



Figure S2 Regulatory impact of variation in *AFT1*, *CCC1*, and *YAP5* between Malaysian and wine/European yeast. Data are as in Figure 4 of the main text except that distributions of expression measurements across experimental replicates are shown. Each panel reports expression of iron-starvation (GASCH *et al.* 2004) and iron-resistance (LIN *et al.* 2011; PIMENTEL *et al.* 2012) genes as ratios of the levels measured in two strains derived from a Malaysian (UWOPS03.461.4) and a wine/European strain (BC187). At left in each row, each bar reports the median across replicates ($n = 2$) of the ratio of expression of the indicated gene between a Malaysian homozygote and a wine/European homozygote. In each remaining panel, each bar reports the median across replicates (*CCC1*, $n = 8$; *AFT1*, $n = 4$; *YAP5*, $n = 4$) of the ratio of expression of the indicated gene between a Malaysian-wine/European hemizygote bearing the Malaysian allele of the indicated variant locus, and the hemizygote bearing the wine/European allele. Error bars report 95% confidence intervals. (A) Iron-starvation genes in synthetic complete medium. (B) Iron-resistance genes in synthetic complete medium supplemented with 5 mM FeSO₄. Raw data are reported in Supplementary Data Sets 7 and 8.