A practical guide to
METHAMPHETAMINE CHARACTERIZATION / IMPURITY PROFILING:
Method procedures, mass spectral data of selected impurities, and literature references

prepared by
Scientific Section
Division for Operations and Analysis
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1. INTRODUCTION

1.1. Background

Clandestine manufacture, trafficking and abuse of methamphetamine and the involvement of large-scale organized criminal groups in these activities, are increasing around the world, particularly in East and South East Asia, and North America. With ever larger consignments of clandestinely manufactured methamphetamine being intercepted, law enforcement authorities require enhanced capacity to identify the sources of supply of those drugs and to establish trafficking routes / distribution patterns and conspiracy links. A tool which adds valuable, scientific information in support of law enforcement intelligence gathering and operational work is drug characterization / impurity profiling, i.e., the systematic characterization of seized drug samples by physical and chemical means.

Worldwide, characterization / impurity profiling of seized drugs is increasingly viewed as a valuable complement to routine law enforcement investigative work. Chemical links between samples can be established, material from different seizures can be classified into groups of related samples, and the origin of samples can be identified. This information can be used for evidential purposes, or it can be used as a source of more general intelligence to identify trafficking patterns and distribution networks. Drug characterization / impurity profiling may also assist in the identification of output from new illicit laboratories, and in the monitoring of common methods used for drug manufacture, which, in turn, may provide information helpful to other intelligence gathering tools, for instance in precursor monitoring programmes. Finally, drug characterization studies may provide supportive evidence in cases where a differentiation of illicitly manufactured drugs from those diverted from legitimate sources is required.

1.2. UNDCP activities

In response to a mandate by the Commission on Narcotic Drugs (CND)\(^1\), which recognized the need for a cohesive international strategy in this field, UNDCP’s Scientific Section (Laboratory) has in recent years devoted resources to developing standard “methods for the profiling / signature analysis of key narcotic drugs and psychotropic substances”. Activities are aimed at developing methods for the characterization / impurity profiling of those substances, at supporting basic research to assist in the interpretation of analytical results, and at assisting in the development of operational capacity, at national and regional levels, in drug and precursor characterization / impurity profiling. Work has concentrated so far on methamphetamine and its main precursor ephedrine, substances which were also specifically addressed in the action plan against illicit manufacture, trafficking and abuse of amphetamine-type stimulants (ATS) and their precursors, as endorsed by the 20th special session of the United Nations General Assembly (UNGASS) in June 1998. The geographic focus has been South East Asia, a region which is particularly affected by clandestine manufacture, trafficking, and abuse of methamphetamine.

Impurity profiling methods for methamphetamine and ephedrine have been developed by UNDCP. Analysis of samples using the methods developed has enabled the identification and/or confirmation of new trends in illicit manufacture, and the development of operational intelligence by law enforcement personnel in the countries concerned. As a result, more and more countries have shown an interest in the potential of drug characterization / impurity profiling work. The need to develop appropriate tools and operational programmes in this field

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\(^1\) CND Resolution I (XXXIX), adopted at CND 39th session, 1996.
was recently also agreed by participants at a sub-regional technical meeting\(^2\) in Bangkok, Thailand, which discussed, in an operational framework, concepts, value and methods of methamphetamine characterization / impurity profiling.

1.3. Purpose of publication

The present note is intended as a practical guide for laboratories interested to embark on profiling activities\(^3\). It provides details of the method procedures developed by UNDCP\(^4\), mass spectral data of selected impurities found in seized samples of methamphetamine and ephedrine, as well as selected literature references. To facilitate their practical application, method procedures are presented with a high level of detail, i.e., in the form of standard operating procedures (SOP). It should be understood, however, that the exact methods and procedures depend on (i) the intended utilization of results, and (ii) the availability and specifications of the analytical equipment and data processing software available.

