

Metabolic fate and detectability of the new psychoactive substances 2-(4-bromo-2,5-dimethoxyphenyl)-N-[(2-methoxyphenyl)methyl]ethanamine (25B-NBOMe) and 2-(4-chloro-2,5-dimethoxyphenyl)-N-[(2-methoxyphenyl)methyl]ethanamine (25C-NBOMe) in human and rat urine by GC-MS, LC-MSⁿ, and LC-HR-MS/MS approaches

Achim T. Caspar^a, Simon D. Brandt^b, Andreas E. Stoever^c, Markus R. Meyer^a, Hans H. Maurer^{a,*}

^a*Department of Experimental and Clinical Toxicology, Institute of Experimental and Clinical Pharmacology and Toxicology, Saarland University, D-66421 Homburg (Saar), Germany*

^b*School of Pharmacy and Biomolecular Sciences, Liverpool John Moores University, James Parsons Building, Byrom Street, Liverpool L3 3AF, UK*

^c*Institute of Legal Medicine, University of Munich, D-80336 Munich, Germany*

*Corresponding author at Saarland University, Department of Experimental and Clinical Toxicology, Homburg (Saar), Germany

E-mail address: hans.maurer@uks.eu (H.H. Maurer)

ABSTRACT

25B-NBOMe and 25C-NBOMe are potent 5-HT_{2A} receptor agonists that have been associated with inducing hallucinogenic effects in drug users and severe intoxications. This paper describes the identification of their metabolites in rat and human urine by liquid chromatography (LC)-high resolution (HR)-MS/MS, the comparison of metabolite formation in vitro and in vivo and in different species, the general involvement of human cytochrome-P450 (CYP) isoenzymes on their metabolism steps, and their detectability by standard urine screening approaches (SUSAs) using GC-MS, LC-MSⁿ, or LC-HR-MS/MS. Both NBOMe derivatives were mainly metabolized by *O*-demethylation, *O,O*-bis-demethylation, hydroxylation, and combinations as well as by glucuronidation and sulfation of the main phase I metabolites. For 25B-NBOMe, 66 metabolites could be identified and 69 for 25C-NBOMe. After application of low doses of both substances to rats, they were detectable mainly via their metabolites by both LC-based SUSAs. In case of acute intoxication, it was possible to detect 25B-NBOMe and its metabolites in an authentic human urine sample when using the GC-MS SUSA in addition to the LC-based SUSAs. Initial CYP activity screening revealed the involvement of CYP1A2 and CYP3A4 in hydroxylation and CYP2C9 and CYP2C19 in *O*-demethylation. The presented study demonstrated that 25B-NBOMe and 25C-NBOMe were extensively metabolized and detectable by both LC-based SUSAs.

Keywords:

25B-NBOMe

25C-NBOMe

new psychoactive substance

metabolism

cytochrome-P450

LC-MSⁿ

LC-HR-MS/MS

1. Introduction

According to annual drug reports published by the European Monitoring Centre for Drugs and Drug Addiction (EMCDDA) and United Nations Office on Drugs and Crime (UNODC) [1-4], the availability and abuse of new psychoactive substances (NPS) increased during the last few years. Besides synthetic cannabinoids, cathinones, opioids, and tryptamines, the group of phenethylamines gained more importance in the last years [5]. Among others, the so-called 2C-type phenethylamines have been a constant feature in the detection of NPS [6]. They were first described by Alexander Shulgin in his book PIHKAL [7]. Like many phenethylamines, they have powerful psychoactive and stimulating effects [7,8]. Although many of them have been scheduled, new and uncontrolled alternatives have emerged. Structure-activity relationship studies revealed that derivatization of the primary amine of the 2C partial structure with a 2-methoxybenzyl substituent significantly increased the affinity toward the serotonin 5-HT_{2A} receptor, thus, mediating potent hallucinogenic effects [9-12]. The resulting 2C derivatives, the so-called NBOMes (*N*-2-methoxybenzyl phenethylamines), represent a new group of potent phenethylamine hallucinogens with high abuse potential. 2-(4-Bromo-2,5-dimethoxyphenyl)-*N*-[(2-methoxyphenyl)methyl]ethanamine (25B-NBOMe, 2C-B-NBOMe), 2-(4-chloro-2,5-dimethoxyphenyl)-*N*-[(2-methoxyphenyl)methyl]ethanamine (25C-NBOMe, 2C-C-NBOMe), and 2-(4-iodo-2,5-dimethoxyphenyl)-*N*-[(2-methoxyphenyl)methyl]ethanamine, 25I-NBOMe, 2C-I-NBOMe) are among the most prevalent NBOMes. They are consumed depending on desired effects in reported dosages between 200-1,000 µg, administered orally, sublingually, buccally or insufflated as powder or in solution as nose spray [8,13-20]. In recent years, NBOMe consumption was described in the context of acute and severe intoxications and fatalities [8,14-18,21,22]. In some cases, an unintentional intake of NBOMes, sold as LSD or 2Cs, were found to be responsible for adverse events [8,15,19,22]. However, 25B-NBOMe has also been employed in positron emission tomography (PET) in human volunteers to assess binding of this ligand in distinct brain areas and at non-psychoactive dosage levels [23,24].

Due to high receptor affinity and functional activity as full agonists, comparatively low doses, comparable to LSD, are needed to induce psychoactive effects. Consequently, the resulting low blood plasma or urine concentrations can make it challenging to identify and characterize the intake of NBOMes. In urine, the concentration of compounds is generally higher than in blood, but in many cases, metabolites rather than the parent compounds are the targets. Therefore, metabolism studies are needed for the development of urine screening approaches. The comprehensive metabolism study for 25I-NBOMe revealed that it was extensively metabolized and that the parent compound was found in urine only in small amounts [25].

Recently, Wohlfarth et al. [26] described the metabolism of 25C-NBOMe and 25I-NBOMe in mice and human urine as well as in human hepatocytes, and the reported results were consistent with previously published human and rat data for 25I-NBOMe [25]. For 25B-NBOMe, only limited data on its biotransformation are available [27,28], and for both compounds, no comprehensive data appear to be available on their detectability. Therefore, the aims of the present study were to investigate the metabolism of 25B-NBOMe and 25C-NBOMe in rats and humans with LC-HR-MS/MS, to compare the results with in vitro and in vivo data and between different species, and to investigate their detectability by the authors' standard urine screening approaches (SUSA) by GC-MS, LC-MSⁿ, and LC-HR-MS/MS, respectively.

2. Experimental

2.1. Chemicals and reagents

25B-NBOMe hydrochloride and 25C-NBOMe hydrochloride were purchased by LGC Standards (Wesel, Germany). Isolute HXC cartridges (130 mg, 3 mL) were obtained from Biotage (Uppsala, Sweden), isocitrate and isocitrate dehydrogenase from Sigma (Taufkirchen, Germany), NADP⁺ from Biomol (Hamburg, Germany), acetonitrile (LC-MS grade), ammonium formate (analytical grade),

formic acid (LC-MS grade), methanol (LC-MS grade), mixture (100,000 Fishman units/mL) of glucuronidase (EC No. 3.2.1.31) and arylsulfatase (EC No. 3.1.6.1) from *Helix Pomatia*, and all other chemicals and reagents (analytical grade) from VWR (Darmstadt, Germany). The baculovirus-infected insect cell microsomes (Supersomes) containing 1 nmol/mL of human cDNA-expressed CYP1A2, CYP2A6, CYP2B6, CYP2C8, CYP2C9, CYP2C19, CYP2D6, CYP2E1 (2 nmol/mL), CYP3A4, or CYP3A5 (2 nmol/mL) were obtained from Corning (Amsterdam, The Netherlands). After delivery, the CYPs were thawed at 37°C, aliquoted, snap-frozen in liquid nitrogen, and stored at -80°C until use.

2.2. Urine samples

According to an established study design [29], the investigations were performed using rat urine samples from male Wistar rats (Charles River, Sulzfeld, Germany) for toxicological diagnostic reasons according to German law. Both compounds were administered in an aqueous suspension by gastric intubation of a single 10 mg/kg body weight (BW) dose for identification of the metabolites and of 0.1 mg/kg BW for screening (dose calculated based on common single dose reported in trip reports (<https://www.erowid.org>) and scaled by dose-by-factor approach from man to rat according to Sharma and McNeill [30]), respectively. The rats were housed in metabolism cages for 24 h, having water ad libitum. Urine was collected separately from feces over a 24 h period. Blank urine samples were collected before drug administration to verify that the samples were free of interfering compounds. The samples were directly analyzed and then stored at -20°C.

In addition, for 25B-NBOMe, an authentic ante mortem human urine sample after unintentional intake of an unknown dose of 25B-NBOMe (declared as 2C-B) submitted to the authors' laboratory for toxicological diagnostics was also analyzed.

2.3. Sample preparation

2.3.1. Sample preparation for identification of phase I metabolites by LC-HR-MS/MS

According to a published procedure [29], 2 mL of urine was adjusted to pH 5.2 with acetic acid (1 M, approximately 50 μ L) and incubated at 50 °C for 2 h with 50 μ L of a mixture of glucuronidase and arylsulfatase. The urine sample was then loaded on an HCL cartridge previously conditioned with 1 mL of methanol and 1 mL of water. After passage of the sample, the cartridge was washed with 1 mL of water, 1 mL of 0.01 M hydrochloric acid, and again with 1 mL of water. The acidic and neutral compounds were eluted with 1 mL of methanol into a 1.5 mL reaction vial and evaporated to dryness under a stream of nitrogen. In the same reaction vial, the basic compounds were eluted with 1 mL of a freshly prepared mixture of methanol/aqueous ammonia 32% (98:2, v/v). After another evaporation step the residues were reconstituted with 50 μ L of a mixture of eluent A and B (1:1, v/v) for LC-HR-MS/MS analysis. A 5- μ L aliquot was then injected onto the LC-HR-MS/MS.

2.3.2. Sample Preparation for the identification of phase I metabolites and MBPs

According to a published procedure [29], 100 μ L of urine was mixed with 500 μ L of acetonitrile for precipitation. After shaking and centrifugation, the supernatant was gently evaporated to dryness and reconstituted in 50 μ L of a mixture of 10 mM aqueous ammonium formate buffer and acetonitrile (1:1, v/v) and 5 μ L injected onto the LC-HR-MS/MS system.

2.4. Incubations for initial CYP activity screening studies

According to standard procedures [29,31], microsomal incubations were performed at 37°C at a concentration of 25 μ M 25B-NBOMe and 25C-NBOMe, respectively, with the CYP isoenzymes (75 pmol/mL, each) CYP1A2, CYP2A6, CYP2B6, CYP2C8, CYP2C9, CYP2C19, CYP2D6, CYP2E1, CYP3A4, or CYP3A5 for 30 min. Besides enzymes and substrates, the incubation mixtures (final

volume, 50 μ L) contained 90 mM phosphate buffer (pH 7.4), 5 mM Mg^{2+} , 5 mM isocitrate, 1.2 mM $NADP^+$, 0.5 U/mL isocitrate dehydrogenase, and 200 U/mL superoxide dismutase. For incubations with CYP2A6 and CYP2C9, phosphate buffer was replaced with 45 mM or 90 mM Tris buffer, respectively, according to the Gentest manual. Reactions were initiated by addition of the CYP enzymes and stopped with 50 μ L of ice-cold acetonitrile. The solution was centrifuged for 2 min at 14,000 rpm; 70 μ L of the supernatant phase were transferred to an autosampler vial and 5 μ L injected onto the LC-HR-MS/MS system.

2.5. LC-HR-MS/MS instrumentation for identification of phase I and II metabolites and CYP initial screening

According to a published procedure [32], the extracts were analyzed using a ThermoFisher Scientific (TF, Dreieich, Germany) Dionex UltiMate 3000 RS pump consisting of a degasser, a quaternary pump, and an UltiMate autosampler, coupled to a TF Q-Exactive Plus system equipped with a heated electrospray ionization (HESI)-II source. The instrument was used in positive or in negative ionization mode. Mass calibration was performed prior to analysis according to the manufacturer's recommendations using external mass calibration.

Gradient elution was run on a TF Accucore PhenylHexyl column (100 mm x 2.1 mm, 2.6 μ m). The mobile phases consisted of 2 mM aqueous ammonium formate containing formic acid (0.1%, v/v) and acetonitrile (1%, v/v) (pH 3, eluent A) and 2 mM ammonium formate solution with acetonitrile:methanol (50:50, v/v) containing formic acid (0.1%, v/v) and water (1%, v/v) (eluent B). The gradient and flow rate were programmed as follows: 0-1 min hold 99% A, 1-16 min 95% A to 5% A, 16-18 min hold 5% A, and 18-20 min hold 99% A, constantly at 500 μ L/min.

The HESI-II source conditions were as follows: sheath gas, 60 arbitrary units (AU); auxiliary gas, 10 AU; spray voltage, 3.00 (positive polarity) and -4.00 kV (negative polarity); heater temperature, 320°C; ion transfer capillary temperature, 320°C; and S-lens RF level, 60.0. Mass spectrometry was

performed in positive and negative polarity mode using full scan (FS) data and a subsequent data dependent acquisition (DDA) mode with an inclusion list on the masses of interest (phase I or phase II metabolites). Additionally, DDA runs without inclusion list (positive and negative mode) were performed to detect unexpected metabolites.

The settings for FS data acquisition were as follows: resolution, 35,000; microscans, 1; automatic gain control (AGC) target, $1e6$; maximum injection time (IT), 120 ms; and scan range, m/z 100–700.

The settings for the DDA mode with and without an inclusion list were as follows: resolution, 17,500; microscans 1, AGC target, $2e5$; maximum IT, 250 ms; isolation window, 1.0 m/z , HCD with stepped normalized collision energy (NCE), 17.5, 35, and 52.5%; spectrum data type, profile; and underfill ratio, 0.5%. For the run without inclusion list, the five most intense precursor ions were transferred to an exclusion list for 1 s (dynamic exclusion).

For analyzing the initial CYP activity screening, the MS settings and the mobile phases as well as the gradient and flow rate were the same with the same inclusion list as for identification of phase I metabolites.

2.6. Standard urine screening procedures (SUSAs)

The SUSAs were performed as described in the following references: GC-MS SUSA [33,34], LC-MSⁿ SUSA [25,35], and LC-HR-MS/MS SUSA [25,36].

3. Results and discussion

3.1. Identification of metabolites

3.1.1. Identification of 25B-NBOMe and 25C-NBOMe and their phase I metabolites via HR-MS/MS fragmentation

208 The HR-MS/MS fragmentation patterns and metabolite formation of 25B-NBOMe and 25C-
209 NBOMe were similar to those described for 25I-NBOMe [25]. Briefly, and for discussion purposes,
210 the molecules were viewed as two distinct parts, i.e. the 4-halogenated 2,5-
211 dimethoxyphenethylamine (2C) part and the *N*-(2-methoxybenzyl) (NBOMe) part. Due to the high
212 number of metabolites, the fragmentation patterns could not be discussed in detail for all metabolites
213 and only the typical fragment ions used for identification will be discussed.

214 In general, for both compounds and their metabolites, the precursor masses and the most abundant
215 fragment ions formed from unmodified or modified NBOMe parts were used to identify the
216 corresponding metabolites. As expected, the fragment ions formed by the NBOMe part were identical
217 for 25B-NBOMe and 25C-NBOMe. To confirm the predicted chemical structure of the metabolites,
218 the corresponding 2C fragment ions (Table S1 in the electronic supplementary data for 25B-NBOMe
219 and Table S2 for 25C-NBOMe) were used. For the *N*-dealkylated metabolites, no fragment ions of
220 the NBOMe part could be detected, but characteristic 2C fragmentation patterns for the bromo and
221 chloro analogues (Tables S1 and S2). The precursor masses (PM) are given with the calculated exact
222 masses.

223 25B-NBOMe (B1; PM at m/z 380.0856, $M+H^+$) showed a fragmentation pattern characteristic also
224 for most of the detected metabolites. The most abundant fragment ion of m/z 121.0653 represented
225 the cleavage of the NBOMe moiety, followed by the loss of the methoxy group (-30.0105 u)
226 producing the tropylium ion of m/z 91.0548. The fragment ions representing the 2C part showed low
227 abundances of at least 1 % (Table S1). The fragment ion of m/z 258.0124 representing the 2C-B
228 iminium ion resulted from benzyl cleavage. A loss of NH (- 15.0109 u) formed the fragment ion of
229 m/z 243.0021 followed by a loss of a methyl radical (- 15.0235 u) of one of the two methoxy groups in
230 the 2C part resulting in fragment ion of m/z 227.9786. For the MS² spectrum of 25I-NBOMe, a
231 rearrangement was described in the literature [25]. In the parent spectrum of 25B-NBOMe, one
232 fragment ion could also be formed by the same rearrangement. The fragment ion of m/z 363.0596
233 resulted from a loss of ammonia (-17.0263 u) and appeared consistent with the postulated

rearrangement reaction. Few MS² spectra of metabolites also showed possible rearranged fragment ions.

The fragmentation patterns of 25C-NBOMe (C1; PM at m/z 336.1361, M+H⁺) corresponded to those of 25B-NBOMe and 25I-NBOMe. Similarly, the most abundant fragment ions in MS² were formed by cleavage of the NBOMe moiety producing fragment ions of m/z 91.0548 and 121.0653. Also, the fragment ions representing the 2C part showed a lower abundance of about 1 % (Table S2). The fragment ions of m/z 214.0629, m/z 199.0526, and m/z 184.0291 represented the 2C-C iminium ion, the subsequent loss of NH (- 15.0109 u), and the loss of a methyl radical (- 15.0235 u) of one of the two methoxy groups, respectively. In the spectrum of 25C-NBOMe, no fragment ions indicating the rearrangement were detected, possibly due to low relative abundance. However, in the MS² spectra of some 25C-NBOMe metabolites (e.g. *O*-demethyl metabolite isomers 1 and 2, C16 and C17), some rearranged fragment ions could also be detected. Overall, 35 phase I metabolites could be detected for 25B-NBOMe in urine and 36 for 25C-NBOMe, respectively. All phase I metabolites are listed in Tables S1 and S2 in the electronic supplementary data.

For metabolite identification based on the MS² spectra, in most cases the representative fragment ion for the NBOMe part was used. Unmodified NBOMe parts led to a fragment ion of m/z 121.0653. The presence of this fragment ion led to the suggestion that the expected modification took place at the 2C part based on the predicted precursor mass for the metabolite. An unchanged fragment ion of m/z 121.0653 could be seen for the parent compounds (B1 and C1) as well as for mono- and bis-demethylated (B13, B14, B8 and C16, C17, C10), mono-hydroxylated (B29, B32 and C31, C32), bis-hydroxylated (B35), combined mono-demethylated with mono-hydroxylated (B22), dehydrogenated (B20 and C24), dehydrogenated combined with mono-demethylated (B12 and C13), and mono-hydroxylated (B27 and C30) metabolites. On the other hand, the fragment ion of m/z 137.0603 represented mono-hydroxylation at the NBOMe part (B16, B23, B28, B30, B31 and C19, C25, C27, C33, C35). At the NBOMe moiety, bis-hydroxylation led to the fragment ion of m/z 153.0552 (B33, B34 and C34, C36) and *O*-demethylation to fragment ion of m/z 107.0497 (B7, B9, B10, B11, B15,

260 B19, B21 and C8, C9, C11, C12, C14, C18, C21, C23, C26). The fragment ion of m/z 107.0497
261 could also be found for the NBOMe mono-hydroxylated (B16, B23, B28, B31 and C19, C25, C27,
262 C33) or NBOMe bis-hydroxylated (B34 and C34, C35, C36) metabolites, but consistently in
263 combination with the fragment ions of m/z 137.0603 or m/z 153.0552 as mentioned above. Therefore,
264 the absence of the fragment ions of m/z 137.0603 and 153.0552 indicated *O*-demethylation at the
265 NBOMe part. *O*-Demethylation combined with mono-hydroxylation led to fragment ion of m/z
266 123.0446 (B17, B18, B24, B25 and C20, C22, C28, C29). As mentioned above, all *N*-
267 demethoxybenzyl metabolites were identified based on the 2C part fragmentation patterns (B2–B6
268 and C2–C7).

269 In summary, the fragmentation patterns of both NBOMes corresponded to those of 25I-NBOMe.
270 Some compound-related characteristics were found for the bromo and chloro analogues as already
271 described for 25I-NBOMe. All metabolites, which were *O*-demethylated at the NBOMe part (m/z
272 107.0497), showed higher abundances for fragment ions representing the 2C part probably due to a
273 hydrogen bond between the nitrogen and the hydroxy group resulting from *O*-demethylation at the
274 NBOMe part [25]. In addition, for these metabolites, the corresponding 2C fragment ion carrying the
275 nitrogen was represented by the 2C primary amine instead of the 2C iminium ion found for the parent
276 compounds or metabolites, which were not *O*-demethylated at the NBOMe part.

