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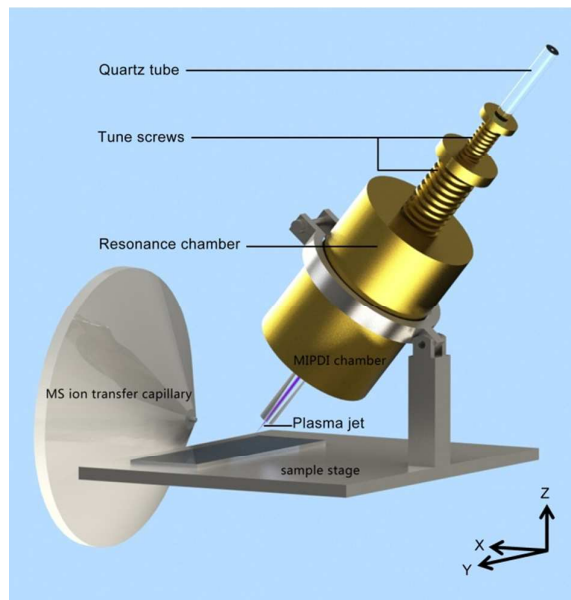
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The quantitative and qualitative behaviors of the MIPDI source were systematically studied for the first time in this work.

1 **Microwave Induced Plasma Desorption**
2 **Ionization (MIPDI) Mass Spectrometry for**
3 **Qualitative and Quantitative Analysis of**
4 **Preservatives in Cosmetics**

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1 **Introduction**

2 Preservatives, which are typically antibacterial agents, are added into cosmetics
3 because the contents of cosmetics are usually organic substances, which are suitable
4 for microbial growth. Also, antioxidants are used in order to retain the effectiveness
5 of the functional substance. Many antibacterial agents contain chlorine or bromine to
6 suppress the growth of microorganisms¹. The halogen is usually allergenic and/or
7 poisonous to organism of human body. To control the use of these preservatives, the
8 additives to cosmetics are strictly limited by the Ministry of Health of P.R.C.
9 National Hygienic Standard for Cosmetics (NHSC). Over 1,200 forbidden additives
10 and 56 permitted preservatives with their maximum dosage are listed by the NHSC
11 to ensure contact safety of products. Excessive adding of cosmetic preservative will
12 increase the potential hazard to health. These substances may lead to acute oral
13 toxicity, acute dermal toxicity, dermal irritation/corrosion, skin sensitization, skin
14 photo toxicity, cellular chromosome aberration, cell gene mutation, teratogenicity
15 and even carcinogenicity. Thus, the analysis of preservatives in cosmetics is a
16 socially relevant and challenging activity. NHSC has drafted a series of regulations
17 and testing protocols. The recommended analysis method from the NHSC is either
18 high performance liquid chromatography (HPLC) or gas chromatography (GC), the
19 most widely used techniques all over the world. These methods provide accurate and
20 plentiful information on the substance of interest. However, the reliable data require
21 appropriate chromatography conditions and long separation time. Furthermore,
22 sample manipulation takes much time and effort. Due to the complex content of

1 cosmetics, an extraction process must be done before chromatography separation is
2 feasible. It has so far been impossible to establish a fast screening and classifying
3 method based on the above techniques. It is therefore worthwhile to search for an
4 alternative method that would avoid the shortcomings of the traditional analysis
5 technologies.

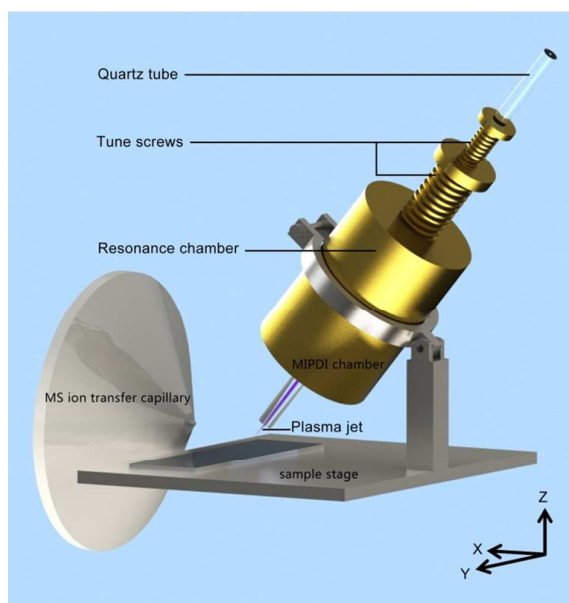
6 The newly emerging technique of ambient desorption mass spectrometry has
7 captured the attention of analysts. The analysis process can be much shortened with
8 the using of a mass spectrometer equipped with an ambient desorption ionization
9 source. Ambient desorption ionization sources for mass spectrometry have been
10 studied for years, extending the use the power of the mass spectrometer. Sample
11 pre-treatments before testing are minimal to none when using ambient desorption
12 ionization sources. Molecules in relatively raw matrices (solid state, liquid state or
13 gaseous state) can be directly ionized, *i.e.* direct analysis of samples is feasible. This
14 means that in-situ analyzing can be realized for workers who want analytes to be
15 kept in their original chemical²⁻⁴ and biological⁵⁻⁹ environment. For these reasons,
16 ambient desorption ionization sources have been applied to the problems of directly
17 analyzing for additives of food^{10,11}, for carcinogenic aromatic amines in textiles^{2,3},
18 for pharmaceuticals¹²⁻¹⁴ and even for metabolites in live cells or tissues¹⁵⁻¹⁷. The
19 ambient desorption ionization sources demonstrated a great reliability of qualitative
20 analysis in these applications. At the same time, the ambient desorption ionization
21 sources provided as low as pg/mm^2 detection limit^{18,19}. The low detect limit makes
22 the ambient desorption ionization sources ideal tools to detect trace substances in

