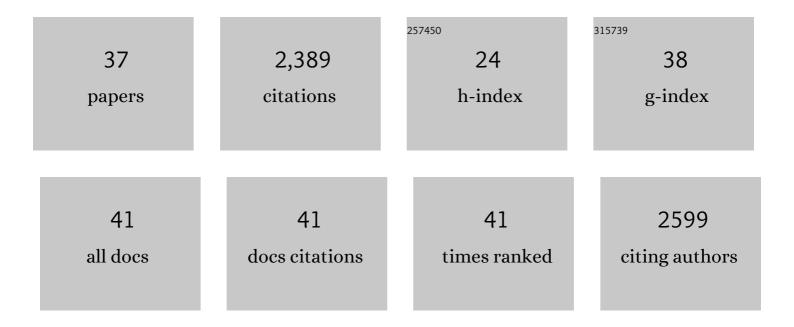
## Sabine Rospert

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Ribosome-bound Get4/5 facilitates the capture of tail-anchored proteins by Sgt2 in yeast. Nature Communications, 2021, 12, 782.	12.8	14
2	Quantitative proteomics identifies the universally conserved ATPase Ola1p as a positive regulator of heat shock response in Saccharomyces cerevisiae. Journal of Biological Chemistry, 2021, 297, 101050.	3.4	6
3	The ribosome-associated complex RAC serves in a relay that directs nascent chains to Ssb. Nature Communications, 2020, 11, 1504.	12.8	21
4	Polyamines and eIF5A Hypusination Modulate Mitochondrial Respiration and Macrophage Activation. Cell Metabolism, 2019, 30, 352-363.e8.	16.2	223
5	A dual role of the ribosome-bound chaperones RAC/Ssb in maintaining the fidelity of translation termination. Nucleic Acids Research, 2019, 47, 7018-7034.	14.5	12
6	The Hsp70 homolog Ssb affects ribosome biogenesis via the TORC1-Sch9 signaling pathway. Nature Communications, 2017, 8, 937.	12.8	22
7	Two chaperones locked in an embrace: structure and function of the ribosome-associated complex RAC. Nature Structural and Molecular Biology, 2017, 24, 611-619.	8.2	50
8	Protein glutaminylation is a yeast-specific posttranslational modification of elongation factor 1A. Journal of Biological Chemistry, 2017, 292, 16014-16023.	3.4	13
9	The yeast Hsp70 homolog Ssb: a chaperone for general de novo protein folding and a nanny for specific intrinsically disordered protein domains. Current Genetics, 2017, 63, 9-13.	1.7	9
10	Ribosomal Protein Rps26 Influences 80S Ribosome Assembly in Saccharomyces cerevisiae. MSphere, 2016, 1, .	2.9	11
11	Interaction of the cotranslational Hsp70 Ssb with ribosomal proteins and rRNA depends on its lid domain. Nature Communications, 2016, 7, 13563.	12.8	49
12	The Hsp70 homolog Ssb and the 14-3-3 protein Bmh1 jointly regulate transcription of glucose repressed genes in <i>Saccharomyces cerevisiae</i> . Nucleic Acids Research, 2016, 44, 5629-5645.	14.5	25
13	Cotranslational Intersection between the SRP and GET Targeting Pathways to the Endoplasmic Reticulum of <i>Saccharomyces cerevisiae</i> . Molecular and Cellular Biology, 2016, 36, 2374-2383.	2.3	15
14	Inefficient SRP Interaction with a Nascent Chain Triggers a mRNA Quality Control Pathway. Cell, 2014, 156, 146-157.	28.9	77
15	Release Factor eRF3 Mediates Premature Translation Termination on Polylysine-Stalled Ribosomes in Saccharomyces cerevisiae. Molecular and Cellular Biology, 2014, 34, 4062-4076.	2.3	40
16	Interaction of Nascent Chains with the Ribosomal Tunnel Proteins Rpl4, Rpl17, and Rpl39 of Saccharomyces cerevisiae. Journal of Biological Chemistry, 2013, 288, 33697-33707.	3.4	35
17	NAC functions as a modulator of SRP during the early steps of protein targeting to the endoplasmic reticulum. Molecular Biology of the Cell, 2012, 23, 3027-3040.	2.1	57
18	Elongation Factor 1A Is the Target of Growth Inhibition in Yeast Caused by Legionella pneumophila Glucosyltransferase Lgt1. Journal of Biological Chemistry, 2012, 287, 26029-26037.	3.4	28

