Maya Schuldiner

List of Publications by Year
in descending order

153 papers	16,569 citations	28274 55 h-index	17105 122 g-index
195 all docs	195 docs citations	195 times ranked	16805 citing authors

#	Article	IF	Citations
1	Differentiation of Human Embryonic Stem Cells into Embryoid Bodies Comprising the Three Embryonic Germ Layers. Molecular Medicine, 2000, 6, 88-95.	4.4	1,377
2	An ER-Mitochondria Tethering Complex Revealed by a Synthetic Biology Screen. Science, 2009, 325, 477-481.	12.6	1,129
3	Functional dissection of protein complexes involved in yeast chromosome biology using a genetic interaction map. Nature, 2007, 446, 806-810.	27.8	806
4	Exploration of the Function and Organization of the Yeast Early Secretory Pathway through an Epistatic Miniarray Profile. Cell, 2005, 123, 507-519.	28.9	804
5	Cotranscriptional Set2 Methylation of Histone H3 Lysine 36 Recruits a Repressive Rpd3 Complex. Cell, 2005, 123, 593-605.	28.9	712
6	Comprehensive Characterization of Genes Required for Protein Folding in the Endoplasmic Reticulum. Science, 2009, 323, 1693-1697.	12.6	646
7	Characterization of the expression of MHC proteins in human embryonic stem cells. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 9864-9869.	7.1	628
8	A comprehensive strategy enabling high-resolution functional analysis of the yeast genome. Nature Methods, 2008, 5, 711-718.	19.0	473
9	The GET Complex Mediates Insertion of Tail-Anchored Proteins into the ER Membrane. Cell, 2008, 134, 634-645.	28.9	460
10	Coming together to define membrane contactÂsites. Nature Communications, 2019, 10, 1287.	12.8	435
11	Induced neuronal differentiation of human embryonic stem cells. Brain Research, 2001, 913, 201-205.	2.2	410
12	A mitochondrial-focused genetic interaction map reveals a scaffold-like complex required for inner membrane organization in mitochondria. Journal of Cell Biology, 2011, 195, 323-340.	5.2	402
13	Establishment of human embryonic stem cell-transfected clones carrying a marker for undifferentiated cells. Current Biology, 2001, 11, 514-518.	3.9	360
14	Definition of a High-Confidence Mitochondrial Proteome at Quantitative Scale. Cell Reports, 2017, 19, 2836-2852.	6.4	346
15	A Dynamic Interface between Vacuoles and Mitochondria in Yeast. Developmental Cell, 2014, 30, 95-102.	7.0	321
16	A Tether Is a Tether Is a Tether: Tethering at Membrane Contact Sites. Developmental Cell, 2016, 39, 395-409.	7.0	315
17	A strategy for extracting and analyzing large-scale quantitative epistatic interaction data. Genome Biology, 2006, 7, R63.	9.6	287
18	Selective Ablation of Human Embryonic Stem Cells Expressing a â€œSuicideâ€•Gene. Stem Cells, 2003, 21, 257-265.	3.2	267

