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Comparison of six derivatizing agents for the determination of nine synthetic cathinones using gas chromatography-mass spectrometry

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Six acylation reagents have been compared for their derivatisation potential towards nine synthetic cathinones by gas chromatography-mass spectrometry (GC-MS). The evaluated reagents were pentafluoropropionic anhydride (PFPA), trifluoroacetic anhydride (TFA), chlorodifluoroacetic anhydride (CLF₂AA), heptafluorobutyric anhydride (HFBA), acetic anhydride (AA) and propionic anhydride (PA). The synthetic cathinones included flephedrone (4-fluoromethcathinone or 4-FMC), mephedrone (4-methylmethcathinone or 4-MMC), pentedrone (also known as α -methylamino-valerophenone), methedrone (4-methoxy-*N*-methcathinone, *p*-methoxymethcathinone), methylone (3,4-methylenedioxy-*N*-methylcathinone or bk-MDMA), butylone (β -keto-*N*-methylbenzodioxolylbutanamine or bk-MBDB), ethylone (3,4-methylenedioxy-*N*-ethylcathinone MDEC or bk-MDEA), pyrovalerone (4-methyl- β -keto-prolintane) and 3,4-methylenedioxyprovalerone (MDPV). The derivatizing agents were optimised for incubation time and temperature with some important validation parameters studied to evaluate derivatisation reactions. The anhydrides studied proved to be suitable for synthetic cathinones – all of them showing RSD and accuracy below 20%. PFPA and HFBA followed by TFA are the best choice of derivatising agents based on validation parameters. Five internal standards were evaluated with good results. Three way ANOVA, interference, fragmentation patterns and high peak area values at a concentration of 0.50 $\mu\text{g ml}^{-1}$ were evaluated and discussed. AA and PA derivatives give high relative abundance for most drugs examined. HFBA gives more ions and multi-fragmentation patterns.

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Introduction

Cathinone (β -Keto-amphetamine) is found in the leaves of the *Catha edulis* (Khat) plant.¹ Synthetic cathinones have similar properties to other stimulants and hallucinogenic drugs including amphetamines and ring-substituted amphetamines. They have pharmacological effects, known as “cardiovascular and neurological side-effects”, and can cause deaths, many of which have occurred in Europe.^{2–5} Despite the misuse potential associated with these compounds, the legislation governing their use is not consistent worldwide. Additionally, due to chemical modifications, these new psychoactive substances (NPS) are rapidly altered to produce new variants to bypass drug legislations of a particular country^{6,7} with detection and identification of these drugs proving difficult because of a lack of reference standards.⁸

A review by Zuba and colleagues discussed pathways and unknown structures of cathinones based on mass spectrometry⁹

with the isomers of substituted cathinones having been investigated using NMR spectroscopy by Kavanagh.¹⁰ More than 70 synthetic cathinones have been reported by the European Monitoring Centre for Drugs and Drug Addiction (EMCDDA) with synthetic cathinones making up the second biggest portion of NPS identified in 2015.^{11,12}

The GC analysis of cathinones generally requires the use of derivatising reagents. To select an appropriate derivatization reagent for GC analysis, the following criteria can be used as guidance:¹³

- The reagent should generate >95% of complete derivatives.
- During derivative formation, the reagent should not alter/rearrange the structure of the compound.
- Loss of sample should not occur during the reaction.
- Derivatives produced should not interact with the GC column.
- A stable derivative should be formed.

To achieve the above goals, acylation was selected for this study instead of other common derivatization methods (*e.g.* silylation and alkylation) since it is a popular derivatising technique widely used to increase the sensitivity, produce excellent fragmentation in mass spectra, improve the chromatographic peak shape and resolution as well as reduction in the polarity of analytes. PFPA, TFA, CLF₂AA, HFBA, AA, and PA

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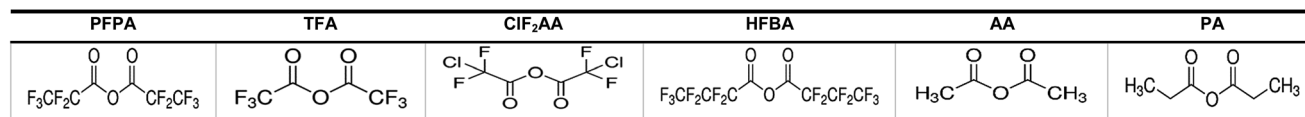


Fig. 1 Chemical structures of selected derivatisation agents.

were the chosen acylation reagents in this study (see Fig. 1 for the chemical structures of reagents).

The first problem encountered with analysis of synthetic cathinones is that during GCMS method development several cathinones have only one or two important ions in the mass spectrum. Most of the remaining ions have smaller abundance (<10%) when compared with the highest abundance ion, consequently resulting in poor detection. Secondly, some cathinones have positional isomers, which produce ambiguous mass spectra. Thirdly, MDPV and pyrovalerone are not derivatised, and the analyst is dependent on a limited number of mass ions. Fourthly, some cathinones have overlap between the high abundance ions and those from internal standards. Taking these factors into account with additional legal implications, derivatisation techniques are necessary to produce many patterns and high abundance resolution of fragmentations aiding correct identification of these compounds.

In this study, the evaluation and comparison of these reagents for selected cathinones were investigated with the focus on the following:

- Effect of time and temperature on the reaction and optimising these conditions.
- Examination of the highest values of peak areas.
- Quality of fragmentation ions *vs.* reagents.
- Quality of mass spectrum based on its relative ion intensities.
- Total ion chromatograms from interference studies.
- Data treatment analysis by running ANOVA.¹⁴
- Choice of best-fit regression between internal standards.

Some validation parameters including the LOD, linearity, accuracy, RSD, and recovery were used.

Mephedrone, flephedrone, pentedrone, methylone, ethylone, methedrone, MDPV, butylone, and pyrovalerone are the most frequently abused cathinones in the UK and in Europe and therefore selected as the target analytes¹⁵ (see Table 1).

