

Evaluation of drug incorporation into hair segments and nails by enantiomeric analysis following controlled single MDMA intakes

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ABSTRACT

Incorporation rates of the enantiomers of 3,4-methylenedioxyamphetamine (MDMA) and its metabolite 3,4-methylenedioxyamphetamine (MDA) into hair and nails were investigated after controlled administration. Fifteen subjects without MDMA use received two doses of 125 mg of MDMA. Hair, nail scrapings and nail clippings were collected 9-77 days after the last administration (median: 20 days). Hair samples were analyzed in segments of 1- to 2-cm length. After chiral derivatization with N-(2,4-dinitro-5-fluorophenyl)-L-valinamide, MDMA and MDA diastereomers were analyzed by liquid chromatography-tandem mass spectrometry. Highest concentrations in hair segments corresponded to the time of MDMA intake. They ranged from 101 to 3,200 pg/mg and 71 to 860 pg/mg for *R*- and *S*-MDMA, and 3.2 to 116 pg/mg and 4.4 to 108 pg/mg for *R*- and *S*-MDA, respectively. MDMA and MDA concentrations in nail scrapings and clippings were significantly lower than in hair samples. There was no significant difference between enantiomeric ratios of *R/S*-MDMA and *R/S*-MDA in hair and nail samples (medians: 2.2-2.4 for MDMA and 0.85-0.95 for MDA). Metabolite concentration ratios of MDA to MDMA were in the same range in hair and nail samples (medians: 0.044-0.055). Our study demonstrates that administration of two representative doses of MDMA was detected in the hair segments corresponding to the time of intake based on average hair growth rates. MDMA was detected in all nail samples regardless of time passed after intake. Comparable *R/S* ratios in hair and nail samples may indicate that incorporation mechanisms for both matrices are comparable.

Keywords: Hair analysis; nail analysis, MDMA enantiomers; controlled administration

Introduction

The analysis of keratinized matrices such as hair has achieved considerable importance in forensic toxicology. Main applications are long-term monitoring of drug and alcohol consumption, e.g. of drug abusers or dealers, in abstinence controls for regranting driver's license, in postmortem investigations or in drug-facilitated crimes [1,2]. In some cases, it is necessary to evaluate whether an individual has taken the drug on a regular basis or whether a single or sporadic intake is more likely. In order to gain time-resolved information on an individual's drug exposure, the Society of Hair Testing (SoHT) recommends analysis of hair segments of 1- to 3-cm [3]. Based on an average growth rate of 1 cm per month (range 0.7 – 1.4 cm per month) the time frame of drug intake can be estimated [1]. Several reports describe the detection of drug not only in hair segments corresponding to the expected time of intake/administration but also in adjacent segments [4-7]. This may render interpretation of analysis results difficult. For amphetamines, a positive correlation after controlled administration of a low and high dose of amphetamine and methamphetamine was demonstrated within subjects but not between subjects [8].

Nails, another keratinized matrix, are being discussed as an alternative to hair because nails also accumulate drugs [9-15]. Investigations on paired hair and nail analysis for amphetamines delineated similar [16] or higher concentrations in hair [17], but one group reported slightly higher amphetamine concentrations in fingernails compared to hair [18].

The exact mechanism for drug incorporation into hair and nails has not been identified [1,10]. It is nowadays recognized that drug disposition into hair takes place not only at the hair follicle but also via diffusion from sweat and sebum [1]. Studies on the incorporation pathways into nails were performed after a single dose of zolpidem [19,20]. The germinal nail matrix and nail bed were identified as incorporation regions

for drugs. Furthermore, sweat presumably contributes to a large extent to drug disposition into nail as drug could already be detected in fingernail clippings 24 hours after drug intake [20]. The investigation of drug stereoisomers could hypothetically provide insights into incorporation mechanisms into hair and nail. It is recognized that enantiomers can display significantly distinct pharmacokinetic and pharmacodynamic behavior implying differences in affinity to metabolizing enzymes or receptor sites [21].

MDMA (3,4-methylenedioxyamphetamine, ecstasy) is an illicit drug mostly traded as racemic mixture [22]. It belongs to the most prevalent drugs involved in forensic and driving under the influence cases [23-27]. MDMA is used for its entactogenic effects and stimulation of the central nervous system [28] which are mainly attributed to the *S*-MDMA enantiomer [29]. The *R*-enantiomer was reported to be more abundant in human tissues and fluids representing its enantioselective metabolism most likely because of the elimination of the *S*-enantiomer from plasma at a higher rate [30-33]. MDMA is primarily metabolized to 3,4-dihydroxymethamphetamine (DHMA) by the cytochrome P450 isoenzyme CYP2D6 [29], which is polymorphic [34]. MDA (3,4-methylenedioxyamphetamine) is a minor metabolite of MDMA formed by *N*-demethylation [29].

The present study aims at analyzing hair and nail samples for enantiomers of MDMA and its metabolite MDA after controlled administration of two doses of 125 mg of racemic MDMA. Segmental analysis of hair samples is performed to study whether drug detection in a hair segment can be correlated to the respective time period of MDMA intake. Nail samples are evaluated as an alternative specimen to hair when testing for single/sporadic MDMA intake. The enantioselective patterns of MDMA and MDA in hair and nails are examined to gain insights whether incorporation mechanisms into these matrices differ. To our knowledge, the present study is the

first report on the disposition of MDMA and MDA enantiomers in hair and nail samples of non-drug using participants following controlled intake of MDMA.

