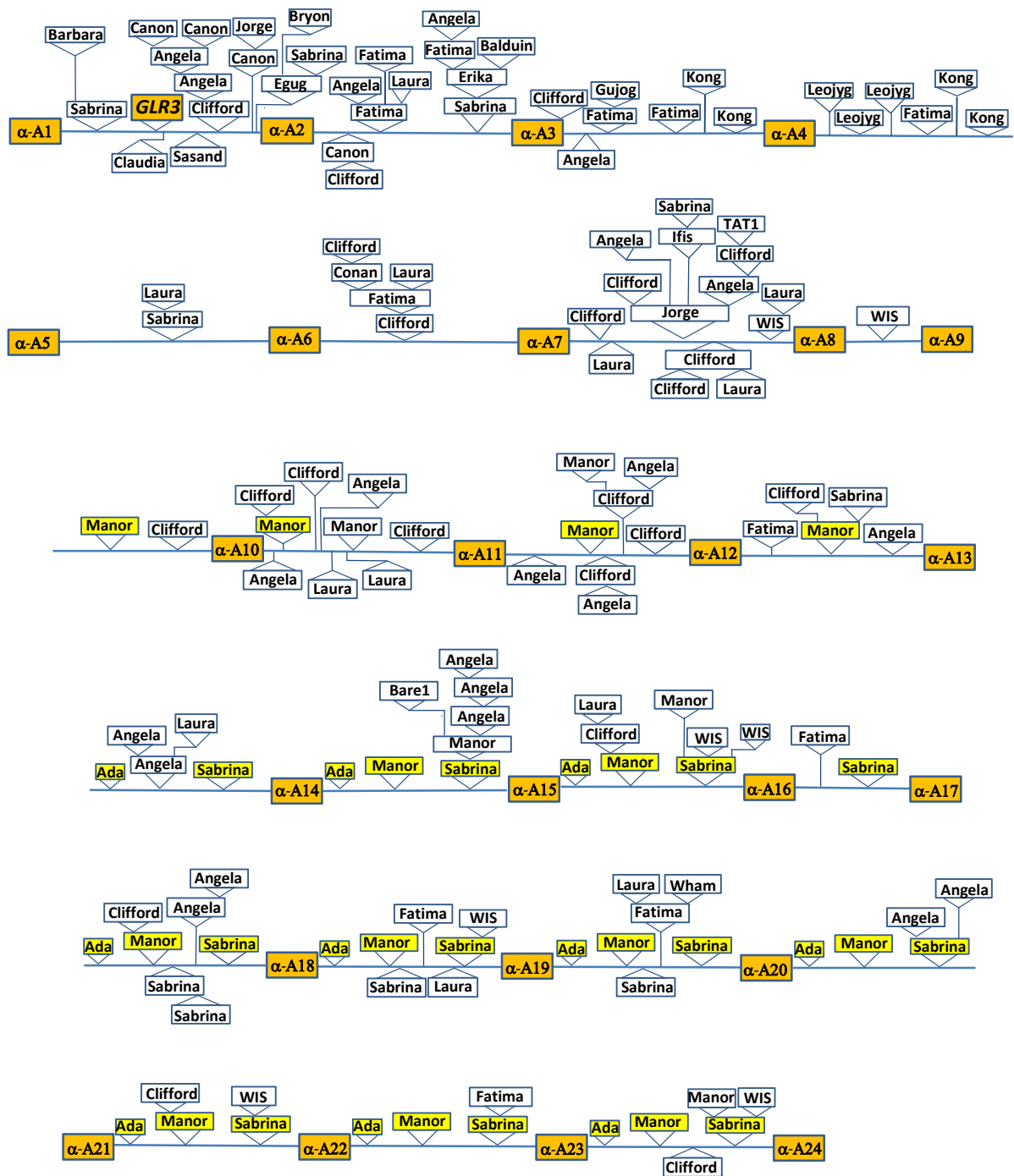


A



B

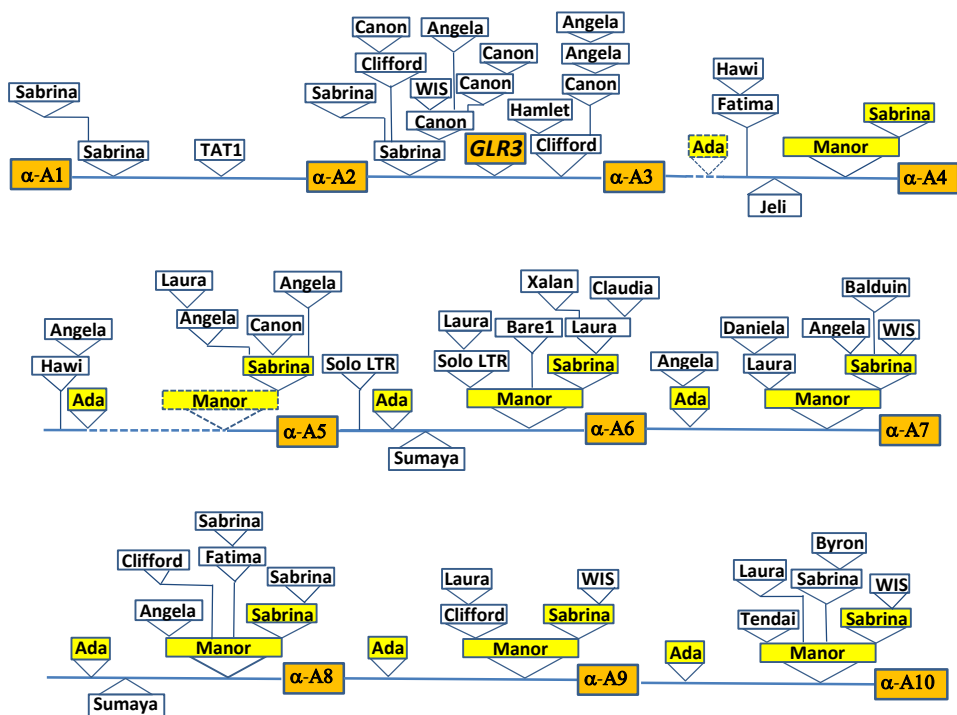


Figure S2. Organization of the α -gliadin locus region in the wild emmer and Chinese Spring A genomes.

The genomic organization of regions containing α -gliadin genes from wild emmer A genome (A) and Chinese Spring A genome (B) (Huo et al., 2018b) are displayed with α -gliadin genes represented by color-filled boxes. Retroelements are boxed with their names, and their insertion positions are indicated by triangles or solid vertical lines. Retroelement insertion sites that are likely shared in the intergenic regions or between two genomes are highlighted with yellow. Dashed lines indicate deleted sequences.