

Supporting Information

In vivo selective capture and rapid identification of luteolin and its metabolites in rat livers by molecularly imprinted solid-phase microextraction

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2 **1. Instrument and Chromatographic Conditions.**

3 Analyses were carried out on an Agilent 1260 Series HPLC system (Agilent
4 Technologies, Palo Alto, California, USA). Chromatographic separation was
5 performed using an Agilent SB-C18 column (150×4.6 mm, particle size 5 μm).
6 HPLC-MS/MS analysis was based on a Nexera X2 HPLC system (Shimadzu, Japan)
7 coupled to an AB Sciex Triple 5600 tandem mass spectrometer (Applied 24
8 Biosystems/MDS Sciex, MA, USA). A Zorbax Eclipse plus C18 column, Rapid
9 Resolution HD (2.1×100mm, 1.8-Micron) was used for separation. JSM-7600F Field
10 emission scanning electron microscope (JEOL, Japan) was used to observe
11 morphology of MIPs. Fourier transform infrared (FT-IR) spectra were obtained via an
12 IR Affinity-1 FT-IR spectrometer (Shimadzu, Japan). The surface-area and the
13 pore-size distribution were obtained via Autosorb - iQ automatic specific surface and
14 pore size distribution analyzer (Autosorb-iQ, America).

15 **2. Surface Treatment of Stainless Steel Fibers**

16 Firstly, stainless steel fibers were cut into 4.0 cm of segments (each about 20.0
17 mg). Then put the cut fibers into acetone and ultrasonic for 15 min to clean the
18 surface. The fibers were washed with methanol for another three times and dried in
19 oven. Secondly, the self-assembly of dopamine was conducted on fibers. The fibers
20 were immersed into 10 mL of Tris-HCl solution containing 6 mg/mL of dopamine
21 (pH8.5) and kept for 24 h in dark. Then, the fibers were washed with methanol for
22 three times to obtain the modified stainless steel fibers. Thirdly, silylation was carried
23 out on the dopamine modified fibers. The fibers were soaked in a mixture of

24 3-MPS/H₂O/Methanol (v/v/v, 1:1:8) and kept for 30 min, then took out the fibers and
25 dried in a vacuum drying oven for 150 min at 150 °C. The dried fibers were washed
26 with methanol for three times and dried under nitrogen.

27 **3. Calibration curve of luteolin analyzed by HPLC**

28 10 mg of luteolin was dissolved in 10.0 mL methanol to prepare the stock
29 solution (1 mg/mL) and working standard solutions were diluted using 55% (v/v)
30 methanol-water with a series of concentrations (1, 5, 10, 20, 40, 60, 80 and 100
31 µg/mL, respectively). The absorbance of different concentration of luteolin was
32 measured using HPLC.

33 **4. Preparation of liver homogenates**

34 Rats decapitated and livers were transferred into a 5 mL beaker. Then livers were
35 washed with saline till becoming a gray. After that, pieces of fresh livers of rats
36 (approximately 0.5 g each group) were transferred into a microcentrifuge tube (2 mL)
37 prior to homogenization with a tissue tearor for 30 s.

38 **5. *Extraction Time Profile***

39 The extractions by using MIPs fibers were performed at 2, 5, 10, 15 and 20 min
40 from prepared liver homogenates spiked with 30 µg/mL of luteolin. Then the fibers
41 were quickly rinsed for 30 s in ultrapure water and dried under nitrogen. After that,
42 fibers were desorbed in 500 µL (minimum volume to ensure complete immersion of
43 the coating into the solvent) of methanol/acetic acid (8:2, v/v) to elute luteolin. Finally,
44 eluants were dried under nitrogen and dissolved in 100 µL methanol for HPLC
45 analysis.

46 **6. Diffusion-Based Calibration: Pre-Determined Sample In Rates**

47 Different concentrations of target luteolin were prepared in 0.5 g prepared liver
48 homogenates (0.1, 0.25, 0.5, 1, 5, 10, 20 and 30 $\mu\text{g/mL}$) to determine sampling rates
49 in matrix-matched samples. On the basis of the extraction time profile (Fig.6A),
50 MIP-SPME sampling was performed for 10 min. Adsorption by MIP-SPME fibers,
51 rinsing and desorption steps were performed using the same conditions as described in
52 section 2.8.2. The total concentrations in each liver homogenates were measured by
53 the following method: after adding the different concentrations of luteolin in 0.5 g
54 prepared liver homogenates (0.1, 0.25, 0.5, 1, 5, 10, 20 and 30 $\mu\text{g/mL}$), 1 mL ethyl
55 acetate were added into liver homogenates to extract luteolin. Liver homogenates
56 solutions then were centrifuged at $13000\times g$ for 20 min at 4°C and the supernatants
57 were obtained. After that, the supernatants were dried under nitrogen and dissolved in
58 100 μL methanol for HPLC analysis.