Finally, it is important to recognize that even in the absence of sophisticated analytical equipment a substantial scientific contribution to law enforcement operational work can be made on the basis of the systematic characterization of seized drug samples, i.e., the systematic comparison of samples based on their physical characteristics. This applies particularly to samples in tablet form. To facilitate the building of appropriate databases in this field, at both national and regional levels, a tablet identification sheet, which was agreed on at the Bangkok meeting, is also included as a model for the collection of standardized data.

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\(^2\) Sub-regional meeting on drug characterization / impurity profiling and its investigational value for law-enforcement authorities; special focus: methamphetamine in South East Asia, Bangkok, 6-8 June 2000.

\(^3\) See also UN publication entitled “Drug characterization / impurity profiling: background and concepts” (ST/NAR/32).

\(^4\) The method described has been tested and used by collaborating laboratories.
2. EXPERIMENTAL PART

2.1. Purpose

The method described below was developed by UNDCP’s Scientific Section as a means for the characterization / impurity profiling of seized samples of methamphetamine\(^5\). It includes: (i) liquid/liquid extraction of neutral / basic impurities from methamphetamine, and gas chromatographic analysis of extracted impurities (“impurity profiling”), (ii) establishment of calibration curves and determination of the contents (“purity”) in illicit samples of methamphetamine hydrochloride, and (iii) data handling and analysis.

Criteria for method development included (i) universality, i.e., the applicability of the method to methamphetamine samples synthesized via different synthesis routes, (ii) simplicity in extraction and analytical techniques, (iii) repeatability and reproducibility, and (iv) a focus on methamphetamine synthesis by-products which are mostly basic or neutral substances, as opposed to diluents, cutting agents, and substances carried over from starting materials or preparations thereof.

To facilitate the building of appropriate databases for methamphetamine tablets a model tablet identification sheet is also included (see Part 3 below).

2.2. Method procedures

2.2.1. Compressed Gases

Nitrogen is employed as carrier gas (>99.999% pure). FID gases are synthetic air (20.5 % \(O_2\) in \(N_2\)) and hydrogen (>99.999% pure).

2.2.2. Solvents, standard substances

- ethyl acetate, spectroscopic grade
- \(n\)-hexane, for chromatography
- phosphate buffer (buffer solution ready for use, pH 7.00, ± 0.02, at 20°C)
- diphenylamine
- (+)-methamphetamine hydrochloride
- \(n\)-tridecane
- \(n\)-tetracosane

2.2.3. Solutions

2.2.3.1. Buffer solution (pH 10.5)

a) 10% (w/v) \(Na_2CO_3\): Weigh 10.0g of \(Na_2CO_3\) in a 100ml volumetric flask. Dissolve in a small amount of distilled water. Fill with distilled water to the mark and shake. Label flask as to date and concentration. Discard solution after six months\(^6\).

b) Buffer solution (pH 10.5): Add 4 parts of phosphate buffer (pH 7) to one part of 10% (v/v) \(Na_2CO_3\). Check pH, adjust to pH 10.5 by adding 10% \(Na_2CO_3\), if

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\(^5\) Necessary adjustments to this method for the impurity profiling of ephedrine samples are also included.

\(^6\) Note: The amount of any of the described solutions to be prepared depends on the sample throughput of individual laboratories. As a general rule, if properly stored, solutions which are not used for quantitation (including aqueous buffer solutions and retention time reference standards) can be used for a maximum of six months, all other solutions should be discarded after three months.

-3-
necessary. Solution can be stored at room temperature in a glass stoppered flask. Label flask as to date and concentration. Discard solution after six months.