1 complex matrices. Benefiting from the direct ionization of molecules from their
2 native environments, the whole analysis time is short enough (more than 1 sample/s)
3 to be applied in situations where a high through-put test is demanded. Besides, the
4 mass spectra resulting from the use of ambient ionization sources are usually clear
5 without fragment ions. Molecular ions can be easily identified from the spectra. The
6 ability of qualitatively saying yes or no to the presence of a certain substance is quite
7 effective, which means fast screening and classification can be achieved. Owing to
8 these very desirable features, many kinds of ambient ionization sources have been
9 developed. These ion sources are mainly divided into two species, ESI based and
10 plasma based. The recently introduced microwave induced plasma desorption
11 ionization (MIPDI) source¹⁹⁻²¹ is the one among many plasma based ambient
12 desorption ionization sources^{18, 22-24} for mass spectrometry. In a MIPDI source, the
13 discharge gas (argon or helium) is ionized or excited by resonating with microwave
14 power. The produced argon/helium ions and high energy neutral species quickly
15 react with the neutral species in air, forming secondary ions and metastable state
16 neutrals. These relatively stable ions and metastable state neutrals (reactants) are
17 ejected from the MIPDI source to the surface of sample with the flow of the plasma
18 jet. Once these reactants contact the target molecules on the surface of the sample,
19 proton transfer reactions or penning ionization happens. The ionized target
20 molecules subsequently enter the ambient pressure interface and are detected by the
21 mass spectrometer. In our previous work, the MIPDI source has been successfully
22 applied to the qualitative detection of the active ingredient of pharmaceuticals. The

1 MIPDI source showed good tolerance to a complex matrix of tablets and ointments
2 and great reliability in qualitatively showing the existence of target molecules.
3 However, the existing research of the MIPDI source has simply examined its
4 qualitative ability; no report has systematically emphasized the qualitative and
5 quantitative direct analysis of cosmetics preservatives or any similar substances
6 without any kind of sample treatment. It is of potentially great value if the accurate
7 quantitative analysis and potential fast screening capability of MIPDI source can be
8 developed to meet the demands of commercial cosmetics preservatives detection.

9 To evaluate the application of the MIPDI source in the fast detection and
10 classification of preservatives in cosmetics, 6 of the most commonly used
11 preservative compounds (Table 1) were selected as representatives of the class as a
12 whole. Detection limits of the MIPDI source to those preservatives were examined.
13 Both liquid and solid state (ointment) cosmetic samples were included in performing
14 the fast screening experiment by the MIPDI source, covering the categories of
15 sunscreen, facial cream and moisturizer. The fast classification capability of the
16 MIPDI source was examined by a blind classifying of 5 commercially available
17 cosmetics according to the added preservatives. The quantitative analysis capability
18 of the MIPDI source had not been studied. The ability has been commonly
19 investigated by workers using ambient ionization sources²⁵⁻²⁹. To remedy this
20 oversight, the quantitative analysis capability of the MIPDI source was
21 systematically examined by using two methods, the standard adding method and the
22 calibration curve method. The isotopically labeled standard method was also verified

1 using a proof-of-concept experiment to study the accurate quantitative analysis
2 ability of the MIPDI source. Finally, the result was referenced by the NHSC
3 accepted HPLC protocols.



4

5
Scheme 1 Overview of the MIPDI source in its experimental configuration6

Experimental section

7 **Chemicals and Reagents.** Preservative samples (Analytically Pure) were purchased
8 from Sigma-Aldrich (Germany). Methanol (HPLC grade) was obtained from
9 Honeywell. D3-caffeine (3H substituted by 3D, M.W. 197) was purchased from
10 Quandao Company (China). Other chemicals were products of Sinopharm Chemical
11 Reagent Company (China) and were used without further purification throughout the
12 experiment. Ultra purified water ($18 \text{ M}\Omega \text{ cm}^{-3}$) was produced using a UP water
13 purification system (Youpu Company, China). The discharge gas, argon (99.999%),
14 was purchased from ShenQi Gas Company (Chengdu, China). Cosmetics samples,

1 including sunscreen, facial cream and moisturizer, were purchased locally. Glass
2 slides (China, 25.4 mm \times 76.2 mm) were the product of Sail Brand Company.

3 **MIPDI Source Configuration.** The MIPDI source is a type of surface wave
4 generator called a surfatron. The source has been described by our group^{20, 21}
5 previously and detailed information about the experimental set-up can be found there.
6 The resonance cavity is in a brass cylinder. In the center of the cavity, a quartz tube
7 is mounted axially. The discharge gas flow passes through the quartz tube. In this
8 device, microwaves travel along the surface of the quartz tube in the cavity, forming
9 an argon plasma inside the quartz tube. The quartz tube is of 0.8 mm i.d., 6 mm o.d.,
10 and is 200 mm long. The plasma is formed along the surface of the inside of the
11 quartz tube and is a quasi-conical jet. The microwave induced plasma jet outside of
12 the quartz tube was needle-like. The color of the plasma was bright purple. The
13 resonance cavity was tuned to ensure it worked in its best configuration *i.e.* at the
14 lowest reverse power. The MIPDI source was installed on a rotating stage. The
15 rotating stage was mounted on a 3D moving stage (X-Y-Z) in front of the mass
16 spectrometer ion transfer capillary (Scheme 1). Thus, the whole MIPDI source can
17 be moved in 3 dimensions and 1 angle. The sample stage was fixed at the same level
18 as the ion transfer capillary through which the ions were transported to the mass
19 analyzer. Over all, the plasma jet is at a 45° angle to the sample stage and at a 135°
20 angle to the ion transfer capillary. The microwave power generator (2.45 GHz, 150
21 W max, NanJing Yanyou Electronic Science and Technology Co. Ltd.) was set to 6 V,
22 equaling to 70 watts. Argon was used as the discharge gas with a flow rate of 850

1 mL/min. The flow was controlled by a mass flow controller (D07-19B, Beijing
2 Sevenstar Electronics Co. Ltd.). Under the above conditions, the plasma jet was
3 about 5 mm out of the MIPDI source.

4 **Mass Spectrometer Conditions.** A 3D ion trap mass spectrometer (LCQ Fleet,
5 Thermo Fisher Scientific; San Jose, CA) was used throughout this work. All the
6 voltage settings of the mass spectrometer were tuned and calibrated with the tuning
7 method using an ESI as the ion source. The tube lens was manually set to 80 volts
8 and the ion transfer capillary was set to 30 volts to satisfy the mass to charge ratio
9 ranging of interest, which ranged from m/z 100 to m/z 500. Target molecules were
10 ionized while desorbing, so the MIPDI process did not need any method for
11 removing the solvent. As a consequence, the ion capillary temperature was decreased
12 to 175°C. The mass spectrometer was working in full scan and in the positive ion
13 mode (normal scan, above m/z 50) in this study. Spray voltage, auxiliary gas, sweep
14 gas and sheath gas were turned off. The max ion injection time was set to 500 ms
15 while the micro scan was twice each full scan. The instrument software “Xcalibur”
16 was used to process the data from the mass spectrometer.