Experimental

Study design and sample collection

Hair and nail samples were obtained from a double-blind, placebo-controlled, cross-over study performed at the University Hospital of Basel, Switzerland, to investigate pharmacokinetic effects of MDMA [35]. The study was conducted in accordance with the Declaration of Helsinki and International Conference on Harmonization Guidelines in Good Clinical Practice and approved by the Ethics Committee of the Canton of Basel, Switzerland, and the Swiss Agency for Therapeutic Products (Swissmedic). All participants provided written informed consent. Only subjects with no prior regular MDMA use were included. Five subjects had previously used MDMA (<5 times) at least 2 months prior to the study start. Drug use during the study was excluded using repeated urine tests [35]. Subjects (n=15) were administered two doses of pharmaceutical grade racemic MDMA (MDMA hydrochloride 125 mg) in capsules. The time interval between the two administrations was different for each subject ranging between 9 to 105 days (median: 18 days). Head hair samples, nail scrapings, and clippings were collected; 13 nail samples were from the ring finger and 3 from the big toe. Nails were wiped twice with acetone prior to sample collection. The time between last MDMA intake and sample collection ranged from 9 to 77 days (median: 20 days). All subjects were pheno- and genotyped for CYP2D6 activity [35]. Fourteen participants were extensive metabolizers and one participant (no. 1) was a poor metabolizer (Table 2).

Materials

Deuterated and undeuterated racemic standards were obtained as calibrated methanolic solutions from Lipomed (Arlesheim, Switzerland). N-(2,4-dinitro-5-fluorophenyl)-L-valinamide (DNPV) for derivatization and all other chemicals were

purchased from Sigma-Aldrich Chemie GmbH (Buchs, Switzerland). Solvents for washing, extraction and analysis by liquid chromatography-tandem mass spectrometry (LC-MS/MS), all of HPLC grade, were obtained from Sigma-Aldrich Chemie GmbH (Buchs, Switzerland). Drug-free hair and nail samples for method validation were provided by drug-naïve volunteers.

Analytical methods

Hair and nail sample preparation

Hair samples were segmented in three 1-cm and one 2-cm-segments from proximal to distal. Hair segments were washed separately with 2 mL water and 2 mL acetone for 2 min, nail clippings were washed once with 1 mL water for 30 sec by vortexing. On average 20 mg hair segments, 5 mg clippings and 1-5 mg scrapings were used for analysis. Hair and nail clippings were pulverized at 25 Hz for 8 min using a mixer mill (MM 200 Retsch, Switzerland). For extraction, 0.1 mL IS solution (50 ng/mL) containing MDMA-d5 and MDA-d5 and 0.9 mL MeOH were added and samples were sonicated for 4 h. Extracts were filtered through a 0.22 μ m PTFE syringe filter (BGB, Boeckten, Switzerland) and 50 μ L formic acid (20%, v/v) were added for evaporation at 50 °C under nitrogen. The residue was reconstituted in 80 μ L carbonate buffer (1 M, pH 9). For derivatization of MDMA and MDA 30 μ L 0.3 % N-(2,4-dinitro-5-fluorophenyl)-L-valinamide (DNPV) in acetone were added and vials were placed in a heating block at 50 °C for 30 min as previously described [36,37]. Derivatization was stopped by the addition of 20 μ L hydrochloric acid (1 M). After the addition of 30 μ L mobile phase mixture (A: 25 M ammonium acetate buffer pH 4 containing 0.1 % of formic acid/B: acetonitrile containing 0.1 % of acetic acid; 1/1; v/v) diastereomers of derivatized MDMA and MDA were analyzed by LC-MS/MS.

LC-MS/MS parameters

Analytes were separated on a Phenomenex Kinetex C18 column (100 x 2.1 mm, 2.6 μ m) using a Thermo Fisher Ultimate 3000 high performance liquid chromatography (UHPLC) system and detected by an Applied Biosystems 5500 Q Trap linear ion trap triple quadrupole mass spectrometer (Sciex, Darmstadt/Germany) with Analyst software (Version 1.6). The following mobile phase gradient of eluent A and B was adjusted: 0-1 min: 2% B; 1-2 min: increase to 40% B; 2-9.3 min: increase to 60% B; 9.3-9.5 min: increase to 98% B; 9.5-10.5 min: holding at 98% B; 10.5-10.8 min: decrease to 2% B; 10.8-12 min: holding at 2% B. The flow rate was 0.4 mL/min. The column temperature was maintained at 35°C and the injection volume was 10 μ L. The MS instrument was operated in positive electrospray ionization and multiple reaction monitoring mode. Quantification was achieved using the most abundant transition of each precursor to the respective product ion and two MRM transitions served for identification. Further MS operation parameters were as previously reported by Steuer et al. [36].

Method validation

Standard solutions of racemic MDMA and MDA for six calibrators and quality controls (QC) at two concentration levels, low and high, were prepared separately by diluting the respective stock solution with acetonitrile. Concentrations are given in Table 1. An internal standard solution of racemic MDMA-d5 and MDA-d5 in acetonitrile was prepared at a concentration of 50 ng/mL. Blank matrix of 20 mg hair and 5 mg nail were spiked with 100 μ L of each calibrator or QC solution. Calibration curves and quality control samples (in duplicate) were analyzed on five days. Quantitation was performed by determining the analyte to IS peak area ratios. A weighted linear regression model (1/concentration) was applied. The lower limit of quantification

(LLOQ) and limit of detection (LOD) was chosen to be the concentration resulting in a peak height with a signal-to-noise ratio of 10 and 3, respectively. Matrix effects were studied according to Matuszewski et al. [38]. The method was validated according to international guidelines[39].

Data analysis

Statistical analyses were carried out with Prism 6 (GraphPad Software, La Jolla, CA, USA). The nonparametric two-sided Wilcoxon matched pairs rank signed test was utilized for comparison of paired nail and hair samples results because no Gaussian distribution was observed. P values between 0.001–0.01 were considered as very significantly different, 0.01–0.05 as significantly different, p values above 0.05 were not significant.

Results

Method validation

The method was validated based on the Guidelines for Quality Control in Forensic-Toxicological Analyses of the GTFCh for the criteria selectivity, LOD, LLOQ, linearity of calibration, accuracy, repeatability, precision, and matrix effects [39]. Validation data are given in Table 1. Calibration curves were linear over the concentration range of 0.5 to 500 pg/mg for hair samples and 2.0 to 2000 pg/mg for nail samples. The criteria for accuracy, repeatability and precision were fulfilled. In case of *S*-MDMA, the deuterated analogue as IS should compensate for matrix effects which were slightly above the acceptable threshold (Table 1).

