

## Supplementary Information for

### Nicotine Binding to Brain Receptors Requires a Strong Cation- $\pi$ Interaction

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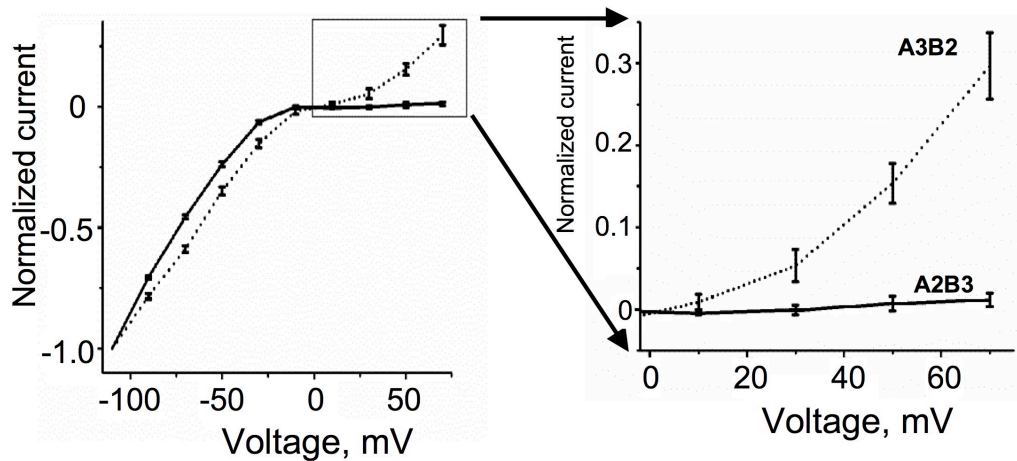
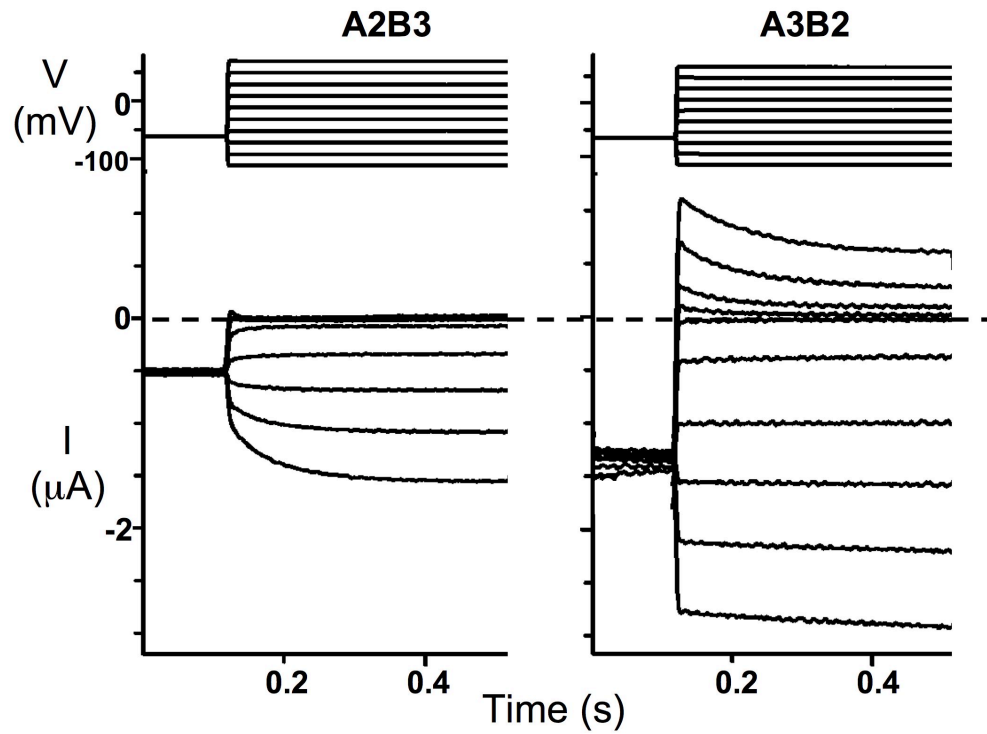
## Supplementary Figures

