Susan Rae Wente

List of Publications by Year in descending order

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52 papers 4,456 citations

32 h-index 53 g-index

55 all docs

55 docs citations

55 times ranked 3571 citing authors

#	Article	IF	CITATIONS
1	Structure and activation mechanism of the yeast RNA Pol II CTD kinase CTDK-1 complex. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118 , .	7.1	6
2	Functions of Gle1 are governed by two distinct modes of self-association. Journal of Biological Chemistry, 2020, 295, 16813-16825.	3.4	7
3	Nuclear envelope–vacuole contacts mitigate nuclear pore complex assembly stress. Journal of Cell Biology, 2020, 219, .	5.2	18
4	Dbp5 associates with RNA-bound Mex67 and Nab2 and its localization at the nuclear pore complex is sufficient for mRNP export and cell viability. PLoS Genetics, 2020, 16, e1009033.	3.5	9
5	MAPK- and glycogen synthase kinase 3–mediated phosphorylation regulates the DEAD-box protein modulator Gle1 for control of stress granule dynamics. Journal of Biological Chemistry, 2019, 294, 559-575.	3.4	20
6	Gle1 mediates stress granule-dependent survival during chemotoxic stress. Advances in Biological Regulation, 2019, 71, 156-171.	2.3	14
7	A role for Gle1, a regulator of DEAD-box RNA helicases, at centrosomes and basal bodies. Molecular Biology of the Cell, 2017, 28, 120-127.	2.1	35
8	Nup100 regulates <i>Saccharomyces cerevisiae</i> replicative life span by mediating the nuclear export of specific tRNAs. Rna, 2017, 23, 365-377.	3.5	21
9	The Plk1 Piece of the Nuclear Envelope Disassembly Puzzle. Developmental Cell, 2017, 43, 115-117.	7.0	2
10	Nup42 and <scp>IP₆</scp> coordinate Gle1 stimulation of Dbp5/ <scp>DDX19B</scp> for <scp>mRNA</scp> export in yeast and human cells. Traffic, 2017, 18, 776-790.	2.7	37
11	Simple rules for passive diffusion through the nuclear pore complex. Journal of Cell Biology, 2016, 215, 57-76.	5.2	337
12	A Novel <i>Saccharomyces cerevisiae</i> FG Nucleoporin Mutant Collection for Use in Nuclear Pore Complex Functional Experiments. G3: Genes, Genomes, Genetics, 2016, 6, 51-58.	1.8	13
13	An amyotrophic lateral sclerosis-linked mutation in GLE1 alters the cellular pool of human Gle1 functional isoforms. Advances in Biological Regulation, 2016, 62, 25-36.	2.3	21
14	Altering nuclear pore complex function impacts longevity and mitochondrial function in <i>S. cerevisiae</i> . Journal of Cell Biology, 2015, 208, 729-744.	5. 2	55
15	Combining Spinach-tagged RNA and gene localization to image gene expression in live yeast. Nature Communications, 2015, 6, 8882.	12.8	55
16	Cytoplasmic hGle1A regulates stress granules by modulation of translation. Molecular Biology of the Cell, 2015, 26, 1476-1490.	2.1	36
17	Nuclear pore complex integrity requires Lnp1, a regulator of cortical endoplasmic reticulum. Molecular Biology of the Cell, 2015, 26, 2833-2844.	2.1	38
18	Deleterious mutations in the essential mRNA metabolism factor, hGle1, in amyotrophic lateral sclerosis. Human Molecular Genetics, 2015, 24, 1363-1373.	2.9	122

