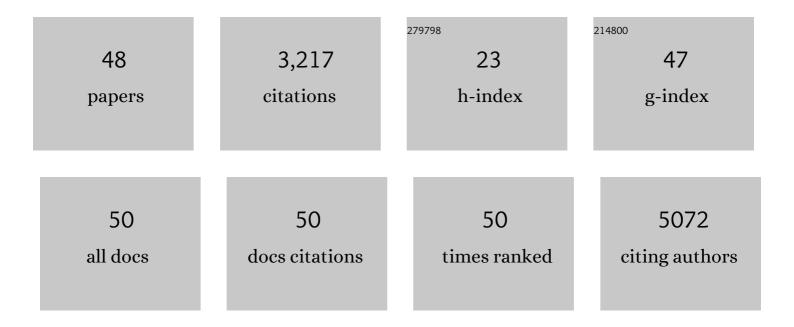
Laura Santucci

List of Publications by Year in descending order

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Ι ΛΗΡΑ SANTUCCI

#	Article	IF	CITATIONS
1	Neutrophil Extracellular Traps Profiles in Patients with Incident Systemic Lupus Erythematosus and Lupus Nephritis. Journal of Rheumatology, 2020, 47, 377-386.	2.0	77
2	Association between maternal omegaâ€3 polyunsaturated fatty acids supplementation and preterm delivery: A proteomic study. FASEB Journal, 2020, 34, 6322-6334.	0.5	5
3	Biological surface properties in extracellular vesicles and their effect on cargo proteins. Scientific Reports, 2019, 9, 13048.	3.3	28
4	Neutrophil Extracellular Traps protein composition is specific for patients with Lupus nephritis and includes methyl-oxidized αenolase (methionine sulfoxide 93). Scientific Reports, 2019, 9, 7934.	3.3	58
5	Proteome of Bovine Mitochondria and Rod Outer Segment Disks: Commonalities and Differences. Journal of Proteome Research, 2018, 17, 918-925.	3.7	14
6	Metabolic Signature of Microvesicles from Umbilical Cord Mesenchymal Stem Cells of Preterm and Term Infants. Proteomics - Clinical Applications, 2018, 12, e1700082.	1.6	26
7	Microvesicles as promising biological tools for diagnosis and therapy. Expert Review of Proteomics, 2018, 15, 801-808.	3.0	28
8	Annexin A1 and Autoimmunity: From Basic Science to Clinical Applications. International Journal of Molecular Sciences, 2018, 19, 1348.	4.1	58
9	Nidogen-1 is a novel extracellular ligand for the NKp44 activating receptor. Oncolmmunology, 2018, 7, e1470730.	4.6	54
10	Urine Proteome Biomarkers in Kidney Diseases. I. Limits, Perspectives, and First Focus on Normal Urine. Biomarker Insights, 2016, 11, BMI.S26229.	2.5	22
11	Urine proteome analysis in Dent's disease shows high selective changes potentially involved in chronic renal damage. Journal of Proteomics, 2016, 130, 26-32.	2.4	9
12	Exosomes from human mesenchymal stem cells conduct aerobic metabolism in term and preterm newborn infants. FASEB Journal, 2016, 30, 1416-1424.	0.5	63
13	Human urinary exosome proteome unveils its aerobic respiratory ability. Journal of Proteomics, 2016, 136, 25-34.	2.4	27
14	Combinatorial Peptide Ligand Library and two dimensional electrophoresis: New frontiers in the study of peritoneal dialysis effluent in pediatric patients. Journal of Proteomics, 2015, 116, 68-80.	2.4	8
15	The human urinary exosome as a potential metabolic effector cargo. Expert Review of Proteomics, 2015, 12, 425-432.	3.0	41
16	Widening and Diversifying the Proteome Capture by Combinatorial Peptide Ligand Libraries via Alcian Blue Dye Binding. Analytical Chemistry, 2015, 87, 4814-4820.	6.5	15
17	2DE Maps in the Discovery of Human Autoimmune Kidney Diseases: The Case of Membranous Glomerulonephritis. Methods in Molecular Biology, 2015, 1243, 127-138.	0.9	2
18	The Latest Advancements in Proteomic Two-dimensional Gel Electrophoresis Analysis Applied to Biological Samples. Methods in Molecular Biology, 2015, 1243, 103-125.	0.9	5

