

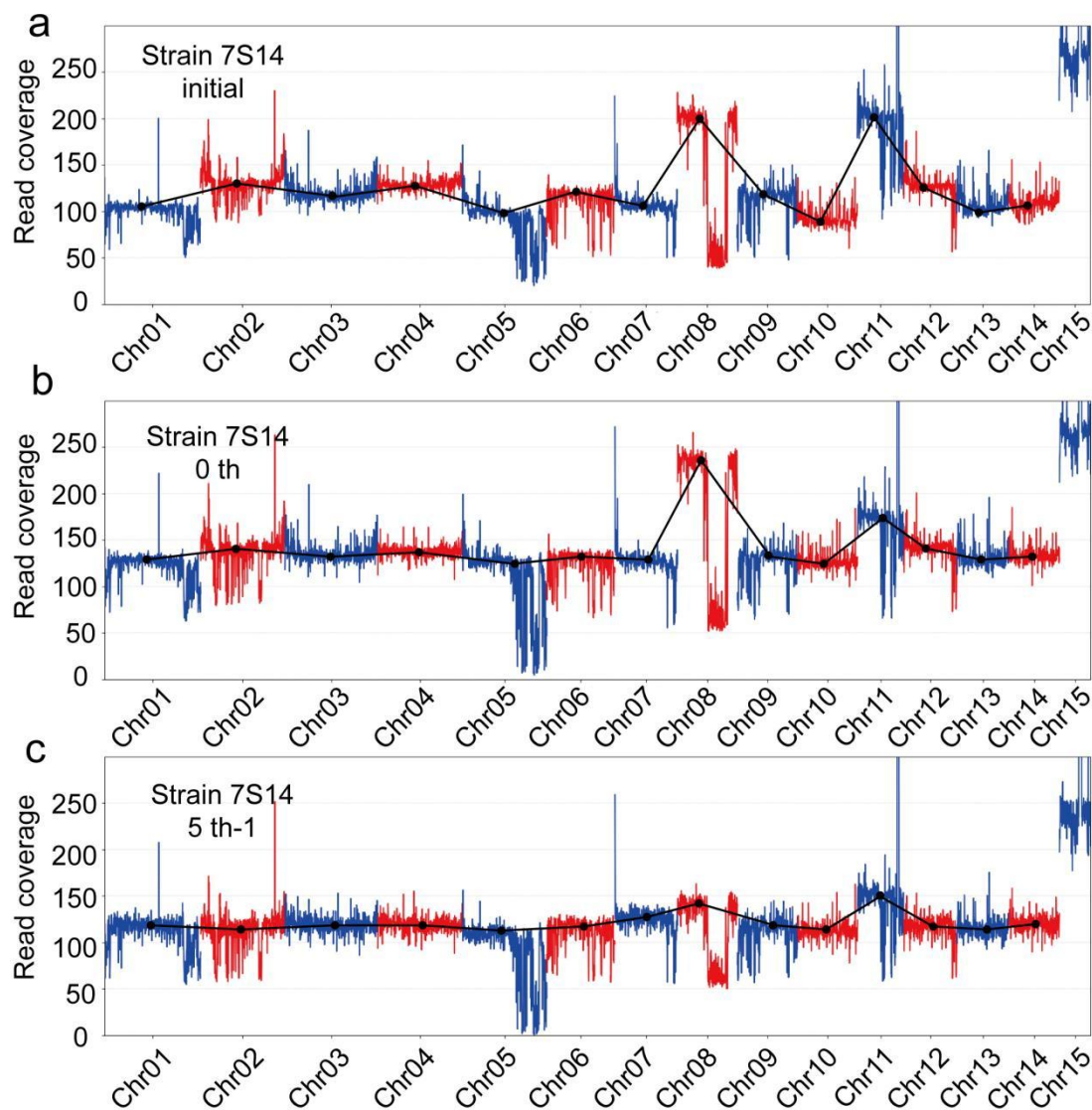


Figure S28. Chromosome-wide coverage of 7S14 strains from different generations, mapped to the 7S14_set1 genome. The resequencing data were the same as those used in Supplementary Fig. 4 (a) and Fig. 6b (b, c). The strain shown in panel a is sequenced after single-spore isolation followed by only one vegetative passage, whereas the strain shown in panel c is derived from the strain in panel b after five additional passages.