



S3 Fig. Molecular phylogenetic analysis of domains of the SAGA complex components. Amino acid sequences of domains of the SAGA complex subunits were used from *At*, *A. thaliana*; *Dm*, *D. melongaster*; *Hs*, *H. sapiens*; *Os*, *O. sativa*; *Mm*, *M. musculus*; *Sc*, *S. cerevisiae*. **(A)** Protein domain of ADAs group; **(B)** Protein domain of SPTs group; **(C)** Protein domain of DUBm group; **(D)** Protein domain of CHD subunit; **(E)** Protein domain of SGF29 subunit; **(F)** Protein domain of TRA1 subunit; **(G)** Protein domain of TAFs group.