

# Complementary classification of solid oral dosage forms in ambient conditions by desorption electrospray ionization mass spectrometry and transmission Raman spectroscopy

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# Abstract

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Discrepancies or defects in active ingredients, excipients and coatings that form solid oral dosage forms can both impact product quality and provide hallmarks of off-brand or counterfeit products. There is therefore a need for rapid and continuous analytical techniques that can assess and classify product differences of intact samples at- or near the production line, or in analytical labs, ideally without resorting to product dissolution.

Here we test the ability of two rapid ambient chemical characterization methods to discriminate between solid dosage forms: desorption electrospray ionization mass spectrometry and transmission Raman spectroscopy. These two techniques are highly complementary, offering greater sensitivity to the analysis of the surface and the tablet bulk, respectively. The data sets generated were then used to test a variety of classification algorithms including linear discriminate analysis, tree-based methods, a simple neural network, and support vector machines (SVM). The highest performing algorithms for DESI-MSI were the SVM, with an additional performance boost when used with a polynomial kernel. For transmission Raman data, a linear discriminant analysis (LDA) model was found to be the most effective.

## Introduction

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Inconsistencies in active ingredients, excipients, the thickness and integrity of coatings and the presence of impurities in solid oral dosage forms all negatively affect their performance. Inferior quality attributes can be useful to identify off-brand or counterfeit products. There is a need for rapid and continuous analytical techniques that can assess and classify product differences of intact samples at- or near the production line, or in analytical labs, ideally without resorting to product dissolution [1]. Rapid measurement tools are particularly important to enable continuous monitoring, necessary to support the change from batch to continuous manufacturing. Analytical methods are required to monitor both the actives, coatings and consistency of the product: For example, in addition to the total API content, insight is also needed on degradation products, impurities, (co-) crystallinity/presence of polymorphs, and content uniformity. The ability to monitor tablet coating thickness and integrity is of great importance, particularly for functional coatings, such as gastro-resistance, which would be compromised by insufficient thickness, or the occurrence of cracks in the film [2].

Quantitative analysis of pharmaceutical tablets is routinely performed by HPLC which offers accurate and sensitive measurements of the active ingredient(s) and excipients, in addition to the presence of any contaminants. However, solution-based analytical methods are destructive and labor-intensive.

Mass spectrometric methods can provide unlabeled identification, both of expected ingredients in known samples and of contaminants or components of unknown formulations. Ambient ionization mass spectrometry approaches including DESI (desorption electrospray ionisation) and DART (direct analysis in real time) facilitate the desorption and ionization from the surface of samples at atmospheric conditions, without dissolution or additional sample preparation. They are therefore potentially useful tools for rapid assessment of solid oral dosage forms

Optical spectroscopy techniques offer rapid, non-destructive analysis, including polymorphic identification [3], and are also able to measure insoluble ingredients. They have consequently been exploited for in-line process analytical testing and as quality control tools [1]. For example, infra-red-based techniques (FTIR and DESI [4]; and NIR and Mid-IR spectroscopy classification of MDMA containing tablets [5]) Near infrared is the most commonly used process analytical tool [1], however Raman spectroscopy provides complementary information and has grown in popularity in recent

years, since it provides more distinct spectral features, and is better-suited to analysis in aqueous environments owing to the relatively weak strength of the Raman O-H band. Technological advancements have facilitated miniaturization, increased speed and reduced cost, resulting in more widespread implementation. [6]

## Ambient ionisation mass spectrometry of tablets

Desorption electrospray ionization uses a charged electrospray of organic solvent which, when directed at the sample surface in proximity to the mass spectrometry inlet, desorbs ions from the sample which may be taken up into a mass spectrometer [7]. As this process takes place at ambient pressure and with a flexible geometry, the technique is suited for the analysis of a wide range of samples including explosives on surfaces [8], fingerprints [9], plants [10] and tissues [11,12]

One of the early descriptions of DESI-MSI was in the profiling of tablets [13]. Chen et al demonstrated the use of DESI-MS to profile tablets containing loratadine, folic acid, acetaminophen (paracetamol), aspirin, melatonin or caffeine. Optimization of DESI parameters including voltage, solvent delivery and capillary temperature facilitated analysis at up to three scans per second. Subsequent studies using DESI-MS of tablets have focused on targeted analysis for active ingredients. For example, the identification MDMA and amphetamine derivatives in ecstasy tablets [14], counterfeit artesunate antimalarial tablets [15,16] and antiviral capsules [17].

