

Analytical Methods

Accepted Manuscript



This is an *Accepted Manuscript*, which has been through the Royal Society of Chemistry peer review process and has been accepted for publication.

Accepted Manuscripts are published online shortly after acceptance, before technical editing, formatting and proof reading. Using this free service, authors can make their results available to the community, in citable form, before we publish the edited article. We will replace this *Accepted Manuscript* with the edited and formatted *Advance Article* as soon as it is available.

You can find more information about *Accepted Manuscripts* in the [Information for Authors](#).

Please note that technical editing may introduce minor changes to the text and/or graphics, which may alter content. The journal's standard [Terms & Conditions](#) and the [Ethical guidelines](#) still apply. In no event shall the Royal Society of Chemistry be held responsible for any errors or omissions in this *Accepted Manuscript* or any consequences arising from the use of any information it contains.

1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
22
23
24
25
26
27
28
29
30
31
32
33
34
35
36
37
38
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60

1 **A practical method for the rapid detection and**
2 **structural characterization of major constituents**
3 **from traditional Chinese medical formulas: Analysis**
4 **Multiple Constituents in Yinchenhao Decoction**

5 Zhiwen Fu^{a, b}, Zhixiong Li^a, Pei Hu^a, Qin Feng^b, Rui Xue^a, Yiyang Hu^{b*}, Chenggang Huang^{a*}

6
7 ^a *Shanghai Institute of Materia Medica, Chinese Academy of Sciences, 501 Haik Rd., Shanghai, 201203,*

8 *P.R. China*

9 ^b *Department of Clinical Pharmacology, Shuguang hospital affiliated to Shanghai University of*

10 *Traditional Chinese Medicine, 528 Zhangheng Rd., Shanghai, 201203, China*

11
12 *Correspondence to:

13 Prof. Chenggang Huang

14 Shanghai Institute of Materia Medica, Chinese Academy of Sciences, 501 Haik Rd., Shanghai, 201203

15 Tel: 86-21-20231963; Fax: 86-21-20231963; E-mail: cghsimm2309@126.com.

16
17 Prof. Yiyang Hu

18 Department of Clinical Pharmacology, Shuguang hospital affiliated to Shanghai University of

19 Traditional Chinese Medicine, E-mail: yhuliver@163.com

1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
22
23
24
25
26
27
28
29
30
31
32
33
34
35
36
37
38
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60

21 **Abstract:**

22 The components of traditional Chinese medical formulas (TCMFs) are usually extremely complex,
23 belonging to hundreds of compound classes with diverse chemical and physical properties. Rapid and
24 effective characterization of TCMFs is imperative in order to provide a first step toward determining
25 which components are responsible for the therapeutic effects of a particular medical plant. A practical
26 method based on Find by Formula (FBF) algorithms is described which provides the rapid detection and
27 structural characterization of major constituents from TCMFs. Results are presented by analysis a
28 classical TCMFs of Yinchenhao Decoction (YCHD). The data of the YCHD extract was acquired using
29 high performance liquid chromatography/quadrupole time-of-flight mass spectrometry
30 (HPLC-Q/TOF-MS/MS) system which features high resolution, mass accuracy, and sensitivity. Based on
31 a prepared simple database file and requisite filter template, the FBF approach was developed and
32 employed to rapidly pick out the peaks of target compounds from mass chromatograms and the particular
33 FBF spectrum with its MS/MS information were listed simultaneously. By this approach, 77 compounds
34 from YCHD were rapidly characterized. The present results demonstrate that this strategy possesses the
35 enormous advantage. This approach is adaptable to analyze complex components from TCMFs and
36 even available to investigate all potential metabolites of one compound with specific molecular
37 formula.

38
39 **Keywords:** Structural characterization; Yinchenhao Decoction; FBF approach; HPLC-Q/TOF-MS/MS

41 Introduction

42 The rapid detection and structural characterization of the major constituents have become an integral part
43 of quality control and research on active principles of traditional Chinese medical formulas (TCMFs) in
44 order to promote the modernization that move toward internationalization [1-3]. TCMFs consisting of
45 multiple herbal medicines have complicated chemical combinations, making the detection and
46 identification a huge challenge, due to significant interference from other ingredients. Although we have
47 acquired sufficient compounds information after decades of phytochemical efforts, we currently need a
48 rapid and effective instrument or method to characterizing of these compounds.

49 Up to now, a large number of detection and technique methods have been reported for characterizing
50 the chemical profiling of TCMFs, including the liquid chromatography-diode array detection
51 (LC-DAD), gas chromatography-mass spectrometry (GC-MS), liquid chromatography-nuclear magnetic
52 resonance (LC-NMR), electrophoresis chromatography-mass spectrometry (EC-MS), and LC-MS [4-7].

53 The LC-MS technique, especially high performance liquid chromatography-electrospray
54 ionization-quadrupole-time of-flight mass spectrometry (HPLC-ESI-Q-TOF-MS), offering rapid and
55 efficient separation and detection methods with accurate mass measurement and tandem mass
56 spectrometry (MS/MS), has been the most widely applied [8, 9]. However, in most previous reports, the
57 inspection of the MS or MS/MS chromatograms was usually performed manually one by one to screen
58 the components from TCMFs. Obviously, the manual action is labor-intensive, low-effective and
59 error-prone. Moreover, it was often difficult to distinguish the relatively small signals from the complex
60 chemical background in full-scan mass chromatograms.

61 Find by Formula (FBF) is a targeted feature finding algorithm using a simple imported database,
62 enabling lower detection levels with higher accuracy. It can find compounds by extracting ions that are

1
2
3
4 63 calculated from a formula in a database of compounds, including user-specified charge carriers (such as
5
6 64 $[M+Na]^+$, $[M+Cl]^-$ and $[M+HCOO]^-$ etc) and multimers (such as $[2M-H]^-$). Furthermore, the
7
8
9 65 corresponding MS/MS spectrums can be shown simultaneously so that we confirm the structural
10
11 66 rationality of compounds. Following this idea, a strategy involving the establishment of a database of
12
13 67 analogous TCMFs constituents was developed in the present study. The resulting simplified data could
14
15
16 68 facilitate the identification of structural analogues in TCMFs.
17

18
19 69 Therefore, the main objective of the present study was to apply the FBF algorithms to the detection and
20
21 70 characterization of major constituents from TCMFs. The effectiveness, rapidity and rationality of the
22
23 71 FBF approach was evaluated by analyzing a typical TCMF, Yinchenhao Decoction (YCHD). As
24
25 72 described in the *Shanghanlun* [10], a classic traditional Chinese medicine publication, YCHD is a
26
27 73 famous prescription which consists of Artemisia capillaries, Fructus Gardeniae, and Rhizoma et Radix
28
29 74 Rhei, and it is used clinically in China and Japan for the treatment of dampness-heat and jaundice [11,
30
31 75 12]. We highlighted the practical and reliable qualities of the FBF approach in the rapid analysis of major
32
33 76 constituents from YCHD. Importantly, we supposed that this methodology could be envisioned to exhibit
34
35 77 a wide application for the identification of complicated components from various complex mixtures and
36
37 78 even screening the possible metabolites of a compound with specific molecular formula *in vivo* or *in*
38
39 79 *vitro*.
40
41
42
43
44
45

46 80 **Experimental**

47 81 **Chemicals and materials**

48
49
50 82 HPLC-MS grade acetonitrile was purchased from Dikma Technologies Inc. (Dikma, CA, USA) and
51
52
53 83 MS grade formic acid was supplied by Sigma-Aldrich (Steinheim, Germany). All other solvents and
54
55
56
57
58
59
60

1
2
3
4 84 chemicals were of analytical grade and were purchased from Sinopharm Chemical Reagent Co. Ltd.
5
6 85 (Shanghai, China). Ultra-pure water was prepared using a Milli-Q system (Millipore, Billerica, MA,
7
8 86 USA).

9
10
11 87 The reference compounds geniposidic acid (no. 111828-201102), caffeic acid (no. 110885-200102),
12
13 88 chlorogenic acid (no. 110753-201314), geniposide (no. 110749-201115), quercetin (100081-200907),
14
15
16 89 rutin (100080-201408), crocin-II (no. 121009-200502), rhein (no. 110757-200206) and emodin (no.
17
18 90 110756-200110) were purchased from the National Institutes for Food and Drug Control (Beijing,
19
20 91 China). The genipin gentiobioside was isolated previously from Fructus Gardeniae in our laboratory.
21
22 92 Their structures were confirmed by HRMS, 1D and 2D NMR comparison of these data with those
23
24 93 reported in the literature [13]. The purity of all reference compounds were determined to be over 98%
25
26 94 by HPLC analysis.

27
28
29
30
31 95 The crude herbs of *Artemisia capillaries*, *Fructus Gardeniae* and *Rhizoma et Radix Rhei* were
32
33 96 purchased from Shanghai Kangqiao Traditional Chinese Medicine Co. Ltd. (Shanghai, China), and
34
35 97 identified by Professor Jingui Shen, Shanghai Institute of Materia Medica, Chinese Academy of
36
37 98 Sciences.

