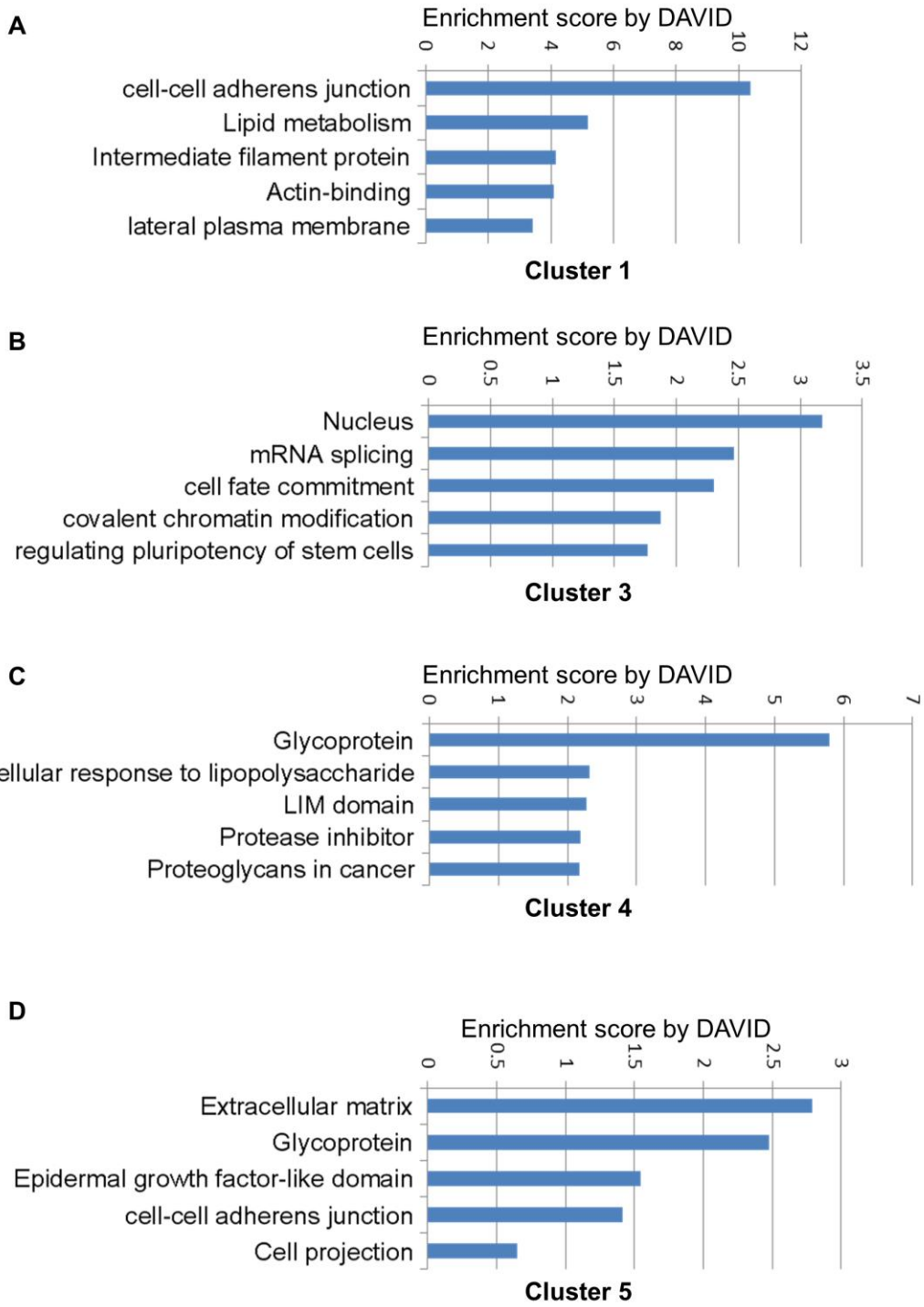
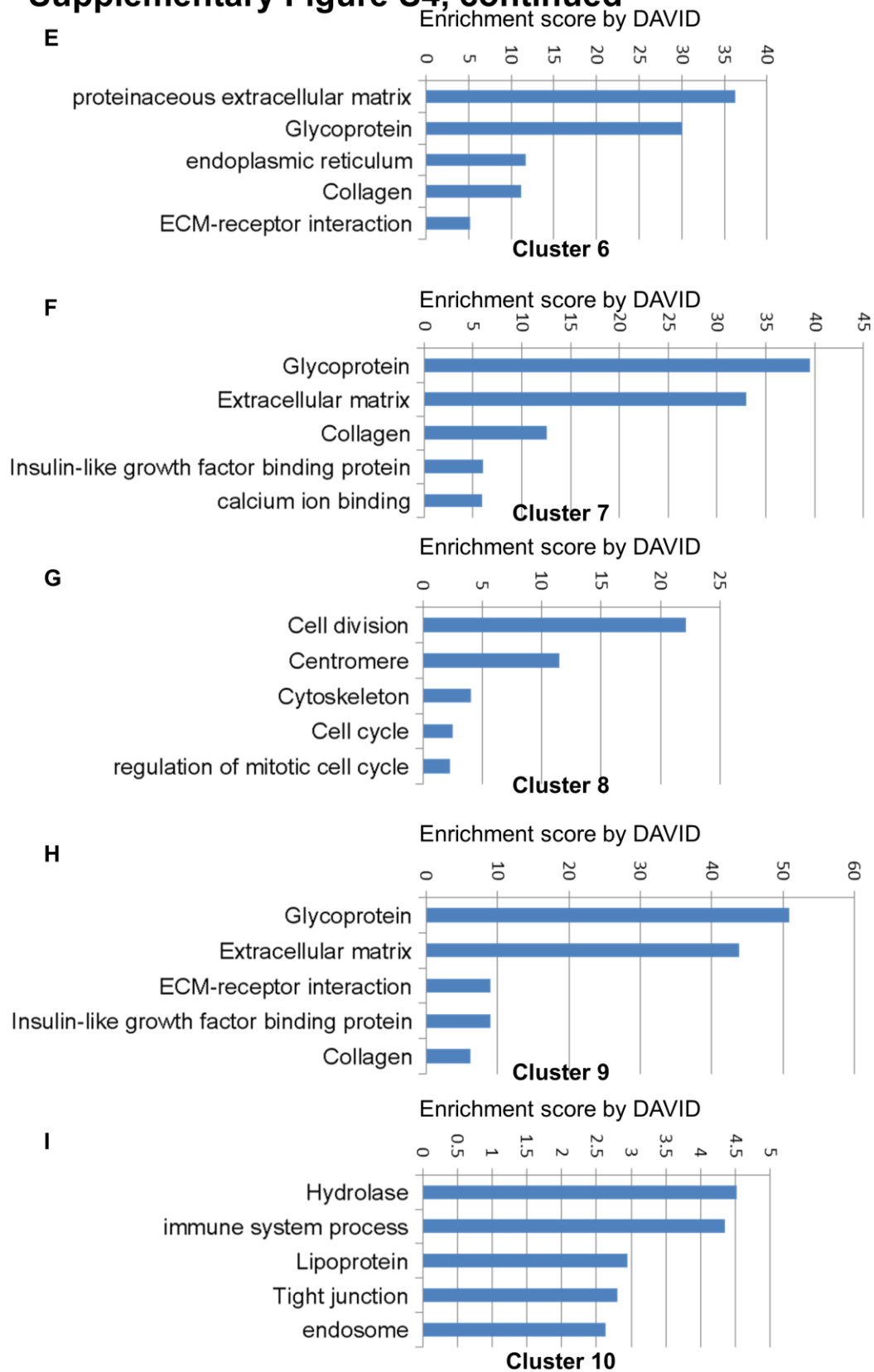


Supplementary Figure S4



Supplementary Figure S4, continued



Supplementary Figure S4. Gene annotation enrichment analysis of each cluster showed functional differences between clusters. Enrichment score was analyzed using DAVID.