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19	Systematic mapping of contact sites reveals tethers and a function for the peroxisome-mitochondria contact. Nature Communications, 2018, 9, 1761.	12.8	222
20	A novel single-cell screening platform reveals proteome plasticity during yeast stress responses. Journal of Cell Biology, 2013, 200, 839-850.	5.2	210
21	Staying in touch: the molecular era of organelle contact sites. Trends in Biochemical Sciences, 2011, 36, 616-623.	7.5	195
22	Confinement to Organelle-Associated Inclusion Structures Mediates Asymmetric Inheritance of Aggregated Protein in Budding Yeast. Cell Reports, 2012, 2, 738-747.	6.4	173
23	Lam6 Regulates the Extent of Contacts between Organelles. Cell Reports, 2015, 12, 7-14.	6.4	173
24	One library to make them all: streamlining the creation of yeast libraries via a SWAp-Tag strategy. Nature Methods, 2016, 13, 371-378.	19.0	171
25	The SND proteins constitute an alternative targeting route to the endoplasmic reticulum. Nature, 2016, 540, 134-138.	27.8	168
26	Modeling for Lesch-Nyhan Disease by Gene Targeting in Human Embryonic Stem Cells. Stem Cells, 2004, 22, 635-641.	3.2	167
27	A Network of Cytosolic Factors Targets SRP-Independent Proteins to the Endoplasmic Reticulum. Cell, 2013, 152, 1134-1145.	28.9	166
28	A different kind of love â€“ lipid droplet contact sites. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 2017, 1862, 1188-1196.	2.4	160
29	Backup without redundancy: genetic interactions reveal the cost of duplicate gene loss. Molecular Systems Biology, 2007, 3, 86.	7.2	143
30	Genome-wide SWAp-Tag yeast libraries for proteome exploration. Nature Methods, 2018, 15, 617-622.	19.0	134
31	An ER surface retrieval pathway safeguards the import of mitochondrial membrane proteins in yeast. Science, 2018, 361, 1118-1122.	12.6	129
32	No peroxisome is an island â€” Peroxisome contact sites. Biochimica Et Biophysica Acta - Molecular Cell Research, 2016, 1863, 1061-1069.	4.1	126
33	Ergosterol content specifies targeting of tail-anchored proteins to mitochondrial outer membranes. Molecular Biology of the Cell, 2012, 23, 3927-3935.	2.1	119
34	Targeting and translocation of proteins to the endoplasmic reticulum at a glance. Journal of Cell Science, 2017, 130, 4079-4085.	2.0	111
35	OM14 is a mitochondrial receptor for cytosolic ribosomes that supports co-translational import into mitochondria. Nature Communications, 2014, 5, 5711.	12.8	106
36	Advanced Methods for High-Throughput Microscopy Screening of Genetically Modified Yeast Libraries. Methods in Molecular Biology, 2011, 781, 127-159.	0.9	101

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37	A defect in the RNA-processing protein HNRPDL causes limb-girdle muscular dystrophy 1G (LGMD1G). Human Molecular Genetics, 2014, 23, 4103-4110.	2.9	101
38	The PH gene determines fruit acidity and contributes to the evolution of sweet melons. Nature Communications, 2014, 5, 4026.	12.8	100
39	Identification of seipin-linked factors that act as determinants of a lipid droplet subpopulation. Journal of Cell Biology, 2018, 217, 269-282.	5.2	99
40	Peroxisomes are juxtaposed to strategic sites on mitochondria. Molecular BioSystems, 2014, 10, 1742-1748.	2.9	95
41	The Protease Ste24 Clears Clogged Translocons. Cell, 2016, 164, 103-114.	28.9	93
42	Rapid creation and quantitative monitoring of high coverage shRNA libraries. Nature Methods, 2009, 6, 443-445.	19.0	92
43	Lipid Droplets Are Essential for Efficient Clearance of Cytosolic Inclusion Bodies. Developmental Cell, 2015, 33, 603-610.	7.0	92
44	APOL1â€“Mediated Cell Injury Involves Disruption of Conserved Trafficking Processes. Journal of the American Society of Nephrology: JASN, 2017, 28, 1117-1130.	6.1	88
45	A Systematic Approach to Pair Secretory Cargo Receptors with Their Cargo Suggests a Mechanism for Cargo Selection by Erv14. PLoS Biology, 2012, 10, e1001329.	5.6	87
46	The emergence of proteome-wide technologies: systematic analysis of proteins comes of age. Nature Reviews Molecular Cell Biology, 2014, 15, 453-464.	37.0	80
47	Cytosolic Events in the Biogenesis of Mitochondrial Proteins. Trends in Biochemical Sciences, 2020, 45, 650-667.	7.5	79
48	Formation and dissociation of proteasome storage granules are regulated by cytosolic pH. Journal of Cell Biology, 2013, 201, 663-671.	5.2	76
49	The Back and Forth of Cargo Exit from the Endoplasmic Reticulum. Current Biology, 2014, 24, R130-R136.	3.9	75
50	Ground control to major TOM: mitochondriaâ€“nucleus communication. FEBS Journal, 2017, 284, 196-210.	4.7	75
51	LoQAtEâ€"Localization and Quantitation ATlas of the yeast proteomE. A new tool for multiparametric dissection of single-protein behavior in response to biological perturbations in yeast. Nucleic Acids Research, 2014, 42, D726-D730.	14.5	74
52	Natural genetic variation for expression of a <scp>SWEET</scp> transporter among wild species of <i>Solanum lycopersicum</i> (tomato) determines the hexose composition of ripening tomato fruit. Plant Journal, 2018, 96, 343-357.	5.7	74
53	The Role of Djp1 in Import of the Mitochondrial Protein Mim1 Demonstrates Specificity between a Cochaperone and Its Substrate Protein. Molecular and Cellular Biology, 2013, 33, 4083-4094.	2.3	68
54	Characterization of proteome dynamics in oleate reveals a novel peroxisome targeting receptor. Journal of Cell Science, 2016, 129, 4067-4075.	2.0	63