59 **7. HPLC analysis**

60 For the evaluation of the adsorption capacity of MIPs and the selectivity of the
61 MIPs toward structural analogues in section 2.7, the gradient elution process of
62 methanol (solvent A) and 0.2% phosphoric acid aqueous solution (solvent B) was as
63 followed: 0-10 min, 45%-55% A; 10-15 min, 55% - 60% A; 15-20 min, 60% - 80% A;
64 20-25 min, 80% A; 25-30 min, 80% - 45% A. The flow rate was set at 0.8 mL/min.
65 The column temperature was maintained at 30°C and injection volume was 10 μL .
66 The detective wavelength was set at 350 nm.

67 For the analysis of luteolin in liver samples, the gradient elution process of

68 methanol (solvent A) and 0.2% phosphoric acid aqueous solution (solvent B) was as
69 followed: 0-50 min, 40%-42% A; 50-60 min, 42% A; 60-75 min, 42%-52% A; 75-77
70 min, 52%-40% A. The flow rate was set at 0.8 mL/min. The column temperature was
71 maintained at 30 °C and injection volume was 10 µL. The detective wavelength was
72 set at 350 nm.

73 For the analysis of metabolites of luteolin in liver sample, high-performance
74 liquid chromatography-tandem mass spectrometry (MS/MS) analysis was based on an
75 Nexera X2 HPLC system (Shimadzu, Japan) coupled to an AB Sciex Triple
76 5600 tandem mass spectrometer (Applied Biosystems/MDS Sciex, MA, USA). A
77 Zorbax Eclipse plus C18 column, Rapid Resolution HD (2.1×100mm, 1.8-Micron)
78 was used for separation. An aqueous solution containing 0.1% formic acid (A) and (B)
79 acetonitrile were used as the mobile phases, the gradient elution process was as
80 followed: 0-2 min, 5% B→25% B; 2-5 min, 25% B→38% B; 5-30 min, 38% B→100%
81 B; 30-32 min, 100% B; 32-32.1 min, 100%→5% B; 32.1-35 min, 15% B→stop. The
82 flow rate was set at 0.3 mL·min⁻¹, the column temperature was maintained at 30 °C
83 and injection volume was 2 µL.

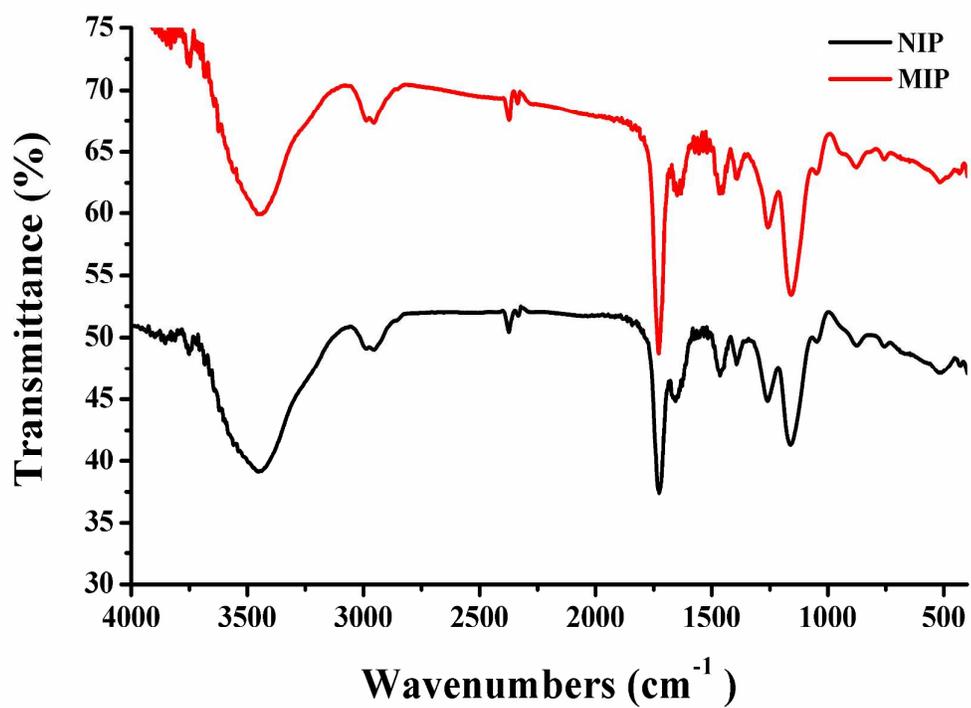
84 High resolution mass spectrometry (MS) detections were performed on AB Sciex
85 Triple 5600 tandem mass spectrometer (Applied Biosystems/MDS Sciex, MA,
86 USA), connecting to the HPLC. The conditions of MS/MS spectra were as followed:
87 ion source: electronic electrospray ionization (ESI) source; Ionization ways: positive
88 ion scanning; TOF - MS quality scan range (m/z): 100-2000 da. Quality of TOF
89 MS/MS scan range (100-2000 m/z) da; auxiliary gas (GS1 and GS2), 60 and 60 psi,