For ambient mass spectrometry to be deployable in the field for counterfeiting applications, or in manufacturing environments for QA/QC, the mass spectrometer must be compact. Several designs for small field-deployable mass spectrometers have been demonstrated with DESI MS sources [18,19].

Each of these applications has targeted expected components of the tablet of interest, predominantly active ingredients or excipients. However, in manufacturing QA/QC and counterfeit-detection applications, additional information on unexpected changes in active or excipient source or quality, as well as the introduction of contaminants may be of importance. Untargeted multivariate and machine learning approaches are therefore of interest to determine differences between samples using all spectral information.

Classification approaches for mass spectrometry applications are proving powerful in a range of applications. The two most widespread applications of classification in mass spectrometry are in disease diagnosis and determination of bacterial type [20]. A range of classification algorithms have been applied to mass spectrometry and spectroscopy data. PLS-DA is most commonly reported, although a range of algorithms including neural networks, and support vector machines [21] have been reported. Several publications have evaluated different classification algorithms but unsurprisingly the optimal algorithm depends greatly on the nature of the input data. A summary of classification and other data analysis for proteomics can be found here [22] and for metabolomics here [23]. Classification approaches are becoming more accessible through modeling tools with consistent grammar and data structure, and their integration into mass spectrometry software [SciLSLab, Waters software] [23].

Notably, classification of rapid evaporative ionization MS enables real-time classification of tissue types during surgery [24]. Classification of REIMS data has also found applications in food security [25] and bacterial speciation [26]. Classification approaches have also been widely employed in mass spectrometry imaging data, particularly in the classification of cancerous tissue [27]

## Raman spectroscopy analysis of tablets

Raman spectroscopy exploits the inelastic scattering of light by the sample to reveal valuable chemical and structural information. Information can be obtained from the sample in a non-destructive manner, making it a popular process analytical technology tool. Raman spectroscopy can be performed in a variety of sampling configurations/geometries, the most appropriate depending on the application. Confocal Raman microscopy can provide detailed chemical mapping with high spatial resolution, however this is generally reserved for forensic investigation rather than continuous monitoring, since it requires lengthy acquisition times. Sub-sampling issues associated with conventional backscattered Raman can be overcome by strategies such as sample rotation in conjunction with spectral averaging, or simultaneous wide angle illumination [28].

Matousek et al demonstrated the ability of transmission Raman spectroscopy to probe deep into turbid materials such as pharmaceutical tablets and provide information on their bulk properties [29,30]. In contrast to conventional backscattered Raman, in transmission Raman spectroscopy the beam passes through the full thickness of the tablet, sampling a much larger volume of the material, and consequently provides more representative sampling [31]. Although Raman scattering intensity is linear with concentration within the same confocal plane, transmission Raman signal intensity is slightly biased towards the bulk of the tablet relative to the exterior due to internal scattering [32]. In contrast, DESI-MSI sampling is biased towards the surface/coating composition. Therefore, in combination, these two techniques should provide a powerful toolkit with which to assess compositional differences between pharmaceutical tablet formulations.

Raman spectra of complex mixtures such as solid dosage forms often have complicated spectra with overlapping peaks. For this reason, multivariate techniques are often applied to help identify the components of interest and changes in chemistry. The selection and use of unsupervised and/or supervised techniques on Raman spectra rely on factors such as prior knowledge of the raw component spectra, and the quantity and complexity of the spectra.[33]

As with mass spectrometry, classification of Raman spectroscopy data has been primarily focused on disease diagnostics [34,35,36] and bacterial analysis [37,38]. Other noteworthy examples of the use of classification in Raman spectroscopy include differentiation of narcotics [39], pharmaceuticals [40], [doi:<https://www.sciencedirect.com/science/article/abs/pii/S0731708506000926?>], and counterfeit tablets [41].

There have been relatively few comparisons of different classification methods for Raman spectroscopy data. Zheng et al. compared SVM, LDA and k-nearest neighbours (KNN) methods to classify renin hypertension from Raman data from serum [42]. They found that SVM and LDA performed similarly, and both outperformed the KNN algorithm. Partial least squares (PLS) and PLS discriminant analysis are also commonly used methods in characterizing tablets, however care is required depending on the data quantity and the pre-processing performed [43]. Qun et al. tested the classification of expired drugs using PLS-DA, SVM and KNN, and reported that SVM gave the strongest performance [44]. Fransson et al tested the performance of multivariate methods including PLS, classical least squares (CLS) and multivariate curve resolution (MCR) for classification of pharmaceutical tablets [45].

