

Corrections

DEVELOPMENTAL BIOLOGY. In the article “Transcription repression by *Xenopus* ET and its human ortholog TBX3, a gene involved in ulnar-mammary syndrome” by Ming-liang He, Leng Wen, Christine E. Campbell, Jane Y. Wu, and Yi Rao, which appeared in number 18, August 31, 1999, of *Proc. Natl. Acad. Sci. USA* (96,

10212–10217), the following changes are noted. In the legend to Fig. 1, the text explaining panels *D* and *E* was omitted due to a printer’s error. Fig. 1 and the complete legend are reproduced below.

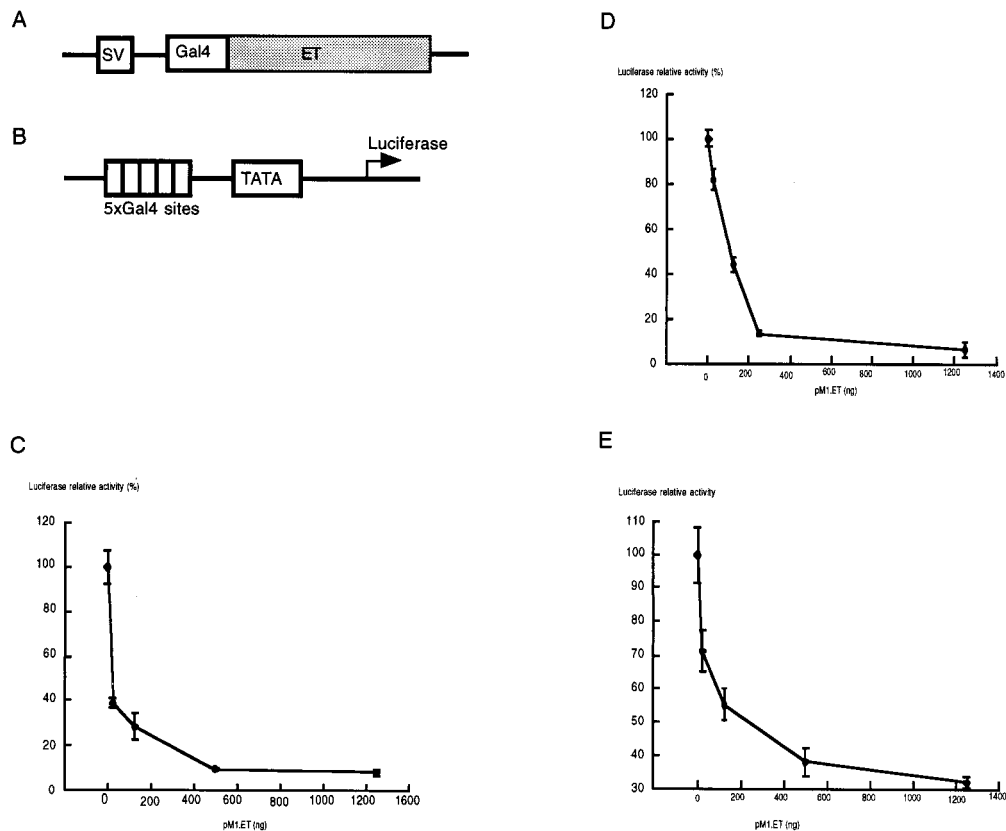


Fig. 1. Repression of basal transcription by ET. (A) A diagram of the plasmid expressing Gal4–ET fusion protein under the SV40 promoter. (B) A diagram of the reporter plasmid with five copies of the Gal4-binding sites upstream of a promoter driving the expression of luciferase. (C) ET can repress the luciferase expression driven by herpes simplex virus thymidine kinase promoter. Reporter plasmid (0.1 μ g) was cotransfected with varying amounts (0, 25, 125, 250, 500, or 1,250 ng) of Gal4–ET expression plasmid into 293T cells. The luciferase activity without Gal4–ET was defined as 100%. The relative luciferase activities with increasing amounts of Gal4–ET were 100 \pm 7.2 (mean \pm SD), 39 \pm 2.0, 28.5 \pm 5.9, 9.8 \pm 0.4, 8.5 \pm 1.1, respectively. (D) ET can repress adenovirus major late promoter (AdMLP); the relative luciferase activities were 100 \pm 8.5, 71.2 \pm 6.3, 55.5 \pm 4.7, 38.3 \pm 4.2, 32.1 \pm 1.9, respectively. (E) ET can repress SV40 promoter in the presence of Gal4-binding sites. The relative luciferase activities were 100 \pm 8.5, 71.2 \pm 6.3, 55.5 \pm 4.7, 38.3 \pm 4.2, 32.1 \pm 1.9, respectively.

POPULATION BIOLOGY. In the article “Origin of a new *Phytophthora* pathogen through interspecific hybridization” by C. M. Brasier, D. E. L. Cooke, and J. M. Duncan, which appeared in number 10, May 11, 1999, of *Proc. Natl. Acad. Sci. USA* 96, 5878–5883, the authors wish to note that all of the Genbank accession numbers were missing one digit. The correct numbers are as follows: AF139366, AF139367, AF139368, AF139369, and AF139370.