



Figure S2. HK genes selected from different pathways span a wider range of expression levels. The expression ranges of HK genes in each of the 7 HK-enriched pathways for two tissues are shown by mean value (small solid circles) and standard errors (error bars) of their expression levels in log₂ scale. The Kendall's tau, $\bar{\tau}_{gg'}$ for HK genes selected from different pathways was computed to be 0.75 ± 0.00 for 100 runs of random sampling of 7 HK genes, each from one of the 7 HK-enriched pathways.