

Corrections

Pallitto, Monica, and Regina M. Murphy. 2001. *Biophys. J.* 81:1805–1822.

On p. 1813, there is an error in Eq. 19. The correct equation is:

$$\frac{d\lambda_{f2}}{dt} = k_p[I]\{(n+1)^2[N] + 2\lambda_{f1} + \lambda_{f0}\} - k_{-p}\{n^2[f_{n+1}] + (2\lambda_{f1} - \lambda_{f0})\} - pk_{la}\lambda_{f2}\lambda_{f0}^{q-1} + \bar{k}'_{ij}\lambda_{f1}^2 \quad (19)$$

On p. 1815, there are errors in Table 2. The correct Table 2 is printed below.

TABLE 2 Model parameters

Parameter	Value
K_{MD}	$0.64 \pm 0.08 \mu M^{-1}$
k_M/k_I	$80 \pm 30 \mu M$
k_D/k_I	0.65 ± 0.15
k_n/k_p	$1.7 \pm 0.1 \times 10^{-9} \mu M^{-4}$
k_{la}	$4.7 \pm 0.3 \times 10^{-2} \mu M^{-2} h^{-1}$
$\delta\omega_{fil}$	$9.8 \pm 0.9 \times 10^{-10} cm \cdot rad$
$\delta\omega_{fib}$	$1.06 \pm 0.03 \times 10^{-8} cm \cdot rad$
n	6
p	6
q	3

Palm, Thomas, Sarah Graboski, Sarah E. Hitchcock-DeGregori, and Norma J. Greenfield. 2001. *Biophys. J.* 81:2827–2837.

On p. 2829, there are errors in Table 1. The correct Table 1 is printed below.

TABLE 1 Oligonucleotide primers used to prepare Gly-hcTnT₇₀₋₁₇₀ and to introduce FHC mutations

Primer	Primer sequence	
A	5'-CATATGTCGTACTACCACCATCACCATCACGATTACGATATCCAAACGACCGAAAACC TGTATTTCAAGGCATGCTGACATAGAAGAGGTGGTGG-3'	
B	5'-P-TATATCTCCTCTTAAAGTTAACAAAATTATTC-3'	
C	5'-TAATAGGATCCATGCATTTGGGGTTACATCCAG-3'	
D	5'-P-ATCCTCAGCCTTCCCTGTTCTCC-3'	
E	5'-CCAACGACCGAAAAACCTGTATTTCAGGGCTCATGCCAACCTGGTGCCCTCCAAGATC-3'	
F	5'-GCTAGTTATTGCTCAGCGGTGGCAG-3'	
G	5'-CGATCCCGGAAATAATACGACTCAC-3'	
H	5'-GATCTGGGAGGCACCAAGTGGCATGAAGCCCTGAAAATACAGGTTTCGGTCGTTGG-3'	
Mutation	Coding primer sequence*	Restriction site
I79N	5'-CCCAACTTGGTACCTCCAAGAACCCGATGGAGAG-3'	<i>Kpn</i> I
R92Q	5'-GACTTTGATGATATCCACCA <u>Q</u> GAAGCGCATGGAG-3'	<i>Eco</i> RV
R92W	5'-GACTTTGATGATATCCAC <u>T</u> GAAGCGCATGGAG-3'	<i>Eco</i> RV
R92L	5'-GACTTTGATGATATCCAC <u>C</u> TAAGCGCATGGAG-3'	<i>Eco</i> RV
R94L	5'-CATCCACCGGAAG <u>C</u> TATGGAGAAGGACCTG-3'	<i>Hind</i> III
A104V	5'-GACCTGAATGAGCTGCAGGTGCTGATCGAGGCTAC <u>A</u> TTGAGAACAGG-3'	<i>Pst</i> I
F110I	5'-GACCTGAATGAGCTGCAGGC <u>I</u> CTGATCGAGGCTCAC <u>A</u> TTGAGAACAGG-3'	<i>Pst</i> I
ΔE160	5'-GCTGAAGAGAGAG <u>CT</u> CGACGA <u>G</u> AGGAGGAGAACAGGAGGAAG-3'	<i>Sac</i> I
E163K	5'-GCTGAAGAGAGAG <u>CT</u> CGAC <u>G</u> AGGAGGAG <u>A</u> GAACAGGAGG-3'	<i>Sac</i> I