Materials and methods

Materials

Reference standards of the nine synthetic cathinones (1 mg ml⁻¹) – flephedrone, mephedrone, pentedrone, methedrone, methylone, butylone, ethylone, pyrovalerone, and MDPV; five internal standards (0.10 mg ml⁻¹) – mephedrone-d₃, methylone-d₃, butylone-d₃, ethylone-d₅ and MDPV-d₈ as their hydrochloride salts; seven derivatization agents – pentafluoropropionic anhydride (PFPA) ≥ 99%, trifluoro-acetic anhydride (TFA) ≥ 99%, chloro di-fluoro acetic anhydride (ClF₂AA) ≥ 98%, heptafluoro-butyric anhydride (HFBA) ≥ 99%, acetic anhydride (AA) ≥ 99%, propionic anhydride (PA) ≥ 99%, and

butyric anhydride (BA) ≥ 98% were purchased from Sigma-Aldrich, Gillingham, UK. Ethyl acetate (EtOAc), methanol, ammonium hydroxide, sodium phosphate dibasic, sodium chloride, sodium phosphate monobasic, dichloromethane (DCM), isopropanol (IPA), ammonium hydroxide (NH₄OH), and acetic acid were obtained from VWR International, East Grinstead, UK. Blank blood was supplied by the Scottish National Blood Transfusion Service based at Gartnavel Hospital, Glasgow. Phosphate buffer and sodium phosphate were purchased from Fisher Scientific, Loughborough, UK. Solid phase extraction columns (200 mg clean screen® part number ZSDAU20 manufactured by United Chemical Technologies) were purchased from Chromatography Direct, Runcorn, UK.

Methods

Preparation of standards

Stock solutions (100 µg ml⁻¹) of the nine drugs were prepared by dilution of the purchased standards *via* 1 : 10 dilution in methanol. The working solution of each standard was prepared by dilution of stock solutions 100 µg ml⁻¹ *via* 1 : 10 methanol to reach 10 µg ml⁻¹. Working internal standards of the deuterated standards were similarly prepared taking into account the different concentrations of the supplied standards.

Optimisation of temperature and incubation time study

For optimisation of the reaction temperature and incubation time within the study, cathinones were derivatised at the same time in the following way: 50 µl of the 10 µg ml⁻¹ standard drug mixture and 50 µl of 2 µg ml⁻¹ of mixture of internal standards were added to samples. Then the mixture was evaporated to dryness at Room Temperature (RT) under a stream of nitrogen followed by derivatisation with 50 µl of PFPA and EtOAc (2 : 1); 50 µl of TFA and EtOAc (2 : 1); 50 µl ClF₂AA and EtOAc (2 : 1); 65 µl of HFBA and EtOAc (3 : 2), AA and EtOAc (3 : 2), and PA and pyridine (2 : 1). All samples were capped and vortexed immediately for 15 seconds and then incubated for specific times (5–10–15–20–25–30–35–40 min) and at specific temperatures (RT, 40 °C, 55 °C, 70 °C). The samples were evaporated under a stream of nitrogen with the hot block set at RT, 40 °C and 50 °C thereafter reconstituted in 50 µl of ethyl acetate. The top layer of EtOAc was transferred to an auto-sampler vial for GC-MS analysis. The GC syringe was washed three times before injection in EtOAc. A volume of 1.0 µl was injected at 225 °C and GC-MS was run under the conditions outlined below.

Samples were prepared in triplicate on eight days at concentrations of 0.50 µg ml⁻¹ and 0.10 µg ml⁻¹ for internal standards. From day one to four, 72 samples (18 samples for



Table 1 Chemical structures studied

Usual names	Chemical names	Structure	Usual names	Chemical names	Structure
Flephedrone or 4-fluoromethcathinone or 4-FMC	2-(Methylamino)-1-(4-fluorophenyl)-1-propanone		Butylone or β-keto-N-methylbenzodioxolylbutanamine (bk-MBDB)	1-(1,3-Benzodioxol-5-yl)-2-(methylamino)butan-1-one	
Mephedrone or 4-methylmethcathinone or 4-MMC	2-(Methylamino)-1-(4-methylphenyl)-1-propanone		Ethylone or bk-MDEA	2-(Ethylamino)-1-(3,4-methylenedioxyphenyl)-1-propanone	
Pentadrone or α-methylamino-valerophenone	(±)-1-Phenyl-2-(methylamino)pentan-1-one		Pyrovalerone Centroton, 4-methyl-β-keto-prolintane, Thymergix	(RS)-1-(4-methylphenyl)-2-(1-pyrrolidiny)pentan-1-one	
Methedrone or para-methoxymethcathinone, 4-methoxymethcathinone	(RS)-1-(4-Methoxyphenyl)-2-(methylamino)propan-1-one		MDPV or MDPK	(RS)-1-(Benzodioxol[1,3]dioxol-5-yl)-2-(pyrrolidin-1-yl)pentan-1-one	
Methylone or MDMA or bk-MDMA	2-Methylamino-1-[3,4-methylenedioxyphenyl]-1-propanone				



Table 2 Fragment ions and relative ion intensities in SIM mode (fragment ions in bold (quantification ions) were used in the calculation of peak area ratios; the remaining ions were used as qualification ions (confirmation ions); under-lined ions are the unique ions; the ions given in the brackets are the target ions of internal standards; the main ions that have 100% intensity were used to calculate the highest peak areas; the italicized ones are the molecular ions)

