

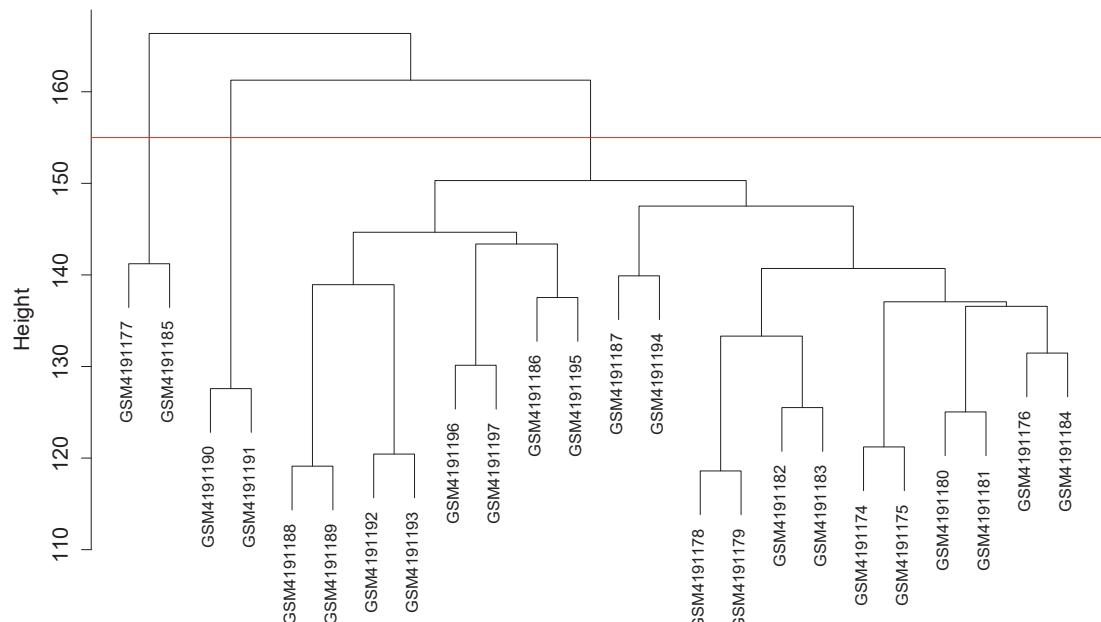
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Supplementary Figure S1

A

Sample clustering to detect outliers (GSE140947)



B

Sample clustering to detect outliers (GSE7084)

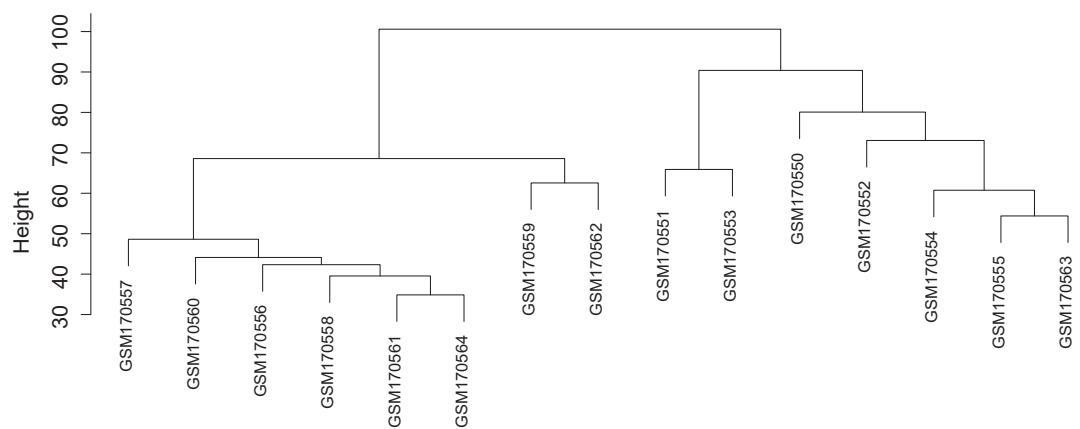


Figure S1: Clustering dendrogram of GSE140947 and GSE7084 datasets in WGCNA analysis. **(A)** GSE140947 dataset; **(B)** GSE7048 dataset.