*Only the coding primer is shown in this table. These primers were used together with primer F in one primary PCR reaction. The reverse complements of these primers were used together with primer G in the second primary PCR. FHC mutations are underlined, mutations that introduce a silent restriction site are in italics.

Haugh, Jason M. 2002. *Biophys. J.* 82:591–604.

On p. 594, Table 1 is aligned incorrectly. The correct Table 1 is printed below.

TABLE 1 Dimensionless model parameters

Parameter	Definition*		Brief description	Estimated range†
	Two-state	Regulated supply		
κ		k_{RE}/D	Enzyme reaction rate constant	$10^{-3}-10^3$
η_R		$s^2 n_R$	Activated receptor density	$10^{-8}-10^{-1}$
η_{RE}		$s^2 n_{RE}$	Receptor-enzyme density	$10^{-8}-10^{-1}$
τ_{RE}		$D t_{RE}/s^2$	Receptor-enzyme lifetime	$10-10^7$
Da	$(k_a + k_i)s^2/D$	$k_c s^2/D$	Bulk membrane rate constant	$10^{-7}-10^{-1}$
β	0	$(n_{RT}/n_R) k_{RT}/s^2 R_{T,0}$	Enhancement of substrate supply	No estimate

*See Fig. 1 for illustrations of the various rate processes.

†Parameter ranges are calculated as follows: k_{RE} is estimated using a k_{cat}/K_M range of 10^4-10^8 (Ms) $^{-1}$ and dividing by a confinement layer of $\sim 3-10$ nm; n_R and n_{RE} are estimated as $1-10^6$ molecules in a $10^3\text{-}\mu\text{m}^2$ membrane; the rate constants k_i , k_c , and t_{RE}^{-1} are given a range spanning $0.01-100$ s $^{-1}$; other estimates are $s \sim 3-10$ nm, $D \sim 0.1-1$ $\mu\text{m}^2/\text{s}$.

On p. 595, Eq. 14 is incorrect. The correct Eq. 14 is:

$$\alpha = \kappa \frac{\Psi_{ss}(1)}{\bar{\Psi}_{ss}} + \frac{8}{\pi \tau_{RE}} \left[1 - \frac{\beta Da^*}{\kappa(1 + \beta \eta_R)} \right] \times \int_0^\infty \frac{[1 - e^{-(\lambda^2 + Da^*)\tau_{RE}}](\lambda^2 + Da^*)^{-2}\lambda d\lambda}{\left[J_0(\lambda) + \frac{2\pi\lambda J_1(\lambda)}{\kappa} \right]^2 + \left[Y_0(\lambda) + \frac{2\pi\lambda Y_1(\lambda)}{\kappa} \right]^2}. \quad (14)$$

On p. 598, Eq. 17 is incorrect. The correct Eq. 17 is:

$$\frac{\bar{n}_{S^*}}{n_{S,\text{tot}}} = \frac{k_a + k_{RE}^{\text{eff}} n_{RE}}{k_a + k_i + k_{RE}^{\text{eff}} n_{RE}} = \frac{\frac{k_a s^2}{D} + \alpha \eta_{RE}}{Da + \alpha \eta_{RE}}. \quad (17)$$

On p. 600, Eq. 18 is incorrect. The correct Eq. 18 is:

$$\frac{\text{rate}}{R_{T,0}} = \alpha \eta_{RE} \bar{\Psi}_{ss}; \bar{\Psi}_{ss} = \frac{1 + \beta \eta_R}{Da + \alpha \eta_{RE}}. \quad (18)$$