

CORRECTIONS

The primary structure of the calcium ion-transporting adenosine triphosphatase protein of rabbit skeletal sarcoplasmic reticulum: peptides derived from digestion with cyanogen bromide, and the sequences of three long extramembranous segments

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p. 614, Fig. 9, sequence 3:

for $\overset{100}{\text{B}}\overset{110}{\text{S}}\overset{120}{\text{L}}\text{LDFNETKGVYEKVGEA}\overset{120}{\text{D}}\overset{120}{\text{E}}\overset{120}{\text{T}}\overset{120}{\text{A}}$

read $\overset{100}{\text{B}}\overset{110}{\text{S}}\overset{120}{\text{L}}\text{LDFNETKGVYEKVGEA}\overset{120}{\text{T}}\overset{120}{\text{E}}\overset{120}{\text{T}}\overset{120}{\text{A}}$

The kinetic properties and reaction mechanism of histamine methyltransferase from human skin

D. M. FRANCIS, M. F. THOMPSON and M. W. GREAVES

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p. 825, Table 2, column 2:

for *S*-Adenosylmethione *read* *S*-Adenosylmethionine

p. 821, para. 1, l. 18:

for Substrae kinetic *read* substrate kinetics

The combination of the viscosity increment with the harmonic mean rotational relaxation time for determining the conformation of biological macromolecules in solution

S. E. HARDING

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p. 359, Eqn. 1:

$$\text{for } \frac{\tau_h}{\tau_0} = \frac{1}{3} \frac{\tau_a}{\tau_0} + \frac{2\tau_b}{\tau_0} \equiv \frac{\tau_h kT}{3\eta_0 V_e}$$

$$\text{read } \frac{\tau_h}{\tau_0} = \frac{3}{\left(\frac{\tau_0}{\tau_a} + \frac{2\tau_0}{\tau_b}\right)} \equiv \frac{\tau_h kT}{3\eta_0 V_e}$$

p. 360, Eqn. 5:

$$\text{for } \beta = \frac{N_A^\dagger}{(16\,200\pi^2)} v^\dagger \left(\frac{f_0}{f}\right) \equiv \frac{N_A s[\eta]^\dagger \eta_0}{M_r^\dagger (1 - \bar{v}\rho_0) 100^\dagger}$$

$$\text{read } \beta = \frac{N_A^\dagger}{(16\,200\pi^2)^\dagger} v^\dagger \left(\frac{f_0}{f}\right) \equiv \frac{N_A s[\eta]^\dagger \eta_0}{M_r^\dagger (1 - \bar{v}\rho_0) 100^\dagger}$$