Paul L Fox

List of Publications by Year in descending order

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		23567	33894
120	10,315	58	99
papers	citations	h-index	g-index
121	121	121	13853
all docs	docs citations	times ranked	citing authors

ΡΑΠΙ ΓΟΧ

#	Article	IF	CITATIONS
1	Translational control by the $3\hat{\epsilon}^2$ -UTR: the ends specify the means. Trends in Biochemical Sciences, 2003, 28, 91-98.	7.5	448
2	Identification of a Novel Family of Oxidized Phospholipids That Serve as Ligands for the Macrophage Scavenger Receptor CD36. Journal of Biological Chemistry, 2002, 277, 38503-38516.	3.4	389
3	Apolipoprotein A-I is a selective target for myeloperoxidase-catalyzed oxidation and functional impairment in subjects with cardiovascular disease. Journal of Clinical Investigation, 2004, 114, 529-541.	8.2	333
4	An abundant dysfunctional apolipoprotein A1 in human atheroma. Nature Medicine, 2014, 20, 193-203.	30.7	316
5	Regulated Release of L13a from the 60S Ribosomal Subunit as A Mechanism of Transcript-Specific Translational Control. Cell, 2003, 115, 187-198.	28.9	288
6	TGF-β-mediated phosphorylation of hnRNP E1 induces EMT via transcript-selective translational induction of Dab2 and ILEI. Nature Cell Biology, 2010, 12, 286-293.	10.3	269
7	Preferential Iron Trafficking Characterizes Glioblastoma Stem-like Cells. Cancer Cell, 2015, 28, 441-455.	16.8	249
8	Role of Hypoxia-inducible Factor-1 in Transcriptional Activation of Ceruloplasmin by Iron Deficiency. Journal of Biological Chemistry, 2000, 275, 21048-21054.	3.4	248
9	Aminoacylâ€ŧRNA synthetases in medicine and disease. EMBO Molecular Medicine, 2013, 5, 332-343.	6.9	234
10	A stress-responsive RNA switch regulates VEGFA expression. Nature, 2009, 457, 915-919.	27.8	231
11	Myeloperoxidase, paraoxonase-1, and HDL form a functional ternary complex. Journal of Clinical Investigation, 2013, 123, 3815-3828.	8.2	226
12	Noncanonical Function of Glutamyl-Prolyl-tRNA Synthetase. Cell, 2004, 119, 195-208.	28.9	224
13	Role of Ceruloplasmin in Cellular Iron Uptake. Science, 1998, 279, 714-717.	12.6	193
14	Cancer Stem Cell-Secreted Macrophage Migration Inhibitory Factor Stimulates Myeloid Derived Suppressor Cell Function and Facilitates Glioblastoma Immune Evasion. Stem Cells, 2016, 34, 2026-2039.	3.2	189
15	The GAIT system: a gatekeeper of inflammatory gene expression. Trends in Biochemical Sciences, 2009, 34, 324-331.	7.5	187
16	Programmed Translational Readthrough Generates Antiangiogenic VEGF-Ax. Cell, 2014, 157, 1605-1618.	28.9	184
17	Regulation of production of a platelet-derived growth factor-like protein by cultured bovine aortic endothelial cells. Journal of Cellular Physiology, 1984, 121, 298-308.	4.1	163
18	Aminoacyl-tRNA synthetases as therapeutic targets. Nature Reviews Drug Discovery, 2019, 18, 629-650.	46.4	162

