**Correction.** In the article "Induction of ornithine decarboxylase activity by insulin and growth factors is mediated by amino acids" by Clifford A. Rinehart, Jr., and Evangelos S. Canellakis, which appeared in number 13, July 1985, of Proc. Natl. Acad. Sci. USA (82, 4365-4368), the following error should be noted. In the last line of the legend to Fig. 5 on p. 4367, the symbols should be reversed; the last line should read "○, Asparagine alone; ●, asparagine with insulin."

**Correction.** In the article "DNA polymorphic loci mapped to human chromosomes 3, 5, 9, 11, 17, 18, and 22" by S. L. Naylor, A. Y. Sakaguchi, D. Barker, R. White, and T. B. Shows, which appeared in number 8, April 1984, of Proc. Natl. Acad. Sci. USA (81, 2447-2451), the authors request the following changes be noted. Table 2 erroneously indicated that the REX-12 hybrid contained the X/22 translocation chromosome. This cell hybrid actually contains the 22/X translocation chromosome (22pter→22q13: Xq22→Xqter). The REX-12 hybrid is positive for sequences detected by the probe pMS3-18, which places pMS3-18 in the pter→q13 region of chromosome 22. Also, in Table 3, the regional location of p12-32 is incorrectly listed as 3q21→qter. p12-32 (gene symbol D3S2) is located in the p21→q21 region of chromosome 3 as the data indicate.

**Correction.** In the article "High-affinity-receptor-mediated uptake and degradation of glucose-modified proteins: A potential mechanism for the removal of senescent macromolecules" by Helen Vlassara, Michael Brownlee, and Anthony Cerami, which appeared in number 17, September 1985, of Proc. Natl. Acad. Sci. USA (82, 5588-5592), the authors wish that the following correction be noted. In both the abstract (line 16) and the second line from the bottom of the second column of text on p. 5590, the value of the affinity constant should be $1.75 \times 10^7 \text{M}^{-1}$.

**Correction.** In the article "Accurate transcription of cloned Neurospora RNA polymerase II-dependent genes in vitro by homologous soluble extracts" by Brett M. Tyler and Norman H. Giles, which appeared in number 16, August 1985, of Proc. Natl. Acad. Sci. USA (82, 5450-5454), the authors request that the following correction be noted. On p. 5450, in the right-hand column, line 8 of text from the bottom of the page, the dialysis buffer was incorrectly described as containing 10 mM K$_2$EDTA instead of 10 mM K$_2$EGTA. The correct composition of the dialysis buffer should read "20 mM HEPES, pH 7.9/10 mM K$_2$EGTA/10 mM MgSO$_4$/15% (vol/vol) glycerol/5 mM dithiothreitol." Substitution of K$_2$EDTA for K$_2$EGTA could lead to transcription extracts with poor activity.