Target compounds	PPPA			TFA			CLF ₂ AA			HPBA			AA			PA		
	RT	<i>m/z</i>	Relative ion intensity (%)	RT	<i>m/z</i>	Relative ion intensity (%)	RT	<i>m/z</i>	Relative ion intensity (%)	RT	<i>m/z</i>	Relative ion intensity (%)	RT	<i>m/z</i>	Relative ion intensity (%)	RT	<i>m/z</i>	Relative ion intensity (%)
Flephedrone	10.22	204	100	10.26	154	100	12.40	170	100	10.63	254	100	12.87	58	100	13.61	58	100
		<u>123</u>	54		<u>123</u>	74		<u>123</u>	51		<u>123</u>	42		100	67		114	63
		160	32		110	29		<u>95</u>	<u>29</u>		210	27		<u>95</u>	<u>23</u>		<u>95</u>	<u>24</u>
		95	26		<u>95</u>	<u>26</u>		<u>75</u>	<u>11</u>		<u>95</u>	<u>17</u>		<u>123</u>	<u>19</u>		<u>75</u>	<u>12</u>
Mephedrone	11.81	119	100	11.93	119	100	14.00	119	100	12.16	119	100	14.58	58	100	15.25	58	100
		204 (207)	29		91	25		170	32		254	36		100	83		114	70
		91.1	24		154	21		91	26		(257)	—		(103)	—		(61)	—
		160	17		(157)	—		65	11		<u>91</u>	<u>23</u>		91	23		91	23
Pentredone	12.02	232	100	12.17	182	100	14.20	198	100	12.36	282	100	14.73	86	100	15.34	86	100
		190	66		<u>140</u>	<u>68</u>		156	46		240	58		128	61		142	48
		<u>105</u>	41		<u>105</u>	<u>55</u>		<u>105</u>	<u>37</u>		<u>103</u>	30		<u>77</u>	<u>17</u>		<u>77</u>	<u>13</u>
		77	27		77	31		77	29		<u>79</u>	<u>16</u>		105	10		105	11
Methedrone	13.60	135	100	13.77	135	100	15.74	135	100	13.89	135	100	16.34	58	100	16.93	58	100
		204	12		77.1	10		170	10		254	13		100	77		114	66
		77	11		91	8		77	10		210	7		135	34		135	28
		160	8		154	8		—	—		<u>169</u>	<u>7</u>		235	6		77	18
Methylone	14.61	149	100	14.77	149	100	16.72	149	100	14.89	149	100	17.29	58	100	17.85	58	100
		204	19		154	14		170	20		254	22		100	65		114	57
		(207)	—		(157)	—		121	12		(257)	—		(61)	—		(117)	—
		160	13		121	13		<u>319</u>	<u>6</u>		121	11		149	24		149	21
Butylone	15.15	149	100	15.37	149	100	17.25	149	100	15.41	149	100	17.80	72	100	18.34	72	100
		218	27		168	19		184	27		268	31		114	55		128	48
		(221)	—		(171)	—		(187)	—		(271)	—		(75)	—		—	—
		367	10		121	12		121	12		210	10		149	14		149	16
Ethylone	15.30	149	100	15.55	149	100	17.42	149	100	15.51	149	100	17.87	72	100	18.38	72	100
		218	37		168	28		184	37		268	40		114	60		128	48
		(223)	—		(173)	—		(338)	—		417	3		—	—		—	—
		190	18		140	13		333	12		417	5		236	8		277	3
Pyrovalerone	15.68	126	100	15.68	126	100	15.68	126	100	15.68	126	100	15.68	126	100	15.68	126	100
		149	9		91	7		127	6		127	9		91	7		91	5
MDPV	18.23	126	100	18.23	126	100	18.23	126	100	18.23	126	100	18.23	126	100	18.23	126	100
		(134)	—		(134)	—		(134)	—		(134)	—		(134)	—		(134)	—



each temperature) were added each day at RT, 40 °C, 55 °C and 70 °C respectively and set of the same time at 10, 20, 30, and 40 minutes for each derivatization reagent. From day five until day eight, samples were set in the same way as previous days; however, the times were changed to 5, 15, 25, and 35 minutes. Each temperature and each incubation time were analysed in triplicate. The samples were evaporated under a stream of nitrogen at RT for all reagents on days 1, 2, 3 and 4. The temperature on days 5, 6, 7 and 8 for PFPA, TFA and HFBA was RT, while 40 °C was used for AA and ClF₂AA and 50 °C for PA.

The 72 samples (18 samples for each temperature) under nitrogen gas were also evaluated using a TurboVap® in the following way: triplicate samples were run in one day when the hot block was set at 50 °C and the period of incubation was 20 minutes under RT, 40, 55 and 70 °C.

A different procedure was carried out to examine the effect of pyridine as a solvent in BA and PA in the following way: 200 µl of 10 µg ml⁻¹ from the mixtures of cathinones was added followed by evaporation at RT. The triplicates of 18 derivatized samples of BA and PA were closed and vortexed for 15 seconds and then incubated at 90 °C for 30 minutes and then in the evaporation step the samples were set at RT, 40 and 50 °C.

54 samples were set in the same way as mentioned above in one day to evaluate the reaction at RT, 55 °C, and 70 °C in 30 minutes (18 samples for each temperature). The evaporation step was set at RT. Again, each temperature was analysed in triplicate.

Optimisation procedure

The optimisation procedure for the incubation time and the temperature of the hot block were as follows: PFPA and TFA were set at RT and 40 °C, respectively, for 20 min; the hot block was set at RT; ClF₂AA and HFBA were set at 55 °C for 25 min; the hot block was set at 40 °C; AA and PA were set at 70 °C for 20 min; and the hot block was set at 50 °C.

The above procedures were used in this study to calculate peak areas, relative standard deviation (RSD), accuracy values and significant differences between the reaction temperature and time using ANOVA.

Table 4 Three way ANOVA shows if there is significant difference or not when incubation time and temperatures were changed in the procedure

Drug name/derv.	PFPA	TFA	ClF ₂ AA	HFBA	AA	PA
Flephedrone	Yes	No	No	No	No	Yes
Mephedrone	Yes	No	Yes	No	Yes	Yes
Pentadrone	No	No	Yes	No	No	Yes
Methedrone	No	No	No	Yes	Yes	Yes
Methylone	No	No	Yes	Yes	No	Yes
Butylone	No	No	Yes	Yes	No	Yes
Ethylone	No	No	No	Yes	No	Yes
Pyrovalerone	No	No	No	No	No	Yes
MDPV	No	No	No	No	No	Yes

Study of linearity, LOD, recovery and internal standards

For linearity, the samples were prepared in triplicate and spiked with cathinones at seven concentrations (2, 1, 0.75, 0.50, 0.25, 0.10, and 0.05) µg ml⁻¹, covering the range in which a common stimulant (amphetamine) is commonly encountered within toxicological samples.

For the LOD, the samples were prepared in triplicate and spiked at seven concentrations (250, 100, 50, 25, 10, 5, and 1 ng ml⁻¹).

SPE method used for recovery study

1 ml of whole blood of each sample was added and mixed with 1 ml of 0.10 M phosphate buffer (pH = 6), and all samples were then mixed and centrifuged. The extraction column was conditioned using 3 ml of methanol, followed by 3 ml deionized water and then 1 ml of 0.10 M phosphate buffer at pH 6 for washing the cartridges and removing unwanted substances. The samples were added and allowed to pass through the columns completely. Washing consisted of addition of 3 ml of deionized water, followed by 1 ml of 100 mM acetic acid and then 3 ml methanol followed by drying under full vacuum for 5 minutes. The samples were eluted with 3 ml of DCM : IPA : NH₄OH (78 : 20 : 2), and then evaporated under a stream of nitrogen at RT until dry. The dried extracts were then derivatised in the same way as mentioned above in the optimisation procedure.

Table 3 Optimisation of temperature and incubation time. This is according to the average of the highest values of peak areas at a concentration of 0.50 µg ml⁻¹. The temperature between the brackets is the optimised temperature of the evaporation step (after derv.)

