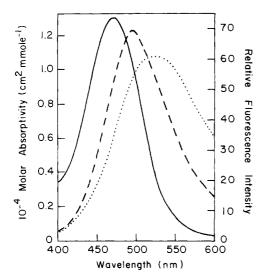
## **CORRECTIONS**

Spatial Relationship of the σ Subunit and the Rifampicin Binding Site in RNA Polymerase of *Escherichia coli*, by Cheng-Wen Wu,\* Lynwood R. Yarbrough, Felicia Y.-H. Wu, and Zaharia Hillel, Volume 15, Number 10, May 18, 1976, pages 2097–2104.

The numerical scale and unit of Figure 7 are erroneous. The corrected figure is enclosed (other data and the conclusions in the paper are unaffected).



Cooperative and Noncooperative Binding of Pyridoxal 5'-Phosphate to Tryptophan Synthase from *Escherichia coli*, by Peter Bartholmes, Kasper Kirschner,\* and Hans-Peter

Gschwind, *Biochemistry*, Volume 15, Number 21, October 19, 1976, pages 4712-4717.

Equation 9 under Appendix should read:

$$\lim \left(\frac{\mathrm{d}(\nu/P)}{\mathrm{d}\nu}\right)_{P\to\infty} = -n\Psi_n/\Psi_{n-1} = -\frac{1}{K_{\mathrm{d},n}} \left(\mathrm{not} - K_{\mathrm{d},n}\right)$$

Adrenocorticotropic Hormone Regulation of Adrenal RNA Polymerases. Stimulation of Nuclear RNA Polymerase III, by Shella A. Fuhrman and Gordon N. Gill,\* Volume 15, Number 25, December 14, 1976, pages 5520–5527.

Page 5522, first column, last paragraph, the sentence beginning on line 3 should read: Data derived from seven experiments using three different nuclear preparations were compiled; the activity in control nuclei was  $33.3 \pm 9.8$  fmol of UMP incorporated per  $\mu g$  of nuclear DNA and the activity in ACTH nuclei was  $72.1 \pm 9.0$  fmol of UMP incorporated per  $\mu g$  of nuclear DNA ( $\rho < 0.01$ ).

Evidence for Negative Cooperativity in the Adsorption of Phosphorylase b on Hydrophobic Agaroses, by Herbert P. Jennissen, Volume 15, Number 26, December 28, 1976, pages 5683-5692.

On page 5686 in Table III the adsorption constant  $\alpha$  for Butyl-Sepharose, buffer A, 18 °C, should read 16.26. On page 5690 the footnote should read:  ${}^2K^*_{0.5}$  denotes the reciprocal of the alkyl residue density, i.e., concentration in (mol of alkyl residue/ml of packed Sepharose)<sup>-1</sup> symbolized by  $M_s^{-1}$  at half-maximal saturation with ligand.