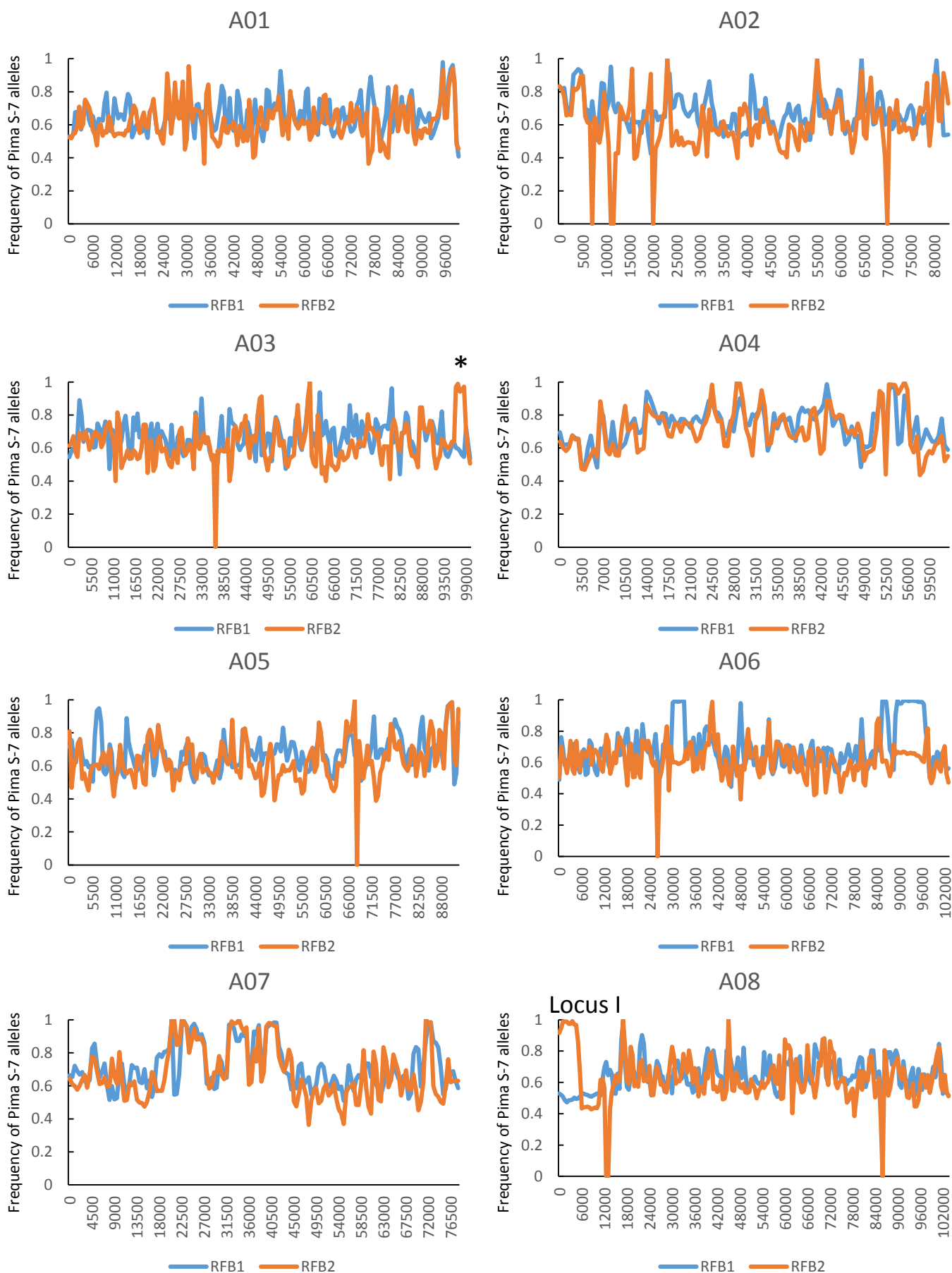
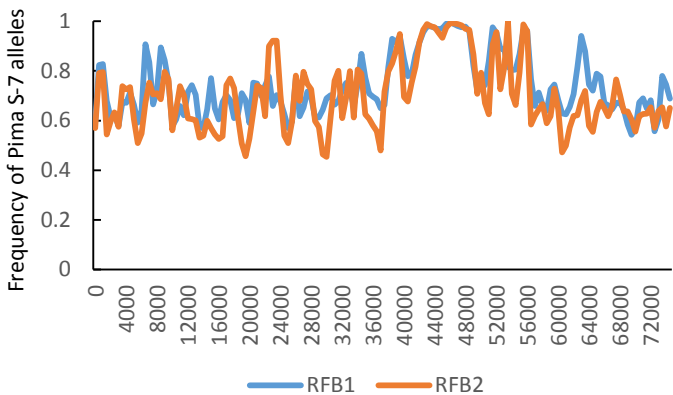