Drug name/derv.	PFPA (RT)	TFA (RT)	ClF ₂ AA (40 °C)	HFBA (40 °C)	AA (50 °C)	PA (50 °C)
Flephedrone	20 min RT	20 min 40 °C	20 min 40 °C	20 min 40 °C	25 min 55 °C	25 min 70 °C
Mephedrone	10 min RT	20 min 40 °C	25 min 70 °C	25 min 55 °C	20 min 40 °C	25 min 70 °C
Pentadrone	20 min RT	20 min 40 °C	25 min 55 °C	20 min 40 °C	25 min 55 °C	25 min 55 °C
Methedrone	20 min 40 °C	20 min 40 °C	25 min 55 °C	25 min 70 °C	25 min 55 °C	25 min 70 °C
Methylone	35 min 70 °C	20 min 40 °C	25 min 55 °C	20 min 40 °C	25 min 55 °C	25 min 70 °C
Butylone	20 min 40 °C	20 min 40 °C	25 min 55 °C	20 min 40 °C	25 min 70 °C	25 min 70 °C
Ethylone	20 min 40 °C	20 min 40 °C	25 min 55 °C	20 min 40 °C	15 min 70 °C	25 min 70 °C
Pyrovalerone	35 min 70 °C	25 min 70 °C	25 min 70 °C	25 min 55 °C	25 min 70 °C	25 min 70 °C
MDPV	35 min 70 °C	25 min 70 °C	25 min 70 °C	25 min 55 °C	15 min 70 °C	25 min 70 °C
Optimisation	20 min RT	20 min 40 °C	25 min 55 °C	20 min 55 °C	25 min 70 °C	25 min 70 °C



For ISDs, the procedure outlined in the linearity study was used.

GC-MS conditions

Gas chromatography-mass spectrometry (GC-MS) was carried out using a 7890A GC/5975C MSD equipped with a split/splitless inlet and a DB-5ms (5% phenyl/95 methylsiloxane; 30 m × 0.25 mm, 0.25 μm film thickness) separation column (All Agilent Technologies, Waldbronn, Germany). Helium was used as a carrier gas (99.99% purity). Splitless injection at 225 °C was employed. The MS transfer line temperature was

maintained at 250 °C. The MS was operated in the electron impact ionization mode (70 eV). The ion source was maintained at 200 °C. MS data acquisition was initiated at 7 minutes and was performed in selected ion monitoring (SIM) mode and scan mode. The column temperature program was initially started at 70 °C and then increased by 10 °C per minute to reach 280 °C with a final hold time of 23 minutes. The mass spectrometer was operated in full scan mode (m/z 40–500) to study ion and peak interference. Selected ion monitoring (SIM) mode was used to study the linearity, limit of detection (LOD), recoveries, and peak areas.

Table 5 Examination of the quality of (R^2) in the target ions of selected internal standards of each drug. All drugs are listed according to their elution in chromatograms (tR)

Compound with ISDs/deriv.	PFPA (R^2)	TFA (R^2)	CLF ₂ AA (R^2)	HFBA (R^2)	AA (R^2)	PA (R^2)
Flephedrone- <i>ISD</i> mephadrone d ₃	0.998 ^a	0.999 ^a	B.R	0.998 ^a	0.997 ^a	0.999 ^a
Flephedrone- <i>ISD</i> methylone d ₃	0.996	0.999	0.999 ^a	B.R ^b	0.990	0.991
Flephedrone- <i>ISD</i> butylone d ₃	0.997	1.000	0.999	0.995	0.995	B.R
Flephedrone- <i>ISD</i> ethylone d ₅	0.995	0.990	0.995	B.R	B.R	B.R
Flephedrone- <i>ISD</i> MDPV d ₈	0.994	0.998	B.R	0.996	0.991	0.999
Mephadrone- <i>ISD</i> mephadrone d ₃	0.999 ^a	0.999 ^a	B.R	0.999 ^a	0.997 ^a	1.000 ^a
Mephadrone- <i>ISD</i> methylone d ₃	0.997	0.997	0.997 ^a	B.R	0.942	0.995
Mephadrone- <i>ISD</i> butylone d ₃	0.997	0.996	1.000	0.994	0.959	B.R
Mephadrone- <i>ISD</i> ethylone d ₅	0.995	0.996	0.998	B.R	B.R	B.R
Mephadrone- <i>ISD</i> MDPV d ₈	0.994	0.989	B.R	0.994	0.941	0.988
Pentedrone- <i>ISD</i> mephadrone d ₃	0.998 ^a	0.998 ^a	B.R	0.998 ^a	0.997 ^a	0.997 ^a
Pentedrone- <i>ISD</i> methylone d ₃	0.995	0.995	0.997 ^a	B.R	0.955	0.978
Pentedrone- <i>ISD</i> butylone d ₃	0.995	0.995	0.997	0.997	0.967	B.R
Pentedrone- <i>ISD</i> ethylone d ₅	0.994	0.995	0.994	B.R	B.R	B.R
Pentedrone- <i>ISD</i> MDPV d ₈	0.994	0.988	B.R	0.998	0.954	0.986
Methadrone- <i>ISD</i> mephadrone d ₃	1.000 ^a	0.999 ^a	B.R	1.000 ^a	0.999 ^a	0.996
Methadrone- <i>ISD</i> methylone d ₃	0.996	1.000	0.999 ^a	0.999	0.999	0.999 ^a
Methadrone- <i>ISD</i> butylone d ₃	0.994	0.999	1.000	0.996	0.999	B.R
Methadrone- <i>ISD</i> ethylone d ₅	0.998	0.999	0.998	B.R	B.R	B.R
Methadrone- <i>ISD</i> MDPV d ₈	0.996	0.996	B.R	B.R	0.999	0.997
Methylone- <i>ISD</i> mephadrone d ₃	0.998	0.999	B.R	0.998	0.999	0.995
Methylone- <i>ISD</i> methylone d ₃	0.999 ^a	0.999 ^a	0.998 ^a	0.999 ^a	0.998 ^a	1.000 ^a
Methylone- <i>ISD</i> butylone d ₃	0.999	0.999	1.000	0.998	0.999	B.R
Methylone- <i>ISD</i> ethylone d ₅	0.999	0.998	0.999	B.R	B.R	B.R
Methylone- <i>ISD</i> MDPV d ₈	0.993	0.998	B.R	B.R	0.997	0.997
Butylone- <i>ISD</i> mephadrone d ₃	0.997	0.999	B.R	0.996	0.999	0.996
Butylone- <i>ISD</i> methylone d ₃	0.999	1.000	0.995	0.999	0.997	1.000 ^a
Butylone- <i>ISD</i> butylone d ₃	0.999 ^a	1.000 ^a	0.999 ^a	1.000 ^a	1.000 ^a	B.R
Butylone- <i>ISD</i> ethylone d ₅	0.999	0.999	0.997	B.R	B.R	B.R
Butylone- <i>ISD</i> MDPV d ₈	0.995	0.997	B.R	B.R	0.996	0.996
Ethylone- <i>ISD</i> mephadrone d ₃	0.994	0.999	B.R	B.R	0.995	0.996
Ethylone- <i>ISD</i> methylone d ₃	0.998	1.000	0.996	0.942	0.998	1.000 ^a
Ethylone- <i>ISD</i> butylone d ₃	0.999	1.000	0.999	0.978 ^a	0.994 ^a	B.R
Ethylone- <i>ISD</i> ethylone d ₅	0.999 ^a	0.999 ^a	0.999 ^a	B.R	B.R	B.R
Ethylone- <i>ISD</i> MDPV d ₈	0.998	0.997	B.R	B.R	0.997	0.996
Pyrovalerone- <i>ISD</i> mephadrone d ₃	0.992	0.994	B.R	0.998	0.995	0.986
Pyrovalerone- <i>ISD</i> methylone d ₃	0.996	0.995	0.997	0.997	0.998	0.998
Pyrovalerone- <i>ISD</i> butylone d ₃	0.997	0.995	0.996 ^a	0.996 ^a	0.994	B.R
Pyrovalerone- <i>ISD</i> ethylone d ₅	0.997	0.994	0.992	B.R	B.R	B.R
Pyrovalerone- <i>ISD</i> MDPV d ₈	0.997 ^a	0.999 ^a	B.R	B.R	0.998 ^a	0.994 ^a
MDPV- <i>ISD</i> mephadrone d ₃	0.990	0.992	B.R	0.998	0.995	0.982
MDPV- <i>ISD</i> methylone d ₃	0.996	0.993	B.R	0.993	0.995	0.996
MDPV- <i>ISD</i> butylone d ₃	0.997	0.993	0.909 ^a	0.994 ^a	0.988	B.R
MDPV- <i>ISD</i> ethylone d ₅	0.997	0.992	B.R	B.R	B.R	B.R
MDPV- <i>ISD</i> MDPV d ₈	0.999 ^a	0.999 ^a	B.R	B.R	0.995 ^a	1.000 ^a

