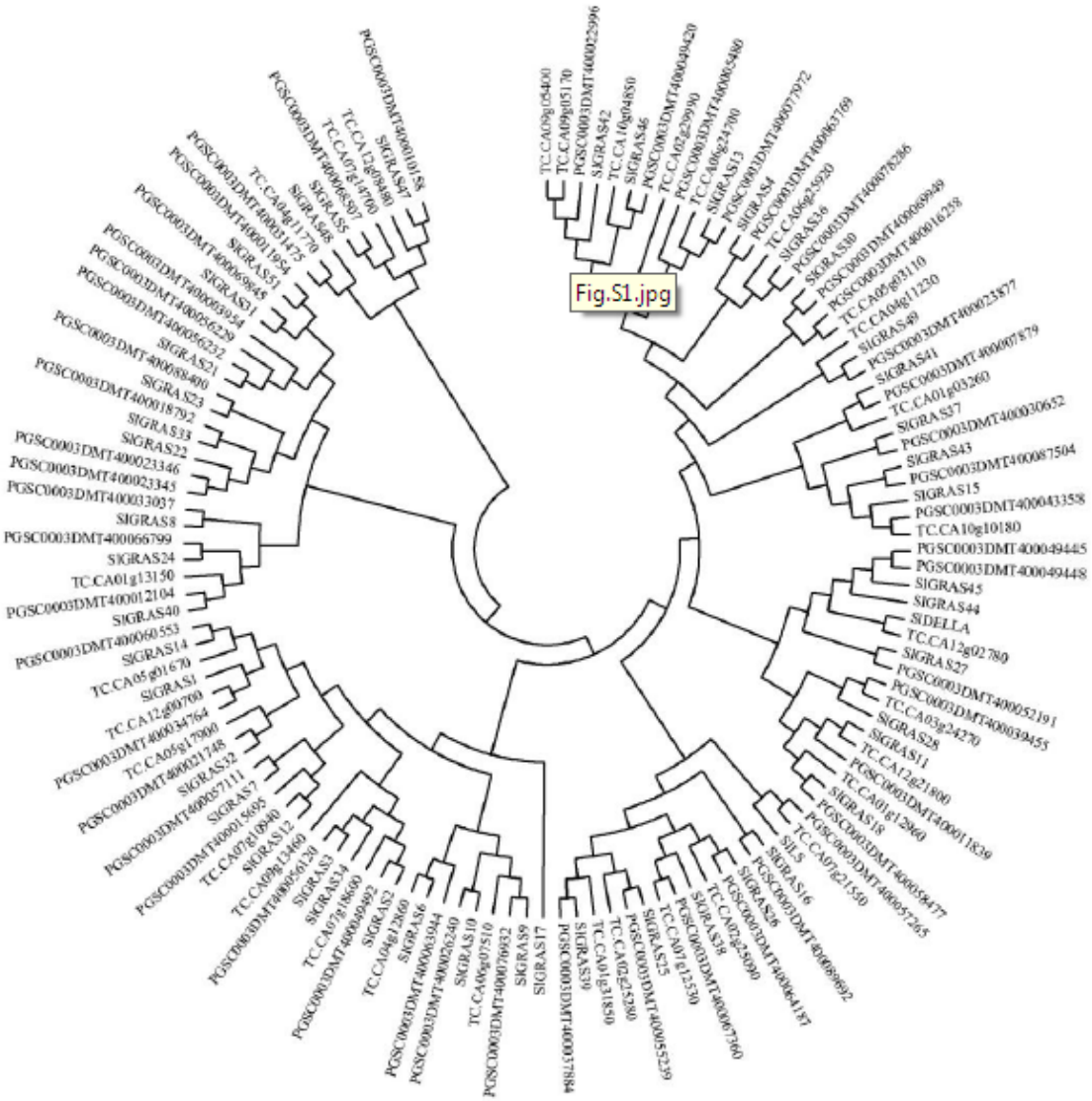Additional file 4: Phylogenetic analysis of GRAS proteins of tomato, potato, and pepper. The phylogenetic tree was generated by Neighbor-Joining method derived from ClustalX alignment of 48, 50, and 30 GRAS amino acid sequences from tomato, potato, and pepper, respectively.