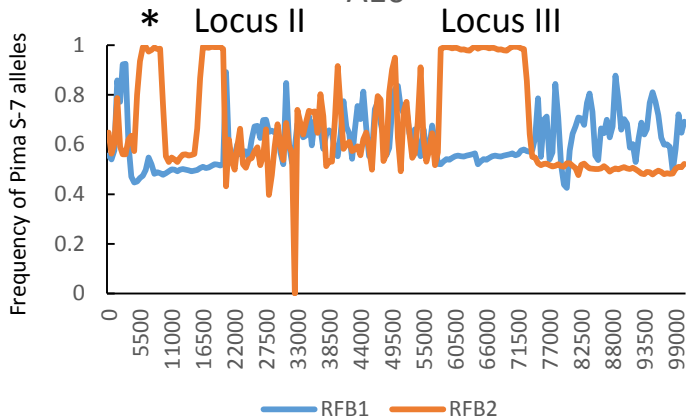
Figure S4 Distribution of the Pima S-7 allele frequency across the 26 cotton chromosomes in the NILs showing fuzzless (RFB2) and segregating fuzz phenotype (RFB1). * indicates the potential candidate regions used in association analysis. Locus I, II, III, IV and V represent the five loci associated with fuzz development.



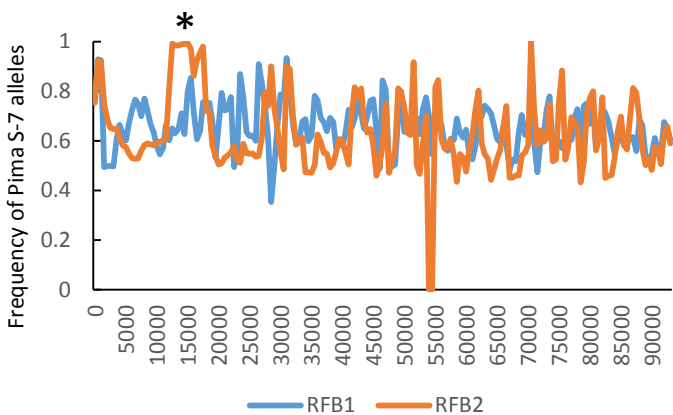
A09



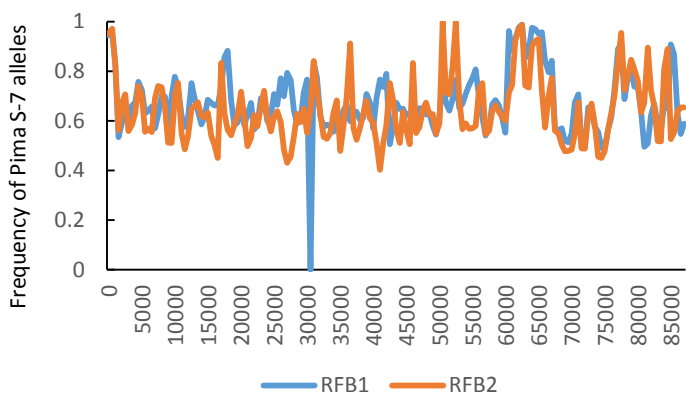
A10



A11



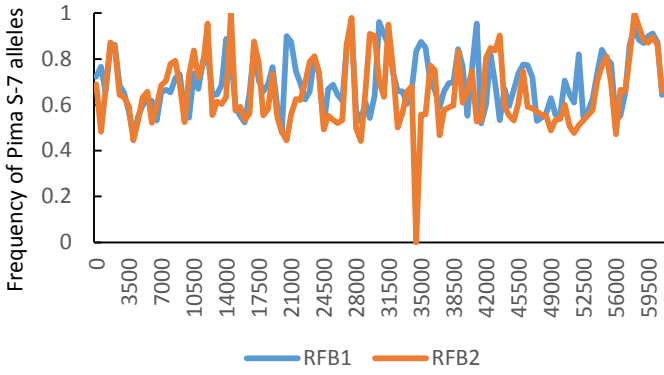
A12



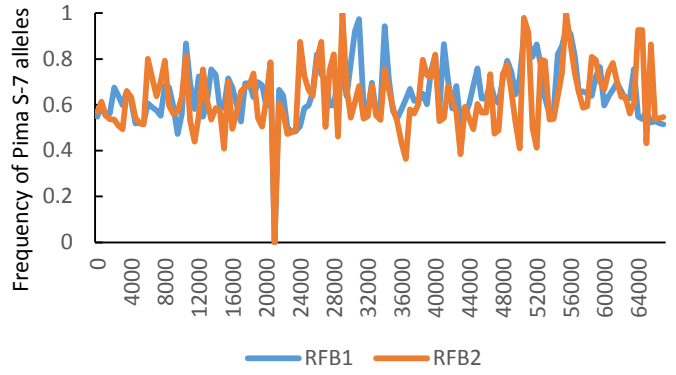
A13



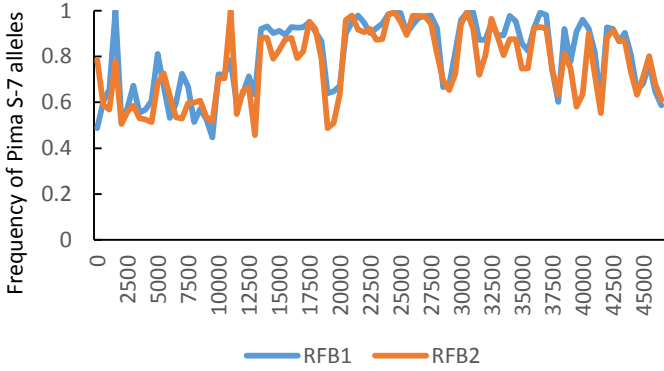
D01



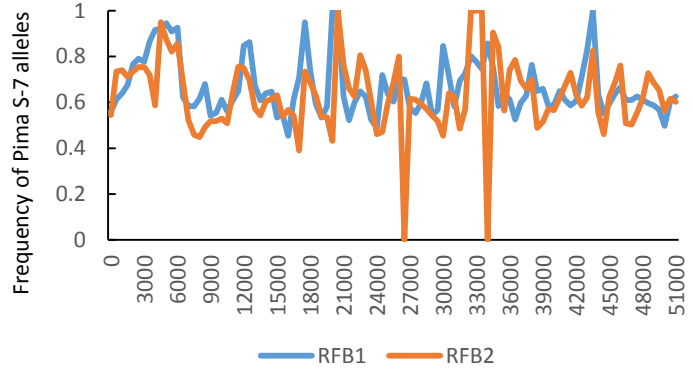
D02



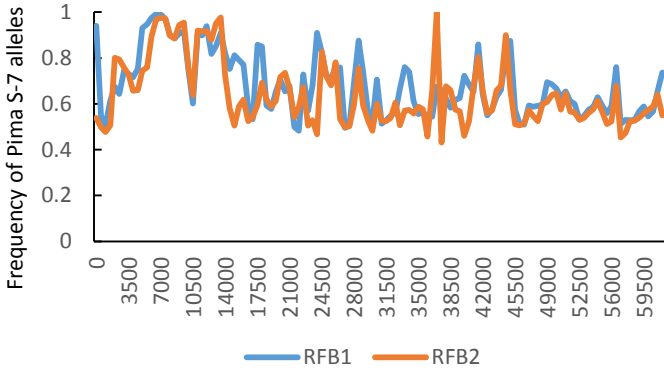
D03



