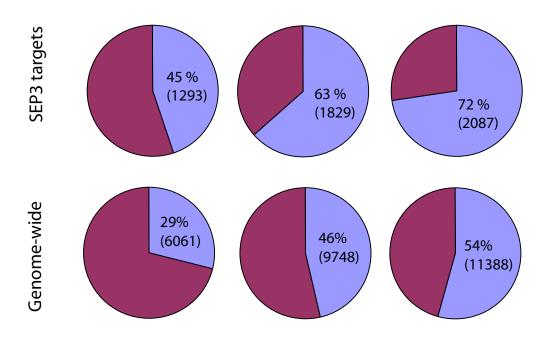
A. Global gene expression analysis

Floral homeotic mutants Developmental stages Combined data



B. Genes correlated with correlated expression to SEP3