277 It was not possible in this study to identify the demethylated position of the methoxy group (2'- or 5'-
278 position) or the position at which the NBOMe part was hydroxylated. Nevertheless, Wohlfarth et al.
279 [26] synthesized six potential 25C-NBOMe metabolites (2'- and 5'-*O*-demethyl-25C-NBOMe and 3-
280 /4-/5- and 6-hydroxy-25C-NBOMe) to confirm the exact position of the metabolic reaction. They
281 observed that both in vivo samples (mouse and human urine) showed prevalence for *O*-demethylation
282 at the 5'-position. Furthermore, they observed that the most intense signal for a mono-hydroxylated
283 metabolite was detected for 5-hydroxy-25C-NBOMe in human urine as well as in mouse urine. In
284 general, Wohlfarth et al. described the same main metabolic steps compared to the present study and

25I-NBOMe [25]. In accordance, Leth-Petersen et al. [28] described that the main metabolic step of 25B-NBOMe was also the 5'-*O*-demethylation in humans and pigs.

3.1.2. Identification of 25B-NBOMe and 25C-NBOMe and their phase I metabolites via HR-MS/MS fragmentation

The phase II metabolite formation and fragmentation patterns were very similar for both compounds and comparable with those described for 25I-NBOMe. For both compounds, the precursor masses and the most abundant fragment ions formed from unmodified or modified NBOMe part were used to identify the corresponding phase II metabolites. 2C fragment ions were used to confirm the predicted metabolites.

Overall, 31 phase II metabolites could be identified for 25B-NBOMe and 33 for 25C-NBOMe. A list of all phase II metabolites is given in Tables S3 and S4 in the electronic supplementary data. All glucuronides eliminated glucuronic acid (- 176.0321 u) and all sulfates sulfuric acid (- 79.9568 u). Thus, the rest of the spectra of phase II conjugates was in accordance with the spectrum of the corresponding phase I metabolite. Also, for some phase II metabolites, fragment ions formed by conjugated partial structures could be used to elucidate the position of the conjugation.

As already described for 2C derivatives [37] and 25I-NBOMe the metabolites formed after *N*-demethoxybenzylation could further be conjugated by acetylation, glucuronidation, sulfation, or even combinations of them. Furthermore, in accordance to 25I-NBOMe, an *O,O*-bis-demethylation of the 2C part led to a hydroquinone partial structure, which could further be conjugated with glutathione (GSH). The degradation products of GSH conjugated metabolites could be found for both compounds. Also the described conjugation catalyzed by catechol-*O*-methyl-transferase (COMT) could be found for both NBOMes forming *O*-methyl metabolites (B33ME and C24ME, C36ME), after bis-hydroxylation at the NBOMe part (m/z 167.0708) producing a catecholic partial structure.

3.2. Initial CYP activity screening

For identification of the CYPs catalyzing the initial metabolic steps, the ten most abundant human hepatic CYPs were incubated under conditions allowing a statement on the general involvement of a particular CYP enzyme. It should be kept in mind that these qualitative data did not reflect a quantitative contribution of a CYP to the hepatic clearance that would require the collection of enzyme kinetic data [38], which was beyond the scope of this study. As summarized in Tables 1 and 2, CYP2C9 and CYP2C19 were involved in *O*-demethylation for both, 25B-NBOMe and 25C-NBOMe, respectively, CYP1A2 and CYP3A4 in hydroxylation, and CYP3A4 in *N*-demethoxybenzylation.

3.3. Proposed metabolic pathways

According to the 25B-NBOMe metabolites identified in human and rat urine after cleavage of conjugates and 25C-NBOMe metabolites identified in rat urine after cleavage of conjugates (Tables S1 and S2), the following metabolic pathways, depicted in Figs. 1 and 2, could be proposed.

As expected, both compounds underwent the same main metabolic steps. *O*-Demethylation led to the most abundant peaks in human and rat urine followed by *O*-bis-demethylation and /or by *O*-demethylation plus hydroxylation. *N*-Demethoxybenzylation led to only small peaks in both species. However, the relative abundance of the different metabolites varied between the species, but it should also be kept in mind that the rat urines were pooled over 24 h and the human urine was collected at an unknown time after administration of an unknown dose. Finally, the relation of the metabolites may vary over the time of excretion.

For both derivatives, the following phase I pathways could be found: mono-demethylation (B13–B15 and C16–C18), bis-demethylation (B8–B10 and C10–C12), tris-demethylation (B7 and C8) of the methoxy groups, mono- and bis-hydroxylation (B29–B32, B34, B35, and C31–C33, C35, C36), *N*-

demethoxybenzylation (B6 and C5), and combinations of mono-hydroxylation with mono-demethylation (B22–B25 and C26–C29), and bis-demethylation (B16–B19 and C19–C23) as well as bis-hydroxylation with mono-demethylation (B33 and C34), and *N*-demethoxybenzylation with mono-demethylation (B3, B4 and C2, C3) followed by oxidative deamination (B2) and oxidation to the corresponding carboxylic acid (B5 and C4). In addition, for 25C-NBOMe also *N*-demethoxybenzylation with mono-hydroxylation (C7), and oxidation forming an amide structure (C6) could be predicted. Also, dehydro metabolites (B20 and C24) were found for both compounds. The presence of this metabolic step was already described for 25I-NBOMe [25].

Nielsen et al. [39] described dehydrogenation as a CYP-catalyzed reaction. The resulting double bond was located at the 2C moiety and not between the nitrogen and the α -carbon of the 2C moiety as confirmed with reference standard of the 25I-NBOMe imine. These compounds could further be metabolized by mono-demethylation (B11, B12 and C13–C15), bis-demethylation (C9), hydroxylation (B26–B28 and C30), and combination of mono-demethylation and hydroxylation (B21 and C25). However, the possibility could not be excluded that the dehydro compound could also be formed by artificial dehydration of the corresponding hydroxy metabolite. If hydroxylation took place at the α -position to the nitrogen forming an unstable hemiaminal, then this metabolite could further eliminate water under the ESI conditions described above. In summary, the metabolic pathways for 25B-NBOMe and 25C-NBOMe corresponded to those described for 25I-NBOMe, i.e. showing the same main phase I metabolism reactions.

The following phase II pathways could be proposed for humans and/or rats as given in Tables S3 and S4 and Figs. 1 and 2: sulfation (S) glucuronidation (G) and/or of the *O*-demethyl metabolites (B13/14S, B15S, B13G–B15G and C16/17S, C18S, C16G–C18G), of the *O,O*-bis-demethyl metabolites (B8S, B9/10S, B8G, B9/10G and C10S–C12S, C10G–C12G), of *O,O,O*-tris-demethyl metabolite (B7S, B7G and C8S, C8G), of the *O*-demethyl-hydroxy metabolites (B22S, B24/25S, B22G, B23, B24/25G and C22S, C27S, C28/29S), of the *O,O*-bis-demethyl-hydroxy metabolites (B16S, B17/18S, B16G, B19G and C20S, C19G–C22G), and of the hydroxy metabolites (B30G,

B31G and C31/32G, C33G). Glutathione (GSH) conjugation could be proposed for the *O,O*-bis-demethyl metabolite isomer 1 (B8-GSH-1, B8-GSH-2 and C10-GSH-1, C10-GSH-2), *N*-acetylation (AC) for the *N*-demethoxybenzyl-*O*-demethyl metabolites (B3, B4 and C2, C3) followed by further sulfation and/or glucuronidation (B3AC+S, B4AC+S, B3/4AC+G and C3/4AC+S, C2/3AC+G), and *O*-methylation (ME) of the bis-hydroxy metabolite (C36ME) and the *O*-demethyl-bis-hydroxy metabolites (B33ME and C34ME). In summary, all phase II pathways could be proposed for both species except for glutathione conjugation, which was observed only in rats after administration of the high dose.

3.4. Comparison of metabolite formation in vitro and in vivo and in different species

In contrast to the development of new therapeutics drug, pharmacokinetic data are not routinely collected for NPS before emergence on the market. For ethical reasons, controlled human studies are not possible. Therefore, animal studies under controlled conditions are common in combination with human in vivo assays as described e.g. in refs. [25,26]. Both data can be confirmed by authentic human samples of e.g. intoxication cases. For development of urine screening approaches, it is important to know the possible target. Thus, any metabolites identified first in animal urine can become the main target in human urine considering e.g. inter-species and/or genetic variations in drug metabolism and transport. For this reason, Tables 3 and 4 list the phase I and II metabolites identified in this study compared to those detected in human liver microsomes (HLM) incubation, porcine liver microsomes (PLM) incubation, mouse urine (MU), authentic human urines (HU), or human hepatocyte (HP) incubation. Differences could be explained by species differences, higher doses, and/or different sampling time after administration.

3.4.1. 25B-NBOMe

Boumrah et al. [27] described 21 phase I and II metabolites of 25B-NBOMe identified only in vitro after incubation with HLM and cofactors for CYPs and glucuronyl transferases. Leth-Petersen et al. [28] compared formation of phase I metabolites in HLM and PLM incubations. In the present study, 35 phase I and 31 phase II metabolites have been identified in human and rat urine. According to Table 3, in both urine samples, various metabolites could be identified not described by Boumrah et al. or Leth-Petersen et al. Most of them were isomers of metabolites formed by combined metabolic reactions such as mono- and bis-*O*-demethylation with hydroxylation or *O*-demethylation with *N*-dealkylation. Species differences occurred for the hydroxylation step because rats seemed to prefer hydroxylation at the 2C part whereas human biotransformation might result in preferential hydroxylation at the NBOMe part. Concerning phase II metabolism, Boumrah et al. investigated only the glucuronide formation. In the present study, sulfation, *N*-acetylation, and *O*-methylation were found in rat and human urine as further reactions. In addition, rats showed GSH conjugation and combinations of *N*-acetylation with sulfation or glucuronidation. In contrast, the *N*-glucuronide of the parent compound detected in HLM could not be found in the human or rat urine.

3.4.2. 25C-NBOMe

Table 4 summarizes the data obtained in rat urine and those in human hepatocytes and urines of humans and mice [26]. Concerning phase I metabolism, most metabolites were common for all species while the highest number was found in the rat urine probably due to the high dosage, urine collection time, sample preparations, and/or chromatographic separation. Some metabolites were only detected in rat urine such as the combined *N*-dealkylated and *O*-demethylated metabolites or various isomers of *O,O*-bis-demethyl-hydroxy metabolites. Wohlfarth et al. [26] described *N*-oxidation and carbonylation in the hepatocyte incubation although it was not clear why this could not be found in their human and mice urine. As already described for 25B-NBOMe, rats seemed to preferentially hydroxylated at the 2C part and humans at the NBOMe part. Most phase II pathways

could be proposed for all three species with the exception of *O*-acetylation, *N*-acetylation, GSH conjugation, and *O*-methylation. Again, the highest number of metabolites was identified in rat urine probably due to the reasons described above.

3.5. Toxicological detection of 25B-NBOMe and 25C-NBOMe by SUSAs

3.5.1. GC-MS SUSAs

Unfortunately, 25B-NBOMe and 25C-NBOMe and/or their metabolites could not be detected in rat urine after low dose administration (0.1 mg/kg BW). However, 25B-NBOMe and metabolites (Table 5) could be detected in the human urine sample by GC-MS SUSAs. The compound ingested by the user was believed to be 2C-B, which typically requires a ten-fold higher dose compared to 25B-NBOMe [7]. Therefore, for acute and/or severe poisonings with NBOMes an intake could also be detected by GC-MS SUSAs. 25C-NBOMe could only be detected after the high dose, enzymatic cleavage of conjugates, solid-phase extraction, and acetylation according to Welter et al. [31].

3.5.2. LC-MSⁿ SUSAs

The LC-MSⁿ approach could detect 25B-NBOMe and 25C-NBOMe and/or their metabolites in rat urine after low dosage (0.1 mg/kg BW) as well as in the authentic human urine sample. A list of the detected metabolites is given in Table 6. As already mentioned above, the differences of detected analytes in the human and rat urine samples could be caused by different doses and urine collection times.

3.5.3. LC-HR-MS/MS SUSAs

As expected, this approach was also able to reveal 25B-NBOMe and 25C-NBOMe and/or their metabolites in rat urine after low dosage (0.1 mg/kg BW) as well as in the authentic human urine sample. A list of the detected metabolites is given in Table 7. Again, the differences of detected analytes in the human and rat urine samples could be caused by different doses and urine collection times. Mostly due to the lethal overdose, the parent compound gave one of the most abundant signals in the human urine sample. However, low dose rat urine studies showed that the parent compound should not be expected in high amounts after recreational use. Therefore, it should not be used as the only target for NBOMe urine screening.

4. Conclusions

Both, 25B-NBOMe and 25C-NBOMe were extensively metabolized similar to 25I-NBOMe including *O*-demethylation, *O,O*-bis-demethylation, and hydroxylations as predominant pathways in humans and rats. This was in accordance to published human and animal in vitro and in vivo data. Several CYP isoenzymes were involved in formation of the main metabolites. An intake could be detected mainly via their metabolites by low and high resolution LC-MS SUSAs and by GC-MS SUSA only in overdose cases.

Acknowledgements

The authors like to thank Julian A. Michely, Andreas G. Helfer, Sascha K. Manier, Lilian H. J. Richter, Lea Wagmann, Dr. Jessica Welter-Lüdecke, Carsten Schröder, Gabriele Ulrich, and Armin A. Weber for support and/or helpful discussion.

Conflict of interest

The authors declare that there are no conflicts of interest.

References

- [1] European Monitoring Centre for Drugs and Drug Addiction (EMCDDA), New psychoactive substances in Europe. An update from the EU Early Warning System, 2015.
- [2] United Nations Office on Drugs and Crime (UNODC), World Drug Report 2014, http://www.unodc.org/documents/data-and-analysis/WDR2014/World_Drug_Report_2014_web.pdf, 2014.
- [3] United Nations Office on Drugs and Crime (UNODC), World Drug Report 2015, https://www.unodc.org/documents/wdr2015/World_Drug_Report_2015.pdf, 2015.
- [4] United Nations Office on Drugs and Crime (UNODC), World Drug Report 2016, https://www.unodc.org/doc/wdr2016/WORLD_DRUG_REPORT_2016_web.pdf, 2016.
- [5] H.H. Maurer, Chemistry, Pharmacology, and Metabolism of Emerging Drugs of Abuse [review], *Ther. Drug Monit.* 32 (2010) 544-549.
- [6] L.A. King, New phenethylamines in Europe, *Drug Test. Anal* (2013) DOI 10.1002/dta.1570.
- [7] A. Shulgin, Pihkal, A Chemical Love Story, Transform Press, Berkley (CA), 1991.
- [8] J. Suzuki, M.A. Dekker, E.S. Valenti, F.A. Arbelo Cruz, A.M. Correa, J.L. Poklis, A. Poklis, Toxicities associated with NBOMe ingestion-a novel class of potent hallucinogens: a review of the literature, *Psychosomatics* 56 (2015) 129-139.
- [9] M.R. Braden, J.C. Parrish, J.C. Naylor, D.E. Nichols, Molecular interaction of serotonin 5-HT_{2A} receptor residues Phe339(6.51) and Phe340(6.52) with superpotent N-benzyl phenethylamine agonists, *Mol. Pharmacol.* 70 (2006) 1956-1964.
- [10] M. Hansen, K. Phonekeo, J.S. Paine, S. Leth-Petersen, M. Begtrup, H. Brauner-Osborne, J.L. Kristensen, Synthesis and structure-activity relationships of N-benzyl phenethylamines as 5-HT_{2A/2C} agonists, *ACS Chem. Neurosci.* 5 (2014) 243-249.

- 491 [11] A.L. Halberstadt, M.A. Geyer, Effects of the hallucinogen 2,5-dimethoxy-4-
492 iodophenethylamine (2C-I) and superpotent N-benzyl derivatives on the head twitch
493 response, *Neuropharmacology* 77 (2014) 200-207.
- 494 [12] D.E. Nichols, M.F. Sassano, A.L. Halberstadt, L.M. Klein, S.D. Brandt, S.P. Elliott, W.J.
495 Fiedler, N-Benzyl-5-methoxytryptamines as Potent Serotonin 5-HT₂ Receptor Family
496 Agonists and Comparison with a Series of Phenethylamine Analogues, *ACS. Chem.*
497 *Neurosci.* 6 (2015) 1165-1175.
- 498 [13] W. Lawn, M. Barratt, M. Williams, A. Horne, A. Winstock, The NBOMe hallucinogenic drug
499 series: Patterns of use, characteristics of users and self-reported effects in a large international
500 sample, *J. Psychopharmacol.* 28 (2014) 780-788.
- 501 [14] D. Zuba, K. Sekula, A. Buczek, 25C-NBOMe - New potent hallucinogenic substance
502 identified on the drug market, *Forensic Sci. Int.* 227 (2013) 7-14.
- 503 [15] S.L. Hill, T. Doris, S. Gurung, S. Katebe, A. Lomas, M. Dunn, P. Blain, S.H. Thomas, Severe
504 clinical toxicity associated with analytically confirmed recreational use of 25I-NBOMe: case
505 series, *Clin. Toxicol. (Phila)* 51 (2013) 487-492.
- 506 [16] J.L. Poklis, K.G. Devers, E.F. Arbefeville, J.M. Pearson, E. Houston, A. Poklis, Postmortem
507 detection of 25I-NBOMe [2-(4-iodo-2,5-dimethoxyphenyl)-N-[(2-
508 methoxyphenyl)methyl]ethanamine] in fluids and tissues determined by high performance
509 liquid chromatography with tandem mass spectrometry from a traumatic death, *Forensic Sci.*
510 *Int.* 234 (2014) e14-e20.
- 511 [17] S.J. Stellpflug, S.E. Kealey, C.B. Hegarty, G.C. Janis, 2-(4-Iodo-2,5-dimethoxyphenyl)-N-
512 [(2-methoxyphenyl)methyl]ethanamine (25I-NBOMe): clinical case with unique
513 confirmatory testing, *J. Med. Toxicol.* 10 (2014) 45-50.
- 514 [18] M.H. Tang, C.K. Ching, M.S. Tsui, F.K. Chu, T.W. Mak, Two cases of severe intoxication
515 associated with analytically confirmed use of the novel psychoactive substances 25B-
516 NBOMe and 25C-NBOMe, *Clin. Toxicol. (Phila)* 52 (2014) 561-565.

- [19] J. Suzuki, J.L. Poklis, A. Poklis, "My friend said it was good LSD": a suicide attempt following analytically confirmed 25I-NBOMe ingestion, *J. Psychoactive. Drugs.* 46 (2014) 379-382.
- [20] F.S. Bersani, O. Corazza, G. Albano, G. Valeriani, R. Santacroce, P.F. Bolzan Mariotti, E. Cinosi, P. Simonato, G. Martinotti, G. Bersani, F. Schifano, 25C-NBOMe: preliminary data on pharmacology, psychoactive effects, and toxicity of a new potent and dangerous hallucinogenic drug, *Biomed. Res. Int.* 2014 (2014) 734749.
- [21] J.L. Poklis, C.R. Nanco, M.M. Troendle, C.E. Wolf, A. Poklis, Determination of 4-bromo-2,5-dimethoxy-N-[(2-methoxyphenyl)methyl]-benzeneethanamine (25B-NBOMe) in serum and urine by high performance liquid chromatography with tandem mass spectrometry in a case of severe intoxication, *Drug Test. Anal.* 6 (2014) 764-769.
- [22] J.P. Walterscheid, G.T. Phillips, A.E. Lopez, M.L. Gonsoulin, H.H. Chen, L.A. Sanchez, Pathological findings in 2 cases of fatal 25I-NBOMe toxicity, *Am. J. Forensic Med. Pathol.* 35 (2014) 20-25.
- [23] A. Ettrup, S. da Cunha-Bang, B. McMahon, S. Lehel, A. Dyssegaard, A.W. Skibsted, L.M. Jorgensen, M. Hansen, A.O. Baandrup, S. Bache, C. Svarer, J.L. Kristensen, N. Gillings, J. Madsen, G.M. Knudsen, Serotonin 2A receptor agonist binding in the human brain with [(1)(1)C]Cimbi-36, *J. Cereb. Blood. Flow. Metab.* 34 (2014) 1188-1196.
- [24] A. Ettrup, C. Svarer, B. McMahon, S. da Cunha-Bang, S. Lehel, K. Moller, A. Dyssegaard, M. Ganz, V. Beliveau, L.M. Jorgensen, N. Gillings, G.M. Knudsen, Serotonin 2A receptor agonist binding in the human brain with [(11)C]Cimbi-36: Test-retest reproducibility and head-to-head comparison with the antagonist [(18)F]altanserin, *Neuroimage* 130 (2016) 167-174.
- [25] A.T. Caspar, A.G. Helfer, J.A. Michely, V. Auwaerter, S.D. Brandt, M.R. Meyer, H.H. Maurer, Studies on the metabolism and toxicological detection of the new psychoactive designer drug 2-(4-iodo-2,5-dimethoxyphenyl)-N-[(2-methoxyphenyl)methyl]ethanamine

(25I-NBOMe) in human and rat urine using GC-MS, LC-MSⁿ, and LC-HR-MS/MS, *Anal. Bioanal. Chem.* 407 (2015) 6697-6719.

