Supplemental Figure S4A.

Variability in gene expression was biased towards TATA box genes. Transcriptome profiles were obtained for *Saccharomyces cerevisiae* strains 06L3FF02, 06L6FF20, AEB Fermol Rouge, Lalvin ICV D254, Lalvin EC-1118, J940047 and S288C during fermentation in synthetic wine must, at stages from T1 to T6 (see Manuscript for details). The average deviation from the mean relative transcript abundance of every examined ORF was taken as a measure of the respective variability in expression. The distribution of TATA box genes (per interval of 100 genes) was determined in a list of ORFs, ranked from the lowest to the highest average deviation value. TATA box genes were identified according to Basehoar *et al.*, 2004. *Cell* 116: 699-709. For reference, the dashed line indicates the approximate percentage of TATA box genes in the genome of strain S288C.
Variability in gene expression was biased towards TATA box genes. Analysis was performed only with wine strains 06L3FF02, 06L6FF20, AEB Fermol Rouge, Lalvin ICV D254, Lalvin EC-1118. Remaining legend as for Supplemental Figure S4A.