

Figure S2 A) As in Supplemental Figure 1, for *C. elegans* (*fog-2*) data. Plots are separated by X (left panels) and autosomes (right panels). B) For all species, boxplots indicate the range of expression (calculated as the log10 FPKM) of each sex-bias category. Male expression is plotted for male-biased genes (left 3 boxes); female expression is plotted for female-biased genes (right 3 boxes). Overall, sex-specific genes are expressed at low levels in the corresponding sex. (MS – Male-Specific, HM – High Male, LM – Low Male, LF – Low Female, HF – High Female, FS – Female Specific).