

# Yu-Ling Shih

## List of Publications by Year in descending order

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31  
papers

1,953  
citations

394421

19  
h-index

501196

28  
g-index

32  
all docs

32  
docs citations

32  
times ranked

1803  
citing authors

#	ARTICLE	IF	CITATIONS
1	Probing bacterial cell wall growth by tracing wall-anchored protein complexes. <i>Nature Communications</i> , 2021, 12, 2160.	12.8	6
2	Harnessing Fluorescent Moenomycin A Antibiotics for Bacterial Cell Wall Imaging Studies. <i>ChemBioChem</i> , 2021, 22, 3462-3468.	2.6	1
3	Effector loading onto the VgrG carrier activates type VI secretion system assembly. <i>EMBO Reports</i> , 2020, 21, e47961.	4.5	47
4	Peptidoglycan Endopeptidase Spr of Uropathogenic <i>Escherichia coli</i> Contributes to Kidney Infections and Competitive Fitness During Bladder Colonization. <i>Frontiers in Microbiology</i> , 2020, 11, 586214.	3.5	5
5	Active Transport of Membrane Components by Self-Organization of the Min Proteins. <i>Biophysical Journal</i> , 2019, 116, 1469-1482.	0.5	17
6	Quantitative inner membrane proteome datasets of the wild-type and the $\Delta$ min mutant of <i>Escherichia coli</i> . <i>Data in Brief</i> , 2016, 8, 304-307.	1.0	1
7	Involvement of type VI secretion system in secretion of iron chelator pyoverdine in <i>Pseudomonas taiwanensis</i> . <i>Scientific Reports</i> , 2016, 6, 32950.	3.3	60
8	A Multivalent Marine Lectin from <i>Crenomytilus grayanus</i> Possesses Anti-cancer Activity through Recognizing Globotriose Gb3. <i>Journal of the American Chemical Society</i> , 2016, 138, 4787-4795.	13.7	51
9	Quantitative Proteomics Analysis Reveals the Min System of <i>Escherichia coli</i> Modulates Reversible Protein Association with the Inner Membrane. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 1572-1583.	3.8	22
10	Study of Min Protein-Induced Membrane Waves in vitro. <i>Biophysical Journal</i> , 2015, 108, 78a.	0.5	0
11	Mitochondrial Genome Maintenance 1 (Mgm1) Protein Alters Membrane Topology and Promotes Local Membrane Bending. <i>Journal of Molecular Biology</i> , 2015, 427, 2599-2609.	4.2	25
12	Atomic Force Microscopy Characterization of Protein Fibrils Formed by the Amyloidogenic Region of the Bacterial Protein MinE on Mica and a Supported Lipid Bilayer. <i>PLoS ONE</i> , 2015, 10, e0142506.	2.5	17
13	Self-Assembly of MinE on the Membrane Underlies Formation of the MinE Ring to Sustain Function of the <i>Escherichia coli</i> Min System. <i>Journal of Biological Chemistry</i> , 2014, 289, 21252-21266.	3.4	18
14	Spatial control of the cell division site by the Min system in <i>Escherichia coli</i> . <i>Environmental Microbiology</i> , 2013, 15, 3229-3239.	3.8	27
15	Mgm1 Alters Membrane Topology and Promotes Local Membrane Bending to Drive Mitochondrial Membrane Fusion. <i>Biophysical Journal</i> , 2013, 104, 98a.	0.5	0
16	The N-Terminal Amphipathic Helix of the Topological Specificity Factor MinE Is Associated with Shaping Membrane Curvature. <i>PLoS ONE</i> , 2011, 6, e21425.	2.5	39
17	Direct MinE-membrane interaction contributes to the proper localization of MinDE in <i>E. coli</i> . <i>Molecular Microbiology</i> , 2010, 75, 499-512.	2.5	82
18	Assembly of the MreA-associated cytoskeletal ring of <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2009, 72, 170-182.	2.5	79

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19	Analyses of dynamic properties of the MinD cytoskeleton. <i>FASEB Journal</i> , 2008, 22, 262-262.	0.5	0
20	Polar positional information in <i>Escherichia coli</i> spherical cells. <i>Biochemical and Biophysical Research Communications</i> , 2007, 353, 493-500.	2.1	11
21	The Bacterial Cytoskeleton. <i>Microbiology and Molecular Biology Reviews</i> , 2006, 70, 729-754.	6.6	225
22	The MreB and Min cytoskeletal-like systems play independent roles in prokaryotic polar differentiation. <i>Molecular Microbiology</i> , 2005, 58, 917-928.	2.5	103
23	Spatial control of bacterial division-site placement. <i>Nature Reviews Microbiology</i> , 2005, 3, 959-968.	28.6	249
24	Division site selection in <i>Escherichia coli</i> involves dynamic redistribution of Min proteins within coiled structures that extend between the two cell poles. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 7865-7870.	7.1	331
25	Division site placement in <i>E.coli</i> : mutations that prevent formation of the MinE ring lead to loss of the normal midcell arrest of growth of polar MinD membrane domains. <i>EMBO Journal</i> , 2002, 21, 3347-3357.	7.8	106
26	The hexA gene of <i>Erwinia carotovora</i> encodes a LysR homologue and regulates motility and the expression of multiple virulence determinants. <i>Molecular Microbiology</i> , 2002, 28, 705-717.	2.5	106
27	Polar Explorers. <i>Cell</i> , 2001, 106, 13-16.	28.9	55
28	The MinE ring required for proper placement of the division site is a mobile structure that changes its cellular location during the <i>Escherichia coli</i> division cycle. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 980-985.	7.1	126
29	Structural basis for the topological specificity function of MinE. <i>Nature Structural Biology</i> , 2000, 7, 1013-1017.	9.7	75
30	The hexY genes of <i>Erwinia carotovora</i> ssp. <i>carotovora</i> and ssp. <i>atroseptica</i> encode novel proteins that regulate virulence and motility co-ordinately. <i>Environmental Microbiology</i> , 1999, 1, 535-547.	3.8	27
31	Generalized transduction in the potato blackleg pathogen <i>Erwinia carotovora</i> subsp. <i>atroseptica</i> by bacteriophage M1. <i>Microbiology (United Kingdom)</i> , 1997, 143, 2433-2438.	1.8	41