Study samples

Hair specimens were obtained from 13 participants. Hair samples were analyzed in segments of 1 + 1 + 1 + 2 cm length from proximal to distal. All subjects (n=15) provided nail samples; 13 clippings from fingernails and three from toenails, and 12 scrapings from fingernails and two from toenails were collected, respectively. All samples were tested for MDMA and MDA enantiomers. The sum of MDA and MDMA enantiomers, respectively, was calculated. Concentrations for 5-cm-hair segments were calculated as an average of the four hair segments considering segment lengths. Concentrations of MDMA and MDA enantiomers and their sum in all samples are given in Figure 1 and Table 2, respectively.

MDMA and MDA in hair samples

All hair segments were tested positive for MDMA and MDA enantiomers. *R*- and *S*-MDMA concentrations ranged from 101 to approximately 3,200 pg/mg (median: 470 pg/mg) and 71 to approximately 860 pg/mg (median: 221 pg/mg), respectively

(Figure 1A, B). The highest *R*-MDMA concentration (identified as outlier by Grubbs' test, $\alpha < 0.01$) was observed in the hair sample of participant no. 1 who was genotyped a CYP2D6 poor metabolizer. *R*-MDMA was detected to a significantly higher extent into hair than *S*-MDMA (Figure 1A) whereas no difference was observed for MDA enantiomers (Figure 1B). *R*- and *S*-MDA concentrations ranged from 3.2 to 116 pg/mg (median: 21 pg/mg) and 4.4 to 108 pg/mg (median: 14 pg/mg), respectively (Figure 1B).

In all cases, the highest concentration of each MDMA and MDA enantiomer was found in hair segments corresponding to the time of MDMA intake based on an average hair growth rate of 1 cm per month. Accordingly, concentrations of the sum of MDMA enantiomers in hair segments corresponding to the time of intake ranged from 173 to 3'770 pg/mg (median: 694 pg/mg). The sum of concentrations of MDMA enantiomers in the four hair segments and the time of MDMA intake are presented in Figure 2. In all samples MDMA and MDA could also be detected in hair segments adjacent to the one corresponding to time of intake, however, at much lower levels. Calculated concentrations of the sum of MDMA and MDA enantiomers, respectively, in 5-cm-segments varied from 78 to 850 pg/mg (median: 370 pg/mg) and 3.9 to 46 pg/mg (median: 19 pg/mg), respectively (Table 2).

MDMA and MDA in nail samples

All nail clippings were tested positive for MDMA and MDA enantiomers regardless of time past MDMA intake (Table 2) (Figure 1C-F). *R*- and *S*-MDMA concentrations in nail clippings ranged from 62 to approximately 990 pg/mg (median: 116 pg/mg) and from 20 to approximately 600 pg/mg (median: 56 pg/mg), respectively. *R*- and *S*-MDA concentrations in nail clippings ranged from 2.0 to 28 pg/mg (median: 5.8 pg/mg) and from 2.9 to 48 pg/mg (median: 6.2), respectively. By far highest *R*-/*S*-

MDMA and *R/S*-MDA concentrations were detected in the toenail clipping from participant no. 12 (Figure 1C, D). All nail scrapings were tested positive for *R*- and *S*-MDMA (Figure 1E) with concentrations ranging from 4.3 to 303 pg/mg (median: 93 pg/mg) and from 2.0 to 180 pg/mg (median: 44 pg/mg). *R*- and *S*-MDA could only be detected in seven of 13 scrapings at low concentrations ranging from 2.3 to 12.9 pg/mg for both enantiomers (Figure 1F).

There was no statistical difference in the calculated sum of concentrations of MDMA and MDA enantiomers in scrapings and clippings; medians were 134 and 167 pg/mg for MDMA, and 6.3 and 12 pg/mg for MDA, respectively (Table 2). However, the sum of MDMA and MDA enantiomers were significantly lower in nail samples compared to hair samples considering a segment length of 5 cm (Table 2).

Enantiomer ratios of MDMA and MDA in hair and nail samples

R/S-MDMA concentration ratios were higher than 1 in all hair samples (Figure 3A). There was no significant difference between hair segments corresponding (median: 2.1) and not corresponding (median: 1.9) to the time of intake (Figure 3A). *R/S*-MDMA concentration ratios were higher than 1 in all nail clippings and scrapings (medians: 2.2 and 2.4, respectively) (Figure 3A). The highest *R/S*-MDMA ratio of 5.7 and 22 was observed in a hair segment and nail scraping of participant no. 1 who was a CYP2D6 poor metabolizer (Figure 3A).

S-MDA was detected at slightly higher amounts in hair segments in most subjects, median *R/S* ratios were 0.84 and 0.90, respectively (Figure 3B). In nail clippings and scrapings, the median concentration ratios for *R/S*-MDA were 0.77 and 0.95, respectively (Figure 3B). There was no significant difference for enantiomeric ratios of MDMA and MDA in nail and hair samples (Figure 3).

Two trends of *R/S*-MDMA concentration ratios along the four hair segments of each participant were observed: decreasing (no. 1, 2, 4, 7, 11, 12, 13, 16, 17, and 18) and increasing (no. 6 and 10) from proximal to distal, respectively (Figure 4). There was no obvious trend in samples from participant no. 14 and 18 (Figure 4).

Metabolite ratios of MDA to MDMA in hair and nail samples

MDA to MDMA ratios in corresponding and not corresponding hair segments ranged from 0.031 to 0.074 (median: 0.047) and from 0.029 to 0.11 (median: 0.054), respectively (Figure 5). In nail clippings and scrapings, metabolite ratios ranged from 0.041 to 0.13 (median: 0.055) and from 0.041 and 0.084 (median: 0.044), respectively (Figure 5). Overall, there was no statistical difference in metabolite concentration ratios of the sum of concentrations of MDA to MDMA enantiomers in hair segments and nail samples.