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19	Target-Selective Protein S-Nitrosylation by Sequence Motif Recognition. Cell, 2014, 159, 623-634.	28.9	158
20	Quantitative H2S-mediated protein sulfhydration reveals metabolic reprogramming during the integrated stress response. ELife, 2015, 4, e10067.	6.0	154
21	Activation of Cytosolic Phospholipase A by Basic Fibroblast Growth Factor via a p42 Mitogen-activated Protein Kinase-dependent Phosphorylation Pathway in Endothelial Cells. Journal of Biological Chemistry, 1995, 270, 2360-2366.	3.4	150
22	The Oxidation of Lipoproteins by Monocytes-Macrophages. Journal of Biological Chemistry, 1999, 274, 25959-25962.	3.4	148
23	Ceruloplasmin and cardiovascular disease. Free Radical Biology and Medicine, 2000, 28, 1735-1744.	2.9	144
24	The copper-iron chronicles: the story of an intimate relationship. BioMetals, 2003, 16, 9-40.	4.1	141
25	Unexpected role of ceruloplasmin in intestinal iron absorption. Cell Metabolism, 2005, 2, 309-319.	16.2	133
26	EPRS is a critical mTORC1–S6K1 effector that influences adiposity in mice. Nature, 2017, 542, 357-361.	27.8	130
27	DAPK-ZIPK-L13a Axis Constitutes a Negative-Feedback Module Regulating Inflammatory Gene Expression. Molecular Cell, 2008, 32, 371-382.	9.7	128
28	T cell–intrinsic ASC critically promotes TH17-mediated experimental autoimmune encephalomyelitis. Nature Immunology, 2016, 17, 583-592.	14.5	127
29	Ceruloplasmin Ferroxidase Activity Stimulates Cellular Iron Uptake by a Trivalent Cation-specific Transport Mechanism. Journal of Biological Chemistry, 1999, 274, 1116-1123.	3.4	125
30	GAPDH regulates cellular heme insertion into inducible nitric oxide synthase. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 18004-18009.	7.1	125
31	Macromolecular complexes as depots for releasable regulatory proteins. Trends in Biochemical Sciences, 2007, 32, 158-164.	7.5	117
32	Two-Site Phosphorylation of EPRS Coordinates Multimodal Regulation of Noncanonical Translational Control Activity. Molecular Cell, 2009, 35, 164-180.	9.7	115
33	Transcript-Selective Translational Silencing by Gamma Interferon Is Directed by a Novel Structural Element in the Ceruloplasmin mRNA 3′ Untranslated Region. Molecular and Cellular Biology, 2003, 23, 1509-1519.	2.3	113
34	Repression of VEGFA by CA-rich element-binding microRNAs is modulated by hnRNP L. EMBO Journal, 2011, 30, 1324-1334.	7.8	111
35	WHEP Domains Direct Noncanonical Function of Glutamyl-Prolyl tRNA Synthetase in Translational Control of Gene Expression. Molecular Cell, 2008, 29, 679-690.	9.7	110
36	Inflammation mobilizes copper metabolism to promote colon tumorigenesis via an IL-17-STEAP4-XIAP axis. Nature Communications, 2020, 11, 900.	12.8	108

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37	Structure, oxidant activity, and cardiovascular mechanisms of human ceruloplasmin. Life Sciences, 1995, 56, 1749-1758.	4.3	107
38	IRAK-M mediates Toll-like receptor/IL-1R-induced NFκB activation and cytokine production. EMBO Journal, 2013, 32, 583-596.	7.8	103
39	Function and Distribution of Apolipoprotein A1 in the Artery Wall Are Markedly Distinct From Those in Plasma. Circulation, 2013, 128, 1644-1655.	1.6	98
40	A post-transcriptional pathway represses monocyte VEGF-A expression and angiogenic activity. EMBO Journal, 2007, 26, 3360-3372.	7.8	96
41	IL-17-receptor-associated adaptor Act1 directly stabilizes mRNAs to mediate IL-17 inflammatory signaling. Nature Immunology, 2018, 19, 354-365.	14.5	91
42	L13a Blocks 48S Assembly: Role of a General Initiation Factor in mRNA-Specific Translational Control. Molecular Cell, 2007, 25, 113-126.	9.7	88
43	Coding Region Polyadenylation Generates a Truncated tRNA Synthetase that Counters Translation Repression. Cell, 2012, 149, 88-100.	28.9	87
44	Regulation and dysregulation of 3′UTR-mediated translational control. Current Opinion in Genetics and Development, 2013, 23, 29-34.	3.3	87
45	Role of Ceruloplasmin in Macrophage Iron Efflux during Hypoxia. Journal of Biological Chemistry, 2003, 278, 44018-44024.	3.4	86
46	Site-specific Nitration of Apolipoprotein A-I at Tyrosine 166 Is Both Abundant within Human Atherosclerotic Plaque and Dysfunctional. Journal of Biological Chemistry, 2014, 289, 10276-10292.	3.4	84
47	HuR Is Required for IL-17–Induced Act1-Mediated CXCL1 and CXCL5 mRNA Stabilization. Journal of Immunology, 2013, 191, 640-649.	0.8	83
48	Profilin-1 phosphorylation directs angiocrine expression and glioblastoma progression throughÂHIF-1α accumulation. Nature Cell Biology, 2014, 16, 445-456.	10.3	83
49	Infection-specific phosphorylation of glutamyl-prolyl tRNA synthetase induces antiviral immunity. Nature Immunology, 2016, 17, 1252-1262.	14.5	76
50	Membrane microviscosity regulates endothelial cell motility. Nature Cell Biology, 2002, 4, 894-900.	10.3	75
51	Protection of Extraribosomal RPL13a by GAPDH and Dysregulation by S-Nitrosylation. Molecular Cell, 2012, 47, 656-663.	9.7	74
52	HOXA9 Methylation by PRMT5 Is Essential for Endothelial Cell Expression of Leukocyte Adhesion Molecules. Molecular and Cellular Biology, 2012, 32, 1202-1213.	2.3	72
53	MyD88-dependent interplay between myeloid and endothelial cells in the initiation and progression of obesity-associated inflammatory diseases. Journal of Experimental Medicine, 2014, 211, 887-907.	8.5	70
54	Inactivation of the Enzyme GSK3α by the Kinase IKKi Promotes AKT-mTOR Signaling Pathway that Mediates Interleukin-1-Induced Th17 Cell Maintenance. Immunity, 2012, 37, 800-812.	14.3	69

