

Transmission Mode Desorption Electrospray Ionization

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A new mode of operation for desorption electrospray ionization (DESI) analysis of liquids or solid residues from evaporated solvents is presented. Unlike traditional DESI, the electrospray is not deflected off of a surface but instead is transmitted through a sampling mesh at a 0° angle between the electrospray tip, sample mesh, and capillary inlet of a mass spectrometer. In this configuration, deposited samples can be analyzed rapidly without rigorous optimization of spray distances or angles and without the preparation time associated with solvent evaporation. The new transmission mode desorption electrospray ionization (TM-DESI) technique is not applicable to bulk materials, but instead is a method designed to simplify the sample preparation process for liquid samples and sample extracts. The technique can reduce analysis time to seconds while consuming only microliters of sample. The results presented summarize the optimization of the technique, highlight key figures of merit for several model compounds, and illustrate potential applications to high throughput screening of liquid mixtures in both extraction solvents and biological matrices. (J Am Soc Mass Spectrom 2008, 19, 1612–1620) © 2008 American Society for Mass Spectrometry

Desorption electrospray ionization (DESI) [1–40], direct analysis in real time (DART) [41], desorption atmospheric pressure chemical ionization (DAPCI) [12, 22, 42], electrospray assisted laser desorption ionization (ELDI) [43], atmospheric solids analysis probe (ASAP) [44], and flowing afterglow atmospheric pressure glow discharge ionization (FA-APGD) [45, 46] are among the recently developed ambient ionization techniques that have revolutionized mass spectrometric analysis by facilitating direct and rapid analysis of both bulk materials (e.g., tissues, pharmaceuticals) and samples deposited from a solution onto a sampling surface. In the case of DESI, ions are produced by directing charged solvent droplets from an electrospray source toward a sample. Analytes on the surface are ionized and desorbed by the incoming plume before mass analysis. DESI is one of the more universal techniques since it can be used to analyze larger biomolecules and exploit reactive chemistry via alteration of the solvent and solvent additives [5]. However, DESI requires optimization of an array of experimental variables including DESI spray solvent composition, desorption angle and distances, sampling angle and distances [5]. Optimization is especially critical as the chemical identity of the target analytes varies [5].

Since its discovery, DESI has found many applications, including forensic analysis [5, 14, 25, 29], explosives detection [3, 4, 27], chemical warfare agent detection [13, 14, 24, 33], imaging [12, 26], pharmaceutical

analysis [2, 7–9, 15, 22, 29, 31, 38], natural product characterization [5, 6], polymer analysis [16, 35], metabolomics [5, 11, 18, 30], proteomics [5, 10, 32], and recent progress toward glycomics [40]. In some cases, samples are ionized directly in their native environment (e.g., imaging, pharmaceutical analysis) without any pretreatment. However, in many others the sample is extracted or otherwise prepared in a suitable solvent and ultimately deposited onto a surface (e.g., glass slide, paper, metal, plastic, TLC plate). For DESI, this sample preparation process often requires the complete evaporation of the solvent before analysis since the typical incident angle of the electrospray nebulizing plume tends to rapidly erode liquids and dissolved solids from the surface before they are ionized. When the solvent is highly volatile (e.g., methanol), the evaporation time may be minimal; however, the low surface tension of the solvent makes reproducible sample deposition more challenging. In cases where the solvent is less volatile (e.g., water), the slow evaporation time of the solvent may reduce sample throughput.

Historically, a major factor in the successful application of DESI has been the tuning of the geometry of the experiment. The numerous degrees of freedom associated with a freely adjustable sample stage and spray tip relative to a fixed capillary inlet allow incredible flexibility, but in doing so add inherent complexity. Key adjustable parameters, including incident and collection angles, sample height and spray heights, and plume impact to inlet distance are typically optimized, but often differ among sample types. Recent efforts have been aimed at simplifying the DESI experiment by fixing the geometric arrangement of the sample surface,

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spray tip, and capillary inlet [34]. In these studies, the dependence on angles and distances is minimized by enclosing the sampling surface in a small chamber and spraying the surface with both the spray tip and capillary held near 90° to it. The ion cloud is created within the sample chamber and transferred into the mass spectrometer due to the pressure differential of the vacuum. In other work, solid-phase microextraction (SPME) fibers used to extract either vapors from the headspace of samples or analytes from solution have been inserted into an electrospray plume for DESI analysis [24, 39]. In these instances, the surface area was sufficiently small relative to the electrospray plume and ionization of adsorbed surface analytes occurred as the plume traveled around the fiber.