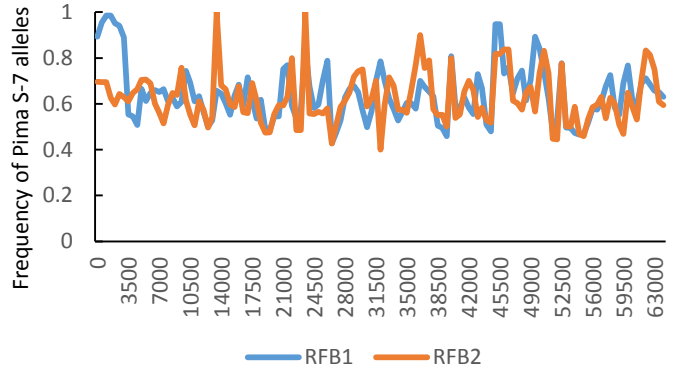
D04



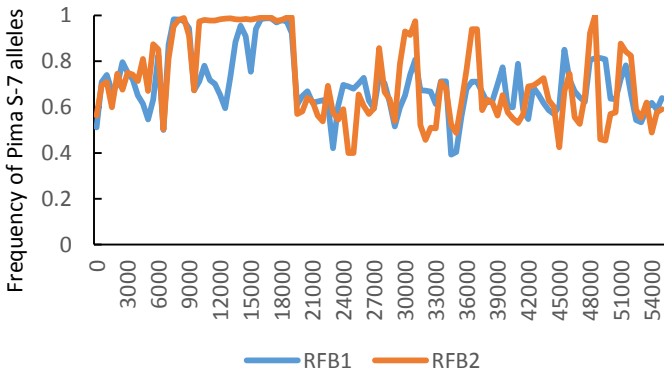
D05



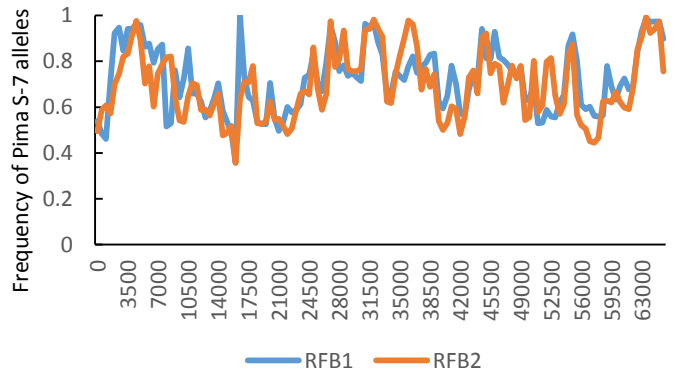
D06



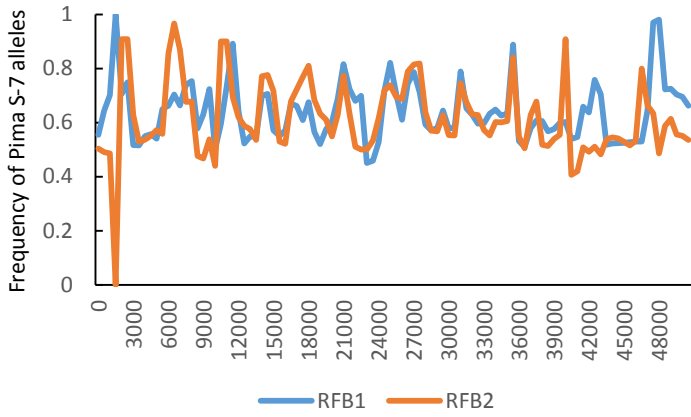
Locus IV D07



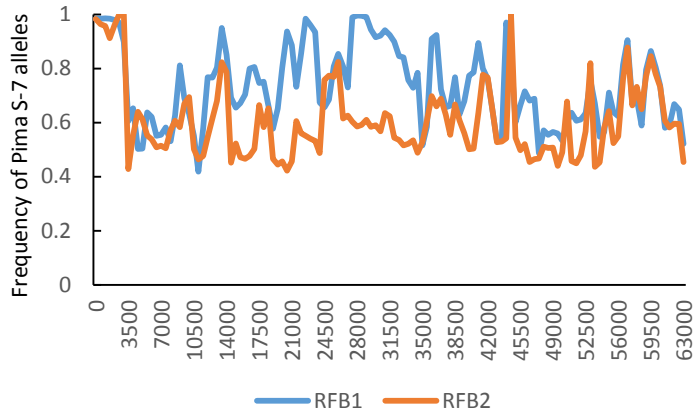
D08



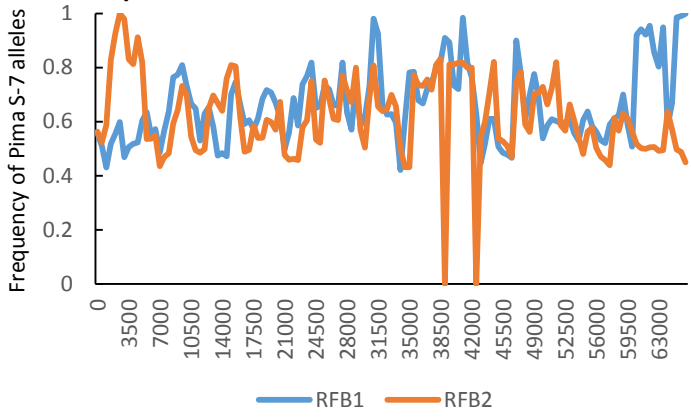
D09



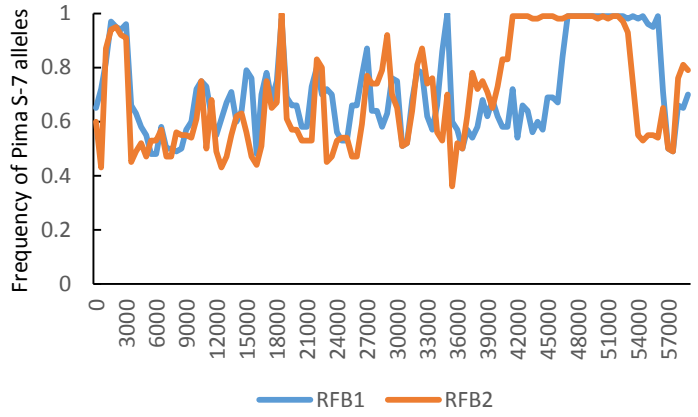
D10



D11



D12



D13

