## Distinct mechanisms of inhibition of Kv2 potassium channels by tetraethylammonium and RY785

## **Supplementary Information**

Shan Zhang<sup>1</sup>, Robyn Stix<sup>1,2</sup>, Esam A. Orabi<sup>1</sup>, Nathan Bernhardt<sup>1</sup>, José D. Faraldo-Gómez<sup>1\*</sup>

<sup>1</sup>Theoretical Molecular Biophysics Laboratory, National Heart, Lung and Blood Institute, National Institutes of Health, Bethesda, MD

<sup>2</sup>Molecular and Cell Biology Graduate Program, Johns Hopkins University, Baltimore, MD

\*Correspondence should be addressed to: jose.faraldo@nih.gov

July 25<sup>th</sup>, 2024



**Figure S1.** Chemical structures of RY785 and tetraethylammonium. Atom names and partial electronic charges are indicated.



**Figure S2.** Potential-energy curve for the  $C_{46}C_{42}N_{43}$  angle between 100° and 135°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S3.** Potential-energy curve for the  $C_{42}C_{39}N_{38}$  angle between 90° and 145°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S4.** Potential-energy curve for the  $C_{39}C_{42}N_{43}$  angle between 95° and 140°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S5.** Potential-energy curve for  $C_{42}C_{39}N_{40}$  angle between  $105^{\circ}$  and  $150^{\circ}$ , calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S6.** Potential-energy curve for the  $N_{40}C_{39}N_{38}$  angle between 95° and 135°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S7.** Potential-energy curve for the  $C_{25}N_{27}C_{29}C_{30}$  angle between 0° and 180°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S8.** Potential-energy curve for the  $N_{40}C_{39}C_{42}N_{43}$  angle between -200° and 0°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S9.** Potential-energy curve for the  $C_{39}C_{42}N_{43}C_{44}$  angle between  $120^{\circ}$  and  $240^{\circ}$ , calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S10.** Potential-energy curve for the  $C_{46}C_{42}N_{43}C_{44}$  angle between -60° and 60°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S11.** Potential-energy curve for the  $C_{42}C_{39}N_{40}C_{33}$  angle between -240° and -120°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S12.** Potential-energy curve for the  $N_{38}C_{39}N_{40}C_{33}$  angle between -60° and 60°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S13.** Potential-energy curve for the  $C_{42}C_{39}N_{38}C_{32}$  angle between 120° and 240°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S14.** Potential-energy curve for the  $C_{42}C_{39}N_{38}H_{41}$  angle between -60° and 60°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S15.** Potential-energy curve for the  $N_{40}C_{39}N_{38}C_{32}$  angle between -60° and 60°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S16.** Potential-energy curve for the  $N_{40}C_{39}N_{38}H_{41}$  angle between -240° and -120°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S17.** Potential-energy curve for the  $S_{45}C_{46}C_{42}C_{39}$  angle between -240° and -120°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray) by scanning. The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S18.** Potential-energy curves for the  $S_{45}C_{46}C_{42}N_{43}$  angle between -60° and 60°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S19.** Potential-energy curve for the  $H_{48}C_{46}C_{42}N_{43}$  angle between 120° and 240°, calculated with MP2/6-31G(d) (blue) and with default CGengFF (orange) or with our optimized forcefield (gray). The Y axis shows the energy (in kcal/mol) relative to the lowest-energy conformer.



**Figure S20.** Ab-initio optimized geometries of RY785-water complexes used in the calibration of our molecular-mechanics forcefield for RY785.



**Figure S21.** Interaction energies (in kcal/mol) for the RY785-water complexes shown in Fig. S20. QM data (Y-axis) are correlated with MM values calculated with default CGenFF (blue) and with our optimized forcefield (orange).

Table S1. Interaction distances (in Å) and Interaction energies (in kcal/mol) for the RY785-water
complexes shown in Fig. S20. QM data are compared with MM values calculated with default
CGenFF and with our optimized forcefield.

Complex	Eqm	r <sub>QM</sub>	EMM, def	<b>r</b> мм, def	Eмм, opt	<b>r</b> <sub>MM, opt</sub>
Α	-1.83	2.58	-1.76	2.60	-1.86	2.59
В	-2.12	2.51	-1.82	2.60	-1.90	2.60
С	-1.71	2.61	-3.56	2.53	-2.07	2.64
D	-2.34	2.47	-3.83	2.51	-2.32	2.62
E	-4.92	1.99	-6.70	1.83	-4.22	1.92
F	-3.35	2.14	-5.53	1.87	-3.88	1.94
G	-8.17	2.06	-7.15	1.88	-7.89	1.87
Н	-9.44	2.04	-7.11	1.88	-7.24	1.88
I	-5.56	2.09	-7.75	1.85	-3.90	1.97
J	-3.84	2.07	-9.76	1.83	-4.75	1.95
К	-3.81	2.32	-3.38	2.23	-3.84	2.22
L	-3.93	2.30	-3.34	2.24	-3.82	2.22
Μ	-2.06	2.31	-1.86	2.26	-2.04	2.25
Ν	-2.81	2.22	-2.20	2.22	-2.51	2.21
0	-0.67	2.86	-0.38	2.60	-0.40	2.62
Р	-0.59	2.92	-0.49	2.59	-0.48	2.60
Q	-0.40	2.94	-0.12	2.66	-0.29	2.67
R	-0.82	2.80	-0.29	2.58	-0.37	2.57