	Loop A					Loop B					Loop C																
$\alpha$ 1 mouse	W	R	P	D	V	V	L	Y	W	T	Y	D	G	S	V	V	Y	S	C	C	P	T	T	P	Y	L	D
$\alpha$ 1 human	W	R	P	D	L	V	L	Y	W	T	Y	D	G	S	V	V	Y	S	C	C	P	D	T	P	Y	L	D
$\alpha$ 2 human	W	I	P	D	I	V	L	Y	W	T	Y	D	K	A	K	I	Y	D	C	C	A	E	-	I	Y	P	D
$\alpha$ 4 human	W	R	P	D	I	V	L	Y	W	T	Y	D	K	A	K	I	Y	E	C	C	A	E	-	I	Y	P	D
$\alpha$ 4 rat	W	R	P	D	I	V	L	Y	W	T	Y	D	K	A	K	I	Y	E	C	C	A	E	-	I	Y	P	D
$\alpha$ 3 human	W	K	P	D	I	V	L	Y	W	S	Y	D	K	A	K	I	Y	N	C	C	E	E	-	I	Y	P	D
$\alpha$ 6 human	W	K	P	D	I	V	L	Y	W	T	Y	D	K	A	E	I	Y	N	C	C	E	E	-	I	Y	T	D
$\alpha$ 7 human	W	K	P	D	I	L	L	Y	W	S	Y	G	G	W	S	L	Y	E	C	C	K	E	-	P	Y	P	D
$\alpha$ 7 rat	W	K	P	D	I	L	L	Y	W	S	Y	G	G	W	S	L	Y	E	C	C	K	E	-	P	Y	P	D
$\alpha$ 9 human	W	R	P	D	I	V	L	Y	W	T	Y	N	G	N	Q	V	Y	G	C	C	S	E	-	P	Y	P	D

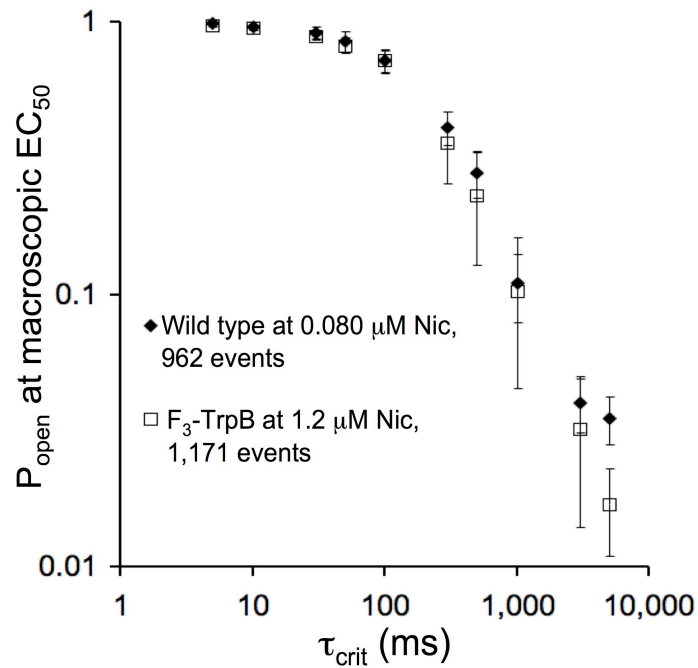
  

	Loop D					
$\gamma$ mouse	W	I	E	M	Q	W
$\gamma$ human	W	I	E	M	Q	W
$\delta$ mouse	W	I	D	H	A	W
$\delta$ human	W	I	E	H	G	W
$\beta$ 2 human	W	L	T	Q	E	W
$\beta$ 2 rat	W	L	T	Q	E	W
$\beta$ 3 human	W	L	K	Q	E	W
$\beta$ 4 human	W	L	K	Q	E	W
$\alpha$ 7 human	W	L	Q	M	S	W
$\alpha$ 7 rat	W	L	Q	M	S	W
$\alpha$ 9 human	W	I	R	Q	I	W

**Supplementary Figure 1.** Sequence alignment for Loops A, B, C, and D in the vicinity of the aromatic binding box. The five residues of the aromatic box: TyrA, TrpB, TryC1, TyrC2, and TrpD are highlighted in green. They are universally conserved in these subunits. G135 ( $\alpha$ 1) is the fourth residue after TrpB, highlighted in blue.



