

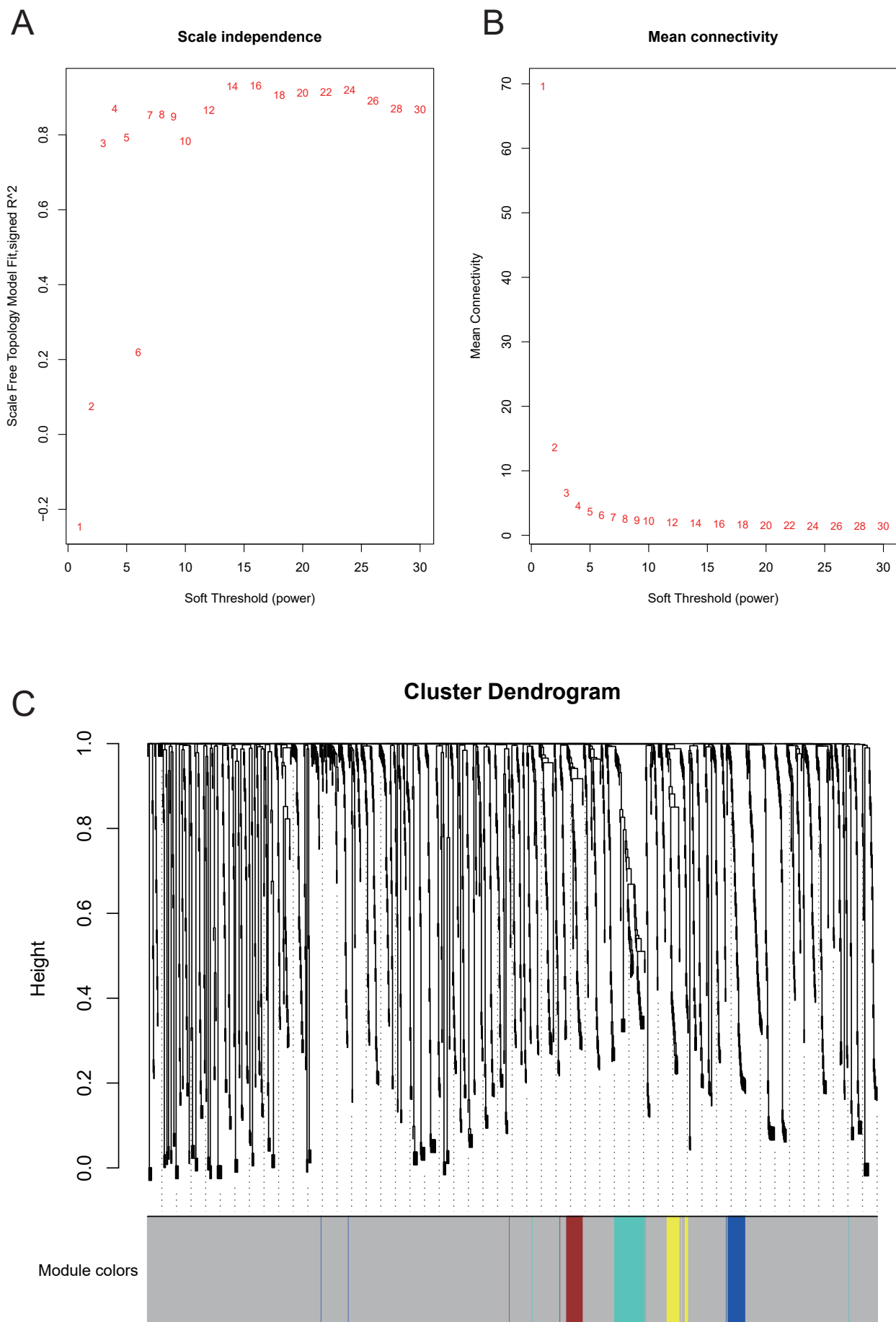


Fig. S1 MicroRNAs co-expression network construction. A The relationship between the scale-free fit index and various soft-thresholding powers. B The mean connectivity under different weighting coefficients. C Clustering dendrogram of miRNAs, various colors represent different modules.

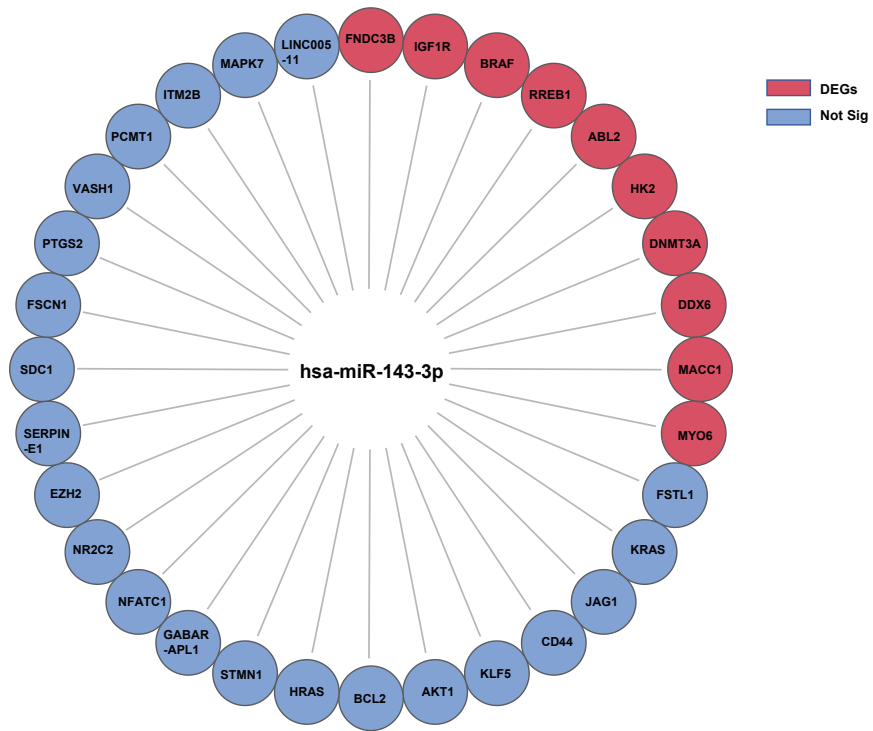


Fig. S2 Ten target genes of miR-143-3p were differentially expressed between the prefrontal cortex of patients with bipolar disorder and healthy controls. DEGs differentially expressed genes. Not Sig Not significant differentially expressed genes.