17 **Sample Treatments and preparations.** Both filter paper and glass slides were
18 examined as potential sample holders. Results showed that filter paper gave more
19 background ions. Also, the sample area was hard to confine due to the
20 chromatography phenomenon. That is the actual sample distribution area is not as
21 large as the solvent ring. Glass slides were therefore used as the sample support
22 system to avoid the shortcomings of filter paper. A 5 μ L analyte sample was pipetted

1 onto the glass slide. The pipetted sample quickly spread out, forming a sample ring.
2 The sample solution dispersed evenly on the glass slide surface. The area of the
3 sample ring was about 10 mm². Experiments were carried out as the methanol
4 vaporized. Two main experiments were performed using the MIPDI source in this
5 study: 1) qualitative analysis of preservatives in commercial cosmetics and 2)
6 quantitative analysis of preservatives in commercial cosmetics. The samples were
7 divided into two types, the solid state and the liquid state. The sample preparation
8 was quite different in those two experiments.

9 For qualitative analysis of preservatives in commercial cosmetics, the sample
10 preparation was simple. The liquid cosmetic sample was diluted 1:1 (v/v) using
11 methanol as the solvent. The diluted solution was pipetted onto the glass slides and
12 analyzed by the MIPDI source. The solid state cosmetic sample, *i.e.* sun screen (5
13 mg), was evenly smeared directly onto the glass slide. The doped glass slides were
14 then ready to be analyzed by the MIPDI source.

15 For quantitative analysis of preservatives in commercial cosmetics, the sample
16 treatment process depended on the state of the sample and the quantitative analysis
17 method. The standard adding method was used to quantitatively analyze liquid
18 cosmetic sample. The liquid cosmetic was 1:1 (v/v) diluted and the solution was
19 used without any further treatment. The calibration curve method was used to
20 quantitatively analyze solid state cosmetic sample, so the sample was extracted by
21 methanol to prepare a sample stock solution. The extraction routine was the same as
22 recommended in the NHSC. A 1.00 g sample of the cosmetic was dissolved in 10 mL

1 of methanol. The solution was energetically shaken for 15 min. After centrifugation,
2 the mixture was filtered through a 0.45 μm filter membrane (organic phase, Hengxin
3 Company). The filtered solution was used as the sample stock solution, which was
4 ideal to going through the quantitative analysis with calibration curve method.

5 **Isotopically Labeled Standards method**

6 The use of isotopic labeled standards method for the MIPDI quantitative analysis
7 was validated in this work. Due to the lack of an isotopically labeled standard
8 preservative, caffeine was selected as a substitute model preservative in order to do a
9 proof of concept experiment. Commercial cosmetics were used as a blank matrix. To
10 produce 125 $\mu\text{g}/\text{mL}$ “matrix matched cosmetic samples”, a 2 mL aliquot of cosmetic
11 matrix was added by 20 μL of 12.5 mg/mL caffeine solution in methanol. These
12 “matrix matched cosmetic samples” were then spiked with known amounts of
13 D3-caffeine (8.6 mg/mL) 5 μL , 10 μL , 15 μL and 20 μL . The volume difference was
14 adjusted to standard by adding 15 μL , 10 μL , 5 μL and 0 μL methanol, respectively,
15 as reference. The D3-caffeine concentration was 21.5 $\mu\text{g}/\text{mL}$, 43.0 $\mu\text{g}/\text{mL}$, 64.5
16 $\mu\text{g}/\text{mL}$ and 86.0 $\mu\text{g}/\text{mL}$, respectively.

17 **Liquid Chromatography Reference Method.** To ascertain the quantitative analysis
18 accuracy of the MIPDI source, high performance liquid chromatography (HPLC)
19 was performed as a reference according to the NHSC standards. An HPLC (Agilent
20 1220 Infinity) was equipped with an Agilent zorbax eclipse XDB-C18 4.6*250mm
21 5-micron C_{18} reversed-phase analytical column and with a UV diode array detector.
22 The wavelength 280 nm was chosen for the analysis, as recommended by the NHSC.

1 Sample treatment before HPLC was as follows: a 1.00 g sample was placed into a
2 cuvette. A water bath was used to remove the volatile solvent. Samples were diluted
3 to 10 mL with methanol. The solution was shaken for 15 min with ultrasonic
4 extraction. After centrifugation, the solutions were filtered through a 0.45 μm
5 membrane to produce the sample stock solution. The calibration solution was
6 prepared by directly dissolving the target preservatives in methanol. The mobile
7 phase, a solution of 50% 0.05 M sodium dihydrogen phosphate, 35% methanol and
8 15% acetonitrile was prepared, ignoring the slight volume changes on mixing.
9 Hexadecane trimethylamine chloride was dissolved in this solvent to the
10 concentration of 0.002 M. The pH of the final solution was adjusted to 3.5 with
11 phosphate buffer. During the HPLC analysis, the mobile phase flow rate was set to
12 1.5 mL/min and the column temperature was ambient. A calibration curve was
13 constructed with 3.5 μL of 2-methyl-4-isothiazolin-3-one (MIT, standard
14 preservative, 100 mg/L, 500 mg/L and 1000 mg/L).

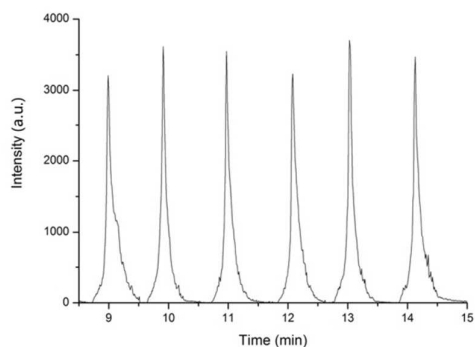
15 **Safety Considerations.** Microwave radiation can be hazardous to people and may
16 lead to pathological changes. Furthermore, electrical shock may happen when
17 igniting the plasma with a slender metal wire. Precautions such as aluminum foil
18 clothing, safety glasses, and electrically insulating gloves should be worn.