38 39 40 41 42 99 **HPLC conditions**

43
44
45 100 HPLC analysis was performed on a Agilent 1260 Series (Agilent, Santa Clara, CA, USA) LC system,
46
47 101 equipped with a binary pump, an online degasser, an auto plate-sampler, a thermostatically controlled
48
49 102 column compartment and a DAD. Samples were eluted on an Agilent poroshell 120 EC-C₁₈ column
50
51 103 (100 mm × 2.1 mm, 2.7 μm; Agilent, CA, USA). The column and auto-sampler temperature were
52
53 104 maintained at 35 °C and 4 °C, respectively. A mobile phase consisting of 0.1% formic acid in water
54
55 105 (solvent A) and 0.1% formic acid in acetonitrile (solvent B) was applied with the optimized gradient
56
57
58
59
60

1
2
3
4 106 program as follows: 3-15%B (0–12 min), 15–18% B (12–15 min), 18–20% B (15–18 min), 20–40% B
5
6 107 (18–28 min) and 40–90% B (28–45 min). The flow rate was set at 0.35 mL·min⁻¹.
7
8

9
10 108 **Mass spectrometry conditions**

11
12 109 Mass spectrometry was performed on an Agilent 6530 Q-TOF mass spectrometer (Agilent, Santa Clara,
13
14 110 CA, USA) equipped with an electrospray ionization (ESI) interface, and was operated in negative ion
15
16 111 mode with parameters set as follows: capillary voltage, 3500 V; fragmentor, 150 V; skimmer, 65 V;
17
18 112 OCT 1 RF V_{pp}, 750 V; pressure of nebulizer, 35 psi; drying gas temperature, 300 °C; sheath gas
19
20 113 temperature, 350 °C. Nitrogen was used as sheath and drying gas at a flow rate of 8.0 L·min⁻¹ and 11.0
21
22 114 L·min⁻¹, respectively. The collision energy (CE) set values of 12 V and 26 V. An external calibration
23
24 115 solution (Agilent calibration solution A) was continuously sprayed in the ESI source of the Q-TOF
25
26 116 system, employing the ions with *m/z* 112.9855 (TFA anion) and 1033.9881 [HP-0921(TFA adduct)] to
27
28 117 recalibrate the mass axis ensuring mass accuracy and reproducibility throughout the chromatographic
29
30 118 run.
31
32
33
34
35
36
37
38

39 119 **Sample preparation**

40
41 120 *Reference compound solutions*

42
43 121 Known amounts of reference compounds were dissolved in methanol to obtain ten stock solutions
44
45 122 (about 1.0 mg/mL) and stored at 4 °C. All the working solutions were prepared by serial dilution of the
46
47 123 stock solutions with methanol. A mixture containing all ten references compounds was also prepared in
48
49 124 methanol.
50
51

52
53 125 *YCHD samples*

54
55 126 Artemisia capillaries (18 g) was boiled in 1.2 L distilled water until the volume of water reduced to
56
57
58
59
60

1
2
3
4 127 about 600 mL, then 15 g Fructus Gardeniae and 6 g Rhizoma et Radix Rhei were added, and kept it
5
6 128 boiling for concentrating to 300 mL. The extracted solution was centrifuged at 12,000 rpm for 10 min
7
8
9 129 at 4 °C and the supernatant was concentrated under reduced pressure to afford 30 mL residue, and 1 mL
10
11 130 of the residue was filtered through a 0.22 µm microporous membrane before use. An aliquot of 1.0 µL
12
13
14 131 sample was injected into the LC-MS instrument for analysis.

15
16
17
18 132 **Establishment of FBF approach to detect the characteristic components**

19
20 133 Before processing, the following limits were applied to collect the data: the acquisition type set as Auto
21
22 134 MS/MS and the mass range was m/z 50-1200 for both MS and MS/MS while the acquisition rates were 2
23
24
25 135 spectra per second and 1 spectra per second, respectively. Besides, the reference ions of m/z 112.9855
26
27 136 and 1033.9881 as well as the inherent interference ions in mobile phase were excluded in the
28
29
30 137 acquisition lists all the time.

31
32
33 138 The strategy for complex constituents identification was implemented using the following three
34
35 139 steps:

36
37 140 **Step 1.** Establish a specialized *.csv file (See *YCHD.csv* in the supplemental data) as the database of
38
39 141 chemical constituents of Artemisia capillaries, Fructus Gardeniae and Rhizoma et Radix Rhei according
40
41
42 142 to the previous reported literatures. The reported compounds that were isolated from the plants in the
43
44
45 143 formulae were searched for in the following databases: PubMed of the U.S. National Library of
46
47 144 Medicine, SciFinder Scholar of the American Chemical Society and the Chinese National Knowledge
48
49
50 145 Infrastructure (CNKI) of Tsinghua University. The database should contain the molecular formula, the
51
52 146 exact molecular weight, and the compound name. Without doubt, it would be better if we could acquire
53
54
55 147 the retention time.

56
57
58 148 **Step 2.** Open the acquired mass spectrometry data file of YCHD on the software of Agilent MassHunter
59
60

1
2
3
4 149 Qualitative Analysis (Version B 06.01), and import the database of YCHD.csv file in the **Method**
5
6 150 **Editor**. Then set up the important requisite filter templates: a. match tolerance, masses for ± 5 ppm and
7
8 151 retention time for ± 0.1 min (if available); b. charge carriers, $[M-H]^-$ / $[M+Cl]^-$ / $[M-H+HCOOH]^-$ and
9
10 152 activate the aggregates of $[2M-H]^-$; .c. results, automatically extract EIC, FBF spectrum and separate
11
12 153 MS/MS spectrum per CE.
13
14
15
16 154 **Step 3.** Click the **Run** button in the toolbar, and then we can get all the candidates which met the set
17
18 155 conditions in the compound list. Importantly, the MS/MS information of candidates and the processed
19
20 156 extract ion chromatography (p-EIC) could be acquired simultaneously.
21
22
23

24 157 **Results and discussion**

25 26 27 28 158 **Optimization of LC and MS conditions**

29
30 159 Although MS can distinguish overlapping peaks through extracted ion chromatography (EIC), the
31
32 160 isomers have to be identified by their retention behaviors if the co-eluting components have the same
33
34 161 m/z values. Considering the complex constituents of the sample, a Poroshell 120-EC18 column was
35
36 162 selected here to get better separation. Its packing has a solid core of $1.7 \mu\text{m}$ in size with a $0.5 \mu\text{m}$ thick
37
38 163 porous outer layer and a total particle size of $2.7 \mu\text{m}$ providing high efficiency at lower column
39
40 164 pressures [14]. In addition, appropriate concentration of formic acid (0.1%) was introduced into the
41
42 165 mobile phase during HPLC to alleviate the peak tailing and increase the ionization effect. In order to
43
44 166 obtain the most sensitive ionization method for analytes, positive and negative ion modes were
45
46 167 investigated with the same HPLC mobile phase at a flow rate of $0.35 \text{ mL}\cdot\text{min}^{-1}$. As can be seen in
47
48
49
50
51 168 Supplement Information of **Fig. S1**, more representative fragmentation ions information of the structures
52
53
54
55
56 169 were obtained in the negative ion mode.
57
58
59
60

170 On-line characterization of major constituents from YCHD

171 The optimal HPLC-Q/TOF-MS/MS method was applied to analysis the YCHD. The total ion
172 chromatograms (TIC) in negative ESI mode was shown in Fig. 1 (a). After being optimized, FBF
173 approach was used to analysis the full-scan chromatogram of YCHD, the p-EIC was shown in Fig. 1
174 (b). In order to detect the low abundance peak, the p-EIC was scaled to largest for each chromatogram
175 as shown in Fig. 1(c). It was used for clear labeling and it exhibited that the FBF approach could
176 perform rapid and effective analysis of major constituents in YCHD in comparison with manual
177 inspection. As a result, a total of 77 peaks with information on accurate molecular weight were
178 extracted from the full-scan chromatogram and assigned rapidly to the major constituents in YCHD.

179 In the p-EIC of YCHD samples, the peaks of compounds 5, 19, 20, 23, 25, 32, 51, 53, 71 and 73
180 could be unambiguously identified as geniposidic acid, chlorogenic acid, caffeic acid, genipin
181 gentiobioside, geniposide, rutin, quercetin, crocin-II, rhein and emodin by comparing their retention
182 times, molecular weights and fragment ions with standards. The retention times, molecular weights,
183 fragment ions of the 77 compounds plausibly identified are presented in Table 1.

184 The constituents of YCHD could be mainly classified into five types: organic acid, anthraquinones,
185 iridoid glycosides, crocin glycosides and flavonoids. Some representative characteristic fragmentation
186 behaviors of these five categories of compounds were discussed below. These results would be helpful
187 to give chemical information for later research of any combination of these herbs or other herbs which
188 contain these constituents.

189 *Characterization of organic acid*

190 Peaks 8, 10 and 21 in the first-order mass spectrum yielded the same $[M-H]^-$ ions at m/z 353.0878
191 ($C_{16}H_{17}O_9^-$) with the reference compound of 3-O-caffeoylquinic acid (3-CQA), revealing that these

1
2
3
4 192 three compounds were isomers of 3-CQA. However, it was possible to distinguish them according to
5
6 193 the clear differences of their fragmentation ions. As described in the literatures [15, 16], peak 21
7
8 194 afforded the product ion of m/z 173, a unique fragmentation ion to 4-CQA, as its base peak, thus peak
9
10 195 21 could be identified as 4-CQA. In addition, unlike peak 19, the relatively intense product ions at
11
12 196 m/z 179 (51.1% base peak) and 135 (62.5% base peak) of peak 10 could indicate it was a 5-CQA
13
14 197 (Incidentally, as shown in SciFinder Scholar, some serial numbers in literature were incorrect, authors
15
16 198 treated 3-CQA and 5-CQA opposite). The MS/MS information were not possible to reliably distinguish
17
18 199 between peaks 8 and 19. Fortunately, peak 19 of 3-CQA, namely chlorogenic acid, was readily
19
20 200 available from commercial source, so, in practice, the discrimination was straightforward. The related
21
22 201 MS/MS spectra were shown in Fig. 2.