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55	The Yeast P5 Type ATPase, Spf1, Regulates Manganese Transport into the Endoplasmic Reticulum. PLoS ONE, 2013, 8, e85519.	2.5	62
56	<scp>hS</scp>nd2 protein represents an alternative targeting factor to the endoplasmic reticulum in human cells. FEBS Letters, 2017, 591, 3211-3224.	2.8	55
57	Identification of yeast proteins necessary for cell-surface function of a potassium channel. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 18079-18084.	7.1	53
58	A pathway of targeted autophagy is induced by DNA damage in budding yeast. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E1158-E1167.	7.1	52
59	Compartmentalized Synthesis of Triacylglycerol at the Inner Nuclear Membrane Regulates Nuclear Organization. Developmental Cell, 2019, 50, 755-766.e6.	7.0	52
60	Promethin Is a Conserved Seipin Partner Protein. Cells, 2019, 8, 268.	4.1	52
61	Get3 is a holdase chaperone and moves to deposition sites for aggregated proteins when membrane targeting is blocked. Journal of Cell Science, 2013, 126, 473-483.	2.0	50
62	Assessment of GFP Tag Position on Protein Localization and Growth Fitness in Yeast. Journal of Molecular Biology, 2019, 431, 636-641.	4.2	49
63	The Yeast ER-Intramembrane Protease Ypf1 Refines Nutrient Sensing by Regulating Transporter Abundance. Molecular Cell, 2014, 56, 630-640.	9.7	48
64	The chaperone-binding activity of the mitochondrial surface receptor Tom70 protects the cytosol against mitoprotein-induced stress. Cell Reports, 2021, 35, 108936.	6.4	47
65	Starvation-Dependent Regulation of Golgi Quality Control Links the TOR Signaling and Vacuolar Protein Sorting Pathways. Cell Reports, 2015, 12, 1876-1886.	6.4	46
66	Primersâ€•4â€•Yeast: a comprehensive web tool for planning primers for <i>Saccharomyces cerevisiae</i>. Yeast, 2014, 31, 77-80.	1.7	41
67	Making Sense of the Yeast Sphingolipid Pathway. Journal of Molecular Biology, 2016, 428, 4765-4775.	4.2	41
68	Getting the whole picture: combining throughput with content in microscopy. Journal of Cell Science, 2011, 124, 3743-3751.	2.0	40
69	A cytosolic degradation pathway, prERAD, monitors pre-inserted secretory pathway proteins. Journal of Cell Science, 2014, 127, 3017-23.	2.0	40
70	YeastRGB: comparing the abundance and localization of yeast proteins across cells and libraries. Nucleic Acids Research, 2019, 47, D1245-D1249.	14.5	39
71	The NADH Dehydrogenase Nde1 Executes Cell Death after Integrating Signals from Metabolism and Proteostasis on the Mitochondrial Surface. Molecular Cell, 2020, 77, 189-202.e6.	9.7	39
72	Mind the Organelle Gap â€“ Peroxisome Contact Sites in Disease. Trends in Biochemical Sciences, 2018, 43, 199-210.	7.5	36