Discussion

The analytical method applying chiral derivatization and LC-MS/MS analysis proved to be a valuable tool for the quantification of MDMA and MDA enantiomers in hair samples, nail scrapings and nail clippings. The samples were not tested for further MDMA metabolites such as DHMA and HMMA. Preliminary studies on hair samples positive for MDMA were negative for DHMA and negative for HMMA or only traces of HMMA. The method is specific and highly sensitive as low LLOQ were achieved with a sample weight of as low as 20 mg of hair and 5 mg of nail, respectively. In forensic investigations, it is often crucial to identify the time period in which a drug was taken. Therefore, controlled drug administration studies are needed for better understanding of disposition profiles into hair and nails.

MDMA and MDA in hair samples

As recently demonstrated, the time point of amphetamine intake could be estimated by the analysis of 0.5-cm-hair segments after controlled administration of 50 mg amphetamine. However, only five out of nine study participants were tested positive [5]. In our study, all hair samples were positive for MDMA and MDA enantiomers after two representative intakes of 125 mg MDMA (Table 2, Figure 1). Segmental hair analysis demonstrated that the highest enantiomeric (data not shown) and the highest calculated sum of concentrations of MDMA enantiomers (Figure 2) in hair segments correlated well with time of MDMA intake based on an average hair growth rate of 1 cm per month. However, the inter-individual variability in MDMA hair concentrations was large.

In seven of the 13 participants (no. 4, 6, 11, 12, 16-18) one 1-cm-segment covered the time of both MDMA doses. Five participants (no. 2, 7, 10, 13, 14) received the

two MDMA doses with a time interval long enough to allow for monitoring by two hair segments (Figure 2). Intra-individual MDMA concentrations in these two hair segments of these participants were very similar (Figure 2, Table 2).

The hair sample of participant no. 1 was too short to cover the time of the first MDMA intake (Table 2). The highest MDMA concentration (determined as outlier by Grubbs test, $p > 0.05$) was observed for subject no. 1 who was a CYP2D6 poor metabolizer involving a reduced elimination rate of MDMA compared to extensive metabolizers. The lowest sum of MDMA enantiomer concentrations was detected in a light brown hair sample (no. 11) (Table 2). However, there was no significant difference between (light) brown and dark brown hair samples in MDMA concentrations in 1-, 2- and 5-cm-hair segments (data not shown). Also Mieczkowski and Newel who reviewed different data sets stated that there is no relationship of hair colour and MDMA concentrations in hair [40].

For CYP2D6 extensive metabolizers, the median sum of MDMA enantiomer concentration in the corresponding 1-cm and the 5-cm-hair segments was 670 and 340 pg/mg, respectively (Table 2). Applying the SoHT cut-off of 200 pg/mg for amphetamines [3], 12 of 13 hair samples with a 1-cm- segment and 10 of 13 hair samples with a 5-cm-segment would have tested positive after two representative MDMA intakes (each 125 mg) within a time period of about a maximum of 3 months (94 days, median: 42 days) (Table 2). Hair samples of participants (no. 2, 7, 10, 13, 14) in which the two intakes were represented by two different hair segments would theoretically have tested positive for MDMA only after one single intake (Figure 2). Our findings demonstrate that the length of a hair segment should be considered for the interpretation of hair analysis results and application of cut-offs.

MDMA and MDA in nail samples and comparison to hair

In two previous studies, it was demonstrated that a single drug dose is detectable in fingernail clippings [20,19]. The present study describes nail findings following two MDMA administrations. All nail samples were positive for MDMA enantiomers with a first and second administration ranging from 9 to 77 and 26 to 128 days, respectively, prior to sample collection. MDMA concentrations were below 500 pg/mg in all nail samples (Table 2). There was no relationship between drug concentrations in nail samples and days passed after the first or second MDMA intake. Nail incorporation takes place via nail matrix and nail bed [20,19]. In nail clipping drugs can be detected earlier after incorporation via nail bed than via nail matrix. Drugs can be detected in nail scrapings as soon as outgrowth of incorporated drug at the nail matrix starts reaching the exposed nail surface. Moreover, sweat/sebum contribute to drug disposition into the nail leading to drug detection already within hours after the intake. In most subjects, MDMA and MDA concentrations of enantiomers and their sum were higher in nail clippings than in scrapings (Table 2). In a recent study with cocaine users, cocaine concentrations were higher in scrapings compared to clippings which was explained by external contamination from drug residues in the environment [12]. In the present study, contamination can be excluded as intake was performed in the scope of a controlled study. Lower levels in scrapings may be explained by drug extraction during daily hygiene as the nail surface which is collected for scrapings is more exposed to drug extraction [12].

Our study included the analysis of 13 fingernail and 3 toenail samples. In two toenail samples (no. 10, 16), concentrations of MDMA and MDA enantiomers and their sum were in the same range as in fingernail samples (Table 2). Highest MDMA and MDA concentrations were detected in the toenail clipping of participant no. 12. Interestingly, the MDMA concentration in the respective toenail scraping was not

significantly higher compared to other samples and also MDA was not detectable (Figure 1C-F, Table 2). Because toenails grow slower than fingernails [10,11,9], they represent a longer window of detection for drugs. Therefore, higher drug concentrations in toenail samples may be explained by drug use long time prior to the study. The 5-cm long hair sample of participant no. 12 was too short to verify this assumption.