23
24
25
26
27
28
29 202 Similarly, peaks 35, 36, 37 and 42 showed the same $[M-H]^-$ ion at m/z 515.1207 in line with the
30
31 203 $C_{25}H_{24}O_{12}$ formula, indicating that these four compounds were isomers of dicaffeoylquinic acid
32
33 204 (di-CQA). ESI-MS/MS spectra provided considerably more structural information; the MS/MS spectra
34
35 205 data of these four peaks are displayed in Table 1. Based on these fragmentation behaviors and the
36
37 206 literature, peaks 35, 36, 37 and 42 were tentatively identified as 1,4-diCQA, 4,5-diCQA, 3,5-diCQA
38
39 207 and 3,4-diCQA, respectively [15, 16]. First, the characteristic diagnostic product ion at m/z 173,
40
41 208 presented in the MS/MS ion spectra, is indicative of acylation at 4-position. Thus, peak 37 was
42
43 209 tentatively identified as 3,5-diCQA. Besides, 1,4-diCQA was also easily distinguished by its unique
44
45 210 MS/MS fragments including comparatively intense ions at m/z 203 (57.5% base peak) and 299 (21.1%
46
47 211 base peak) supported by less intense fragments at m/z 255 and 317. As for 3,4-diCQA and 4,5-diCQA,
48
49 212 4,5-diCQA had an intense product ion at m/z 335 (25.3% base peak) while it was almost not seen in
50
51 213 MS/MS spectrum of 3,4-diCQA. The related MS/MS spectra were shown in Fig. 3.
52
53
54
55
56
57
58
59
60

214 *Characterization of anthraquinones*

215 Anthraquinones, as the major bioactive constituents of rhubarb, include free anthraquinones,
216 anthraquinone glycosides, and sennosides. The MS/MS spectra of five representative free
217 anthraquinones were shown in Fig. 4. In the MS/MS spectrum of chrysophanol, a product ion at m/z
218 225.0556 was observed as the base peak, which generating from the direct neutral loss of CO (28 Da)
219 from m/z 253. Due to the carbonyl group at C-9 position was prone to produce the intramolecular
220 hydrogen bonding with the α -OH at C-1 and C-8, we believed the CO elimination might occur at C-10
221 firstly [17]. Next, the ion at m/z 225 further lost a $\text{CH}_3\cdot$ (15 Da) and a CO (28 Da), affording the ions at
222 m/z 210.0365 and 182.0393, respectively.

223 Emodin and aloe-emodin, along with rhein and physcion, could be differentiated by their high
224 resolution mass data and MS/MS information. In particular, rhein and physcion, with the distinct
225 precursor ions of m/z 283.0247 and 283.0610, respectively, thus, were facile to identify. In addition,
226 rhein gave the product ions at m/z 239, 211 and 183 in its MS/MS spectrum, obviously indicating the
227 existence of carboxyl group, while physcion afforded the ions at m/z 240 and 212. As for emodin and
228 aloe-emodin, emodin was initiated by the elimination a carbonyl group to produce the ion at m/z 241
229 and followed by loss of a hydroxyl group to give the ion at m/z 225. However, the $[\text{M}-\text{H}]^-$ ion of
230 aloe-emodin, produced the ions at m/z 240.0416, 211.0401, and 183.0437 [17]. Therefore, the distinct
231 difference of MS/MS information between these isomers sufficed their differentiation, and even were
232 valuable for subsequent identification of their corresponding glycosides.

233 *Characterization of iridoid glycosides*

234 Iridoid glycosides are the bioactive components in Fructus Gardeniae which usually give $[\text{M}+\text{Cl}]^-$
235 ion as their precursor ions in negative ESI mode. As described in our group previous study, the

1
2
3
4 236 fragment patterns were shown that the glycosidic bond of iridoid glycosides was prone to break with
5
6 237 neutral losses of 162 Da and always further lost a H₂O. Subsequently, it yielded the characteristic
7
8
9 238 fragment ion at m/z 123.0452 (C₇H₇O₂⁻) by the retro-Diels–Alder (RDA) reaction. When the C-4
10
11 239 position of iridoid glycosides was connected with the carboxyl group, it was easy to lose a molecule of
12
13 240 carbon dioxide (CO₂) with neutral losses of 44 Da following deglycosylation, while the C-4 position
14
15 241 was connected with the carbomethoxy, it always produced the characteristic fragment ion at m/z
16
17 242 101.0244 (C₄H₅O₃⁻) via the bond cleavage between C₁-O₂ and C₄-C₅. Furthermore, if any groups were
18
19 243 substituted at chains of glucose, it was prone to obtain [M_s+glucosyl-H-H₂O]⁻ and [M_s-H]⁻ of the
20
21 244 substituent [18].
22
23
24

25
26 245 For instance, as shown in Fig. 5, peak 44 at the retention time of 20.784 min (exact molecular weight
27
28 246 696.2265) gave the precursor ion of [M+Cl]⁻ at m/z 731.1947 and afforded the typical product ions at
29
30 247 m/z 469.1347 ([coumaroyl+gentiobiosyl-H-H₂O]⁻), 225.0769 (C₁₁H₁₃O₅⁻), 163.0397
31
32 248 ([coumaroyl-H]⁻), 145.0294 ([coumaroyl-H-H₂O]⁻), 123.0451 (C₇H₇O₂⁻) and 101.0246 (C₄H₅O₃⁻),
33
34 249 which was confirmed as 6''-O-*trans*-coumaroylgenipin gentiobioside [19].
35
36
37
38

39 250 *Characterization of crocin glycosides*

40
41 251 Crocin glycosides was the diester formed from the disaccharide gentiobiose and the dicarboxylic acid
42
43 252 crocetin. Therefore, they showed the product ions of [M-162*n]⁻, [crocetin-H]⁻ and
44
45 253 [crocetin-H-CO₂]⁻ in the MS/MS spectra according to the successive losses of glycosides and
46
47 254 carboxyl group. Besides, the crocin glycosides in YCHD seemed to have a *cis-trans* isomer of
48
49 255 13-position. They exhibited the identical MS/MS data except for distinct retention time. Based on the
50
51 256 conclusions summarized, it would be possible to consider that the 13-*cis* isomers took a longer
52
53 257 retention time on the reverse phase packing. For crocin-I (retention time of 22.892 min) and
54
55
56
57
58
59
60

1
2
3
4 258 13-*cis*-crocin I (retention time of 28.035 min), the $[M+Cl]^-$ ion at m/z 1011.3479 with the formula
5
6 259 $[(C_{38}H_{54}O_{19})+Cl]^-$ was selected as the precursor ion in the MS/MS experiment to give fragmentation
7
8
9 260 information (Fig. 6). The ions at m/z 651.2650 ($C_{32}H_{43}O_{14}^-$) and 327.1596 ($C_{20}H_{23}O_4^-$) corresponded to
10
11 261 the successive losses of 324 Da (disaccharide) from the deprotonated molecule. In addition, for the ion
12
13 262 at m/z 283.1696 ($C_{19}H_{23}O_2^-$), the loss of 44Da could be assigned as CO_2 originating from m/z 327 [20].
14
15

16 263 *Characterization of flavonoids*

17
18
19 264 Flavonoids, The mass spectrum of quercetin (peak 51), which was detected at retention time 23.566
20
21 265 min, gave an $[M-H]^-$ ion at m/z 301.0347 ($C_{15}H_9O_7^-$). As shown in Table 1, its fragmentation patterns
22
23 266 were quite clear as a flavonoid. Characteristic product ions at m/z 151.0028 ($C_7H_3O_4^-$) and 107.0135
24
25 267 ($C_6H_3O_2^-$), originating from two pathways of the RDA reaction, could be found in MS/MS spectra. In
26
27 268 addition, ions at m/z 83.0144 ($C_4H_3O_2^-$), 65.0039 (C_4HO^-) and 178.9974 ($C_8H_3O_5^-$) were also
28
29 269 observed.
30
31
32

33
34 270 Peaks 31 and 34 both exhibited $[M-H]^-$ ions at m/z 463.0876 ($C_{21}H_{19}O_{12}^-$). Moreover, they had the
35
36 271 similar fragment ions of m/z 301 and 151 in MS/MS spectra, but different retention times even without
37
38 272 regular retention behavior. They could be identified as hyperoside or isoquercitrin, but not precisely
39
40 273 identified by the method above, and need isolation and identification by NMR in the future.
41
42
43
44

45 274 **The discussion of FBF algorithms**

46
47 275 Although the limitations of mass spectrum for identifying exact structures, it is a convenient way to
48
49 276 rapidly analyze mass spectrum character to speculate tentative structures. And it is very imperative to
50
51 277 apply to TCMFs without isolation and purification each compound. The HPLC-Q/TOF-MS/MS
52
53 278 method combine with FBF algorithms requires only a simple *csv* file and the rapid and effective
54
55 279 characterization will be performed facilely. However, the deficiency of FBF algorithms should be noted,
56
57
58
59
60

1
2
3
4 280 we can just characterize the known compounds reported previously using FBF algorithms. As a famous
5
6 281 traditional Chinese medicinal materials, the previous phytochemical isolations have been accomplished
7
8
9 282 maturely and the related compounds information are also acquired sufficiently, hence, the compounds in
10
11 283 *csv* file will be counted comprehensively.

