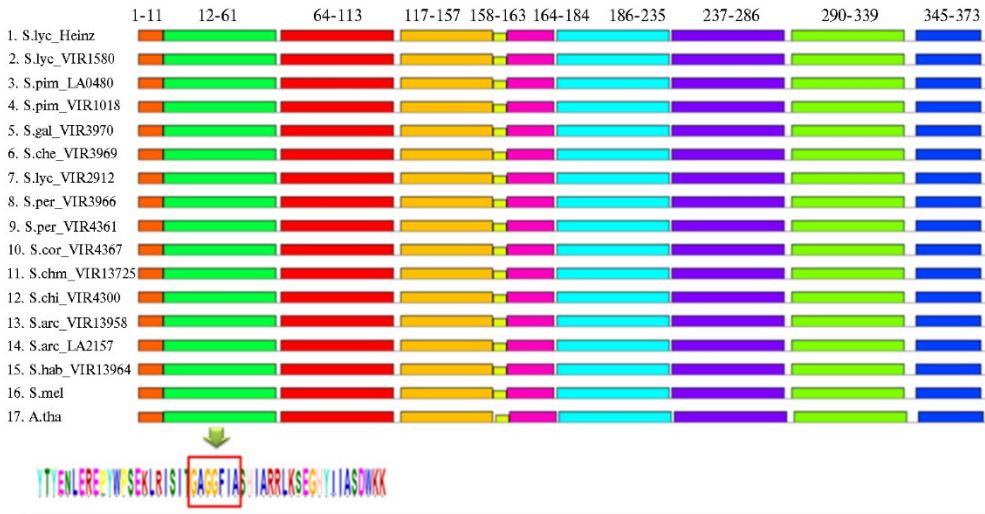


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Ten conserved motifs identified by the amino acid sequence analysis (the MEME 5.3.2 program) for *GME1* in the studied tomato (*Solanum L.*) species and in *GME1* homologous genes from other species.