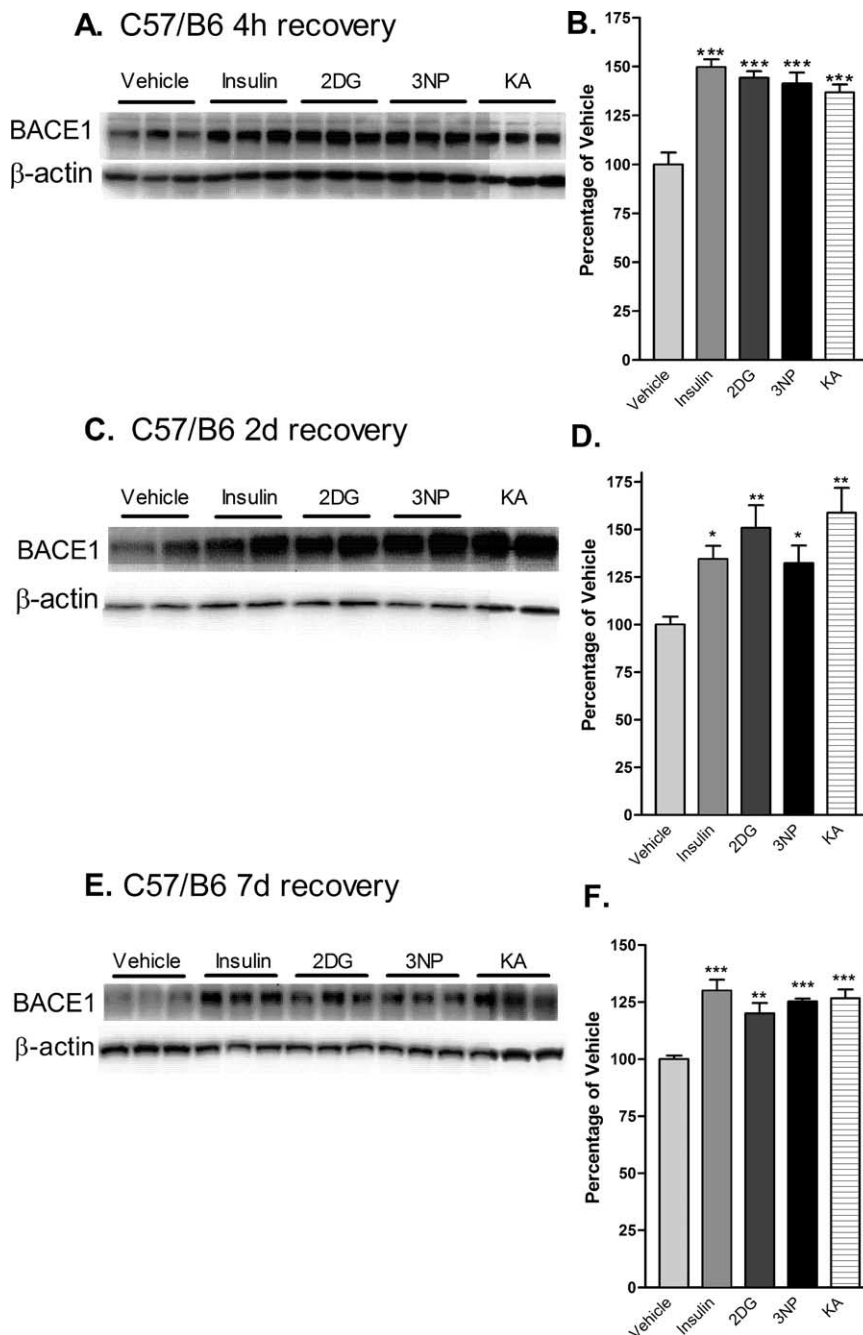
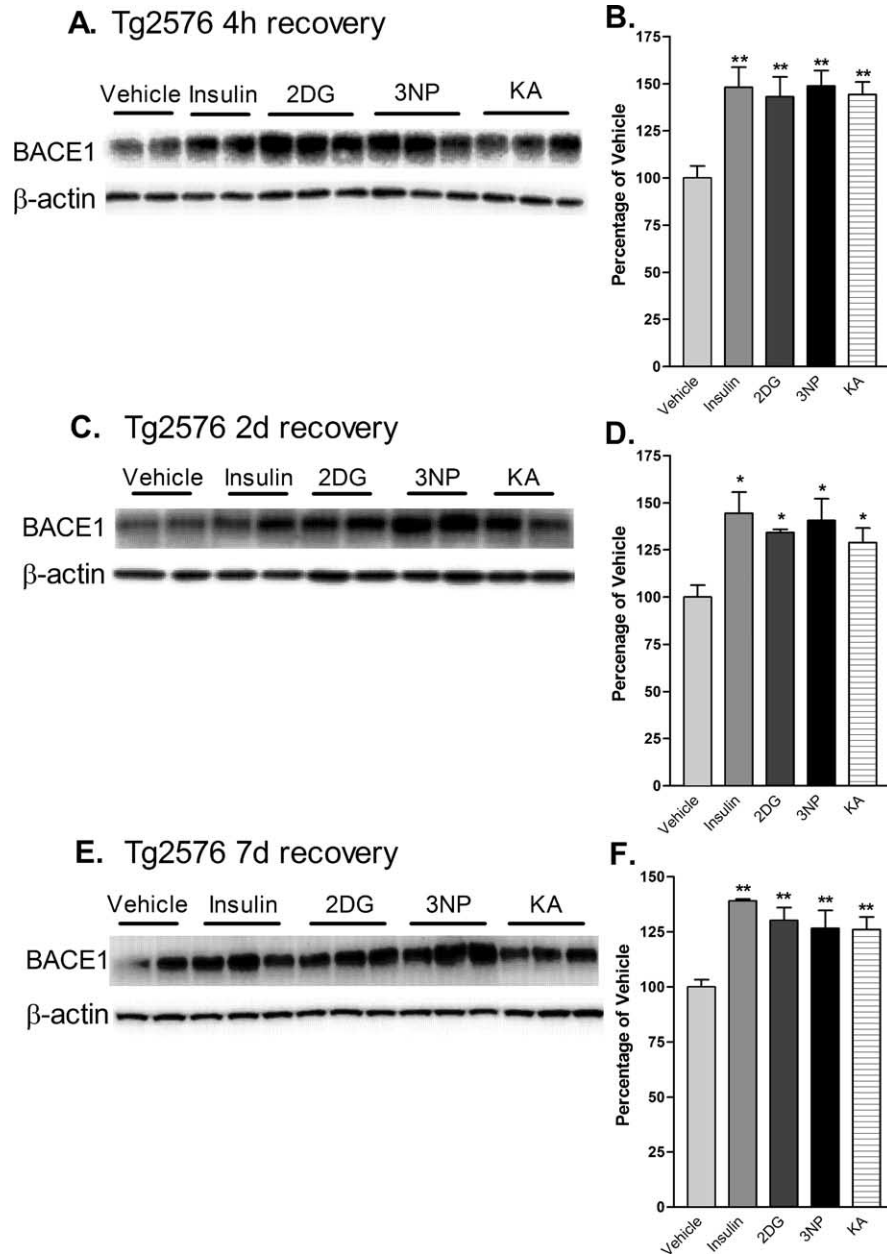


# Erratum

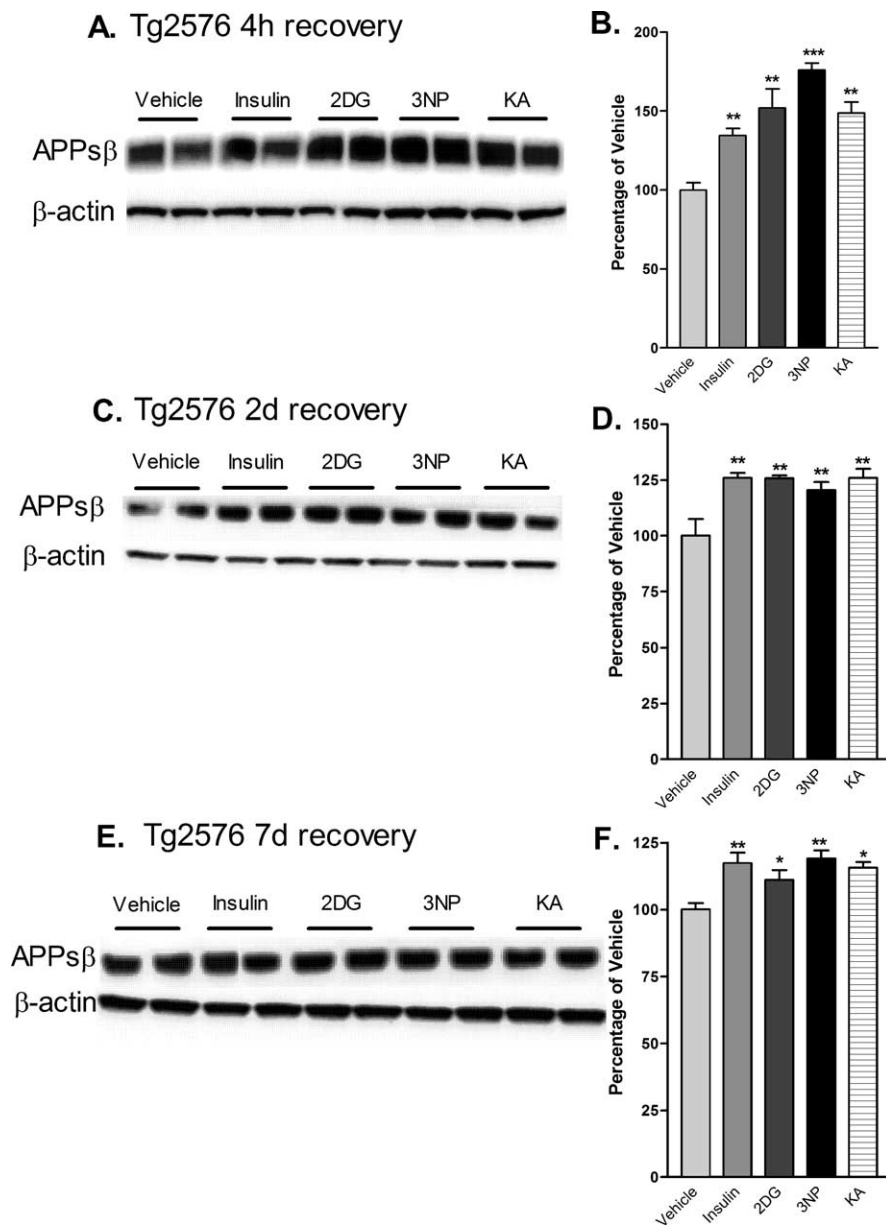
In the article “Energy Inhibition Elevates  $\beta$ -Secretase Levels and Activity and Is Potentially Amyloidogenic in APP Transgenic Mice: Possible Early Events in Alzheimer’s Disease Pathogenesis,” by Rodney A. Velliquette, Tracy O’Connor, and Robert Vassar, which appeared on pages 10874–10883 of the November 23, 2005 issue, a low-resolution version of Figures 3, 4, and 5 was used in the printed issue. These figures and their legends are reprinted in this issue.



