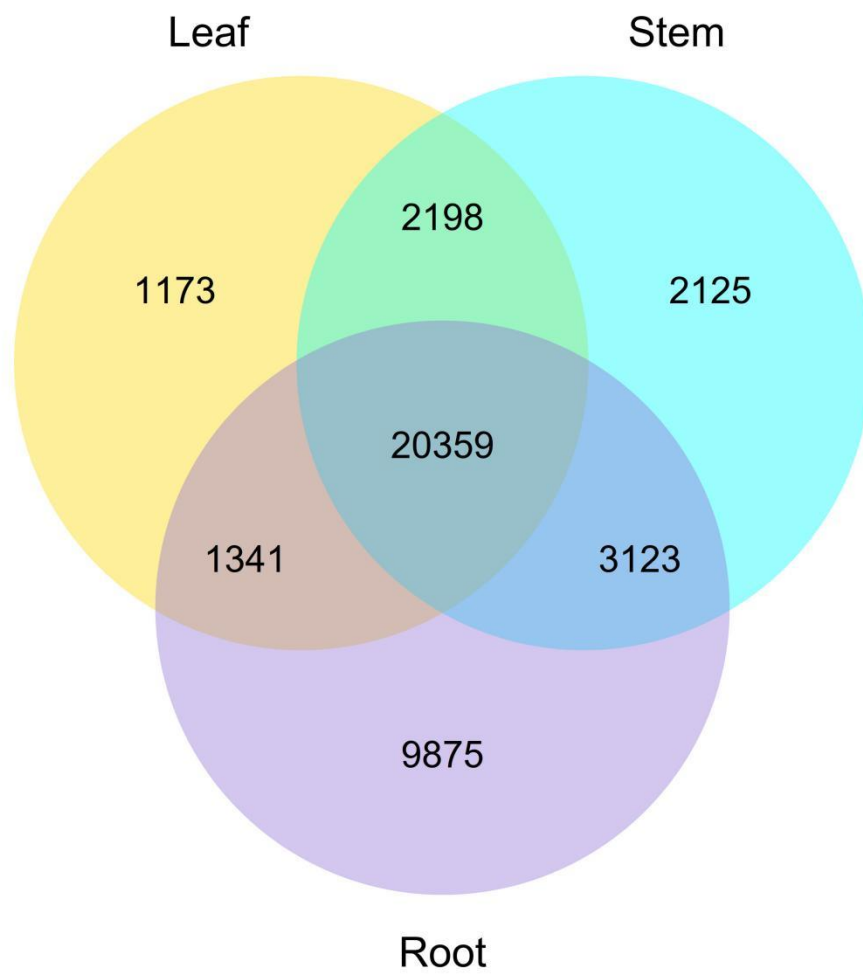
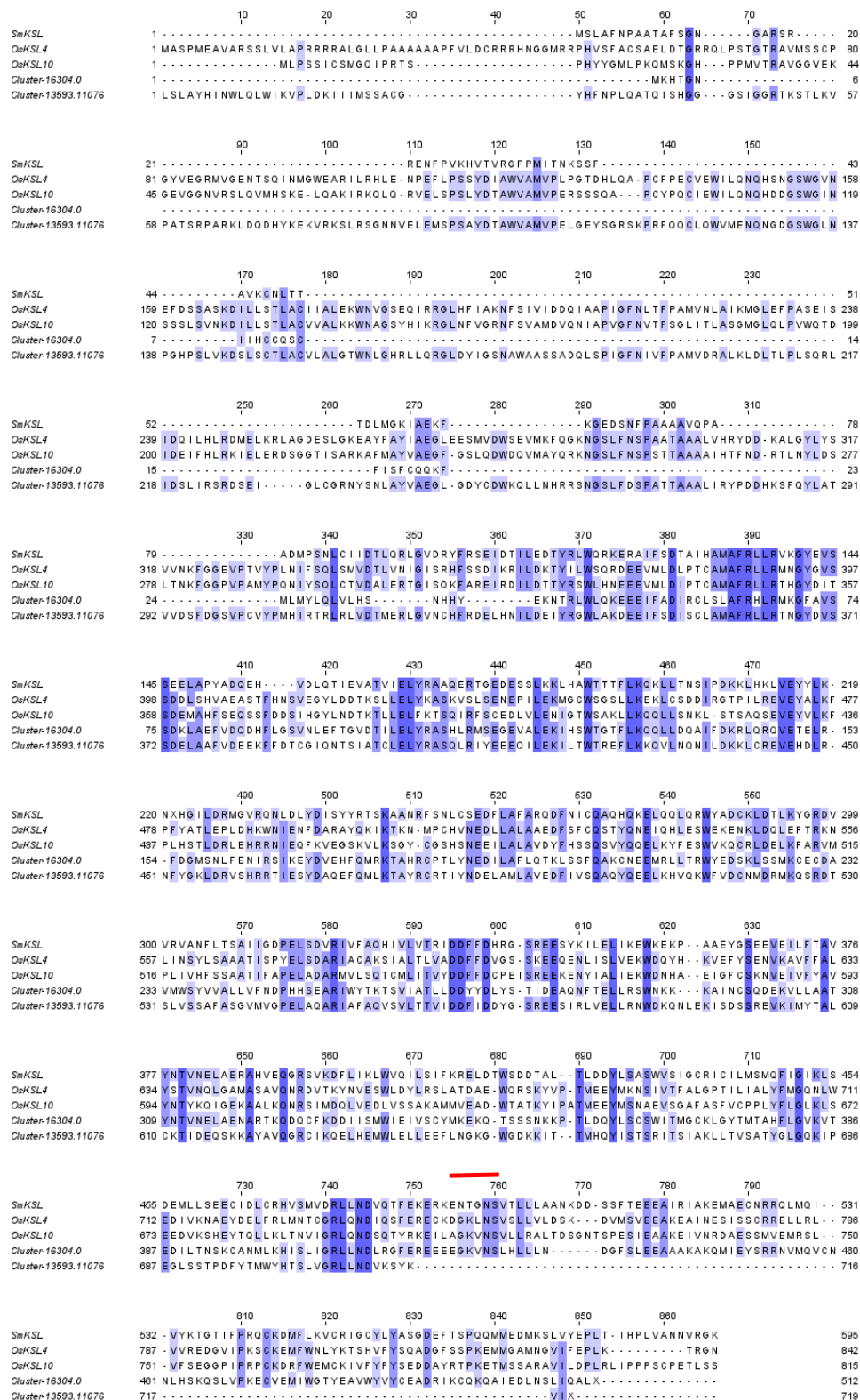


**Supplementary Figure S1.** Database annotation classification. **(A)** Species distribution of transcriptomic unigenes against NR database. **(B)** KOG classification.



**Supplementary Figure S2.** Venn diagram of coexpressive unigenes in leaves, stems and roots of *A. paniculata*.



**Supplementary Figure S3.** Two sequence comparison of the *A. paniculata* kaurene synthase-like (KSL-like) enzymatic family with three previously characterized KSL enzymes from *Oryza sativa* and *salvia miltiorrhiza*. Underline indicates the characteristic DDxxD motif.