2.2.3.2. Instrument performance solution (‘test mixture’)
Weigh out 7.0 mg (±5%) of (+)-methamphetamine hydrochloride into a 5 ml screw-capped glass tube, and add 1 ml of buffer (pH 10.5). Screw up tube and place it on the platform of a circular shaker. Shake for 5 minutes. Add 2 ml of n-hexane using a volumetric glass pipette. Shake for another 5 minutes. Centrifuge for 5 minutes at 3000 rpm and room temperature (approx. 21°C). Transfer 1.0 ml of the n-hexane phase into a 100ml glass stoppered volumetric flask using a volumetric glass pipette. Add 3.50 mg (±5%) of diphenylamine and 2.50 mg (±5%) of n-tetracosane. Record actual weight to the tenth of a milligram. Dilute to the mark with hexane. Shake final solution. Concentrations obtained: approx. 35mg/l of (+)-methamphetamine hydrochloride, 35mg/l of diphenylamine and 25mg/l of n-tetracosane.
Label the flasks as to date and concentrations. Seal with Parafilm™, and store in the refrigerator. Discard solution after three months. When removed from the cold, fill an autosampler vial, and let the solution equilibrate to ambient temperature.

2.2.3.3. Internal standard solution for methamphetamine profiling (‘ISTD-prof’)

a) **Stock internal standard solutions**: Weigh out 25.0 mg (±5%) of n-tridecane (C_{13}), 35.0 mg (±5%) of diphenylamine and 20.0 mg (±5%) of n-tetracosane (C_{24}) into separate 10 ml volumetric flasks using an analytical balance. Record the actual weight to the tenth of a milligram. Dissolve each substance in a small amount of ethyl acetate. Dilute with ethyl acetate to the mark of each volumetric flask. Shake final solutions. Label each stock solution as to date and concentration. Discard solutions after three months. When not in use, stopper, seal the top of the flasks with Parafilm™, and store in refrigerator.

b) **Internal standard solution (‘ISTD-prof’)**: Remove the stock internal standard solutions prepared above from the cold and allow them to equilibrate to ambient temperature. Transfer as much of each stock internal standard solution into a 100ml glass stoppered volumetric flask as to get the following concentrations: n-C_{13} at 25 mg/l, diphenylamine at 35 mg/l and n-C_{24} at 20 mg/l. Dilute with ethyl acetate to the mark. Shake final solution. Label the flask as to date and concentration. Discard solution after three months. When not in use, stopper, seal the top of the flask with Parafilm™, and store in refrigerator. Before use, when removed from the cold, allow the solution to equilibrate to ambient temperature.

2.2.3.4. Internal standard solution for calibration curve / methamphetamine purity determinations (‘ISTD-purity’)
Weigh out 50.00 mg (±5%) of diphenylamine into a glass stoppered 100ml flask using an analytical balance. Record the actual weight to the hundredth of a milligram. Dissolve in a small amount of ethyl acetate. Dilute with ethyl acetate to the mark and shake. Label solution as to date and concentration. Discard solution after three months. When not in use, stopper, seal the top of the flask with Parafilm™, and store in refrigerator. Before use, when removed from the cold, allow the solution to equilibrate to ambient temperature.

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7 The two hydrocarbon ISTDs are used to bracket the impurity peaks of interest, to enable synchronization of the time axis of different chromatograms.
2.2.3.5. Methamphetamine calibration standards

a) **Methamphetamine stock solution**: Weigh out 30.00 mg (±5%) of (+)-methamphetamine hydrochloride into a 15 ml screw-capped glass vial using an analytical balance. Record the actual weight to the hundredth of a milligram. Add 1 ml of buffer solution (pH 10.5). Screw up tube and place it on the platform of a circular shaker. Shake for 5 min. Add 12 ml of ‘ISTD-purity’ solution. Shake for another 5 min. Centrifuge sample for 5 min at 3000 rpm and room temperature (approx. 21°C). Transfer 10 ml of organic layer into a screw-capped glass vial using a 10ml volumetric pipette. Label flask as to date and concentration of methamphetamine hydrochloride (approx. 2.5 mg/ml). Discard solution after preparation of the different calibration standards.

b) **Methamphetamine calibration standards**: Transfer the required amount of stock solution into a 1 ml volumetric flask using a 100 µl-syringe. Add ‘ISTD-purity’ solution using the same syringe (i.e., wash it into flask), then dilute with ‘ISTD-purity’ solution to the mark and shake. Each concentration is prepared directly from the stock solution. Keep calibration standards for the period of use of the corresponding calibration curve (see 2.5.1, below).