^a ISD used to study validation parameters. ^b B.R is bad response = <0.900.



Results and discussion

Fragment ions and relative ion intensities

The observed fragment ions and relative ion intensities for the different cathinone derivatives are summarised in Table 2. The target ions in bold were used to calculate accuracy and RSD values. The highest abundance ion values were used to calculate the peak area values.

Temperature, reaction time, and three way ANOVA study

The optimum temperature and reaction time for each compound using each reagent are shown in Table 3.

The optimum time and temperature in Table 3 were chosen for the mixture of synthetic cathinones to develop a method that works for the drug substances in each reagent. Therefore, the combination of information from Tables 3 and 4 illustrates the optimal conditions for reagents and drugs. Using the PFPA derivative of flephedrone as an example, the reaction conditions of RT for 20 min duration were chosen from Table 3 in combination with the ANOVA results from Table 4.

The optimal derivatization conditions for each compound were chosen according to the average of the highest values of peak areas at a concentration of $0.50 \mu\text{g ml}^{-1}$.

The peak area values of the target ions of cathinones were more evident using reaction conditions of 25 minutes at 70°C with the exception of PFPA and TFA derivatives

which showed excellent responses from RT derivatisation conditions. It should be noted that AA and PA are preferable for most of the cathinones when a high temperature of 70°C is applied. The cathinones generally require high temperatures for most of the derivatisation reagents which may be due to properties including the boiling point of each reagent and its molecular weight. It may be concluded that the higher the boiling points of reagents the higher the temperatures for reactions. PA, AA, HFBA, ClF_2AA , TFA and PFPA have the boiling points of 167°C , 139.8°C , 120°C , $96\text{--}97^\circ\text{C}$, 72.4°C , and $69\text{--}70^\circ\text{C}$, respectively. Mephedrone and flephedrone are more volatile compounds than other drugs because they have a lower molecular weight. However, the responses are improved when the reaction occurs at high temperatures when PFPA and TFA were excluded.

Due to the high boiling point (198°C) of butyric anhydride the reaction is not successful at 70°C . The excess reagent is not evaporated under nitrogen even when the temperature is higher than 70°C for 20 minutes. Additionally, this reagent provided a poor response for all compounds except when applied at a high concentration ($5 \mu\text{g ml}^{-1}$). For the above reasons this reagent was not investigated further.

The R programming language was used to perform a three-way ANOVA considering three factors (temperature and reaction time during incubation and the temperature of the hot block during the evaporation step) as independent

Table 6 Accuracy and precision, *the mean is the average of the highest peak area values of quantification ions at a concentration of $0.50 \mu\text{g ml}^{-1}$

Derv./drug name		PFPA	TFA	ClF_2AA	HFBA	AA	PA
Flephedrone	Mean*	2 283 223	1 178 147	1 881 598	3 563 229	2 407 035	1 263 952
	RSD (%)	4.07%	3.41%	10%	14%	1.13%	5.5%
	Accuracy	1.81%	−9.8%	−19%	−4.83%	3.67%	1.81%
Mephedrone	Mean	4 467 040	3 657 740	3 698 786	3 702 338	1 086 728	2 523 269
	RSD (%)	1.96%	0.99%	6.4%	2.02%	2.71%	11%
	Accuracy	4.79%	3.47%	−12%	−0.36%	−9.0%	−12%
Pentadrone	Mean	2 714 988	1 860 552	368 017	2 582 720	2 145 452	3 099 143
	RSD (%)	1.51%	2.37%	2.20%	4.33%	2.59%	12%
	Accuracy	10%	4.09%	−9.3%	11%	−9.3%	10%
Methedrone	Mean	7 144 720	6 822 530	6 657 846	6 353 019	574 827	2 489 097
	RSD (%)	4.49%	2.89%	4.43%	7.7%	2.59%	0.18%
	Accuracy	5.62%	12%	−7.1%	13%	5.2%	−12%
Methylone	Mean	6 296 421	9 973 042	5 487 420	3 591 150	2 099 231	1 157 643
	RSD (%)	1.46%	1.76%	0.45%	0.98%	1.76%	0.06%
	Accuracy	−11%	−1.11%	−12%	−2.55%	−6.8%	1.42%
Butylone	Mean	5 835 783	5 881 945	5 132 108	4 476 375	2 139 855	5 185 680
	RSD (%)	1.96%	7.6%	8.2%	9.7%	2.43%	5.6%
	Accuracy	−3.28%	−13%	−16%	−7.5%	8.5%	−3.28%
Ethylone	Mean	4 630 147	4 161 097	4 026 282	1 914 781	63 541.29	5 185 680
	RSD (%)	1.14%	1.81%	7.9%	6.8%	3.82%	5.6%
	Accuracy	2.09%	−12%	−9.0%	14%	−17%	0.44%
Pyrovalerone	Mean	5 801 857	2 929 385	5 626 518	7 895 943	6 658 780	4 976 504
	RSD (%)	12%	1.01%	4.74%	6.8%	10%	12%
	Accuracy	−13%	15%	−19%	14%	3.43%	13%
MDPV	Mean	4 735 925	3 709 708	4 519 016	6 421 153	4 600 039	5 523 840
	RSD (%)	14%	2.54%	10%	11%	12%	15%
	Accuracy	9.5%	7.5%	−16%	−19%	0.39%	−5.8%