**Figure 3.** Acute energy inhibition increases BACE1 protein levels in the brains of C57/B6 mice. **A–F**, Two- to 3-month-old C57/B6 mice were given a single intraperitoneal injection of 18 U/kg insulin, 1 g/kg 2DG, 100 mg/kg 3NP, 30 mg/kg KA, or vehicle and were then allowed to recover for 4 h (**A, B**), 2 d (**C, D**) or 7 d (**E, F**). Brains were then isolated and homogenized, and samples (15  $\mu$ g/lane) were subjected to immunoblot analysis for BACE1 protein using anti-BACE1 antibody PA1–757. Blots were stripped and reprobed with anti- $\beta$ -actin antibody as a loading control. **A, C, E**, Representative BACE1 (top panels) and  $\beta$ -actin (bottom panels) immunoblots for the various treatments and recovery times are shown. **B, D, F**, The intensities of BACE1 band signals were quantified on a PhosphorImager (Eastman Kodak), normalized against the  $\beta$ -actin immunosignals for each sample, and then expressed as percentages of the mean of the vehicle control for a given recovery time. Note that the energy-inhibitor treatments elevated cerebral BACE1 protein levels to 125–150% of vehicle control values for all recovery times ( $p < 0.05$ ,  $**p < 0.01$ , and  $***p < 0.001$ , one-way ANOVA with Newman-Keuls multiple-comparison test). **A–D**, Data represent mean  $\pm$  SEM;  $n = 9$  mice/treatment (**A, B**),  $n = 5$  mice/treatment (**C, D**), and  $n = 4$  mice/treatment (**E, F**).



**Figure 4.** Acute energy inhibition increases BACE1 protein levels in the brains of Tg2576 mice. *A–F*, Two- to 3-month-old Tg2576 mice were injected with 18 U/kg insulin, 1 g/kg 2DG, 100 mg/kg 3NP, 30 mg/kg KA, or vehicle and were allowed to recover for 4 h (*A, B*), 2 d (*C, D*), or 7 d (*E, F*). *A, C, E*, Representative immunoblots of brain samples for BACE1 (top panels) and  $\beta$ -actin (bottom panels). *B, D, F*, BACE1 immunosignals were quantified, normalized against  $\beta$ -actin signals, and expressed as percentages of vehicle controls, as before. Similar to the effects observed in C57/B6 mice, energy-inhibitor treatments in Tg2576 mice caused cerebral BACE1 levels to increase to 125–150% of vehicle controls for all recovery times ( $*p < 0.05$  and  $**p < 0.01$ , one-way ANOVA with Newman-Keuls multiple-comparison test). Data represent mean  $\pm$  SEM;  $n = 6$  mice/treatment (*A, B*), and  $n = 4$  mice/treatment (*C–F*).



**Figure 5.** Cerebral levels of APPs(sw) are elevated after acute inhibition of energy production in Tg2576 mice. Brain homogenates from Tg2576 mice treated with 18 U/kg insulin, 1 g/kg 2DG, 100 mg/kg 3NP, 30 mg/kg KA, or vehicle were subjected to immunoblot analysis using an antibody raised against the C-terminal neopeptide generated by BACE1 cleavage of APPsw, which recognizes APPsβ(sw) (Seubert et al., 1993; Vassar et al., 1999; Cole et al., 2005). **A, C, E**, Representative APPsβ(sw) (top panels) and β-actin (bottom panels) immunoblots of brain samples from treated Tg2576 mice. Recovery times were for 4 h (**A, B**), 2 d (**C, D**) or 7 d (**E, F**). **B, D, F**, APPsβ(sw) immunosignals were normalized against β-actin signals and expressed as percentages of vehicle controls. Note that cerebral APPsβ(sw) levels were increased to ~125–175% of vehicle controls after 4 h of recovery from energy inhibition (**B**), and they continued to be significantly elevated after 2 d (**D**) and 7 d (**F**) of recovery (\* $p < 0.05$ , \*\* $p < 0.01$ , and \*\*\* $p < 0.001$ , one-way ANOVA with Newman-Keuls multiple-comparison test). Data represent mean ± SEM;  $n = 6$  mice/treatment (**A, B**), and  $n = 4$  mice/treatment (**C–F**).