## Objective

In this study we set out to explore the potential of DESI-MSI and transmission Raman spectroscopy to distinguish commercially available pharmaceutical tablets with similar or different formulations. Pairing DESI with transmission Raman was of particular interest due to their complementarity and relative abilities to sensitively probe the surface vs the bulk of the tablets. Classification of tablets based on both active ingredients and excipients has the potential to be used for in-line quality control measures during pharmaceutical manufacturing, and for rapid counterfeit testing. As such we have

tested a range of classification algorithms on their capability to differentiate these tablets using a range of pre-processing methods to determine the best approaches to use in different applications.

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Right-aligned text

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1. Ordered list item
2. Ordered list item
  - a. Sub-item
  - b. Sub-item
    - i. Sub-sub-item
3. Ordered list item
  - a. Sub-item

- List item
- List item
- List item

subscript: H<sub>2</sub>O is a liquid

superscript: 2<sup>10</sup> is 1024.

[unicode superscripts](#)<sup>0123456789</sup>

[unicode subscripts](#)<sub>0123456789</sub>

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Putting each sentence on its own line has numerous benefits with regard to [editing](#) and [version control](#).

Line break without starting a new paragraph by putting two spaces at end of line.

## Document organization

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Document section headings:

# Heading 1

## Heading 2

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### Heading 3

#### Heading 4

##### Heading 5

###### Heading 6

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Horizontal rule:

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Heading 1's are recommended to be reserved for the title of the manuscript.

Heading 2's are recommended for broad sections such as *Abstract*, *Methods*, *Conclusion*, etc.

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Multiple citations can be put inside the same set of brackets [\[46,50,52\]](#). Manubot plugins provide easier, more convenient visualization of and navigation between citations [\[47,48,52,53\]](#).

Citation tags (i.e. aliases) can be defined in their own paragraphs using Markdown's reference link syntax:

## Referencing figures, tables, equations

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Figure [1](#)

Figure [2](#)



Figure [3](#)

Figure [4](#)

Table [1](#)

Equation [1](#)

Equation [2](#)

## Quotes and code

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Quoted text

Quoted block of text

Two roads diverged in a wood, and I—  
I took the one less traveled by,  
And that has made all the difference.

Code `in the middle` of normal text, aka `inline code`.

Code block with Python syntax highlighting:

```
from manubot.cite.doi import expand_short_doi

def test_expand_short_doi():
    doi = expand_short_doi("10/c3bp")
    # a string too long to fit within page:
    assert doi == "10.25313/2524-2695-2018-3-vliyanie-enhansera-copia-i-
insulyatora-gypsy-na-sintez-ernk-modifikatsii-hromatina-i-
svyazyvanie-insulyatornyh-belkov-vtransfetsirovannyh-geneticheskikh-
konstruktsiyah"
```

Code block with no syntax highlighting:

```
Exporting HTML manuscript
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```

## Figures

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**Figure 1: A square image at actual size and with a bottom caption.** Loaded from the latest version of image on GitHub.



**Figure 2: An image too wide to fit within page at full size.** Loaded from a specific (hashed) version of the image on GitHub.



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**Figure 4: A vector `.svg` image loaded from GitHub.** The parameter `sanitize=true` is necessary to properly load SVGs hosted via GitHub URLs. White background specified to serve as a backdrop for transparent sections of the image.

## Tables

**Table 1:** A table with a top caption and specified relative column widths.

<i>Bowling Scores</i>	Jane	John	Alice	Bob
Game 1	150	187	210	105
Game 2	98	202	197	102
Game 3	123	180	238	134

**Table 2:** A table too wide to fit within page.

	Digits 1-33	Digits 34-66	Digits 67-99	Ref.
pi	3.14159265358979323 846264338327950	28841971693993751 0582097494459230	78164062862089986 2803482534211706	<a href="#">piday.org</a>
e	2.71828182845904523 536028747135266	24977572470936999 5957496696762772	40766303535475945 7138217852516642	<a href="#">nasa.gov</a>

**Table 3:** A table with merged cells using the `attributes` plugin.

	Colors	
Size	Text Color	Background Color
big	blue	orange
small	black	white

## Equations

A LaTeX equation:

$$\int_0^\infty e^{-x^2} dx = \frac{\sqrt{\pi}}{2} \tag{1}$$

An equation too long to fit within page:

$$x = a + b + c + d + e + f + g + h + i + j + k + l + m + n + o + p + q + r + s + t + u + v + w + x + y + z + 1 + 2 + 3 + 4 + 5 + 6 + 7 + 8 + 9 \tag{2}$$

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