19 **Results and discussion**

20 **Analytical Performance of MIPDI-MS.**

21 The detection limits of 6 preservatives were investigated with optimized source

1 parameters (discharge gas flow rate 850 sccm, microwave input power 6 V) by the
2 MIPDI-MS. These samples were pure preservatives dissolved in methanol. The
3 concentration of these samples ranged from 30 to 70 mg/mL. To discover the
4 detection limit, each preservative, contained in 6 identical samples, was tested six
5 times repeatedly. Abnormal values were evaluated with the 4d inspection method³⁰.
6 The detailed results of detect limits and basic information of the preservatives are
7 summarized in Table 1. The ions selected for calculating the detection limits were
8 the protonated molecular ion or the molecular ion. The detection limits of the
9 MIPDI-MS of the selected preservatives were as low as 3.0 pg/mm² and in the range
10 of 3.0 - 5.7 pg/mm² except for the case of climbazole. The relatively poor detection
11 limit of climbazole (400 pg/mm²) was due to its high vaporization point. This high
12 vaporization point makes the climbazole harder to be desorbed/ionized. Overall, the
13 performance of the MIPDI source is remarkable and can easily fulfill the analysis
14 requirements of preservatives for cosmetics under the requirements of the NHSC.



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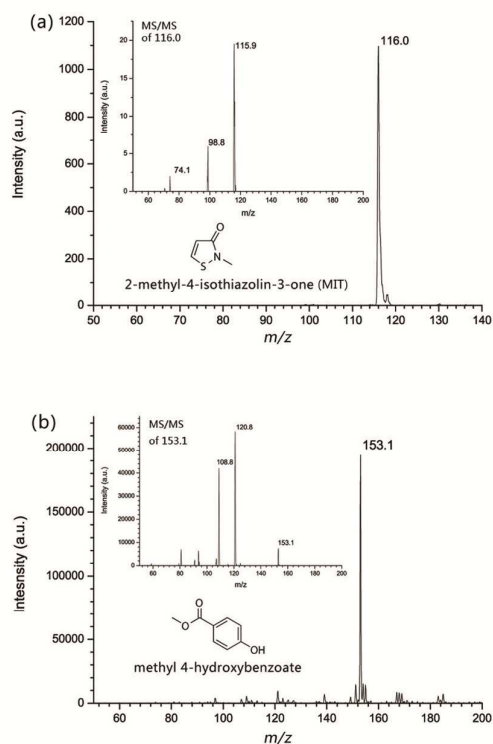
16 Figure 1. Selected ion current (m/z 116) of 6 continuous MIT analysis, RSD of the
17 peak area is 6.7%

18 To evaluate the accuracy of the MIPDI source when used in the detection of

1 preservatives, it is necessary to check the relative standard deviation of the
2 individual test. An acceptable RSD is a basic requirement for the possibility of
3 further quantitative analysis. In RSD test experiment, MIT was tested repeatedly
4 with a set of six identical samples, as described previously. Integrated peak areas
5 (ion current) of the set of samples were used to evaluate the deviation among the
6 tests. Fig. 1 is the selected ion current of MIT (m/z 116 \pm 0.5) of full scan total ion
7 current of the 6 individual tests. The RSD of the 6 individual MIT tests was 6.7%.
8 The RSD of the other preservatives are listed in Table 1. The relatively less satisfied
9 RSD of 4-phenylphenol (m/z 170) is the result of the unstable formation process of
10 dimers (m/z 338, 339). The formation of dimers depends heavily on the ionization
11 temperature. However, due to the open ion source, the air flow in the laboratory
12 severely affected the temperature stability of the desorption point. As a consequence,
13 the product ion ratio of the molecular ion and the dimer ions differed between each
14 sample run, *i.e.* leading to the less satisfactory RSD.

15 **Fast Screening and Classifying Preservatives in Cosmetics with MIPDI-MS.** The
16 standard spectra of two preservatives, MIT (Fig. 2 (a)) and methyl
17 4-hydroxybenzoate (M.W. 152, Fig. 2 (b)), were obtained along, with their collision
18 induced dissociation (CID) spectra by the MIPDI source. These standard spectra
19 were compared with those obtained from commercial cosmetic samples to further
20 assure confidence in the finding of MIT and methyl 4-hydroxybenzoate in order to
21 avoid false positive test. The insets of Fig.2 are the CID spectrum of the
22 corresponding protonated molecular ion. It can be inferred that both MIT (Fig. 2 (a)),

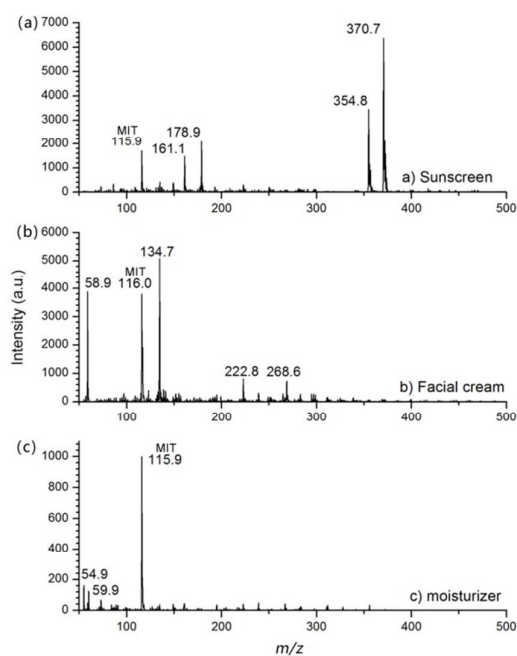
1 $[M+H]^+$, m/z 116) and methyl 4-hydroxybenzoate (Fig. 2 (b), $[M+H]^+$, m/z 153) gave
2 their respective protonated molecular ion. A 25 a.u. collision energy was applied.
3 The featured fragment ions of MIT are m/z 98.8 and m/z 74.0. For methyl
4 4-hydroxybenzoate, a 20 a.u. collision energy was applied to obtain the fragment
5 ions. The featured fragment ions of methyl 4-hydroxybenzoate are m/z 120.8 and m/z
6 108.8. These feature fragment ions were used for the further proof of the existence of
7 those two substances.



