

Two-way communication between SecY and SecA suggests a mechanism for protein translocation

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Composition and Activity of the Non-canonical Gram-positive SecY2 Complex. <i>Journal of Biological Chemistry</i> , 2016, 291, 21474-21484.	1.6	10
2	Membrane protein insertion and assembly by the bacterial holo-translocon SecYEG- SecDF- YajC- YidC. <i>Biochemical Journal</i> , 2016, 473, 3341-3354.	1.7	61
3	Protein Translocation: SecA- SecY Conformational Crosstalk Opens Channel. <i>Current Biology</i> , 2016, 26, R811-R813.	1.8	1
4	Allosteric Signaling Is Bidirectional in an Outer-Membrane Transport Protein. <i>Biophysical Journal</i> , 2016, 111, 1908-1918.	0.2	16
5	Biphasic actions of SecA inhibitors on Prl/Sec suppressors: Possible physiological roles of SecA-only channels. <i>Biochemical and Biophysical Research Communications</i> , 2017, 482, 296-300.	1.0	2
6	<sc>S</sc>ec<sc>A</sc> functions <i>in vivo</i> as a discrete anti-parallel dimer to promote protein transport. <i>Molecular Microbiology</i> , 2017, 103, 439-451.	1.2	18
7	Targeting and Insertion of Membrane Proteins. <i>EcoSal Plus</i> , 2017, 7, .	2.1	63
8	Structural and Mechanistic Insights into Protein Translocation. <i>Annual Review of Cell and Developmental Biology</i> , 2017, 33, 369-390.	4.0	258
9	Substrate translocation involves specific lysine residues of the central channel of the conjugative coupling protein TrwB. <i>Molecular Genetics and Genomics</i> , 2017, 292, 1037-1049.	1.0	6
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11	Transmembrane β -barrels: Evolution, folding and energetics. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2017, 1859, 2467-2482.	1.4	49
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14	SecA- a New Twist in the Tale. <i>Journal of Bacteriology</i> , 2017, 199, .	1.0	8
15	Protein export through the bacterial Sec pathway. <i>Nature Reviews Microbiology</i> , 2017, 15, 21-36.	13.6	332
16	Dissecting structures and functions of SecA-only protein-conducting channels: ATPase, pore structure, ion channel activity, protein translocation, and interaction with SecYEG/SecDF- YajC. <i>PLoS ONE</i> , 2017, 12, e0178307.	1.1	3
17	Structurally detailed coarse-grained model for Sec-facilitated co-translational protein translocation and membrane integration. <i>PLoS Computational Biology</i> , 2017, 13, e1005427.	1.5	22
18	Mass spectrometry-enabled structural biology of membrane proteins. <i>Methods</i> , 2018, 147, 187-205.	1.9	69

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19	Driving Forces of Translocation Through Bacterial Translocon SecYEG. <i>Journal of Membrane Biology</i> , 2018, 251, 329-343.	1.0	27
20	Structure-based working model of SecDF, a proton-driven bacterial protein translocation factor. <i>FEMS Microbiology Letters</i> , 2018, 365, .	0.7	41
21	Single-molecule observation of nucleotide induced conformational changes in basal SecA-ATP hydrolysis. <i>Science Advances</i> , 2018, 4, eaat8797.	4.7	23
22	Large conformational changes of a highly dynamic pre-protein binding domain in SecA. <i>Communications Biology</i> , 2018, 1, 130.	2.0	14
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24	Specific cardiolipin- SecY interactions are required for proton-motive force stimulation of protein secretion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 7967-7972.	3.3	65
25	Dynamic action of the Sec machinery during initiation, protein translocation and termination. <i>ELife</i> , 2018, 7, .	2.8	52
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41	Structural Basis of the Sec Translocon and YidC Revealed Through X-ray Crystallography. <i>Protein Journal</i> , 2019, 38, 249-261.	0.7	13
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43	SecA-Mediated Protein Translocation through the SecYEG Channel. , 0, , 13-28.		0
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