2.2.4. Apparatus and equipment

2.2.4.1. Gas chromatograph
The gas chromatograph (GC) is equipped with a GC autosampler controller, an autosampler tray, an automatic injector, and a flame ionization detector (FID).

A fused silica capillary column, 25 m x 0.2 mm x 0.33 µm, crosslinked, 5% phenylmethylsilicone, is used for analysis.

Split/splitless injection port liners are used. To avoid any loss of chromatographic performance, liners should be continuously checked for contamination and replaced at least once a month. Used liners can be cleaned with chromic sulphuric acid (by immersion overnight). They are then rinsed with distilled water and dried. Silanization is achieved by immersion overnight in a solution of 5% DMDCS (dimethyldichlorosilane) in toluene. Liners are then rinsed twice with toluene, and immediately afterwards with absolute methanol. They are dried under clean nitrogen.

The septum should be replaced after approx. 100 injections.

2.2.4.2. Vials
5ml screw-capped glass vials with 1.2 cm outer diameter\(^8\) are used for the extraction procedures (methamphetamine and ephedrine samples for impurity profiling).

Clear 2 ml autosampler vials with microvolume glass inserts (100 µl, with polymer support feet) and plastic open-top screw-caps with silicone / teflon septa are used for GC analysis.

15 ml screw-capped glass vials are used for the preparation of the methamphetamine stock solution for calibration standards.

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\(^8\) A small diameter of the tube is desirable as it determines the height (and thus, the ease of recovery) of the organic phase.
2.2.4.3. Analytical balance
The balance used has a readability of 0.1 mg and 0.01 mg, respectively. The balance is calibrated daily using the manufacturer’s internal calibration procedure.

2.2.4.4. Shaker
The shaker is a circular shaker with a foam-top platform.

2.2.4.5. Other
Automatic pipette with 5 ml and 2.5 ml tips.
Disposable Pasteur pipettes, 20 cm.
pH Indicator stripes (pH 7.5-14 and pH 5.0-10.0, respectively).

2.2.5. Instrument parameters
2.2.5.1. Gas flows
The FID gases air and hydrogen are set to 2.5 bar (±5%) and 1.4 bar (±10%), respectively. The corresponding flow rates are approx. 300 ml/min and 30 ml/min, respectively (measured with a bubble flow meter).

The carrier gas nitrogen is set to 150 kPa (approx. 22 psi), corresponding to a column flow of 1.4 ml/min at 50°C oven temperature (measured with a bubble flow meter). The total nitrogen flow (carrier gas plus make-up gas) is 30 ml/min.

2.2.5.2. Injection mode
Splitless injection is used, the valve is closed for 1 min. Split vent is set to 30 ml/min.

2.2.5.3. Parameters for automatic injector
Sample washes 1
Sample pumps 5
Injection volume 1.0 µl
Syringe size 10.0 µl
Post injection washes 6
Viscosity delay 2 seconds
Plunger speed fast

2.2.5.4. Temperature programming
Injector temperature 250°C
Detector temperature 300°C
Initial temperature 50°C (for 1 min)
Final temperature 300°C (for 4 min)
Temperature rate 10°C/min
Oven equilibrium time 2 min
The total run time is 30 min.