[26] A. Wohlfarth, M. Roman, M. Andersson, F.C. Kugelberg, X. Diao, J. Carlier, C. Eriksson, X. Wu, P. Konradsson, M. Josefsson, M.A. Huestis, R. Kronstrand, 25C-NBOMe and 25I-NBOMe metabolite studies in human hepatocytes, in vivo mouse and human urine with high-resolution mass spectrometry, *Drug Test. Anal.* (2016) DOI: 10.1002/dta.2044.

[27] Y. Boumrah, L. Humbert, M. Phanithavong, K. Khimeche, A. Dahmani, D. Allorge, In vitro characterization of potential CYP- and UGT-derived metabolites of the psychoactive drug 25B-NBOMe using LC-high resolution MS, *Drug Test. Anal.* 8 (2016) 248-256.

[28] S. Leth-Petersen, C. Gabel-Jensen, N. Gillings, S. Lehel, H.D. Hansen, G.M. Knudsen, J.L. Kristensen, Metabolic fate of hallucinogenic NBOMes, *Chem. Res. Toxicol.* (2015) DOI: 10.1021/acs.chemrestox.5b00450.

[29] J. Welter, P. Kavanagh, M.R. Meyer, H.H. Maurer, Benzofuran analogues of amphetamine and methamphetamine: studies on the metabolism and toxicological analysis of 5-APB and 5-MAPB in urine and plasma using GC-MS and LC-(HR)-MSⁿ techniques, *Anal. Bioanal. Chem.* 407 (2015) 1371-1388.

[30] V. Sharma, J.H. McNeill, To scale or not to scale: the principles of dose extrapolation, *Br. J. Pharmacol.* 157 (2009) 907-921.

[31] J. Welter, M.R. Meyer, E. Wolf, W. Weinmann, P. Kavanagh, H.H. Maurer, 2-Methiopropamine, a thiophene analogue of methamphetamine: studies on its metabolism and detectability in the rat and human using GC-MS and LC-(HR)-MS techniques, *Anal. Bioanal. Chem.* 405 (2013) 3125-3135.

[32] L.H.R. Richter, Y.R. Kaminski, F. Noor, M.R. Meyer, H.H. Maurer, Metabolic fate of desomorphine elucidated using rat urine, pooled human liver preparations, and human hepatocyte cultures as well as its detectability using standard urine screening approaches, *Anal. Bioanal. Chem.* 408 (2016) 6283-6294.

- [33] M.R. Meyer, C. Lindauer, J. Welter, H.H. Maurer, Dimethocaine, a synthetic cocaine derivative: Studies on its in vivo metabolism and its detectability in urine by LC-HR-MSⁿ and GC-MS using a rat model, *Anal. Bioanal. Chem.* 406 (2014) 1845-1854.
- [34] H.H. Maurer, K. Pfleger, A.A. Weber, Mass spectral data of drugs, poisons, pesticides, pollutants and their metabolites, Wiley-VCH, Weinheim (Germany), 2016.
- [35] D.K. Wissenbach, M.R. Meyer, D. Remane, A.A. Philipp, A.A. Weber, H.H. Maurer, Drugs of abuse screening in urine as part of a metabolite-based LC-MS(n) screening concept, *Anal. Bioanal. Chem.* 400 (2011) 3481-3489.
- [36] A.G. Helfer, J.A. Michely, A.A. Weber, M.R. Meyer, H.H. Maurer, Orbitrap technology for comprehensive metabolite-based liquid chromatographic-high resolution-tandem mass spectrometric urine drug screening - exemplified for cardiovascular drugs, *Anal. Chim. Acta* 891 (2015) 221-233.
- [37] D.S. Theobald, G. Fritschi, H.H. Maurer, Studies on the toxicological detection of the designer drug 4-bromo-2,5-dimethoxy-beta-phenethylamine (2C-B) in rat urine using gas chromatography-mass spectrometry, *J. Chromatogr. B* 846 (2007) 374-377.
- [38] C.S.D. Wink, G.M.J. Meyer, M.R. Meyer, H.H. Maurer, Toxicokinetics of lefetamine and derived diphenylethylamine designer drugs - Contribution of human cytochrome P450 isozymes to their main phase I metabolic steps, *Toxicol. Lett.* 238 (2015) 39-44.
- [39] L.M. Nielsen, N.B. Holm, S. Leth-Petersen, J.L. Kristensen, L. Olsen, K. Linnet, Characterization of the hepatic cytochrome P450 enzymes involved in the metabolism of 25I-NBOMe and 25I-NBOH, *Drug Test. Anal.* (2016) DOI: 10.1002/dta.2031.

Table 1 General involvement of the CYP isoenzymes on the formation of the given 25B-NBOMe metabolites, numbering according to Table S1

Metabolite	CYP 1A2	CYP 2A6	CYP 2B6	CYP 2C8	CYP 2C9	CYP 2C19	CYP 2D6	CYP 2E1	CYP 3A4	CYP 3A5
<i>N</i> -Demethoxybenzyl (B6)	+		+						+	+
<i>O</i> -Demethyl isomer 1 (B13)	+				+	+			+	
<i>O</i> -Demethyl isomer 2 (B14)	+				+	+	+		+	
<i>O</i> -Demethyl isomer 3 (B15)	+			+	+	+	+		+	
Dehydro- (B20)						+	+		+	
Hydroxy isomer 2 (B30)	+					+			+	+
Hydroxy isomer 3 (B31)	+					+			+	+
Hydroxy isomer 4 (B32)			+						+	

Table 2 General involvement of the CYP isoenzymes on the formation of the given 25C-NBOMe metabolites, numbering according to Table S2

Metabolite	CYP 1A2	CYP 2A6	CYP 2B6	CYP 2C8	CYP 2C9	CYP 2C19	CYP 2D6	CYP 2E1	CYP 3A4	CYP 3A5
<i>N</i> -Demethoxybenzyl (C5)	+		+						+	+
<i>O</i> -Demethyl isomer 1 (C16)	+				+	+			+	
<i>O</i> -Demethyl isomer 2 (C17)	+				+	+	+		+	
<i>O</i> -Demethyl isomer 3 (C18)	+			+	+	+	+		+	
Dehydro- (C24)						+	+		+	
Hydroxy isomer 3 (C33)	+					+			+	+

Table 3 25B-NBOMe phase I and II metabolites detected in rat (RU) and human (HU) urine compared to those detected in human liver microsome (HLM) incubation published by Boumrah et al. [27] and in HLM and porcine liver microsome (PLM) incubations published by Leth-Petersen et al. [28]. Numbering according to Tables S1 and S3, * = metabolite only described in references [27,28]

No.	Metabolite	RU	HU	HLM [27]	HLM [28]	PLM [28]
B1	25B-NBOMe	+	+	+	+	+
B2	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-deamino-hydroxy-)	+	+			
B3	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) isomer 1	+	+			
B4	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) isomer 2	+	+			
B5	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-deamino-carboxy-)	+	+			
B6	25B-NBOMe-M (<i>N</i> -demethoxybenzyl-)	+	+	+	+	+
B7	25B-NBOMe-M (<i>O,O,O</i> -tris-demethyl-)	+	+			
B8	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 1	+	+	+	+	+
B9	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 2	+	+	+	+	
B10	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 3	+	+	+	+	+
B11	25B-NBOMe-M (<i>O</i> -demethyl-dehydro-) isomer 1	+	+			
B12	25B-NBOMe-M (<i>O</i> -demethyl-dehydro-) isomer 2	+				
B13	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 1	+	+	+	+	+
B14	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 2	+	+	+	+	+
B15	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 3	+	+	+	+	+
B16	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 1	+				
B17	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 2	+	+			
B18	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 3	+	+			
B19	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 4	+				
B20	25B-NBOMe-M (dehydro-)	+	+			+
B21	25B-NBOMe-M (<i>O</i> -demethyl-dehydro-hydroxy-)	+				
B22	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 1	+				+
B23	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 2	+	+		+	+
B24	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 3		+	+	+	+
B25	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 4	+		+	+	+
B26	25B-NBOMe-M (dehydro-hydroxy-) isomer 1		+			
B27	25B-NBOMe-M (dehydro-hydroxy-) isomer 2	+				
B28	25B-NBOMe-M (dehydro-hydroxy-) isomer 3	+	+			
B29	25B-NBOMe-M (hydroxy-) isomer 1	+				
B30	25B-NBOMe-M (hydroxy-) isomer 2		+	+	+	+
B31	25B-NBOMe-M (hydroxy-) isomer 3	+	+	+		
B32	25B-NBOMe-M (hydroxy-) isomer 4		+	+	+	
B33	25B-NBOMe-M (<i>O</i> -demethyl-bis-hydroxy-)	+				
B34	25B-NBOMe-M (<i>bis</i> -hydroxy-) isomer 1		+			
B35	25B-NBOMe-M (<i>bis</i> -hydroxy-) isomer 2	+	+			
M11	25B-NBOMe-M (carbonyl) *					+
B3 AC	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl isomer 1	+	+			
B4 AC	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl isomer 2	+	+			

B3 AC+S	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl sulfate isomer 1	+				
B4 AC+S	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl sulfate isomer 2	+				
B8 GSH-1	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) <i>S</i> -methyl	+				
B33 ME	25B-NBOMe-M (<i>O</i> -demethyl-bis-hydroxy-) <i>O</i> -methyl	+	+			
B7 S	25B-NBOMe-M (<i>O,O,O</i> -tris-demethyl-) sulfate	+				
B3/4 G	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) glucuronide	+				
B8 S	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) sulfate isomer 1	+	+			
B9/10 S	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) sulfate isomer 2		+			
B13/14 S	25B-NBOMe-M (<i>O</i> -demethyl-) sulfate isomer 1	+	+			
B15 S	25B-NBOMe-M (<i>O</i> -demethyl-) sulfate isomer 2	+				
B16 S	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) sulfate isomer 1	+				
B17/18 S	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) sulfate isomer 2		+			
B22 S	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) sulfate isomer 1	+				
B24/25 S	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) sulfate isomer 2	+	+			
B3/4 AC+G	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl glucuronide	+				
B8 GSH-2	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) acetylcysteine	+				
B7 G	25B-NBOMe-M (<i>O,O,O</i> -tris-demethyl-) glucuronide	+				
B8 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 1	+	+	+		
B9/10 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 2	+	+	+		
B13 G	25B-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 1	+	+	+		
B14 G	25B-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 2	+	+	+		
B15 G	25B-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 3	+	+	+		
B16 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) glucuronide isomer 1	+				
B19 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) glucuronide isomer 2	+				
B23 G	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 1	+	+			
B22 G	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 2	+				
B24/25 G	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 3		+	+		
B30 G	25B-NBOMe-M (hydroxy-) glucuronide isomer 1		+	+		
B31 G	25B-NBOMe-M (hydroxy-) glucuronide isomer 2		+	+		
M21	25B-NBOMe-M <i>N</i> -glucuronide *			+		

Table 4 25C-NBOMe phase I and II metabolites detected in rat (RU) urine compared to those in authentic human urines (HU), mouse urine (MU) and human hepatocyte (HP) incubation as published by Wohlfarth et al. [26]. Numbering according to Tables S2 and S4, * = metabolite only described in reference [26]

No.	Metabolite	RU	HU [26]	MU [26]	HP [26]
C1	25C-NBOMe	+	+	+	+
C2	25C-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) isomer 1	+			
C3	25C-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) isomer 2	+			
C4	25C-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-deamino-carboxy-)	+			
C5	25C-NBOMe-M (<i>N</i> -demethoxybenzyl-)	+	+		+
C6	25C-NBOMe-M (<i>N</i> -demethoxybenzyl-oxo-)	+			
C7	25C-NBOMe-M (<i>N</i> -demethoxybenzyl-hydroxy-)	+			
C8	25C-NBOMe-M (<i>O,O,O</i> -tris-demethyl-)	+			
C9	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-dehydro-)	+			
C10	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 1	+		+	
C11	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 2	+	+	+	
C12	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 3	+	+	+	
C13	25C-NBOMe-M (<i>O</i> -demethyl-dehydro-) isomer 1	+			
C14	25C-NBOMe-M (<i>O</i> -demethyl-dehydro-) isomer 2	+			
C15	25C-NBOMe-M (<i>O</i> -demethyl-dehydro-) isomer 3	+			
C16	25C-NBOMe-M (<i>O</i> -demethyl-) isomer 1	+	+	+	+
C17	25C-NBOMe-M (<i>O</i> -demethyl-) isomer 2	+	+	+	+
C18	25C-NBOMe-M (<i>O</i> -demethyl-) isomer 3	+	+		+
C19	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 1	+			
C20	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 2	+		+	
C21	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 3	+			
C22	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 4	+			
C23	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) isomer 5	+			
C24	25C-NBOMe-M (dehydro-)	+			
C25	25C-NBOMe-M (<i>O</i> -demethyl-dehydro-hydroxy-)	+			
C26	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 1	+			
C27	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 2	+	+	+	
C28	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 3	+	+	+	+
C29	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) isomer 4	+			
C30	25C-NBOMe-M (dehydro-hydroxy-)	+			
C31	25C-NBOMe-M (hydroxy-) isomer 1	+			
C32	25C-NBOMe-M (hydroxy-) isomer 2	+			
C33	25C-NBOMe-M (hydroxy-) isomer 3	+	+	+	+
C34	25C-NBOMe-M (<i>O</i> -demethyl-bis-hydroxy-)	+	+		
C35	25C-NBOMe-M (<i>bis</i> -hydroxy-) isomer 1	+			
C36	25C-NBOMe-M (<i>bis</i> -hydroxy-) isomer 2	+			
C-Hp-21	25C-NBOMe-M (<i>N</i> -oxide) *				+
C-Hp-22	25C-NBOMe-M (carbonyl) *				+
C2	25C-NBOMe-M				
AC	(<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl isomer 1	+			
C3	25C-NBOMe-M				
AC	(<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl isomer 2	+			
C7	25C-NBOMe-M				
AC	(<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-hydroxy-) <i>N</i> -acetyl	+			
C3/4	25C-NBOMe-M				
AC+S	(<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl sulfate	+			
C10	25C-NBOMe-M				
GSH-1	(<i>O,O</i> -bis-demethyl-) S-methyl	+			
C34	25C-NBOMe-M	+			

ME	(<i>O</i> -demethyl- <i>bis</i> -hydroxy-) <i>O</i> -methyl				
C8 S	25C-NBOMe-M (<i>O,O,O</i> - <i>tris</i> -demethyl-) sulfate	+			
C2/3 G	25C-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) glucuronide	+			
C36 ME	25C-NBOMe-M (<i>bis</i> -hydroxy-) <i>O</i> -methyl	+			
C10 S	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) sulfate isomer 1	+			
C11 S	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) sulfate isomer 2	+			
C12 S	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) sulfate isomer 3	+	+		+
C16/17 S	25C-NBOMe-M (<i>O</i> -demethyl-) sulfate isomer 1	+	+	+	+
C18 S	25C-NBOMe-M (<i>O</i> -demethyl-) sulfate isomer 2	+			+
C20 S	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-hydroxy-) sulfate	+			
C22 S	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) sulfate	+			
C2/3 AC+G	25C-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl glucuronide	+			
C10 GSH-2	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) acetylcysteine	+			
C8 G	25C-NBOMe-M (<i>O,O,O</i> - <i>tris</i> -demethyl-) glucuronide	+		+	
C10 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) glucuronide isomer 1	+	+	+	
C11 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) glucuronide isomer 2	+	+	+	
C12 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-) glucuronide isomer 3	+		+	
C16 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 1	+	+	+	+
C17 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 2	+			+
C18 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 3	+			+
C19 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-hydroxy-) glucuronide isomer 1	+			
C20 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-hydroxy-) glucuronide isomer 2	+			
C21 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-hydroxy-) glucuronide isomer 3	+			
C22 G	25C-NBOMe-M (<i>O,O</i> - <i>bis</i> -demethyl-hydroxy-) glucuronide isomer 4	+			
C27 G	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 1	+			
C28/29 G	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 2	+			+
C31/32 G	25C-NBOMe-M (hydroxy-) glucuronide isomer 1	+			
C33 G	25C-NBOMe-M (hydroxy-) glucuronide isomer 2	+	+	+	+
C-Hp-6	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer *				+
C-Hp-8	25C-NBOMe-M (hydroxy-) glucuronide isomer *				+
C-Hp-10	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer *				+

C-Hp-18	25C-NBOMe-M (hydroxy-) sulfate *				+
C-Hp-19	25C-NBOMe-M (hydroxy-) sulfate *				+
C-MH-21	25C-NBOMe-M (<i>O</i> -demethyl-) <i>O</i> -acetyl *			+	

Table 5 25B-NBOMe and its metabolites, molecular mass, five most abundant EI-GC-MS fragment ions, retention indices (RI), and detectability in rat urine (RU) or human urine (HU) by GC-MS SUSA. The numbers correspond to those of Table S1.

No.	Target for SUSA	Molecular mass, u	GC-MS fragment ions, m/z and their relative intensities, %	RI	Detected in urine sample
B1	25B-NBOMe AC	421	121 (100), 150 (9), 229 (12), 242 (33), 421 (2)	2920	HU
B2	25B-NBOMe-M (<i>N</i> -demethoxybenzyl-deamino- <i>O</i> -demethyl-hydroxy-) 2AC	330	215 (55), 228 (100), 246 (10), 288 (15), 330 (4)	2160	HU
B3/B4	25B-NBOMe-M (<i>N</i> -demethoxybenzyl - <i>O</i> -demethyl-) isomer 1 / isomer 2 2AC	329	215 (17), 228 (100), 270 (10), 287 (21), 329 (8)	2440	HU
B6	25B-NBOMe-M (<i>N</i> -demethoxybenzyl -) AC	301	148 (39), 199 (12), 229 (31), 242 (100), 301 (15)	2180	HU
B9/10	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 2 / isomer 3 3AC	477	107 (78), 178 (100), 228 (42), 270 (12), 477 (1)	3020	HU
B13/14	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 1 / isomer 2 2AC	449	121 (100), 192 (22), 228 (19), 270 (3), 449 (2)	3000	HU

Table 6 25B-NBOMe, 25C-NBOMe, and their metabolites, protonated precursor ions, characteristic MS² and MS³ fragment ions, retention time (RT), and detectability in rat urine (RU) or human urine (HU, 25B-NBOMe) by LC-MSⁿ SUSA. The numbers correspond to those of Tables S1-S4.

