

Figure S1: Box plots of novel associations

Box-and-whisker diagrams include smallest gene expression values, lower quartiles, medians, upper quartiles, largest gene expression values, and outliers of one representative association for each of the 31 trait genes uniquely identified in this study

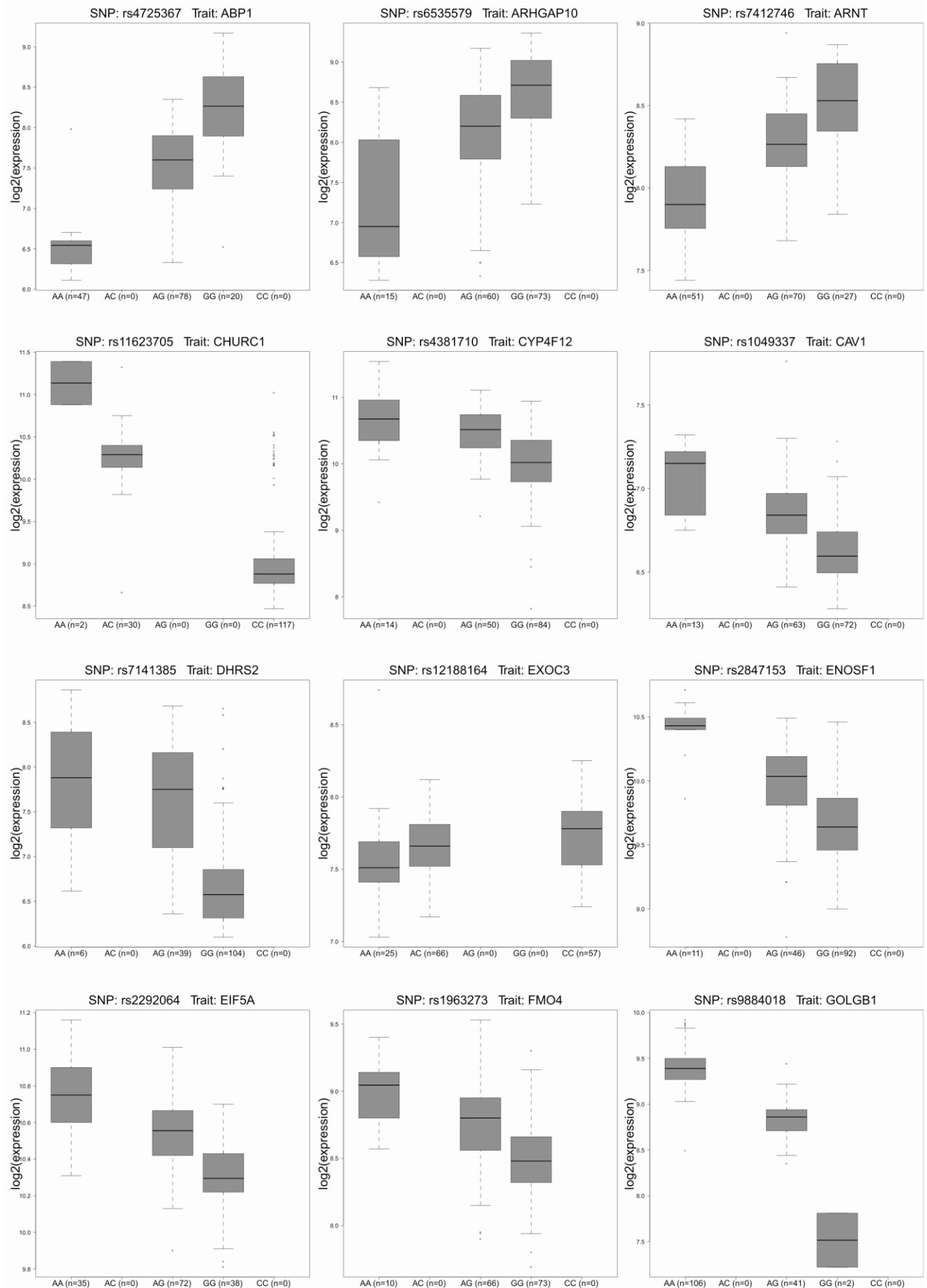


Figure S1 (continued)