2.2.5.5. Signal parameters
Peak width 0.013
Sampling rate 10 Hz
Signal plot 10% offset
2.2.5.6. Sequence of injection

- solvent (instrument blank)
- reference (check) sample in duplicate \(^9\) (i.e., replicate extractions)
- 2 method blanks (i.e., replicate blank extractions) \(^10\)
- 1 sample in duplicate \(^11\) (i.e., replicate extractions of the same sample)
- solvent (instrument blank)
- 1 sample in duplicate
- solvent (instrument blank)
- 1 sample in duplicate
- solvent (instrument blank)
- reference (check) sample
- solvent (instrument blank)
- 1 sample in duplicate
- solvent (instrument blank)
- 1 sample in duplicate
- solvent (instrument blank)
- 1 sample in duplicate
- solvent (instrument blank)
- reference (check) sample

This sequence is recommended provided that proper instrument performance (i.e., checking by injection of 'test mixture') is ensured before starting the sequence (see 2.3. below).

The maximum number of samples (including blanks, check samples, etc.) should ideally not exceed 26 \(^12\).

2.3. Procedure to check instrument performance

2.3.1. To check basic column performance, perform analysis of the 'test mixture' daily, for example, each time after turning on the GC.

Shewhart charts (see Figure 1) are created based on date and peak area ratios of methamphetamine / diphenylamine and n-C\(_{24}\) / diphenylamine, respectively. The standard deviation (STDEV) is determined based on the mean of 10 consecutive injections of the instrument performance solution ('test mixture') on the same day. The warning limit of instrument performance is within two STDEVs, the action limit within 3 STDEVs of the area ratios of

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\(^9\) A reference (check) sample is prepared from a stock of methamphetamine sample available in sufficient quantity to allow for repeated analyses over a prolonged period of time to ensure comparability of results.

\(^10\) As part of every sequence, two method blanks should be prepared and analyzed. To this end, steps 2.4.2 to 2.4.8 (see below) should be carried out, without methamphetamine being present.

\(^11\) Two injections of the same sample without a solvent run in between require that it has been previously determined that none of the impurities of interest causes a memory effect. Memory of methamphetamine itself does not pose a problem since this peak is of no interest in the comparison of impurity profiles.

\(^12\) Sequences longer than 18 hours should be avoided as samples late in the sequence, kept at room temperature, may degrade.
methamphetamine / diphenylamine, and n-tetracosane / diphenylamine, respectively. The acceptance criteria are drawn as follows (see Figure 1):
- warning limit: \( \text{mean} \pm 2 \text{ STDEV} \)
- action limit: \( \text{mean} \pm 3 \text{ STDEV} \).

If acceptance criteria are not met, the column must be conditioned (see 2.3.3. below) or a new column used.

2.3.2. To check specific instrument performance (i.e., suitability of instrument status for methamphetamine profiling), perform analysis of a methamphetamine check sample as part of every sequence, as described in 2.2.5.6. Repeated GC impurity profiles of the check sample have to show similarities, based on the same criteria as overall sample comparison, which are acceptable within the chosen comparison criteria (i.e., they should show the closest similarities).

![Figure 1](image-url)

Figure 1. ‘Test mixture’ analyzed 10 times on the same day. The upper and lower lines bracket the range of acceptance (warning limit and action limit, respectively) of the area ratio of methamphetamine / diphenylamine.

2.3.3. If the instrument performance solution (‘test mixture’) does not meet the acceptance criteria (action limit), especially after a change of the column, the new column can be conditioned using injections of pyridine, until criteria are met. The ultimate acceptance criterion is the repeatability of the check sample (see 2.3.2 above).
2.4. Extraction of methamphetamine samples for impurity profiling

2.4.1. Homogenize illicit sample material in a mortar and weigh 30.0 mg (±5%) of illicit sample material into a 5 ml screw-capped glass tube using an analytical balance. Record the actual weight to the tenth of a milligram.

2.4.2. Add 1 ml of buffer solution (pH 10.5) using an automatic pipette with a 5 ml tip.

2.4.3. Screw up tube, and place it on the platform of a circular shaker. Shake the sample for 5 min. Check and record pH with pH indicator stripes (pH 7.5-14), record colour of water phase, and the appearance of a precipitate or insoluble material, if any.