**Supplementary Figure 2.** Rectification behaviors of A2B3 and A3B2 L9'A  $\alpha 4\beta 2$  nAChR. Upper: Representative voltage traces and current responses for voltage jump experiments. Lower: I-V curves for A2B3 (solid line) and A3B2 (dotted line). The inset shows positive voltages, where A2B3 and A3B2 exhibit markedly different behavior.



**Supplementary Figure 3.** Comparison of  $P_{open}$  at macroscopic  $EC_{50}$  for wild type and F<sub>3</sub>-TrpB  $\alpha 4\beta 2$  (A2B3) over a range of  $\tau_{crit}$  values between 5 ms and 5000 ms reveals essentially equivalent gating behaviors. Error bars are mean  $\pm$  s.e.m. and are smaller than the symbol when not shown.

<b><math>\alpha 4(L9'A)\beta 2</math></b>						
<b>Mutation</b>	<b>ACh</b>	<b>n<sub>H</sub></b>	<b>Nicotine</b>	<b>n<sub>H</sub></b>	<b>Norm. I (+70mV)</b>	
<b>Wild type</b>						
<b>A2B3</b>	0.42 ± 0.01	1.2 ± 0.1	0.08 ± 0.01	1.2 ± 0.1	0.041 ± 0.005	
<b>A3B2</b>	0.023 ± 0.001	1.3 ± 0.1	0.01 ± 0.001	1.7 ± 0.2	0.297 ± 0.041	
<b>TyrA (Tyr98) A2B3</b>						
<b>Tyr</b>	0.42 ± 0.03	1.2 ± 0.1	0.08 ± 0.01	1.7 ± 0.3	0.023 ± 0.009	
<b>Phe</b>	12 ± 1	1.3 ± 0.1	0.77 ± 0.05	2.1 ± 0.3	0.064 ± 0.011	
<b>MeO-Phe</b>	2.3 ± 0.2	1.2 ± 0.1	0.40 ± 0.02	1.7 ± 0.2	0.054 ± 0.032	
<b>F-Phe</b>	15 ± 1	1.2 ± 0.1	0.32 ± 0.03	1.4 ± 0.2	-0.076 ± 0.046	
<b>F<sub>2</sub>-Phe</b>	16 ± 2	1.8 ± 0.3	0.39 ± 0.05	1.8 ± 0.4	0.028 ± 0.005	
<b>F<sub>3</sub>-Phe</b>	14 ± 1	1.2 ± 0.1	0.53 ± 0.04	1.4 ± 0.1	0.044 ± 0.010	
<b>Br-Phe</b>	3.3 ± 0.2	1.2 ± 0.1	0.54 ± 0.04	1.5 ± 0.1	-0.003 ± 0.031	
<b>CN-Phe</b>	73 ± 4	1.7 ± 0.1	8.8 ± 0.9	1.5 ± 0.2	0.075 ± 0.008	
<b>TrpB (Trp 154) A2B3</b>						
<b>Trp</b>	0.44 ± 0.03	1.3 ± 0.1	0.09 ± 0.01	1.5 ± 0.1	0.006 ± 0.014	
<b>F-Trp</b>	1.9 ± 0.1	1.2 ± 0.1	0.26 ± 0.02	1.3 ± 0.1	-0.065 ± 0.047	
<b>F<sub>2</sub>-Trp</b>	2.0 ± 0.1	1.3 ± 0.1	0.32 ± 0.04	1.3 ± 0.1	0.032 ± 0.025	
<b>F<sub>3</sub>-Trp</b>	13 ± 1	1.3 ± 0.1	1.2 ± 0.1	1.4 ± 0.2	-0.073 ± 0.029	
<b>F<sub>4</sub>-Trp</b>	29 ± 2	1.1 ± 0.1	4.2 ± 0.4	1.3 ± 0.2	-0.027 ± 0.023	
<b>CN-Trp</b>	12 ± 1	1.2 ± 0.1	0.90 ± 0.07	1.4 ± 0.1	0.009 ± 0.017	
<b>Br-Trp</b>	1.1 ± 0.1	1.3 ± 0.1	0.20 ± 0.02	1.3 ± 0.2	0.020 ± 0.005	
<b>TyrC1 (Tyr195) A2B3</b>						
<b>Tyr</b>	0.42 ± 0.03	1.5 ± 0.1	0.07 ± 0.01	1.3 ± 0.1	0.042 ± 0.014	
<b>Phe</b>	53 ± 4	1.3 ± 0.1	3.3 ± 0.2	1.2 ± 0.1	0.059 ± 0.014	
<b>MeO-Phe</b>	48 ± 5	1.4 ± 0.2	2.8 ± 0.4	1.2 ± 0.2	0.064 ± 0.028	
<b>CN-Phe</b>	210 ± 10	1.6 ± 0.1	19 ± 2	1.6 ± 0.2	0.057 ± 0.011	
<b>TyrC2 (Tyr202) A2B3</b>						
<b>Tyr</b>	0.42 ± 0.03	1.3 ± 0.1	0.09 ± 0.01	1.6 ± 0.1	0.057 ± 0.016	
<b>Phe</b>	0.32 ± 0.02	1.4 ± 0.1	0.14 ± 0.01	1.4 ± 0.1	0.014 ± 0.010	
<b>MeO-Phe</b>	0.33 ± 0.02	1.3 ± 0.1	0.097 ± 0.006	1.7 ± 0.2	0.034 ± 0.033	
<b>CN-Phe</b>	0.42 ± 0.04	1.4 ± 0.2	0.11 ± 0.01	1.6 ± 0.2	0.066 ± 0.046	
<b>Thr (B+1) (Thr 155) A2B3</b>						
<b>Thr</b>	0.41 ± 0.02	1.4 ± 0.1	0.09 ± 0.01	1.6 ± 0.1	0.044 ± 0.007	
<b>Tah</b>	0.37 ± 0.02	1.3 ± 0.1	1.71 ± 0.14	1.2 ± 0.1	0.018 ± 0.013	
<b>Muscle-type Receptor<sup>a</sup></b>						
<b>Thr (B+1) (Thr150)<sup>b</sup></b>						
<b>Thr</b>	0.83 ± 0.04	1.8 ± 0.1	57 ± 2	2.1 ± 0.1	ND	
<b>Tah</b>	0.25 ± 0.01	1.4 ± 0.1	92 ± 4	1.7 ± 0.1	ND	

