Supplemental Figure 2

FIGURE S2.—Northern analysis of Ty1 host cofactor mutants required for Ty1 mobility. Total RNA was extracted from 22 deletion mutants that showed altered Gag levels compared to the wild-type strain, and subjected to Northern analysis using $^{32}$P-labeled RT probe. The signals were quantified by phosphorimage analysis and normalized to the total amount of 25S and 18S rRNA. The numbers below each lane indicate the fold change in the total Ty1 mRNA relative to the wild-type strain, which is normalized to 1.