**Figure S8.**—Boxplots showing the distributions of e_HC values for genes within human-specific duplications that do not overlap CNVs in humans (H-), chimpanzee-specific duplications that do not overlap CNVs in chimpanzee (-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.