12
13
14 284 More importantly, based on the principles of FBF approach, we can not only apply it conveniently to
15
16 285 analyze the major constituents in TCMFs like YCHD in this work, but also utilize it as important tools to
17
18 286 investigate all potential metabolites of one compound with specific molecular formula. We can list all
19
20
21 287 possible forms of metabolites, their molecular formula and exact molecular weight in the *csv* file. Once
22
23
24 288 the database file is established, it will be a major breakthrough to avoid the matrix interference when we
25
26 289 performed a qualitative analysis metabolic process in biological sample.

29 **Conclusions**

30
31
32 291 A primary challenge in herbal analysis was to enable global qualitative performance rapidly and
33
34 292 accurately. In the present study, a practical method based on the FBF algorithms was developed for
35
36
37 293 rapid detection and structural characterization of major constituents from YCHD. As a result, a total of
38
39
40 294 77 compounds were structural characterized. Compared with the conventional manual inspection and
41
42 295 fragmentation-based method, the FBF approach enabled the original data to be analyzed much faster
43
44 296 and more accurately by reducing the potential interferences of matrix ions. More importantly,
45
46
47 297 according to the principle of FBF, this method can also be used for the rapid analysis the possible
48
49
50 298 metabolites of a compound with specific molecular formula. Meanwhile, it should be noted that the
51
52
53 299 major limitation of this study was that only the known chemical composition was used to validate the
54
55 300 characterizing results.

1
2
3
4 301 **Acknowledgement**
5

6 302 This work was supported by the Program of National Natural Science Foundation of China (NO.
7
8 303 81173404, 81374031) and the Program of Science and Technology Commission of Shanghai
9
10 304 Municipality (NO. 13401900303).
11

12
13
14 305 **References**
15

- 16
17 306 [1] M. Holcapek, R. Jirasko, M. Lisa, *J. Chromatogr. A*, 2012, **1259**, 3-15.
18
19 307 [2] J.L. Zhou, L.W. Qi, P. Li, *J. Chromatogr. A*, 2009, **1216**, 7582-7594.
20
21 308 [3] M. Yang, J.H. Sun, Z.Q. Lu, G.T. Chen, S.H. Guan, X. Liu, B.H. Jiang, M. Ye, D.A. Guo, *J.*
22
23 309 *Chromatogr. A*, 2009, **1216**, 2045-2062.
24
25 310 [4] S.L. Li, S.F. Lai, J.Z. Song, C.F. Qiao, X. Liu, Y. Zhou, H. Cai, B.C. Cai, H.X. Xu, *J. Pharm.*
26
27 311 *Biomed. Anal.*, 2010, **53**, 946-957.
28
29 312 [5] Y.Q. Wang, Z.M. Guo, Y. Jin, X.L. Zhang, L. Wang, X.Y. Xue, X.M. Liang, *J. Pharm. Biomed.*
30
31 313 *Anal.*, 2010, **51**, 606-616.
32
33 314 [6] G.L. Yan, H. Sun, W.J. Sun, L. Zhao, X.C. Meng, X.J. Wang, *J. Pharm. Biomed. Anal.*, 2010, **53**,
34
35 315 421-431.
36
37 316 [7] T. Xie, Y. Liang, H. Hao, J. A, L. Xie, P. Gong, C. Dai, L. Liu, A. Kang, X. Zheng, G. Wang, *J.*
38
39 317 *Chromatogr. A*, 2012, **1227**, 234-244.
40
41 318 [8] X.L. Zhang, L.F. Liu, L.Y. Zhu, Y.J. Bai, Q. Mao, S.L. Li, S.L. Chen, H.X. Xu, *J. Pharm. Biomed.*
42
43 319 *Anal.*, 2014, **88**, 391-400.
44
45 320 [9] I.M. Abu-Reidah, D. Arráez-Román, R. Quirantes-Piné, S. Fernández-Arroyo, A. Segura-Carretero,
46
47 321 A. Fernández-Gutiérrez, *Food Research International*, 2012, **46**, 108-117.
48
49 322 [10] Z.J. Zhang, *Shanhanlun*, 2013, ch.3, 69.
50
51
52
53
54
55
56
57
58
59
60

- 1
2
3
4 323 [11] T.Y. Lee, H.H. Chang, J.H. Chen, M.L. Hsueh, J.J. Kuo, *J. Ethnopharmacol.*, 2007, **109**, 318-324.
5
6 324 [12] T.Y. Lee, H.H. Chang, J.J. Kuo, J.J. Shen, *International Journal of Molecular Medicine*, 2009, **23**,
7
8 325 477-484.
9
10
11 326 [13] H. Inouye, Y. Takeda, H. Nishimura, *Phytochemistry*, 1974, **13**, 2219-2224.
12
13
14 327 [14] F. Gritti, G. Guiochon, *J. Chromatogr. A*, 2011, **1218**, 907-921.
15
16 328 [15] M.N. Clifford, K.L. Johnston, S. Knight, N. Kuhnert, *J. Agric. Food Chem.*, 2003, **51**, 2900-2911.
17
18 329 [16] M.N. Clifford, S. Knight, B. Surucu, N. Kuhnert, *J. Agric. Food Chem.*, 2006, **54**, 1957-1969.
19
20
21 330 [17] M. Ye, J. Han, H.B. Chen, J.H. Zheng, D.A. Guo, *J. Am. Soc. Mass. Spectrom.*, 2007, **18**, 82-91.
22
23
24 331 [18] Z.W. Fu, R. Xue, Z.X. Li, M.C. Chen, Z.L. Sun, Y.Y. Hu, C.G. Huang, *Biomed. Chromatogr.*, 2014,
25
26 332 **28**, 1795-1807.
27
28
29 333 [19] Z.W. Fu, Y. Ling, Z.X. Li, M.C. Chen, Z.L. Sun, C.G. Huang, *Biomed. Chromatogr.*, 2014, **28**,
30
31 334 475-485.
32
33
34 335 [20] X. Zhao, Z.M. Long, J.N. Dai, K.S. Bi, X.H. Chen, *Rapid Commun. Mass Spectrom.*, 2012, **26**,
35
36 336 2443-2453.
37
38
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60

337

338 **Figure captions**

339 **Fig. 1.** The extract ion chromatogram (EIC) of YCHD in negative ion mode (a); the processed extract
340 ion chromatogram (p-EIC) of YCHD in negative ion mode (b); the scaled to largest for each
341 chromatogram of p-EIC (c).

342 **Fig. 2.** The MS/MS spectra of the $[M-H]^-$ ions at m/z 353.0878 for 1-CQA (a); 5-CQA (b); 3-CQA (c)
343 and 4-CQA (d).

344 **Fig. 3.** The MS/MS spectra of the $[M-H]^-$ ions at m/z 515.1207 for 1,4-dicaffeoylquinic acid (a);
345 4,5-dicaffeoylquinic acid (b); 3,5-dicaffeoylquinic acid (c) and 3,4-dicaffeoylquinic acid (d).

346 **Fig. 4.** The MS/MS spectra of the $[M-H]^-$ ions at m/z 253.0500 for chrysophanol (a); m/z 283.0610 for
347 physcion (b); m/z 283.0247 for rhein (c); m/z 269.0450 for emodin (d) and m/z 269.0450 for
348 aloe-emodin (e).

349 **Fig. 5.** The FBF spectrum for 6''-O-*trans*-coumaroyl genipin gentiobioside (a) and its MS/MS spectrum
350 of the $[M+Cl]^-$ ion at m/z 731.1947 (b).

351 **Fig. 6.** The EIC of m/z 1011 in negative ion mode (a), the FBF spectrum for crocin-I (b); the MS/MS
352 spectra of the $[M+Cl]^-$ ion at m/z 1011.3468 for crocin-I (c) and m/z 1011.3475 for 13-*cis*-crocin-I (d).

Table 1. Chemical constituents identified in an extract of YCHD by HPLC-Q/TOF-MS/MS in negative ion mode

Peak No.	t_R (min)	Molecular formula	Theoretical [M-H] ⁻	Measured [M-H] ⁻	Error ^a (ppm)	Characteristic MS/MS fragmentation ions (% base peak) (m/z)	Proposed compounds
1	0.803	C ₁₂ H ₂₂ O ₁₁	377.0856 ^b	377.0857	0.27	341 (100), 215 (11.1), 179 (20.5), 119 (18.1), 89 (49.6), 59 (34.1)	gentiobiose
2	0.972	C ₅ H ₇ NO ₃	128.0353	128.0364	8.59	101.0257 (17.8), 88.0426 (54.1), 85.0308 (100)	pyroglutamic acid
3	1.309	C ₇ H ₆ O ₅	169.0137	169.0136	-0.59	125.0238 (100), 107.0133 (2.2), 79.0193 (12.2)	gallic acid
4	1.646	C ₁₉ H ₂₆ O ₁₅	493.1199	493.1189	-2.03	313.0564 (63.2), 271.0455 (66.2), 211.0240 (21.9), 169.0137 (100), 125.0237 (30.7)	6-O-Galloylsucrose
5	1.815	C ₁₆ H ₂₂ O ₁₀	373.1140	373.1146	1.61	211.0612 (27.4), 193.0502 (5.8), 167.0712 (13.6), 149.0604 (78.9), 123.0453 (100)	geniposidic acid
6	2.068	C ₁₃ H ₁₆ O ₉	315.0722	315.0722	0.00	153.0190 (32.4), 152.0115 (47.9), 109.0291 (34.4), 108.0227 (100)	protocatechuic acid-3-glucoside
7	2.237	C ₁₆ H ₂₆ O ₉	397.1271 ^b	397.1265	-1.51	361.1496 (100), 317.1589 (1.9), 181.0862 (31.7), 137.0964 (11.2), 89.0245 (12.4), 59.0144 (20.6)	villosolside
8	2.405	C ₁₆ H ₁₈ O ₉	353.0878	353.0878	0.00	191.0538 (100), 85.0284 (7.9)	1-CQA
9	2.405	C ₁₆ H ₂₄ O ₁₁	391.1246	391.1250	1.02	229.0713 (24.5), 211.0611 (17.1), 185.0823 (75.2), 167.0712 (93.1), 149.0616 (100), 141.0549 (42.5), 123.0449 (57.9)	shanzhiside
10	3.401	C ₁₆ H ₁₈ O ₉	353.0878	353.0878	0.00	191.0538 (100), 179.0326 (51.1), 135.0429 (62.5)	neochlorogenic acid (5-COA)
11	3.67	C ₁₇ H ₂₄ O ₁₁	439.1013 ^b	439.1007	-1.37	403.1247 (8.9), 241.0719 (100), 223.0610 (12.3), 139.0397 (43.7), 101.0239 (31.8)	scandoside methyl ester
12	3.754	C ₁₆ H ₂₆ O ₈	381.1322 ^b	381.1319	-0.79	345.1548 (50.6), 179.0547 (15.9), 165.0913 (100), 121.1017 (12.9), 119.0345 (19.0), 89.0245 (56.9)	picrocrocinic acid
13	3.754	C ₁₆ H ₂₂ O ₁₁	389.1089	389.1088	-0.26	227.0566 (24.3), 209.0450 (48.9), 183.0667 (51.1),	deacetylasperulosidic acid

