Rats. Stroke, 2001, 32, 753-760.	2.0	93
5	Characterization of the Protein Unfolding Processes Induced by Urea and Temperature. Biophysical Journal, 2008, 94, 2241-2251.	0.5	85
6	Potent inhibitors of human LAT1 (SLC7A5) transporter based on dithiazole and dithiazine compounds for development of anticancer drugs. Biochemical Pharmacology, 2017, 143, 39-52.	4.4	72
7	Proteins of rat serum: I. Establishing a reference two-dimensional electrophoresis map by immunodetection and microbore high performance liquid chromatography-electrospray mass spectrometry. Electrophoresis, 1998, 19, 1484-1492.	2.4	67
8	Novel insights into the transport mechanism of the human amino acid transporter LAT1 (SLC7A5). Probing critical residues for substrate translocation. Biochimica Et Biophysica Acta - General Subjects, 2017, 1861, 727-736.	2.4	64
9	Biosynthesis of Flavin Cofactors in Man: Implications in Health and Disease. Current Pharmaceutical Design, 2013, 19, 2649-2675.	1.9	61
10	Rabphilin 3A retains NMDA receptors at synaptic sites through interaction with GluN2A/PSD-95 complex. Nature Communications, 2015, 6, 10181.	12.8	59
11	Hypocholesterolaemic effects of soya proteins: results of recent studies are predictable from the Anderson meta-analysis data. British Journal of Nutrition, 2007, 97, 816-822.	2.3	58
12	Coordinated and reversible reduction of enzymes involved in terminal oxidative metabolism in skeletal muscle mitochondria from a riboflavin-responsive, multiple acyl-CoA dehydrogenase deficiency patient. Electrophoresis, 2006, 27, 1182-1198.	2.4	55
13	Reference maps of mouse serum acute-phase proteins: Changes with LPS-induced inflammation and apolipoproteinâ€A-I and A-II transgenes. Proteomics, 2005, 5, 4245-4253.	2.2	53
14	In silico identification of new ligands for GPR17: a promising therapeutic target for neurodegenerative diseases. Journal of Computer-Aided Molecular Design, 2011, 25, 743-752.	2.9	53
15	With or without you â€" Proteomics with or without major plasma/serum proteins. Journal of Proteomics, 2016, 140, 62-80.	2.4	53
16	Role of the <scp>GM</scp> 1 ganglioside oligosaccharide portion in the TrkAâ€dependent neurite sprouting in neuroblastoma cells. Journal of Neurochemistry, 2017, 143, 645-659.	3.9	53
17	Reorganization in apo- and holo-β-lactoglobulin upon protonation of Glu89: Molecular dynamics and pKa calculations. Proteins: Structure, Function and Bioinformatics, 2004, 54, 744-758.	2.6	50
18	Serum protein pattern during cow pregnancy: Acute-phase proteins increase in the peripartum period. Electrophoresis, 2006, 27, 1617-1625.	2.4	50

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19	Monitoring the effects of drug treatment in rat models of disease by serum protein analysis. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 771, 107-130.	2.3	48
20	Proteins of rat serum: III. Gender-related differences in protein concentration under baseline conditions and upon experimental inflammation as evaluated by two-dimensional electrophoresis. Electrophoresis, 1999, 20, 836-845.	2.4	46
21	A Computational Approach to Evaluate the Androgenic Affinity of Iprodione, Procymidone, Vinclozolin and Their Metabolites. PLoS ONE, 2014, 9, e104822.	2.5	46
22	Oxysterols act as promiscuous ligands of class-A GPCRs: In silico molecular modeling and in vitro validation. Cellular Signalling, 2014, 26, 2614-2620.	3.6	46
23	Proteins of rat serum: II. Influence of some biological parameters of the two-dimensional electrophoresis pattern. Electrophoresis, 1998, 19, 1493-1500.	2.4	43
24	Redox regulation of cyclophilin A by glutathionylation. Proteomics, 2006, 6, 817-825.	2.2	43
25	Analysis of <i>Lupinus albus </i> Storage Proteins by Two-Dimensional Electrophoresis and Mass Spectrometry. Journal of Agricultural and Food Chemistry, 2005, 53, 4599-4606.	5.2	40
26	Predicting estrogen receptor binding of chemicals using a suite of in silico methods $\hat{a} \in ``Complementary approaches of (Q)SAR, molecular docking and molecular dynamics. Toxicology and Applied Pharmacology, 2019, 378, 114630.$	2.8	37