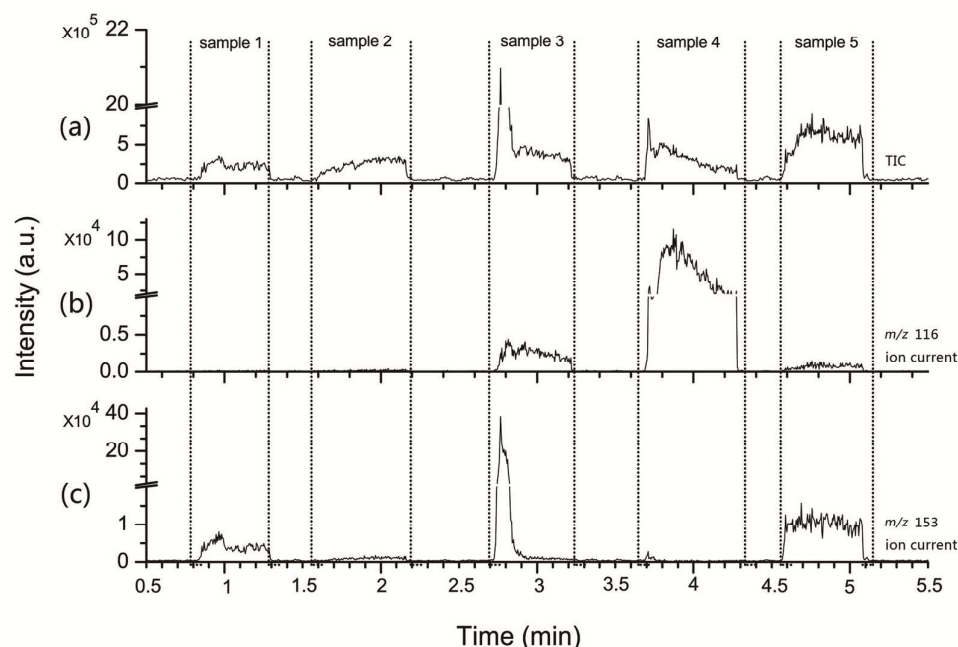
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9 Figure 2. Standard spectra of (a) MIT and (b) methyl 4-hydroxybenzoate, the insets
10 are the CID spectra corresponding to the protonated molecular ions, collision energy
11 are 25 a.u. and 20 a.u. respectively.

12 Locally purchased sunscreen, facial cream and moisturizer were chosen to perform
13 the qualitative analysis experiment. The sample preparation was done by smearing
14 the cosmetics on glass slides. No further pre-treatment was done. The cosmetics

1 samples were desorbed by the MIPDI source directly. The original mass spectra of
2 the samples, corresponding to sunscreen, facial cream and moisturizer, are show in
3 Fig. 3. What can be found in the spectra is that, despite the complex matrix, the
4 presence of MIT ($[M+H]^+$) is still clearly indicated by the m/z 116. Other peaks in
5 the spectra show the ions of ingredient substance in the cosmetics samples desorbed
6 along with MIT.



7
8 Figure 3. Full scan spectra of commercial cosmetics samples: a) sunscreen; b) facial
9 cream; c) moisturizer. The spectra were obtained from direct desorption ionization.



1

2 Figure 4. (a) TIC of fast screening test (no pre-treatment) of 5 commercially
3 available cosmetic analyses by MIPDI. (b) Extracted ion current at m/z 116 (MIT
4 dosage). (c) Extracted ion current at m/z 153 (methyl 4-hydroxybenzoate dosage).

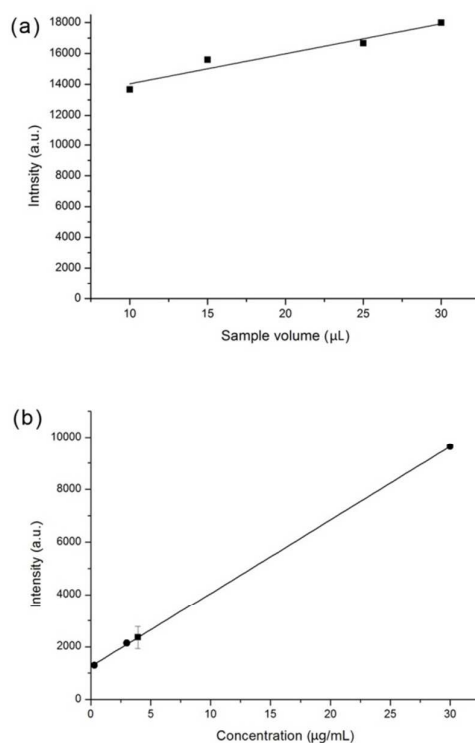
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6 A deficiency that vexes the users of traditional chromatography is the limited
7 sampling speed when chromatography is performed in the qualitative application of
8 bulk samples. The MIPDI source is advantageous compared with chromatography
9 method, because the MIPDI source is exactly suitable to instantly and qualitatively
10 say “yes/no” with regard to one or several substances in a sample, *i.e.* the MIPDI
11 source possesses great potential in fast screening applications. As a very crucial
12 characteristic of the MIPDI source, the fast classification capability cosmetics
13 samples were examined in this research. In the section, a classification process of
14 commercially available cosmetics according to the added preservatives is described
15 using MIPDI source. There were 5 commercially available cosmetics samples from
local cosmetic shop used in this experiment. The selected cosmetics possibly contain

1 MIT and methyl 4-hydroxybenzoate. The cosmetics samples applied to glass slides
2 were randomly analyzed. Depicted in fig. 4 (a) is the real time total ion current (TIC)
3 of the 5 individual cosmetic samples analyzed with the MIPDI source. Fig. 4 (b) and
4 Fig. 4 (c) are the extracted ion currents of m/z 116 (MIT, $[M+H]^+$) and m/z 153
5 (methyl 4-hydroxybenzoate, $[M+H]^+$) respectively. The time axes of the three
6 chromatograms are parallel. The TIC was indicative of the time over which the
7 samples were desorbed. The extracted ion currents were used to determine if there
8 was corresponding ion appears when the samples were desorbed. Accordingly, it is
9 self-evident that the 5 samples were classified into two groups. Sample 3, 4 and 5
10 appear to use MIT as a preservative. Sample 1, 2, 3 and 5 appear to contain methyl
11 4-hydroxybenzoate as a preservative. This was a surprise, since the ingredient lists
12 from the cosmetic packages have it that only 3 samples contain methyl
13 4-hydroxybenzoate and one sample contain both of the preservatives. There must be
14 a false positive result in sample 3 or 5 with respect to the existence of methyl
15 4-hydroxybenzoate.