						165.0551 (100), 139.0395 (53.9),	
14	4.344	C ₁₆ H ₂₄ O ₉	395.1114 ^b	395.1111	-0.76	359.1346 (95.4), 197.0813 (100), 125.0612 (62.4), 107.0492 (28.9)	ixoroside
15	4.85	C ₁₇ H ₂₆ O ₁₁	405.1402	405.1395	-1.73	225.0766 (29.3), 179.0566 (57.6), 141.0555 (82.2), 123.0447 (48.5), 119.0349 (30.4), 101.0239 (100)	shanzhiside methyl ester
16	4.934	C ₁₇ H ₂₄ O ₁₁	439.1013 ^b	439.1015	0.46	403.1249 (30.5), 241.0714 (100), 223.0615 (14.1), 139.0399 (87.3), 121.0299 (46.8), 101.0242 (24.1)	gardenoside
17	5.778	C ₁₇ H ₂₄ O ₁₁	439.1013 ^b	439.1005	-1.82	403.1246 (12.6), 241.0710 (100), 223.0617 (40.8), 139.0394 (71.9), 101.0248 (52.6)	deacetylasperulosidic acid methyl ester
18	6.368	C ₁₅ H ₁₄ O ₆	289.0718	289.0710	-2.77	245.0816 (45.1), 203.0706 (35.3), 137.0242 (30.25), 123.0450 (78.6), 109.0294 (100),	catechin
19	6.436	C ₁₆ H ₁₈ O ₉	353.0878	353.0878	0.00	191.0547 (100), 85.0294 (6.1)	chlorogenic acid (3-CQA)
20	6.874	C ₉ H ₈ O ₄	179.0350	179.0350	0.00	135.0437 (100), 89.0395 (5.7)	caffeic acid
21	7.632	C ₁₆ H ₁₈ O ₉	353.0878	353.0878	0.00	191.0535 (60.7), 179.0324 (70.3), 173.0426 (100), 135.0431 (80.9), 93.0335 (27.6)	cryptochlorogenic acid (4-CQA)
22	7.885	C ₁₆ H ₂₄ O ₁₀	375.1297	375.1303	1.60	213.0764 (39.1), 195.0660 (38.4), 169.0877 (47.9), 151.0770 (95.2), 125.0605 (80.7), 107.0510 (100)	mussaenosidic acid
23	8.307	C ₂₃ H ₃₄ O ₁₅	585.1592 ^b	585.1584	-1.37	549.1815 (12.3), 225.0769 (69.3), 207.0657 (13.4), 123.0449 (100), 101.0243 (64.9)	genipin gentiobioside
24	8.56	C ₈ H ₈ O ₂	135.0452	135.0453	0.74	120.0244 (5.5), 108.0206 (10.8), 92.0279 (100)	4'-hydroxyacetophenone
25	9.487	C ₁₇ H ₂₄ O ₁₀	423.1063 ^b	423.1070	1.65	387.1282 (5.5), 225.0762 (100), 207.0653 (13.5), 123.0447 (85.3), 101.0243 (74.5)	geniposide
26	10.161	C ₁₆ H ₂₆ O ₈	381.1322 ^b	381.1309	-3.41	345.1464 (11.9), 183.1020 (86.1), 165.0908 (100), 109.0290 (46.7), 89.0231 (58.9)	jasminoside B
27	11.173	C ₂₇ H ₃₀ O ₁₅	593.1512	593.1520	1.35	549.1610 (52.4), 343.1053 (11.6), 265.0722 (19.1), 207.0295 (100), 163.0397 (21.6)	rheinoside D/rheinoside C

28	11.763	C ₂₇ H ₃₀ O ₁₅	593.1512	593.1522	1.69	549.1606 (45.2), 343.1052 (13.9), 265.0719 (14.8), 207.0299 (100), 163.0401 (12.7)	rheinoside C/rheinoside D
29	12.606	C ₂₂ H ₂₂ O ₁₂	477.1038	477.1020	-3.77	313.0554 (40.3), 169.0133 (100), 163.0388 (29.33), 125.0237 (13.77)	<i>p</i> -coumaroyl-O-galloyl-glucose
30	13.454	C ₂₅ H ₂₈ O ₁₂	519.1508	519.1511	0.58	355.1029 (8.9), 307.0824 (2.3), 211.0606 (18.2), 193.0504 (22.4), 167.0716 (27.8), 163.0397 (84.3), 149.0601(33.2), 145.0299 (100), 123.0445 (52.2)	6'- <i>O</i> - <i>trans</i> -coumaroyl geniposidic acid
31	15.562	C ₂₁ H ₂₀ O ₁₂	463.0882	463.0876	-1.30	301.0330 (64.82), 300.0265 (100), 283.0252 (1.3), 271.0240 (37.85), 255.0293 (16.3), 151.0022 (8.6)	hyperoside/isoquercitrin
32	15.726	C ₂₇ H ₃₀ O ₁₆	609.1461	609.1457	-0.66	301.0357 (95.9), 300.0274 (100), 151.0028 (5.3)	rutin
33	15.894	C ₂₁ H ₁₈ O ₁₁	445.0776	445.0773	-0.67	283.0238 (80.8), 239.0347 (100), 211.0413 (18.7)	rhein monoglucoside
34	16.068	C ₂₁ H ₂₀ O ₁₂	463.0882	463.0874	-1.73	301.0331 (49.3), 300.0271 (100), 271.0244 (36.2), 255.0297 (17.2), 151.0034 (8.5)	isoquercitrin/hyperoside
35	17.417	C ₂₅ H ₂₄ O ₁₂	515.1195	515.1207	2.33	353.0878 (100), 317.0665 (18.7), 299.0552 (21.1), 255.0658 (15.6), 203.0349 (57.5), 191.0558 (12.9), 179.0354 (49.5), 173.0456 (57.6), 135.0467 (7.4)	1,4-dicaffeoylquinic acid
36	17.923	C ₂₅ H ₂₄ O ₁₂	515.1195	515.1207	2.33	353.0872 (100), 335.0766 (25.3), 191.0561 (47.6), 179.0354 (58.3), 173.0460 (71.9), 135.0452 (10.3)	4,5-dicaffeoylquinic acid
37	18.26	C ₂₅ H ₂₄ O ₁₂	515.1195	515.1207	2.33	353.0874 (100), 191.0551 (78.8), 179.0349 (29.3), 135.0445 (8.5)	3,5-dicaffeoylquinic acid
38	18.424	C ₂₇ H ₃₄ O ₁₃	601.1693 ^b	601.1695	0.33	565.1893 (14.6), 385.1127 (24.8), 325.0924 (100), 295.0824 (54.1), 265.0727 (79.7), 223.0604 (55.7), 205.0507 (25.3)	11-(6- <i>O</i> - <i>trans</i> -sinapoyl glucopyranosyl) gardendiol
39	18.424	C ₂₂ H ₃₆ O ₁₂	527.1901 ^b	527.1897	-0.76	491.2130 (13.7), 167.1064 (100), 89.0238 (23.6)	jasminoside H/jasminoside I
40	18.929	C ₂₂ H ₂₂ O ₁₂	477.1038	477.1037	-0.21	357.0598 (14.1), 314.0421 (100), 285.0438 (42.9), 271.0237 (36.9), 243.0325 (50.3)	isorhamnetin-3-glucoside

