



FIGURE S2.—EM93 is the source of the BY/S288C *sal1-1*, *CAT5-911*, and *MIP1-661A* alleles. The diploid EM93 strain was sporulated. Each spore of a tetrad was sequenced for a region encompassing the relevant polymorphisms in the *SAL1*, *CAT5*, and *MIP1* genes. Each trace picture shows the relevant DNA (top) and protein (underneath) sequence. The black box in the *SAL1* column highlights the heterozygosity of the *sal1-1* frameshift mutation (present in spores c & d). The amino acid differences in *CAT5* and *MIP1* are also boxed in black.