16 To identify the potential false positive in the measurement, another verification
17 experiment was conducted. A 20 a.u. collision energy was applied to the ion m/z 153
18 in sample 3 and 5 to obtain the CID spectra. The obtained CID spectra were
19 compared to the standard CID spectrum (20 a.u. collision energy) of methyl
20 4-hydroxybenzoate (Fig. 2(b)). Two major fragments m/z 120.8 and m/z 108.8 of
21 methyl 4-hydroxybenzoate in the standard CID spectrum did not exist in the CID
22 spectrum of sample 5. Therefore, sample 5 was excluded from the methyl

1 4-hydroxybenzoate containing list.
2 The whole classification procedure took less than 5 minutes including sample
3 preparations, equaling to 1 sample per minute. The classification time can possibly
4 be significantly reduced further if an automatic sampler were to be used. As an
5 efficient fast screening method, the MIPDI source has proven its usefulness in real
6 sample pre-classification before going through a more quantitative analysis such as
7 HPLC or GC. The pre-classification ability of the MIPDI source made it possible to
8 do a preliminary screening and classification of an unknown sample with one scan.



9
10 Figure 5. a) Calibration curve of liquid state commercial cosmetics sample using
11 Standard Addition Method. The quantitation equation is $y = 194x + 12104$, $R^2=0.915$.
12 b) Calibration curve of solid state commercial cosmetics sample extracts using
13 Standard Calibration Curve Method. The quantitation equation is $y = 2.79x + 1254$,
14 $R^2=0.999$. The round symbol indicates the concentration used to establish the
15 calibration curve and the square symbol indicates the sample concentration found on
16 the curve with standard deviations.

1 **Quantitative Analysis of Preservatives in Cosmetics.** The quantitative analysis
2 ability of the MIPDI-MS was explored for the first time in the present investigation
3 through the standard adding method and standard calibration curve method. To
4 simulate real world analysis conditions, both solid state and liquid state commercial
5 cosmetics were used. The target preservative was MIT. The standard adding method
6 was used for the quantitative analysis of liquid samples and the standard calibration
7 curve method was used for the quantitative analysis of solid samples. Each data
8 point was acquired with a set of 6 individual repetitions in the following experiment.
9 When using the standard adding method, the standard substance is required to be
10 uniformly mixed into the sample substrate. Liquid state sample is suitable to use this
11 method for its favorable dispersion of preservative solution. Consequently, standard
12 adding method was used to quantify the concentration of MIT in the commercial
13 moisturizer which is a kind of liquid state cosmetics. This method can efficiently
14 eliminate the matrix effect, *i.e.* matrix suppression or enhancing can be ignored. To
15 prepare the gradient solution, different volumes (10 μL , 20 μL , 40 μL , 50 μL) of a
16 standard MIT solution (59.6 mg/mL), used as internal standard, were added into 5
17 mL samples of a 1:1 methanol diluted commercial sample respectively. The
18 difference of solvent volume was compensated by adding methanol (40 μL , 30 μL ,
19 10 μL , 0 μL respectively). The obtained gradient sample solutions were directly
20 analyzed with the MIPDI source as is described. Fig. 5 (a) is the quantitative analysis
21 curve ($R^2 = 0.91$). The X-axis represents the volume of added standard MIT solution.
22 The Y-axis represents the integration of ion current intensity. If the curve is extended

1 to reach the X-axis, the absolute value of X on the intersection point is the equivalent
2 volume of the standard MIT (59.6 mg/mL) in original sample. Therefore, the weight
3 of the MIT in the cosmetics sample can be calculated. The MIPDI source
4 quantitative analysis result was 0.14% (w/w). It should be noted that the RSD of the
5 result was the average RSD of the points used to establish the quantitative analysis
6 curve. The relative standard deviation was 8.3%.

7 The above standard adding method is not suitable for the quantitative analysis of
8 solid samples because a different volume of an internal standard was hard to be
9 uniformly added into the sample matrix. Consequently, an extraction procedure was
10 demanded. The extracted sample solution is clean with no interference by the sample
11 matrix. After extraction, the sample solution had a simple matrix (methanol as
12 solvent), so the calibration curve method was suitable for use in this situation. The
13 extraction procedure was the same as recommended in the NHSC standard for the
14 preparation of HPLC sample. The extracted solutions were diluted to 1% (w/w) and
15 analyzed with MIPDI source. At the same time, a calibration curve was established
16 with MIPDI source using pure MIT standard solution. Fig. 5 (b) is the calibration
17 curve ($R^2 = 0.99$). The circles which were used for establishing the calibration curve
18 represented the MIPDI source results of analyzing the standard solutions. The
19 squares represent the extraction solution signal. To obtain a reliable result, each data
20 point was repeated 6 times. The signal from extract is shown with the corresponding
21 standard error. The MIPDI-MS quantitative analysis result was 39 ppm (Table 2) and
22 the relative standard deviation of the 6 tests was 5.9%.

