

Electronic Supplementary Information

High efficiency screening of nine lipid-lowering adulterants in herbal dietary supplements by using thin layer chromatography coupled with surface enhanced Raman spectroscopy

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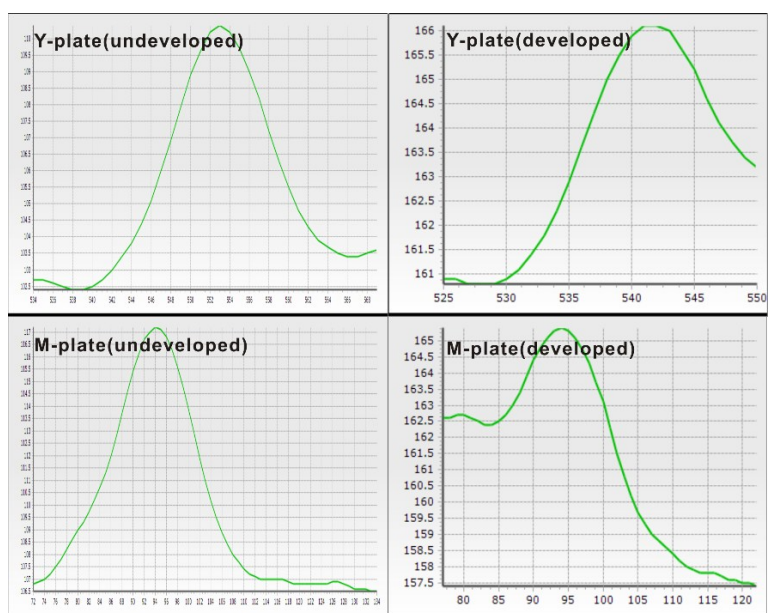


Fig. S1. The chromatographic peaks of undeveloped and developed spots on different TLC plates.

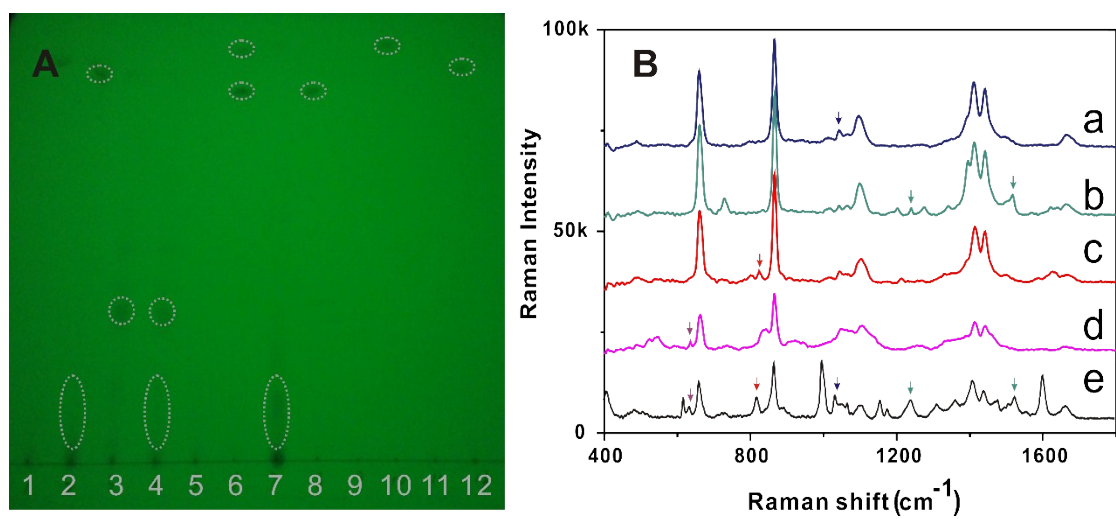


Fig. S2. (A) The TLC image of HDS matrix developed by dichloromethane-methanol-water 8:2:0.2 (v/v) and (B) the SERS spectra of matrix (a-d) and ATO (e).



Fig. S3. The TLC image of HDS matrix developed by petroleum ether-acetic ether - acetic acid 5.5:2.8:1(v/v).

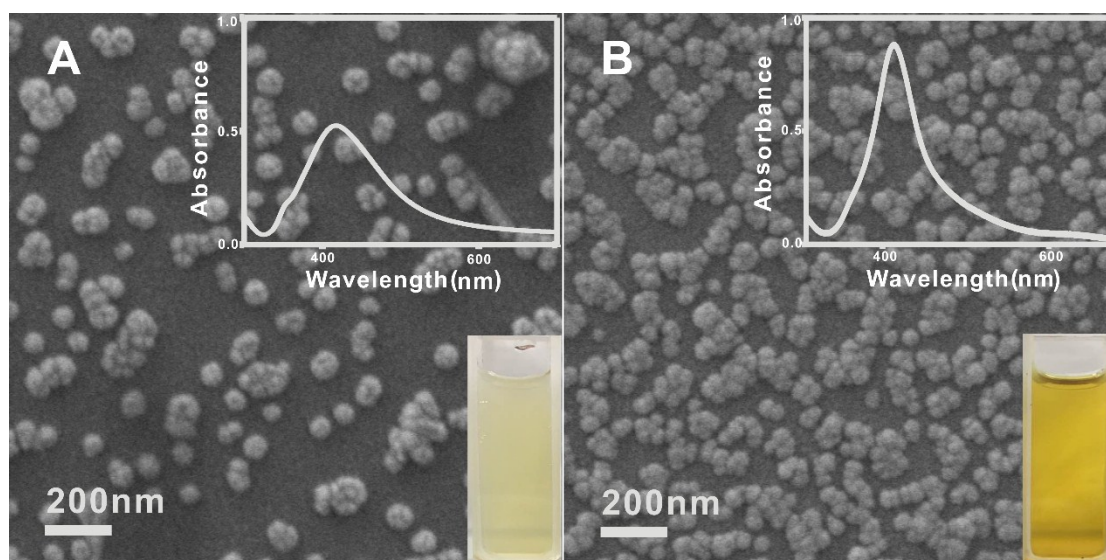


Fig. S4. UV-vis spectra and SEM image of the prepared silver colloids: (A) L-colloids and (B) D-colloids. The inset shows the accompanying color of the prepared silver colloids.

