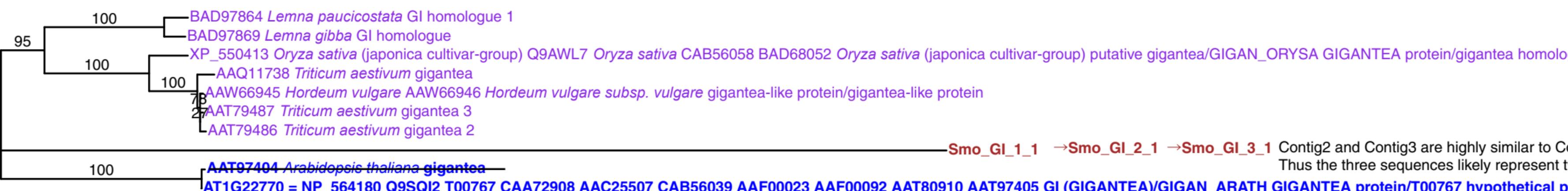


GI (AT1G22770): global tree
File name: GI.njtree0.svg
672 aa residues 10 OTU included.
Results of 2 partial trees included
Assemble Job Id: T1HaFO
Tree job id: uAaOaO
BP values out of 100
Analyzed by aono@nibb.ac.jp
Date: Tue Jun 13 16:24:34 JST 2006



Smo_GI_1_1 → Smo_GI_2_1 → Smo_GI_3_1 Contig2 and Contig3 are highly similar to Contig1 and have no overlap.
Thus the three sequences likely represent two alleles.