The present investigation reports an alternate method for simplifying the geometry dependence of DESI experiments that require sample deposition. In this method, the sample is not deposited onto a continuous solid surface but rather onto a sampling mesh. Instead of deflecting the electrospray plume off of the surface and into the mass spectrometer, the incident spray angle is reduced to 0° and the spray is transmitted through the sample (Figure 1). This "transmission-mode" DESI technique allows for rapid analysis of both deposited residues and solutions without rigorous optimization of spray distances or angles and without the preparation time associated with solvent evaporation.

Experimental

Materials

Nicotine, cotinine, bradykinin, quercetin, apigenin, myricetin, and rhodamine 6G were purchased from Sigma Aldrich (St. Louis, MO). Neurotensin and Substance P were purchased from Bachem (Torrence, CA). All standards were used without further purification and prepared in high purity solvent (e.g., water, meth-

Table 1. Sample mesh characteristics

Mesh material	Open space (μm)	Strand diameter (μm)	Transmittance (%) ^a
PEEK	300	200	36.0
Nylon	350	240	35.0
Polyester	350	250	34.0
Polypropylene	297	215	33.5
Stainless steel	381	250	36.0

^aThe percent transmittance was calculated as the square of the open space divided by the square of the sum of the open space and strand diameter.

anol, hexane, or acetonitrile) obtained from Fisher Scientific (Hampton, NH). Red and blue permanent markers (Fine Sharpie; Sanford Corp., Oak Brook, IL) were also used as sources of the easily ionized dyes rhodamine 6G and Basic Blue 7.

Five different sheets of mesh material with similar characteristics (Table 1) were purchased from Small Parts Inc. (Miramar, FL) and cut into 5 mm × 10 mm rectangular pieces. Mesh pieces were rinsed with a mixture of water, methanol, and acetone and allowed to dry before use. Blank measurements were taken before sample preparation to ensure that the pieces were free of analyte or any detectable chemical interference.

Mass Spectrometry

An Omni Spray ion source (Prosolia, Inc., Indianapolis, IN) was mounted to a Thermo Fisher Scientific LTQ XL mass spectrometer (Thermo Fisher Scientific Inc., Waltham, MA) and modified to allow a 0° angle between the electrospray tip and capillary inlet to the mass spectrometer. Samples were affixed to the sample slide arm of the Omni Spray ion source using a sample holder constructed of two rectangular pieces of high density polyethylene (HDPE) that held the sample screen on one end. Mass spectra were acquired by scrolling the sample mesh perpendicularly into the electrospray plume between the spray tip and the capillary inlet, thereby allowing transmission of the ionizing spray through the mesh. Specific studies (summarized in the text and discussed in further detail in the Supplementary Material, which can be found in the electronic version of this article) were carried out to determine the optimum position of the sample mesh and electrospray tip relative to the capillary inlet to the mass spectrometer.

Mass spectra were acquired in either positive or negative mode, depending on the ionization affinity of the sample, with the ion accumulation time set to 10 ms and signal averaging set for 4 microscans. The temperature of the heated capillary was held at 250 °C for all sample analysis. Nitrogen was used as the DESI nebulizing gas. A syringe pump (Harvard Apparatus, Holliston, MA) was used to deliver water; methanol; acetonitrile; a 50:50 (vol:vol) mixture of water and methanol; or a 50:50:0.05 (vol:vol:vol) mixture of water,

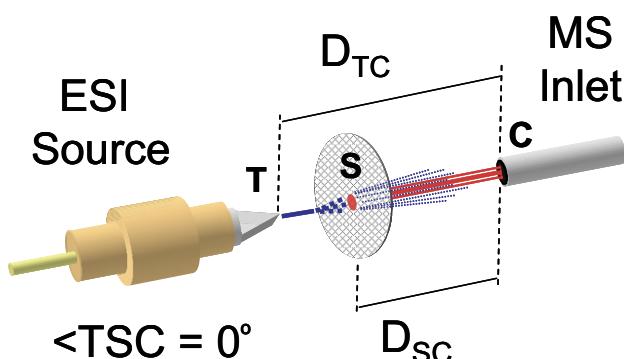


Figure 1. Transmission mode desorption electrospray ionization (TM-DESI) geometry. Sample (S) is deposited on a mesh screen and analyzed by passing an electrospray through it. The angle between the electrospray tip (T), sample (S), and capillary inlet (C) to the mass spectrometer is set to 0°. D_{TC} is the distance from the electrospray tip to the capillary and D_{SC} is the distance from the sample to the capillary.

methanol, and formic acid as the electrospray solvent. Specific studies (summarized in the text and discussed in more detail in the Supplementary material) were carried out to determine the appropriate range of nitrogen pressures, electrospray solvent flow rates, and electrospray voltages.