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19	Ribosome-Associated Complex and Ssb Are Required for Translational Repression Induced by Polylysine Segments within Nascent Chains. Molecular and Cellular Biology, 2012, 32, 4769-4779.	2.3	40
20	The Chaperone Network Connected to Human Ribosome-Associated Complex. Molecular and Cellular Biology, 2011, 31, 1160-1173.	2.3	77
21	The ribosome-bound Hsp70 homolog Ssb of Saccharomyces cerevisiae. Biochimica Et Biophysica Acta - Molecular Cell Research, 2010, 1803, 662-672.	4.1	57
22	Molecular chaperones and intracellular protein transport. Biochimica Et Biophysica Acta - Molecular Cell Research, 2010, 1803, 639-640.	4.1	1
23	Transcriptional activation of polycomb-repressed genes by ZRF1. Nature, 2010, 468, 1124-1128.	27.8	127
24	A Targeted Analysis of Cellular Chaperones Reveals Contrasting Roles for Heat Shock Protein 70 in Flock House Virus RNA Replication. Journal of Virology, 2010, 84, 330-339.	3.4	30
25	A signal-anchor sequence stimulates signal recognition particle binding to ribosomes from inside the exit tunnel. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 1398-1403.	7.1	113
26	The Hsp70 homolog Ssb is essential for glucose sensing via the SNF1 kinase network. Genes and Development, 2009, 23, 2102-2115.	5.9	36
27	Ribosome-associated Complex Binds to Ribosomes in Close Proximity of Rpl31 at the Exit of the Polypeptide Tunnel in Yeast. Molecular Biology of the Cell, 2008, 19, 5279-5288.	2.1	78
28	Association of Protein Biogenesis Factors at the Yeast Ribosomal Tunnel Exit Is Affected by the Translational Status and Nascent Polypeptide Sequence*. Journal of Biological Chemistry, 2007, 282, 7809-7816.	3.4	116
29	Functional Characterization of the Atypical Hsp70 Subunit of Yeast Ribosome-associated Complex. Journal of Biological Chemistry, 2007, 282, 33977-33984.	3.4	38
30	Distinct yet linked: chaperone networks in the eukaryotic cytosol. Genome Biology, 2006, 7, 208.	9.6	7
31	The Hsp70 Ssz1 modulates the function of the ribosome-associated J-protein Zuo1. Nature Structural and Molecular Biology, 2005, 12, 497-504.	8.2	112
32	The chaperones MPP11 and Hsp70L1 form the mammalian ribosome-associated complex. Proceedings of the United States of America, 2005, 102, 10064-10069.	7.1	121
33	The Ribosome-Bound Chaperones RAC and Ssb1/2p Are Required for Accurate Translation in Saccharomyces cerevisiae. Molecular and Cellular Biology, 2004, 24, 9186-9197.	2.3	87
34	Ribosome Function: Governing the Fate of a Nascent Polypeptide. Current Biology, 2004, 14, R386-R388.	3.9	24
35	The Yeast Nα-Acetyltransferase NatA Is Quantitatively Anchored to the Ribosome and Interacts with Nascent Polypeptides. Molecular and Cellular Biology, 2003, 23, 7403-7414.	2.3	201
36	A functional chaperone triad on the yeast ribosome. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 4209-4214.	7.1	149

#	Article	IF	CITATIONS
37	Nascent Polypeptide–associated Complex Stimulates Protein Import into Yeast Mitochondria. Molecular Biology of the Cell, 1999, 10, 3289-3299.	2.1	145