

Biochemistry. In the article "A common ancestry for multiple catalytic antibodies generated against a single transition-state analog" by Hideaki Miyashita, Tomoko Hara, Ryuji Tamimura, Fujie Tanaka, Masakazu Kikuchi, and Ikuro Fujii, which appeared in number 13, June 1994, of *Proc. Natl.*

Acad. Sci. USA (91, 6045–6049), the following correction should be noted. Due to a printer's error, Figs. 2 and 3 on p. 6047 were transposed; the correct figures and their legends are shown below.

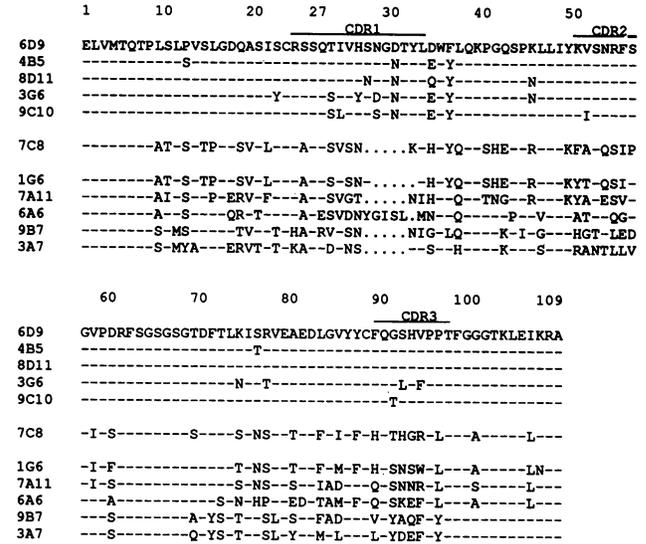
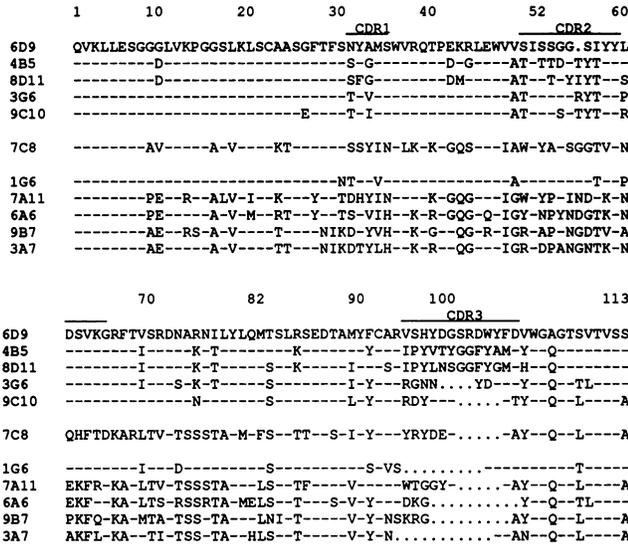


FIG. 2. Deduced amino acid sequence comparison of V_H cDNA generated from 11 anti-CPP monoclonal antibodies. Dashes denote identity to antibody 6D9 and dots, used to align sequences, represent no amino acid residue at a position. CDR designations and amino acid residue numbering are as described by Kabat *et al.* (14).

FIG. 3. Comparison of the deduced amino acid sequences of V_L for 11 anti-CPP monoclonal antibodies. See legend to Fig. 2 for symbol designation.

Biophysics. In the article "Excited-state intramolecular proton transfer as a fluorescence probe for protein binding-site static polarity" by Alexander Sytnik and Michael Kasha, which appeared in number 18, August 1994, of *Proc. Natl. Acad. Sci. USA* (91, 8627–8630), the following acknowledgement was inadvertently omitted by the authors: "This research was supported under Contract DE-FG05-87ER60517 between the Division of Biomedical and Environmental Research and the Florida State University."

Cell Biology. In the article "An essential gene of *Saccharomyces cerevisiae* coding for an actin-related protein" by Masahiko Harata, Anneliese Karwan, and Ulrike Wintersberger, which appeared in number 17, August 16, 1994, of *Proc. Natl. Acad. Sci. USA* (91, 8258–8262), the authors request that the following change be noted. On page 8258, in line 17 of the *Abstract*, "hydrophobic" should be replaced by "hydrophilic."