Sample Preparation

Depending on the experiment, samples were prepared either by spotting 1 μL of solution onto a sample mesh using a 5 μL syringe (SGE, Austin, TX) or by immersing the mesh in solution for several seconds. Specific studies (discussed in detail in the text) were carried out to assess the impact of the sample deposition solvent, the sample substrate, and the drying time (i.e., wet versus dry analysis).

Results and Discussion

Transmission mode desorption electrospray ionization depends on 10 experimental variables that may be subdivided into three categories: those that define the geometry of the experiment, those that characterize the electrospray, and those that govern the desorption/ionization chemistry at the sample surface.

Geometric Variables

With a fixed electrospray angle of 0° (Figure 1), the geometric variables of the experiment are reduced from seven [37] to two: the distance between the sample mesh and the capillary inlet to the mass spectrometer, D_{SC} , and the distance between the electrospray tip and the capillary inlet, D_{TC} . Defining these two distances necessarily dictates the position of the electrospray tip relative to the sample mesh.

Experiments using rhodamine 6G and Basic Blue 7 were conducted to determine the optimal range for the geometric variables D_{SC} and D_{TC} . With the solvent flow rate set to 10 $\mu\text{L}/\text{min}$ and the nebulizing gas pressure set to 100 psi, D_{SC} and D_{TC} were varied incrementally throughout their possible range (i.e., 5 mm < D_{TC} < 21 mm and 1 mm < D_{SC} < 20 mm) and an average peak area for the protonated species at each position was used to construct a contour plot of the response (see Supplementary material, Figure S1). Under these electrospray conditions, the largest responses were observed when the sample mesh was placed between 8 and 10 mm from the capillary inlet and the electrospray tip was held 2 to 3 mm from the mesh (i.e., D_{TC} between 10 and 12 mm). These findings correlate with what is already known about the optimal distance between an electrospray tip and a capillary inlet in a standard ESI experiment [47] and the distance between the ESI tip and a sample surface in a DESI experiment [5]. Furthermore, additional experiments conducted at lower flow rates (e.g., 3 $\mu\text{L}/\text{min}$) favored D_{SC} values of less than 8 mm, thereby indicating that a more compact geometry

that is similar to traditional DESI may also be appropriate for TM-DESI analysis of dried residues. However, when extended to the analysis of wet samples, the response was lower at these shorter distances, a result most likely due to the incomplete desolvation of the analyte ions. Finally, the experimental results also illustrate that the TM-DESI technique is only moderately sensitive to variations in these distances under given electrospray conditions, and that slight differences in sample placement are unlikely to result in major fluctuations in signal intensities.

Electrospray Variables

Variables that characterize the electrospray in TM-DESI are identical to those in standard electrospray experiments, namely the electrospray voltage, the pressure or flow rate of the nebulizing gas, and the flow rate of the sample (i.e., the flow rate of the spray solvent). Experiments using rhodamine 6G and Basic Blue 7 were conducted to determine the impact of varying these variables on the TM-DESI response. With the geometry set at D_{SC} equal to 8 mm and D_{TC} equal to 10 mm, the maximum average peak area was observed when the solvent flow was 10 $\mu\text{L}/\text{min}$ and the nitrogen pressure was 100 psi, although other pairings also produced strong responses (see Supplementary material, Figure S2). Variation of the electrospray voltage showed a maximum response at 4.0 kV, with a range of values greater than 3.0 kV providing adequate response (see Supplementary material, Figure S3). While the results for nebulizing gas pressure and electrospray voltages are typical for both standard electrospray experiments and DESI analyses [5], the optimal flow rate of 10 $\mu\text{L}/\text{min}$ concurred with several values reported for DESI [7, 17], but was higher than others performed with lower flow rates of only 2–5 $\mu\text{L}/\text{min}$ [5, 12]. As discussed in the previous section, this optimal flow rate is likely an artifact of the chosen geometry and would be reduced if the geometry were more compact.

