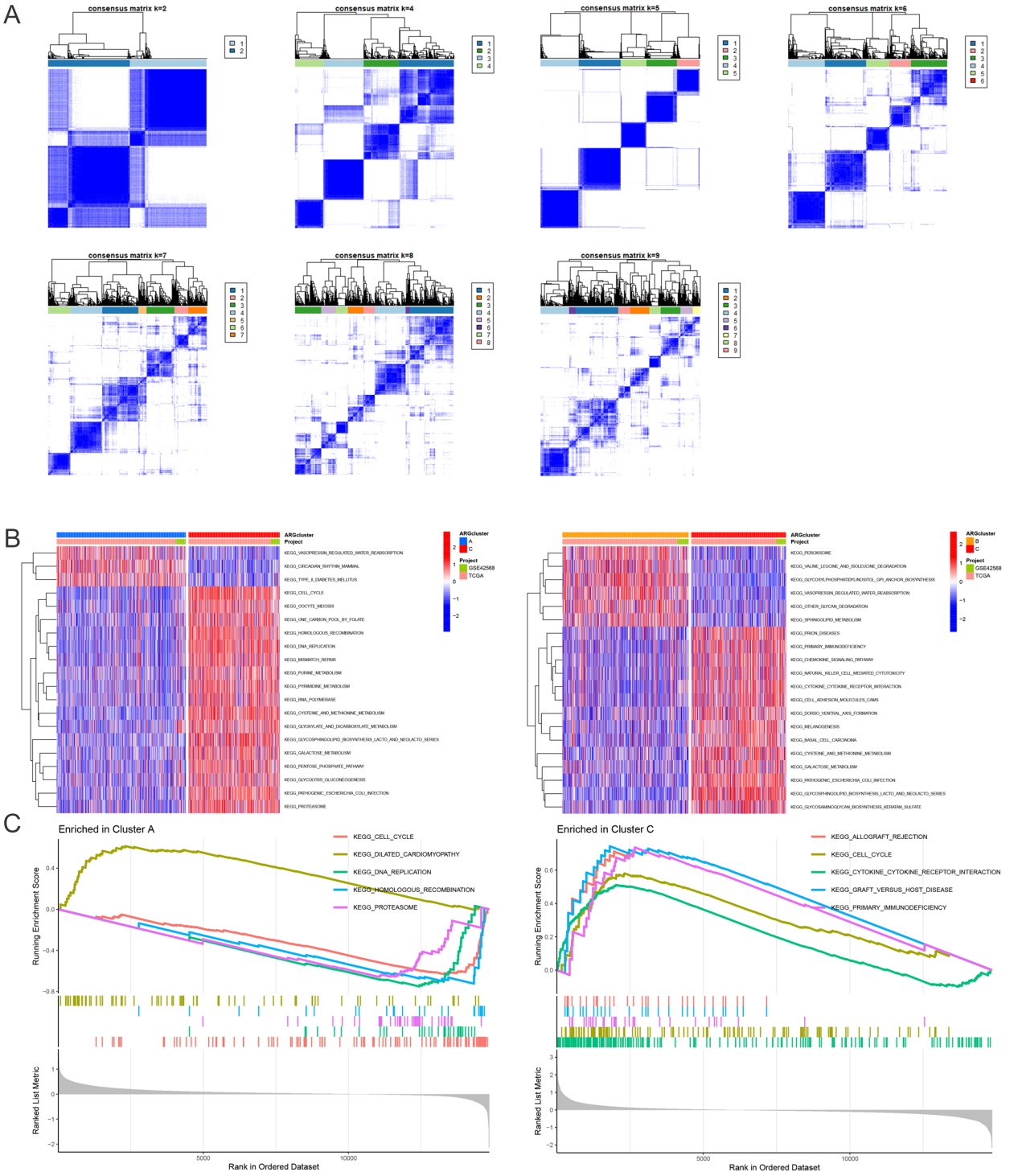

Figure S1 A, Identification of clusters of breast cancer based on prognosis-related DEGs using consensus clustering for k = 2,4,5, 6,7,8,9. B. Differential enrichment of KEGG pathways between clusters A and B, B and C analyzed by GSVA. C, Enriched KEGG pathways in cluster A and C.