

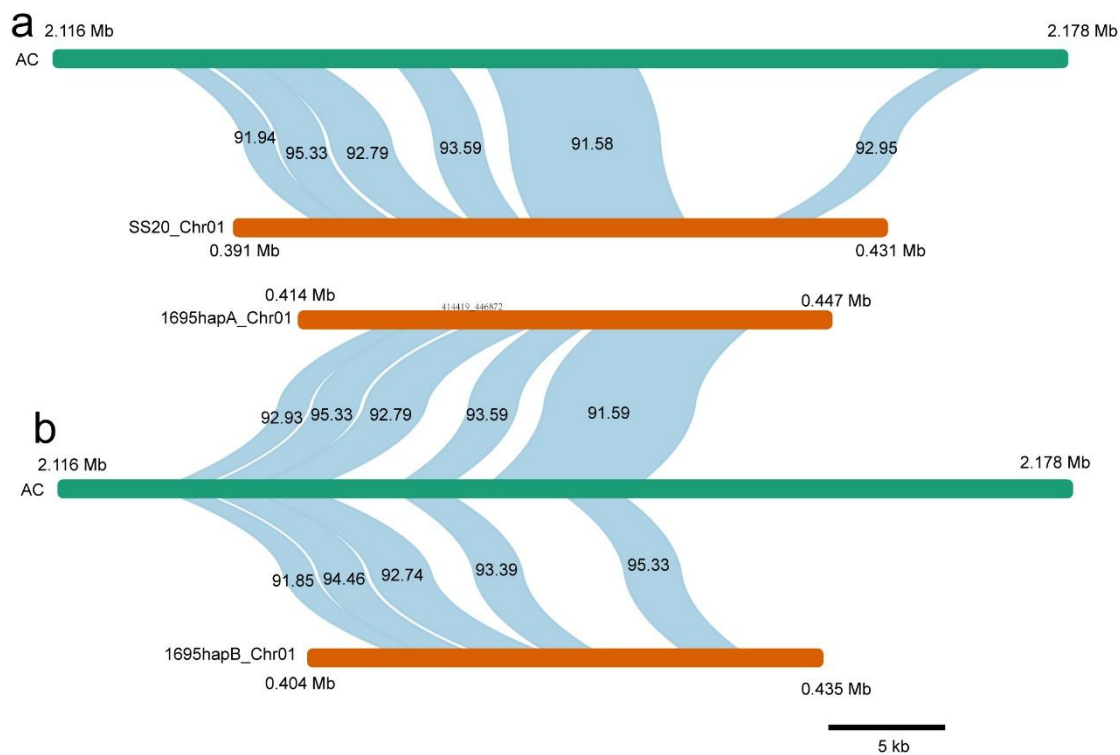


Figure S27. Collinearity of a putative ancestral chromosome breakage-derived segment in the AC with the Chr01 in SS20_T2T, 1695_hapA and 1695_hapB. Shaded ribbons connect syntenic regions, with sequence identity indicated.