

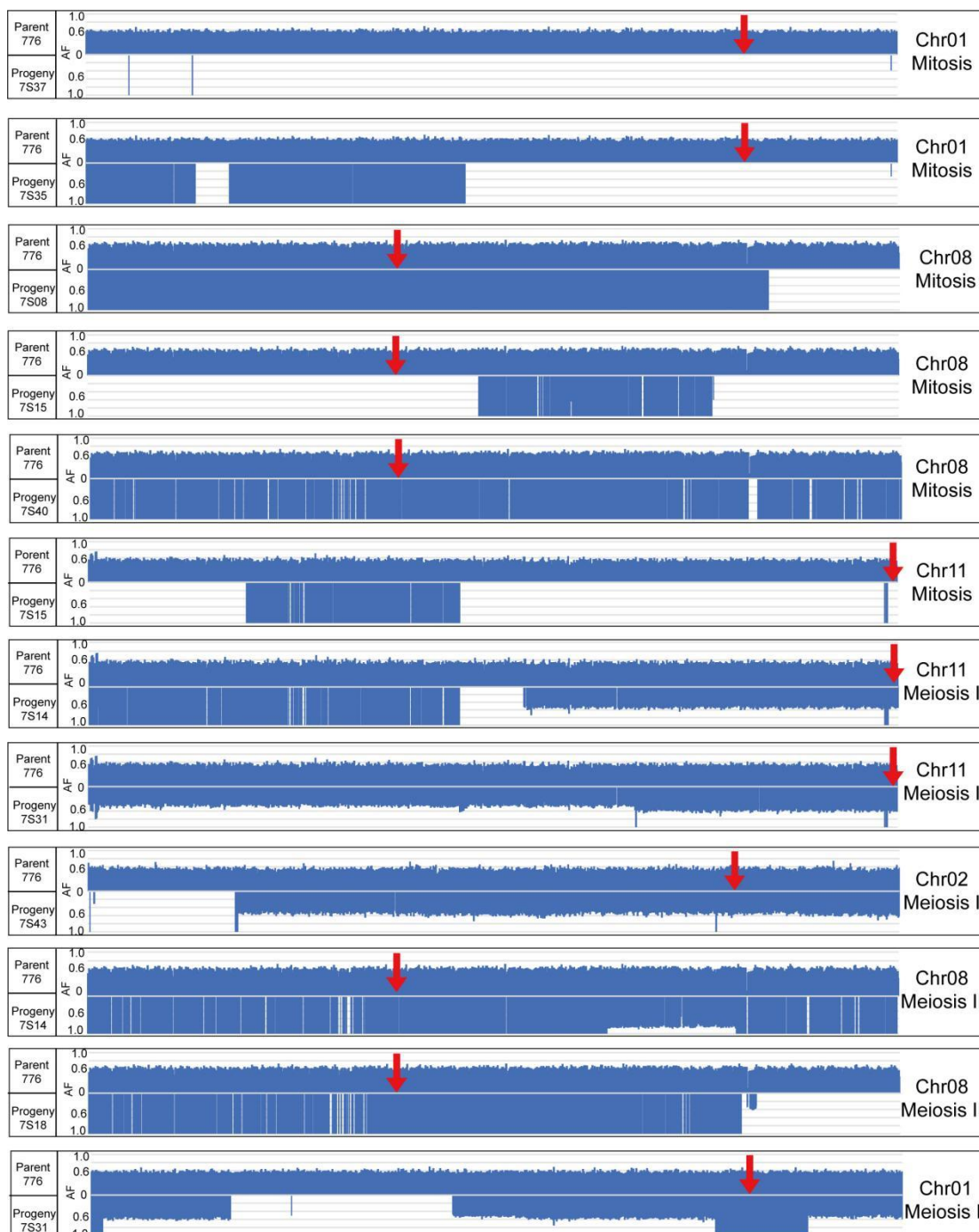


Figure S15. SNP allele frequencies of 12 CC aneuploidy events in sexual progeny derived from strain 776. Red arrows indicate putative centromere regions. Inferred segregation stages are shown on the right.