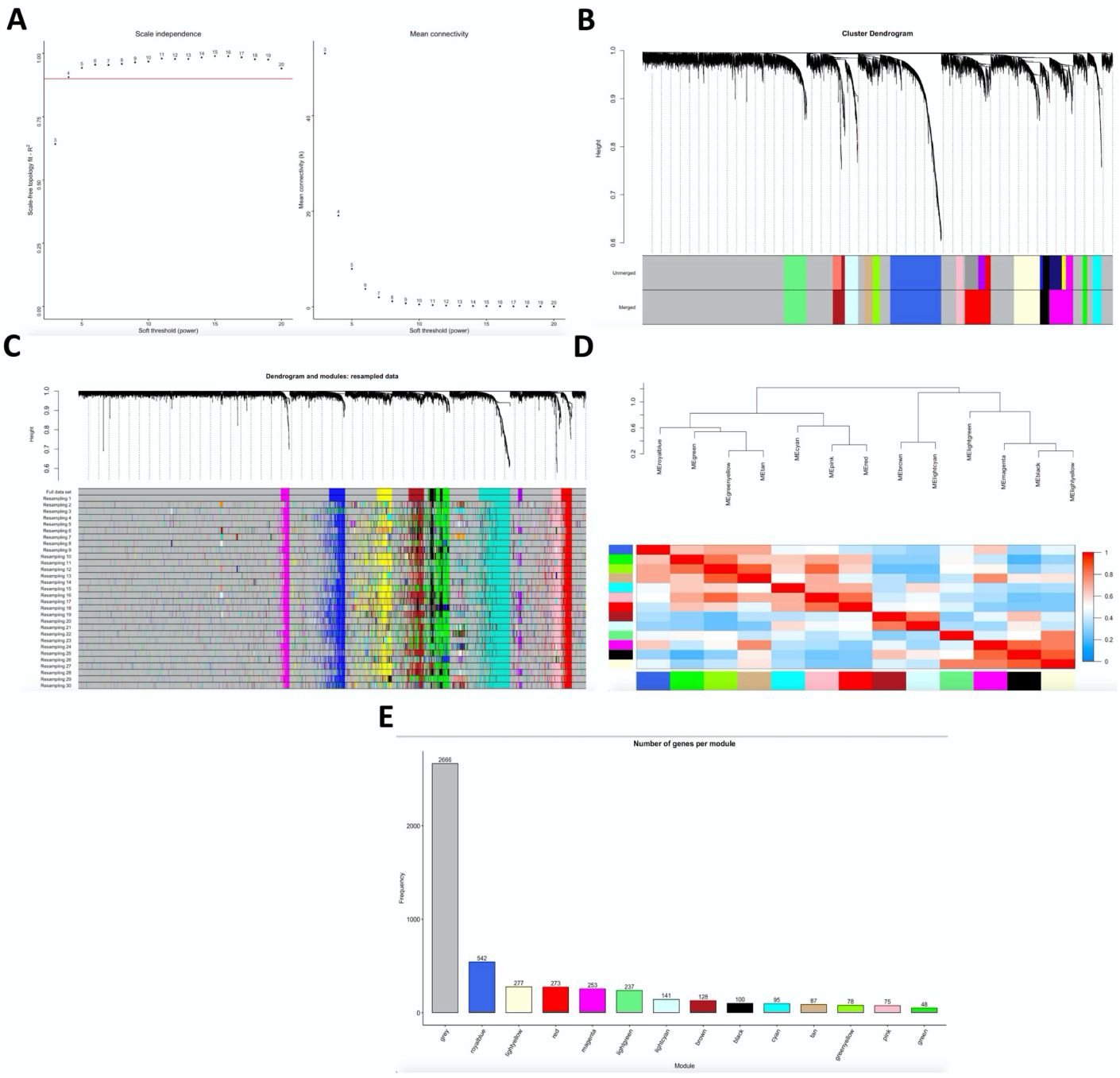
A
Scale independence
Mean connectivity
Scale-free topology fit - R²
Mean connectivity (k)
Soft threshold (power)
Soft threshold (power)
B
Cluster Dendrogram
Height
Unmerged
Merged
C
Dendrogram and modules: resampled data
Height
Full data set
Resampling 1
Resampling 2
Resampling 3
Resampling 4
Resampling 5
Resampling 6
Resampling 7
Resampling 8
Resampling 9
Resampling 10
Resampling 11
Resampling 12
Resampling 13
Resampling 14
Resampling 15
Resampling 16
Resampling 17
Resampling 18
Resampling 19
Resampling 20
Resampling 21
Resampling 22
Resampling 23
Resampling 24
Resampling 25
Resampling 26
Resampling 27
Resampling 28
Resampling 29
Resampling 30
D
MEroyalblue
MEgreen
MEgreenyellow
MEtan
MEcyan
MEpink
MEred
MEbrown
MElightcyan
MElightgreen
MEmagenta
MEblack
MEdarkyellow
E
Number of genes per module
Frequency
Module
grey
royalblue
lightyellow
red
magenta
lightgreen
lightcyan
brown
black
cyan
tan
greenyellow
pink
green
2666
542
277
273
253
237
141
128
100
95
87
78
75
48