

**Biochemistry.** In the article “Oxidative damage during aging targets mitochondrial aconitase” by Liang-Jun Yan, Rodney L. Levine, and Rajindar S. Sohal, which appeared in number 21, October 14, 1997, of *Proc. Natl. Acad. Sci. USA* (**94**, 11168–11172), the authors request that the following correction be noted. On page 11169, in the paragraph on enzyme activity assay, line 8, “isocitrate” should be “citrate.” The next sentence on method 2 should read: “Method 2 is based on the formation of both citrate and isocitrate from cis-aconitate . . . .” This same change applies to the description of the enzyme activity assay in the legend of Fig. 2A.

**Cell Biology.** In the article “The  $\alpha$  chain of laminin-1 is independently secreted and drives secretion of its  $\beta$ - and  $\gamma$ -chain partners” by Peter D. Yurchenco, Yong Quan, Holly Colognato, Todd Mathus, David Harrison, Yoshihiko Yamada, and Julian J. O’Rear, which appeared in number 19, September 16, 1997, of *Proc. Natl. Acad. Sci. USA* (**94**, 10189–10194), the following corrections should be noted. (i) In Fig. 2a and c, on the left side of the immunoblots, migration position “ $\alpha/\gamma$ ” should read “ $\beta/\gamma$ .” (ii) In Fig. 2a, eighth lane, the lower label (medium fraction, anti-E8- $\beta\gamma$ ) should read “ $\alpha \rightarrow \beta^*$ ” (i.e., transient transfection of a  $\beta$  stable clone with  $\alpha$  cDNA).

**Cell Biology.** The title of the article “Smad8 mediates the signaling of the receptor serine kinase” by Yan Chen, Anil Bhushan, and Wylie Vale, which appeared in number 24, November 25, 1997, of *Proc. Natl. Acad. Sci. USA* (**94**, 12938–12943), appeared incorrectly due to a printer’s error. The correct title is “Smad8 mediates the signaling of the ALK-2 receptor serine kinase.”

**Developmental Biology.** In the article “Proximo–distal specification in the wing disc of *Drosophila* by the *nubbin* gene” by Francisco J. Cifuentes and Antonio García-Bellido, which appeared in number 21, October 14, 1997, of *Proc. Natl. Acad. Sci. USA* (**94**, 11405–11410), the authors request that the following corrections to Fig. 6 be noted. A higher-contrast figure and its legend appear below.

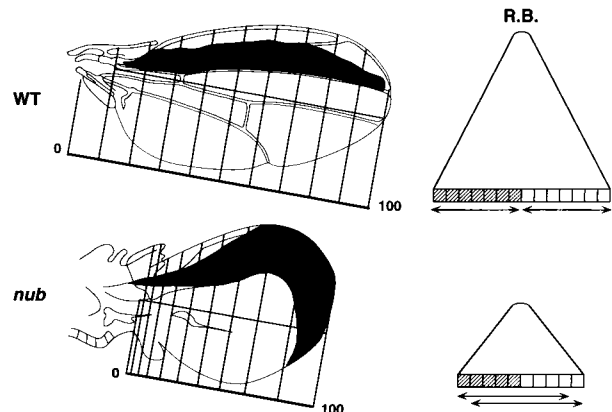


FIG. 6. Proximal–distal axis positional values and formation of compartment borders. In wild-type wings, the positional values along the proximal–distal axis (vertical bars) are correctly established by normal cells and the clones are elongated along the proximal–distal axis. Compartment boundaries (restriction boundaries = RB) in wild-type wings are the consequence of the high positional values reached by the cells in characteristic regions associated with the confrontation of cells expressing different selector genes (represented as squares with differential shading). The clones (horizontal arrows) never cross these compartment boundaries. In *nub* wings the proximal–distal positional values are not correctly established and *nub* cells are unable to reach the highest distal values, provoking the growth of the clones perpendicular to the A–P boundary. Clones in *nub* wings cross the compartment boundaries as a consequence of the inability of *nub* cells to meet the high positional values required for clonal restrictions.

**Evolution.** In the article “Accurate reconstruction of a known HIV-1 transmission history by phylogenetic tree analysis” by Thomas Leitner, David Escanilla, Christer Franzén, Mathias Uhlén, and Jan Albert, which appeared in number 20, October 1, 1996, of *Proc. Natl. Acad. Sci. USA* (**93**, 10864–10869), the authors request that the following correction be noted. When the epidemiological data for this investigation were collected, the study subjects were assigned codes to protect their identity. Unfortunately, one individual was not assigned the correct code, thus p11 should be p10. The phylogeny that describes the known transmission history has been corrected and should read (((p8.822, p8.159),(p10.113, p10.9939)), p9.256), (p7.6760,(p5.317, p6.6767),(p2.135,(p3.529, p3.105))), (p1.719, p1.136))))). The GenBank sequence files (U68496–U68521) also have been updated with the corrected labels and the correct topology. Furthermore, all results presented in the paper have been reevaluated and were found to be virtually identical to the published results. Thus, the published findings and conclusions remain unchanged. A detailed account for the reanalyses can be obtained from the authors. Dr. Leitner can be reached at his current address: Theoretical Biology and Biophysics, Group T-10, Mail Stop K710, Los Alamos National Laboratory, Los Alamos, NM 87545. e-mail: tk1@t10.lanl.gov.