Supplementary Figure S2

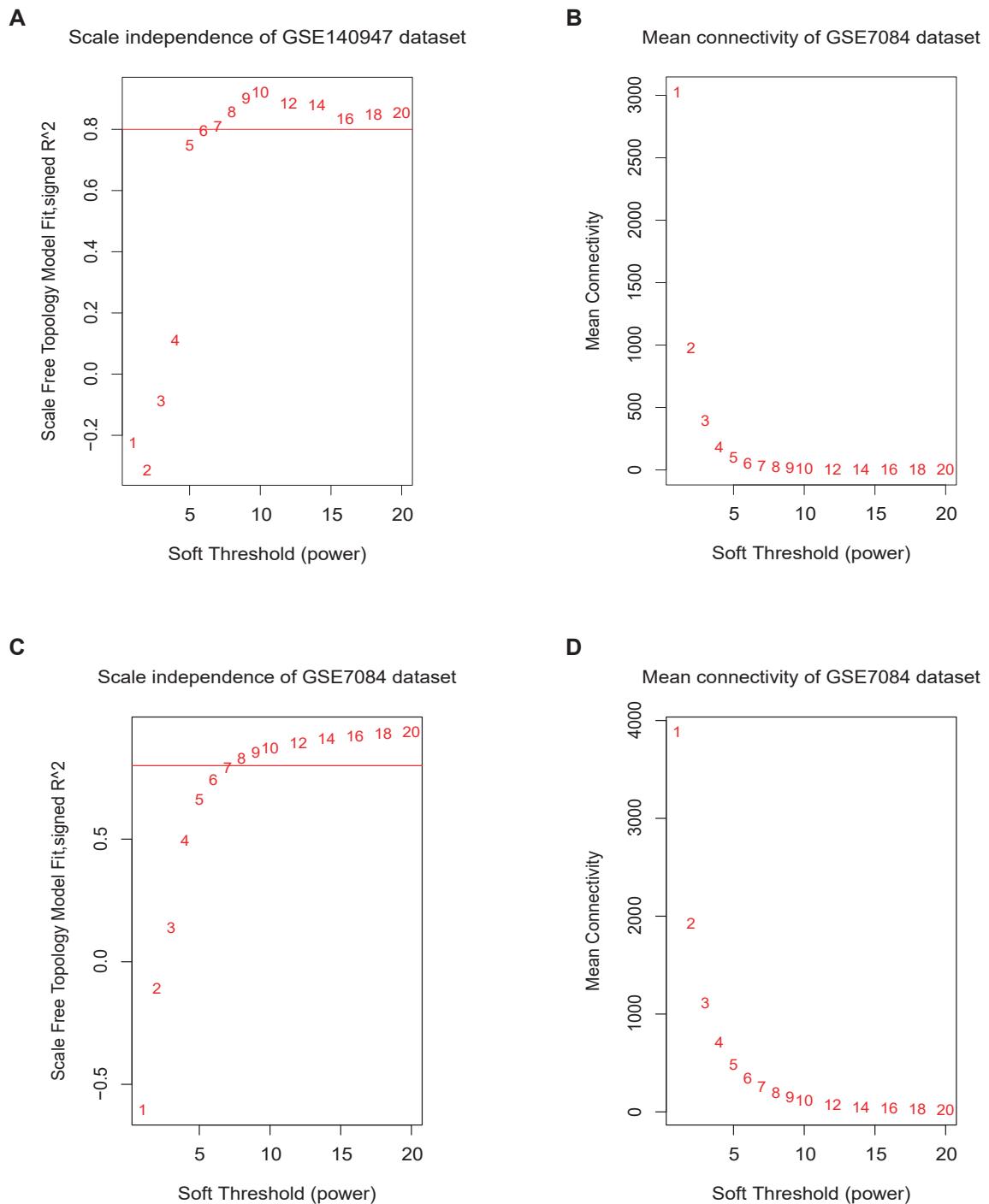


Figure S2: Identifying the soft threshold power in WGCNA analysis. The approximate scale-free fit index is achieved at the soft threshold power of 7 for both GSE140947 (**A,B**) and GSE7084 (**C,D**) datasets.

Supplementary Figure S3



Figure S3: The overlapping genes between aging and inflammation, considered as inflammaging-related genes.