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73	Mice lacking WRB reveal differential biogenesis requirements of tail-anchored proteins in vivo. Scientific Reports, 2016, 6, 39464.	3.3	35
74	Iron affects Ire1 clustering propensity and the amplitude of endoplasmic reticulum stress signaling. Journal of Cell Science, 2017, 130, 3222-3233.	2.0	35
75	The Endoplasmic Reticulum-Mitochondria Encounter Structure Complex Coordinates Coenzyme Q Biosynthesis. Contact (Thousand Oaks (Ventura County, Calif)), 2019, 2, 251525641882540.	1.3	35
76	All roads lead to Rome (but some may be harder to travel): SRP-independent translocation into the endoplasmic reticulum. Critical Reviews in Biochemistry and Molecular Biology, 2013, 48, 273-288.	5.2	34
77	Embracing the voidâ€"how much do we really know about targeting and translocation to the endoplasmic reticulum?. Current Opinion in Cell Biology, 2014, 29, 8-17.	5.4	34
78	Pex35 is a regulator of peroxisome abundance. Journal of Cell Science, 2017, 130, 791-804.	2.0	34
79	The GET pathway can increase the risk of mitochondrial outer membrane proteins to be mistargeted to the ER. Journal of Cell Science, 2018, 131, .	2.0	34
80	Interactions of subunit CCT3 in the yeast chaperonin CCT/TRiC with Q/N-rich proteins revealed by high-throughput microscopy analysis. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18833-18838.	7.1	32
81	An unrecognized function for COPII components in recruiting a viral replication protein to the perinuclear ER. Journal of Cell Science, 2016, 129, 3597-3608.	2.0	32
82	Accurate, Model-Based Tuning of Synthetic Gene Expression Using Introns in S. cerevisiae. PLoS Genetics, 2014, 10, e1004407.	3.5	31
83	Heterosis as a consequence of regulatory incompatibility. BMC Biology, 2017, 15, 38.	3.8	31
84	Genome-Wide Screens in <i>Saccharomyces cerevisiae</i> Highlight a Role for Cardiolipin in Biogenesis of Mitochondrial Outer Membrane Multispan Proteins. Molecular and Cellular Biology, 2015, 35, 3200-3211.	2.3	30
85	The yeast oligopeptide transporter Opt2 is localized to peroxisomes and affects glutathione redox homeostasis. FEMS Yeast Research, 2014, 14, n/a-n/a.	2.3	29
86	Cnm1 mediates nucleusâ€"mitochondria contact site formation in response to phospholipid levels. Journal of Cell Biology, 2021, 220, .	5.2	29
87	Saccharomyces cerevisiae cells lacking Pex3 contain membrane vesicles that harbor a subset of peroxisomal membrane proteins. Biochimica Et Biophysica Acta - Molecular Cell Research, 2017, 1864, 1656-1667.	4.1	28
88	Modularity and directionality in genetic interaction maps. Bioinformatics, 2010, 26, i228-i236.	4.1	27
89	Mitochatting â€" If only we could be a fly on the cell wall. Biochimica Et Biophysica Acta - Molecular Cell Research, 2017, 1864, 1469-1480.	4.1	27
90	Temporal profiling of redox-dependent heterogeneity in single cells. ELife, 2018, 7, .	6.0	27

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91	Defining the Mammalian Peroxisomal Proteome. Sub-Cellular Biochemistry, 2018, 89, 47-66.	2.4	26
92	Yeast ceramide synthases, Lag1 and Lac1, have distinct substrate specificity. Journal of Cell Science, 2019, 132, .	2.0	26
93	New horizons in mitochondrial contact site research. Biological Chemistry, 2020, 401, 793-809.	2.5	24
94	Stepping outside the comfort zone of membrane contact site research. Nature Reviews Molecular Cell Biology, 2018, 19, 483-484.	37.0	21
95	AMPK regulates ESCRT-dependent microautophagy of proteasomes concomitant with proteasome storage granule assembly during glucose starvation. PLoS Genetics, 2019, 15, e1008387.	3.5	21
96	A piggybacking mechanism enables peroxisomal localization of the glyoxylate cycle enzyme Mdh2 in yeast. Journal of Cell Science, 2020, 133, .	2.0	21
97	Pex14p Phosphorylation Modulates Import of Citrate Synthase 2 Into Peroxisomes in Saccharomyces cerevisiae. Frontiers in Cell and Developmental Biology, 2020, 8, 549451.	3.7	20
98	Beyond rare disorders: A new era for peroxisomal pathophysiology. Molecular Cell, 2022, 82, 2228-2235.	9.7	19
99	Factors Controlling Human Embryonic Stem Cell Differentiation. Methods in Enzymology, 2003, 365, 446-461.	1.0	18
100	From rags to riches â€" The history of the endoplasmic reticulum. Biochimica Et Biophysica Acta - Molecular Cell Research, 2013, 1833, 2389-2391.	4.1	18
101	Protein Topology Prediction Algorithms Systematically Investigated in the Yeast <i>Saccharomyces cerevisiae</i>. BioEssays, 2019, 41, e1800252.	2.5	18
102	High-throughput ultrastructure screening using electron microscopy and fluorescent barcoding. Journal of Cell Biology, 2019, 218, 2797-2811.	5.2	18
103	The ER protein Ema19 facilitates the degradation of nonimported mitochondrial precursor proteins. Molecular Biology of the Cell, 2021, 32, 664-674.	2.1	18
104	ER-SURF: Riding the Endoplasmic Reticulum Surface to Mitochondria. International Journal of Molecular Sciences, 2021, 22, 9655.	4.1	18
105	Peroxisome function relies on organelle-associated mRNA translation. Science Advances, 2022, 8, eabk2141.	10.3	18
106	Peroxisome Mini-Libraries: Systematic Approaches to Study Peroxisomes Made Easy. Methods in Molecular Biology, 2017, 1595, 305-318.	0.9	17
107	Uncovering targeting priority to yeast peroxisomes using an in-cell competition assay. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 21432-21440.	7.1	17
108	Noncanonical regulation of phosphatidylserine metabolism by a Sec14-like protein and a lipid kinase. Journal of Cell Biology, 2020, 219, .	5.2	16