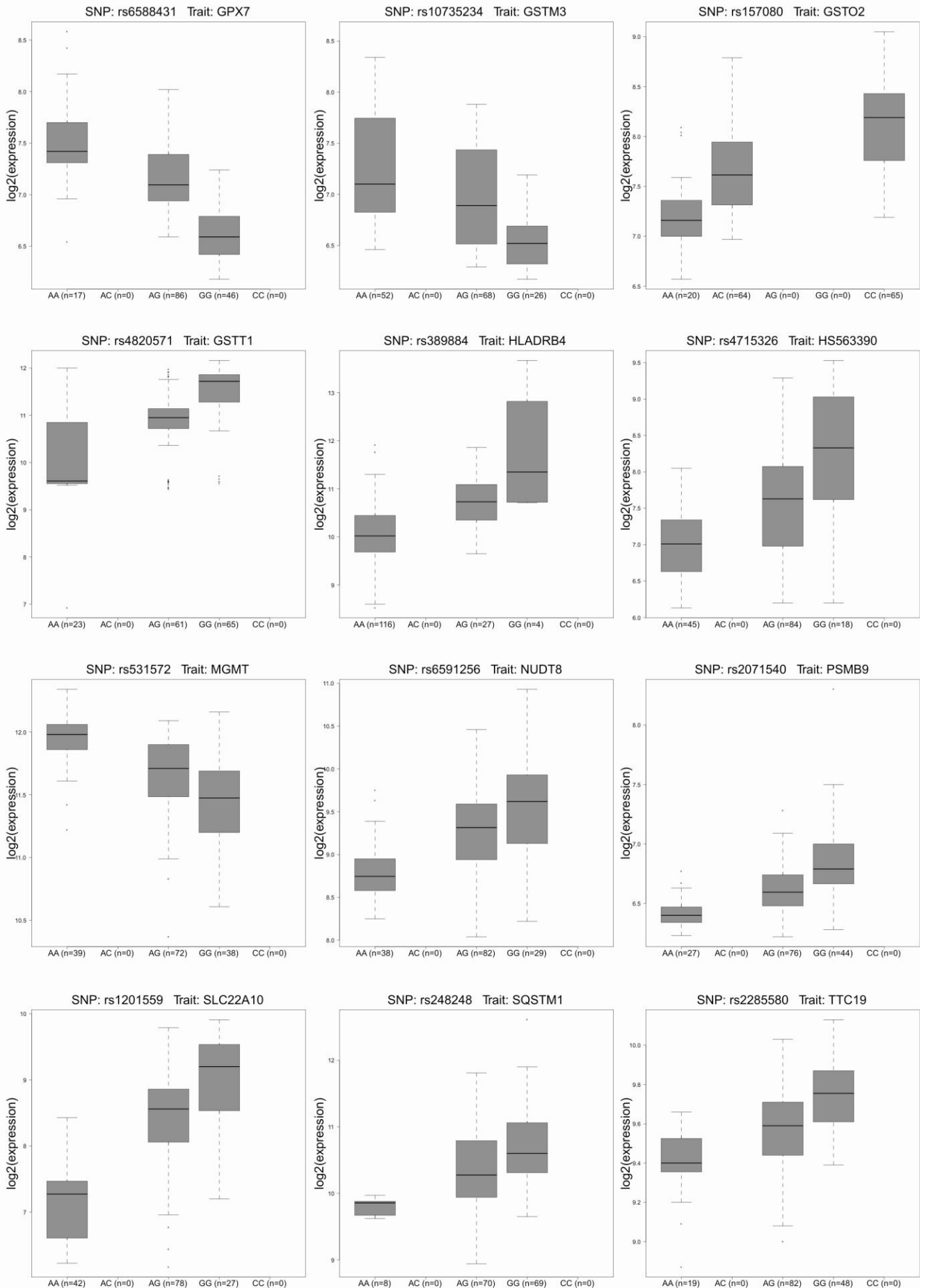


Figure S1 (continued)