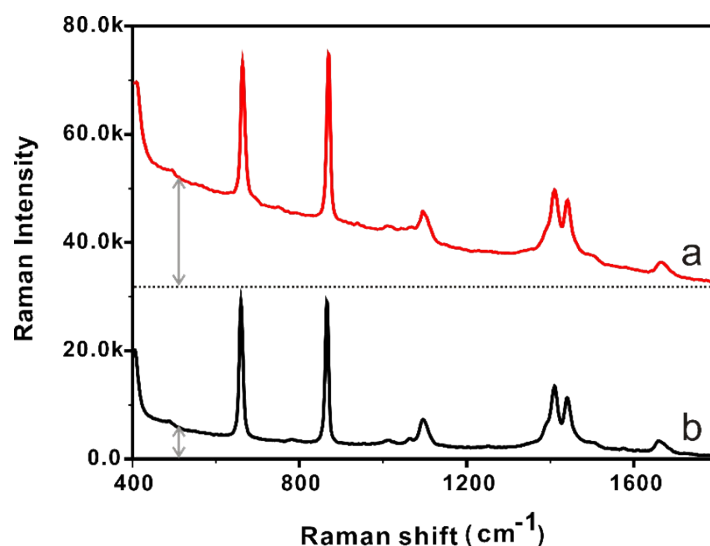


Fig. S5. The fluorescence interference from different HPTLC plates: (a) Y-plate and (b) M-plate.

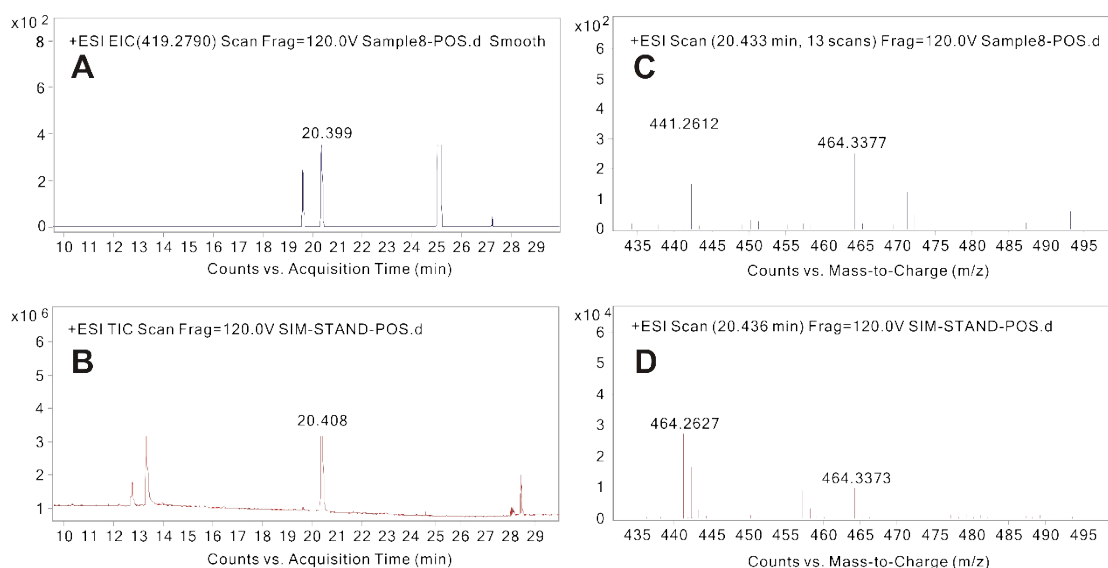


Fig. S6. HPLC spectra of (A) sample 8 and (B) standards of SIM; MS spectra of (C) sample 8 and standard of (D) SIM.

LC-MS verification conditions: ACQUITY UPLCTM BEH C18 column (2.1 mm×100 mm, 1.7μm, Waters, Milford, MA) was used with a mobile-phase gradient prepared from formic acid (component A) and 0.1% formic acid in acetonitrile (component B). The gradient was: held at 5% B for 2 min, linear increase from 5-25% B in 7 min, linear increase from 25-55% B in 11 min, linear increase from 60-95% B in 3 min, and held at 95 % B for 2 min. The flow rate was 0.35 mL/min, the injection volume was 4 μL, and the column temperature was 40°C. The following MS conditions were applied: ion source was the ESI source, using the positive ion mode (detection of PRO with negative ion mode). MS scanning range was from 100 to 1100 m/z.

Table S1 LOD comparison of six analytes by LC-MS/MS method and TLC-SERS method (unit: μg).

	NIC	PRA	ATO	FLU	BEZ	SIM	FEN
LC-MS/MS	0.02	0.012	0.004	0.004	0.02	0.002	0.002
TLC-SERS	0.0025	0.01	0.0025	0.01	0.001	0.05	0.005

Table S2 Validation of SIM in sample 8 by UPLC-QTOF/MS.

Name	Identification formula [M+H]⁺	Mass	Error (ppm)
SIM standard	$\text{C}_{25}\text{H}_{38}\text{O}_5$	418.5719	-0.38
Sample 8	$\text{C}_{25}\text{H}_{38}\text{O}_5$	418.5714	-0.12