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55	Ceruloplasmin Enhances Smooth Muscle Cell- and Endothelial Cell-mediated Low Density Lipoprotein Oxidation by a Superoxide-dependent Mechanism. Journal of Biological Chemistry, 1996, 271, 14773-14778.	3.4	67
56	Stimulus-dependent phosphorylation of profilin-1 inÂangiogenesis. Nature Cell Biology, 2012, 14, 1046-1056.	10.3	66
57	Polarization of Plasma Membrane Microviscosity during Endothelial Cell Migration. Developmental Cell, 2004, 6, 29-41.	7.0	65
58	Palmitoylation of Caveolin-1 in Endothelial Cells Is Post-translational but Irreversible. Journal of Biological Chemistry, 2001, 276, 15776-15782.	3.4	63
59	Cx26 drives self-renewal in triple-negative breast cancer via interaction with NANOG and focal adhesion kinase. Nature Communications, 2018, 9, 578.	12.8	60
60	Translational Silencing of Ceruloplasmin Requires the Essential Elements of mRNA Circularization: Poly(A) Tail, Poly(A)-Binding Protein, and Eukaryotic Translation Initiation Factor 4G. Molecular and Cellular Biology, 2001, 21, 6440-6449.	2.3	58
61	Spatial Coordination of Actin Polymerization and ILK–Akt2 Activity during Endothelial Cell Migration. Developmental Cell, 2009, 16, 661-674.	7.0	58
62	IRAKMâ€Mincle axis links cell death to inflammation: Pathophysiological implications for chronic alcoholic liver disease. Hepatology, 2016, 64, 1978-1993.	7.3	55
63	Ceruloplasmin Copper Induces Oxidant Damage by a Redox Process Utilizing Cell-Derived Superoxide as Reductantâ€. Biochemistry, 1998, 37, 14222-14229.	2.5	54
64	Delayed Translational Silencing of Ceruloplasmin Transcript in Gamma Interferon-Activated U937 Monocytic Cells: Role of the 3′ Untranslated Region. Molecular and Cellular Biology, 1999, 19, 6898-6905.	2.3	54
65	Phosphorylation of glutamyl-prolyl tRNA synthetase by cyclin-dependent kinase 5 dictates transcript-selective translational control. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 1415-1420.	7.1	54
66	Clinical and Genetic Association of Serum Ceruloplasmin With Cardiovascular Risk. Arteriosclerosis, Thrombosis, and Vascular Biology, 2012, 32, 516-522.	2.4	54
67	The HILDA Complex Coordinates a Conditional Switch in the 3′-Untranslated Region of the VEGFA mRNA. PLoS Biology, 2013, 11, e1001635.	5.6	51
68	Expression of Sorsby's Fundus Dystrophy Mutations in Human Retinal Pigment Epithelial Cells Reduces Matrix Metalloproteinase Inhibition and May Promote Angiogenesis. Journal of Biological Chemistry, 2002, 277, 13394-13400.	3.4	50
69	Protein Kinase Cδ–Dependent Phosphorylation of Syndecan-4 Regulates Cell Migration. Circulation Research, 2005, 97, 674-681.	4.5	49
70	Myo1c facilitates G-actin transport to the leading edge of migrating endothelial cells. Journal of Cell Biology, 2012, 198, 47-55.	5.2	48
71	Antiangiogenic VEGF-Ax: A New Participant in Tumor Angiogenesis. Cancer Research, 2015, 75, 2765-2769.	0.9	48
72	Protective role of macrophage-derived ceruloplasmin in inflammatory bowel disease. Gut, 2013, 62, 209-219.	12.1	47

