

Thomas Nyström

List of Publications by Year in descending order

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Version: 2024-02-01

73
papers

7,365
citations

66234

42
h-index

88477

70
g-index

160
all docs

160
docs citations

160
times ranked

7315
citing authors

#	ARTICLE	IF	CITATIONS
1	Large organellar changes occur during mild heat shock in yeast. <i>Journal of Cell Science</i> , 2022, 135, .	1.2	16
2	Correlative single-molecule fluorescence barcoding of gene regulation in <i>Saccharomyces cerevisiae</i> . <i>Methods</i> , 2021, 193, 62-67.	1.9	8
3	Differential role of cytosolic Hsp70s in longevity assurance and protein quality control. <i>PLoS Genetics</i> , 2021, 17, e1008951.	1.5	12
4	An Hsp90 co-chaperone links protein folding and degradation and is part of a conserved protein quality control. <i>Cell Reports</i> , 2021, 35, 109328.	2.9	15
5	Genome-wide imaging screen uncovers molecular determinants of arsenite-induced protein aggregation and toxicity. <i>Journal of Cell Science</i> , 2021, 134, .	1.2	11
6	Comparison of endogenously expressed fluorescent protein fusions behaviour for protein quality control and cellular ageing research. <i>Scientific Reports</i> , 2021, 11, 12819.	1.6	8
7	Nuclear envelope budding is a response to cellular stress. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	28
8	FMN reduces Amyloid- β toxicity in yeast by regulating redox status and cellular metabolism. <i>Nature Communications</i> , 2020, 11, 867.	5.8	50
9	Hitchhiking on vesicles: a way to harness age-related proteopathies?. <i>FEBS Journal</i> , 2020, 287, 5068-5079.	2.2	4
10	Peroxiredoxin promotes longevity and H ₂ O ₂ -resistance in yeast through redox-modulation of protein kinase A. <i>ELife</i> , 2020, 9, .	2.8	20
11	Syntaxin 5 Is Required for the Formation and Clearance of Protein Inclusions during Proteostatic Stress. <i>Cell Reports</i> , 2019, 28, 2096-2110.e8.	2.9	30
12	Studying Spatial Protein Quality Control, Proteopathies, and Aging Using Different Model Misfolding Proteins in <i>S. cerevisiae</i> . <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 249.	1.4	28
13	Mitochondrial Translation Efficiency Controls Cytoplasmic Protein Homeostasis. <i>Cell Metabolism</i> , 2018, 27, 1309-1322.e6.	7.2	85
14	Restricted access: spatial sequestration of damaged proteins during stress and aging. <i>EMBO Reports</i> , 2017, 18, 377-391.	2.0	104
15	The Upsides and Downsides of Organelle Interconnectivity. <i>Cell</i> , 2017, 169, 24-34.	13.5	82
16	Differential effects of soluble and aggregating polyQ proteins on cytotoxicity and type-1 myosin-dependent endocytosis in yeast. <i>Scientific Reports</i> , 2017, 7, 11328.	1.6	6
17	Role of the ribosomal quality control machinery in nucleocytoplasmic translocation of polyQ-expanded huntingtin exon-1. <i>Biochemical and Biophysical Research Communications</i> , 2017, 493, 708-717.	1.0	17
18	How and why do toxic conformers of aberrant proteins accumulate during ageing?. <i>Essays in Biochemistry</i> , 2017, 61, 317-324.	2.1	18