#	Article	IF	Citations
19	Nucleoporin FG Domains Facilitate mRNP Remodeling at the Cytoplasmic Face of the Nuclear Pore Complex. Genetics, 2014, 197, 1213-1224.	2.9	28
20	Casein Kinase II Regulation of the Hot1 Transcription Factor Promotes Stochastic Gene Expression. Journal of Biological Chemistry, 2014, 289, 17668-17679.	3.4	9
21	From Hypothesis to Mechanism: Uncovering Nuclear Pore Complex Links to Gene Expression. Molecular and Cellular Biology, 2014, 34, 2114-2120.	2.3	40
22	Insights into mRNA export-linked molecular mechanisms of human disease through a Gle1 structure–function analysis. Advances in Biological Regulation, 2014, 54, 74-91.	2.3	29
23	Gle1 Functions during mRNA Export in an Oligomeric Complex that Is Altered in Human Disease. Cell, 2013, 155, 582-593.	28.9	70
24	Trafficking to uncharted territory of the nuclear envelope. Current Opinion in Cell Biology, 2012, 24, 341-349.	5 . 4	40
25	Dbp5, Gle1-IP ₆ and Nup159. Nucleus, 2011, 2, 540-548.	2.2	91
26	The Dbp5 cycle at the nuclear pore complex during mRNA export II: nucleotide cycling and mRNP remodeling by Dbp5 are controlled by Nup159 and Gle1. Genes and Development, 2011, 25, 1065-1077.	5.9	98
27	The Dbp5 cycle at the nuclear pore complex during mRNA export I: <i>dbp5</i> mutants with defects in RNA binding and ATP hydrolysis define key steps for Nup159 and Gle1. Genes and Development, 2011, 25, 1052-1064.	5.9	99
28	Gle1 Is a Multifunctional DEAD-box Protein Regulator That Modulates Ded1 in Translation Initiation. Journal of Biological Chemistry, 2011, 286, 39750-39759.	3.4	40
29	Spatial and temporal impacts on a career in science. Molecular Biology of the Cell, 2011, 22, 3923-3925.	2.1	1
30	Zebrafish inositol polyphosphate kinases: New effectors of cilia and developmental signaling. Advances in Enzyme Regulation, 2010, 50, 309-323.	2.6	4
31	The Nuclear Pore Complex and Nuclear Transport. Cold Spring Harbor Perspectives in Biology, 2010, 2, a000562-a000562.	5. 5	569
32	The Mitogen-Activated Protein Kinase Slt2 Regulates Nuclear Retention of Non-Heat Shock mRNAs during Heat Shock-Induced Stress. Molecular and Cellular Biology, 2010, 30, 5168-5179.	2.3	48
33	ER membrane–bending proteins are necessary for de novo nuclear pore formation. Journal of Cell Biology, 2009, 184, 659-675.	5.2	137
34	Inositol polyphosphates: a new frontier for regulating gene expression. Chromosoma, 2008, 117, 1-13.	2.2	91
35	The mRNA Export Factor Gle1 and Inositol Hexakisphosphate Regulate Distinct Stages of Translation. Cell, 2008, 134, 624-633.	28.9	134
36	Nuclear mRNA export requires specific FG nucleoporins for translocation through the nuclear pore complex. Journal of Cell Biology, 2007, 178, 1121-1132.	5. 2	111

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37	The Karyopherin Kap95 Regulates Nuclear Pore Complex Assembly into Intact Nuclear Envelopes In Vivo. Molecular Biology of the Cell, 2007, 18, 886-898.	2.1	47
38	The DEAD-Box Protein Dbp5 Controls mRNA Export by Triggering Specific RNA:Protein Remodeling Events. Molecular Cell, 2007, 28, 850-859.	9.7	200
39	Inositol hexakisphosphate and Gle1 activate the DEAD-box protein Dbp5 for nuclear mRNA export. Nature Cell Biology, 2006, 8, 711-716.	10.3	270
40	Interaction between the Shuttling mRNA Export Factor Gle1 and the Nucleoporin hCG1: A Conserved Mechanism in the Export of Hsp70 mRNA. Molecular Biology of the Cell, 2005, 16, 4304-4315.	2.1	70
41	Nuclear Export of the Yeast mRNA-binding Protein Nab2 Is Linked to a Direct Interaction with Gfd1 and to Gle1 Function. Journal of Biological Chemistry, 2004, 279, 35384-35391.	3.4	32
42	The mRNA Export Factor Human Gle1 Interacts with the Nuclear Pore Complex Protein Nup155. Molecular and Cellular Proteomics, 2004, 3, 145-155.	3.8	52
43	Minimal nuclear pore complexes define FG repeat domains essential for transport. Nature Cell Biology, 2004, 6, 197-206.	10.3	337
44	The Ran GTPase cycle is required for yeast nuclear pore complex assembly. Journal of Cell Biology, 2003, 160, 1041-1053.	5.2	104
45	An essential role for hGle1 nucleocytoplasmic shuttling in mRNA export. Journal of Cell Biology, 2003, 160, 1029-1040.	5.2	54
46	The GLFG Regions of Nup116p and Nup100p Serve as Binding Sites for Both Kap95p and Mex67p at the Nuclear Pore Complex. Journal of Biological Chemistry, 2001, 276, 6445-6452.	3.4	109
47	Nuclear transport: never-ending cycles of signals and receptors. Essays in Biochemistry, 2000, 36, 89-103.	4.7	11
48	A Novel Fluorescence-based Genetic Strategy Identifies Mutants of Saccharomyces cerevisiae Defective for Nuclear Pore Complex Assembly. Molecular Biology of the Cell, 1998, 9, 2439-2461.	2.1	45
49	The Integral Membrane Protein Snl1p Is Genetically Linked to Yeast Nuclear Pore Complex Function. Molecular Biology of the Cell, 1998, 9, 355-373.	2.1	51
50	A Nuclear Export Signal in Kap95p Is Required for Both Recycling the Import Factor and Interaction with the Nucleoporin GLFG Repeat Regions of Nup116p and Nup100p. Journal of Cell Biology, 1997, 137, 797-811.	5.2	84
51	An RNA-export mediator with an essential nuclear export signal. Nature, 1996, 383, 357-360.	27.8	224
52	Pores for thought: nuclear pore complex proteins. Trends in Cell Biology, 1994, 4, 357-365.	7.9	276