27	Inhibition of SIRT1 deacetylase and p53 activation uncouples the anti-inflammatory and chemopreventive actions of NSAIDs. British Journal of Cancer, 2019, 120, 537-546.	6.4	37
28	Computational and experimental approaches assess the interactions between bovine $\hat{l}^2$ -lactoglobulin and synthetic compounds of pharmacological interest. Journal of Molecular Graphics and Modelling, 2008, 26, 1004-1013.	2.4	35
29	C2238 Atrial Natriuretic Peptide Molecular Variant Is Associated With Endothelial Damage and Dysfunction Through Natriuretic Peptide Receptor C Signaling. Circulation Research, 2013, 112, 1355-1364.	4.5	34
30	Proteins of rat serum V: Adjuvant arthritis and its modulation by nonsteroidal anti-inflammatory drugs. Electrophoresis, 2000, 21, 2170-2180.	2.4	32
31	Human FAD synthase is a bi-functional enzyme with a FAD hydrolase activity in the molybdopterin binding domain. Biochemical and Biophysical Research Communications, 2015, 465, 443-449.	2.1	29
32	Gender differences in endothelial function and inflammatory markers along the occurrence of pathological events in stroke-prone rats. Experimental and Molecular Pathology, 2007, 82, 33-41.	2.1	28
33	Identification of small molecules uncoupling the Notch::Jagged interaction through an integrated high-throughput screening. PLoS ONE, 2017, 12, e0182640.	2.5	28
34	Nimesulide binding site in the BOAT1 (SLC6A19) amino acid transporter. Mechanism of inhibition revealed by proteoliposome transport assay and molecular modelling. Biochemical Pharmacology, 2014, 89, 422-430.	4.4	27
35	Bacterial Production, Characterization and Protein Modeling of a Novel Monofuctional Isoform of FAD Synthase in Humans: An Emergency Protein?. Molecules, 2018, 23, 116.	3.8	26
36	pH and Ionic Strength Dependence of Protein (Un)Folding and Ligand Binding to Bovine β-Lactoglobulins A and Bâ€. Biochemistry, 2002, 41, 15415-15422.	2.5	25

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37	In between â€" Proteomics of dog biological fluids. Journal of Proteomics, 2014, 106, 30-45.	2.4	24
38	Stereoselective reduction of aromatic ketones by a new ketoreductase from Pichia glucozyma. Applied Microbiology and Biotechnology, 2016, 100, 193-201.	3.6	24
39	Surface Plasmon Resonance as a Tool for Ligand Binding Investigation of Engineered GPR17 Receptor, a G Protein Coupled Receptor Involved in Myelination. Frontiers in Chemistry, 2019, 7, 910.	3.6	24
40	Other than IPGâ€DALT: 2â€DE variants. Proteomics, 2010, 10, 586-610.	2.2	23
41	Inhibition of Pancreatic α-amylase by Resveratrol Derivatives: Biological Activity and Molecular Modelling Evidence for Cooperativity between Viniferin Enantiomers. Molecules, 2019, 24, 3225.	3.8	23
42	Enzymatic reduction of acetophenone derivatives with a benzil reductase from Pichia glucozyma (KRED1-Pglu): electronic and steric effects on activity and enantioselectivity. Organic and Biomolecular Chemistry, 2016, 14, 3404-3408.	2.8	21
43	The scaffold protein RACK1 is a target of endocrine disrupting chemicals (EDCs) with important implication in immunity. Toxicology and Applied Pharmacology, 2017, 325, 37-47.	2.8	20
44	Apolipoprotein A†breakdown is induced by thrombolysis in coronary patients. Annals of Medicine, 2007, 39, 306-311.	3.8	19
45	Conformational and dynamics changes induced by bile acids binding to chicken liver bile acid binding protein. Proteins: Structure, Function and Bioinformatics, 2008, 71, 1889-1898.	2.6	18
46	A Novel SEMA3G Mutation in Two Siblings Affected by Syndromic GnRH Deficiency. Neuroendocrinology, 2021, 111, 421-441.	2.5	18
47	Hemolymph proteins: An overview across marine arthropods and molluscs. Journal of Proteomics, 2021, 245, 104294.	2.4	18
48	Molecular Mechanism of Inhibition of the Mitochondrial Carnitine/Acylcarnitine Transporter by Omeprazole Revealed by Proteoliposome Assay, Mutagenesis and Bioinformatics. PLoS ONE, 2013, 8, e82286.	2.5	18