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19	A genome-wide imaging-based screening to identify genes involved in synphilin-1 inclusion formation in <i>Saccharomyces cerevisiae</i> . <i>Scientific Reports</i> , 2016, 6, 30134.	1.6	12
20	Asymmetric Inheritance of Aggregated Proteins and Age Reset in Yeast Are Regulated by Vac17-Dependent Vacuolar Functions. <i>Cell Reports</i> , 2016, 16, 826-838.	2.9	66
21	Lifespan Control by Redox-Dependent Recruitment of Chaperones to Misfolded Proteins. <i>Cell</i> , 2016, 166, 140-151.	13.5	120
22	Spatial sequestration and detoxification of Huntingtin by the ribosome quality control complex. <i>ELife</i> , 2016, 5, .	2.8	57
23	Selective protein degradation ensures cellular longevity. <i>ELife</i> , 2016, 5, .	2.8	0
24	The dual role of a yeast metacaspase: What doesn't kill you makes you stronger. <i>BioEssays</i> , 2015, 37, 525-531.	1.2	32
25	Stress Granule-Defective Mutants Deregulate Stress Responsive Transcripts. <i>PLoS Genetics</i> , 2014, 10, e1004763.	1.5	40
26	Essential Genetic Interactors of SIR2 Required for Spatial Sequestration and Asymmetrical Inheritance of Protein Aggregates. <i>PLoS Genetics</i> , 2014, 10, e1004539.	1.5	73
27	Protein quality control in time and space - links to cellular aging. <i>FEMS Yeast Research</i> , 2014, 14, 40-48.	1.1	32
28	The good and the bad of being connected: the integrons of aging. <i>Current Opinion in Cell Biology</i> , 2014, 26, 107-112.	2.6	115
29	The mystery of aging and rejuvenation—a budding topic. <i>Current Opinion in Microbiology</i> , 2014, 18, 61-67.	2.3	62
30	Life-span extension by a metacaspase in the yeast <i>Saccharomyces cerevisiae</i> . <i>Science</i> , 2014, 344, 1389-1392.	6.0	113
31	Opposing roles of Ubp3-dependent deubiquitination regulate replicative life span and heat resistance. <i>EMBO Journal</i> , 2014, 33, 747-761.	3.5	49
32	Aging: Filtering Out Bad Mitochondria. <i>Current Biology</i> , 2013, 23, R1037-R1039.	1.8	6
33	A Proximal Promoter Element Required for Positive Transcriptional Control by Guanosine Tetraphosphate and DksA Protein during the Stringent Response. <i>Journal of Biological Chemistry</i> , 2013, 288, 21055-21064.	1.6	46
34	Deletion of the mitochondrial Pim1/Lon protease in yeast results in accelerated aging and impairment of the proteasome. <i>Free Radical Biology and Medicine</i> , 2013, 56, 9-16.	1.3	62
35	Enhancing protein disaggregation restores proteasome activity in aged cells. <i>Aging</i> , 2013, 5, 802-812.	1.4	75
36	Peroxiredoxins, gerontogenes linking aging to genome instability and cancer. <i>Genes and Development</i> , 2012, 26, 2001-2008.	2.7	84

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37	The Role of Mitochondria in the Aging Processes of Yeast. <i>Sub-Cellular Biochemistry</i> , 2011, 57, 55-78.	1.0	43
38	Segregation of Protein Aggregates Involves Actin and the Polarity Machinery. <i>Cell</i> , 2011, 147, 959-961.	13.5	71
39	Absence of Mitochondrial Translation Control Proteins Extends Life Span by Activating Sirtuin-Dependent Silencing. <i>Molecular Cell</i> , 2011, 42, 390-400.	4.5	74
40	Life Span Extension and H ₂ O ₂ Resistance Elicited by Caloric Restriction Require the Peroxiredoxin Tsa1 in <i>Saccharomyces cerevisiae</i> . <i>Molecular Cell</i> , 2011, 43, 823-833.	4.5	93
41	Spatial protein quality control and the evolution of lineage-specific ageing. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 71-75.	1.8	26
42	Sir2-dependent asymmetric segregation of damaged proteins in <i>ubp10</i> null mutants is independent of genomic silencing. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2010, 1803, 630-638.	1.9	20
43	The Polarisome Is Required for Segregation and Retrograde Transport of Protein Aggregates. <i>Cell</i> , 2010, 140, 257-267.	13.5	305
44	Increased RNA polymerase availability directs resources towards growth at the expense of maintenance. <i>EMBO Journal</i> , 2009, 28, 2209-2219.	3.5	45
45	Identical, Independent, and Opposing Roles of ppGpp and DksA in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2007, 189, 5193-5202.	1.0	144
46	A Bacterial Kind of Aging. <i>PLoS Genetics</i> , 2007, 3, e224.	1.5	60
47	Sir2p-dependent protein segregation gives rise to a superior reactive oxygen species management in the progeny of <i>Saccharomyces cerevisiae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 10877-10881.	3.3	115
48	Accelerated aging and failure to segregate damaged proteins in Sir2 mutants can be suppressed by overproducing the protein aggregation-remodeling factor Hsp104p. <i>Genes and Development</i> , 2007, 21, 2410-2421.	2.7	328
49	Conditional and replicative senescence in <i>Escherichia coli</i> . <i>Current Opinion in Microbiology</i> , 2006, 9, 612-618.	2.3	45
50	Oxidative Damage and Cellular Senescence: Lessons from Bacteria and Yeast. , 2006, , 473-484.		1
51	Oxidation of Bacterial Proteome in Response to Starvation. <i>Methods of Biochemical Analysis</i> , 2005, 49, 89-95.	0.2	4
52	Role of oxidative carbonylation in protein quality control and senescence. <i>EMBO Journal</i> , 2005, 24, 1311-1317.	3.5	683
53	Differential Roles of the Universal Stress Proteins of <i>Escherichia coli</i> in Oxidative Stress Resistance, Adhesion, and Motility. <i>Journal of Bacteriology</i> , 2005, 187, 6265-6272.	1.0	299
54	Growth versus maintenance: a trade-off dictated by RNA polymerase availability and sigma factor competition?. <i>Molecular Microbiology</i> , 2004, 54, 855-862.	1.2	200

