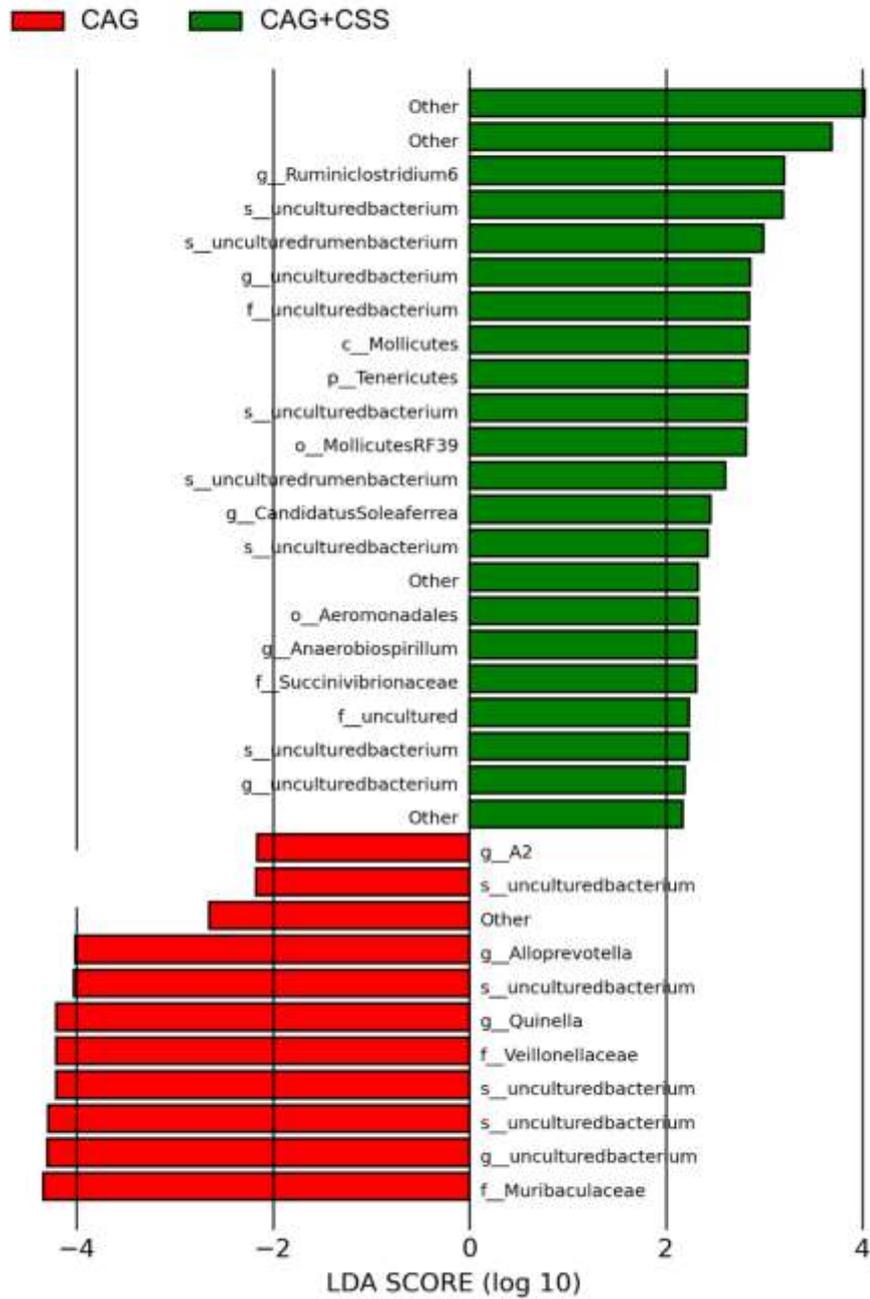


Supplementary Figure 1 The herb-active ingredient interaction network. A: Blue nodes represent the herbs in CSS, orange nodes represent the active ingredients, and each edge represents the interaction between the node; B: The active ingredients-targets interaction network. Blue nodes represent the active ingredients, green nodes represent the targets, and each edge represents the interaction between the node; C: GO enrichment histogram of MF, CC, BP. Each histogram shows the top 20 entries.



Supplementary Figure 2 LDA Score Distribution Histogram.

Supplementary Table 1 Molecular docking analysis

	CurPocket ID	Vina score
Baicalin-I κ B α	C1	-8.7
	C2	-7.3
	C3	-7.7
	C4	-8.9
	C5	-8.3
Glabrone-I κ B α	C1	-9.1
	C2	-7.0
	C3	-7.8
	C4	-8.7
	C5	-8.4
Glycyrrhiza flavonol A-I κ B α	C1	-8.5
	C2	-7.8
	C3	-8.2
	C4	-9.3
	C5	-8.7
Licochalcone B-I κ B α	C1	-8.1
	C2	-7.4
	C3	-6.4
	C4	-8.4
	C5	-6.4
Marmin-I κ B α	C1	-7.5
	C2	-9.0
	C3	-6.7
	C4	-7.8
	C5	-7.4
Licoisoflavone B-I κ B α	C1	-8.6
	C2	-7.2
	C3	-7.8
	C4	-8.7

C5

-8.6