41	19.908	C ₂₂ H ₃₆ O ₁₂	527.1901 ^b	527.1899	-0.38	491.2130 (25.5), 167.1066 (100), 89.0234 (31.9)	jasminoside I/jasminoside H
42	20.031	C ₂₅ H ₂₄ O ₁₂	515.1195	515.1207	2.33	353.0880 (100), 191.0569 (11.43), 179.0354 (34.4), 173.0458 (50.8), 135.0455 (5.1)	3,4-dicaffeoylquinic acid
43	20.278	C ₂₇ H ₃₄ O ₁₃	601.1693 ^b	601.1697	0.67	565.1907 (28.9), 385.1119 (25.1), 325.0927 (100), 295.0815 (62.2), 265.0711 (91.3), 223.0609 (64.4), 205.0504 (25.2)	10-(6-O- <i>trans</i> -sinapoyl glucopyranosyl) gardendiol
44	20.784	C ₃₂ H ₄₀ O ₁₇	731.1960 ^b	731.1947	-1.78	695.2186 (100), 469.1347 (10.1), 367.1018 (3.4), 225.0769 (12.7), 163.0397 (9.6), 145.0294 (9.4), 123.0451 (20.6), 101.0246 (13.4)	6"-O- <i>trans</i> -coumaroyl genipin gentiobioside
45	21.121	C ₂₂ H ₂₂ O ₁₁	461.1089	461.1082	-1.52	313.0552 (100), 169.0129 (41.4), 147.0437 (52.4), 125.0234 (8.1), 103.0544 (12.4)	2-cinnamoyl-1-galloylglucos e
46	21.374	C ₃₄ H ₄₄ O ₁₉	791.2171	791.2168	-0.38	755.2393 (100), 529.1560 (9.6), 427.1242 (1.6), 225.0763 (16.6), 223.0607 (11.6), 207.0657 (3.1), 205.0498 (6.3), 123.0449 (21.4), 101.0252 (13.4)	6"-O- <i>trans</i> -sinapoyl genipin gentiobioside
47	22.217	C ₃₁ H ₃₂ O ₁₆	659.1618	659.1618	0.00	497.1291 (100), 353.0863 (40.2), 335.0829 (69.3), 273.0965 (10.5), 233.0653 (14.9), 191.0548 (82.9), 161.0440 (62.8)	3,5-di-O-caffeoyl-4-O-(3-hy droxy-3-methyl)-glutaroylqui nic acid
48	22.639	C ₂₇ H ₂₈ O ₁₃	559.1457	559.1453	-0.72	397.1131 (100), 223.0602 (11.2), 173.0446 (71.6)	5-O-caffeoyl-4-O-sinapoylqu inic acid
49	22.892	C ₄₄ H ₆₄ O ₂₄	1011.3482 ^b	1011.3479	-0.30	651.2650 (100), 327.1596 (39.9), 283.1696 (8.6), 179.0556(3.5)	crocin-I
50	23.061	C ₂₈ H ₃₄ O ₁₄	629.1643 ^b	629.1640	-0.48	593.1878 (100), 367.1036 (9.7), 225.0766 (23.8), 223.0609 (27.5), 207.0659 (10.3), 205.0507 (64.3), 123.0451 (46.8), 101.0249 (31.7)	6'-O- <i>trans</i> -sinapoyl geniposide
51	23.566	C ₁₅ H ₁₀ O ₇	301.0354	301.0347	-2.33	178.9974 (40.2), 151.0028 (100), 121.0293 (31.9), 107.0135 (35.7), 65.0039 (46.7)	quercetin

52	23.735	C ₂₂ H ₂₂ O ₁₁	461.1089	461.1091	0.43	313.0561 (100), 169.0138 (40.8), 147.0446 (53.2), 125.0243 (8.1), 103.0555 (13.6)	6-cinnamoyl-1-galloylglucose
53	24.325	C ₃₈ H ₅₄ O ₁₉	849.2959 ^b	849.2951	-0.94	651.2651 (100), 489.2115 (25.5), 327.1587 (74.5), 283.1711 (23.7)	crocin-II
54	24.41	C ₂₆ H ₃₀ O ₁₂	533.1664	533.1659	-0.94	325.0926 (56.2), 307.0819 (37.7), 225.0767 (81.9), 207.0669 (27.3), 163.0406 (100), 145.0287 (33.6), 123.0455 (64.1), 101.0241 (58.7)	6'-O-trans-coumaroyl geniposide
55	24.41	C ₂₉ H ₂₆ O ₁₅	613.1199	613.1209	1.63	465.0664 (100), 313.0564 (33.8), 271.0458 (28.2), 169.0142 (41.6)	cinnamoyl-di-O-galloyl-glucose
56	24.747	C ₁₆ H ₁₀ O ₇	313.0354	313.0354	0.00	269 (100), 241 (10.1), 225 (11.2)	laccaic acid D
57	25.084	C ₂₁ H ₂₀ O ₁₀	431.0989	431.0987	-0.46	269 (100), 240 (54.1), 223 (17.4)	aloe-emodin monoglucoside
58	25.506	C ₃₂ H ₄₀ O ₁₆	715.2010 ^b	715.2002	-1.12	679 (100), 531 (20.9), 453 (11.2), 225 (82.2), 207 (22.6), 147 (28.7), 123 (35.1), 101 (16.5)	6"-O-trans-cinnamoyl genipin gentiobioside
59	26.433	C ₂₄ H ₂₂ O ₁₃	517.0988	517.0978	-1.93	473 (20.2), 269 (100)	emodin-8-O-(6'-O-malonyl)-glucoside
60	26.601	C ₄₈ H ₆₄ O ₂₃	1007.3766	1007.3761	-0.50	683 (100), 327 (6.9), 283 (6.4)	crocin-I-acid
61	27.023	C ₁₅ H ₁₀ O ₄	253.0506	253.0509	1.19	225 (100), 210 (17.1), 195 (16.8), 182 (28.8), 154 (8.1)	chrysophanol
62	27.192	C ₂₂ H ₂₂ O ₁₀	481.0907 ^b	481.0911	0.83	283 (100), 240 (30.3)	physcion monoglucoside
63	27.445	C ₂₁ H ₂₀ O ₁₀	431.0984	431.0989	1.16	269 (100), 225 (12.5)	emodin monoglucoside
64	27.866	C ₃₂ H ₄₄ O ₁₄	687.2425 ^b	687.2415	-1.46	651 (13.9), 327 (100), 283 (70.6), 239 (20.3)	crocin-III
65	28.035	C ₄₄ H ₆₄ O ₂₄	1011.3482 ^b	1011.3479	-0.30	651 (100), 327 (39.4), 283 (8.3), 179 (3.1)	13-cis-crocin-I
66	28.288	C ₁₆ H ₁₀ O ₆	297.0405	297.0408	1.01	253 (100), 225 (58.16), 182 (15.6)	3-(1,8-Dihydroxy-6-methyl)anthraquinonecarboxylic acid
67	28.541	C ₁₉ H ₂₄ O ₄	315.1607	315.1599	-2.54	297 (43.2), 271 (33.7), 241 (100), 186 (15.6)	Capillartemisin A
68	28.878	C ₁₆ H ₁₂ O ₅	283.0617	283.0610	-2.47	240 (100), 212 (12.5), 184 (7.3)	physcion

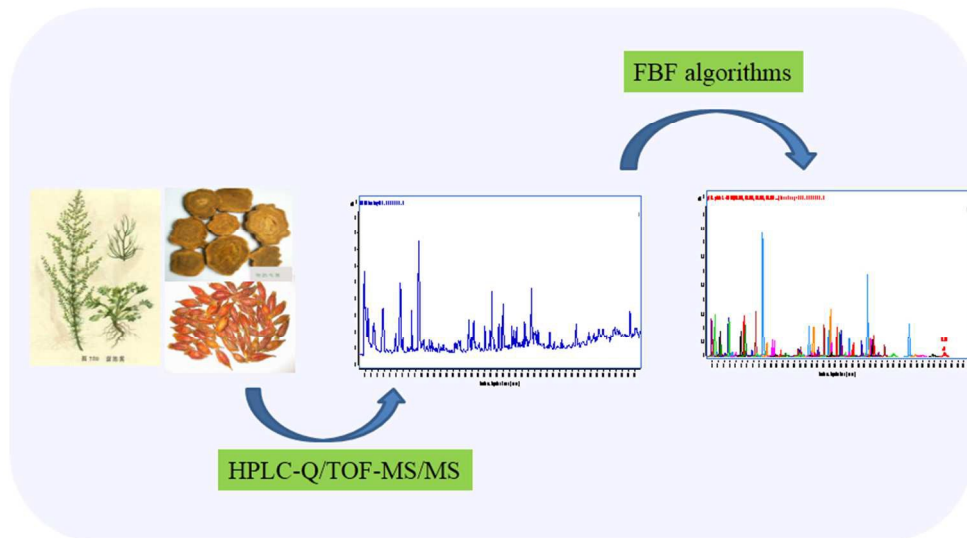
69	29.215	C ₁₅ H ₁₀ O ₅	269.0455	269.0450	-1.86	240 (100), 223 (20.2), 211 (78.1), 195 (14.5), 183 (61.7), 167 (26.4)	aloe-emodin
70	29.384	C ₃₈ H ₅₄ O ₁₉	849.2953 ^b	849.2954	0.12	651 (85.5), 489 (29.7), 327 (100), 283 (19.6)	13- <i>cis</i> -crocin-II
71	30.395	C ₁₅ H ₈ O ₆	283.0248	283.0247	-0.35	239 (100), 211 (44.1), 183 (91.5)	rhein
72	31.829	C ₃₂ H ₄₄ O ₁₄	687.2425 ^b	687.2422	-0.44	651 (12.3), 489 (2.9), 327 (100), 283 (46.8), 239 (13.4)	13- <i>cis</i> -crocin-III
73	34.511	C ₁₅ H ₁₀ O ₅	269.0455	269.0450	-1.86	241 (34.6), 225 (100), 210 (20.7), 195 (36.5), 105 (21.9)	emodin
74	35.628	C ₃₀ H ₂₂ O ₈	509.1242	509.1247	0.98	254 (100)	palmidin A
75	36.808	C ₃₀ H ₂₀ O ₉	523.1035	523.1037	0.38	485 (4.4), 254 (100), 224 (9.84)	rheidin
76	38.916	C ₃₀ H ₂₂ O ₇	493.1293	493.1290	-0.61	476 (20.1), 462 (52.5), 254 (100), 238 (88.9), 152 (24.6)	palmidin B
77	40.602	C ₃₀ H ₂₂ O ₇	493.1293	493.1294	0.20	254 (100), 152 (5.68)	palmidin C

^a Differences between the measured and theoretical values, ppm.