In 8 of 13 subjects, the sum of concentrations of MDMA enantiomers were lower in nail clippings compared to the 5-cm-hair segment, in two participants comparable concentrations were found (Table 2). Cirimele et al. reported for one case slightly higher MDMA and MDA concentrations in fingernail scrapings compared to hair [18]. It should be noted that hair and nail samples represent different time frames: about 5 months based on an average growth rate if the hair sample is 5 cm long, for nail clippings a median window of detection of 3.5 months was determined [20]. To our knowledge, there is no report on windows of detection for scrapings. Further, the bizonal drug incorporation mechanism into nails makes time-resolved analysis impossible unless clippings are collected continuously over a longer period [20]. Despite of these limitations, our findings illustrate that nail clippings can be suitably used for retrospective monitoring of drug history alternatively to hair, especially in cases where hair is not long enough or cosmetically treated, as a positive result for MDMA and MDA proves MDMA intake. Nail scrapings seem less suitable as not all samples were positive for the metabolite MDA after two single MDMA intakes.

Enantiomer ratios of MDMA and MDA in hair and nail samples

R-MDMA concentrations were significantly higher than *S*-MDMA concentrations in all hair and nail samples whereas no difference was observed for concentrations of MDA enantiomers (Figure 1, Figure 3). Participant no. 1, who was a CYP2D6 poor

metabolizer, had the highest *R/S*-MDMA ratio in the hair segment corresponding to the time of intake and the nail scraping whereas the ratio in the nail clipping was in the range of the extensive CYP2D6 metabolizers (Figure 3A). To our knowledge, there are two reports on chiral analysis of MDMA and MDA in hair samples from suspected or self-declared MDMA users [41-43]. Enantiomeric ratios for MDMA and MDA from our controlled intake study are in concordance with hair findings reported by Martins et al. from a study with self-declared MDMA users after controlled MDMA administration [41]. Fallon et al. demonstrated that plasma and urine concentrations of *R*-MDMA exceed those of *S*-MDMA at all time-points analyzed over 24 h after administration whereas *S*-MDA plasma concentrations exceeded those of *R*-MDA [31]. Our study results display the same trend for hair and nail samples.

Similar enantiomeric ratios of MDMA and MDA in hair, nail clippings and scrapings may suggest that drug disposition into these three matrices are comparable.

Within subject comparison revealed two patterns for *R/S*-MDMA concentration ratios along the hair segments: a decreasing or increasing trend from proximal to distal segments. The decreasing and increasing trend corresponded to decreasing and increasing sum of concentrations of MDMA enantiomers along the hair segments, respectively, except for participant no. 18 (Figure 3 and 4). The highest *R/S*-MDMA ratio in the proximal (participant no.1, 12, 16, 4, 17, 2, 11,13, 7) or distal (participant no. 6 and 10) segments corresponded to recent or earlier MDMA administration, respectively (Figure 4). In case of recent MDMA intake lower *R/S*-MDMA ratios in distal segments may be explained by sweat incorporation. There was no trend for MDA enantiomeric ratios along the hair segments (data not shown).

Participant no. 14 received the two MDMA doses with a time interval of 2 months, which was monitored by significantly increased MDMA concentrations in both

corresponding hair segments (Figure 2). The longer time interval of intakes resulted in comparable *R/S*-MDMA ratios along the hair segments (Figure 4B).

Metabolite ratios of MDA to MDMA in hair and nail samples

The determination of metabolite to parent drug ratios was recommended for the discrimination between drug intake and external contamination of a hair sample [44] and has proved to be useful [45-47]. For this purpose, metabolite ratios have to be determined after controlled intake of drug in a scope where contamination can be excluded. As routine analysis is typically achiral, the sum of the respective enantiomer concentrations were used for the metabolite ratios. In our study, median ratios of MDA to MDMA concentrations in hair following administration of pharmaceutical racemic MDMA were 0.047 in corresponding and 0.054 in not corresponding hair segments (Figure 5). Other authors reported similar metabolic ratios after street MDMA use (0.04 to 0.06) [48,49] or controlled administration (< 0.09) [41].

Conclusion

A suitable analytical method for the chiral analysis of MDMA and MDA in hair and nail samples was presented. The study demonstrates that administrations of two representative single doses of MDMA can be detected in the hair segments corresponding to the time of intake and in all nail samples. As all nail clippings were positive for MDMA and the metabolite MDA irrespective of days passed after intake, clippings appears a suitable alternative to hair for prove of MDMA intake. Since no significant difference was observed for enantiomeric ratios in hair and nail samples incorporation mechanisms into both keratinized matrices may be comparable.