Table 7 LOD and linearity. $n = 3$ for each point

Deriv./drug name	Flephedrone		Mephedrone		Pentadrone		Methedrone		Methylone		Butylone		Ethylone		Pyrovalerone		MDPV	
	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)	LOD	(R^2)
PFPA	1	0.998	1	0.999	1	0.997	1	0.999	5	0.999	5	0.999	5	0.998	50	0.956	50	0.944
TFA	5	0.995	5	0.999	5	0.998	5	0.999	5	0.999	5	0.997	10	0.999	50	0.912	50	0.985
ClF ₂ AA	10	0.992	5	0.995	5	0.997	5	0.994	25	0.998	50	0.998	25	0.996	50	0.955	50	0.978
HFBA	1	0.998	1	0.999	1	0.998	5	0.999	10	1.000	5	0.999	100	0.935	100	0.912	100	0.905
AA	50	0.996	25	0.990	1	0.996	50	0.999	25	0.998	250	0.998	100	0.996	25	0.998	25	0.995
PA	10	0.997	10	0.981	5	0.997	10	1.000	5	0.999	5	0.999	5	0.998	5	0.996	5	0.994

variables. The dependent variables (54 different ANOVA = 9 drugs \times 6 reagents) were the mean of peak area values at each specific time and temperature for each drug and for each derivatisation reagent alone (5184 tests of peak area values were produced; 5184 tests = 8 days \times 72 samples per day \times 9 drugs). In order to infer that there was a difference in the results it was expected to see at least one of the three independent variables to appear as statistically significant within the 5% level of confidence. If the probability factor (F) was higher than 5%, this means that the difference between peak area values, produced by altering the three variables noted above, was statistically significant. The data in Table 4 demonstrate that we should run the sample under a strict procedure or under specific conditions if there is significant difference (yes) in the derivatised drug. For example, the samples should follow the optimised procedure in the case of flephedrone and mephedrone derivatised by PFPA to get the best response; if not the peak area values will significantly change above the 95% confidence limit then, as the consequence will give a bad response. In the case of TFA derivatisation, the probabilities for all drugs to give the same values of peak areas even with changes in time or temperature within the 95% limit confidence are significantly the same. Therefore, many incubation times and temperatures are appropriate for this reagent. All substances derivatised with PA should follow the optimised procedure specifically the temperature of the hot block in the evaporation step. PA samples may need more than an hour to evaporate at RT.

The uncertainty studies may require answering the question: why do we have no significant differences?

It may be the effects of many factors such as losing the drug during the evaporation step or as a result of thermal decomposition of derivatised drugs in the injector port.¹⁶

Table 8 Recovery studies

Drug name/deriv.		PFPA	TFA	ClF ₂ AA	HFBA	AA
Flephedrone	Recovery	69%	69%	100%	59%	81%
	RSD (%)	7.4%	12%	17%	6.2%	2.14%
Mephedrone	Recovery	107%	104%	94%	64%	121%
	RSD (%)	7.1%	1.43%	9.7%	20%	7.6%
Pentadrone	Recovery	70%	112%	92%	43%	68%
	RSD (%)	2.63%	3.48%	17%	20%	5.2%
Methedrone	Recovery	107%	129%	100%	110%	94%
	RSD (%)	9.6%	7.9%	7.5%	19%	10%
Methylone	Recovery	101%	98%	98%	126%	82%
	RSD (%)	0.75%	2.37%	3.59%	16%	2.35%
Butylone	Recovery	145%	51%	37%	53%	75%
	RSD (%)	5.3%	0.84%	56%	1.80%	18%
Ethylone	Recovery	229%	117%	97%	14%	119%
	RSD (%)	32%	1.27%	5.4%	7.3%	15%
Pyrovalerone	Recovery	77%	19%	64%	52%	187%
	RSD (%)	1.90%	11%	23%	15%	15%
MDPV	Recovery	58%	122%	63%	134%	106%
	RSD (%)	1.13%	2.86%	20%	3.76%	3.24%



Internal standards, RSD and accuracy, linearities, limit of detection, and recovery studies

The internal standard results are shown in Table 5. The purpose of this procedure was to evaluate the application of internal standards.

- Can we use one or two internal standards (ISDs) for all studied cathinones?
- Which ISD has the best-fit regression?

To answer the above points, we applied each of the five internal standards to all nine drugs. All internal standards worked well and gave more than 0.990 when PFPA and TFA were applied. The ISDs that gave a poor response were avoided in all experiments.

The RSD (%), accuracy, linearity, LOD and recovery data were only calculated according to optimal conditions.

The RSD (%) values were calculated from the procedure of optimal methods only (the mean of SD \div the mean of peak area ratio) \times 100 at a concentration of 0.5 $\mu\text{g ml}^{-1}$. According to the RSD values of peak areas at 0.5 $\mu\text{g ml}^{-1}$, the best results were given by ClF₂AA followed by PFPA then AA, HFBA, TFA and PA respectively.

The accuracy values were calculated from (the mean of calculation of concentration – true values \div true values) \times 100 at a concentration of 0.5 $\mu\text{g ml}^{-1}$. The best results were given by PFPA then HFBA, TFA, PA, ClF₂AA and lastly AA. The anhydrides proved to be suitable for cathinone derivatization because none exceeded 20% for both RSD and accuracy, which is recommended in ref. 14 (see Table 6).

Linear correlation coefficients (R^2) were calculated from the triplicate samples at seven concentrations (2, 1, 0.75, 0.5, 0.25, 0.1, and 0.05 $\mu\text{g ml}^{-1}$). All R^2 values were greater than 0.905. The best results were obtained with PFPA and HFBA; all values were higher than 0.998 followed by PA, AA, TFA, and then ClF₂AA (pyrovalerone and MDPV were excluded).