No.	Target for SUSA	Precursor ions, <i>m/z</i>	MS ² fragment ions, <i>m/z</i> and relative intensity, %	MS ³ fragment ions, <i>m/z</i> and relative intensity, % on the ion given in bold	RT, min	Detected in urine sample
B1	25B-NBOMe	380	121 (100), 179 (10), 243 (10), 255 (18), 258 (14), 269 (10), 284 (15)	121 : 91 (30), 93 (100) 255 : 148 (10), 176 (100), 225 (44)	14.6	HU
B9	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 2	352	107 (1), 229 (56), 246 (100)	229 : 135 (5), 150 (100) 246 : 135 (3), 150 (51), 214 (100)	9.7	HU
B13	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 1	366	121 (100), 229 (3), 241 (7), 244 (12), 270 (26)	121 : 91 (24), 93 (100) 270 : 145 (6), 224 (7), 239 (100)	11.5	HU
B14	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 2	366	121 (88), 241 (100), 257 (92), 258 (37), 270 (51)	241 : 147 (5), 162 (100) 257 : 149 (46), 162 (55), 225 (100)	12.3	HU, RU
B8 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 1	528	227 (8), 244 (4), 335 (7), 352 (100)	352 : 121 (100), 227 (55), 244 (18), 256 (21), 273 (7)	5.9	RU
B14 G	25B-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 2	542	244 (2), 270 (2), 349 (3), 366 (100)	349 : 241 (41), 255 (22), 270 (100) 366 : 121 (100), 241 (4), 244 (6), 270 (17)	9.4	HU, RU
C16	25C-NBOMe-M (<i>O</i> -demethyl-) isomer 1	322	91 (9), 121 (100), 197 (9), 200 (11), 214 (5)	121 : 91 (22), 93 (100)	13.1	RU
C10 G	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 1	484	183 (15), 200 (4), 291 (11), 308 (100)	291 : 121 (47), 183 (100), 255 (23) 308 : 121 (100), 183 (51), 200 (21)	6.4	RU
C11 G	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 2	484	185 (38), 202 (47), 308 (100), 378 (20)	202 : 150 (21), 157 (10), 170 (100) 308 : 185 (84), 202 (100)	8.2	RU
C17 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 2	498	185 (1), 200 (2), 305 (2), 322 (100)	322 : 121 (100), 197 (11), 200 (10), 214 (5)	9.9	RU
C28/29 G	25C-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 2	514	216 (2), 321 (5), 338 (100)	338 : 121 (100), 198 (5), 216 (3), 230 (1), 303 (7)	8.6	RU

Table 7 25B-NBOMe, 25C-NBOMe, and their metabolites, calculated masses of their precursor ions, retention times (RT) recorded in rat urine or human urine (25B-NBOMe, 25C-NBOMe not tested, n.t.) by LC-HR-MS/MS SUSA. The numbers correspond to those of Tables S1-S4 (D = detection of the precursor ion in MS¹, I = identification via MS¹ and MS²).

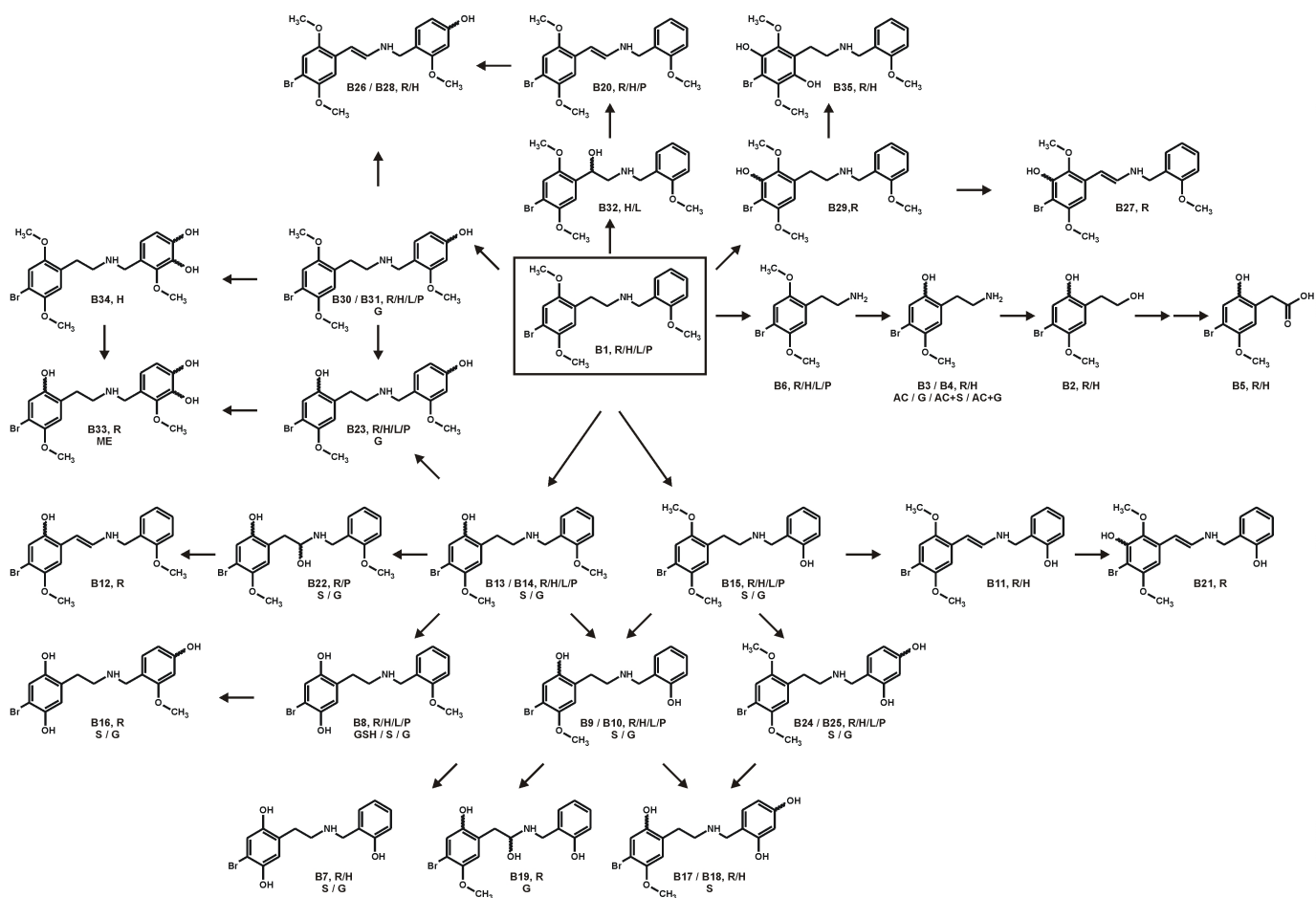
No.	Targets in SUSA	Calculated exact masses of precursor ions, <i>m/z</i>	RT, min	Human urine	Rat urine 0.1 mg/kg BW
B1	25B-NBOMe	380.0856	6.0	I	
B5	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-deamino-carboxy-)	258.9606	5.3	D	D
B8	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 1	352.0543	5.0	I	D
B9	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 2	352.0543	5.3	I	D
B13	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 1	366.0699	5.3	I	
B14	25B-NBOMe-M (<i>O</i> -demethyl-) isomer 2	366.0699	5.8	I	I
B30	25B-NBOMe-M (hydroxy-) isomer 2	396.0805	5.4	D	
B31	25B-NBOMe-M (hydroxy-) isomer 3	396.0805	5.9	D	
B3 AC	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl isomer 1	288.0230	5.4		D
B4 AC	25B-NBOMe-M (<i>N</i> -demethoxybenzyl- <i>O</i> -demethyl-) <i>N</i> -acetyl isomer 2	288.0230	5.5		D
B8 S	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) sulfate isomer 1	432.0111	4.9	D	D
B9/10 S	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) sulfate isomer 2	432.0111	5.7	I	
B13/14 S	25B-NBOMe-M (<i>O</i> -demethyl-) sulfate isomer 1	446.0267	5.8	I	
B24/25 S	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) sulfate isomer 2	462.0217	5.5	D	
B7 G	25B-NBOMe-M (<i>O,O,O</i> -tris-demethyl-) glucuronide	514.0707	3.8		D
B8 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 1	528.0864	4.2	I	I
B9/10 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 2	528.0864	4.8	D	D
B13 G	25B-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 1	542.1020	4.6	I	
B14 G	25B-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 2	542.1020	5.2	I	I
B19 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) glucuronide isomer 2	544.0813	4.5		D
B23 G	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 1	558.0969	4.7		D

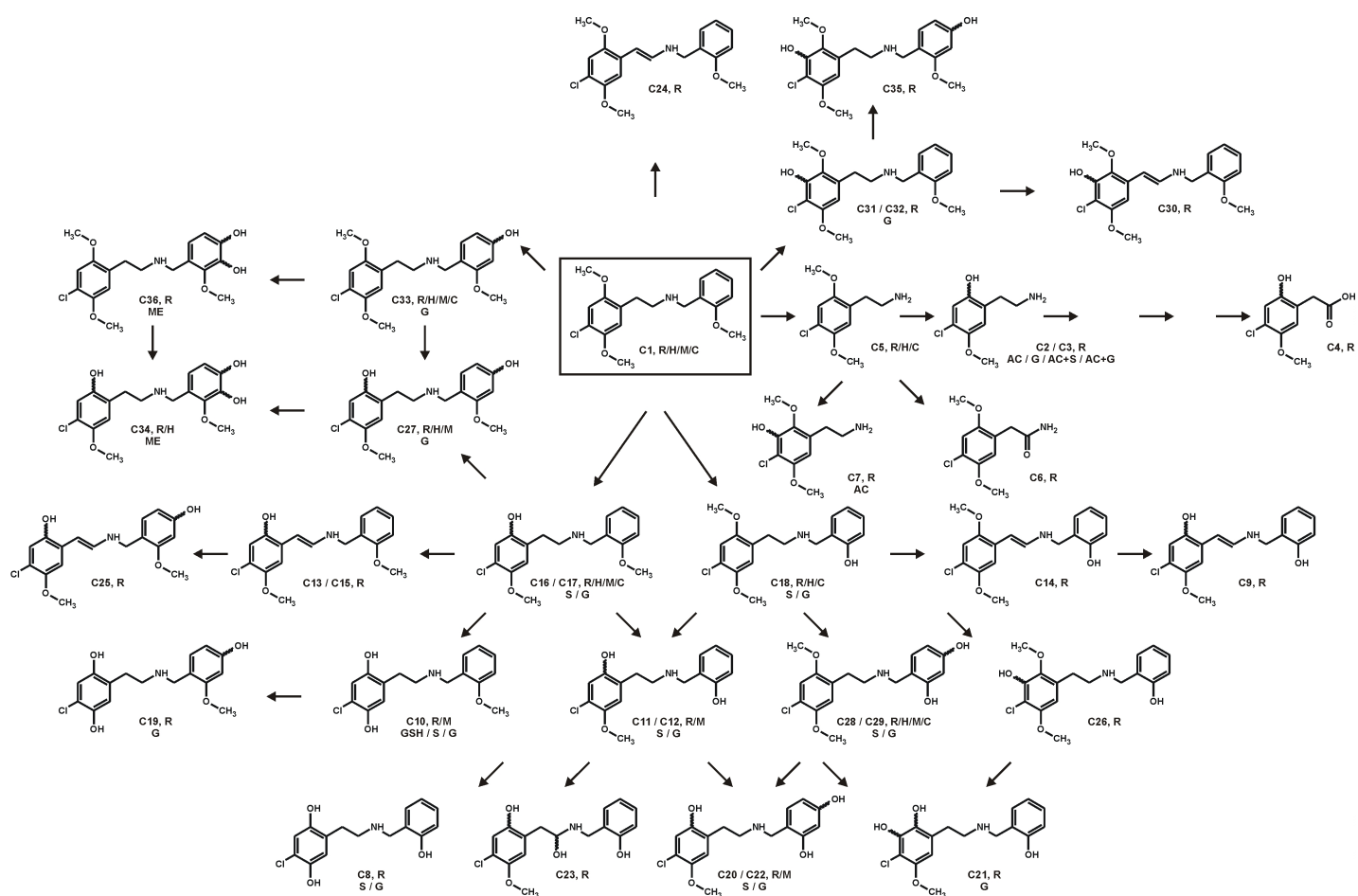
B22 G	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 2	558.0969	4.8		D
C11	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 2	308.1048	4.6	n.t.	D
C12	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) isomer 3	308.1048	4.8	n.t.	D
C17	25C-NBOMe-M (<i>O</i> -demethyl-) isomer 2	322.1204	5.1	n.t.	D
C8 G	25C-NBOMe-M (<i>O,O,O</i> -tris-demethyl-) glucuronide	470.1212	3.5	n.t.	I
C10 G	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 1	484.1369	3.8	n.t.	I
C11 G	25C-NBOMe-M (<i>O,O</i> -bis-demethyl-) glucuronide isomer 2	484.1369	4.4	n.t.	I
C16 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 1	498.1525	4.3	n.t.	D
C17 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 2	498.1525	4.8	n.t.	I
C18 G	25C-NBOMe-M (<i>O</i> -demethyl-) glucuronide isomer 3	498.1525	5.2	n.t.	D
C21 G	25B-NBOMe-M (<i>O,O</i> -bis-demethyl-hydroxy-) glucuronide isomer 3	500.1318	4.1	n.t.	D
C28/29 G	25B-NBOMe-M (<i>O</i> -demethyl-hydroxy-) glucuronide isomer 2	514.1475	4.5	n.t.	I
C31/32 G	25B-NBOMe-M (hydroxy-) glucuronide isomer 1	528.1631	4.1	n.t.	D

Legends to Figures

Fig. 1 Metabolic pathways of 25B-NBOMe studied in rat (R) or human (H) urine as well as in incubations with human (L) or porcine (P) liver microsomes. Phase II metabolites: glucuronides (G), sulfates (S), glutathione conjugates (GSH), acetyl conjugates (AC), *O*-methyl conjugates (ME), acetyl conjugates combined with glucuronidation (AC+G), acetyl conjugates combined with sulfation (AC+S). Undefined position of *O*-demethylation or hydroxylation indicated by tildes. Numbering according to Tables S1 and S3.

Fig. 2 Metabolic pathways of 25C-NBOMe studied in rat (R), mouse (M), or human (H) urine as well as in incubations with human hepatocytes (C). Phase II metabolites: glucuronides (G), sulfates (S), glutathione conjugates (GSH), acetyl conjugates (AC), *O*-methyl conjugates (ME), acetyl conjugates combined with glucuronidation (AC+G), acetyl conjugates combined with sulfation (AC+S). Undefined position of *O*-demethylation or hydroxylation indicated by tildes. Numbering according to Tables S2 and S4.





Highlights

- First detailed Orbitrap-based study on the metabolism of two New Psychoactive Substances (NPS) and on detectability in urine by GC-MS and low and high resolution LC-MS techniques.
- The analytical novelty consists of the description of the identification power of various GC-MS and LC-(HR) MS techniques.
- The corresponding reference spectra and their interpretation are basis for routine drug testing worldwide of these NPS and thus of great relevance for all toxicologists.
- First comparison of metabolism data obtained from in vivo studies with three different species and from human in cellulo and in vitro studies.

Electronic Supplementary Data

Metabolic fate and detectability of the new psychoactive substances 2-(4-bromo-2,5-dimethoxyphenyl)-*N*-[(2-methoxyphenyl)methyl]ethanamine (25B-NBOMe) and 2-(4-chloro-2,5-dimethoxyphenyl)-*N*-[(2-methoxyphenyl)methyl]ethanamine (25C-NBOMe) in human and rat urine by GC-MS, LC-MSⁿ, and LC-HR-MS/MS approaches

Achim T. Caspar, Simon D. Brandt, Andreas E. Stoever, Markus R. Meyer, Hans H. Maurer

Table S1 List of 25B-NBOMe and its phase I metabolites together with the precursor mass (PM) recorded in MS¹, the corresponding characteristic fragment ions (FI) in MS², the calculated exact masses, the corresponding elemental composition, the deviation of the measured from the calculated masses, given as errors in parts per million (ppm), and the retention times (RT) in minutes (min). The metabolites were sorted by mass and RT.

No.	Metabolite and characteristic ions Measured accurate mass, <i>m/z</i>	Relative intensity in MS ² , %	Calculated exact mass, <i>m/z</i>	Elemental composition	Error, ppm	RT, min
B1	25B-NBOMe					8.8
	MS ¹ PM at <i>m/z</i> 380.0859 (M+H)	7	380.0856	C ₁₈ H ₂₅ O ₃ NBr	0.84	
	MS ² FI at <i>m/z</i> 91.0549	58	91.0548	C ₇ H ₇	1.37	
	FI at <i>m/z</i> 121.0651	100	121.0653	C ₈ H ₉ O	-1.98	
	FI at <i>m/z</i> 227.9777	1	227.9786	C ₉ H ₉ O ₂ Br	-3.91	
	FI at <i>m/z</i> 243.0010	1	243.0021	C ₁₀ H ₁₂ O ₂ Br	-4.39	
	FI at <i>m/z</i> 258.0126	0.2	258.0124	C ₁₀ H ₁₃ O ₂ NBr	0.71	
	FI at <i>m/z</i> 363.0597	0.3	363.0596	C ₁₈ H ₂₀ O ₃ Br	0.33	
B2	25B-NBOMe-M (N-demethoxybenzyl-O-demethyl-deamino-hydroxy-)					6.5
	MS ¹ PM at <i>m/z</i> 244.9820 (M-H)	5	244.9820	C ₉ H ₁₀ O ₃ Br	2.73	
	MS ² FI at <i>m/z</i> 78.9176	100	78.9183	Br	-9.33	
	FI at <i>m/z</i> 199.9474	9	199.9473	C ₇ H ₅ O ₂ Br	0.55	
	FI at <i>m/z</i> 211.9476	12	211.9473	C ₈ H ₅ O ₂ Br	1.46	
	FI at <i>m/z</i> 229.9584	76	229.9584	C ₈ H ₇ O ₃ Br	2.37	
B3	25B-NBOMe-M (N-demethoxybenzyl-O-demethyl-) isomer 1					3.9
	MS ¹ PM at <i>m/z</i> 246.0123 (M+H)	1	246.0124	C ₉ H ₁₃ O ₂ NBr	-0.47	
	MS ² FI at <i>m/z</i> 135.0442	24	135.0446	C ₈ H ₇ O ₂	-3.00	
	FI at <i>m/z</i> 150.0677	47	150.0681	C ₉ H ₁₀ O ₂	-2.53	
	FI at <i>m/z</i> 213.9626	82	213.9629	C ₈ H ₇ O ₂ Br	-1.59	
	FI at <i>m/z</i> 228.9861	100	228.9864	C ₉ H ₁₀ O ₂ Br	-1.38	
B4	25B-NBOMe-M (N-demethoxybenzyl-O-demethyl-) isomer 2					4.0
	MS ¹ PM at <i>m/z</i> 246.0130 (M+H)	3	246.0124	C ₉ H ₁₃ O ₂ NBr	2.37	
	MS ² FI at <i>m/z</i> 135.0442	22	135.0446	C ₈ H ₇ O ₂	-3.00	
	FI at <i>m/z</i> 150.0676	40	150.0681	C ₉ H ₁₀ O ₂	-3.20	
	FI at <i>m/z</i> 213.9625	90	213.9629	C ₈ H ₇ O ₂ Br	-2.06	
	FI at <i>m/z</i> 228.9861	100	228.9864	C ₉ H ₁₀ O ₂ Br	-1.38	
B5	25B-NBOMe-M (N-demethoxybenzyl-O-demethyl-deamino-carboxy-)					6.3
	MS ¹ PM at <i>m/z</i> 258.9610 (M-H)	1	258.9606	C ₉ H ₈ O ₄ Br	1.56	
	MS ² FI at <i>m/z</i> 78.9176	100	78.9183	Br	-9.33	
	FI at <i>m/z</i> 199.9473	81	199.9473	C ₇ H ₅ O ₂ Br	0	
	FI at <i>m/z</i> 214.9709	5	214.9708	C ₈ H ₈ O ₂ Br	0.62	
B6	25B-NBOMe-M (N-demethoxybenzyl-)					5.6
	MS ¹ PM at <i>m/z</i> 260.0273 (M+H)	1	260.0281	C ₁₀ H ₁₅ O ₂ NBr	-2.95	
	MS ² FI at <i>m/z</i> 164.0830	22	164.0837	C ₁₀ H ₁₂ O ₂	-4.45	
	FI at <i>m/z</i> 212.9543	39	212.9551	C ₈ H ₆ O ₂ Br	-3.83	
	FI at <i>m/z</i> 227.9776	100	227.9786	C ₉ H ₉ O ₂ Br	-4.35	
	FI at <i>m/z</i> 243.0013	90	243.0021	C ₁₀ H ₁₂ O ₂ Br	-3.15	
B7	25B-NBOMe-M (O,O,O-tris-demethyl-)					5.1
	MS ¹ PM at <i>m/z</i> 338.0392 (M+H)	9	338.0386	C ₁₅ H ₁₇ O ₃ NBr	1.68	
	MS ² FI at <i>m/z</i> 107.0496	100	107.0497	C ₇ H ₇ O	-0.84	
	FI at <i>m/z</i> 136.0520	33	136.0524	C ₈ H ₈ O ₂	-3.16	
	FI at <i>m/z</i> 214.9703	81	214.9708	C ₈ H ₈ O ₂ Br	-2.17	
	FI at <i>m/z</i> 231.9968	32	231.9968	C ₈ H ₁₁ O ₂ NBr	0	
B8	25B-NBOMe-M (O,O-bis-demethyl-) isomer 1					6.0