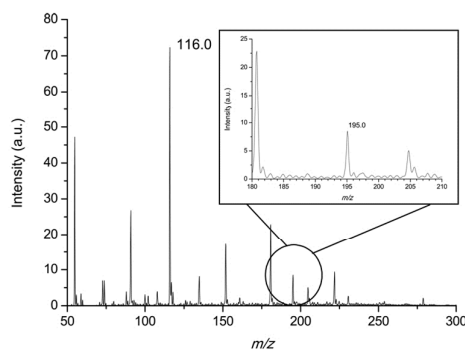
1 The liquid and solid state cosmetics samples were successfully analyzed
2 quantitatively by the MIPDI source. However, the accuracy of the MIPDI result had
3 to be taken into consideration. A comparison of quantitative analysis results was
4 made between HPLC and MIPDI-MS (Table 2). The HPLC quantification result was
5 obtained using the method recommended by the NHSC. The results of each
6 quantitative analysis of each of the samples are listed in Table 2. The liquid sample
7 analyzed with MIPDI-MS had an error of -66% when compared to the HPLC result.
8 The solid sample analyzed by MIPDI-MS had a relative error of -49%. Negative
9 deviation was introduced by the MIPDI-MS quantitative analysis process. The
10 accuracy of the quantitative analysis of the MIPDI-MS is not ideal. The deviation of
11 the quantitative analysis data made it difficult to accurately quantify the
12 preservatives in cosmetics. However, the MIPDI-MS quantitative analysis values
13 were in the same order of magnitude as that of the HPLC method which means that
14 the amount present can be roughly estimated with the MIPDI source *i.e.*
15 semi-quantitative analysis is feasible using the above two methods. The solid state
16 sample had better quantitative result by MIPDI-MS because these samples had a
17 much simpler matrix after extraction. For the same reason, the RSD of solid state
18 sample quantitative analysis is better during analysis.

19 **Quantitative Analysis Using an Isotopically Labeled Standard**

20 The standard adding method and standard calibration curve method had been proved
21 not to be ideal in accurate quantitative analysis when using the MIPDI-MS. The
22 accuracy is acceptable for semi-quantitative analysis (order of magnitude level

1 accuracy). However, the performance should be improved further if the MIPDI
2 method is applied in accurate quantitative applications. Moreover, both of the above
3 quantitative analysis methods require complex sample pre-treatment. For the reasons
4 above, those two quantitative analysis methods for MIPDI-MS have no superiority
5 compared with traditional HPLC method no matter in operational convenience or
6 quantitative analysis accuracy.

7 As an accurate and fast quantitative method, the isotopic labeled standard²⁸ method
8 has been widely used in HPLC-MS. In a few cases, the isotopic labeled standard
9 method was used in the quantitative analysis^{31,32} of ambient ionization source mass
10 spectrometry and obtained superior accuracy and reliable results. In this method, the
11 quantitative analysis process does not merely measure the absolute intensity of the
12 analyte ion, but also the relative intensity between analyte and isotopically labeled
13 standard. In this way, the sampling imprecision and matrix interference can be
14 efficiently reduced. Sampling imprecision is an essential deficiency that strongly
15 affect the quantitative accuracy of the MIPDI source because the desorption process
16 randomly affected by the surroundings due to the open ion source. The quantitative
17 analysis accuracy will be improved if the isotopic labeled standard method is applied
18 in the quantitative analysis of the MIPDI source. Furthermore, the sample
19 preparation is simplified by using isotopically labeled method. All that needs to be
20 done is to spike a known concentration of isotopic labeled standard into the sample.
21 It is necessary to study the application of isotopically labeled standard method in the
22 MIPDI quantitative analysis.



1

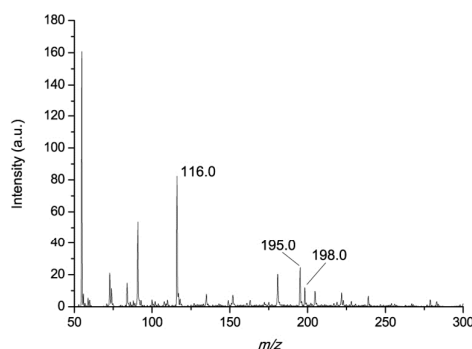
2 Figure 6. Mass spectrum of “matrix matched cosmetic samples” spiked with caffeine;
3 the inset is the detailed mass spectrum around m/z 195.0 (protonated caffeine).

4 To demonstrate the accurate quantitative analysis ability of the MIPDI source, a
5 conceptual experiment was conducted using isotopic labeled standards method. The
6 confirmatory experiment aimed at validating if the isotopic labeled standard method
7 can be used in the quantitative detection of preservative in cosmetic matrix through
8 MIPDI source. During the experiment, caffeine was regarded as preservatives and
9 was added into cosmetics which were called the “matrix matched cosmetic samples”.

10 It was essential to check if the peak of reference D3-caffeine (m/z 198.0) was
11 affected by the matrix before quantitative analysis, because spectral overlap can lead
12 to negative deviation, so blank “matrix matched cosmetic samples” were analyzed
13 by the MIPDI source. As can be seen in Fig. 6, there is no interference peak at the
14 position of m/z 198.0 in the “matrix matched cosmetic samples”. The average
15 intensity at m/z 198.0 is 0.4 a.u.. Such weak signal intensity cannot affect the signal
16 accuracy of D3-caffeine, i.e. the further quantitation process cannot be affected by
17 spectral overlap.

18 The “matrix matched cosmetic samples” were prepared as described in the

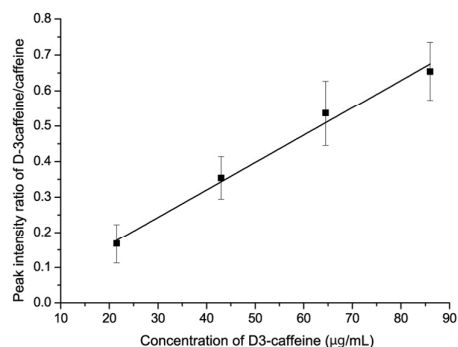
1 experimental section. To quantify the concentration of the “unknown” concentration
2 of caffeine (125 $\mu\text{g/mL}$) in the cosmetic matrices, a comparison was made between
3 the peak height of the caffeine (m/z 195.0, $[\text{M}+\text{H}]^+$) and that of the spiked isotopic
4 labeled standard D3-caffeine (m/z 198.0, $[\text{M}+\text{H}]^+$) which are ionized at the same
5 time through the MIPDI source. The peak height ratio equals the concentration ratio,
6 and thus the concentration of caffeine can be calculated. Fig. 7 is one of the MIPDI
7 mass spectra used for quantitative analysis. The peaks of caffeine (m/z 195.0,
8 $[\text{M}+\text{H}]^+$) and D3-caffeine (m/z 198.0, $[\text{M}+\text{H}]^+$) can be clearly identified along with
9 the ionized matrices. The main preservative, MIT ($[\text{M}+\text{H}]^+$), can also be clearly
10 identified at m/z 116.0. The quantitative analysis experiments were performed with 4
11 different concentrations of reference D3-caffeine, ranging from 21.5 $\mu\text{g/mL}$ to 86.0
12 $\mu\text{g/mL}$. Measurement of each concentration was repeated 10 times. Results show
13 that, the accuracy of MIPDI quantitative analysis is within -3.9% to +5.1%. RSD
14 was as low as 5.5%. The detailed information is generalized in Table 3, including ion
15 intensity ratio, quantitation result, relative standard deviation and accuracy.