<b><math>\alpha 1(\text{G153K})</math></b>													
<b>Trp</b>	0.019	±	0.001	1.5	±	0.1	0.59	±	0.04	1.8	±	0.2	ND
<b>F-Trp</b>	0.094	±	0.004	1.6	±	0.1	2.8	±	0.1	1.3	±	0.1	ND
<b>F<sub>2</sub>-Trp</b>	0.079	±	0.004	1.3	±	0.1	2.3	±	0.1	1.3	±	0.1	ND
<b>F<sub>3</sub>-Trp</b>	1.05	±	0.03	1.3	±	0.1	11	±	1	1.5	±	0.1	ND
<b>F<sub>4</sub>-Trp</b>	7.5	±	0.5	1.2	±	0.1	32	±	4	1.5	±	0.2	ND
<b>CN-Trp</b>	2.4	±	0.1	1.5	±	0.1	36	±	3	1.7	±	0.2	ND
<b>Br-Trp</b>	0.047	±	0.001	1.4	±	0.1	4.45	±	0.42	1.2	±	0.1	ND

**Supplementary Table 1.** EC<sub>50</sub> values ( $\mu\text{M}$ ), Hill coefficients ( $n_H$ ) and current size at +70 mV (normalized to current size at -110 mV). ND = not determined. a. All studies of the muscle-type receptor contain a L9'S mutation in the  $\beta$  subunit. b. These values were previously reported<sup>25</sup>.

## Supplementary Discussion

### **Controlling the Stoichiometry of $\alpha 4\beta 2$ Receptors**

As in the case of previous studies, we find that the stoichiometry of  $\alpha 4\beta 2$  receptors can be controlled by altering the ratio of the subunits of mRNA during injection. Our criteria for defining a pure population of A2B3  $\alpha 4(L9'A)\beta 2$  receptors are whole-cell dose-response curves that fit a single component and very strong inward rectification such that  $(I_{\max}$  at +70 mV)/(I<sub>max</sub> at -110 mV) < 0.1. An alternative analysis which can also demonstrate a mixed population of receptors is the production of intermediate EC<sub>50</sub> values when fit to a single component. As shown below, by a 3:1  $\alpha 4:\beta 2$  mRNA ratio, the EC<sub>50</sub> value has reached the higher EC<sub>50</sub> value, which is the A2B3 stoichiometry.