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FIGURES

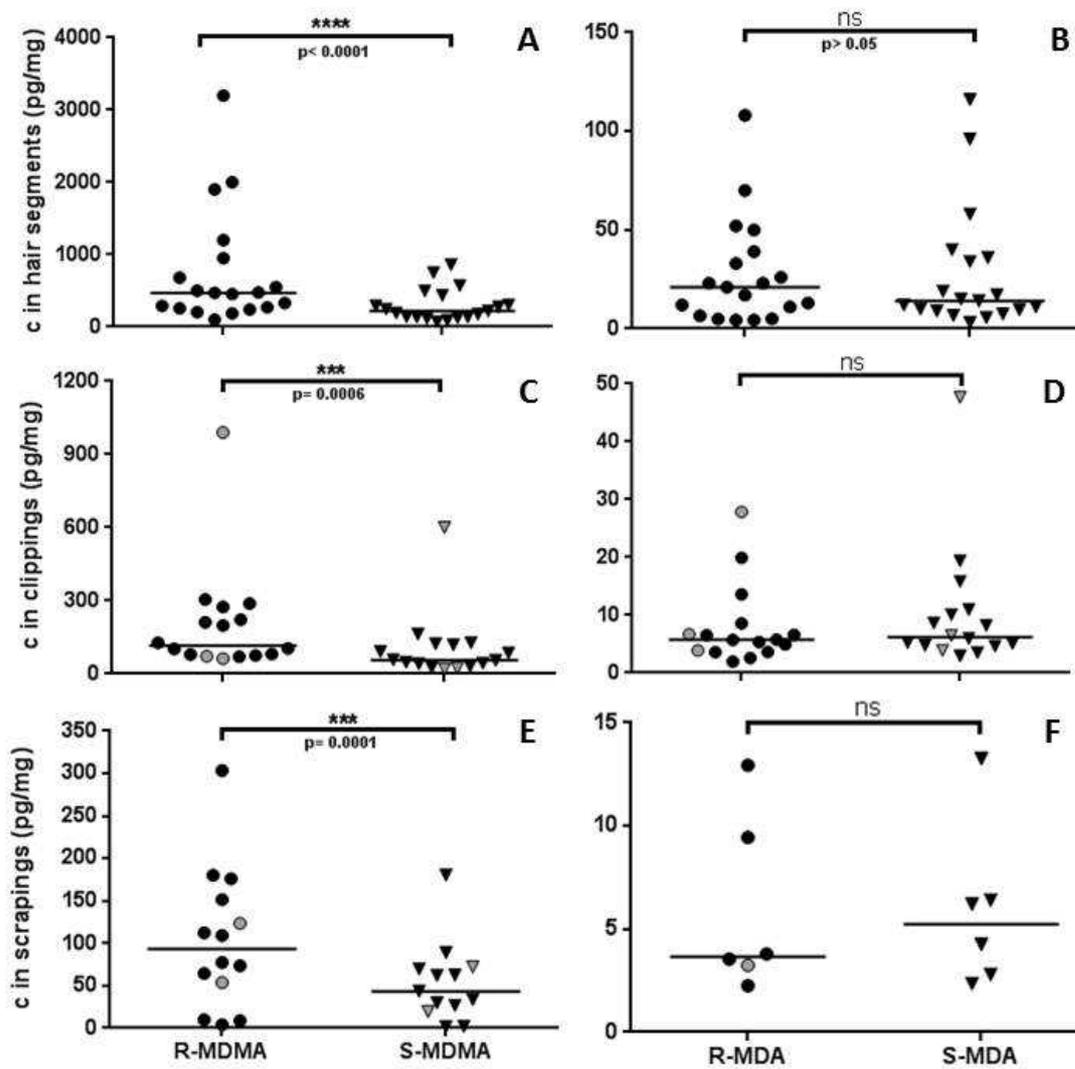


Figure 1. Measured R/S-MDMA and R/S-MDA concentrations in hair segments corresponding to the calculated time of intake (A+B), in nail clippings (C+D) and nail scrapings (E+F); black circle/triangle= specimen from fingernail, grey circle/triangle= specimen from toenail. Statistical analysis by paired, two-tailed, non-parametric t-test; $p < 0.001$: very significant (***), $p > 0.05$: not significant (ns).

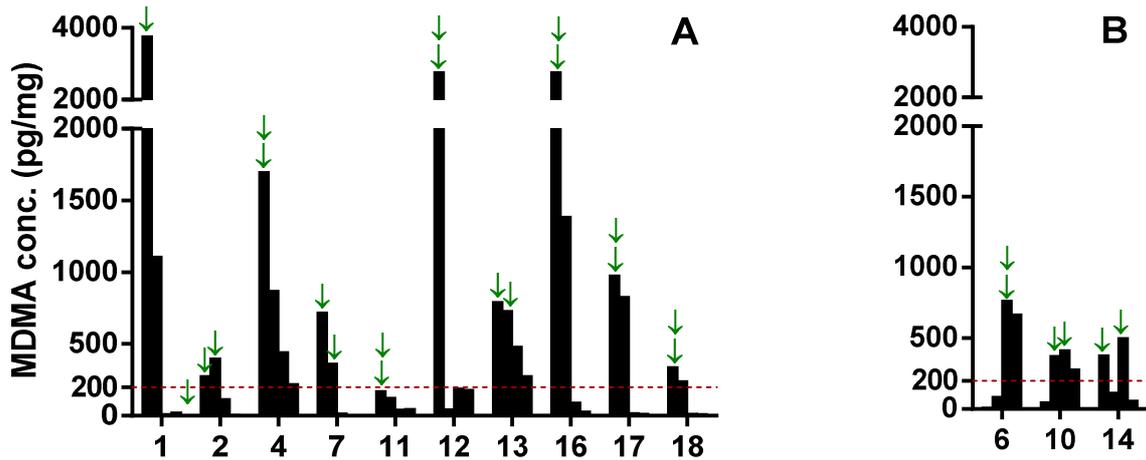


Figure 2. Calculated sum of concentrations of MDMA enantiomers in four hair segments (1+1+1+2 cm, from proximal to distal) from participants with recent (A), earlier (participant no. 6 and 10) and recent with earlier (participant no. 14) intake (B). Green arrows indicate hair segments corresponding to the time of MDMA intake. Red dotted line: SoHT cut-off for MDMA in hair samples. Participant no. 1 was phenotyped a CYP2D6 poor metabolizer.

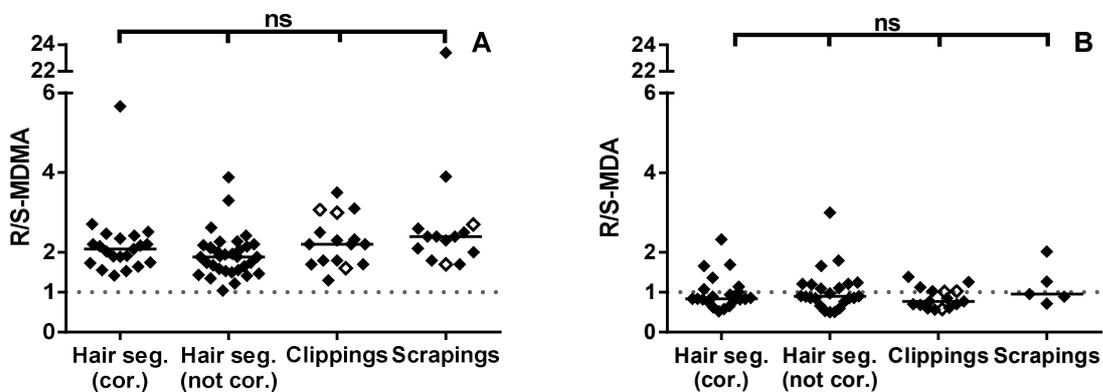


Figure 3. Enantiomeric concentration ratios of *R/S*-MDMA (A) and *R/S*-MDA (B) in hair segments (seg.) corresponding (cor.) and not corresponding (not cor.) to the time of intake, nail clippings and scrapings; black square= fingernail, white square= toenail. Statistical analysis by parametric one-way ANOVA test; $p > 0.05$: not significant (ns). Dotted line represents racemic 1:1 ratio. Outlier with *R/S*-MDMA ratio of 5.6 and 23 (A) was phenotyped a CYP2D6 poor metabolizer (participant no. 1).

