Figure S2A PhyML phylogenetic relations and composition of domains for TTG1 (AT5G24520) homologues of representative plant species.

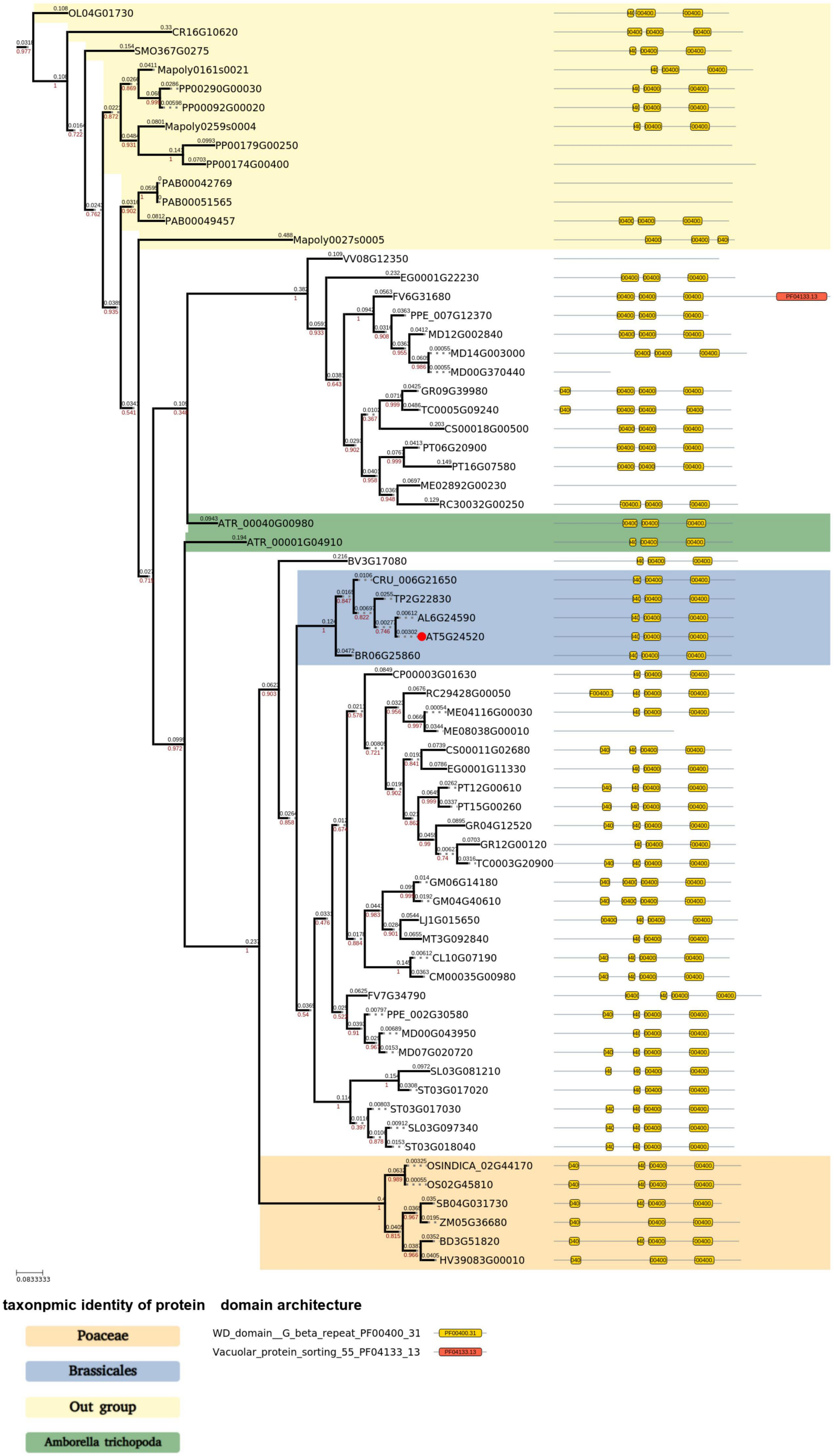

Figure S2B Evolutionary changes in protein structure of the TTG1 (AT5G24520) homologous proteins being studied branch reflects from right to left: disorder (2 residue types), secondary structure (3 types), secondary structure (8 types), globule surface (3 residue types), rare (comparing with protein specific model) amino acid substitutions
Color scheme:
black- outer branch (not analyzed);
colours define branch lengths quartile: blue – Q1; green – Q2; orange –Q3; red –Q4.