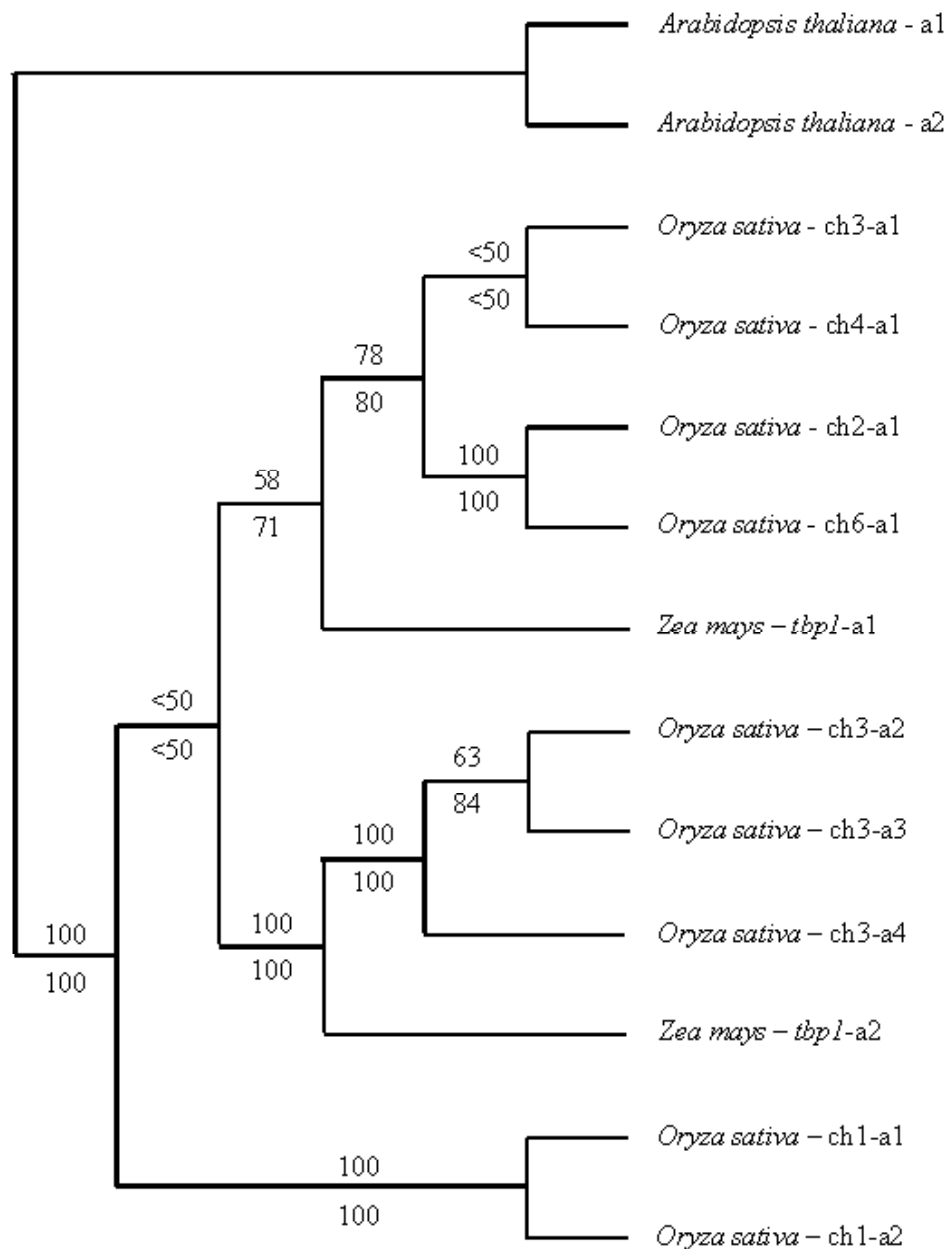


Additional file 2 for “Retrotranspositions in orthologous regions of closely related grass species” by Du et al. BMC Evolutionary Biology #4916306119647011.



Cladogram of the auxin-related genes resulting from maximum parsimony analysis using branch-and-bound search option. Numbers at the internodes refer to bootstrap proportions (upper number based on data set with missing characters excluded, 220 total; lower number based on data including all characters, 1044 total; heuristic search performed with 1000 pseudoreplicates).