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73	The <scp>GAIT</scp> translational control system. Wiley Interdisciplinary Reviews RNA, 2018, 9, e1441.	6.4	46
74	Reactive Oxygen Species Regulate Ceruloplasmin by a Novel mRNA Decay Mechanism Involving Its 3′-Untranslated Region. Journal of Biological Chemistry, 2009, 284, 1873-1883.	3.4	44
75	The Critical Role of IL-1 Receptor-Associated Kinase 4-Mediated NF-κB Activation in Modified Low-Density Lipoprotein-Induced Inflammatory Gene Expression and Atherosclerosis. Journal of Immunology, 2011, 186, 2871-2880.	0.8	44
76	A CC′ Loop Decoy Peptide Blocks the Interaction Between Act1 and IL-17RA to Attenuate IL-17– and IL-25–Induced Inflammation. Science Signaling, 2011, 4, ra72.	3.6	44
77	Heterotrimeric GAIT Complex Drives Transcript-Selective Translation Inhibition in Murine Macrophages. Molecular and Cellular Biology, 2012, 32, 5046-5055.	2.3	41
78	Anemia and impaired stress-induced erythropoiesis in aceruloplasminemic mice. Blood Cells, Molecules, and Diseases, 2004, 33, 346-355.	1.4	40
79	Pellino 2 Is critical for Toll-like Receptor/Interleukin-1 Receptor (TLR/IL-1R)-mediated Post-transcriptional Control. Journal of Biological Chemistry, 2012, 287, 25686-25695.	3.4	39
80	Oxidative stress inhibits caveolin-1 palmitoylation and trafficking in endothelial cells. Biochemical Journal, 2002, 361, 681-688.	3.7	37
81	Multisite Phosphorylation of S6K1 Directs a Kinase Phospho-code that Determines Substrate Selection. Molecular Cell, 2019, 73, 446-457.e6.	9.7	36
82	Regulation of ceruloplasmin in human hepatic cells by redox active copper: identification of a novel AP-1 site in the ceruloplasmin gene. Biochemical Journal, 2007, 402, 135-141.	3.7	34
83	Human Colon Tumors Express a Dominant-Negative Form ofÂSIGIRR That Promotes Inflammation and Colitis-Associated Colon Cancer in Mice. Gastroenterology, 2015, 149, 1860-1871.e8.	1.3	33
84	Interplay between miR-574-3p and hnRNP L regulates VEGFA mRNA translation and tumorigenesis. Nucleic Acids Research, 2017, 45, 7950-7964.	14.5	33
85	Dual Role of Insulin in Transcriptional Regulation of the Acute Phase Reactant Ceruloplasmin. Journal of Biological Chemistry, 2002, 277, 27903-27911.	3.4	32
86	Letâ€7aâ€regulated translational readthrough of mammalian <i> <scp>AGO</scp> 1 </i> generates a micro <scp>RNA</scp> pathway inhibitor. EMBO Journal, 2019, 38, e100727.	7.8	30
87	Evolution of Function of a Fused Metazoan tRNA Synthetase. Molecular Biology and Evolution, 2011, 28, 437-447.	8.9	28
88	Non-catalytic Regulation of Gene Expression by Aminoacyl-tRNA Synthetases. Topics in Current Chemistry, 2013, 344, 167-187.	4.0	28
89	Cancer-predicting transcriptomic and epigenetic signatures revealed for ulcerative colitis in patient-derived epithelial organoids. Oncotarget, 2018, 9, 28717-28730.	1.8	28
90	3-Dimensional architecture of the human multi-tRNA synthetase complex. Nucleic Acids Research, 2020, 48, 8740-8754.	14.5	27