49	Computational and experimental approaches for assessing the interactions between the model calycin β-lactoglobulin and two antibacterial fluoroquinolones. Proteins: Structure, Function and Bioinformatics, 2006, 65, 555-567.	2.6	16
50	Mapping the 5–50-kDa fraction of human amniotic fluid proteins by 2-DE and ESI-MS. Proteomics - Clinical Applications, 2007, 1, 167-175.	1.6	16
51	Unfolding of betaâ€lactoglobulin on the surface of polystyrene nanoparticles: Experimental and computational approaches. Proteins: Structure, Function and Bioinformatics, 2014, 82, 1272-1282.	2.6	16
52	Development of the first in vivo GPR17 ligand through an iterative drug discovery pipeline: A novel disease-modifying strategy for multiple sclerosis. PLoS ONE, 2020, 15, e0231483.	2.5	16
53	Perivascular carotid collar placement induces neointima formation and outward arterial remodeling in mice independent of apolipoprotein E deficiency or Western-type diet feeding. Atherosclerosis, 2007, 195, e112-e124.	0.8	15
54	Raloxifene inhibits matrix metalloproteinases expression and activity in macrophages and smooth muscle cells. Pharmacological Research, 2007, 56, 160-167.	7.1	15

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55	Structural and dynamic roles of permanent water molecules in ligand molecular recognition by chicken liver bile acid binding protein. Journal of Molecular Recognition, 2008, 21, 348-354.	2.1	15
56	SARS-CoV-2 infection among asymptomatic homebound subjects in Milan, Italy. European Journal of Internal Medicine, 2020, 78, 161-163.	2.2	14
57	Simulation of urea-induced protein unfolding: A lesson from bovine $\hat{I}^2$ -lactoglobulin. Journal of Molecular Graphics and Modelling, 2011, 30, 24-30.	2.4	13
58	A proteomic portrait of atherosclerosis. Journal of Proteomics, 2013, 82, 92-112.	2.4	13
59	A promiscuous recognition mechanism between GPR17 and SDF-1: Molecular insights. Cellular Signalling, 2016, 28, 631-642.	3.6	13
60	Design, synthesis, molecular modelling and in vitro cytotoxicity analysis of novel carbamate derivatives as inhibitors of Monoacylglycerol lipase. Bioorganic and Medicinal Chemistry, 2018, 26, 2561-2572.	3.0	13
61	Strategic single point mutation yields a solvent- and salt-stable transaminase from Virgibacillus sp. in soluble form. Scientific Reports, 2018, 8, 16441.	3.3	13
62	Propiconazole is an activator of AHR and causes concentration additive effects with an established AHR ligand. Archives of Toxicology, 2018, 92, 3471-3486.	4.2	13
63	In silico Description of LAT1 Transport Mechanism at an Atomistic Level. Frontiers in Chemistry, 2018, 6, 350.	3.6	13
64	SLC6A14, a Pivotal Actor on Cancer Stage: When Function Meets Structure. SLAS Discovery, 2019, 24, 928-938.	2.7	13
65	Activation of Naturally Occurring Lecithin:Cholesterol Acyltransferase Mutants by a Novel Activator Compound. Journal of Pharmacology and Experimental Therapeutics, 2020, 375, 463-468.	2.5	13
66	Some more about dogs: Proteomics of neglected biological fluids. Journal of Proteomics, 2020, 218, 103724.	2.4	13
67	A Model Structure for the Heterodimer apoA-IMilano–apoA-II Supports Its Peculiar Susceptibility to Proteolysis. Biophysical Journal, 2006, 91, 3043-3049.	0.5	12
68	Proteomics of lung physiopathology. Proteomics, 2008, 8, 5053-5073.	2.2	12
69	In silico prediction and characterization of protein post-translational modifications. Journal of Proteomics, 2016, 134, 65-75.	2.4	12
70	Gender proteomics I. Which proteins in non-sexual organs. Journal of Proteomics, 2018, 178, 7-17.	2.4	12
71	Distinguishing mode of action of compounds inducing craniofacial malformations in zebrafish embryos to support dose-response modeling in combined exposures. Reproductive Toxicology, 2020, 96, 114-127.	2.9	12
72	Proteomics of rat biological fluids â€" The tenth anniversary update. Journal of Proteomics, 2012, 75, 3113-3128.	2.4	10