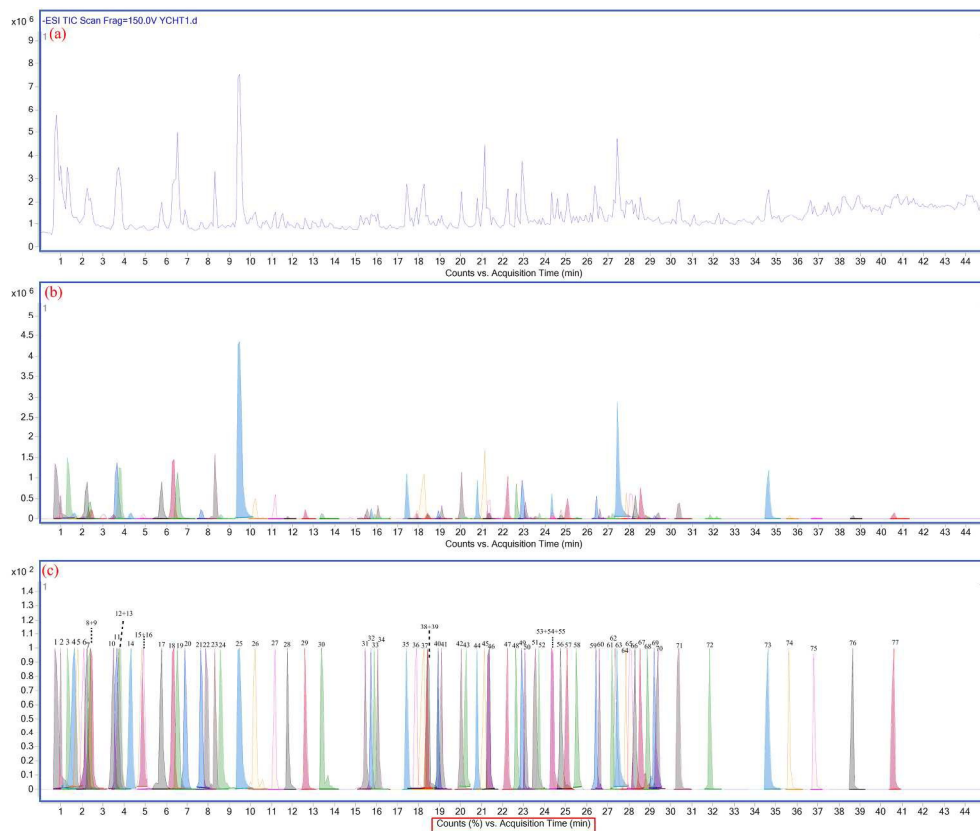
^d gave [M+Cl]⁻ as precursor ions

* identified with the reference compounds

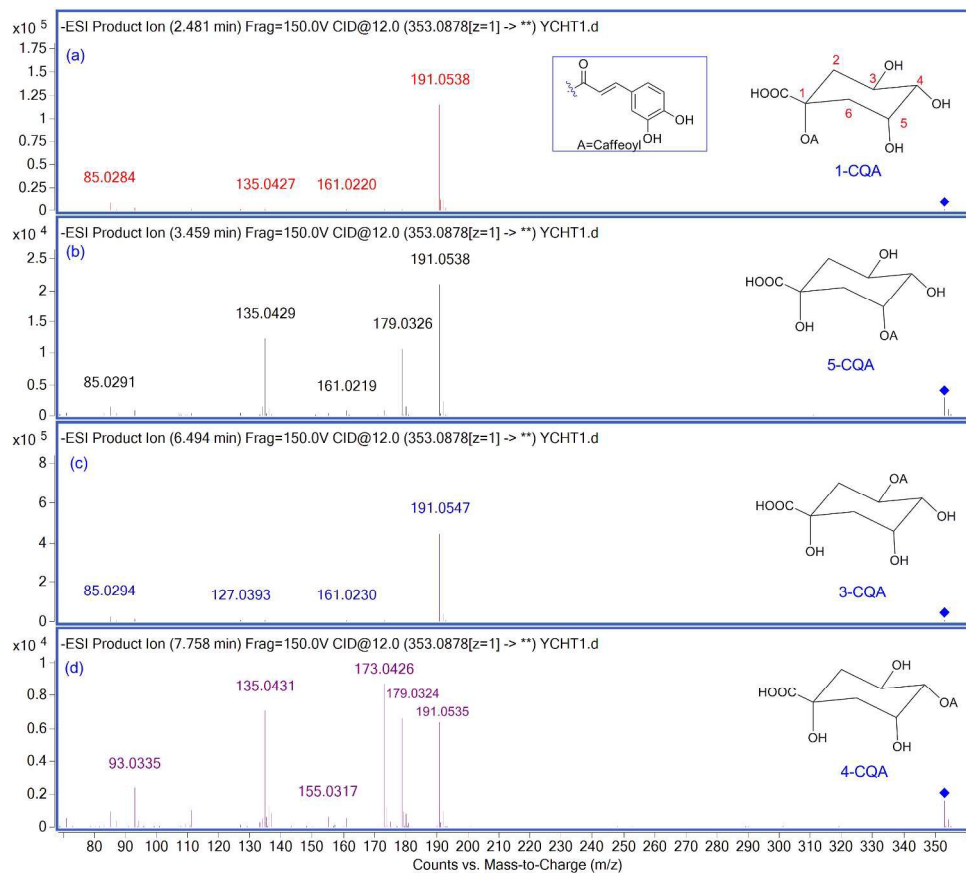
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
22
23
24
25
26
27
28
29
30
31
32
33
34
35
36
37
38
39
40
41
42
43
44
45
46
47
48
49
50
51
52
53
54
55
56
57
58
59
60



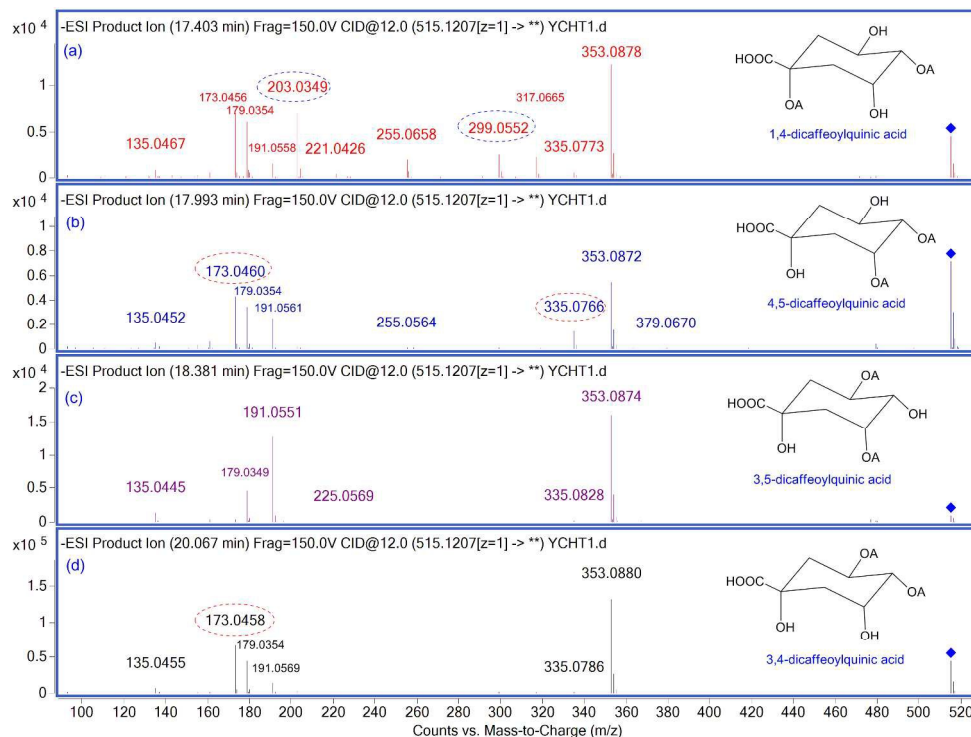
254x190mm (96 x 96 DPI)