2.4.4. Add 200 µl of 'ISTD-prof' solution using an automatic pipette with a 2.5 ml tip. Screw up.

2.4.5. Place tube on the shaker platform, and shake it for another 5 min. Record colour of organic phase.

2.4.6. Centrifuge for 5 min at 3000 rpm and room temperature (approx. 21°C).

2.4.7. Transfer the organic layer with a disposable Pasteur pipette into a 2 ml autosampler vial with glass insert, and close it with silicone/teflon caps.

2.4.8. Analyze the sample on the same day of extraction.

2.5. Determinations of drug content (“purity”) 14

Calibration standards should be prepared so that they cover the range of concentrations expected in the unknown samples. It is useful to prepare at least 6 calibration standards (six-point calibration) across a range of 50% to 150% of the expected sample concentration. 15 Each calibration point is calculated based on two injections of each standard. The calibration should be linear across the range of likely use, i.e., correlation coefficient > 0.999. The calibration points are constructed by calculating the amount ratio (concentration of component divided by concentration of internal standard) and response ratio (area of component divided by area of internal standard). The equation for the curve through the calibration points is calculated using the following type of curve fit:

- Curve type: linear
- Origin: included
- Weight: equal

2.5.1. Extraction of methamphetamine samples for drug content determinations

Weigh out approx. 2.50 mg (±5%) of seized methamphetamine into a 5 ml screw-capped glass tube using an analytical balance. Record the actual weight to the hundredth of a milligram. Add 2 ml of buffer solution (pH 10.5) using an automatic pipette with a 5 ml tip. Screw up tube, and place it on the platform of a circular shaker.

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13 A suitable extraction procedure for ephedrine samples is as follows:

- 100 mg homogenized sample material
- + 1 ml buffer (pH 7.0), shake, centrifuge
- + 0.2 ml ethyl acetate containing three ISTDs ('ISTD-Prof', see 2.2.3.3 above) shake, centrifuge

Analytical conditions are the same as for methamphetamine (see 2.2.4.1 and 2.2.5 above).

14 As an easy means for the quick comparison of illicit methamphetamine samples based on the ratios of their methamphetamine, ephedrine, and caffeine 'contents', manual integration of the three peaks, and calculation of the respective peak area percentages, can be used.

15 To cover the range of concentrations of methamphetamine in illicit tablets from Thailand, calibration standards starting at a concentration of approx. 30 ng/µl, and additional standards in 150 ng/µl-steps, have proven to be suitable. For methamphetamine samples of very high purity ('ice'), a separate calibration curve should be prepared, covering a concentration range from approx. 0.6 µg/µl to 2 µg/µl.
Shake for 5 min. Add 2 ml of ‘ISTD-purity’ solution using a 2 ml volumetric glass pipette, and screw up. Place tube on the shaker platform, and shake it for another 5 min. Centrifuge for 5 min at 3000 rpm and room temperature (approx. 21°C). Transfer organic layer with a disposable Pasteur pipette into a 2ml autosampler vial, and close it with a silicone/teflon cap. Analyze the sample on the same day of extraction. Every fourth sample in the GC sequence should be a ‘purity check sample’, i.e., an aliquot of an original methamphetamine calibration standard. The quantitative results (in ng/µl) of these samples should be within ± 5% of the true value.

2.5.2 Calculation of drug content of unknown samples
The amount of methamphetamine hydrochloride (in ng/µl) is calculated based on an internal standard report. Conversion into percent (%) purity is based on the amount of methamphetamine weighed out initially. In addition, for tablets, the total amount (in mg) of methamphetamine hydrochloride is calculated.

2.6 Data handling and analysis

The sophistication of impurity data analysis can range from comparison and evaluation of a limited number of peak area ratios to multivariate data analysis, or a combination thereof. Visual comparison of chromatograms for final conclusions is recommended.

