

# Cryo-EM Structure of a KCNQ1/CaM Complex Reveals I Syndrome

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Ca <sup>2+</sup> -Calmodulin and PIP2 interactions at the proximal C-terminus of Kv7 channels. <i>Channels</i> , 2017, 11, 686-695.	1.5	28
2	A comprehensive structural model for the human KCNQ1/KCNE1 ion channel. <i>Journal of Molecular Graphics and Modelling</i> , 2017, 78, 26-47.	1.3	10
3	Calmodulin confers calcium sensitivity to the stability of the distal intracellular assembly domain of Kv7.2 channels. <i>Scientific Reports</i> , 2017, 7, 13425.	1.6	7
4	Chansporter complexes in cell signaling. <i>FEBS Letters</i> , 2017, 591, 2556-2576.	1.3	18
5	KCNE1 and KCNE3 modulate KCNQ1 channels by affecting different gating transitions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E7367-E7376.	3.3	38
6	Electron cryo-microscopy structure of a human TRPM4 channel. <i>Nature</i> , 2017, 552, 200-204.	13.7	161
7	PIP2 mediates functional coupling and pharmacology of neuronal KCNQ channels. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E9702-E9711.	3.3	42
8	Two missense mutations in KCNQ1 cause pituitary hormone deficiency and maternally inherited gingival fibromatosis. <i>Nature Communications</i> , 2017, 8, 1289.	5.8	33
9	“Divide and conquer” approach to the structural studies of multidomain ion channels by the example of isolated voltage sensing domains of human Kv2.1 and Nav1.4 channels. <i>Russian Journal of Bioorganic Chemistry</i> , 2017, 43, 634-643.	0.3	3
10	Human In Silico Drug Trials Demonstrate Higher Accuracy than Animal Models in Predicting Clinical Pro-Arrhythmic Cardiotoxicity. <i>Frontiers in Physiology</i> , 2017, 8, 668.	1.3	227
11	Phosphatidylinositol-4,5-bisphosphate is required for KCNQ1/KCNE1 channel function but not anterograde trafficking. <i>PLoS ONE</i> , 2017, 12, e0186293.	1.1	9
12	Inactivation gating of Kv7.1 channels does not involve concerted cooperative subunit interactions. <i>Channels</i> , 2018, 12, 89-99.	1.5	4
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14	Cryo-EM structure of the polycystic kidney disease-like channel PKD2L1. <i>Nature Communications</i> , 2018, 9, 1192.	5.8	45
15	Kv7.3 Compound Heterozygous Variants in Early Onset Encephalopathy Reveal Additive Contribution of C-Terminal Residues to PIP2-Dependent K <sup>+</sup> Channel Gating. <i>Molecular Neurobiology</i> , 2018, 55, 7009-7024.	1.9	21
16	Exploiting ion channel structure to assess rare variant pathogenicity. <i>Heart Rhythm</i> , 2018, 15, 890-894.	0.3	4
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18	A Calmodulin C-Lobe Ca <sup>2+</sup> -Dependent Switch Governs Kv7 Channel Function. <i>Neuron</i> , 2018, 97, 836-852.e6.	3.8	63

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20	How to Connect Cardiac Excitation to the Atomic Interactions of Ion Channels. <i>Biophysical Journal</i> , 2018, 114, 259-266.	0.2	9
21	Mechanisms of KCNQ1 channel dysfunction in long QT syndrome involving voltage sensor domain mutations. <i>Science Advances</i> , 2018, 4, eaar2631.	4.7	64
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29	Human Calmodulin Mutations. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 396.	1.4	81
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36	Four drug-sensitive subunits are required for maximal effect of a voltage sensor-targeted KCNQ opener. <i>Journal of General Physiology</i> , 2018, 150, 1432-1443.	0.9	19

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