	MS¹	PM at <i>m/z</i> 352.0542 (M+H)	5	352.0543	C ₁₆ H ₁₉ O ₃ NBr	-0.23	
	MS²	FI at <i>m/z</i> 91.0548	57	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0650	100	121.0653	C ₈ H ₉ O	-2.81	
		FI at <i>m/z</i> 226.9700	1	226.9708	C ₉ H ₈ O ₂ Br	-3.37	
		FI at <i>m/z</i> 335.0267	0.5	335.0283	C ₁₆ H ₁₆ O ₃ Br	-4.72	
B9	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) isomer 2						6.8
	MS¹	PM at <i>m/z</i> 352.0540 (M+H)	11	352.0543	C ₁₆ H ₁₉ O ₃ NBr	-0.80	
	MS²	FI at <i>m/z</i> 107.0495	100	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 213.9630	31	213.9629	C ₈ H ₇ O ₂ Br	0.28	
		FI at <i>m/z</i> 228.9858	90	228.9864	C ₉ H ₁₀ O ₂ Br	-2.69	
		FI at <i>m/z</i> 246.0124	29	246.0124	C ₉ H ₁₃ O ₂ NBr	0	
B10	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) isomer 3						6.9
	MS¹	PM at <i>m/z</i> 352.0541 (M+H)	12	352.0543	C ₁₆ H ₁₉ O ₃ NBr	-0.52	
	MS²	FI at <i>m/z</i> 107.0495	100	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 213.9623	32	213.9629	C ₈ H ₇ O ₂ Br	-3.00	
		FI at <i>m/z</i> 228.9858	89	228.9864	C ₉ H ₁₀ O ₂ Br	-2.69	
		FI at <i>m/z</i> 246.0124	29	246.0124	C ₉ H ₁₃ O ₂ NBr	0	
B11	25B-NBOMe-M (<i>O</i>-demethyl-dehydro-) isomer 1						6.7
	MS¹	PM at <i>m/z</i> 364.0540 (M+H)	25	364.0543	C ₁₇ H ₁₉ O ₃ NBr	-0.77	
	MS²	FI at <i>m/z</i> 107.0495	100	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 228.9857	16	228.9864	C ₉ H ₁₀ O ₂ Br	-3.13	
		FI at <i>m/z</i> 258.0122	71	258.0124	C ₁₀ H ₁₃ O ₂ NBr	-0.84	
B12	25B-NBOMe-M (<i>O</i>-demethyl-dehydro-) isomer 2						7.4
	MS¹	PM at <i>m/z</i> 364.0540 (M+H)	3	364.0543	C ₁₇ H ₁₉ O ₃ NBr	-0.77	
	MS²	FI at <i>m/z</i> 91.0548	57	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0651	100	121.0653	C ₈ H ₉ O	-1.98	
		FI at <i>m/z</i> 227.9655	13	227.9655	C ₈ H ₇ O ₂ NBr	0	
		FI at <i>m/z</i> 242.9890	12	242.9890	C ₉ H ₁₀ O ₂ NBr	0	
B13	25B-NBOMe-M (<i>O</i>-demethyl-) isomer 1						7.6
	MS¹	PM at <i>m/z</i> 366.0700 (M+H)	4	366.0699	C ₁₇ H ₂₁ O ₃ NBr	0.19	
	MS²	FI at <i>m/z</i> 91.0548	58	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0650	100	121.0653	C ₈ H ₉ O	-2.81	
		FI at <i>m/z</i> 228.9861	0.5	228.9864	C ₉ H ₁₀ O ₂ Br	-1.38	
		FI at <i>m/z</i> 257.1167	1	257.1178	C ₁₆ H ₁₇ O ₃	-4.16	
		FI at <i>m/z</i> 349.0433	0.6	349.0439	C ₁₇ H ₁₈ O ₃ Br	-1.81	
B14	25B-NBOMe-M (<i>O</i>-demethyl-) isomer 2						7.7
	MS¹	PM at <i>m/z</i> 366.0703 (M+H)	6	366.0699	C ₁₇ H ₂₁ O ₃ NBr	1.01	
	MS²	FI at <i>m/z</i> 91.0548	59	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0649	100	121.0653	C ₈ H ₉ O	-3.63	
		FI at <i>m/z</i> 228.9855	0.4	228.9864	C ₉ H ₁₀ O ₂ Br	-4.00	
		FI at <i>m/z</i> 243.9961	0.3	243.9968	C ₉ H ₁₁ O ₃ NBr	-2.73	
		FI at <i>m/z</i> 270.1254	0.1	270.1256	C ₁₇ H ₁₈ O ₃	-0.72	
B15	25B-NBOMe-M (<i>O</i>-demethyl-) isomer 3						8.0
	MS¹	PM at <i>m/z</i> 366.0696 (M+H)	3	366.0699	C ₁₇ H ₂₁ O ₃ NBr	-0.91	
	MS²	FI at <i>m/z</i> 107.0495	100	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 227.9779	50	227.9786	C ₉ H ₉ O ₂ Br	-3.03	
		FI at <i>m/z</i> 243.0014	92	243.0021	C ₁₀ H ₁₂ O ₂ Br	-2.74	
		FI at <i>m/z</i> 260.0280	26	260.0281	C ₁₀ H ₁₃ O ₂ NBr	-0.26	
B16	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 1						5.1
	MS¹	PM at <i>m/z</i> 368.0495 (M+H)	4	368.0492	C ₁₆ H ₁₉ O ₄ NBr	0.82	
	MS²	FI at <i>m/z</i> 107.0496	44	107.0497	C ₇ H ₇ O	-0.84	
		FI at <i>m/z</i> 137.0598	100	137.0603	C ₈ H ₉ O ₂ Br	-3.32	
		FI at <i>m/z</i> 228.9858	1	228.9864	C ₉ H ₁₀ O ₂ Br	-2.69	
		FI at <i>m/z</i> 351.0238	0.4	351.0232	C ₁₆ H ₁₆ O ₄ Br	1.72	
B17	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 2						5.6
	MS¹	PM at <i>m/z</i> 368.0481 (M+H)	9	368.0492	C ₁₆ H ₁₉ O ₄ NBr	-2.98	
	MS²	FI at <i>m/z</i> 123.0441	100	123.0446	C ₇ H ₇ O ₂	-4.10	
		FI at <i>m/z</i> 213.9620	26	213.9629	C ₈ H ₇ O ₂ Br	-4.40	
		FI at <i>m/z</i> 228.9853	96	228.9864	C ₉ H ₁₀ O ₂ Br	-4.87	

		FI at m/z 246.0122	30	246.0130	$C_9H_{13}O_2NBr$	-3.11	
B18	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 3						6.3
	MS¹	PM at m/z 368.0489 (M+H)	13	368.0492	$C_{16}H_{19}O_4NBr$	-0.91	
	MS²	FI at m/z 123.0443	100	123.0446	$C_7H_7O_2$	-2.48	
		FI at m/z 213.9624	24	213.9629	$C_8H_9O_2Br$	-2.53	
		FI at m/z 228.9859	72	228.9864	$C_9H_{10}O_2Br$	-2.25	
		FI at m/z 246.0124	22	246.0130	$C_9H_{13}O_2NBr$	-2.30	
B19	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 4						6.6
	MS¹	PM at m/z 368.0495 (M+H)	10	368.0492	$C_{16}H_{19}O_4NBr$	0.82	
	MS²	FI at m/z 107.0495	100	107.0497	C_7H_7O	-1.77	
		FI at m/z 229.9573	61	229.9579	$C_8H_7O_3Br$	-2.42	
		FI at m/z 244.9808	86	244.9813	$C_9H_{10}O_3Br$	-2.17	
		FI at m/z 262.0074	33	262.0073	$C_9H_{13}O_3NBr$	0.26	
B20	25B-NBOMe-M (dehydro-)						7.4
	MS¹	PM at m/z 378.0697 (M+H)	16	378.0699	$C_{18}H_{21}O_3NBr$	-0.61	
	MS²	FI at m/z 91.0548	58	91.0548	C_7H_7	0	
		FI at m/z 121.0650	100	121.0653	C_8H_9O	-2.81	
		FI at m/z 239.9647	1	239.9655	$C_9H_7O_2NBr$	-3.19	
		FI at m/z 255.9966	8	255.9968	$C_{10}H_{11}O_2NBr$	-0.65	
B21	25B-NBOMe-M (<i>O</i>-demethyl-dehydro-hydroxy-)						5.7
	MS¹	PM at m/z 380.0494 (M+H)	20	380.0494	$C_{17}H_{19}O_4NBr$	0.54	
	MS²	FI at m/z 107.0496	100	107.0497	C_7H_7O	-0.84	
		FI at m/z 244.9809	9	244.9813	$C_9H_{10}O_3Br$	-1.76	
		FI at m/z 274.0072	55	274.0073	$C_{10}H_{13}O_3NBr$	-0.48	
B22	25B-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 1						6.2
	MS¹	PM at m/z 382.0665 (M+H)	7	382.0648	$C_{17}H_{21}O_4NBr$	4.33	
	MS²	FI at m/z 91.0548	53	91.0548	C_7H_7	0	
		FI at m/z 121.0650	100	121.0653	C_8H_9O	-2.81	
		FI at m/z 228.9859	4	228.9864	$C_9H_{10}O_3Br$	-2.25	
		FI at m/z 365.0403	0.2	365.0388	$C_{10}H_{15}O_3NBr$	3.98	
B23	25B-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 2						6.7
	MS¹	PM at m/z 382.0643 (M+H)	7	382.0648	$C_{17}H_{21}O_4NBr$	-1.43	
	MS²	FI at m/z 107.0494	44	107.0497	C_7H_7O	-2.71	
		FI at m/z 137.0597	100	137.0603	$C_8H_9O_2$	-4.05	
		FI at m/z 228.9861	1	228.9864	$C_9H_{10}O_2Br$	-1.38	
		FI at m/z 243.9975	0.2	243.9968	$C_9H_{11}O_3NBr$	3.01	
		FI at m/z 365.0375	0.1	365.0388	$C_{17}H_{18}O_4Br$	-3.69	
B24	25B-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 3						6.8
	MS¹	PM at m/z 382.0644 (M+H)	11	382.0648	$C_{17}H_{21}O_4NBr$	-1.17	
	MS²	FI at m/z 123.0442	85	123.0446	$C_7H_7O_2$	-3.29	
		FI at m/z 227.9780	55	227.9786	$C_9H_9O_2Br$	-2.59	
		FI at m/z 243.0014	100	243.0021	$C_{10}H_{12}O_2Br$	-2.74	
		FI at m/z 260.0281	28	260.0281	$C_{10}H_{15}O_2NBr$	0	
B25	25B-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 4						7.2
	MS¹	PM at m/z 382.0646 (M+H)	8	382.0648	$C_{17}H_{21}O_4NBr$	-0.65	
	MS²	FI at m/z 123.0442	100	123.0446	$C_7H_7O_2$	-3.29	
		FI at m/z 227.9779	56	227.9786	$C_9H_9O_2Br$	-3.03	
		FI at m/z 243.0014	99	243.0021	$C_{10}H_{12}O_2Br$	-2.74	
		FI at m/z 260.0280	28	260.0281	$C_{10}H_{15}O_2NBr$	-0.26	
B26	25B-NBOMe-M (dehydro-hydroxy-) isomer 1						6.5
	MS¹	PM at m/z 394.0652 (M+H)	30	394.0648	$C_{18}H_{21}O_4NBr$	0.90	
	MS²	FI at m/z 109.0651	100	109.0653	C_7H_9O	-2.20	
		FI at m/z 137.0597	26	137.0603	$C_8H_9O_2$	-4.05	
		FI at m/z 239.9647	1	239.9655	$C_9H_7O_2NBr$	-3.19	
		FI at m/z 255.9965	9	255.9968	$C_{10}H_{11}O_2NBr$	-1.04	
B27	25B-NBOMe-M (dehydro-hydroxy-) isomer 2						6.6
	MS¹	PM at m/z 394.0650 (M+H)	19	394.0648	$C_{18}H_{21}O_4NBr$	0.39	

	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 256.9681 FI at <i>m/z</i> 271.9916	56 100 2 7	91.0548 121.0653 256.9681 271.9917	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₈ O ₃ NBr C ₁₀ H ₁₁ O ₃ NBr	0 -2.81 -0.41 -0.30	
B28	25B-NBOMe-M (dehydro-hydroxy-) isomer 3						6.7
	MS¹	PM at <i>m/z</i> 394.0658 (M+H)	6	394.0648	C ₁₈ H ₂₃ O ₄ NBr	2.42	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 137.0597 FI at <i>m/z</i> 239.9657 FI at <i>m/z</i> 255.9967	38 100 1 6	107.0497 137.0603 239.9655 255.9968	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₉ H ₇ O ₂ NBr C ₁₀ H ₁₁ O ₃ NBr	-1.77 -4.05 0.97 -0.26	
B29	25B-NBOMe-M (hydroxy-) isomer 1						7.1
	MS¹	PM at <i>m/z</i> 396.0806 (M+H)	12	396.0805	C ₁₈ H ₂₃ O ₄ NBr	0.26	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 243.9732 FI at <i>m/z</i> 258.9967	62 100 4 11	91.0548 121.0653 243.9735 258.9970	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₈ O ₃ Br C ₁₀ H ₁₂ O ₃ Br	0 -2.81 -1.25 -1.08	
B30	25B-NBOMe-M (hydroxy-) isomer 2						7.3
	MS¹	PM at <i>m/z</i> 396.0800 (M+H)	17	396.0805	C ₁₈ H ₂₃ O ₄ NBr	-1.25	
	MS²	FI at <i>m/z</i> 109.0651 FI at <i>m/z</i> 137.0596 FI at <i>m/z</i> 243.0015 FI at <i>m/z</i> 258.0123	100 23 4 1	109.0653 137.0603 243.0021 258.0124	C ₇ H ₉ O C ₈ H ₉ O ₂ C ₁₀ H ₁₂ O ₂ Br C ₁₀ H ₁₃ O ₂ NBr	-2.20 -4.78 -2.33 -0.45	
B31	25B-NBOMe-M (hydroxy-) isomer 3						7.8
	MS¹	PM at <i>m/z</i> 396.0805 (M+H)	10	396.0805	C ₁₈ H ₂₃ O ₄ NBr	0	
	MS²	FI at <i>m/z</i> 107.0494 FI at <i>m/z</i> 137.0597 FI at <i>m/z</i> 243.0010 FI at <i>m/z</i> 258.0137	53 100 2 1	107.0497 137.0603 243.0021 258.0124	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₁₀ H ₁₂ O ₂ Br C ₁₀ H ₁₃ O ₂ NBr	-2.71 -4.05 -4.39 4.97	
B32	25B-NBOMe-M (hydroxy-) isomer 4						8.4
	MS¹	PM at <i>m/z</i> 396.0800 (M+H)	6	396.0805	C ₁₈ H ₂₃ O ₄ NBr	-1.25	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0651 FI at <i>m/z</i> 258.0128 FI at <i>m/z</i> 378.0704	54 100 4 2	91.0548 121.0653 258.0124 378.0699	C ₇ H ₇ C ₈ H ₉ O C ₁₀ H ₁₃ O ₂ NBr C ₁₈ H ₂₁ O ₃ NBr	0 -1.98 1.49 1.24	
B33	25B-NBOMe-M (O-demethyl-bis-hydroxy-)						6.2
	MS¹	PM at <i>m/z</i> 398.0590 (M+H)	1	398.0598	C ₁₇ H ₂₁ O ₅ NBr	-1.91	
	MS²	FI at <i>m/z</i> 138.0312 FI at <i>m/z</i> 153.0547 FI at <i>m/z</i> 228.9861 FI at <i>m/z</i> 246.0123	39 100 71 25	138.0317 153.0552 228.9864 246.0124	C ₇ H ₆ O ₃ C ₈ H ₉ O ₃ C ₉ H ₁₀ O ₂ Br C ₉ H ₁₃ O ₂ NBr	-3.59 -3.07 -1.38 -0.47	
B34	25B-NBOMe-M (bis-hydroxy-) isomer 1						7.4
	MS¹	PM at <i>m/z</i> 412.0750 (M+H)	1	412.0754	C ₁₈ H ₂₃ O ₅ NBr	-1.00	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 153.0544 FI at <i>m/z</i> 243.0010 FI at <i>m/z</i> 260.0282	54 63 100 32	107.0497 153.0552 243.0021 260.0281	C ₇ H ₇ O C ₈ H ₉ O ₃ C ₁₀ H ₁₂ O ₂ Br C ₁₀ H ₁₃ O ₂ NBr	-1.77 -5.03 -4.39 0.51	
B35	25B-NBOMe-M (bis-hydroxy-) isomer 2						7.7
	MS¹	PM at <i>m/z</i> 412.0756 (M+H)	6	412.0754	C ₁₈ H ₂₃ O ₅ NBr	0.46	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 274.9907 FI at <i>m/z</i> 290.0018	60 100 1 0.3	91.0548 121.0653 274.9919 290.0022	C ₇ H ₇ C ₈ H ₉ O C ₁₀ H ₁₂ O ₄ Br C ₁₀ H ₁₃ O ₄ NBr	0 -2.81 -4.35 -1.54	

Table S2 List of 25C-NBOMe and its phase I metabolites together with the precursor mass (PM) recorded in MS¹, the corresponding characteristic fragment ions (FI) in MS², the calculated exact masses, the corresponding elemental composition, the deviation of the measured from the calculated masses, given as errors in parts per million (ppm), and the retention times (RT) in minutes (min). The metabolites were sorted by mass and RT.