Surface Desorption/Ionization Variables

Variables that influence the surface ionization in TM-DESI include the identity of the spray solvent, the identity of the sample deposition solvent, the composition and physical characteristics of the substrate material, the identity of the target analyte, and the surface density of the target analyte deposited on the substrate. The chemistry of the desorption/ionization mechanism has been the subject of several reports [12, 21, 23, 37]. TM-DESI depends on four variables that are in common with traditional DESI: the composition of the substrate material, the identity of the electrospray solvent, the identity of the target molecule, and the surface density of the target molecules on the substrate. Since TM-DESI of liquid samples is also possible, the identity of the sample deposition solvent is an additional variable that may directly influence the desorption and ionization in

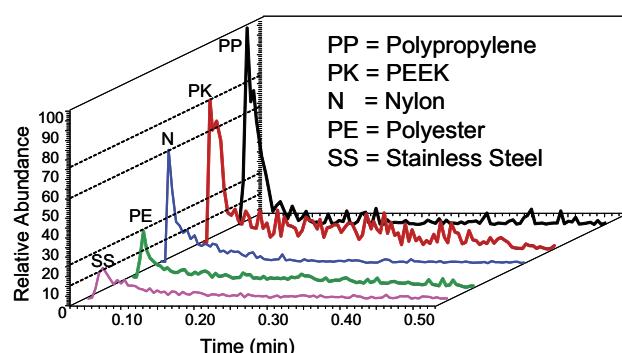


Figure 2. Relative TM-DESI signal of 10 pg of rhodamine 6G from five different mesh materials with similar transmission characteristics (i.e., strand diameters of ~200–250 μm and open space of ~300–400 μm). Samples were spotted using 1 μL of methanol and allowed to dry prior to analysis. No additional potential was applied to the surfaces during the analysis.

these analyses. While a complete investigation of surface parameters was outside the scope of this work, exploratory studies were performed to investigate the potential influence of several of the principal variables.

Sample Substrate

Several attributes of the sample substrate, including the material of construction, the mesh stand size, and mesh open space (i.e., the mesh transmittance), govern how efficiently analytes are suspended on and removed from the surface. Five sample meshes of differing materials but with similar physical characteristics were investigated in this work (Table 1). As an initial study, the effect of the substrate material on the lifetime of the TM-DESI signal was investigated for rhodamine 6G, nicotine, and bradykinin at both low and high concentrations and for both wet (i.e., solvated) and dry analysis (i.e., analysis after evaporation of the deposition solvent).

As exemplified in Figure 2 for rhodamine 6G, the substrate material had a noticeable impact on the response, especially at lower concentrations. In this case the polypropylene, PEEK, and nylon meshes produced the largest responses while the polyester and stainless steel meshes produced much lower responses. It should be noted that no additional potential was applied to the surfaces during analysis. Therefore, the relatively poor ionization efficiency for the stainless steel mesh may be caused by charge dispersion at the conductive surface. For all of the compounds studied, both wet and dry analysis of low concentration samples produced sharp initial peaks, as observed in Figure 2, followed by rapid signal decay. When the concentration was increased, a sharp initial peak was still observed, but decay was at a much slower rate. In general, results obtained for nicotine and bradykinin were comparable to those for rhodamine 6G and look very similar to what has been previously reported for other dyes by DESI [25].

Aside from the differences observed between the mesh materials, a discrepancy was also noted between

wet and dry analysis of samples at higher concentrations. In this case, the signal decay for dry analysis was much more gradual as analyte was continually desorbed from the surface. In contrast, the wet analysis produced slightly larger but more rapidly decaying responses since the analyte was already solvated and thus easier to desorb from the surface. Overall, these results suggest that the identity of the surface substrate may influence the performance of TM-DESI. Therefore, additional studies that more comprehensively evaluate the performance of a larger variety of meshes are underway.

Electrospray Solvent

The importance of the electrospray solvent composition on DESI analysis has been reported numerous times [5, 12]. These studies have suggested that the optimum solvent is analyte dependent and that the efficacy of a particular solvent is a function of its polarity and the solubility of the target analyte. Various common electrospray solvents including methanol, water, acetonitrile, and mixtures of these solvents have been utilized in DESI analyses [5]. In the present study, experiments were undertaken to study the effect of varying the electrospray solvent on the response of nicotine and bradykinin. In each case, 1 μL of solution (1 ng/ μL of nicotine in methanol or 50 μM bradykinin in methanol) was deposited onto a nylon mesh and allowed to dry completely before analysis.

