

1 Supporting Information

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3 **A Novel Strategy for Extracted Ion Chromatogram Extraction to**  
4 **Improve Peak Detection in UPLC–HRMS**

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## 20 **1. Experimental Part.**

### 21 **Dataset 1.**

22 *Sample treatment for tobacco plant and tea samples.* Tobacco plant and tea samples are  
23 grounded to powder before analysis. About 20 mg sample was weighted in a 2.0 mL tube and  
24 added with 1.5 mL solvent of CH<sub>3</sub>OH and H<sub>2</sub>O (70/30, v/v). A 2 min volute, followed by a 30  
25 min ultra-sounded with 40 KHz treatment utilized for components extraction. The tube was  
26 centrifuged at 13000 rpm for 10 min, and 1 μL of supernatant was injected into an Agilent  
27 1290-6540 UPLC-QTOF.

28 An ultra-performance Agilent C18 column was used for tobacco plant sample analysis.  
29 The mobile phase consists of (A) water and (B) ACN with 0.1% Formic acid. A flow of 0.2  
30 mL min<sup>-1</sup> was used. The phase (A) is started from 85% and reduces to 0% within 15 min and  
31 maintains for 5 min. QTOF parameters are: Gas temperature, 350 °C; gas flow, 12 L min<sup>-1</sup>;  
32 nebulizer, 40 psi; sheath gas temperature, 350 °C; sheath gas flow, 10 L min<sup>-1</sup>; Vcap, 3500 V;  
33 and mass range, 50–1000.

34 An ultra-performance Waters T3 column was used for tea sample analysis. The mobile  
35 phase is the same as tobacco plant sample. The mobile phase started with 100% A and then  
36 declined to 70% within 10min, to 10% within 15 min, followed by a decrement to 0% at 17  
37 min and held for 4 min. The QTOF parameters are the same as tobacco plant sample.

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39 *Sample treatment for wine sample.* 100 μL wine sample are pooled with 900 μL solvent  
40 (CH<sub>3</sub>OH/H<sub>2</sub>O, 70/30, v/v). 1 μL sample was injected into the Agilent 1290-6545 UPLC-  
41 QTOF with C18 column. Instrumental condition is the same as tobacco plant sample.

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### 43 **Dataset 2.**

44 *Calibration samples.* Nineteen compounds, including *Metformin hydrochloride*, *Phenformin*  
45 *hydrochloride*, *Telmisartan*, *Clonidine hydrochloride*, *Atenolol*, *Metoprolol*, *Tolbutamide*,

46 *Lovastatin, Simvastatin, Gliclazide, Felodipine, Rosiglitazone maleate, Prazosin*  
47 *Hydrochloride, Candesartan, Cilxetil, Gliquidone, Reserpine, and Nifedipine* were prepared  
48 in methanol. Calibration samples were obtained by diluting the mixture with CH<sub>3</sub>OH.  
49 Concentrations for each compound are 8, 12, 16, 20, 24, and 28 ng mL<sup>-1</sup>. These calibration  
50 samples were analyzed on an Agilent 1290-6545 UPLC-QTOF. The instrumental parameters  
51 are the same as tobacco plant sample.

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53 **Tea mixture samples.** The standard compound solutions were mixture with tea extraction (see  
54 tea sample treatment) to test quantification capability of AiCN-EIC for analyzing complex  
55 samples. Concentrations in these samples are the same as calibration samples. Tea mixture  
56 samples were analyzed on an Agilent 1290-6545 UPLC-QTOF with instrumental parameters  
57 optimized the same as tobacco plant sample.

58

## 59 **2. Data Analysis Part**

60 **Data analysis parameters for XCMS Online, Mzmine2/ADAP, and MS-DIAL.** The  
61 following parameters were used for these methods:

<b>Method</b>	<b>Parameters</b>
XCMS Online	Using parameters that have been optimized for UPLC-QTOF on the web version.
Mzmine2/ADAP	<b>Ion chromatogram extraction.</b> <i>ADAP Chromatogram Builder; m/z tolerance: 0.01 Da or 20 ppm.</i> <b>Peak Detection.</b> <i>Peak deconvolution: wavelets (ADAP); Min highest intensity: 100.</i> <b>Peak Alignment.</b> <i>method: RANSAC; m/z tolerance: 0.01 Da or 20 ppm; RT tolerance: 0.5 min.</i>
MS-DIAL	<b>Data Collection.</b> <i>Centroid parameter: MS1 tolerance 0.01 Da.</i> <b>Peak Detection.</b> <i>Mass slice width: 0.1 Da;</i> <b>Peak Alignment.</b> <i>Retention time tolerance: 0.5 min; MS1 tolerance: 0.01 Da.</i>