<b><math>\alpha 4:\beta 2</math> ratio</b>	<b>EC<sub>50</sub> (<math>\mu</math>M ACh)</b>
100:1	0.023 $\pm$ 0.002
10:1	0.023 $\pm$ 0.001
6:1	0.15 $\pm$ 0.02
3:1	0.44 $\pm$ 0.03
1:1	0.40 $\pm$ 0.01
1:10	0.43 $\pm$ 0.02

Injection of an mRNA ratio  $\alpha 4(L9'A):\beta 2$  of 10:1 or higher produces pure populations of A3B2, while a ratio of 1:3 or lower guarantees a pure population of A2B3. In the experiments described here, we injected a 1:3 ratio of mRNA.

Note that the  $\alpha 4(L9'A)$  mutation lowers EC<sub>50</sub> in a multiplicative fashion, depending on how many  $\alpha 4$  subunits are present. As such, our A3B2 receptor (with three L9'A mutations) actually has a lower EC<sub>50</sub> than our A2B3 receptor (with two L9'A mutations), even though the binding site from the A2B3 stoichiometry is clearly that of the high sensitivity receptor.

### **TyrA, TyrC1, and TyrC2 Display Similar Interactions in Muscle-type and $\alpha 4\beta 2$**

In addition to TrpB, we have performed extensive studies of other aromatic residues in and around the aromatic box (Supplementary Table 1). Briefly, when comparing  $\alpha 4\beta 2$  to the muscle-type receptor, very similar results are seen. TyrC1 is very sensitive to substitution, establishing a key role for this residue, likely in receptor gating. TyrA appears to be a hydrogen

bond donor (large effects for Phe and MeO-Phe substitutions), and while it is generally more sensitive to perturbations in the neuronal receptor, the basic trends are the same. TyrC2 is very permissive in both the muscle-type and  $\alpha 4\beta 2$  receptors.

### **Single-Channel Recording and Analysis**

Here we have used single-channel measurements to convincingly establish that the fluorination approach is changing agonist binding, not channel gating. Macroscopic data establish the large successive shift in function ( $EC_{50}$ ) upon fluorination, and single-channel data establish that gating is unperturbed, since the probability that the channel is open,  $P_{open}$ , is essentially indistinguishable for wild type and  $F_3$ -TrpB at corresponding points on the dose-response relation. At saturating agonist concentrations,  $P_{open,max}$  approaches  $\Theta/(\Theta+1)$ . Our analysis starts by comparing the  $P_{open}$  values at the macroscopic  $EC_{50}$ . The  $P_{open}$  values that we report are directly related to the gating equilibrium constant,  $\Theta$ , by  $\frac{1}{2} * \Theta/(\Theta+1)$ .

#### *Definition of clusters and calculation of $P_{open}$*

Because (a) single-channel channel measurements of  $P_{open}$  are seldom reported for  $\alpha 4\beta 2$  receptors, and (b) we find that  $P_{open}$  depends strongly on the value chosen as the critical closed duration,  $\tau_{crit}$ , we report  $P_{open}$  values for the range  $5 \text{ ms} \leq \tau_{crit} \leq 5000 \text{ ms}$  using two different methods to identify  $\tau_{crit}$  (below and Supplementary Fig. 3, above). The first is the commonly used method: the longest one or more components of the closed dwell time histogram are considered as sojourns in the desensitized state for all of the channels in the patch<sup>32</sup>. The value for  $\tau_{crit}$  was defined based on the closed dwell time histograms fitted with multiple components, as previously described<sup>33</sup>. These components are similar for wild type and  $F_3$ -TrpB, resulting in similar  $\tau_{crit}$  values:  $\tau_{crit1}$  of 1470 vs 1530 ms and  $\tau_{crit2}$  of 42 vs 52 ms, respectively. The similarity of the closed dwell time histograms for these receptors (and the resultant  $\tau_{crit}$  values) can be taken as evidence that fluorination does not significantly impact desensitization. Moreover, whole-cell data show that the wild type and the  $F_3$ -TrpB receptors exhibit similar extent and kinetics of macroscopic desensitization (data not shown). When either of the  $\tau_{crit}$  values calculated from the closed dwell time histogram is applied,  $P_{open}$  is essentially indistinguishable