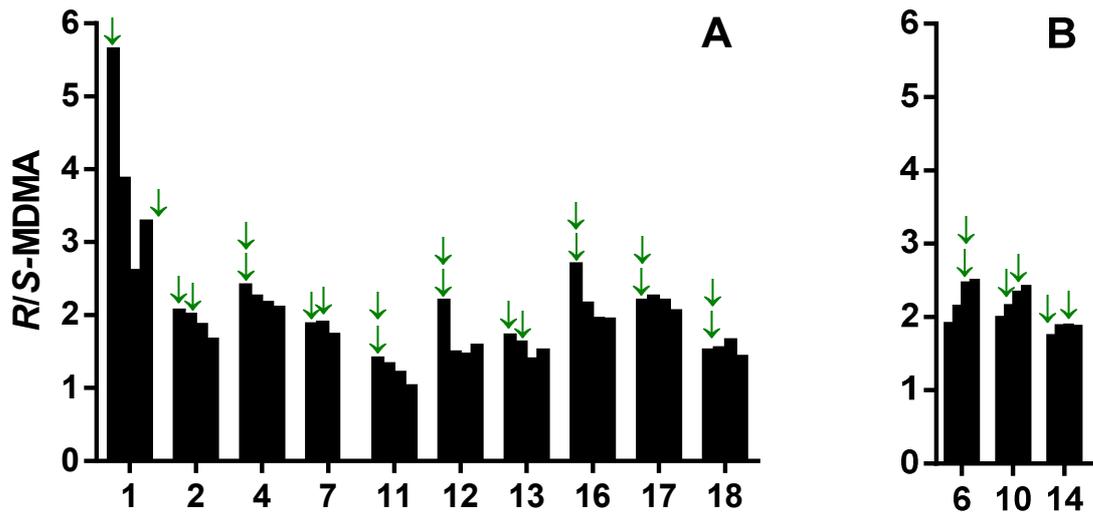


Figure 4. Enantiomeric R/S-MDMA concentration ratios in four hair segments (1+1+1+2 cm, from proximal to distal) from participants with recent (A), earlier (B; no. 6 and 10) and recent with earlier intake (B; no. 14) intake. Green arrows indicate hair segments corresponding to the time of MDMA intake. Participant no. 1 was phenotyped a CYP2D6 poor metabolizer.

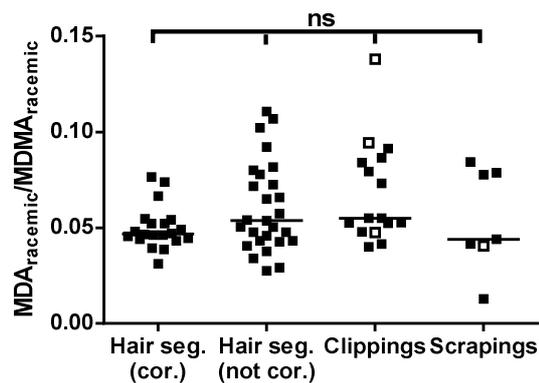


Figure 5. Metabolic ratios of sum of concentrations of MDA and MDMA enantiomers in hair segments corresponding (cor.) and not corresponding (not cor.) to the time of intake, nail clippings and scrapings; black square= fingernail, white square= toenail. Statistical analysis by non-paired, two-tailed, non-parametric t-test; $p > 0.05$: not significant (ns).

TABLES

Table 1. Validation data for analysis of nail and hair samples.

Analyte (IS)	R-MDMA (R-MDMA-D5)		S-MDMA (S-MDMA-D5)		R-MDA (R-MDA-D5)		S-MDA (S-MDA-D5)	
Hair analysis								
LOD (pg/mg)	0.125							
LLOQ (pg/mg)	0.5							
Linearity (pg/mg)	0.5; 50; 200; 300; 400; 500							
QC level	LOW	HIGH	LOW	HIGH	LOW	HIGH	LOW	HIGH
QC conc. (pg/mg)	0.625	450	0.625	450	0.625	450	0.625	450
Accuracy (%)	-9.8	6.1	-15.9	6.7	-10.5	5.2	-1.3	-2.4
Repeatability (%)	8.6	4.7	4.2	4.7	3.7	5.1	8.0	5.8
Precision (%)	8.6	5.8	11.7	5.6	11.7	9.4	8.6	8.5
ME, mean	97	114	106	134	124	127	124	111
±SD (%)	±2.8	±3.7	±6.4	±9.8	±2.8	±7.3	±5.4	±17.6
Nail analysis								
LOD (pg/mg)	0.5							
LLOQ (pg/mg)	2.0							
Calibrators (pg/mg)	2.0; 200; 800; 1200; 1600; 2000							
QC level	LOW	HIGH	LOW	HIGH	LOW	HIGH	LOW	HIGH
QC conc. (pg/mg)	2.5	1800	2.5	1800	2.5	1800	2.5	1800
Accuracy (%)	6.5	-1.8	0.1	-1.2	3.9	0.2	-6.1	2.5
Repeatability (%)	7.4	3.7	14.3	7.5	6.3	4.2	6.1	6.8
Precision (%)	11.2	11.8	15.0	14.9	7.5	14.3	8.5	10.6
ME, mean	116	106	126	130	122	107	113	103
±SD (%)	±4.1	±2.1	±6.2	±2.3	±9.4	±4.1	±14.4	±6.1

