

Structure of V-ATPase from the mammalian brain

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Structures of a Complete Human V-ATPase Reveal Mechanisms of Its Assembly. <i>Molecular Cell</i> , 2020, 80, 501-511.e3.	4.5	88
2	The importance of the membrane for biophysical measurements. <i>Nature Chemical Biology</i> , 2020, 16, 1285-1292.	3.9	25
3	Comparative Multiplexed Interactomics of SARS-CoV-2 and Homologous Coronavirus Nonstructural Proteins Identifies Unique and Shared Host-Cell Dependencies. <i>ACS Infectious Diseases</i> , 2020, 6, 3174-3189.	1.8	92
4	Cryo-EM structures of intact V-ATPase from bovine brain. <i>Nature Communications</i> , 2020, 11, 3921.	5.8	46
5	An affinity change model to elucidate the rotation mechanism of V1-ATPase. <i>Biochemical and Biophysical Research Communications</i> , 2020, 533, 1413-1418.	1.0	1
6	Lysosome as a Central Hub for Rewiring PH Homeostasis in Tumors. <i>Cancers</i> , 2020, 12, 2437.	1.7	44
7	The interaction partners of (pro)renin receptor in the distal nephron. <i>FASEB Journal</i> , 2020, 34, 14136-14149.	0.2	7
8	A "Sugar-Coated" Proton Pump Comes into Focus: High-Resolution Structure of a Human V-ATPase. <i>Molecular Cell</i> , 2020, 80, 379-380.	4.5	3
9	Native mass spectrometry "A valuable tool in structural biology. <i>Journal of Mass Spectrometry</i> , 2020, 55, e4578.	0.7	45
10	The Lysosome at the Intersection of Cellular Growth and Destruction. <i>Developmental Cell</i> , 2020, 54, 226-238.	3.1	77
11	Haploinsufficiency of <i>ATP6V0C</i> possibly underlies 16p13.3 deletions that cause microcephaly, seizures, and neurodevelopmental disorder. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 196-202.	0.7	9
12	Loss of <i>Furin</i> in β -Cells Induces an mTORC1-ATF4 Anabolic Pathway That Leads to β -Cell Dysfunction. <i>Diabetes</i> , 2021, 70, 492-503.	0.3	20
13	Transformative Network Modeling of Multi-omics Data Reveals Detailed Circuits, Key Regulators, and Potential Therapeutics for Alzheimer's Disease. <i>Neuron</i> , 2021, 109, 257-272.e14.	3.8	108
14	Native Mass Spectrometry-Based Screening for Optimal Sample Preparation in Single-Particle Cryo-EM. <i>Structure</i> , 2021, 29, 186-195.e6.	1.6	19
15	Expanding the clinical and molecular spectrum of <i>ATP6V1A</i> related metabolic cutis laxa. <i>Journal of Inherited Metabolic Disease</i> , 2021, 44, 972-986.	1.7	7
16	DOORS syndrome and a recurrent truncating <i>ATP6V1B2</i> variant. <i>Genetics in Medicine</i> , 2021, 23, 149-154.	1.1	11
17	Endolysosomal Disorders Affecting the Proximal Tubule of the Kidney: New Mechanistic Insights and Therapeutics. <i>Reviews of Physiology, Biochemistry and Pharmacology</i> , 2021, , 233-257.	0.9	2
18	Characterization of the T4 gp32 ssDNA complex by native, cross-linking, and ultraviolet photodissociation mass spectrometry. <i>Chemical Science</i> , 2021, 12, 13764-13776.	3.7	3

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19	Membrane Proteins Structure and Mechanism of the Vacuolar H ⁺ -ATPase. , 2021, , 581-593.		2
21	Cross-linking mass spectrometry uncovers protein interactions and functional assemblies in synaptic vesicle membranes. Nature Communications, 2021, 12, 858.	5.8	26
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27	Strong ion difference: Questionable stewardship. Acta Physiologica, 2021, 233, e13667.	1.8	4
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31	RAVE and Rabconnectin-3 Complexes as Signal Dependent Regulators of Organelle Acidification. Frontiers in Cell and Developmental Biology, 2021, 9, 698190.	1.8	21
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42	Recent developments in gastroesophageal mesenchymal tumours. <i>Histopathology</i> , 2021, 78, 171-186.	1.6	9
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62	Detection and quantification of the vacuolar H ⁺ -ATPase using the <i>Legionella</i> effector protein SidK. <i>Journal of Cell Biology</i> , 2022, 221, .	2.3	16
63	Defective Cystinosin, Aberrant Autophagy, Endolysosome Pathways, and Storage Disease: Towards Assembling the Puzzle. <i>Cells</i> , 2022, 11, 326.	1.8	2

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65	Cryo-EM of the Yeast V _O Complex Reveals Distinct Binding Sites for Macrolide V-ATPase Inhibitors. <i>ACS Chemical Biology</i> , 2022, 17, 619-628.	1.6	4
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