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91	IRAK2 directs stimulus-dependent nuclear export of inflammatory mRNAs. ELife, 2017, 6, .	6.0	22
92	IL-17–induced HIF1α drives resistance to anti–PD-L1 via fibroblast-mediated immune exclusion. Journal of Experimental Medicine, 2022, 219, .	8.5	21
93	The RNA-Binding Protein HuR Posttranscriptionally Regulates IL-2 Homeostasis and CD4+ Th2 Differentiation. ImmunoHorizons, 2017, 1, 109-123.	1.8	20
94	Regulation of macrophage ceruloplasmin gene expression: one paradigm of 3'-UTR-mediated translational control. Molecules and Cells, 2005, 20, 167-72.	2.6	20
95	Condensin II and GAIT complexes cooperate to restrict LINE-1 retrotransposition in epithelial cells. PLoS Genetics, 2017, 13, e1007051.	3.5	19
96	Origin and Evolution of Glutamyl-prolyl tRNA Synthetase WHEP Domains Reveal Evolutionary Relationships within Holozoa. PLoS ONE, 2014, 9, e98493.	2.5	19
97	Computational Modeling and Analysis of Iron Release from Macrophages. PLoS Computational Biology, 2014, 10, e1003701.	3.2	15
98	Citric acid cycle and the origin of MARS. Trends in Biochemical Sciences, 2013, 38, 222-228.	7.5	14
99	Aminoacyl-tRNA synthetases of the multi-tRNA synthetase complex and their role in tumorigenesis. Translational Oncology, 2022, 19, 101392.	3.7	13
100	Ceruloplasmin has two nearly identical sites that bind myeloperoxidase. Biochemical and Biophysical Research Communications, 2014, 453, 722-727.	2.1	12
101	Unexpected metabolic function of a tRNA synthetase. Cell Cycle, 2017, 16, 2239-2240.	2.6	12
102	Metabolic origin of the fused aminoacyl-tRNA synthetase, glutamyl-prolyl-tRNA synthetase. Journal of Biological Chemistry, 2018, 293, 19148-19156.	3.4	11
103	Protein S-Nitrosylation of Human Cytomegalovirus pp71 Inhibits Its Ability To Limit STING Antiviral Responses. Journal of Virology, 2020, 94, .	3.4	10
104	Translational Control Mechanisms in Angiogenesis and Vascular Biology. Current Atherosclerosis Reports, 2015, 17, 506.	4.8	9
105	Experimental approaches for investigation of aminoacyl tRNA synthetase phosphorylation. Methods, 2017, 113, 72-82.	3.8	8
106	Discovery and investigation of the GAIT translational control system. Rna, 2015, 21, 615-618.	3.5	7
107	Structural control of caspase-generated glutamyl-tRNA synthetase by appended noncatalytic WHEP domains. Journal of Biological Chemistry, 2018, 293, 8843-8860.	3.4	7
108	Bidirectional Tumor-Promoting Activities of Macrophage Ezrin. International Journal of Molecular Sciences, 2020, 21, 7716.	4.1	7

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109	Aminoacyl-tRNA synthetases in cell signaling. The Enzymes, 2020, 48, 243-275.	1.7	6
110	Cotranslational interaction of human EBP50 and ezrin overcomes masked binding site during complex assembly. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	6
111	A truncated tRNA synthetase directs a "translational trickle―of gene expression. Cell Cycle, 2012, 11, 1868-1869.	2.6	4
112	An optimized protocol for in vitro and in cellulo structural determination of the multi-tRNA synthetase complex by cross-linking mass spectrometry. STAR Protocols, 2022, 3, 101201.	1.2	4
113	The zinc-binding domain of mammalian prolyl-tRNA synthetase is indispensable for catalytic activity and organism viability. IScience, 2021, 24, 102215.	4.1	3
114	Mix 'n' match metalloproteins. Blood, 2004, 103, 4378-4379.	1.4	0
115	Screening of CRISPR-Cas9-generated point mutant mice using MiSeq and locked nucleic acid probe PCR. STAR Protocols, 2021, 2, 100785.	1.2	0
116	Translation Inhibition of Vascular Endothelial Growth Factor mRNA by the GAIT Translational Silencing Complex. FASEB Journal, 2006, 20, A537.	0.5	0
117	Ribosomal protein L13a inhibits translation by blocking the formation of 80S complex on the GAIT element containing mRNA: Dependence on the translation initiation factor eIF4G. FASEB Journal, 2006, 20, A108.	0.5	0
118	Serine phosphorylation of the linker domain of bifunctional glutamylâ€prolyl tRNA synthetase is critical for transcriptâ€specific translational silencing. FASEB Journal, 2006, 20, A496.	0.5	0
119	Cyclinâ€dependent kinase 5 mediated phosphorylation of GluProRS induces translational silencing of inflammatory gene expression. FASEB Journal, 2008, 22, 638.2.	0.5	0
120	Metabolic Origin of the Fused Aminoacyl tRNA Synthetase, Glutamylâ€Prolyl tRNA Synthetase (EPRS). FASEB Journal, 2019, 33, 351.2.	0.5	0