No.	Metabolite and characteristic ions Measured accurate mass, <i>m/z</i>	Relative intensity in MS ² , %	Calculated exact mass, <i>m/z</i>	Elemental composition	Error, ppm	RT, min
C1	25C-NBOMe					8.5
	MS ¹ PM at <i>m/z</i> 336.1360 (M+H)	8	336.1361	C ₁₈ H ₂₅ O ₃ NCl	-0.29	
	MS ² FI at <i>m/z</i> 91.0548	56	91.0548	C ₇ H ₇	0	
	FI at <i>m/z</i> 121.0651	100	121.0653	C ₈ H ₉ O	-1.98	
	FI at <i>m/z</i> 184.0286	0.3	184.0291	C ₉ H ₉ O ₂ Cl	-2.76	
	FI at <i>m/z</i> 199.0522	1	199.0526	C ₁₀ H ₁₂ O ₂ Cl	-1.92	
	FI at <i>m/z</i> 214.0627	0.2	214.0629	C ₁₀ H ₁₃ O ₂ NCl	-1.09	
C2	25C-NBOMe-M (N-demethoxybenzyl-O-demethyl-) isomer 1					3.4
	MS ¹ PM at <i>m/z</i> 202.0629 (M+H)	1	202.0629	C ₉ H ₁₃ O ₂ NCl	0	
	MS ² FI at <i>m/z</i> 135.0441	6	135.0446	C ₈ H ₇ O ₂	-3.74	
	FI at <i>m/z</i> 150.0676	23	150.0681	C ₉ H ₁₀ O ₂	-3.20	
	FI at <i>m/z</i> 170.0129	83	170.0135	C ₈ H ₇ O ₂ Cl	-3.28	
	FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
C3	25C-NBOMe-M (N-demethoxybenzyl-O-demethyl-) isomer 2					3.5
	MS ¹ PM at <i>m/z</i> 202.0628 (M+H)	1	202.0629	C ₉ H ₁₃ O ₂ NCl	-0.66	
	MS ² FI at <i>m/z</i> 135.0441	5	135.0446	C ₈ H ₇ O ₂	-3.74	
	FI at <i>m/z</i> 150.0675	18	150.0681	C ₉ H ₁₀ O ₂	-3.86	
	FI at <i>m/z</i> 170.0129	84	170.0135	C ₈ H ₇ O ₂ Cl	-1.84	
	FI at <i>m/z</i> 185.0364	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.88	
C4	25C-NBOMe-M (N-demethoxybenzyl-O-demethyl-deamino-carboxy-)					5.9
	MS ¹ PM at <i>m/z</i> 215.0108 (M+H)	1	215.0111	C ₉ H ₈ O ₂ Cl	-1.46	
	MS ² FI at <i>m/z</i> 155.9976	100	155.9978	C ₇ H ₅ O ₂ Cl	-1.33	
	FI at <i>m/z</i> 171.0210	2	171.0213	C ₈ H ₈ O ₂ Cl	-1.65	
C5	25C-NBOMe-M (N-demethoxybenzyl-)					5.2
	MS ¹ PM at <i>m/z</i> 216.0781 (M+H)	1	216.0786	C ₁₀ H ₁₅ O ₂ NCl	-2.24	
	MS ² FI at <i>m/z</i> 164.0829	20	164.0837	C ₁₀ H ₁₂ O ₂	-5.06	
	FI at <i>m/z</i> 169.0048	28	169.0056	C ₈ H ₆ O ₂ Cl	-4.93	
	FI at <i>m/z</i> 184.0284	100	184.0291	C ₉ H ₉ O ₂ Cl	-3.85	
	FI at <i>m/z</i> 199.0517	86	199.0526	C ₁₀ H ₁₂ O ₂ Cl	-4.44	
C6	25C-NBOMe-M (N-demethoxybenzyl-oxo-)					6.6
	MS ¹ PM at <i>m/z</i> 230.0581 (M+H)	1	230.0578	C ₁₀ H ₁₃ O ₃ NCl	1.09	
	MS ² FI at <i>m/z</i> 155.0257	42	155.0264	C ₈ H ₈ OCl	-4.31	
	FI at <i>m/z</i> 173.0364	2	173.0369	C ₈ H ₁₀ O ₂ Cl	-3.08	
	FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
C7	25C-NBOMe-M (N-demethoxybenzyl-hydroxy-)					3.8
	MS ¹ PM at <i>m/z</i> 232.0730 (M+H)	1	232.0735	C ₁₀ H ₁₅ O ₃ NCl	-2.15	
	MS ² FI at <i>m/z</i> 185.0000	42	185.0005	C ₈ H ₆ O ₃ Cl	-2.96	
	FI at <i>m/z</i> 200.0234	30	200.0240	C ₈ H ₉ O ₃ Cl	-3.11	
	FI at <i>m/z</i> 215.0469	100	215.0475	C ₁₀ H ₁₂ O ₃ Cl	-2.78	
C8	25C-NBOMe-M (O,O,O-tris-demethyl-)					5.0
	MS ¹ PM at <i>m/z</i> 294.0879 (M+H)	6	294.0891	C ₁₃ H ₁₇ O ₃ NCl	-4.25	
	MS ² FI at <i>m/z</i> 107.0494	100	107.0497	C ₇ H ₇ O	-2.71	
	FI at <i>m/z</i> 136.0518	6	136.0524	C ₈ H ₈ O ₂	-4.63	
	FI at <i>m/z</i> 171.0206	80	171.0213	C ₈ H ₈ O ₂ Cl	-3.99	

		FI at m/z 188.0472	27	188.0473	$C_8H_{11}O_2NCl$	-0.44	
C9	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-dehydro-)						6.4
	MS¹	PM at m/z 306.0893 (M+H)	4	306.0891	$C_{16}H_{17}O_3NCl$	0.49	
	MS²	FI at m/z 107.0495	36	107.0497	C_7H_7O	-1.77	
		FI at m/z 184.0160	46	184.0160	$C_8H_7O_2NCl$	0	
		FI at m/z 199.0395	41	199.0395	$C_9H_{10}O_2NCl$	0	
		FI at m/z 200.0474	100	200.0473	$C_9H_{11}O_2NCl$	0.58	
C10	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-) isomer 1						5.8
	MS¹	PM at m/z 308.1046 (M+H)	6	308.1048	$C_{16}H_{19}O_3NCl$	-0.64	
	MS²	FI at m/z 91.0547	51	91.0548	C_7H_7	-0.82	
		FI at m/z 121.0649	100	121.0653	C_8H_9O	-3.63	
		FI at m/z 185.0360	7	185.0369	$C_9H_{10}O_2Cl$	-5.04	
		FI at m/z 202.0624	2	202.0629	$C_9H_{13}O_2NCl$	-2.64	
C11	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-) isomer 2						6.4
	MS¹	PM at m/z 308.1051 (M+H)	13	308.1048	$C_{16}H_{19}O_3NCl$	0.98	
	MS²	FI at m/z 107.0496	100	107.0497	C_7H_7O	-0.84	
		FI at m/z 170.0129	13	170.0135	$C_8H_7O_2Cl$	-3.28	
		FI at m/z 185.0364	77	185.0369	$C_9H_{10}O_2Cl$	-2.88	
		FI at m/z 202.0632	22	202.0629	$C_9H_{13}O_2NCl$	1.32	
C12	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-) isomer 3						6.6
	MS¹	PM at m/z 308.1048 (M+H)	10	308.1048	$C_{16}H_{19}O_3NCl$	0	
	MS²	FI at m/z 107.0496	100	107.0497	C_7H_7O	-0.84	
		FI at m/z 170.0130	25	170.0135	$C_8H_7O_2Cl$	-2.69	
		FI at m/z 185.0365	87	185.0369	$C_9H_{10}O_2Cl$	-2.34	
		FI at m/z 202.0629	22	202.0629	$C_9H_{13}O_2NCl$	0	
C13	25C-NBOMe-M (<i>O</i>-demethyl-dehydro-) isomer 1						5.7
	MS¹	PM at m/z 320.1047 (M+H)	17	320.1048	$C_{17}H_{19}O_3NCl$	-0.31	
	MS²	FI at m/z 91.0548	57	91.0548	C_7H_7	0	
		FI at m/z 121.0651	100	121.0653	C_8H_9O	-1.98	
		FI at m/z 198.0317	5	198.0316	$C_9H_9O_2N$	-3.14	
C14	25C-NBOMe-M (<i>O</i>-demethyl-dehydro-) isomer 2						6.4
	MS¹	PM at m/z 320.1049 (M+H)	24	320.1048	$C_{17}H_{19}O_3NCl$	0.32	
	MS²	FI at m/z 107.0495	100	107.0497	C_7H_7O	-1.77	
		FI at m/z 185.0365	17	185.0369	$C_9H_{10}O_2Cl$	-2.34	
		FI at m/z 214.0629	73	214.0629	$C_{10}H_{13}O_2NCl$	0	
C15	25C-NBOMe-M (<i>O</i>-demethyl-dehydro-) isomer 3						7.1
	MS¹	PM at m/z 320.1050 (M+H)	3	320.1048	$C_{17}H_{19}O_3NCl$	0.63	
	MS²	FI at m/z 91.0548	49	91.0548	C_7H_7	0.27	
		FI at m/z 121.0650	100	121.0653	C_8H_9O	-2.81	
		FI at m/z 184.0160	10	184.0160	$C_8H_7O_2NCl$	0	
		FI at m/z 199.0394	11	199.0395	$C_9H_{10}O_2NCl$	-0.29	
C16	25C-NBOMe-M (<i>O</i>-demethyl-) isomer 1						7.2
	MS¹	PM at m/z 322.1206 (M+H)	10	322.1204	$C_{17}H_{21}O_3NCl$	0.47	
	MS²	FI at m/z 91.0548	54	91.0548	C_7H_7	0	
		FI at m/z 121.0651	100	121.0653	C_8H_9O	-1.98	
		FI at m/z 185.0368	1	185.0369	$C_9H_{10}O_2Cl$	-0.72	
		FI at m/z 200.0473	1	200.0473	$C_9H_{11}O_2NCl$	0	
		FI at m/z 305.0931	0.1	305.0944	$C_{17}H_{18}O_3Cl$	-4.42	
C17	25C-NBOMe-M (<i>O</i>-demethyl-) isomer 2						7.3
	MS¹	PM at m/z 322.1199 (M+H)	6	322.1204	$C_{17}H_{21}O_3NCl$	-1.70	
	MS²	FI at m/z 91.0549	58	91.0548	C_7H_7	1.37	
		FI at m/z 121.0651	100	121.0653	C_8H_9O	-1.98	
		FI at m/z 185.0364	0.3	185.0369	$C_9H_{10}O_2Cl$	-2.88	
		FI at m/z 200.0473	0.2	200.0473	$C_9H_{11}O_2NCl$	0	
		FI at m/z 305.0934	0.1	305.0944	$C_{17}H_{18}O_3Cl$	-3.44	
C18	25C-NBOMe-M (<i>O</i>-demethyl-) isomer 3						7.7
	MS¹	PM at m/z 322.1198 (M+H)	10	322.1204	$C_{17}H_{21}O_3NCl$	-2.01	

	MS²	FI at <i>m/z</i> 107.0494 FI at <i>m/z</i> 184.0285 FI at <i>m/z</i> 199.0518 FI at <i>m/z</i> 216.0784	100 44 90 21	107.0497 184.0291 199.0526 216.0786	C ₇ H ₇ O C ₈ H ₉ O ₂ Cl C ₁₀ H ₁₂ O ₂ Cl C ₁₀ H ₁₅ O ₂ NCl	-2.71 -3.30 -3.93 -0.85	
C19	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 1						5.0
	MS¹	PM at <i>m/z</i> 324.1006 (M+H)	4	324.0997	C ₁₆ H ₁₉ O ₄ NCl	2.73	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 137.0597 FI at <i>m/z</i> 185.0365 FI at <i>m/z</i> 202.0637 FI at <i>m/z</i> 307.0722	44 100 4 1 1	107.0497 137.0603 185.0369 202.0629 307.0737	C ₇ H ₇ O C ₈ H ₉ O ₂ Cl C ₉ H ₁₀ O ₂ Cl C ₁₀ H ₁₃ O ₂ NCl C ₁₆ H ₁₆ O ₄ Cl	-1.77 -4.05 -2.34 3.79 -4.93	
C20	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 2						5.3
	MS¹	PM at <i>m/z</i> 324.0997 (M+H)	12	324.0997	C ₁₆ H ₁₉ O ₄ NCl	0	
	MS²	FI at <i>m/z</i> 123.0443 FI at <i>m/z</i> 170.0129 FI at <i>m/z</i> 185.0365 FI at <i>m/z</i> 202.0630	95 19 100 26	123.0446 170.0135 185.0369 202.0635	C ₇ H ₇ O ₂ C ₈ H ₇ O ₂ Cl C ₉ H ₁₀ O ₂ Cl C ₉ H ₁₃ O ₂ NCl	-2.48 -3.28 -2.34 -2.39	
C21	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 3						5.6
	MS¹	PM at <i>m/z</i> 324.0995 (M+H)	3	324.0997	C ₁₆ H ₁₉ O ₄ NCl	-0.66	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 186.0078 FI at <i>m/z</i> 201.0312 FI at <i>m/z</i> 218.0578	100 41 94 31	107.0497 186.0084 201.0318 218.0578	C ₇ H ₇ O C ₈ H ₇ O ₃ Cl C ₉ H ₁₀ O ₃ Cl C ₉ H ₁₃ O ₃ NCl	-1.77 -3.08 -3.22 0	
C22	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 4						5.8
	MS¹	PM at <i>m/z</i> 324.0995 (M+H)	11	324.0997	C ₁₆ H ₁₉ O ₄ NCl	-0.66	
	MS²	FI at <i>m/z</i> 123.0443 FI at <i>m/z</i> 170.0130 FI at <i>m/z</i> 185.0365 FI at <i>m/z</i> 202.0628	100 12 65 17	123.0446 170.0135 185.0369 202.0635	C ₇ H ₇ O ₂ C ₈ H ₇ O ₂ Cl C ₉ H ₁₀ O ₂ Cl C ₉ H ₁₃ O ₂ NCl	-2.48 -2.69 -2.34 -3.38	
C23	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) isomer 5						6.2
	MS¹	PM at <i>m/z</i> 324.0990 (M+H)	1	324.0997	C ₁₆ H ₁₉ O ₄ NCl	-2.20	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 200.0473 FI at <i>m/z</i> 218.0581 FI at <i>m/z</i> 306.0890	100 99 1 9	107.0497 200.0473 218.0578 306.0891	C ₇ H ₇ O C ₈ H ₁₁ O ₂ NCl C ₉ H ₁₃ O ₃ NCl C ₁₆ H ₁₇ O ₃ NCl	-1.77 0 1.15 -0.49	
C24	25C-NBOMe-M (dehydro-)						7.2
	MS¹	PM at <i>m/z</i> 334.1202 (M+H)	20	334.1204	C ₁₈ H ₂₁ O ₃ NCl	-0.74	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0651 FI at <i>m/z</i> 196.0160 FI at <i>m/z</i> 212.0472	53 100 1 7	91.0548 121.0653 196.0160 212.0473	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₇ O ₂ NCl C ₁₀ H ₁₁ O ₂ NCl	0 -1.98 0 -0.39	
C25	25C-NBOMe-M (<i>O</i>-demethyl-dehydro-hydroxy-)						6.0
	MS¹	PM at <i>m/z</i> 336.0983 (M+H)	3	336.0997	C ₁₇ H ₂₁ O ₄ NCl	-4.21	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 137.0597 FI at <i>m/z</i> 184.0160 FI at <i>m/z</i> 199.0394	42 100 11 11	107.0497 137.0603 184.0160 199.0395	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₈ H ₇ O ₂ NCl C ₉ H ₁₀ O ₂ NCl	-1.77 -4.05 0 -0.29	
C26	25C-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 1						6.1
	MS¹	PM at <i>m/z</i> 338.1145 (M+H)	9	338.1154	C ₁₇ H ₂₁ O ₄ NCl	-2.55	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 200.0235 FI at <i>m/z</i> 215.0469 FI at <i>m/z</i> 232.0733	100 78 86 26	107.0497 200.0240 215.0475 232.0735	C ₇ H ₇ O C ₉ H ₉ O ₃ Cl C ₁₀ H ₁₂ O ₃ Cl C ₁₀ H ₁₅ O ₃ NCl	-1.77 -3.32 -2.78 -0.86	
C27	25C-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 2						6.4
	MS¹	PM at <i>m/z</i> 338.1151 (M+H)	7	338.1154	C ₁₇ H ₂₁ O ₄ NCl	-0.78	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 137.0598 FI at <i>m/z</i> 185.0365 FI at <i>m/z</i> 200.0465	41 100 1 0.2	107.0497 137.0603 185.0369 200.0473	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₉ H ₁₀ O ₂ Cl C ₉ H ₁₁ O ₂ NCl	-1.77 -3.32 -2.34 -3.92	

		FI at m/z 321.0878	0.2	321.0894	$C_{17}H_{18}O_4Cl$	-4.87	
C28	25C-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 3						6.5
	MS¹	PM at m/z 338.1146 (M+H)	10	338.1154	$C_{17}H_{21}O_4NCl$	-2.26	
	MS²	FI at m/z 123.0442	100	123.0446	$C_7H_7O_2$	-3.29	
		FI at m/z 184.0286	47	184.0291	$C_9H_9O_2Cl$	-2.76	
		FI at m/z 199.0521	91	199.0526	$C_{10}H_{12}O_2Cl$	-2.43	
		FI at m/z 216.0788	22	216.0786	$C_{10}H_{13}O_2NCl$	1.00	
C29	25C-NBOMe-M (<i>O</i>-demethyl-hydroxy-) isomer 4						7.1
	MS¹	PM at m/z 338.1164 (M+H)	5	338.1154	$C_{17}H_{21}O_4NCl$	3.07	
	MS²	FI at m/z 123.0443	100	123.0446	$C_7H_7O_2$	-2.48	
		FI at m/z 184.0287	39	184.0291	$C_9H_9O_2Cl$	-2.22	
		FI at m/z 199.0521	76	199.0526	$C_{10}H_{12}O_2Cl$	-2.43	
		FI at m/z 216.0786	15	216.0786	$C_{10}H_{13}O_2NCl$	0	
C30	25C-NBOMe-M (dehydro-hydroxy-)						6.3
	MS¹	PM at m/z 350.1148 (M+H)	17	350.1154	$C_{18}H_{21}O_4NCl$	-1.61	
	MS²	FI at m/z 91.0547	54	91.0548	C_7H_7	-0.82	
		FI at m/z 121.0649	100	121.0653	C_8H_9O	-3.63	
		FI at m/z 213.0185	2	213.0187	$C_9H_8O_3NCl$	-1.05	
		FI at m/z 228.0421	6	228.0422	$C_{10}H_{11}O_3NCl$	-0.43	
C31	25C-NBOMe-M (hydroxy-) isomer 1						6.9
	MS¹	PM at m/z 352.1303 (M+H)	10	352.1310	$C_{18}H_{23}O_4NCl$	-2.03	
	MS²	FI at m/z 91.0546	51	91.0548	C_7H_7	-1.92	
		FI at m/z 121.0649	100	121.0653	C_8H_9O	-3.63	
		FI at m/z 200.0231	2	200.0240	$C_9H_9O_3Cl$	-4.61	
		FI at m/z 215.0465	2	215.0475	$C_{10}H_{12}O_3Cl$	-4.64	
C32	25C-NBOMe-M (hydroxy-) isomer 2						7.1
	MS¹	PM at m/z 352.1304 (M+H)	13	352.1310	$C_{18}H_{23}O_4NCl$	-1.74	
	MS²	FI at m/z 91.0549	61	91.0548	C_7H_7	1.37	
		FI at m/z 121.0651	100	121.0653	C_8H_9O	-1.98	
		FI at m/z 200.0236	3	200.0240	$C_9H_9O_3Cl$	-2.11	
		FI at m/z 215.0472	10	215.0475	$C_{10}H_{12}O_3Cl$	-1.39	
C33	25C-NBOMe-M (hydroxy-) isomer 3						7.6
	MS¹	PM at m/z 352.1305 (M+H)	9	352.1310	$C_{18}H_{23}O_4NCl$	-1.46	
	MS²	FI at m/z 107.0494	43	107.0497	C_7H_7O	-2.71	
		FI at m/z 137.0596	100	137.0603	$C_8H_9O_2$	-4.78	
		FI at m/z 184.0283	2	184.0291	$C_9H_9O_2Cl$	-4.39	
		FI at m/z 199.0520	5	199.0526	$C_{10}H_{12}O_2Cl$	-2.93	
C34	25C-NBOMe-M (<i>O</i>-demethyl-bis-hydroxy-)						5.7
	MS¹	PM at m/z 354.1090 (M+H)	2	354.1103	$C_{17}H_{21}O_3NCl$	-3.61	
	MS²	FI at m/z 107.0494	60	107.0497	C_7H_7O	-2.71	
		FI at m/z 153.0550	60	153.0552	$C_8H_9O_3$	-1.11	
		FI at m/z 185.0362	100	185.0369	$C_9H_{10}O_2Cl$	-3.96	
		FI at m/z 202.0629	23	202.0629	$C_9H_{13}O_2NCl$	0	
C35	25C-NBOMe-M (bis-hydroxy-) isomer 1						6.0
	MS¹	PM at m/z 368.1241 (M+H)	6	368.1259	$C_{18}H_{23}O_3NCl$	-4.97	
	MS²	FI at m/z 107.0495	40	107.0497	C_7H_7O	-1.77	
		FI at m/z 137.0597	100	137.0603	$C_8H_9O_2$	-4.05	
		FI at m/z 215.0467	1	215.0475	$C_{10}H_{12}O_3Cl$	-3.71	
		FI at m/z 230.0570	1	230.0578	$C_{10}H_{13}O_3NCl$	-3.69	
C36	25C-NBOMe-M (bis-hydroxy-) isomer 2						6.9
	MS¹	PM at m/z 368.1247 (M+H)	1	368.1259	$C_{18}H_{23}O_3NCl$	-3.34	
	MS²	FI at m/z 107.0495	48	107.0497	C_7H_7O	-1.77	
		FI at m/z 153.0546	58	153.0552	$C_8H_9O_3$	-3.72	
		FI at m/z 199.0521	100	199.0526	$C_{10}H_{12}O_2Cl$	-2.43	
		FI at m/z 216.0782	17	216.0786	$C_{10}H_{13}O_2NCl$	-1.78	

Table S3 List of all 25B-NBOMe phase II metabolites together with the precursor mass (PM) recorded in MS¹, the corresponding characteristic fragment ions (FI) in MS², the calculated exact masses, the corresponding elemental composition, the deviation of the measured from the calculated masses, given as errors in parts per million (ppm), and the retention times (RT) in minutes (min). The metabolites were sorted by mass and RT. Numbering according to Table 1 (AC = *N*-acetylation, GSH = glutathione conjugation, ME = *O*-methylation, G = glucuronidation, S = sulfation, AC+G = acetylation combined with glucuronidation, AC+S = acetylation combined with sulfation)