IS= internal standard; LOD= limit of detection; LLOQ= limit of quantification; QC= Quality control; conc.= concentration

Table 2. Measured concentrations of MDMA and MDA enantiomers and calculated sum of concentrations of MDMA and MDA enantiomers in hair segments and fingernail (and/or toenail) scrapings and clippings

Subject no.	Days * ¹)	Days * ²)	Hair colour	Hair segments								Nail samples								
				MDMA concentration (pg/mg)											scrapings			clippings		
				A (0-1 cm) proximal		B (1-2 cm)		C (2-3 cm)		D (3-5 cm) distal		0-5 cm (calc.)								
				R	S	R	S	R	S	R	S	Sum R+S (calc.)	R	S	Sum R+S (calc.)	R	S	Sum R+S (calc.)		
01 (PM)	128	23	dark brown	≈3200	≈570	≈880	227	9.3	3.5	19	5.7	980	10	<LOQ	10	306	86	392		
02 (EM)	55	41	brown	187	90	271	134	77	41	3.7	2.2	160	8.8	2.2	11.0	105	48	153		
03 (EM)	41	14	N/A	N/A								303	180	483	222	127	349			
04 (EM)	39	28	dark brown	≈1200	497	≈610	266	303	139	151	71	650	151		214	199	91	290		
06 (EM)	94	77	dark brown	7.1	3.7	59	28	≈550	221	478	190	310	113	44	157	127	55	182		
07 (EM)	42	17	brown	470	249	238	125	11	6.5	N/A		367	65	27	92	81	44	125		
09 (EM)	67	46	N/A	N/A								4.3	2.0	6.3	103	42	145			
10 (EM)	64	49	brown	32	16	257	119	292	124	199	82	220	54* ³)	20* ³)	74	72* ³)	23* ³)	95		
11 (EM)	26	9	light brown	101	71	72	54	25	20	24	23	78	N/A			75	57	132		
12 (EM)	28	17	dark brown	≈1900	≈860	28	18	113	77	111	70	~640	124* ³)	72* ³)	196	990* ³)	600* ³)	1590		
13 (EM)	49	13	light brown	500	291	454	276	281	199	166	109	460	180	89	269	211	123	334		
14 (EM)	78	23	brown	242	139	76	40	331	174	40	21	210	77	34	111	70	33	103		
16 (EM)	35	14	dark brown	≈2000	≈750	≈950	437	60	30	19	10	~850	74	30	104	80	27	107		
													N/A			62* ³)	20* ³)	82		
17 (EM)	35	17	dark brown	≈680	305	≈570	253	13	5.9	9.3	4.5	370	176	70	246	275	118	393		
18 (EM)	35	20	dark brown	206	134	148	95	9.8	5.9	7.7	5.4	120	109	62	171	288	163	451		
				MDA concentration (pg/mg)																
01 (PM)	128	23	dark brown	116	50	38	21	1.0	<LOQ	1.7	<LOQ	46	nd	nd	nd	20	16	36		
02 (EM)	55	41	brown	7.5	4.4	9	6.6	2.7	2.4	1.2	nd	6.7	nd	nd	nd	3.6	4.7	8.4		
03 (EM)	41	14	N/A	N/A								nd	6.2		5.8	8.2	14.0			
04 (EM)	39	28	dark brown	34	33	19	21	11	13	5.3	5.9	28	nd	nd	nd	6.6	8.6	15.2		
06 (EM)	94	77	dark brown	<LOQ	nd	3.1	2.6	19	23	14	17	16	3.5	2.8	6.3	3.6	6.0	9.6		
07 (EM)	42	17	brown	17	21	7.4	8.8	1.5	0.5	N/A		19	nd	nd	nd	5.7	5.1	11		
09 (EM)	67	46	N/A	N/A								nd	nd	nd	5.3	5.2	11			
10 (EM)	64	49	brown	1.8	1.4	9.5	4.4	11	12	9.3	11	12	3.2* ³)	nd* ³)	3.2	6.7* ³)	6.5* ³)	13		
11 (EM)	26	9	light brown	3.2	5.1	2.5	4.7	1.1	1.1	0.6	1.2	3.9	N/A			2.6	4.7	7.3		
12 (EM)	28	17	dark brown	58	70	1.0	<LOQ	3.9	4.3	3.4	4.0	29	nd* ³)	nd* ³)	nd* ³)	28* ³)	48* ³)	76		
13 (EM)	49	13	light brown	15	26	12	23	6.9	14	3.7	5.6	21	9.5	13.2	22.7	6.5	11	18		
14 (EM)	78	23	brown	6.8	11	2.2	3.7	10	13	1.3	1.5	9.9	2.3	2.4	4.7	2.0	2.9	4.9		
16 (EM)	35	14	dark brown	96	108	40	52	3.0	4.0	1.6	1.4	61	3.8	4.3	8.1	4.9	3.5	8.4		
													N/A			3.9* ³)	3.9* ³)	7.8		
17 (EM)	35	17	dark brown	36	39	34	32	1.0	0.8	0.7	<LOQ	29	12.9	6.4	19.3	14	19	33		
18 (EM)	35	20	dark brown	5.7	4.9	4.4	2.7	nd	0.6	nd	1.0	3.9	nd	nd	nd	8.6	10	19		

*¹) Days past after first MDMA intake; *²) Days past after second MDMA intake; *³) toenail; calc.= calculated; ≈: approximately; N/A: no sample available; PM: phenotyped as CYP2D6 poor metabolizer; EM: phenotyped as CYP2D6 extensive metabolizer; nd: not detected; marked bold= time period of MDMA intake

