Figure S2.—Boxplots showing the distributions of e_{HC} values for genes within human-specific duplications (H-), chimpanzee-specific duplications (-C), and all other genes (HC or --), in liver, kidney, and heart. The horizontal dotted line denotes the median of e_{HC} values in the third group (HC or --). Bars represent the 95 percentiles of each distribution and less than 3% of the extreme data in each distribution are not shown. The complete distributions range from -4.42 to 6.04 in liver; from -3.83 to 4.48 in kidney; and from -4.42 to 5.36 in heart.