A rapid protocol to distinguish between Citri Exocarpium Rubrum and Citri Reticulatae Pericarpium based on characteristic fingerprint and UHPLC-Q-TOF MS methods

Liqiang Shi^a, Rongjin Wang^a, Tianshu Liu^a, Jiajie Wu^a, Hongxu Zhang^a, Zhiqiang

Liu^b, Shu Liu^b, Zhongying Liu^{a, *}

^a School of Pharmaceutical Sciences, Jilin University, Changchun 130021, China

^b National Center of Mass Spectrometry in Changchun & Jilin Province Key

Laboratory of Chinese Medicine Chemistry and Mass Spectrometry, Changchun

Institute of Applied Chemistry, Chinese Academy of Sciences, Changchun, 130022,

China

*Corresponding authors: Zhongying Liu

School of Pharmaceutical Sciences, Jilin University, Changchun 130021, China

Tel.: +86-431-85262613; Fax: +86-431-85262236

E-mail address: <u>liuzy@jlu.edu.cn</u> (Z. Y. Liu)

Sample	Sample No.	Origins	hesperidin%	
CER	J1	Jinhua,Zhejiang	3.76	
CER	J2	Jinhua,Zhejiang	4.12	
CER	J3	Jinhua,Zhejiang	4.85	
CER	J4	Jinhua,Zhejiang	3.90	
CER	J5	Jinhua,Zhejiang	4.41	
CER	J6	Chengdu,Sichuan	3.44	
CER	J7	Chengdu,Sichuan	3.06	
CER	J8	Chengdu,Sichuan	3.78	
CER	J9	Chengdu,Sichuan	3.85	
CER	J10	Chengdu,Sichuan	3.38	
CER	J11	Xinhui,Guangdong	3.34	
CER	J12	Xinhui,Guangdong	3.97	
CER	J13	Xinhui,Guangdong	4.14	
CER	J14	Xinhui,Guangdong	4.88	
CER	J15	Xinhui,Guangdong	4.07	
CRP	C1	Linyi,Shandong	3.56	
CRP	C2	Linyi,Shandong	4.18	
CRP	C3	Linyi,Shandong	4.10	
CRP	C4	Linyi,Shandong	3.62	
CRP	C5	Linyi,Shandong	3.73	

Table.S1 The content of hesperidin in 15 batches of CER and CRP.

CRP	C6	Xinhui,Guangdong	4.08
CRP	C7	Xinhui,Guangdong	3.78
CRP	C8	Xinhui,Guangdong	4.46
CRP	С9	Xinhui,Guangdong	4.86
CRP	C10	Xinhui,Guangdong	4.65
CRP	C11	Suizhou,Hubei	3.14
CRP	C12	Suizhou,Hubei	3.52
CRP	C13	Suizhou,Hubei	4.79
CRP	C14	Suizhou,Hubei	4.31
CRP	C15	Suizhou,Hubei	4.04

Sample No.	Origins	Samples	
J1	Jinhua,Zhejiang	Citri Exocarpium Rubrum	
J2	Jinhua,Zhejiang	Citri Exocarpium Rubrum	
J3	Jinhua,Zhejiang	Citri Exocarpium Rubrum	
J4	Jinhua,Zhejiang	Citri Exocarpium Rubrum	
J5	Jinhua,Zhejiang	Citri Exocarpium Rubrum	
J6	Chengdu,Sichuan	Citri Exocarpium Rubrum	
J7	Chengdu,Sichuan	Citri Exocarpium Rubrum	
J8	Chengdu,Sichuan	Citri Exocarpium Rubrum	
J9	Chengdu,Sichuan	Citri Exocarpium Rubrum	
J10	Chengdu,Sichuan	Citri Exocarpium Rubrum	
J11	Xinhui,Guangdong	Citri Exocarpium Rubrum	
J12	Xinhui,Guangdong	Citri Exocarpium Rubrum	
J13	Xinhui,Guangdong	Citri Exocarpium Rubrum	
J14	Xinhui,Guangdong	Citri Exocarpium Rubrum	
J15	Xinhui,Guangdong	Citri Exocarpium Rubrum	
C1	Linyi,Shandong	Citri Reticulatae Pericarpiu	
C2	Linyi,Shandong	Citri Reticulatae Pericarpiun	
C3	Linvi.Shandong	Citri Reticulatae Pericarpiu	