Figure 2: Impurity profile of a typical methamphetamine tablet (IS = internal standard).

Selected impurities include: (1) benzaldehyde, (2) cis-1,2-dimethyl-3-phenylaziridine, (3) amphetamine, (4) 3,4-dimethyl-5-phenyl-oxazolidine\(^{16}\), (5) ethyl vanillin (known to be frequently added as flavouring agent), (6) N-methyl-ephedrine, (7) N-formylmethamphetamine, (8) N-acetylmethamphetamine, (9) N-acetyllephedrine, and (10) methamphetamine dimer.

For MS spectra of these and other impurities found in seized samples of methamphetamine, see Part 4 below.

Note: Asterisks (*) indicate peaks which appear to be useful for the screening of impurity profiles. These peaks were characterized in terms of retention time and MS spectra, but could not yet be identified.

Figure 2 shows the impurity profile of a typical methamphetamine tablet. The selection of peaks for comparison depends on the intended utilization of results and the availability of software for data analysis. If, for example, links are to be made at the level of the source of the methamphetamine powder material (i.e., differentiating at the level of the clandestine methamphetamine laboratories), only peaks of synthesis by-products and not of cutting agents should be used. In general, the following selection criteria should be applied:

\(^{16}\) Ephedrine has the same retention time. However, the distinction between ephedrine and the oxazolidine, based on GC data only, does not pose a problem since the ephedrine peak, if present, is usually quite large. In none of the methamphetamine tablet samples examined, ephedrine was present in concentrations approaching the impurity level.
- high frequency of occurrence in multiple samples (i.e., relevance of peak);
- no artifacts;
- no co-eluting peaks, i.e., peaks should be single compounds;
- good repeatability (ideally: CV < 5-10%); therefore, highly volatile compounds at the beginning of the chromatogram should be avoided;
- if possible, selected peaks should be linked to synthesis routes.

In general, for GC data comparison, both relative retention times and relative peak areas (ratios) should be used. Normalization of peak heights is another suitable option. The selection of integration events should involve the use of a representative, seized methamphetamine sample containing relevant impurities in usually encountered concentrations (e.g., the reference (check) sample). It should follow common integration practice and the GC instrument manufacturer’s procedure.

The establishment of a database on physical (and relevant chemical) data as well as background information is recommended.
3. MODEL TABLET IDENTIFICATION SHEET
# TABLET IDENTIFICATION SHEET

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Lab Code No.: | Case No.:  

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<td>Orange</td>
<td>Yellow</td>
<td>Green</td>
<td>Blue</td>
<td>Violet/Purple</td>
<td>Grey</td>
<td>Brown</td>
</tr>
<tr>
<td>31</td>
<td>32</td>
<td>33</td>
<td>34</td>
<td>35</td>
<td>36</td>
<td>37</td>
<td>38</td>
<td></td>
</tr>
<tr>
<td>Pink</td>
<td>Black</td>
<td>Colourless</td>
<td>Silver</td>
<td>Gold</td>
<td>Bicoloured</td>
<td>Multicoloured</td>
<td>Undecided</td>
<td></td>
</tr>
</tbody>
</table>

### SCORING

<table>
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<th>53</th>
<th>54</th>
<th>55</th>
<th>56</th>
<th>57</th>
<th>58</th>
<th>59</th>
<th>60</th>
<th>61</th>
<th>62</th>
</tr>
</thead>
<tbody>
<tr>
<td>Unscored</td>
<td>½ scored, no mark</td>
<td>½ scored, mark same side</td>
<td>½ scored, mark other side</td>
<td>½ scored, mark both sides</td>
<td>¼ scored, no mark</td>
<td>¼ scored, mark same side</td>
<td>¼ scored, mark other side</td>
<td>¼ scored, mark both sides</td>
<td>Undecided</td>
</tr>
</tbody>
</table>

