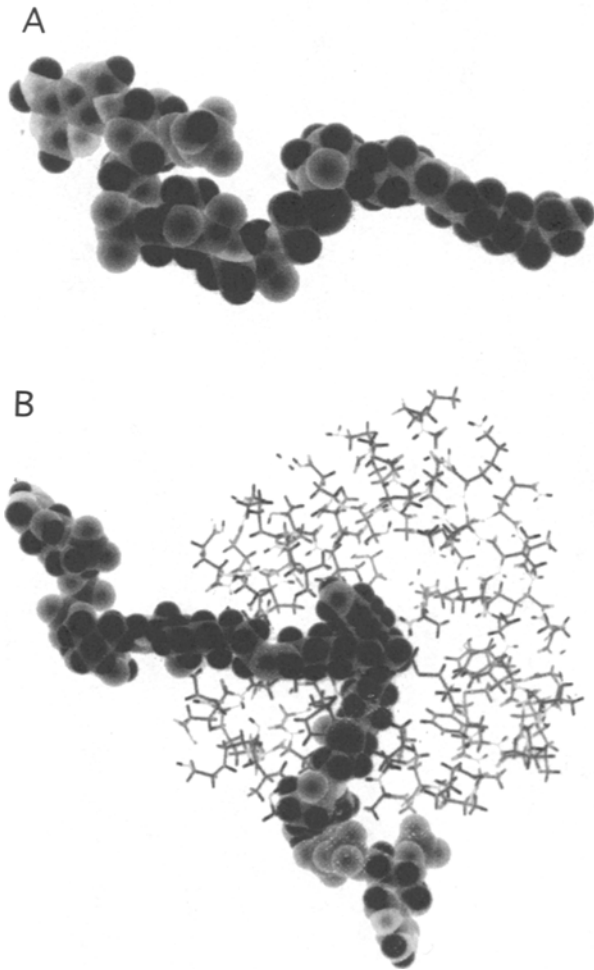
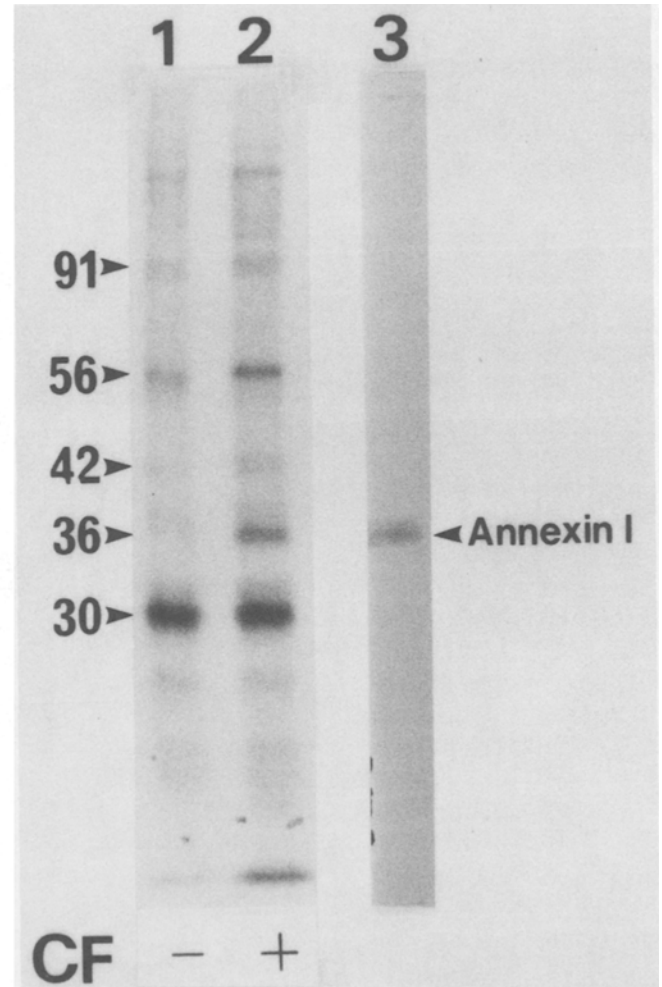


# ERRATUM

Due to a printer error in the September issue of *Lipids*, the photographs on pages 895 and 985 were transposed.



**FIG. 1.** Structure of free (panel A) and liver fatty acid binding protein bound (panel B) oleoyl CoA. These structures were prepared by Silicon Graphics (Mountain View, CA) computer modeling as described elsewhere (Ref. 127).



**FIG. 3.** Western blot analysis of the 36 kDa particulate protein using anti-annexin I. Lanes 1–2, autoradiography; lane 3, the blotting: CF, the protein kinase C cofactors (5  $\mu\text{g}/\text{mL}$  1-oleoyl-2-acetyl-*sn*-glycerol, 25  $\mu\text{g}/\text{mL}$  phosphatidylserine, and 1  $\mu\text{M}$   $\text{Ca}^{2+}$ ).