The LOD was measured in SIM mode using methanol spiked with mixtures of cathinones in the range of 1 to 250 ng ml^{-1} . The signal-to-noise (S/N) ratio was calculated from triplicate

measurements at seven concentrations (250, 100, 50, 25, 10, 5, and 1 ng ml^{-1}). The lowest concentration at which the S/N was greater than 3 was considered to be the LOD. PFPA, PA, TFA, HFBA, ClF₂AA, and AA provided the best results respectively (see Table 7).

The recovery study was completed to check that all drugs can be derivatised after extraction of whole blood. Relative recoveries were calculated using a concentration of 3 $\mu\text{g ml}^{-1}$. The samples were extracted three times without internal standards present until addition prior to the evaporation step under nitrogen. At the same time three unextracted standards were prepared at 3 $\mu\text{g ml}^{-1}$ with internal standards. The recovery of each drug was calculated using the following equation: recovery% = (peak area ratio of extracted standards \div peak area ratio of un-extracted standards) \times 100. The recovery results with precision are shown in Table 8.

Study of interference, fragmentation patterns and the highest peak area values

Mixtures of nine cathinones were examined to study the fragmentation pattern (Fig. 2) and interference (Fig. 3). No co-elution problems were observed except in two cases: butylone could not be separated effectively from ethylone if they were derivatized with AA and PA. However, ethylone has a unique ion allowing the compounds to be distinguished from each other. Butylone and ethylone have the same M.W.; the differences between the fragmentation patterns were discussed by Kerrigan.¹⁶

A number of fragmentation ions were studied for the drugs in each reagent. In general AA followed by PA then HFBA, PFPA, TFA and ClF₂AA respectively give the maximum abundance ions based on ion intensities and greater fragmentation patterns than other reagents (see Fig. 2).

The highest abundance ion values were used to calculate the peak areas. These ions were chosen instead of target ions (quantification ions) because we need to compare the ions that

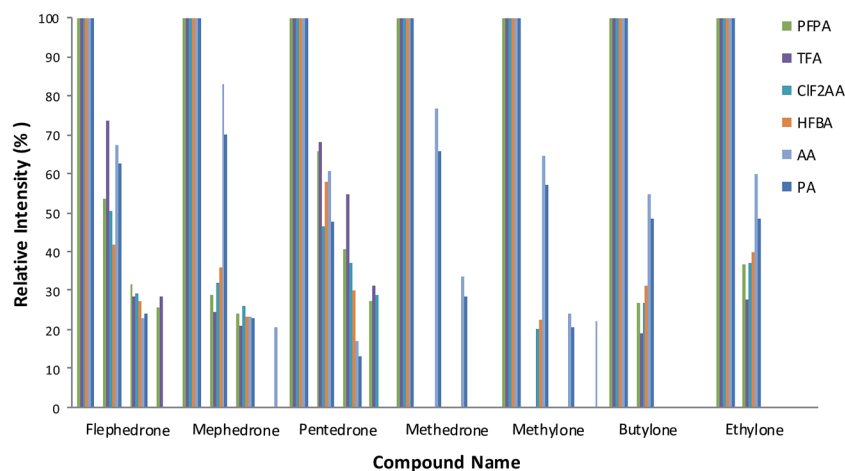


Fig. 2 Fragmentation patterns for each substance applied to selected reagents. Less than 10% fragmentation ions were deleted. The optimised methods were used to show the fragmentation patterns.



