

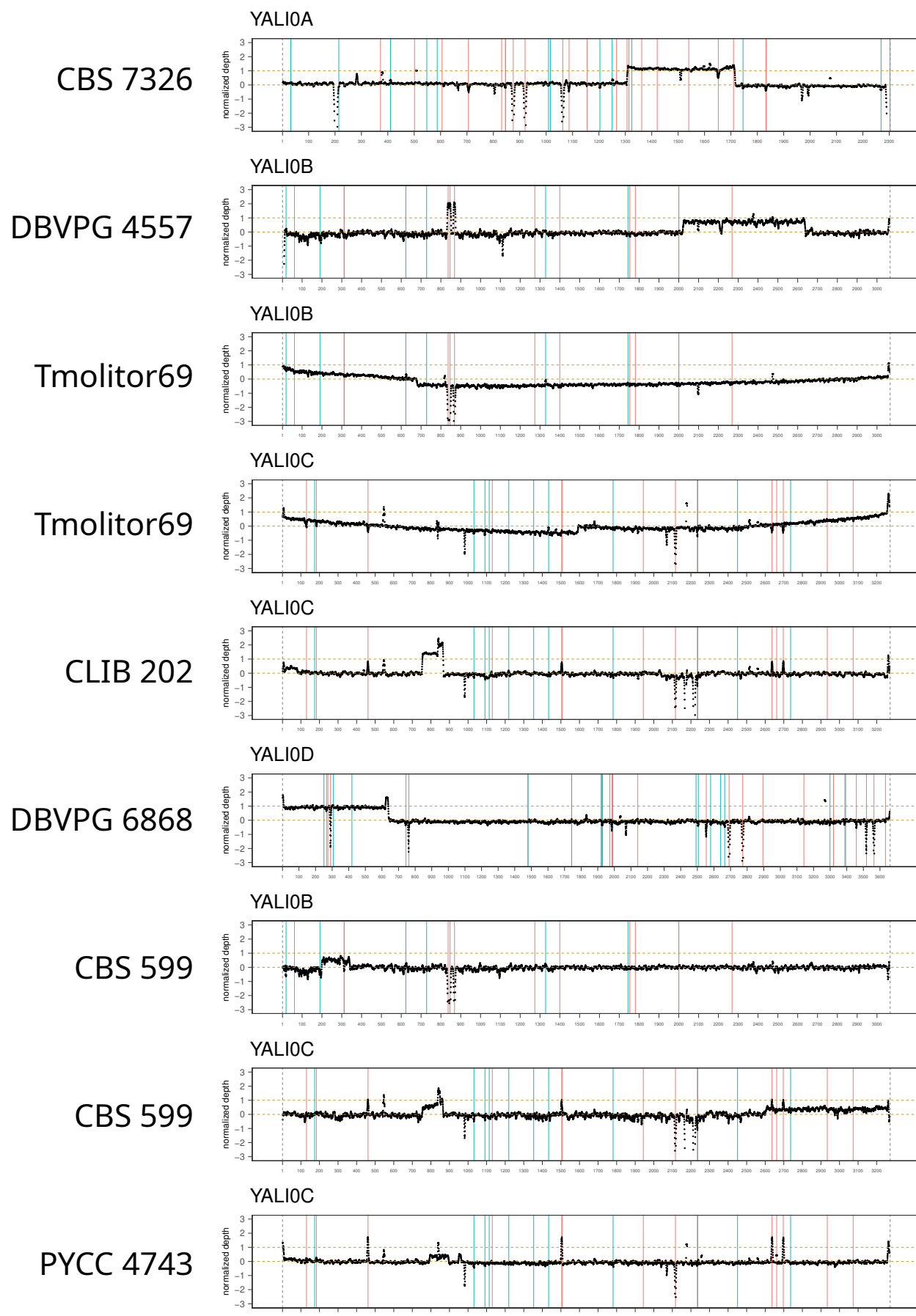


Figure S4: Segmental duplications found in the genome of 7 yeasts. Normalized mapping depth (transformed in logarithm of the base 2) was averaged over 5,000-bp windows (steps = 500 bp). Position of the transposable elements in the genome of *Y. lipolytica* E150 is marked in red and solo LTRs in blue