Integrated transcriptome and single-cell sequencing reveals M2-like tumor-associated macrophages-related prognostic signatures in breast cancer

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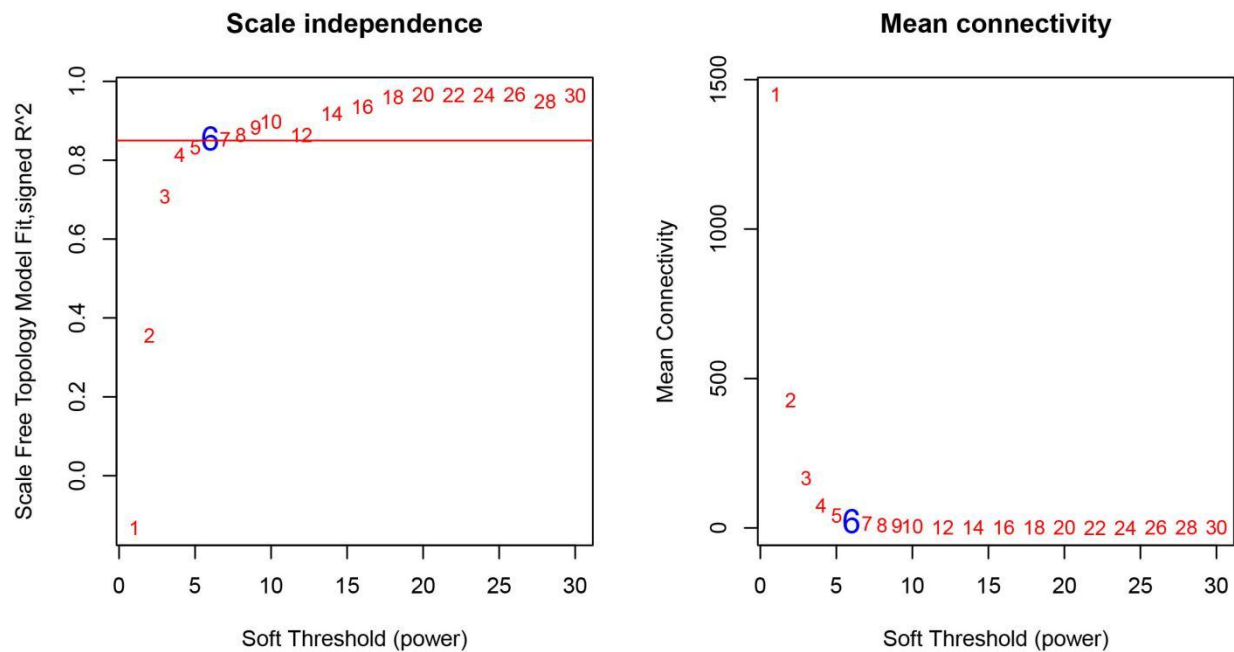
Supplementary Information

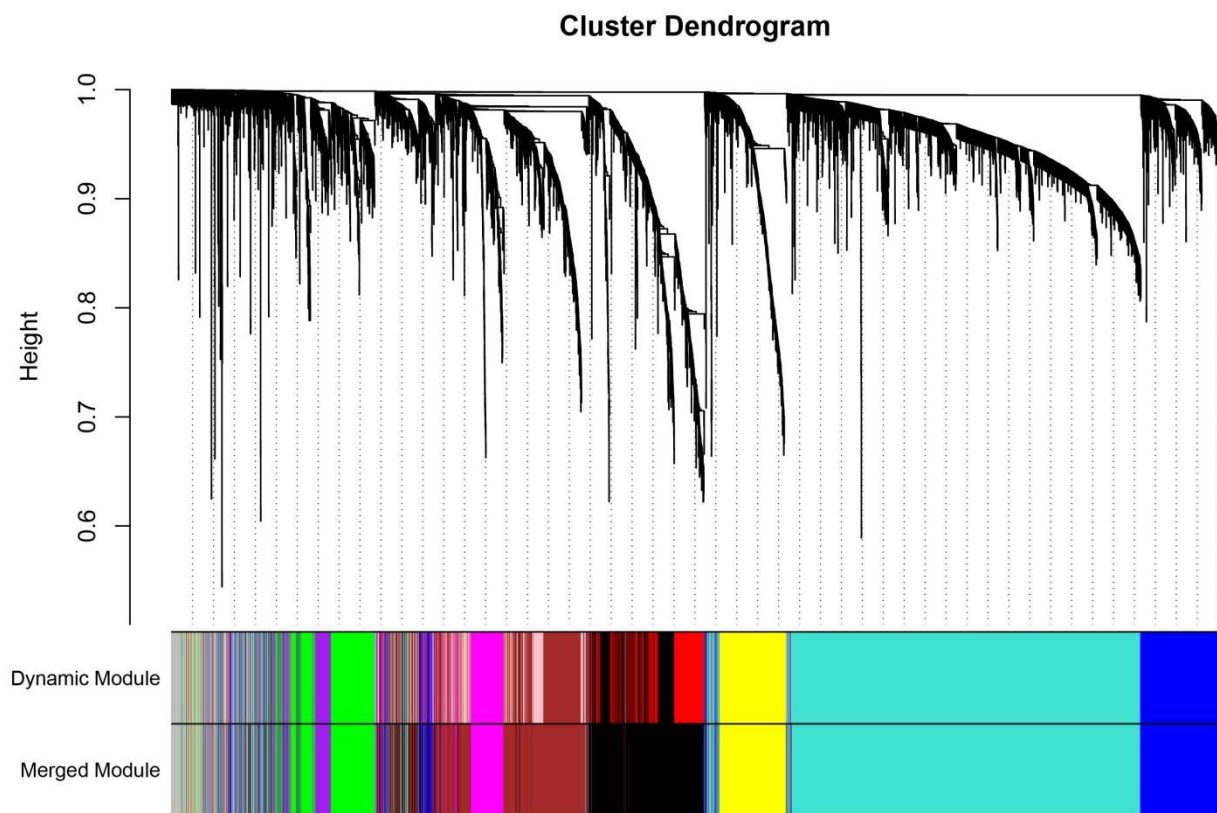**Figure S1.** Screening of the optimal soft threshold by pickSoftThreshold function.

Figure S2. Gene dendrogram and module colors.