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Circle 1							
Step 0							
	scan	m/z	Bleft	Bright	min(Bleft,Bright)	side	cluster
1	1	100.001	Inf	0.001	0.001	right	0
2	5	100.002	0.001	0.0015	0.001	left	0
3	2	100.0035	0.0015	0.0005	0.0005	right	0
4	6	100.004	0.0005	0.0015	0.0005	left	0
5	3	100.0055	0.0015	0.0005	0.0005	right	0
6	7	100.006	0.0005	0.001	0.0005	left	0
7	4	100.007	0.001	0.001	0.001	left	0
8	8	100.008	0.001	Inf	0.001	left	0
i=1							
	scan	m/z	Bleft	Bright	min(Bleft,Bright)	side	cluster
1	1	100.001	Inf	Inf	0		1
2	5	100.002	Inf	0.0015	0.0015	right	1
3	2	100.0035	0.0015	0.0005	0.0005	right	0
4	6	100.004	0.0005	0.0015	0.0005	left	0
5	3	100.0055	0.0015	0.0005	0.0005	right	0
6	7	100.006	0.0005	0.001	0.0005	left	0
7	4	100.007	0.001	0.001	0.001	left	0
8	8	100.008	0.001	Inf	0.001	left	0
<i>When i=2, no merge! continue</i>							
i=3							
	scan	m/z	Bleft	Bright	min(Bleft,Bright)	side	cluster
1	1	100.001	Inf	Inf	0		1
2	5	100.002	Inf	0.0015	0.0015	right	1
3	2	100.0035	0.0015	Inf	0.0015	left	2
4	6	100.004	Inf	0.0015	0.0015	left	2
5	3	100.0055	0.0015	0.0005	0.0005	right	0
6	7	100.006	0.0005	0.001	0.0005	left	0
7	4	100.007	0.001	0.001	0.001	left	0
8	8	100.008	0.001	Inf	0.001	left	0
<i>When i=4, no merge! continue</i>							
i=5							
	scan	m/z	Bleft	Bright	min(Bleft,Bright)	side	cluster
1	1	100.001	Inf	Inf	0	right	1
2	5	100.002	Inf	0.0015	0.0015	right	1
3	2	100.0035	0.0015	Inf	0.0015	left	2
4	6	100.004	Inf	0.0015	0.0015	left	2
5	3	100.0055	0.0015	Inf	0.0015	right	3
6	7	100.006	Inf	0.001	0.001	right	3
7	4	100.007	0.001	0.001	0.001	left	0
8	8	100.008	0.001	Inf	0.001	left	0
i=6							
	scan	m/z	Bleft	Bright	min(Bleft,Bright)	side	cluster
1	1	100.001	Inf	Inf	0	right	1
2	5	100.002	Inf	0.0015	0.0015	right	1
3	2	100.0035	0.0015	Inf	0.0015	left	2
4	6	100.004	Inf	0.0015	0.0015	left	2
5	3	100.0055	0.0015	Inf	0.0015	right	3
6	7	100.006	Inf	Inf	0		3
7	4	100.007	Inf	0.001	0.001	right	3
8	8	100.008	0.001	Inf	0.001	left	0
i=7							
	scan	m/z	Bleft	Bright	min(Bleft,Bright)	side	cluster
1	1	100.001	Inf	Inf	0	right	1
2	5	100.002	Inf	0.0015	0.0015	right	1
3	2	100.0035	0.0015	Inf	0.0015	left	2
4	6	100.004	Inf	0.0015	0.0015	left	2
5	3	100.0055	0.0015	Inf	0.0015	left	3
6	7	100.006	Inf	Inf	0		3
7	4	100.007	Inf	Inf	0		3
8	8	100.008	Inf	Inf	0	left	3

Circle 2							
i=1							
	scan	m/z	Bleft	Bright :ft,Bright)	side	cluster	
1	1	100.001	Inf	Inf	0	right	1
2	5	100.002	Inf	Inf	0		1
3	2	100.004	Inf	Inf	0		1
4	6	100.004	Inf	0.0015	0.0015	right	1
5	3	100.006	0.0015	Inf	0.0015	left	3
6	7	100.006	Inf	Inf	0		3
7	4	100.007	Inf	Inf	0		3
8	8	100.008	Inf	Inf	0	left	3
<i>When i=2, no merge! continue</i>							
<i>When i=3, no merge! continue</i>							
i=4							
	scan	m/z	Bleft	Bright :ft,Bright)	side	cluster	
1	1	100.001	Inf	Inf	0	right	1
2	5	100.002	Inf	Inf	0		1
3	2	100.004	Inf	Inf	0		1
4	6	100.004	Inf	Inf	0		1
5	3	100.006	Inf	Inf	0		1
6	7	100.006	Inf	Inf	0		1
7	4	100.007	Inf	Inf	0		1
8	8	100.008	Inf	Inf	0	left	1
<i>When i=5, no merge! continue</i>							
<i>When i=6, no merge! continue</i>							
<i>When i=7, no merge! continue</i>							
<i>When i=8, no merge! continue</i>							
<b>End</b>							

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64 **Figure S1.** Graphical illustration of the developed EIC construction algorithm. At first, ions  
 65 are sorted based on m/z values. The boundary of each ion is calculated to select matched *side*.

66 Two ions will be merged if the *side* of the  $i$ th is right and the  $i+1$ th is left. For the clustered  
67 ions, the matched *side* is recalculated.