No.	Metabolite and characteristic ions Measured accurate mass, <i>m/z</i>	Relative intensity in MS ² , %	Calculated exact mass, <i>m/z</i>	Elemental composition	Error, ppm	RT, min
B3 AC	25B-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl isomer 1					6.6
	MS¹ PM at <i>m/z</i> 288.0226 (M+H)	5	288.0230	C ₁₁ H ₁₅ O ₃ NBr	-1.32	
	MS² FI at <i>m/z</i> 150.0674	58	150.0681	C ₉ H ₁₀ O ₂	-4.53	
	FI at <i>m/z</i> 213.9621	35	213.9629	C ₈ H ₇ O ₂ Br	-3.93	
	FI at <i>m/z</i> 228.9857	100	228.9864	C ₉ H ₁₀ O ₂ Br	-3.13	
	FI at <i>m/z</i> 246.0123	13	246.0124	C ₉ H ₁₃ O ₂ NBr	-0.47	
B4 AC	25B-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl isomer 2					6.7
	MS¹ PM at <i>m/z</i> 288.0226 (M+H)	6	288.0230	C ₁₁ H ₁₅ O ₃ NBr	-1.32	
	MS² FI at <i>m/z</i> 150.0674	32	150.0681	C ₉ H ₁₀ O ₂	-4.53	
	FI at <i>m/z</i> 213.9621	64	213.9629	C ₈ H ₇ O ₂ Br	-3.93	
	FI at <i>m/z</i> 228.9857	100	228.9864	C ₉ H ₁₀ O ₂ Br	-3.13	
	FI at <i>m/z</i> 246.0123	13	246.0124	C ₉ H ₁₃ O ₂ NBr	-0.47	
B3 AC+S	25B-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl sulfate isomer 1					5.4
	MS¹ PM at <i>m/z</i> 367.9798 (M+H)	4	367.9798	C ₁₁ H ₁₅ O ₆ NBrS	0	
	MS² FI at <i>m/z</i> 228.9858	100	228.9864	C ₉ H ₁₀ O ₂ Br	-2.69	
	FI at <i>m/z</i> 246.0125	8	246.0124	C ₉ H ₁₃ O ₂ NBr	0.34	
	FI at <i>m/z</i> 288.0229	7	288.0230	C ₁₁ H ₁₅ O ₃ NBr	-0.28	
	FI at <i>m/z</i> 308.9424	24	308.9432	C ₉ H ₁₀ O ₃ BrS	-2.70	
	FI at <i>m/z</i> 325.9692	18	325.9692	C ₉ H ₁₃ O ₅ NBrS	0	
B4 AC+S	25B-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl sulfate isomer 2					5.5
	MS¹ PM at <i>m/z</i> 367.9798 (M+H)	4	367.9798	C ₁₁ H ₁₅ O ₆ NBrS	0	
	MS² FI at <i>m/z</i> 228.9858	100	228.9864	C ₉ H ₁₀ O ₂ Br	-2.69	
	FI at <i>m/z</i> 246.0123	12	246.0124	C ₉ H ₁₃ O ₂ NBr	-0.47	
	FI at <i>m/z</i> 288.0226	11	288.0230	C ₁₁ H ₁₅ O ₃ NBr	-1.32	
	FI at <i>m/z</i> 308.9425	16	308.9432	C ₉ H ₁₀ O ₃ BrS	-2.37	
	FI at <i>m/z</i> 325.9692	17	325.9692	C ₉ H ₁₃ O ₅ NBrS	0	
B8 GSH-1	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) <i>S</i>-methyl					7.7
	MS¹ PM at <i>m/z</i> 398.0405 (M+H)	1	398.0420	C ₁₇ H ₂₁ O ₃ NBrS	-3.78	
	MS² FI at <i>m/z</i> 91.0548	53	91.0548	C ₇ H ₇	0	
	FI at <i>m/z</i> 121.0650	100	121.0653	C ₈ H ₉ O	-2.81	
	FI at <i>m/z</i> 259.9503	6	259.9507	C ₉ H ₉ O ₂ BrS	-1.40	
	FI at <i>m/z</i> 381.0149	0.4	381.0160	C ₁₇ H ₁₈ O ₃ BrS	-2.84	
B33 ME	25B-NBOMe-M (<i>O</i>-demethyl-bis-hydroxy-) <i>O</i>-methyl					6.8
	MS¹ PM at <i>m/z</i> 412.0760 (M+H)	1	412.0754	C ₁₈ H ₂₅ O ₃ NBr	1.43	
	MS² FI at <i>m/z</i> 137.0597	39	137.0603	C ₈ H ₉ O ₂	-4.05	
	FI at <i>m/z</i> 167.0704	100	167.0708	C ₉ H ₁₁ O ₃	-2.51	
	FI at <i>m/z</i> 228.9859	18	228.9864	C ₉ H ₁₀ O ₂ Br	-2.25	
	FI at <i>m/z</i> 246.0125	6	246.0124	C ₉ H ₁₃ O ₂ NBr	0.34	
B7 S	25B-NBOMe-M (<i>O,O,O</i>-tris-demethyl-) sulfate					4.7
	MS¹ PM at <i>m/z</i> 417.9959 (M+H)	6	417.9954	C ₁₅ H ₁₇ O ₆ NBrS	1.08	

	MS²	FI at <i>m/z</i> 107.0496 FI at <i>m/z</i> 214.9702 FI at <i>m/z</i> 294.9271 FI at <i>m/z</i> 311.9538 FI at <i>m/z</i> 338.0387	100 80 18 27 10	107.0497 214.9708 294.9276 311.9536 338.0386	C ₇ H ₇ O C ₈ H ₈ O ₂ Br C ₈ H ₈ O ₃ BrS C ₈ H ₁₁ O ₃ NBrS C ₁₅ H ₁₇ O ₃ NBr	-0.84 -2.63 -1.64 0.69 0.20	
B3/4 G	25B-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) glucuronide						3.0
	MS¹	PM at <i>m/z</i> 422.0446 (M+H)	4	422.0445	C ₁₅ H ₁₉ O ₈ NBr	0.22	
	MS²	FI at <i>m/z</i> 150.0674 FI at <i>m/z</i> 213.9623 FI at <i>m/z</i> 228.9858 FI at <i>m/z</i> 246.0126	24 43 100 28	150.0681 213.9629 228.9864 246.0124	C ₈ H ₁₀ O ₂ C ₈ H ₇ O ₂ Br C ₈ H ₁₀ O ₃ Br C ₉ H ₁₃ O ₂ NBr	-3.86 -3.00 -2.69 0.75	
B8 S	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) sulfate isomer 1						6.0
	MS¹	PM at <i>m/z</i> 432.0113 (M+H)	2	432.0111	C ₁₆ H ₁₉ O ₆ NBrS	0.47	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 352.0541	61 100 2	91.0548 121.0653 352.0543	C ₇ H ₇ C ₈ H ₉ O C ₁₆ H ₁₉ O ₃ NBr	0 -2.81 -0.52	
B9/10 S	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) sulfate isomer 2						6.8
	MS¹	PM at <i>m/z</i> 432.0120 (M+H)	24	432.0111	C ₁₆ H ₁₉ O ₆ NBrS	2.09	
	MS²	FI at <i>m/z</i> 107.0496 FI at <i>m/z</i> 228.9860 FI at <i>m/z</i> 308.9428 FI at <i>m/z</i> 325.9696 FI at <i>m/z</i> 352.0546	99 100 34 38 5	107.0497 228.9864 308.9432 325.9692 352.0543	C ₇ H ₇ O C ₉ H ₁₀ O ₂ Br C ₉ H ₁₀ O ₃ BrS C ₉ H ₁₃ O ₃ NBrS C ₁₆ H ₁₉ O ₃ NBr	-0.84 -1.82 -1.40 1.12 0.90	
B13/14 S	25B-NBOMe-M (<i>O</i>-demethyl-) sulfate isomer 1						7.6
	MS¹	PM at <i>m/z</i> 446.0268 (M+H)	2	446.0267	C ₁₇ H ₂₁ O ₆ NBrS	0.12	
	MS²	FI at <i>m/z</i> 91.0549 FI at <i>m/z</i> 121.0651 FI at <i>m/z</i> 349.0431 FI at <i>m/z</i> 366.0703	60 100 1 4	91.0548 121.0653 349.0439 366.0699	C ₇ H ₇ C ₈ H ₉ O C ₁₇ H ₁₈ O ₃ Br C ₁₇ H ₂₁ O ₃ NBr	1.37 -1.98 -2.38 1.01	
B15 S	25B-NBOMe-M (<i>O</i>-demethyl-) sulfate isomer 2						8.5
	MS¹	PM at <i>m/z</i> 446.0264 (M+H)	3	446.0267	C ₁₇ H ₂₁ O ₆ NBrS	-0.78	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 243.0015 FI at <i>m/z</i> 260.0282 FI at <i>m/z</i> 366.0702	59 100 20 30	107.0497 243.0021 260.0281 366.0699	C ₇ H ₇ O C ₁₀ H ₁₂ O ₂ Br C ₁₀ H ₁₅ O ₂ NBr C ₁₇ H ₂₁ O ₃ NBr	-1.77 -2.33 0.51 0.73	
B16 S	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) sulfate isomer 1						4.8
	MS¹	PM at <i>m/z</i> 448.0061 (M+H)	5	448.0060	C ₁₆ H ₁₉ O ₇ NBrS	0.19	
	MS²	FI at <i>m/z</i> 107.0496 FI at <i>m/z</i> 137.0598 FI at <i>m/z</i> 217.0164 FI at <i>m/z</i> 368.0491	41 100 15 5	107.0497 137.0603 217.0171 368.0492	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₈ H ₉ O ₃ S C ₁₆ H ₁₉ O ₄ NBr	-0.84 -3.32 -3.10 -0.26	
B17/18 S	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) sulfate isomer 2						5.6
	MS¹	PM at <i>m/z</i> 448.0052 (M+H)	3	448.0060	C ₁₆ H ₁₉ O ₇ NBrS	-1.82	
	MS²	FI at <i>m/z</i> 123.0443 FI at <i>m/z</i> 228.9860 FI at <i>m/z</i> 308.9426 FI at <i>m/z</i> 325.9695 FI at <i>m/z</i> 368.0499	86 100 32 38 5	123.0446 228.9864 308.9432 325.9692 368.0492	C ₇ H ₇ O ₂ C ₉ H ₁₀ O ₂ Br C ₉ H ₁₀ O ₃ BrS C ₉ H ₁₃ O ₃ NBrS C ₁₆ H ₁₉ O ₄ NBr	-2.48 -1.82 -2.05 0.82 1.91	
B22 S	25B-NBOMe-M (<i>O</i>-demethyl-hydroxy-) sulfate isomer 1						6.2
	MS¹	PM at <i>m/z</i> 462.0212 (M+H)	2	462.0217	C ₁₇ H ₂₁ O ₇ NBrS	-1.00	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 258.9974 FI at <i>m/z</i> 382.0654	65 100 1 4	91.0548 121.0653 258.9970 382.0648	C ₇ H ₇ C ₈ H ₉ O C ₁₀ H ₁₂ O ₃ Br C ₁₇ H ₂₁ O ₄ NBr	1.37 -2.81 1.62 1.45	
B24/25 S	25B-NBOMe-M (<i>O</i>-demethyl-hydroxy-) sulfate isomer 2						7.2
	MS¹	PM at <i>m/z</i> 462.0215 (M+H)	9	462.0217	C ₁₇ H ₂₁ O ₇ NBrS	-0.35	
	MS²	FI at <i>m/z</i> 123.0442 FI at <i>m/z</i> 203.0007 FI at <i>m/z</i> 243.0014	58 9 100	123.0446 203.0014 243.0021	C ₇ H ₇ O ₂ C ₇ H ₇ O ₃ S C ₁₀ H ₁₂ O ₂ Br	-3.29 -3.56 -2.74	

		FI at <i>m/z</i> 260.0282 FI at <i>m/z</i> 382.0640	19 11	260.0281 382.0648	C ₁₀ H ₁₅ O ₂ NBr C ₁₇ H ₂₁ O ₄ NBr	0.51 -2.22	
B3/4 AC+G	25B-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl glucuronide						5.3
	MS¹	PM at <i>m/z</i> 464.0564 (M+H)	1	464.0551	C ₁₇ H ₂₃ O ₉ NBr	2.86	
	MS²	FI at <i>m/z</i> 150.0674 FI at <i>m/z</i> 228.9858 FI at <i>m/z</i> 246.0121 FI at <i>m/z</i> 288.0225	18 100 20 41	150.0681 228.9864 246.0124 288.0230	C ₉ H ₁₀ O ₂ C ₉ H ₁₀ O ₂ Br C ₉ H ₁₃ O ₂ NBr C ₁₁ H ₁₅ O ₃ NBr	-4.53 -2.69 -1.29 -1.67	
B8 GSH-2	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) acetylcysteine						6.4
	MS¹	PM at <i>m/z</i> 513.0695 (M+H)	10	513.0689	C ₂₁ H ₂₆ O ₆ N ₂ BrS	1.08	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 384.0264 FI at <i>m/z</i> 407.0273	60 100 6 1	91.0548 121.0653 384.0264 407.0271	C ₇ H ₇ C ₈ H ₉ O C ₁₆ H ₁₉ O ₃ NBrS C ₁₄ H ₂₀ O ₅ N ₂ BrS	0 -2.81 0 0.53	
B7 G	25B-NBOMe-M (<i>O,O,O</i>-tris-demethyl-) glucuronide						4.0
	MS¹	PM at <i>m/z</i> 514.0714 (M+H)	5	514.0707	C ₂₁ H ₂₅ O ₉ NBr	1.32	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 214.9701 FI at <i>m/z</i> 338.0383 FI at <i>m/z</i> 408.0289	71 100 24 10	107.0497 214.9708 338.0386 408.0289	C ₇ H ₇ O C ₈ H ₈ O ₂ Br C ₁₅ H ₁₇ O ₃ NBr C ₁₄ H ₁₉ O ₈ NBr	-1.77 -3.10 -0.98 0	
B8 G	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) glucuronide isomer 1						4.8
	MS¹	PM at <i>m/z</i> 528.0867 (M+H)	5	528.0864	C ₂₂ H ₂₇ O ₉ NBr	0.62	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 352.0544	47 100 16	91.0548 121.0653 352.0543	C ₇ H ₇ C ₈ H ₉ O C ₁₆ H ₁₉ O ₃ NBr	0 -2.81 0.34	
B9/10 G	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-) glucuronide isomer 2						5.9
	MS¹	PM at <i>m/z</i> 528.0872 (M+H)	5	528.0864	C ₂₂ H ₂₇ O ₉ NBr	1.57	
	MS²	FI at <i>m/z</i> 107.0496 FI at <i>m/z</i> 228.9860 FI at <i>m/z</i> 246.0127 FI at <i>m/z</i> 352.0548 FI at <i>m/z</i> 422.0448	60 100 32 26 7	107.0497 228.9864 246.0124 352.0543 422.0445	C ₇ H ₇ O C ₉ H ₁₀ O ₂ Br C ₉ H ₁₃ O ₂ NBr C ₁₆ H ₁₉ O ₃ NBr C ₁₅ H ₂₁ O ₈ NBr	-0.84 -1.82 1.15 1.47 0.70	
B13 G	25B-NBOMe-M (<i>O</i>-demethyl-) glucuronide isomer 1						5.6
	MS¹	PM at <i>m/z</i> 542.1024 (M+H)	9	542.1020	C ₂₃ H ₂₉ O ₉ NBr	0.70	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 228.9858 FI at <i>m/z</i> 366.0705	57 100 1 12	91.0548 121.0653 228.9864 366.0699	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₁₀ O ₂ Br C ₁₇ H ₂₁ O ₃ NBr	0 -2.81 -2.69 1.55	
B14 G	25B-NBOMe-M (<i>O</i>-demethyl-) glucuronide isomer 2						6.6
	MS¹	PM at <i>m/z</i> 542.1022 (M+H)	4	542.1020	C ₂₃ H ₂₉ O ₉ NBr	0.33	
	MS²	FI at <i>m/z</i> 91.0549 FI at <i>m/z</i> 121.0651 FI at <i>m/z</i> 228.9856 FI at <i>m/z</i> 366.0699	56 100 1 21	91.0548 121.0653 228.9864 366.0699	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₁₀ O ₂ Br C ₁₇ H ₂₁ O ₃ NBr	1.37 -1.98 -3.56 0	
B15 G	25B-NBOMe-M (<i>O</i>-demethyl-) glucuronide isomer 3						7.3
	MS¹	PM at <i>m/z</i> 542.1020 (M+H)	8	542.1020	C ₂₃ H ₂₉ O ₉ NBr	0	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 243.0015 FI at <i>m/z</i> 260.0282 FI at <i>m/z</i> 366.0704	100 45 14 21	107.0497 243.0021 260.0281 366.0699	C ₇ H ₇ O C ₁₀ H ₁₂ O ₂ Br C ₁₀ H ₁₅ O ₂ NBr C ₁₇ H ₂₁ O ₃ NBr	-1.77 -2.33 0.51 1.28	
B16 G	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) glucuronide isomer 1						4.1
	MS¹	PM at <i>m/z</i> 544.0806 (M+H)	4	544.0813	C ₂₂ H ₂₇ O ₁₀ NBr	-1.26	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 137.0597 FI at <i>m/z</i> 368.0493	36 100 12	107.0497 137.0603 368.0492	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₁₆ H ₁₉ O ₄ NBr	-1.77 -4.05 0.28	
B19 G	25B-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) glucuronide isomer 2						5.3
	MS¹	PM at <i>m/z</i> 544.0817 (M+H)	4	544.0813	C ₂₂ H ₂₇ O ₁₀ NBr	0.76	

	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 244.9808 FI at <i>m/z</i> 262.0074 FI at <i>m/z</i> 368.0493 FI at <i>m/z</i> 438.0394	52 100 35 31 3	107.0497 244.9813 262.0073 368.0492 438.0394	C ₇ H ₇ O C ₉ H ₁₀ O ₃ Br C ₉ H ₁₃ O ₃ NBr C ₁₆ H ₁₉ O ₄ NBr C ₁₅ H ₂₁ O ₉ NBr	-1.77 -2.17 0.26 -0.28 0	
B23 G	25B-NBOMe-M (O-demethyl-hydroxy-) glucuronide isomer 1						5.7
	MS¹	PM at <i>m/z</i> 558.0969 (M+H)	4	558.0969	C ₂₃ H ₂₉ O ₁₀ NBr	0	
	MS²	FI at <i>m/z</i> 107.0495 FI at <i>m/z</i> 137.0597 FI at <i>m/z</i> 228.9856 FI at <i>m/z</i> 382.0645	36 100 1 18	107.0497 137.0603 228.9864 382.0648	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₉ H ₁₀ O ₃ Br C ₁₇ H ₂₁ O ₄ NBr	-1.77 -4.05 -2.25 -0.38	
B22 G	25B-NBOMe-M (O-demethyl-hydroxy-) glucuronide isomer 2						5.9
	MS¹	PM at <i>m/z</i> 558.0970 (M+H)	5	558.0969	C ₂₃ H ₂₉ O ₁₀ NBr	0.11	
	MS²	FI at <i>m/z</i> 91.0548 FI at <i>m/z</i> 121.0650 FI at <i>m/z</i> 382.0647	48 100 22	91.0548 121.0653 382.0648	C ₇ H ₇ C ₈ H ₉ O C ₁₇ H ₂₁ O ₄ NBr	0 -2.81 -0.38	
B24/25 G	25B-NBOMe-M (O-demethyl-hydroxy-) glucuronide isomer 3						6.0
	MS¹	PM at <i>m/z</i> 558.0968 (M+H)	8	558.0969	C ₂₃ H ₂₉ O ₁₀ NBr	-0.24	
	MS²	FI at <i>m/z</i> 123.0443 FI at <i>m/z</i> 243.0019 FI at <i>m/z</i> 260.0282 FI at <i>m/z</i> 299.0758 FI at <i>m/z</i> 382.0663	84 100 32 2 5	123.0446 243.0021 260.0281 299.0767 382.0648	C ₇ H ₇ O ₂ C ₁₀ H ₁₂ O ₂ Br C ₁₀ H ₁₅ O ₂ NBr C ₁₃ H ₁₅ O ₈ C ₁₇ H ₂₁ O ₄ NBr	-2.48 -0.68 0.51 -2.99 3.80	
B30 G	25B-NBOMe-M (hydroxy-) glucuronide isomer 1						6.6
	MS¹	PM at <i>m/z</i> 572.1151 (M+H)	7	572.1126	C ₂₄ H ₃₁ O ₁₀ NBr	4.39	
	MS²	FI at <i>m/z</i> 109.0653 FI at <i>m/z</i> 137.0598 FI at <i>m/z</i> 313.0920 FI at <i>m/z</i> 396.0792	100 37 30 5	109.0653 137.0603 313.0923 396.0805	C ₇ H ₉ O C ₈ H ₉ O ₂ C ₁₄ H ₁₇ O ₈ C ₁₈ H ₂₃ O ₄ NBr	-0.37 -3.32 -1.10 -3.27	
B31 G	25B-NBOMe-M (hydroxy-) glucuronide isomer 2						6.9
	MS¹	PM at <i>m/z</i> 572.1134 (M+H)	4	572.1126	C ₂₄ H ₃₁ O ₁₀ NBr	1.42	
	MS²	FI at <i>m/z</i> 107.0496 FI at <i>m/z</i> 137.0599 FI at <i>m/z</i> 313.0918 FI at <i>m/z</i> 396.0808	24 100 26 4	107.0497 137.0603 313.0923 396.0805	C ₇ H ₇ O C ₈ H ₉ O ₂ C ₁₄ H ₁₇ O ₈ C ₁₈ H ₂₃ O ₄ NBr	-0.84 -2.59 -1.74 0.77	

