Molecular evolution of genes that have male- or female-biased expression and non-sex-biased expression. Estimates for $d_N/d_S$, $\alpha$, and $\omega_a$ were calculated using $M. \text{famulus}$ as the ougroup. Error bars are 95% confidence intervals (CIs) obtained by bootstrapping by gene. Two-tailed bootstrap tests were performed to compare $d_N/d_S$, $\alpha$, and $\omega_a$ estimates with the autosomal average (indicated by the dashed line), and between autosomal and X-linked genes of each class. Stars indicate significance for the comparison to the autosomal average (* $P<0.05$, ** $P<0.01$). Signs indicate significance for the comparisons between autosomal and X-linked genes (one sign; $P<0.05$, two signs, $P<0.01$).