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19	From hundreds to thousands: Widening the normal human Urinome (1). Journal of Proteomics, 2015, 112, 53-62.	2.4	43
20	From hundreds to thousands: Widening the normal human Urinome. Data in Brief, 2014, 1, 25-28.	1.0	44
21	Myelin proteomics: the past, the unexpected and the future. Expert Review of Proteomics, 2014, 11, 345-354.	3.0	10
22	Oxidized albumin. The long way of a protein of uncertain function. Biochimica Et Biophysica Acta - General Subjects, 2013, 1830, 5473-5479.	2.4	43
23	Albumin heterogeneity in low-abundance fluids. The case of urine and cerebro-spinal fluid. Biochimica Et Biophysica Acta - General Subjects, 2013, 1830, 5503-5508.	2.4	9
24	Urinary proteome in a snapshot: normal urine and glomerulonephritis. Journal of Nephrology, 2013, 26, 610-616.	2.0	18
25	Combinatorial ligand libraries as a two-dimensional method for proteome analysis. Journal of Chromatography A, 2013, 1297, 106-112.	3.7	18
26	Patients with primary membranous nephropathy lack auto-antibodies against LDL receptor, the homologue of megalin in human glomeruli. CKJ: Clinical Kidney Journal, 2012, 5, 178-179.	2.9	4
27	"Cheek-to-cheek―urinary proteome profiling via combinatorial peptide ligand libraries: A novel, unexpected elution system. Journal of Proteomics, 2012, 75, 796-805.	2.4	27
28	A blue dive: from â€~blue fingers' to â€~blue silver'. A comparative overview of staining methods for in-gel proteomics. Expert Review of Proteomics, 2012, 9, 627-634.	3.0	22
29	Combinatorial peptide ligand libraries for the analysis of lowâ€expression proteins: Validation for normal urine and definition of a first protein MAP. Proteomics, 2012, 12, 509-515.	2.2	22
30	Protein–protein interaction heterogeneity of plasma apolipoprotein A1 in nephrotic syndrome. Molecular BioSystems, 2011, 7, 659-666.	2.9	8
31	Analysis of the oxido-redox status of plasma proteins. Technology advances for clinical applications. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2011, 879, 1338-1344.	2.3	8
32	Proteome profile of peritoneal effluents in children on glucose- or icodextrin-based peritoneal dialysis. Nephrology Dialysis Transplantation, 2011, 26, 308-316.	0.7	9
33	2D-electrophoresis and the urine proteome map: Where do we stand?. Journal of Proteomics, 2010, 73, 829-844.	2.4	57
34	Autoimmunity in Membranous Nephropathy Targets Aldose Reductase and SOD2. Journal of the American Society of Nephrology: JASN, 2010, 21, 507-519.	6.1	190
35	Combinatorial peptide ligand libraries for urine proteome analysis: Investigation of different elution systems. Electrophoresis, 2009, 30, 2405-2411.	2.4	95
36	New iodoâ€acetamido cyanines for labeling cysteine thiol residues. A strategy for evaluating plasma proteins and their oxidoâ€redox status. Proteomics, 2009, 9, 460-469.	2.2	27

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37	The oxido-redox potential of albumin. Journal of Proteomics, 2009, 73, 188-195.	2.4	41
38	Determination of the oxido-redox status of plasma albumin in hemodialysis patients. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 864, 29-37.	2.3	18
39	Proteins and protein fragments in nephrotic syndrome: Clusters, specificity and mechanisms. Proteomics - Clinical Applications, 2008, 2, 956-963.	1.6	14
40	Highâ€resolution 2â€DE for resolving proteins, protein adducts and complexes in plasma. Electrophoresis, 2008, 29, 682-694.	2.4	10
41	Proteomics of Plasma and Urine in Primary Nephrotic Syndrome in Children. , 2008, 160, 17-28.		12
42	ANTI-ATP SYNTHASE Î ² -CHAIN AUTOANTIBODIES. , 2007, , 547-552.		2
43	Protracted remission of proteinuria after combined therapy with plasmapheresis and anti-CD20 antibodies/cyclophosphamide in a child with oligoclonal IgM and glomerulosclerosis. Pediatric Nephrology, 2007, 22, 1953-1956.	1.7	3
44	Transitions of serum albumin in patients with glomerulosclerosis â€~in vivo' characterization by electrophoretic titration curves. Electrophoresis, 2006, 27, 2960-2969.	2.4	12
45	Repetitive Fragmentation Products of Albumin and α1-Antitrypsin in Glomerular Diseases Associated with Nephrotic Syndrome. Journal of the American Society of Nephrology: JASN, 2006, 17, 3139-3148.	6.1	139
46	Circulating anti-actin and anti-ATP synthase antibodies identify a sub-set of patients with idiopathic nephrotic syndrome. Clinical and Experimental Immunology, 2005, 141, 491-499.	2.6	37
47	Proteomic Analysis of Erythrocyte Membranes by Soft Immobiline Gels Combined with Differential Protein Extraction. Journal of Proteome Research, 2005, 4, 1304-1309.	3.7	47
48	Blue silver: A very sensitive colloidal Coomassie G-250 staining for proteome analysis. Electrophoresis, 2004, 25, 1327-1333.	2.4	1,686