Table 2 summarizes the results for these experiments and confirms that electrospray solvent effects are also observed in the transmission mode. The largest response for nicotine was observed when methanol was used as the spray solvent, while much lower responses were observed using water or acetonitrile. In contrast, the results for bradykinin show that the largest response was observed when water was the spray solvent and much lower responses were observed for either pure methanol or pure acetonitrile. Thus, the optimum spray solvent depended on the identity of the target analyte. These results again illustrate that the mechanism of desorption in the transmission mode is likely similar to traditional DESI when the deposition solvent is allowed to evaporate completely before analysis.

Table 2. Relative percent responses of dried nicotine and bradykinin samples using various electrospray solvents^a

Analyte	Methanol/water ^b			
	Methanol	water ^b	Water	Acetonitrile
Nicotine	100.0	60.0	25.8	21.8
Bradykinin	10.7	82.2	100.0	4.2

^aThe relative percent response was calculated as the average response from 10 replicate sample preparations divided by the maximum average response. All samples were prepared on a nylon mesh.

^bThe ratio of methanol to water was 1:1 by volume.

Deposition Solvent

The electrospray angle in most DESI experiments is typically much greater than zero (i.e., 40° to 70°), which causes rapid erosion of liquids from uniform sample surfaces such as glass, plastics, and metal. Thus, analysis of liquids in traditional DESI is most successful from rough substrates such as paper or TLC plates that absorb the solvent and hold the liquid analytes in place. The 0° electrospray in TM-DESI eliminates this erosion effect as the nebulizing gas, desolvated ions, and unionized droplets are directed through the surface at an angle perpendicular to it, thereby reducing the tendency for surface resolvation and spreading of the target analyte on the substrate. Therefore, analysis of liquids from smooth mesh surfaces is possible in TM-DESI and, as a consequence, the deposition solvent plays a more prominent role in the surface chemistry of the analysis.

To investigate the impact of the deposition solvent, the experiments conducted to test the impact of the electrospray solvent were repeated using various deposition solvents. However, unlike the previous experiments, the samples were analyzed while they were suspended as liquid droplets in the mesh, not dried to a solid film. Table 3 summarizes the results obtained for nicotine using three electrospray solvents (i.e., methanol, acetonitrile, water) and four deposition solvents (i.e., methanol, acetonitrile, water, hexane). The most compelling results are the differences observed amongst a set of analyses that utilized the same electrospray solvent, but differing deposition solvents. For example, relative responses varied between 22.8% and 100% when the deposition solvent was varied and methanol was used as the electrospray solvent.

Table 4 summarizes the results obtained for bradykinin using four electrospray solvents (i.e., methanol, acetonitrile, water, methanol/water) and two deposition solvents (i.e., methanol, water). These results correlate well with those presented in Table 2 and also suggest that the efficiency of ionization may be controlled by either the electrospray solvent or the deposition solvent. More specifically, from Table 2, the largest responses for bradykinin (as a dried film) were observed when either water or a mixture of water and methanol was used as the electrospray solvent. Table 4 also shows that the largest responses for solvated brady-

Table 3. Relative percent responses of wet nicotine samples using various electrospray and deposition solvents^a

Deposition solvent	Electrospray solvent		
	Methanol	Acetonitrile	Water
Methanol	100.0	37.7	36.6
Acetonitrile	83.9	34.0	22.7
Water	22.8	10.2	11.5
Hexane	59.6	14.6	4.9

^aThe relative percent response was calculated as the average response from 10 replicate sample preparations divided by the maximum average response. All samples were prepared on a nylon mesh.

Table 4. Relative percent responses of wet bradykinin samples using various electrospray and deposition solvents^a

Deposition solvent	Electrospray solvent			
	Methanol	Methanol/water ^a	Water	Acetonitrile
Methanol	23.3	80.8	96.9	13.2
Water	100.0	64.5	42.0	44.6

^aThe relative percent response was calculated as the average response from 10 replicate sample preparations divided by the maximum average response. All samples were prepared on a nylon mesh.

kinin were observed when water was present, either as the deposition solvent in combination with a methanolic electrospray or as the electrospray solvent in combination with a methanolic deposition. Interestingly, the average responses for either the all-methanol or all-water systems were far lower. Furthermore, the magnitude of the response observed when a mixture of methanol and water was used as the electrospray solvent was between that observed for the pure solvent systems.