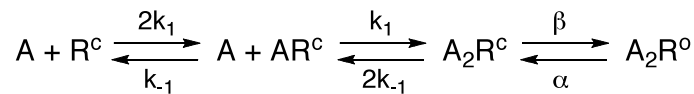
for wild type and F<sub>3</sub>-TrpB. The P<sub>open</sub> values for wild type and F<sub>3</sub>-TrpB at τ<sub>crit1</sub> and τ<sub>crit2</sub> are given here ± s.e.m.

<b>Receptor</b>	<b>P<sub>open</sub> (τ<sub>crit1</sub>)</b>	<b>P<sub>open</sub> (τ<sub>crit2</sub>)</b>
Wild type	0.07 ± 0.02	0.88 ± 0.06
F <sub>3</sub> -TrpB	0.06 ± 0.03	0.82 ± 0.05

Because our recordings are at an intermediate concentration (EC<sub>50</sub>), some closed dwells may reflect agonist dissociation, others may reflect channel closure followed by re-opening without agonist dissociation, while still others may reflect sojourns in desensitized states of varying duration. As a result, a definition of τ<sub>crit</sub> can be distorted by the relatively low number of long non-conducting sojourns. Thus, we also compared calculated P<sub>open</sub> values for a wide range of possible τ<sub>crit</sub> values (3 orders of magnitude), including those calculated from the closed dwell time histogram. Supplementary Fig. 3 shows that, regardless of how we define τ<sub>crit</sub> (5 ms ≤ τ<sub>crit</sub> ≤ 5000 ms), no systematic difference in P<sub>open</sub> is observed between wild type and F<sub>3</sub>-TrpB with nicotine as agonist—their gating behaviors are essentially indistinguishable. Thus, the value chosen for τ<sub>crit</sub> does not affect our conclusion that the gating behavior, as measured by P<sub>open</sub>, is not significantly impacted upon fluorination in the F<sub>3</sub>-TrpB mutant.

*A small shift in the channel open duration does not account for the EC<sub>50</sub> shift of F<sub>3</sub>-TrpB*

Fits to open dwell time histograms reveal that the main component of the channel open duration, which accounted for >90% of the conductance in both wild type and F<sub>3</sub>-TrpB receptors, is shifted 2.4-fold, from 23 ms (wild type) to 9.6 ms (F<sub>3</sub>-TrpB). Because the closed dwell time histograms, fitted with multiple components, displayed similar contributions from the major components for wild type and F<sub>3</sub>-TrpB, interpreting the 2.4-fold shift in open duration in terms of the channel closing rate, α, would imply a modest 2.4-fold shift in Θ in the F<sub>3</sub>-TrpB receptor. We consider these results in terms of a standard, linear 4-state model with two sequential agonist binding steps followed by a gating step:



for which,

$$EC_{50} = \frac{K_D}{\sqrt{\Theta + 2} - 1}$$



where  $K_D$  is the equilibrium agonist dissociation constant ( $k_{-1}/k_1$ ) and  $\Theta$  is the gating equilibrium constant ( $\beta/\alpha$ ). We see that a 2.4-fold change in  $\Theta$  accounts for at most a 1.5-fold shift in  $EC_{50}$ . Thus, both comparison of  $P_{open}$  as well as consideration of kinetics, to the extent possible for data at  $EC_{50}$ , indicate that the overwhelming majority of the 15-fold increase in nicotine's  $EC_{50}$  in the F<sub>3</sub>-TrpB receptor versus wild type is caused by changes to binding rather than the subsequent conformational changes that open the channel. Taken together, macroscopic and single-channel experiments show that fluorination modulates nicotine binding in a way that is systematically correlated to the energy of a cation- $\pi$  interaction.

## Supplementary Notes

- <sup>32</sup> Sakmann, B., Patlak, J., and Neher, E., Single acetylcholine-activated channels show burst-kinetics in presence of desensitizing concentrations of agonist. *Nature* **286** (5768), 71 (1980).
- <sup>33</sup> Jackson, M. B. et al., Successive openings of the same acetylcholine receptor channel are correlated in open time. *Biophysical journal* **42** (1), 109 (1983).