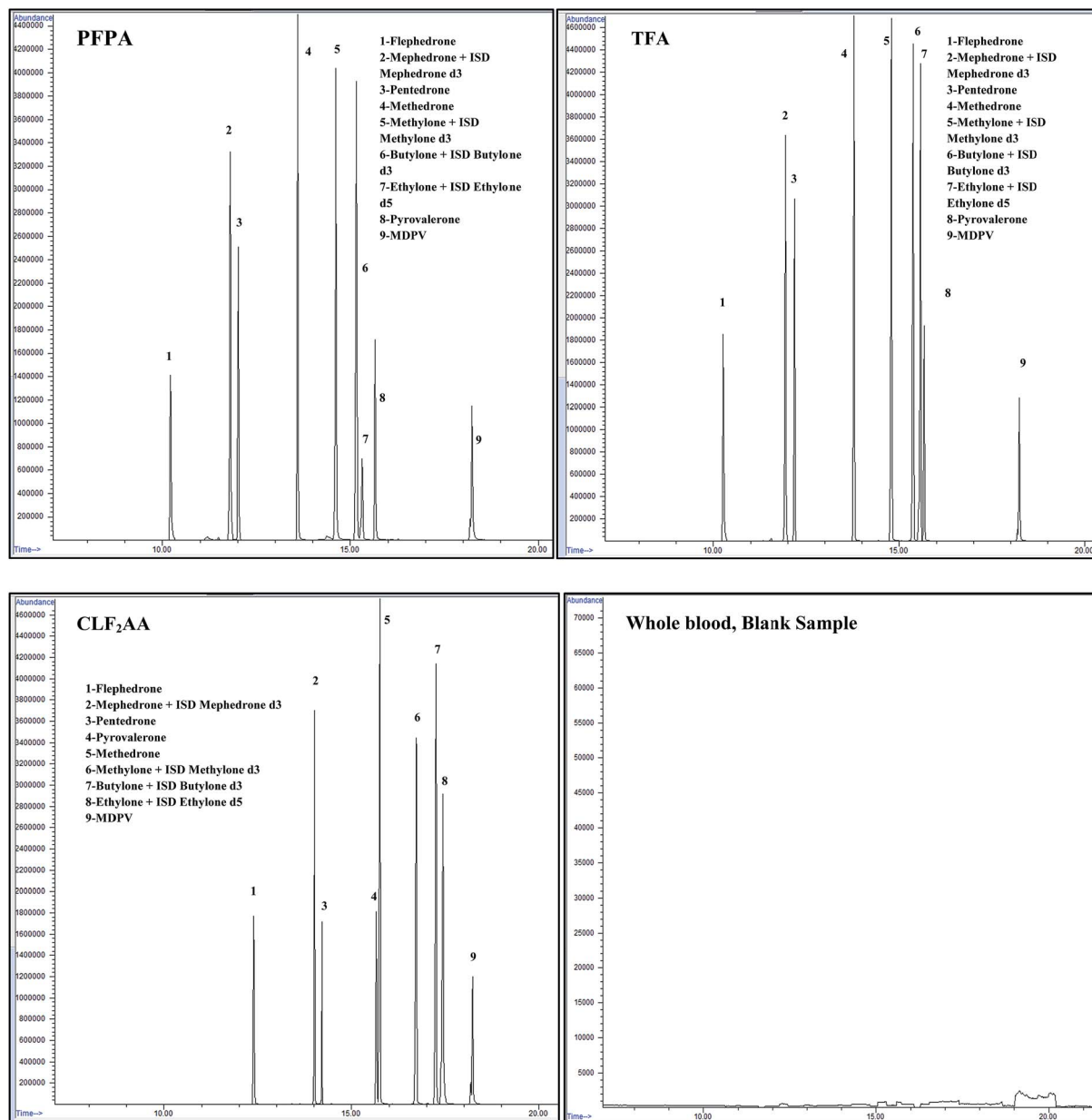


Fig. 3 Chromatograms for three different acetylation derivatives of synthetic cathinones at a concentration of $0.50 \mu\text{g ml}^{-1}$ and blank blood sample.

give 100% of abundance in the background of fragmentation ions for six different reagents. Each one of the reagents has different ions and so we should apply all of them with the same relative ion intensity (100%), as illustrated in Fig. 4. All valid results for derivatised drugs after optimising conditions have good peak areas excluding AA for ethylone and methedrone as well CLF₂AA for pentadrone.

Overview

Table 9 shows which reagent provides the best results under different factors. For example, if a screening method suggests

the presence of flephedrone is positive, we should use TFA for quantification methods due to the following reasons:

- Good fragmentation patterns are evident.
- High quality fragmentation ions are present.
- High response is observed compared to the main ion or remaining ions.
- It has the largest number of unique ions and ions in total.
- It is valid in linearity, accuracy and precision.

The example above can be applied to all figures using a similar explanation.



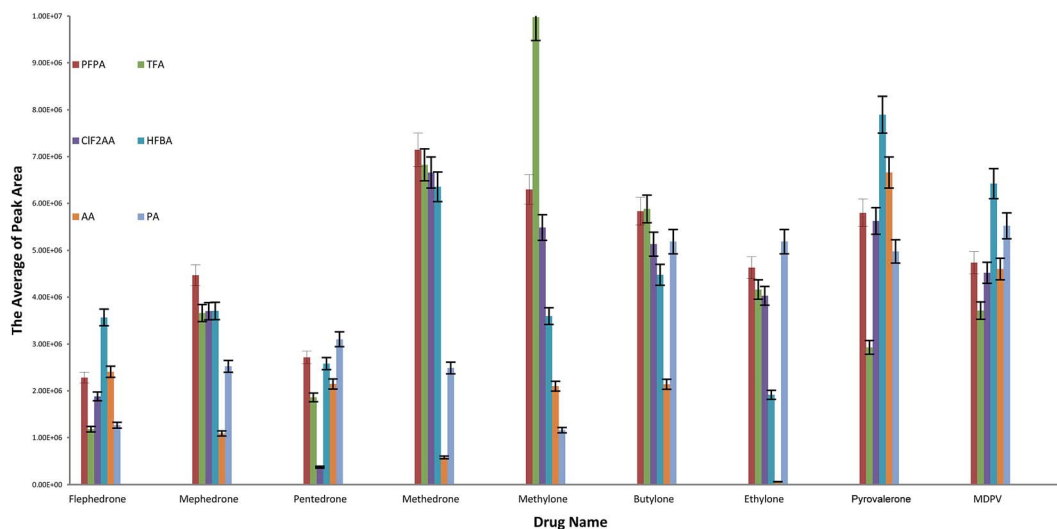


Fig. 4 Mean of the highest peak area values in selected compounds and agents. The optimised methods were used in this calculation.

Table 9 Favoured reagents for each drug depending on selected parameters

Drug name	Ratio between all ions	No. of unique ions	No. of ions	LOD	Linearity	RSD	Accuracy	The highest peak area	Evaporation 25 °C after deriv.	Evaporation 50 °C after deriv.
Flephedrone	TFA	TFA, CLF ₂ A	TFA, HFBA	PFPA, HFBA	All fit	Valid	Valid	HFBA/AA	Valid	Bad response
Mephedrone	AA	PFPA, HFBA, TFA, CLF ₂ A	HFBA	PFPA, HFBA	All fit	Valid	Valid	PFPA/HFBA	Valid	Bad response
Pentedrone	TFA	HFBA, AA	PFPA	PFPA, HFBA, AA	All fit	Valid	Valid	PA/PFPA	Valid	Bad response
Methedrone	PA	HFBA, AA	HFBA	PFPA	All fit	Valid	Valid	PFPA/TFA	Valid	Bad response
Methylone	AA	HFBA, CLF ₂ AA, AA, PA	HFBA	PFPA, TFA, PA	All fit	Valid	Valid	TFA/PFPA	Valid	Bad response
Butylone	AA	No one	HFBA, PA	All except CLF ₂ AA and AA	All fit	Valid	Valid	TFA/PFPA	Valid	Valid
Ethylone	PFPA	All	PA	PFPA, PA	All fit	Valid	Valid	PA/PFPA	Valid	Valid
Pyrovalerone	Underivatized	Underivatized	Underivatized	PA	PA	PA	PA	HFBA/AA	Bad response	Valid
MDPV	Underivatized	Underivatized	Underivatized	PA	PA	PA	PA	HFBA/AA	Bad response	Valid

Conclusion

The mass spectra of nine synthetic cathinones were compared to one another after derivatization with a number of different acylation reagents (PFPA, TFA, CLF₂AA, HFBA, AA and PA). The optimisation of conditions for the incubation time and reaction temperature was discussed, and all anhydrides tested proved to be suitable for synthetic cathinones with RSD and accuracy below 20% under the optimised conditions for the reagents. The independent variables were assessed using a three-way ANOVA approach and demonstrated that the procedure should be strictly followed for combinations of drugs and reagents. In this paper, we have shown that one or two ISDs may be sufficient to provide good linearity in the mixture of cathinones chosen. The overview section shows

some suggestions of which reagent can be chosen for each compound and applied to quantification or semi-quantification methods only. In general, PFPA and HFBA, followed by TFA, are the best choice of derivatising agents. Therefore, our future aim will be to fully validate a method using PFPA. MDPV and pyrovalerone are tertiary amines that were not derivatized with reagents tested. Therefore, we can conclude that several combinations of cathinones and derivatization reagents are suitable for GC-MS analysis.

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