fable.S2 The origins an	d batches of CER and CR	Ρ.
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C4	Linyi,Shandong	Citri Reticulatae Pericarpium		
C5	Linyi,Shandong	Citri Reticulatae Pericarpium		
C6	Xinhui,Guangdong	Citri Reticulatae Pericarpium		
C7	Xinhui,Guangdong	Citri Reticulatae Pericarpium		
C8	Xinhui,Guangdong	Citri Reticulatae Pericarpium		
С9	Xinhui,Guangdong	Citri Reticulatae Pericarpium		
C10	Xinhui,Guangdong	Citri Reticulatae Pericarpium		
C11	Suizhou,Hubei	Citri Reticulatae Pericarpium		
C12	Suizhou,Hubei	Citri Reticulatae Pericarpium		
C13	Suiz hou,Hubei	Citri Reticulatae Pericarpium		
C14	Suizhou,Hubei	Citri Reticulatae Pericarpium		
C15	Suizhou,Hubei	Citri Reticulatae Pericarpium		

Name	Unknow	Unknow	(mg/g) Unknow	Unknow	Unknow	Unknow
	n1	n2	n3	n4	n5	n6
Caffeoylquin ic acid A	0.24	0.66	0.31	0.64	0.68	0.29
Caffeoylquin ic acid B	0.58	0.92	0.44	0.72	0.95	0.36
Caffeoylquin ic acid C	1.15	1.20	0.57	0.81	1.35	0.79
Caffeoylquin ic acid D	1.28	0.66	1.25	1.23	1.29	1.21
Vicenin-2	0.46	0.84	0.51	0.52	0.65	0.49
Melitidin	0.11	0.25	0.09	0.12	0.23	0.11
Nobiletin	0.21	0.30	0.20	0.18	0.29	0.19
3,5,6,7,8,3',4 '- heptemethox yflavone	0.24	0.37	0.23	0.24	0.39	0.23
Natsudaidain	0.04	0.1	0.06	0.05	0.07	0.05
Narirutin 4'- glucoside	1.94	0.86	1.95	1.92	0.77	2.17
Naringin	8.53	2.18	8.10	8.80	2.23	8.92
Hesperidin	6.58	3.94	6.35	6.72	5.23	6.73

Table.S3 The content of 12 chemical markers in 6 unknown samples.



Fig.S1 The UV full-wavelength scanning of CER sample.



Fig. S2 The fragments of caffeoylquinic acid A in negative ion mode.



Fig. S3 The fragments of caffeoylquinic acid B in negative ion mode.



Fig. S4 The fragments of caffeoylquinic acid C in negative ion mode.



Fig. S5 The fragments of caffeoylquinic acid D in negative ion mode.



Fig. S6 The possible fragmentation pattern of nobiletin.



Fig. S7 The possible fragmentation pattern of hesperidin.



Fig. S8 The possible fragmentation pattern of luteolin-6, 8-di-C-glucoside.











Fig. S11 The UV full-wavelength scanning of narirutin 4'-glucoside, naringin and

hesperidin.



Fig. S12 The UV full-wavelength scanning of melitidin, nobiletin, natsudaidain and

3, 5, 6, 7, 8, 3', 4'-heptemethoxyflavone.



Fig. S13 The fragments of vanillic acid in negative ion mode.



Fig. S14 The fragments of caffeic acid in negative ion mode.



Fig. S15 The fragments of luteolin-6, 8-di-C-glucoside in positive ion mode.



Fig. S16 The fragments of luteolin-6, 8-di-C-glucoside in negative ion mode.



Fig. S17 The fragments of vicenin-2 in positive ion mode.



Fig. S18 The fragments of vicenin-2 in negative ion mode.



Fig. S19 The fragments of chrysoeriol-6, 8-di-C-glucoside in positive ion mode.



Fig. S20 The fragments of chrysoeriol-6, 8-di-C-glucoside in negative ion mode.