90 respectively; Air curtain air pressure (CUR): 35 psi. (TEM) ion source temperature:
91 600 °C; Ion spraying voltage (ISVF): 5500 v; To cluster voltage (DP): 100 v, the
92 parent ion collision energy (CE): 10, the child away from the collision energy (CE):
93 50, sub ion collision energy difference (CES): 10. Control of the system, data
94 acquisition and data analysis were all carried out using the Analyst 1.5.2 software
95 (Applied Biosystems).

96 **8. IR analysis**

97 As shown in **Figure S1**, MIPs (after elution) and NIPs coatings had similar IR
98 spectra, which indicated that their backbone structure were similar. The main
99 functional groups of MIPs coatings could be found with corresponding infrared
100 absorption peaks. Four strong infrared absorption peaks of 3448, 2954, 1726, and
101 1664 cm^{-1} were found, which were attributed to hydroxyl groups, methyl groups,
102 carbonyl groups, and alkenes (C=C bonds), respectively. The characteristic peak of
103 C=O at 1726 cm^{-1} in the MIPs spectrum showed that the MIPs were synthesized
104 through the polymerization of EGDMA and AM. Moreover, the special peak intensity
105 of the stretching vibration of C=C (1664 cm^{-1}) in the MIPs was crucial for the
106 chemical bonding between MIP layer and layer during the coating procedures.

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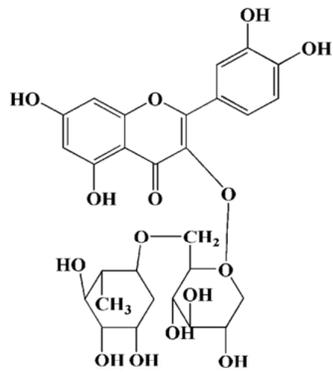
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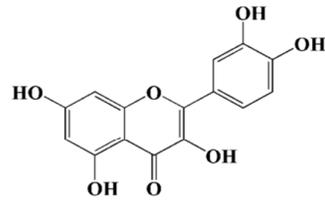
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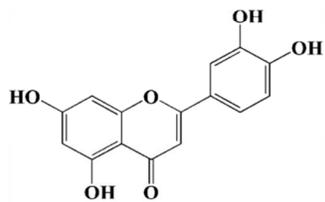
Figure S1. The FT-IR spectra of MIPs (after elution) and NIPs.



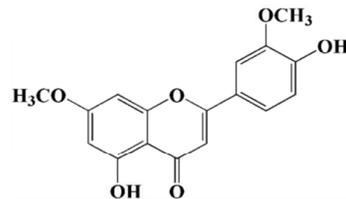
1. rutin



2. quercetin



3. luteolin



4. ombuin

Figure S2. The structure of luteolin and its structural analogues.

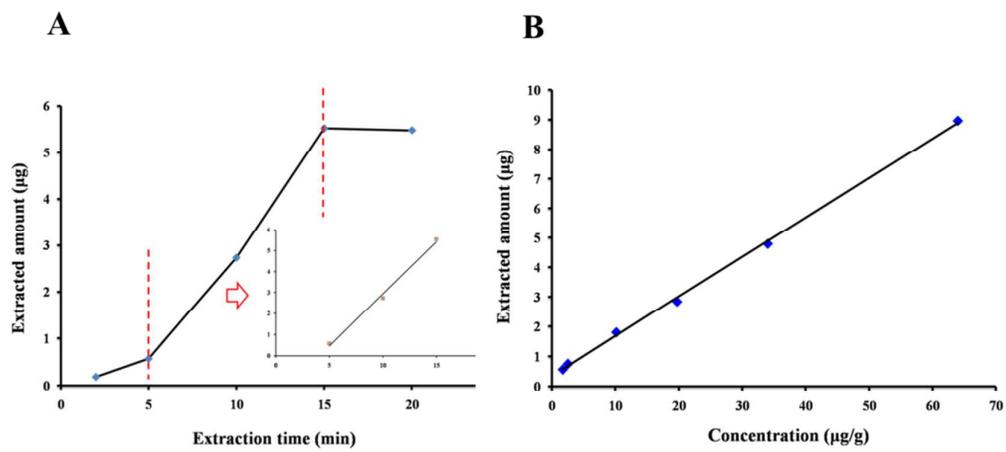


Figure S3. n-t (A) and n-Cs (B) curves of *in vitro* MIP-SPME of rat livers homogenates.

