Supporting information for: Supporting Information: How do aminoadamantanes block the influenza M2 channel and how does resistance develop?

Hadas Leonov ^{a,b}, Peleg Astrahan^a, Miriam Krugliak, and Isaiah T. Arkin^{*}

Department of Biological Chemistry, The Alexander Silberman Institute of Life Sciences, The Hebrew University of Jerusalem, Edmund J. Safra Campus, Jerusalem, 91904 Isreal.

E-mail: arkin@huji.ac.il

^{*a*}These authors contributed equally to the work

^bPresent address: Max Planck Institute for biophysical Chemistry Department of Theoretical and Computational Biophysics Am Faßberg 11 D-37077 Göttingen, Germany.



Figure 1: Structures of 1-aminoadamantane (amantadine, left), 1-(1-adamantyl)ethylamine (rimantadine, center) and 1-hydroxyadamantane (adamantanol, left). Amantadine and rimantadine are shown in their protonated form.

^{*}To whom correspondence should be addressed



Figure 2: The C α -RMSD of the 20ns equilibration MD simulations for two M2 Singapore variants: wild-type (black) and V27A (red).



Figure 3: Growth of bacteria in the presence (black, n = 2) or absence (red, n = 4) of 700 μ M adamantanol. The error represent the standard deviation, and *n* is the number of independent trials.