

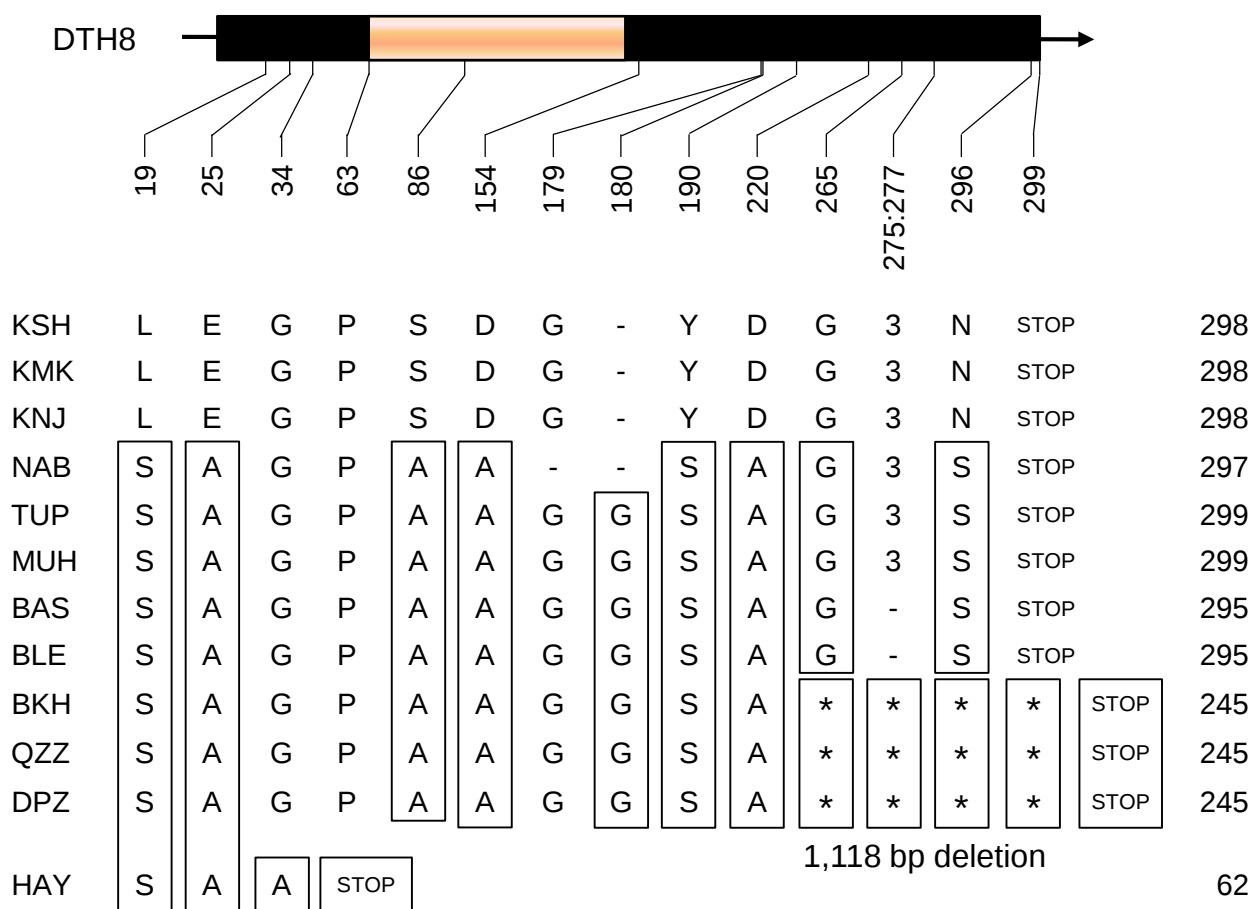


Figure S4. Substitution (boxes) and insertion/deletion (-) polymorphisms of amino acids in the DTH8 protein in 12 diverse accessions of Asian rice. Abbreviations of rice accessions are defined in Table 1. The conserved histon-fold motif domain in DTH8 is indicated in orange. Numbers under the DTH8 diagram indicate the positions of polymorphic sites. Numbers on the right side show the total length of each predicted amino acid sequence. The regions with amino acid changes due to a frame shift are labeled with asterisks. Stop indicates a stop codon. Accession numbers of each sequence are DDBJ: LC016712-LC016721.