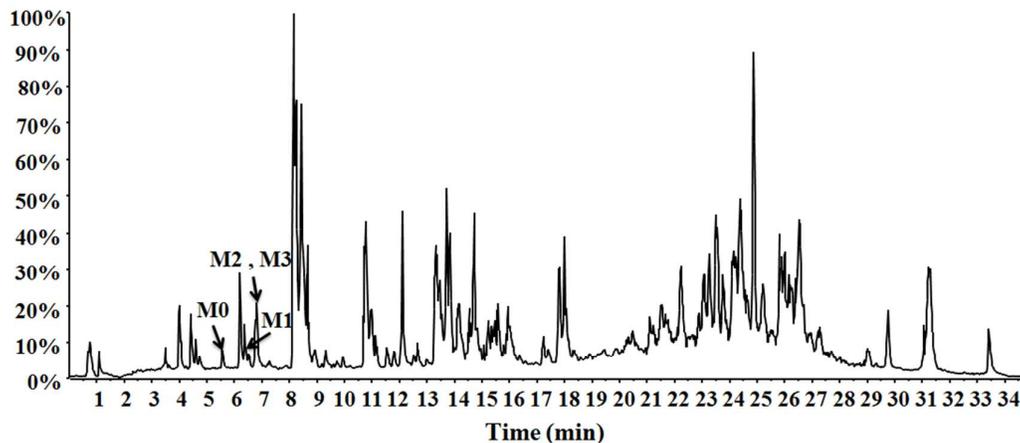


Figure S4. The total ion current of extract extracted by using MIP-SPME sampling method.

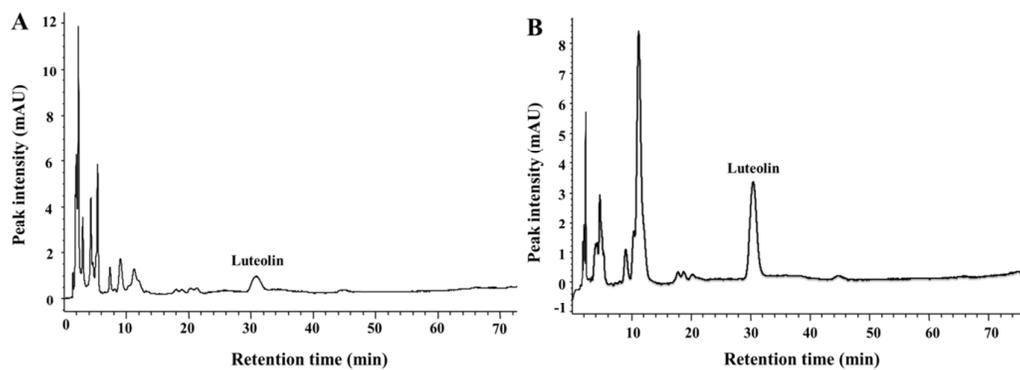


Figure S5. The chromatogram of extract in rat livers at 120 min by using commercial PDMS fiber (A) and DVB fiber.

Table S1 Specific Surface Area, Pore Structure Parameters of MIPs and NIPs Coatings

Samples	BET Surface area (m ² /g)	BJH desorption cumulative volume of pores (cm ³ /g)	BJH desorption average pore radius (nm)
MIP-SPME	320.49	0.367	18.94
NIP-SPME	202.49	0.236	17.01

Table S2 Luteolin and Its Metabolites Identified in Rat Livers Using UPLC-MS/MS

Name	RT/min	[M+H] ⁺	MS/MS fragment ions
M0	5.613	287.0280	269.0195, 258.0264, 241.0240, 213.0315, 161.0029, 152.9990, 135.0262
M1	6.618	271.0339	152.9998
M2	6.860	301.0434	286.0438, 257.0195, 229.0262, 203.0120, 152.9993
M3	6.875	301.0434	286.0207, 257.0186, 229.0255, 203.0116, 152.9993