These results clearly illustrate that the identity of the deposition solvent has an impact on the response in TM-DESI and that the mechanism in the solvated transmission mode analysis is dependent not only on the surface, analyte, and ionizing solvent, but also on the solvent that suspends the target analyte on the mesh. Elucidation of the ionization mechanism must therefore consider not only the interaction of the electrospray solvent with the analyte, but also the interaction of the deposition solvent with the surface and electrospray solvent, the partitioning of the analyte between the deposition solvent and the surface, and the possible partitioning of the analyte between the two solvents.

One possible mechanism could assume that there is no adsorption of the target analyte to the surface and that the ionization results from a partitioning of the analyte between the two solvents. This mechanism would be similar to a liquid–liquid extraction where the miscibility of the two solvents and the corresponding solubility of the target analyte have a large influence on the eventual efficacy of the ionization. Results presented in Table 3 for the analysis of nicotine when hexane was used as the deposition solvent and water was used as the electrospray solvent support this argument. Since the miscibility of these two solvents is minimal, the analyte would have less opportunity to partition into the electrospray solvent and therefore little tendency to be ionized.

Another possible mechanism could involve a “secondary” electrospray of a completely suspended analyte where a suspended analyte droplet is dislodged from the surface by the electrospray plume and subsequently ionized by the desolvation of the analyte amidst the other electrospray solvent ions. Finally, like traditional DESI, the mechanism could involve a droplet pickup mechanism in which the analyte is picked up from the surface or from a thin layer of solvent sur-

Table 5. Figures of merit^a

Analyte	LOD (dry) pg	LOD (wet) pg/ μL	% RSD (wet) ^d
Nicotine ^b	1.0	0.5	11.4
Bradykinin ^c	0.3	0.3	9.6

^aAll analyses were performed using a PEEK mesh and a deposition volume of 1 μL (methanol).

^bMethanol was used as the spray solvent.

^cA mixture of water, methanol, and formic acid (50:50:0.05% by volume) was used as the spray solvent.

^dResult of 10 replicate measurements taken at a concentration 10 times the limit of detection (LOD).

rounding the surface [5, 21, 23]. However, in this case the surface activity of the solute in the various deposition solvents would also be a factor as analytes that favored the surface over the solvent would more likely be available for “pickup” by the electrospray solvent. The data presented here are not sufficient to conclusively elucidate the mechanism of the solvated transmission mode desorption ionization, but they will serve as a guide for subsequent investigations.

Figures of Merit

Table 5 summarizes the detection limit results for nicotine and bradykinin by TM-DESI. Comparison of the results for wet and dry analyses does not suggest a clear benefit in sensitivity using either method. However, it is worth noting that TM-DESI analysis of wet samples consumes the sample more rapidly than dry analysis, since the release of the suspended solvent droplets is nearly instantaneous when the sample intersects the nebulizing plume. Therefore, samples deposited as 1 μL or less and analyzed wet have a “spot” size on the same order as the DESI spray. In contrast, TM-DESI of dry samples produces longer-lived signals because the desorption process must be initiated by resolvation of analytes that have dried on the mesh. In this case, the entire “spot” may not be sampled simultaneously. Therefore, the relative similarity in the sensitivity of the wet and dry analyses may be a balance between these factors. While the reported detection limits cannot necessarily be extended to all analyses, they do suggest that the transmission mode sensitivity is comparable with that of traditional DESI for small molecules like nicotine and peptides such as bradykinin [5].

The qualitative repeatability of TM-DESI was tested by performing 50 replicate analyses of wet and dry samples of both bradykinin and nicotine at sample concentrations twice the limit of detection. For the nicotine samples, nicotine was detected in all 50 samples, and for the bradykinin samples, 49 of 50 samples yielded positive results. In addition, the quantitative reproducibility at concentrations 10 times the limit of detection was also investigated (**Table 5**). The results show ~10% to 12% RSD for the analysis of wet samples deposited as 1 μL in methanol. These results are prom-

ising for qualitative screening, but quantitative analyses may require even better precision.