16

17 Figure 7. Mass spectrum of “matrix matched cosmetic samples” spiked with caffeine
18 ($[\text{M}+\text{H}]^+$, m/z 195.0) and D3-caffeine ($[\text{M}+\text{H}]^+$, m/z 198.0).

1 To further minimize influence of random error, the relative intensity between the
2 caffeine ion and D3-coffeine ion was used to establish a calibration curve. The
3 X-axis is concentration of D3-caffeine and the Y-axis is the peak intensity ratio of
4 D3-caffeine and caffeine (D3-caffeine/caffeine). The spots used to establish the
5 calibration curve are shown with error bar. The linearity (R^2) of the calibration
6 curve is 0.985. The concentration of caffeine in the “matrix matched cosmetic
7 samples” can be calculated through equation $y = 0.0077x + 0.0114$. The calculated
8 x is the concentration of caffeine, when $y=1$. This means when the peak intensity
9 ratio of D3-caffeine and caffeine equals to 1, the concentration of caffeine equals
10 D3-caffeine. In this method, the quantitative analysis result of unknown caffeine is
11 $129 \mu\text{g/mL}$ and accuracy is $+3.2\%$.



12
13 Figure 8. Calibration curve of Isotopic Labeled Standard method. The curve can be
14 described by equation $y = 0.0077x + 0.0114$ ($R^2=0.985$)

15 The above results suggested that the isotopic labeled standard is a potential
16 methodology in the quantitative analysis of cosmetic preservatives with MIPDI-MS.
17 The RSD of the individual test reached 5.5%. The quantitative analysis ability of
18 MIPDI-MS is competitive to those of other ambient ionization source in such a harsh

1 matrix. The RSD value also approached that of a typical HPLC/GC method (5%),
2 which is satisfactory. The average accuracy of the 4 group of tests is 3.6%, meaning
3 that the MIPDI can accurately quantify the concentration of cosmetics preservatives
4 with isotopically labeled standard method. In addition, no sample extraction was
5 done throughout the isotopically labeled standard method quantitative analysis
6 process. All that needed to be done in the sample preparation was to add the
7 isotopically labeled standard.

8 This is the first report of accurately and quantitatively analyzing trace amounts of
9 samples without sample extractions in complex matrices by MIPDI-MS. This report
10 sets an example of the quantitative analysis for MIPDI source, meaning that
11 MIPDI-MS can potentially be applied to other fields such as drugs, foods and
12 environmental analysis. Also, the MIPDI-MS can be a good alternative to the
13 traditional HPLC/GC method.

14 **Conclusions**

15 The microwave induced plasma desorption ionization (MIPDI) source has
16 demonstrated its worthiness and effectiveness in the fast detection of trace amounts
17 of preservatives in various commercial cosmetics. The quantitative and qualitative
18 analysis behavior of the MIPDI source has been investigated for the first time. The
19 MIPDI source provides as low as a 3.0 pg/mm^2 detection limits for 6 commonly
20 used cosmetics preservatives. The relative standard error of individual MIPDI tests
21 was as low as 5.2%. Direct analysis of commercially available cosmetics without

1 any sample pretreatment was also reported, covering solid and liquid state cosmetics.
2 The fast screening capability was investigated by the blind analysis of 5
3 commercially available cosmetics samples. The cosmetics were successfully
4 classified into two categories according to the added preservatives. Quantifying
5 preservatives in cosmetics using the traditional standard adding method and the
6 calibration curve method was proven to be less effective in accurately quantifying
7 the preservatives in cosmetics even though sample pretreatment was done. The
8 standard adding method (liquid state cosmetics, without extraction) led to a 66%
9 negative deviation and the calibration curve method (solid state cosmetics, with
10 extraction) led to a 49% negative deviation, which was validated with HPLC. A
11 conceptual experiment was conducted using an isotopically labeled standard. Using
12 the method, The concentration of caffeine in cosmetics matrices was quantified
13 satisfactorily with no sample pretreatment, with an RSD value of 5.5% and accuracy
14 of 3.6%. The approaches established in this work indicate that the MIPDI source is a
15 promising tool in future applications where rapid qualitative and quantitative
16 analysis is needed.

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- 6
- 7
- 8

1 **Tables**

2

Table 1. Analytical performance of MIPDI

Compound	M.W.	Selected ion (<i>m/z</i>)	D.L.	Solution (w/w)	RSD
Hexamethylenetetramine	140.2	141.0	5.2 pg/mm ²	1.04 ppb	5.2%
4-phenylphenol	170.2	170.1	5.7 pg/mm ²	1.14 ppb	12.4%
2-methyl-4-isothiazolin-3-one	115.2	116.0	3.0 pg/mm ²	0.60 ppb	6.7%
Climbazole	292.8	293.1	5.7 pg/mm ²	1.14 ppb	8.9%
Hexetidine	339.6	354.6	4.3 pg/mm ²	0.86 ppb	5.5%
2-benzyl-4-chlorophenol	218.7	217.2	400 pg/mm ²	80.0 ppb	8.4%

3

Table 2. Semi-quantitation of MIPDI compared with HPLC

Analysis method	Solid state sample (w/w)	Liquid state sample (w/w)
Liquid Chromatography	77 ppm ± 0.30%	0.41% ± 3.7%
MIPDI-MS	39 ppm ± 5.9%	0.14% ± 8.3%
Accuracy	-49%	-66%

4

Table 3. Isotopic labeled standard quantitation result

D3-caffeine label concentration (µg/mL)	Peak height ratio of D3-caffeine/caffeine	Calculated concentration of caffeine (µg/mL)	Accuracy
21.5	0.168 ± 5.5%	128	+2.4%
43.0	0.354 ± 6.0%	122	-2.8%
64.5	0.536 ± 9.1%	120	-3.9%
86.0	0.654 ± 8.1%	131	+5.1%

5