Fig. S21 The fragments of narirutin 4'-glucoside in positive ion mode.

Channel name: High energy : Time 14.2125 +/- 0.0793 minutes



Fig. S22 The fragments of narirutin 4'-glucoside in negative ion mode.







Fig. S24 The fragments of isoorientin in negative ion mode.

Channel name: High energy : Time 15.1474 +/- 0.0920 minutes



Fig. S25 The fragments of orientin in positive ion mode.







Fig. S27 The fragments of eriocitrin in positive ion mode.



Fig. S28 The fragments of eriocitrin in negative ion mode.



Fig. S29 The fragments of rutin in positive ion mode.



Fig. S30 The fragments of rutin in negative ion mode.

Channel name: High energy : Time 18.7205 +/- 0.0920 minutes



Fig. S31 The fragments of lonicerin in positive ion mode.







Fig. S33 The fragments of isolimonic acid in positive ion mode.

Channel name: High energy : Time 19.7095 +/- 0.0920 minutes



Fig. S34 The fragments of deacetylnomilinic acid in positive ion mode.







Fig. S36 The fragments of naringin in negative ion mode.

Channel name: High energy : Time 22.2551 +/- 0.0914 minutes



Fig. S37 The fragments of diosmin in positive ion mode.







Fig. S39 The fragments of hesperidin in positive ion mode.

5.26e6

Channel name: High energy : Time 23.0734 +/- 0.0793 minutes



Fig. S40 The fragments of hesperidin in negative ion mode.



Fig. S41 The fragments of nomilin in positive ion mode.



Fig. S42 The fragments of nomilinic acid in positive ion mode.

Channel name: High energy : Time 29.1440 +/- 0.0914 minutes



Fig. S43 The fragments of poncirin in positive ion mode.







Fig. S45 The fragments of citrusin Ⅲ in positive ion mode.

Channel name: High energy : Time 31.1533 +/- 0.0920 minutes







Fig. S47 The fragments of melitidin in negative ion mode.



Fig. S48 The fragments of citrusin I in positive ion mode.

Channel name: High energy : Time 32.5435 +/- 0.0920 minutes



Fig. S49 The fragments of isosinensetin in positive ion mode.



Fig. S50 The fragments of 3'-hydroxy- 5,6,7,8,4'- pentamethoxyflavanone in positive ion mode.



Fig. S51 The fragments of 5, 7, 8, 3', 4', 5'-hexamethoxyflavone in positive ion mode.

1.06e7



Fig. S52 The fragments of sinensetin in positive ion mode.



Fig. S53 The fragments of monohydroxy- pentamethoxyflavanone in positive ion mode.



Fig. S54 The fragments of 5, 6, 7, 4'-tetramethoxyflavone in positive ion mode.







Fig. S56 The fragments of nobiletin in positive ion mode.



Fig. S57 The fragments of 5, 7, 8, 4'-tetramethoxyflavone in positive ion mode.

Channel name: High energy : Time 36.2042 +/- 0.2761 minutes



Fig. S58 The fragments of 3, 5, 6, 7, 8, 3', 4'-heptemethoxyflavone in positive ion mode.



Fig. 859 The fragments of 5-hydroxy-3, 6, 7, 8, 3', 4'-pentamethoxyflavanone in positive ion mode.



Fig. S60 The fragments of tangeritin in positive ion mode.

Channel name: High energy : Time 38.4580 +/- 0.0920 minutes



Fig. S61 The fragments of hexamethoxyflavone in positive ion mode.



Fig. S62 The fragments of 5-hydroxy- 6, 7, 8, 3', 4'- pentamethoxyflavanone in positive ion mode.



Fig. S63 The fragments of monohydroxy- pentamethoxyflavanone in positive ion mode.

4.29e6

Channel name: High energy : Time 39.5849 +/- 0.0920 minutes







Fig. S65 The fragments of linolic acid in negative ion mode.



Fig. S66 The fragments of palmitic acid in negative ion mode.



Fig. S67 The PCA score plot of 12 chemical markers in CER, CRP and unknown samples.



Fig. S68 Heat map of 12 chemical markers in CER, CRP and unknown samples.