FIGURE S4.—The proportion of genes within species-specific segmental duplications (y-axis), is plotted at different LR cutoffs (x-axis) for classifying genes as differentially expressed between species using the heart expression data (red circles). The proportion of genes within species-specific segmental duplications among genes that are not classified as differentially expressed (based on the different cutoffs) is plotted in blue crosses. The overall proportion of genes within species-specific duplications is shown by the dashed horizontal line.