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109	Toolbox: Creating a systematic database of secretory pathway proteins uncovers new cargo for COPI. Traffic, 2018, 19, 370-379.	2.7	15
110	Incredibly closeâ€"A newly identified peroxisomeâ€"ER contact site in humans. Journal of Cell Biology, 2017, 216, 287-289.	5.2	14
111	Cytotoxicity of 1-deoxysphingolipid unraveled by genome-wide genetic screens and lipidomics in <i>Saccharomyces cerevisiae</i>. Molecular Biology of the Cell, 2019, 30, 2814-2826.	2.1	14
112	Post-ER degradation of misfolded GPI-anchored proteins is linked with microautophagy. Current Biology, 2021, 31, 4025-4037.e5.	3.9	14
113	Two novel effectors of trafficking and maturation of the yeast plasma membrane H⁺â€‹<scp>ATPase</scp>. Traffic, 2017, 18, 672-682.	2.7	13
114	Syp1 regulates the clathrin-mediated and clathrin-independent endocytosis of multiple cargo proteins through a novel sorting motif. Molecular Biology of the Cell, 2017, 28, 2434-2448.	2.1	13
115	Cvm1 is a component of multiple vacuolar contact sites required for sphingolipid homeostasis. Journal of Cell Biology, 2022, 221, .	5.2	13
116	Explorations in topologyâ€"delving underneath the surface of genetic interaction maps. Molecular BioSystems, 2009, 5, 1473.	2.9	12
117	The Contribution of Systematic Approaches to Characterizing the Proteins and Functions of the Endoplasmic Reticulum. Cold Spring Harbor Perspectives in Biology, 2013, 5, a013284-a013284.	5.5	12
118	The Fast and the Furious: Golgi Contact Sites. Contact (Thousand Oaks (Ventura County, Calif)), 2021, 4, 251525642110344.	1.3	12
119	Widespread use of unconventional targeting signals in mitochondrial ribosome proteins. EMBO Journal, 2022, 41, e109519.	7.8	12
120	Yeast phospholipid biosynthesis is linked to mRNA localization. Journal of Cell Science, 2014, 127, 3373-81.	2.0	11
121	Water-Transfer Slows Aging in Saccharomyces cerevisiae. PLoS ONE, 2016, 11, e0148650.	2.5	11
122	Combining Deep Sequencing, Proteomics, Phosphoproteomics, and Functional Screens To Discover Novel Regulators of Sphingolipid Homeostasis. Journal of Proteome Research, 2017, 16, 571-582.	3.7	11
123	Cargo Release from Myosin V Requires the Convergence of Parallel Pathways that Phosphorylate and Ubiquitylate the Cargo Adaptor. Current Biology, 2020, 30, 4399-4412.e7.	3.9	11
124	Double the Fun, Double the Trouble: Paralogs and Homologs Functioning in the Endoplasmic Reticulum. Annual Review of Biochemistry, 2020, 89, 637-666.	11.1	10
125	Unbiased yeast screens identify cellular pathways affected in Niemannâ€"Pick disease type C. Life Science Alliance, 2020, 3, e201800253.	2.8	10
126	Characterization of an M28 metalloprotease family member residing in the yeast vacuole. FEMS Yeast Research, 2013, 13, 471-484.	2.3	9

