Granule-bound starch synthase gene tree. The best-scoring ML tree was selected from 30 GARLI analyses of the GBSSI sequence data set under a GTR+I+Γ model of sequence evolution. Numbers above branches show ML bootstrap support 50%; the 28% value is mentioned in the text. “NA” and “Eu” distinguish North American and Eurasian Elymus species, respectively. Where applicable, numbers following taxon names distinguish individuals within species. Letters following these numbers designate cloned sequences from within individuals.