Reika Watanabe

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avg, IF4.24
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#	Paper	IF	Citations
34	A photoactivable multi-inhibitor nanoliposome for tumour control and simultaneous inhibition of treatment escape pathways. <i>Nature Nanotechnology</i> , 2016 , 11, 378-87	28.7	169
33	Concentration of GPI-anchored proteins upon ER exit in yeast. <i>Traffic</i> , 2009 , 10, 186-200	5.7	134
32	Multiple functions of sterols in yeast endocytosis. <i>Molecular Biology of the Cell</i> , 2002 , 13, 2664-80	3.5	131
31	Expression cloning of PIG-L, a candidate N-acetylglucosaminyl-phosphatidylinositol deacetylase. <i>Journal of Biological Chemistry</i> , 1997 , 272, 15834-40	5.4	118
30	Pig-n, a mammalian homologue of yeast Mcd4p, is involved in transferring phosphoethanolamine to the first mannose of the glycosylphosphatidylinositol. <i>Journal of Biological Chemistry</i> , 1999 , 274, 350	9 ⁵ 9 ⁴ 106	5 ¹⁰⁹
29	The yeast p24 complex regulates GPI-anchored protein transport and quality control by monitoring anchor remodeling. <i>Molecular Biology of the Cell</i> , 2011 , 22, 2924-36	3.5	93
28	Sorting of GPI-anchored proteins into ER exit sites by p24 proteins is dependent on remodeled GPI. <i>Journal of Cell Biology</i> , 2011 , 194, 61-75	7.3	91
27	Yeast ARV1 is required for efficient delivery of an early GPI intermediate to the first mannosyltransferase during GPI assembly and controls lipid flow from the endoplasmic reticulum. <i>Molecular Biology of the Cell</i> , 2008 , 19, 2069-82	3.5	84
26	Sphingolipids are required for the stable membrane association of glycosylphosphatidylinositol-anchored proteins in yeast. <i>Journal of Biological Chemistry</i> , 2002 , 277, 495	3 8 -44	84
25	PIG-C, one of the three human genes involved in the first step of glycosylphosphatidylinositol biosynthesis is a homologue of Saccharomyces cerevisiae GPI2. <i>Biochemical and Biophysical Research Communications</i> , 1996 , 226, 193-9	3.4	81
24	PIG-A and PIG-H, which participate in glycosylphosphatidylinositol anchor biosynthesis, form a protein complex in the endoplasmic reticulum. <i>Journal of Biological Chemistry</i> , 1996 , 271, 26868-75	5.4	72
23	Requirement of PIG-F and PIG-O for transferring phosphoethanolamine to the third mannose in glycosylphosphatidylinositol. <i>Journal of Biological Chemistry</i> , 2000 , 275, 20911-9	5.4	67
22	Structure of LRRK2 in Parkinson's disease and model for microtubule interaction. <i>Nature</i> , 2020 , 588, 344-349	50.4	60
21	The In Situ Structure of Parkinson's Disease-Linked LRRK2. <i>Cell</i> , 2020 , 182, 1508-1518.e16	56.2	57
20	Yeast Ras regulates the complex that catalyzes the first step in GPI-anchor biosynthesis at the ER. <i>Cell</i> , 2004 , 117, 637-48	56.2	54
19	Mammalian PIG-L and its yeast homologue Gpi12p areN-acetylglucosaminylphosphatidylinositol de-N-acetylases essential in glycosylphosphatidylinositol biosynthesis. <i>Biochemical Journal</i> , 1999 , 339, 185	3.8	52
18	Differential ER exit in yeast and mammalian cells. Current Opinion in Cell Biology, 2004, 16, 350-5	9	47

LIST OF PUBLICATIONS

17	Changes in molecular species profiles of glycosylphosphatidylinositol anchor precursors in early stages of biosynthesis. <i>Journal of Lipid Research</i> , 2007 , 48, 1599-606	6.3	40	
16	Preparing samples from whole cells using focused-ion-beam milling for cryo-electron tomography. <i>Nature Protocols</i> , 2020 , 15, 2041-2070	18.8	38	
15	GPI1 stabilizes an enzyme essential in the first step of glycosylphosphatidylinositol biosynthesis. <i>Journal of Biological Chemistry</i> , 1999 , 274, 18582-8	5.4	29	
14	Exit of GPI-anchored proteins from the ER differs in yeast and mammalian cells. <i>Traffic</i> , 2010 , 11, 1017	-3 3 .7	26	
13	Interaction of Escherichia coli RecA protein with ATP and its analogues. <i>Journal of Biochemistry</i> , 1994 , 116, 960-6	3.1	22	
12	GPI biosynthesis is essential for rhodopsin sorting at the trans-Golgi network in Drosophila photoreceptors. <i>Development (Cambridge)</i> , 2013 , 140, 385-94	6.6	19	
11	Apical sorting of lysoGPI-anchored proteins occurs independent of association with detergent-resistant membranes but dependent on their N-glycosylation. <i>Molecular Biology of the Cell</i> , 2013 , 24, 2021-33	3.5	18	
10	The molecular architecture of engulfment during sporulation. <i>ELife</i> , 2019 , 8,	8.9	17	
9	Clathrin and AP1 are required for apical sorting of glycosyl phosphatidyl inositol-anchored proteins in biosynthetic and recycling routes in Madin-Darby canine kidney cells. <i>Traffic</i> , 2018 , 19, 215-228	5.7	12	
8	Stable cell surface expression of GPI-anchored proteins, but not intracellular transport, depends on their fatty acid structure. <i>Traffic</i> , 2014 , 15, 1305-29	5.7	10	
7	Challenges of Integrating Stochastic Dynamics and Cryo-Electron Tomograms in Whole-Cell Simulations. <i>Journal of Physical Chemistry B</i> , 2017 , 121, 3871-3881	3.4	9	
6	Parkinson∄ Disease-linked LRRK2 structure and model for microtubule interaction		9	
5	The in situ structure of Parkinson⊠ disease-linked LRRK2		8	
4	The presence of an ER exit signal determines the protein sorting upon ER exit in yeast. <i>Biochemical Journal</i> , 2008 , 414, 237-45	3.8	5	
3	Visualization of intracellular Ebola virus nucleocapsid assembly by cryo-electron tomography. <i>Microscopy and Microanalysis</i> , 2021 , 27, 1708-1711	0.5	1	
2	Revealing the Native Molecular Architecture of the Nuclear Periphery using Cryo-Focused-Ion-Beam Milling, Light Microscopy and Electron Tomography. <i>Microscopy and Microanalysis</i> , 2017 , 23, 1248-1249	0.5		
1	The in Situ structure of a Parkinson's Disease Mutant LRRK2 Bound to Cellular Microtubules Revealed by Cryo-electron Tomography. <i>Microscopy and Microanalysis</i> , 2020 , 26, 800-802	0.5		