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127	Database for High Throughput Screening Hits (dHITS): a simple tool to retrieve gene specific phenotypes from systematic screens done in yeast. Yeast, 2018, 35, 477-483.	1.7	7
128	Disease-causing mutations in subunits of OXPHOS complex I affect certain physical interactions. Scientific Reports, 2019, 9, 9987.	3.3	7
129	Cellular Consequences of Diminished Protein O-Mannosyltransferase Activity in Bakerâ€™s Yeast. International Journal of Molecular Sciences, 2017, 18, 1226.	4.1	6
130	Transfer of the Septin Ring to Cytokinetic Remnants in ER Stress Directs Age-Sensitive Cell-Cycle Re-entry. Developmental Cell, 2019, 51, 173-191.e5.	7.0	6
131	Weizmann Young PI Forum: The Power of Peer Support. Molecular Cell, 2009, 36, 913-915.	9.7	5
132	Protein Degradation: BAGging Up the Trash. Current Biology, 2011, 21, R692-R695.	3.9	5
133	The mitochondrial intermembrane spaceâ€“facing proteins Mcp2 and Tgl2 are involved in yeast lipid metabolism. Molecular Biology of the Cell, 2019, 30, 2681-2694.	2.1	5
134	Validation of a yeast malate dehydrogenase 2 (Mdh2) antibody tested for use in western blots. F1000Research, 2018, 7, 130.	1.6	5
135	The plasma membrane code. Nature Chemical Biology, 2010, 6, 487-488.	8.0	4
136	Peroxisystem: Harnessing systems cell biology to study peroxisomes. Biology of the Cell, 2015, 107, 89-97.	2.0	4
137	Validation of a yeast malate dehydrogenase 2 (Mdh2) antibody tested for use in western blots. F1000Research, 2018, 7, 130.	1.6	4
138	Translational Regulation of Pmt1 and Pmt2 by Bfr1 Affects Unfolded Protein O-Mannosylation. International Journal of Molecular Sciences, 2019, 20, 6220.	4.1	4
139	Overexpression of branched-chain amino acid aminotransferases rescues the growth defects of cells lacking the Barth syndrome-related gene TAZ1. Journal of Molecular Medicine, 2019, 97, 269-279.	3.9	4
140	Show your true color: Mammalian cell surface staining for tracking cellular identity in multiplexing and beyond. Current Opinion in Chemical Biology, 2022, 66, 102102.	6.1	4
141	Pls1 Is a Peroxisomal Matrix Protein with a Role in Regulating Lysine Biosynthesis. Cells, 2022, 11, 1426.	4.1	3
142	A Similarity-Based Method for Predicting Enzymatic Functions in Yeast Uncovers a New AMP Hydrolase. Journal of Molecular Biology, 2022, 434, 167478.	4.2	2
143	The DNA Damage Road Map. Science, 2010, 330, 1327-1328.	12.6	1
144	Organelle structure and biogenesis. Molecular Biology of the Cell, 2011, 22, 723-723.	2.1	1

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145	Editorial overview: Cell organelles: Organelle communication: new means and new views. Current Opinion in Cell Biology, 2015, 35, v-vi.	5.4	1
146	Maya Schuldiner. Current Biology, 2017, 27, R982-R984.	3.9	0
147	Genetic Interaction mapping of essential genes in <i>Saccharomyces cerevisiae</i>. FASEB Journal, 2007, 21, A1004.	0.5	0
148	Using high content microscopy screening to uncover insertion pathways for transmembrane proteins. FASEB Journal, 2011, 25, 194.3.	0.5	0
149	Novel Regulation of Lipid Metabolism by a Phosphatidylinositol Transfer Protein and a Phosphatidylinositol 4â€Kinase. FASEB Journal, 2019, 33, lb330.	0.5	0
150	Title is missing!. , 2019, 15, e1008387.		0
151	Title is missing!. , 2019, 15, e1008387.		0
152	Title is missing!. , 2019, 15, e1008387.		0
153	Title is missing!. , 2019, 15, e1008387.		0