### Detailed description:

**SIZE**

<table>
<thead>
<tr>
<th>Diameter</th>
<th>Thickness</th>
</tr>
</thead>
<tbody>
<tr>
<td>mm</td>
<td>mm</td>
</tr>
</tbody>
</table>

**WEIGHT**

| mg |

**MARKING**

**DRUG CONTENT**

| % | mg |

**Other observations (e.g., smell):**
4. MASS SPECTRAL DATA OF SELECTED IMPURITIES FOUND IN SEIZED SAMPLES OF METHAMPHETAMINE
(Mass spectral data sorted by molecular weight)

N-methylbenzylamine
C8H11N
MW = 121

trans-1,2-dimethyl-3-phenylaziridine
C10H13N
MW = 147

cis-1,2-dimethyl-3-phenylaziridine
C10H13N
MW = 147

N,N-dimethyl-amphetamine
C11H17N
MW = 163

N-formyl-amphetamine
C10H13NO
MW = 163
erythro-3,4-dimethyl-5-phenyl-oxazolidine
C$_{11}$H$_{15}$NO
MW = 177

N-acetyl-amphetamine
C$_{11}$H$_{15}$NO
MW = 177

N-formyl-methamphetamine
C$_{11}$H$_{15}$NO
MW = 177

(1R,2S)-(-)-N-methylephedrine
C$_{11}$H$_{17}$NO
MW = 179

1-chloro-1-phenyl-2-methylaminopropane
synonym: chloro-ephedrine
C$_{10}$H$_{14}$ClN
MW = 183
N-acetyl-methamphetamine
C12H17NO
MW = 191

N-acetyl-ephedrine
C12H17NO2
MW = 207

N-methyl-2-(1,3-diphenyl)propylamine
synonym: ɣ-benzyl-N-methylphenethylamine
C16H19N
MW = 225

dibenzyldiketone
C16H190
MW = 227

1,3-dimethyl-2-phenyl-naphthalene
synonym: naphthalene A
C18H16
MW = 232
1-benzyl-3-methylnaphthalene
synonym: naphthalene B
C18H16
MW = 232

N-methyl-di-β-phenethylamine
C17H21N
MW = 239

benzfetamine
synonym: N-benzyl-methamphetamine
C17H21N
MW = 239

N,O-diacetyl-ephedrine
C14H19NO3
MW = 249

N,N-di(β-phenylisopropyl)-amine, diastereomer
synonym: diphenylisopropylamine (DPIA)
C18H23N
MW = 253
N-erythro-3,4-dimethyl-2,5-diphenyl-oxazolidine
C_{17}N_{19}O
MW = 253

N-benzyl-ephedrine
C_{17}H_{21}NO
MW = 255

N,N-di(β-phenylisopropyl)-methylamine, diastereomer
synonym: diphenylisopropylmethylamine (DPIMA)
C_{19}H_{25}N
MW = 267

N,N-di(β-phenylisopropyl)-formamide, diastereomer
synonym: N-formyl-diphenylisopropylamine
C_{19}H_{21}NO
MW = 279
N-methyl-N-(α-methylphenethyl)amino-1-phenyl-2-propanone
C19H23NO
MW = 281

methamphetamine dimer
C20H28N2
MW = 296
5. SELECTED BIBLIOGRAPHY

5.1. Tablet characterization, dyes
5.2. Methamphetamine (general)
   5.2.1. Methamphetamine from ephedrine
   5.2.2. Methamphetamine from P2P
5.3. Amphetamine
5.4. Ephedrine / pseudoephedrine
5.5. 1-Phenyl-2-propanone (P2P)
5.6. Miscellaneous chemistry
5.7. Data analysis
5.8. Other (including profiling and clandestine laboratories)
5.1. Tablet characterization, dyes


5.2. Methamphetamine (general)


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5.7. Data analysis


5.8. Other (including profiling and clandestine laboratories)


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