The possibility of using a codeposited internal standard for TM-DESI analysis of liquid samples was explored. In this study, nicotine (10 pg/μL) was added as an internal standard to cotinine standards (50–500 pg/μL) in methanol and analyzed immediately following deposition of 1 μL of sample on a PEEK mesh. Five replicate measurements were taken at four concentrations of cotinine, and the resulting calibration curve is shown in **Figure 3**. The percent RSD for the individual points was reduced from an average of 16% without the internal standard to an average of 6.7% with it. These results are encouraging and suggest that further work should be done to improve the performance. Moreover, results recently reported by Ifa et al. describe similar success and suggest that further improvements may also come from using a deuterated internal standard [38]. Additionally, monitoring of selected transitions in MS-MS analyses or competitive reactions using reactive DESI may also help facilitate better quantitative capability.

Applications

Mass spectra collected for mixtures of peptides and flavonoids obtained via a high throughput adaptation of the TM-DESI analysis are depicted in **Figure 4a** and b. These spectra were acquired using an alternate sampling approach in which the mesh was simply immersed in the sample solution for 1 s, affixed to the sample holder, and scrolled into the electrospray plume. This methodology eliminated precise sample spotting and therefore reduced the sample preparation time. The efficiency was thus greatly increased, and sample analysis, including preparation, was routinely performed in less than 20 s. This high speed qualitative screening technique could find applications where sample volumes are

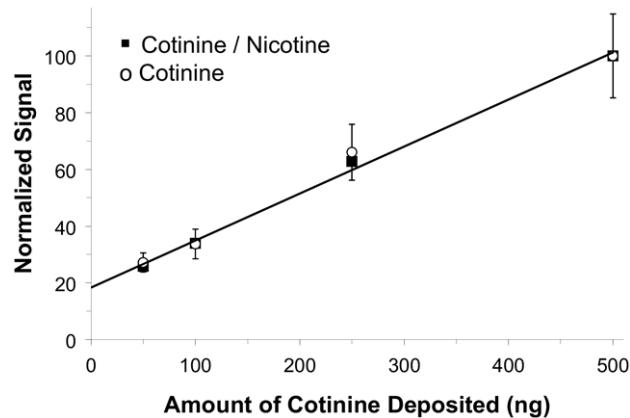


Figure 3. Calibration curve for cotinine with and without the use of nicotine as an internal standard. Error bars indicate one standard deviation of five replicate measurements of cotinine without adjustment to the internal standard.

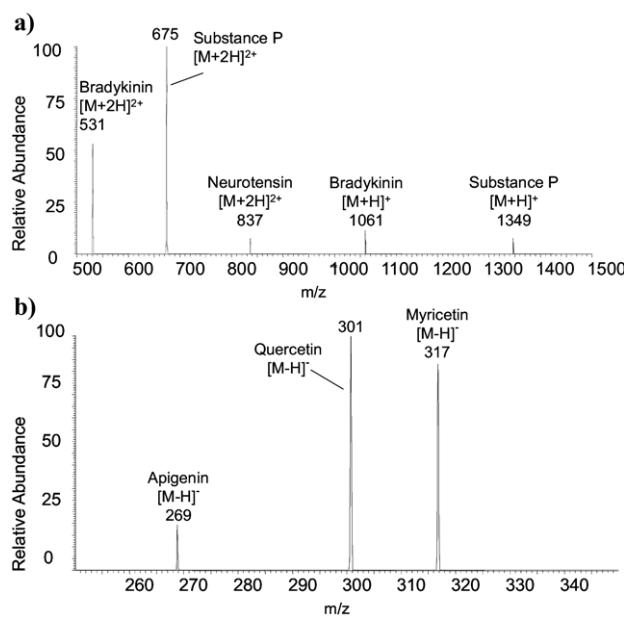


Figure 4. TM-DESI mass spectra for mixtures of (a) bradykinin, neurotensin, and substance P (1 μ M in methanol) and (b) apigenin, quercetin, and myricetin (100 μ g/mL in methanol). Analysis was performed by immersing the mesh in the solution, affixing the mesh to the source, and scrolling the wet mesh into the electrospray plume.

not the limiting factor and quantitative analysis of the solution is not the primary objective.