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55	Stationary-Phase Physiology. <i>Annual Review of Microbiology</i> , 2004, 58, 161-181.	2.9	313
56	The oncogenic RAS2val19 mutation locks respiration, independently of PKA, in a mode prone to generate ROS. <i>EMBO Journal</i> , 2003, 22, 3337-3345.	3.5	101
57	Nonculturable bacteria: programmed survival forms or cells at death's door?. <i>BioEssays</i> , 2003, 25, 204-211.	1.2	75
58	Conditional senescence in bacteria: death of the immortals. <i>Molecular Microbiology</i> , 2003, 48, 17-23.	1.2	99
59	Differential oxidative damage and expression of stress defence regulons in culturable and non-culturable <i>Escherichia coli</i> cells. <i>EMBO Reports</i> , 2003, 4, 400-404.	2.0	156
60	Asymmetric Inheritance of Oxidatively Damaged Proteins During Cytokinesis. <i>Science</i> , 2003, 299, 1751-1753.	6.0	598
61	The bacterial universal stress protein: function and regulation. <i>Current Opinion in Microbiology</i> , 2003, 6, 140-145.	2.3	438
62	Underproduction of σ^{70} Mimics a Stringent Response. <i>Journal of Biological Chemistry</i> , 2003, 278, 968-973.	1.6	43
63	The Role of the Alarmone (p)ppGpp in σ^{70} Competition for Core RNA Polymerase. <i>Journal of Biological Chemistry</i> , 2003, 278, 1494-1503.	1.6	100
64	Regulation of sigma factor competition by the alarmone ppGpp. <i>Genes and Development</i> , 2002, 16, 1260-1270.	2.7	299
65	Translational fidelity, protein oxidation, and senescence: lessons from bacteria. <i>Ageing Research Reviews</i> , 2002, 1, 693-703.	5.0	52
66	Bacterial senescence: protein oxidation in non-proliferating cells is dictated by the accuracy of the ribosomes. <i>EMBO Journal</i> , 2001, 20, 5280-5289.	3.5	156
67	Not quite dead enough: on bacterial life, culturability, senescence, and death. <i>Archives of Microbiology</i> , 2001, 176, 159-164.	1.0	130
68	Emergency derepression: stringency allows RNA polymerase to override negative control by an active repressor. <i>Molecular Microbiology</i> , 2000, 35, 435-443.	1.2	51
69	Expression and role of the universal stress protein, UspA, of <i>Escherichia coli</i> during growth arrest. <i>Molecular Microbiology</i> , 1994, 11, 537-544.	1.2	223
70	The glucose-starvation stimulon of <i>Escherichia coli</i> : induced and repressed synthesis of enzymes of central metabolic pathways and role of acetyl phosphate in gene expression and starvation survival. <i>Molecular Microbiology</i> , 1994, 12, 833-843.	1.2	103
71	Cloning, mapping and nucleotide sequencing of a gene encoding a universal stress protein in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 1992, 6, 3187-3198.	1.2	213
72	Uptake of leucine by a marine Gram-negative heterotrophic bacterium during exposure to starvation conditions. <i>FEMS Microbiology Letters</i> , 1987, 45, 233-241.	0.7	62

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73	Effects of starvation for exogenous carbon on functional mRNA stability and rate of peptide chain elongation in <i>Escherichia coli</i> . , O, .		4