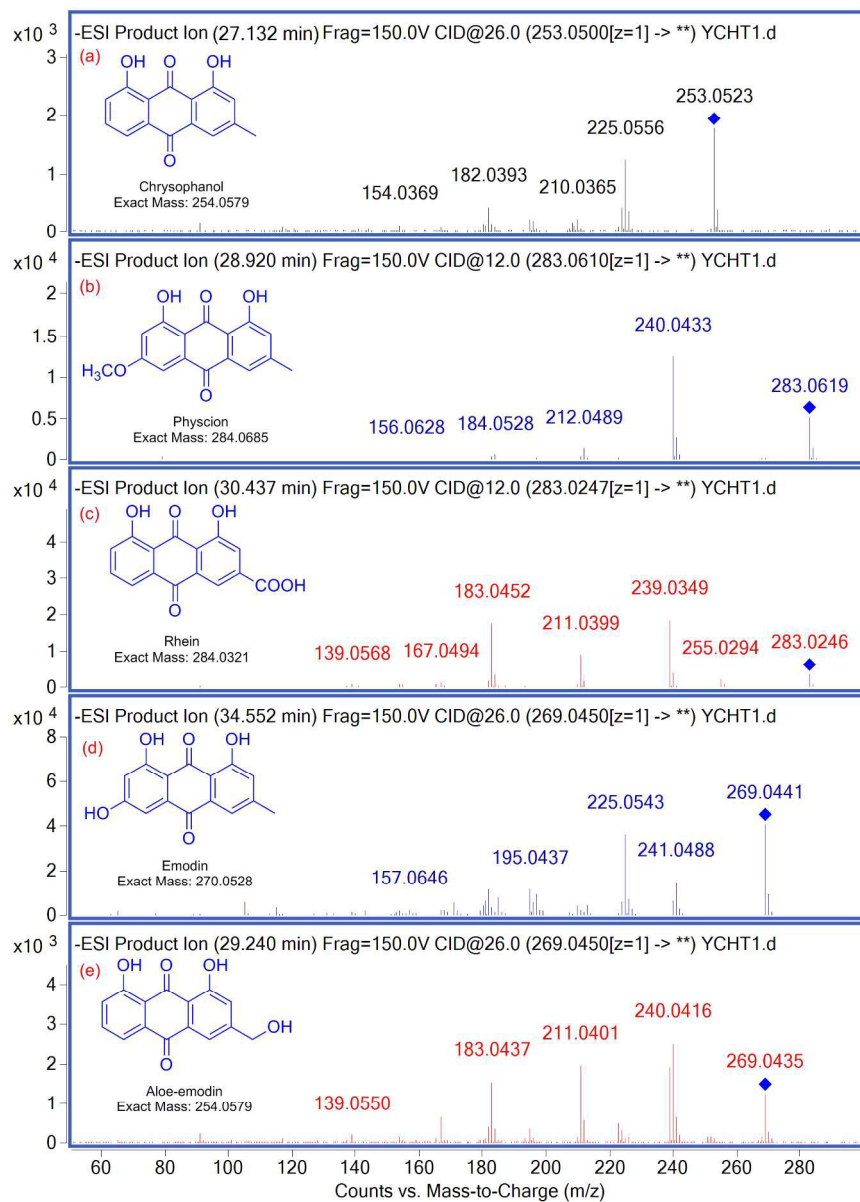
The extract ion chromatogram (EIC) of YCHD in negative ion mode (a); the processed extract ion chromatogram (p-EIC) of YCHD in negative ion mode (b); the scaled to largest for each chromatogram of p-EIC (c).
225x187mm (300 x 300 DPI)



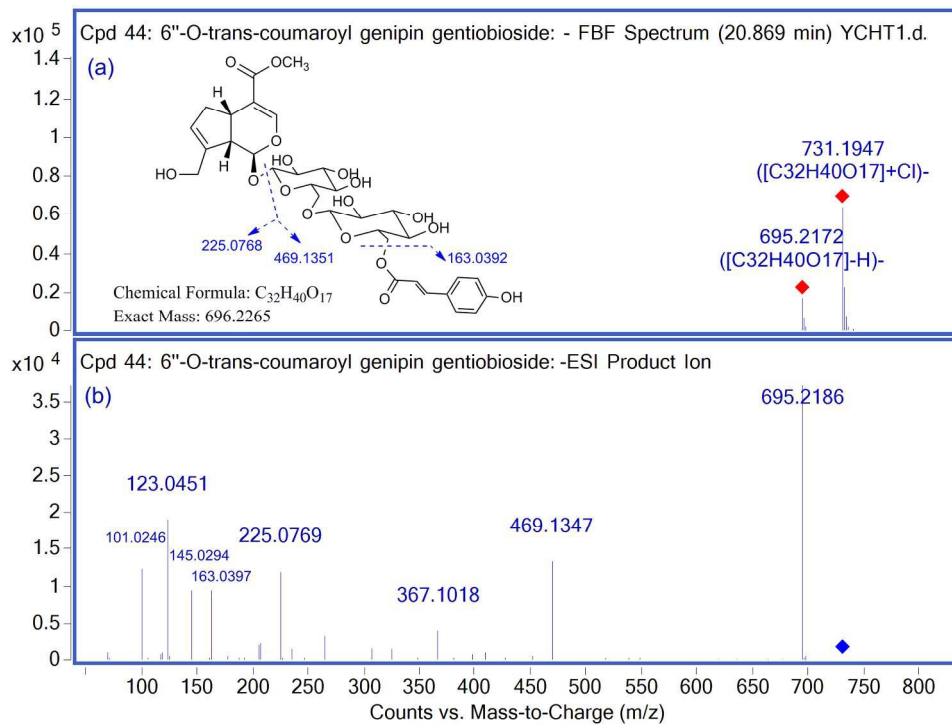
The MS/MS spectra of the $[M-H]^-$ ions at m/z 353.0878 for 1-CQA (a); 5-CQA (b); 3-CQA (c) and 4-CQA (d).
273x239mm (300 x 300 DPI)



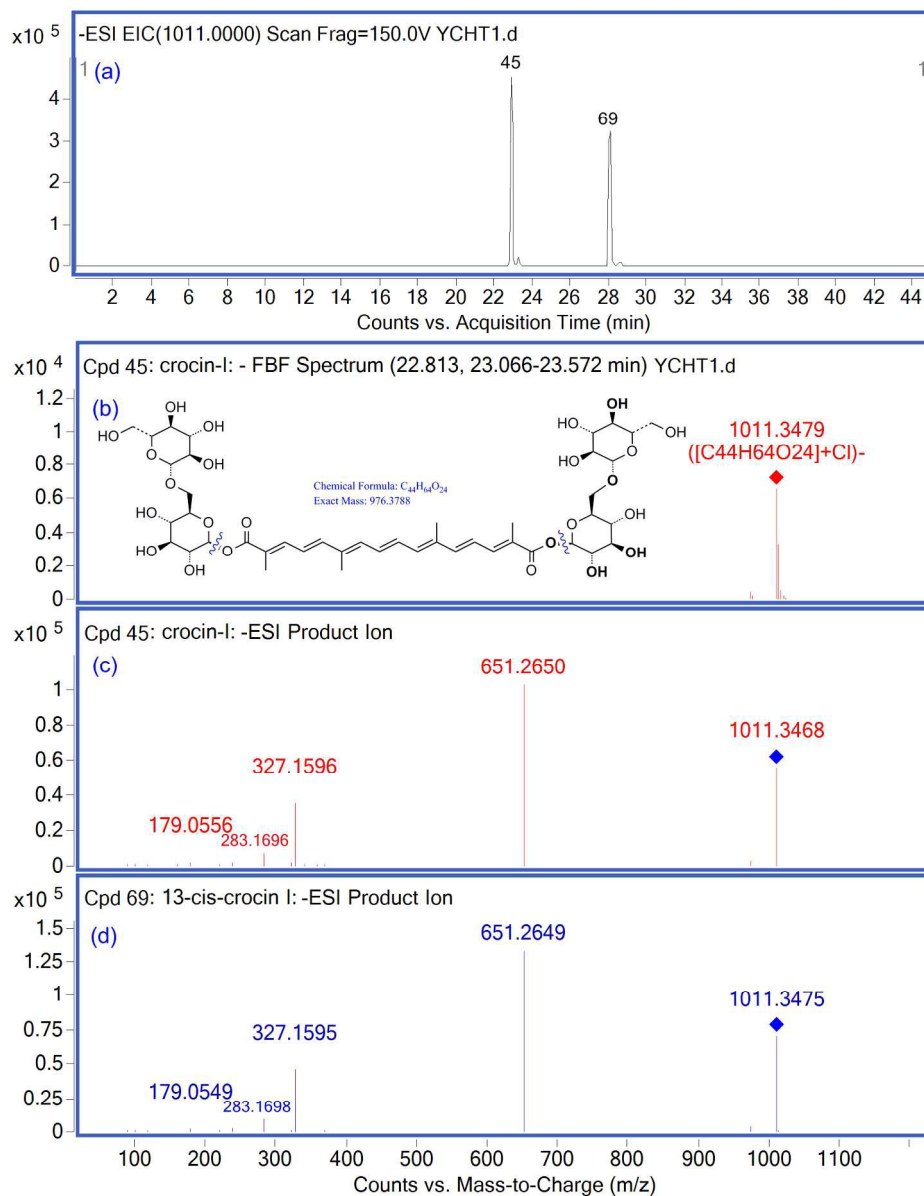
The MS/MS spectra of the $[M-H]^-$ ions at m/z 515.1207 for 1,4-dicaffeoylquinic acid (a); 4,5-dicaffeoylquinic acid (b); 3,5-dicaffeoylquinic acid (c) and 3,4-dicaffeoylquinic acid (d).
273x201mm (300 x 300 DPI)



The MS/MS spectra of the [M-H]⁻ ions at m/z 253.0500 for chrysophanol (a); m/z 283.0610 for physcion (b); m/z 283.0247 for rhein (c); m/z 269.0450 for emodin (d) and m/z 269.0450 for aloë-emodin (e).
184x245mm (300 x 300 DPI)



The FBF spectrum for 6''-O-trans-coumaroyl genipin gentiobioside (a) and its MS/MS spectrum of the $[M+Cl]^-$ ion at m/z 731.1947 (b).
191x139mm (300 x 300 DPI)



The EIC of m/z 1011 in negative ion mode (a), the FBF spectrum for crocin-I (b); the MS/MS spectra of the $[M+Cl]^-$ ion at m/z 1011.3468 for crocin-I (c) and m/z 1011.3475 for 13-cis-crocin-I (d).
188x233mm (300 x 300 DPI)