One potential application of TM-DESI is the analysis of foreign substances and their metabolites in biological matrices (e.g., urine, plasma, saliva). As an example, TM-DESI was used to analyze urine samples for cotinine, the primary urinary metabolite of nicotine. Here, fresh urine samples were collected and spiked with cotinine at a concentration of 100 pg/ μ L. Samples were spotted (1 μ L) either directly onto a PEEK mesh and dried before analysis or diluted 1:100 with methanol and analyzed wet. Similar dilutions or sample extractions have been utilized in some [28], but not all [30, 39], DESI studies involving urine. In the present study, direct analysis of the liquid urine samples without this dilution showed significant ion suppression by the urine matrix, presumably due to the remaining solvated salts. TM-DESI of dried urine samples showed less ion suppression. These results are in accordance with published reports highlighting the high salt tolerance of typical DESI experiments [36], and point toward an ESI-like component in the ionization mechanism of liquid samples via TM-DESI.

Figure 5a depicts a representative mass spectra obtained in the analysis. The ion of m/z 177 corresponds to protonated cotinine, whereas the ion of m/z 199 corresponds to sodium-cationized cotinine. A MS-MS spectrum of the ion of m/z 177 was collected to confirm the presence of cotinine (Figure 5b). The major product ion (m/z 146) corresponds to the loss of CH_3NH_2 . The ions of m/z 98 and 80 stem from cleavage of the bond between the pyridine and pyrrolidine rings.

between the pyridine and pyrrolidine rings. The resulting product ion spectrum matches previously reported results for analysis of cotinine by LC-MS [48, 49] and illustrates that tandem mass spectrometry is also possible in TM-DESI analysis of wet samples, despite the relatively short-lived ion signal. Furthermore, the diluted concentration of 1 pg/ μ L in this analysis is above the detection limit of 0.156 pg/ μ L reported by LC-ESI-MS, but in the vicinity of the limit of quantitation (2.5 pg/ μ L) reported in the same study [48]. While TM-DESI did not produce a lower detection limit than LC-MS, the analysis was performed in seconds compared to the several minutes required for the LC-MS assay.

Conclusions

The results reported here demonstrate that TM-DESI is capable of producing high quality mass spectral data for both solid residues and liquid samples in very short periods of time. The 0° electrospray angle transmits the ionizing plume through the sample surface and effectively reduces the number of geometric experimental variables to two, thus providing a useful simplification to the traditional DESI experiment. The bulk of the variability in the TM-DESI analysis remains defined by the five variables that describe the desorption process: the electrospray solvent, the deposition solvent, the substrate material, the target analyte, and the partition-

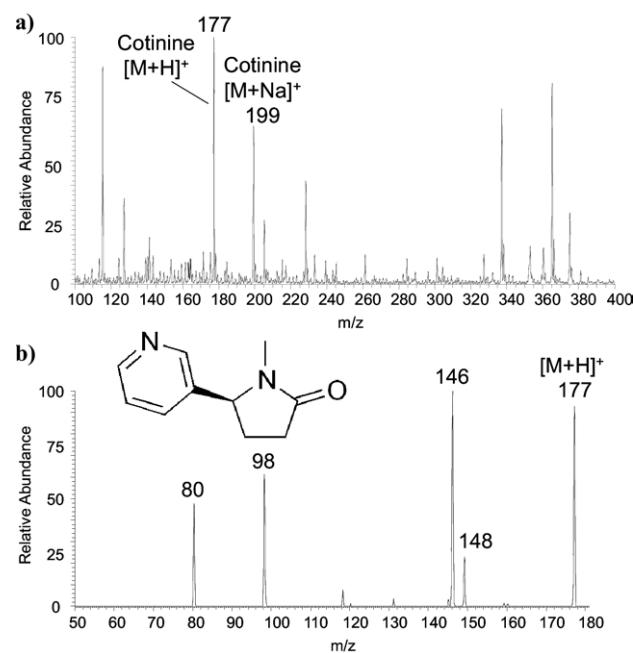


Figure 5. TM-DESI spectrum of (a) urine containing cotinine, the primary urinary metabolite of nicotine, and (b) MS-MS spectrum of the ion of m/z 177 used to confirm the presence of the protonated cotinine species. The major product ion was of m/z 146, which is the loss of CH_3NH_2 from cotinine. The ions of m/z 98 and 80 stem from cleavage of the bond between the pyridine and pyrrolidine rings.

ing of that analyte onto the substrate or into the deposition solvent. Ongoing research aims to move beyond the simple surfaces and solvents tested here and, in doing so, improve the selectivity of the TM-DESI technique by taking advantage of the unique chemistries that can exist in these multicomponent systems.

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Appendix A. Supplementary Material

Supplementary material associated with this article may be found in the online version at doi:10.1016/j.jasms.2008.07.002.

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