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73	Recombinant S. cerevisiae expressing Old Yellow Enzymes from non-conventional yeasts: an easy system for selective reduction of activated alkenes. Microbial Cell Factories, 2014, 13, 60.	4.0	10
74	Glatiramer acetate: A complex drug beyond biologics. European Journal of Pharmaceutical Sciences, 2019, 133, 8-14.	4.0	10
75	Energy matters: Mitochondrial proteomics for biomedicine. Proteomics, 2011, 11, 657-674.	2.2	9
76	Neglected markers: Altered serum proteome in murine models of disease. Proteomics, 2012, 12, 691-707.	2.2	9
77	lgG1 conformational behavior: elucidation of the N-glycosylation role via molecular dynamics. Biophysical Journal, 2021, 120, 5355-5370.	0.5	9
78	Using peripheral blood mononuclear cells to determine proteome profiles in human cardiac failure. European Journal of Heart Failure, 2008, 10, 749-757.	7.1	8
79	Structural features and dynamics properties of human apolipoprotein A-I in a model of synthetic HDL. Journal of Molecular Graphics and Modelling, 2009, 28, 305-312.	2.4	8
80	Electrostatics of folded and unfolded bovine $\hat{l}^2$ -lactoglobulin. Amino Acids, 2012, 42, 2019-2030.	2.7	8
81	Set-Up and Validation of a High Throughput Screening Method for Human Monoacylglycerol Lipase (MAGL) Based on a New Red Fluorescent Probe. Molecules, 2019, 24, 2241.	3.8	8
82	rHDL modeling and the anchoring mechanism of LCAT activation. Journal of Lipid Research, 2021, 62, 100006.	4.2	8
83	Detection of Protein Glutathionylation. Methods in Molecular Biology, 2009, 519, 397-415.	0.9	7
84	Structural and dynamic features of apolipoprotein A-I cysteine mutants, Milano and Paris, in synthetic HDL. Journal of Molecular Graphics and Modelling, 2010, 29, 406-414.	2.4	7
85	Inflammatory serum proteome pattern in mice fed a high-fat diet. Amino Acids, 2013, 44, 1001-1008.	2.7	7
86	<i>In Vitro</i> Chronic Administration of ERbeta Selective Ligands and Prostate Cancer Cell Growth: Hypotheses on the Selective Role of 3beta-Adiol in AR-Positive RV1 Cells. BioMed Research International, 2014, 2014, 1-14.	1.9	7
87	Development of an adverse outcome pathway for cranio-facial malformations: A contribution from in silico simulations and in vitro data. Food and Chemical Toxicology, 2020, 140, 111303.	3.6	6
88	Distant Homology Modeling of LCAT and Its Validation through In Silico Targeting and In Vitro and In Vivo Assays. PLoS ONE, 2014, 9, e95044.	2.5	6
89	Molecular Modelling of NONO and SFPQ Dimerization Process and RNA Recognition Mechanism. International Journal of Molecular Sciences, 2022, 23, 7626.	4.1	6
90	Gender proteomics II. Which proteins in sexual organs. Journal of Proteomics, 2018, 178, 18-30.	2.4	5

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91	ApoAâ€I <sub>Milano</sub> from structure to clinical application. Annals of Medicine, 2008, 40, 48-56.	3.8	4
92	All-Purpose Containers? Lipid-Binding Protein – Drug Interactions. PLoS ONE, 2015, 10, e0132096.	2.5	4
93	What if? Mouse proteomics after gene inactivation. Journal of Proteomics, 2019, 199, 102-122.	2.4	3
94	Wards in the keyway: amino acids with anomalous pK as in calycins. Amino Acids, 2012, 43, 2457-2468.	2.7	1
95	Encore – Sex dependency of the proteome. Journal of Proteomics, 2020, 212, 103579.	2.4	1
96	2,4-Furfurylidene-D-sorbitol and its tetra-methyl ether: synthesis, conformational studies, and radical scavenging activity. Arkivoc, 2017, 2016, 50-68.	0.5	1
97	Pharmacokinetics and pharmacodynamics in the newborn. Veterinary Research Communications, 2008, 32, 77-80.	1.6	O
98	Molecular investigation of riboflavin-responsive multiple acyl-CoA dehydrogenase deficiency (RR-MAD) patients. Biochimica Et Biophysica Acta - Bioenergetics, 2010, 1797, 54.	1.0	0
99	Class A GPCRs: a multifaceted reality. Purinergic Signalling, 2011, 7, 279-281.	2.2	O
100	Editorial: A matter of ingredients. Journal of Proteomics, 2018, 178, 1-6.	2.4	0