

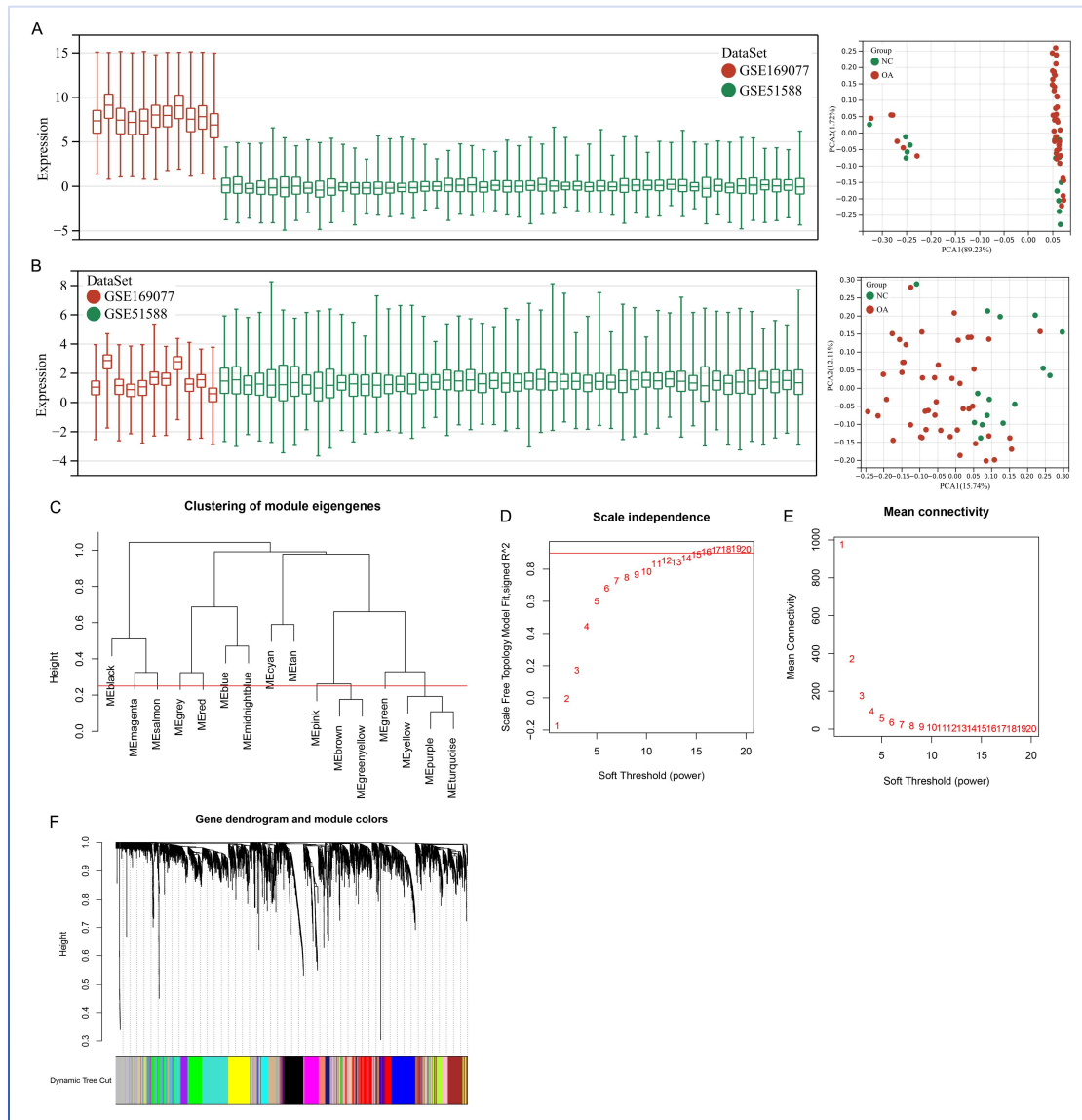


Fig. S1. Met-DCG identification. (A) Batch effect and PCA were conducted respectively on the datasets of GSE51588 and GSE169077 before their combination. (B) Batch effect and PCA were conducted respectively on the datasets of GSE51588 and GSE169077 after their combination. (C) Cluster tree plot analyses of the relationship between 13 module eigengenes and OA, with the red line corresponding to a set height of 0.25. (D, E) Scale independence, mean connectivity, and scale-free topology graphs generated with a soft threshold of 15. (F) A cluster dendrogram for co-expression network modules (1-TOM).