Table S4 List of all 25C-NBOMe phase II metabolites together with the precursor mass (PM) recorded in MS¹, the corresponding characteristic fragment ions (FI) in MS², the calculated exact masses, the corresponding elemental composition, the deviation of the measured from the calculated masses, given as errors in parts per million (ppm), and the retention times (RT) in minutes (min). The metabolites were sorted by mass and RT. Numbering according to Table 2 (AC = *N*-acetylation, GSH = glutathione conjugation, ME = *O*-methylation, G = glucuronidation, S = sulfation, AC+G = acetylation combined with glucuronidation, AC+S = acetylation combined with sulfation)

No.	Metabolite and characteristic ions Measured accurate mass, <i>m/z</i>	Relative intensity in MS ² , %	Calculated exact mass, <i>m/z</i>	Elemental composition	Error, ppm	RT, min
C2 AC	25C-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl isomer 1					6.3
	MS¹ PM at <i>m/z</i> 244.0736 (M+H)	3	244.0735	C ₁₁ H ₁₅ O ₃ NCl	0.42	
	MS² FI at <i>m/z</i> 150.0676	25	150.0681	C ₉ H ₁₀ O ₂	-3.20	
	FI at <i>m/z</i> 170.0130	30	170.0135	C ₈ H ₇ O ₂ Cl	-2.69	
	FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
	FI at <i>m/z</i> 202.0630	9	202.0629	C ₉ H ₁₃ O ₂ NCl	0.33	
C3 AC	25C-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl isomer 2					6.5
	MS¹ PM at <i>m/z</i> 244.0732 (M+H)	4	244.0735	C ₁₁ H ₁₅ O ₃ NCl	-1.22	
	MS² FI at <i>m/z</i> 150.0676	12	150.0681	C ₉ H ₁₀ O ₂	-3.20	
	FI at <i>m/z</i> 170.0129	48	170.0135	C ₈ H ₇ O ₂ Cl	-3.28	
	FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
	FI at <i>m/z</i> 202.0629	11	202.0629	C ₉ H ₁₃ O ₂ NCl	0	
C7 AC	25C-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-hydroxy-) <i>N</i>-acetyl					5.1
	MS¹ PM at <i>m/z</i> 260.0681 (M+H)	4	260.0684	C ₁₁ H ₁₅ O ₄ NCl	-1.21	
	MS² FI at <i>m/z</i> 166.0625	7	166.0630	C ₉ H ₁₀ O ₃	-2.98	
	FI at <i>m/z</i> 186.0078	82	186.0084	C ₈ H ₇ O ₃ Cl	-3.08	
	FI at <i>m/z</i> 201.0312	100	201.0318	C ₉ H ₁₀ O ₃ Cl	-3.22	
	FI at <i>m/z</i> 218.0577	14	218.0578	C ₉ H ₁₃ O ₃ NCl	-0.68	
C3/4 AC+S	25C-NBOMe-M (<i>N</i>-demethoxybenzyl-<i>O</i>-demethyl-) <i>N</i>-acetyl sulfate					5.1
	MS¹ PM at <i>m/z</i> 324.0305 (M+H)	4	324.0303	C ₁₁ H ₁₅ O ₆ NCIS	0.57	
	MS² FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
	FI at <i>m/z</i> 202.0628	8	202.0629	C ₉ H ₁₃ O ₂ NCl	-0.66	
	FI at <i>m/z</i> 244.0736	7	244.0735	C ₁₁ H ₁₅ O ₃ NCl	0.42	
	FI at <i>m/z</i> 264.9933	26	264.9938	C ₉ H ₁₀ O ₃ ClS	-1.70	
	FI at <i>m/z</i> 282.0199	18	282.0198	C ₉ H ₁₃ O ₅ NCIS	0.53	
C10 GSH-1	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-) <i>S</i>-methyl					7.5
	MS¹ PM at <i>m/z</i> 354.0922 (M+H)	6	354.0925	C ₁₇ H ₂₃ O ₃ NCIS	-0.91	
	MS² FI at <i>m/z</i> 91.0547	55	91.0548	C ₇ H ₇	-0.82	
	FI at <i>m/z</i> 121.0650	100	121.0653	C ₈ H ₉ O	-2.81	
	FI at <i>m/z</i> 232.0197	0.5	232.0194	C ₉ H ₁₁ O ₂ NCIS	1.48	
	FI at <i>m/z</i> 337.0650	0.5	337.0665	C ₁₇ H ₁₈ O ₃ ClS	-4.51	
C34 ME	25C-NBOMe-M (<i>O</i>-demethyl-bis-hydroxy-) <i>O</i>-methyl					6.5
	MS¹ PM at <i>m/z</i> 368.1250 (M+H)	1	368.1259	C ₁₈ H ₂₅ O ₅ NCl	-2.52	
	MS² FI at <i>m/z</i> 137.0597	40	137.0603	C ₈ H ₉ O ₂	-4.05	
	FI at <i>m/z</i> 167.0704	100	167.0708	C ₉ H ₁₁ O ₃	-2.51	
	FI at <i>m/z</i> 185.0362	14	185.0369	C ₉ H ₁₀ O ₂ Cl	-3.96	
	FI at <i>m/z</i> 202.0631	2	202.0629	C ₉ H ₁₃ O ₂ NCl	0.82	
C8 S	25C-NBOMe-M (<i>O,O,O</i>-tris-demethyl-) sulfate					4.5

	MS¹	PM at <i>m/z</i> 374.0463 (M+H)	7	374.0460	C ₁₃ H ₁₇ O ₆ NCIS	0.89	
	MS²	FI at <i>m/z</i> 107.0496	100	107.0497	C ₇ H ₇ O	-0.84	
		FI at <i>m/z</i> 171.0210	78	171.0213	C ₈ H ₈ O ₂ Cl	-1.65	
		FI at <i>m/z</i> 250.9778	21	250.9781	C ₈ H ₈ O ₃ ClS	-1.20	
		FI at <i>m/z</i> 268.0043	26	268.0041	C ₈ H ₁₁ O ₅ NCIS	0.74	
		FI at <i>m/z</i> 294.0894	10	294.0891	C ₁₃ H ₁₇ O ₃ NCI	0.85	
C2/3 G	25C-NBOMe-M (N-demethoxybenzyl-O-demethyl-) glucuronide						2.6
	MS¹	PM at <i>m/z</i> 378.0949 (M+H)	5	378.0950	C ₁₃ H ₂₁ O ₈ NCI	-0.33	
	MS²	FI at <i>m/z</i> 150.0675	9	150.0681	C ₉ H ₁₀ O ₂	-3.86	
		FI at <i>m/z</i> 170.0130	35	170.0135	C ₈ H ₇ O ₂ Cl	-2.69	
		FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
		FI at <i>m/z</i> 202.0628	24	202.0629	C ₉ H ₁₃ O ₂ NCI	-0.66	
C36 ME	25C-NBOMe-M (bis-hydroxy-) O-methyl						7.6
	MS¹	PM at <i>m/z</i> 382.1410 (M+H)	1	382.1416	C ₁₀ H ₂₅ O ₃ NCI	-1.51	
	MS²	FI at <i>m/z</i> 137.0596	39	137.0603	C ₈ H ₉ O ₂	-4.78	
		FI at <i>m/z</i> 167.0702	100	167.0708	C ₉ H ₁₁ O ₃	-3.71	
		FI at <i>m/z</i> 199.0519	22	199.0526	C ₁₀ H ₁₂ O ₂ Cl	-3.43	
		FI at <i>m/z</i> 216.0785	6	216.0786	C ₁₀ H ₁₅ O ₂ NCI	-0.39	
C10 S	25C-NBOMe-M (O,O-bis-demethyl-) sulfate isomer 1						5.8
	MS¹	PM at <i>m/z</i> 388.0614 (M+H)	2	388.0616	C ₁₆ H ₁₉ O ₆ NCIS	-0.56	
	MS²	FI at <i>m/z</i> 91.0548	54	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0650	100	121.0653	C ₈ H ₉ O	-2.81	
		FI at <i>m/z</i> 308.1046	2	308.1048	C ₁₆ H ₁₉ O ₃ NCI	-0.64	
C11 S	25C-NBOMe-M (O,O-bis-demethyl-) sulfate isomer 2						6.6
	MS¹	PM at <i>m/z</i> 388.0617 (M+H)	6	388.0616	C ₁₆ H ₁₉ O ₆ NCIS	0.22	
	MS²	FI at <i>m/z</i> 107.0496	96	107.0497	C ₇ H ₇ O	-0.84	
		FI at <i>m/z</i> 185.0366	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-1.80	
		FI at <i>m/z</i> 264.9934	40	264.9938	C ₉ H ₁₀ O ₅ ClS	-1.32	
		FI at <i>m/z</i> 282.0200	42	282.0198	C ₉ H ₁₃ O ₃ NCIS	0.88	
		FI at <i>m/z</i> 308.1047	6	308.1048	C ₁₆ H ₁₉ O ₃ NCI	-0.32	
C12 S	25C-NBOMe-M (O,O-bis-demethyl-) sulfate isomer 3						7.0
	MS¹	PM at <i>m/z</i> 388.0619 (M+H)	11	388.0616	C ₁₆ H ₁₉ O ₆ NCIS	0.73	
	MS²	FI at <i>m/z</i> 107.0496	100	107.0497	C ₇ H ₇ O	-0.84	
		FI at <i>m/z</i> 185.0366	99	185.0369	C ₉ H ₁₀ O ₂ Cl	-1.80	
		FI at <i>m/z</i> 187.0060	6	187.0065	C ₇ H ₇ O ₄ S	-2.71	
		FI at <i>m/z</i> 202.0629	28	202.0629	C ₉ H ₁₃ O ₅ NCIS	0	
		FI at <i>m/z</i> 308.1048	37	308.1048	C ₁₆ H ₁₉ O ₃ NCI	0	
C16/17 S	25C-NBOMe-M (O-demethyl-) sulfate isomer 1						7.4
	MS¹	PM at <i>m/z</i> 402.0778 (M+H)	3	402.0773	C ₁₇ H ₂₁ O ₆ NCIS	1.33	
	MS²	FI at <i>m/z</i> 91.0548	61	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0651	100	121.0653	C ₈ H ₉ O	-1.98	
		FI at <i>m/z</i> 305.0938	1	305.0944	C ₁₇ H ₁₈ O ₃ Cl	-2.12	
		FI at <i>m/z</i> 322.1205	4	322.1204	C ₁₇ H ₂₁ O ₃ NCI	0.16	
C18 S	25C-NBOMe-M (O-demethyl-) sulfate isomer 2						8.2
	MS¹	PM at <i>m/z</i> 402.0785 (M+H)	5	402.0773	C ₁₇ H ₂₁ O ₆ NCIS	3.07	
	MS²	FI at <i>m/z</i> 107.0495	59	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 199.0521	100	199.0526	C ₁₀ H ₁₂ O ₂ Cl	-2.43	
		FI at <i>m/z</i> 216.0787	19	216.0786	C ₁₀ H ₁₅ O ₂ NCI	0.54	
		FI at <i>m/z</i> 322.1205	32	322.1204	C ₁₇ H ₂₁ O ₃ NCI	0.16	
C20 S	25C-NBOMe-M (O,O-bis-demethyl-hydroxy-) sulfate						5.0
	MS¹	PM at <i>m/z</i> 404.0569 (M+H)	9	404.0565	C ₁₆ H ₁₉ O ₇ NCIS	0.91	
	MS²	FI at <i>m/z</i> 123.0443	80	123.0446	C ₇ H ₇ O ₂	-2.48	
		FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
		FI at <i>m/z</i> 264.9934	26	264.9938	C ₉ H ₁₀ O ₅ ClS	-1.32	
		FI at <i>m/z</i> 282.0200	22	282.0198	C ₉ H ₁₃ O ₅ NCIS	0.88	
		FI at <i>m/z</i> 324.0998	8	324.0997	C ₁₆ H ₁₉ O ₄ NCI	0.27	
C22 S	25C-NBOMe-M (O-demethyl-hydroxy-) sulfate						6.9
	MS¹	PM at <i>m/z</i> 418.0726 (M+H)	12	418.0722	C ₁₇ H ₂₁ O ₇ NCIS	1.00	
	MS²	FI at <i>m/z</i> 123.0443	55	123.0446	C ₇ H ₇ O ₂	-2.48	

		FI at m/z 199.0522 FI at m/z 203.0008 FI at m/z 216.0788 FI at m/z 338.1154	100 10 17 11	199.0526 203.0014 216.0786 338.1154	C ₁₀ H ₁₂ O ₂ Cl C ₇ H ₇ O ₂ S C ₁₀ H ₁₅ O ₂ NCl C ₁₇ H ₂₁ O ₄ NCl	-1.92 -3.06 1.00 0	
C2/3 AC+G		25C-NBOMe-M (N-demethoxybenzyl-O-demethyl-) N-acetyl glucuronide					5.0
	MS¹	PM at m/z 420.1060 (M+H)	1	420.1056	C ₁₇ H ₂₃ O ₉ NCl	0.98	
	MS²	FI at m/z 150.0675 FI at m/z 185.0365 FI at m/z 202.0628 FI at m/z 244.0736	6 100 19 36	150.0681 185.0369 202.0629 244.0735	C ₉ H ₁₀ O ₂ C ₉ H ₁₀ O ₂ Cl C ₉ H ₁₃ O ₂ NCl C ₁₁ H ₁₅ O ₃ NCl	-3.86 -2.34 -0.66 0.42	
C10 GSH-2		25C-NBOMe-M (O,O-bis-demethyl-) acetylcysteine					6.2
	MS¹	PM at m/z 469.1200 (M+H)	14	469.1195	C ₂₁ H ₂₆ O ₆ N ₂ ClS	1.14	
	MS²	FI at m/z 91.0548 FI at m/z 121.0650 FI at m/z 340.0770 FI at m/z 363.0781	55 100 6 0.5	91.0548 121.0653 340.0769 363.0776	C ₇ H ₇ C ₈ H ₉ O C ₁₆ H ₁₉ O ₃ NClS C ₁₄ H ₂₀ O ₅ N ₂ ClS	0 -2.81 0.38 1.38	
C8 G		25C-NBOMe-M (O,O,O-tris-demethyl-) glucuronide					3.8
	MS¹	PM at m/z 470.1219 (M+H)	8	470.1212	C ₂₁ H ₂₅ O ₉ NCl	1.41	
	MS²	FI at m/z 107.0495 FI at m/z 171.0208 FI at m/z 294.0891 FI at m/z 364.0796	69 100 28 12	107.0497 171.0213 294.0891 364.0794	C ₇ H ₇ O C ₈ H ₈ O ₂ Cl C ₁₅ H ₁₇ O ₃ NCl C ₁₄ H ₁₉ O ₈ NCl	-1.77 -2.82 0 0.62	
C10 G		25C-NBOMe-M (O,O-bis-demethyl-) glucuronide isomer 1					4.6
	MS¹	PM at m/z 484.1377 (M+H)	10	484.1369	C ₂₂ H ₂₇ O ₉ NCl	1.68	
	MS²	FI at m/z 91.0548 FI at m/z 121.0651 FI at m/z 308.1047	41 100 18	91.0548 121.0653 308.1048	C ₇ H ₇ C ₈ H ₉ O C ₁₆ H ₁₉ O ₃ NCl	0 -1.98 -0.32	
C11 G		25C-NBOMe-M (O,O-bis-demethyl-) glucuronide isomer 2					5.5
	MS¹	PM at m/z 484.1375 (M+H)	23	484.1369	C ₂₂ H ₂₇ O ₉ NCl	1.26	
	MS²	FI at m/z 107.0495 FI at m/z 185.0365 FI at m/z 202.0629 FI at m/z 308.1046 FI at m/z 378.0953	87 100 61 28 30	107.0497 185.0369 202.0629 308.1048 378.0950	C ₇ H ₇ O C ₉ H ₁₀ O ₂ Cl C ₉ H ₁₃ O ₂ NCl C ₁₆ H ₁₉ O ₃ NCl C ₁₅ H ₂₁ O ₈ NCl	-1.77 -2.34 0 -0.64 0.73	
C12 G		25C-NBOMe-M (O,O-bis-demethyl-) glucuronide isomer 3					5.6
	MS¹	PM at m/z 484.1375 (M+H)	8	484.1369	C ₂₂ H ₂₇ O ₉ NCl	1.26	
	MS²	FI at m/z 107.0495 FI at m/z 185.0365 FI at m/z 202.0628 FI at m/z 308.1047 FI at m/z 378.0952	55 100 30 29 9	107.0497 185.0369 202.0629 308.1048 378.0950	C ₇ H ₇ O C ₉ H ₁₀ O ₂ Cl C ₉ H ₁₃ O ₂ NCl C ₁₆ H ₁₉ O ₃ NCl C ₁₅ H ₂₁ O ₈ NCl	-1.77 -2.34 -0.66 -0.32 0.47	
C16 G		25C-NBOMe-M (O-demethyl-) glucuronide isomer 1					5.5
	MS¹	PM at m/z 498.1526 (M+H)	15	498.1525	C ₂₃ H ₂₉ O ₉ NCl	0.12	
	MS²	FI at m/z 91.0548 FI at m/z 121.0651 FI at m/z 185.0365 FI at m/z 322.1205	52 100 1 13	91.0548 121.0653 185.0369 322.1204	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₁₀ O ₂ Cl C ₁₇ H ₂₁ O ₃ NCl	0 -1.98 -2.34 0.16	
C17 G		25C-NBOMe-M (O-demethyl-) glucuronide isomer 2					6.4
	MS¹	PM at m/z 498.1524 (M+H)	8	498.1525	C ₂₃ H ₂₉ O ₉ NCl	-0.28	
	MS²	FI at m/z 91.0548 FI at m/z 121.0650 FI at m/z 185.0365 FI at m/z 322.1205	45 100 1 25	91.0548 121.0653 185.0369 322.1204	C ₇ H ₇ C ₈ H ₉ O C ₉ H ₁₀ O ₂ Cl C ₁₇ H ₂₁ O ₃ NCl	0 -2.81 2.34 0.16	
C18 G		25C-NBOMe-M (O-demethyl-) glucuronide isomer 3					7.0
	MS¹	PM at m/z 498.1525 (M+H)	14	498.1525	C ₂₃ H ₂₉ O ₉ NCl	0	
	MS²	FI at m/z 107.0496 FI at m/z 199.0522 FI at m/z 216.0788 FI at m/z 322.1206	100 46 14 26	107.0497 199.0526 216.0786 322.1204	C ₇ H ₇ O C ₁₀ H ₁₂ O ₂ Cl C ₁₀ H ₁₅ O ₂ NCl C ₁₇ H ₂₁ O ₃ NCl	-0.84 -1.92 1.00 0.47	

C19 G	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) glucuronide isomer 1						3.9
	MS¹	PM at <i>m/z</i> 500.1320 (M+H)	6	500.1318	C ₂₂ H ₂₇ O ₁₀ NCl	0.39	
	MS²	FI at <i>m/z</i> 107.0495	36	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 137.0598	100	137.0603	C ₈ H ₉ O ₂	-3.32	
		FI at <i>m/z</i> 324.0998	15	324.0997	C ₁₆ H ₁₉ O ₄ NCl	0.27	
C20 G	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) glucuronide isomer 2						4.5
	MS¹	PM at <i>m/z</i> 500.1318 (M+H)	12	500.1318	C ₂₂ H ₂₇ O ₁₀ NCl	0	
	MS²	FI at <i>m/z</i> 123.0442	89	123.0446	C ₇ H ₇ O ₂	-3.29	
		FI at <i>m/z</i> 185.0365	100	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
		FI at <i>m/z</i> 202.0628	32	202.0629	C ₉ H ₁₃ O ₂ NCl	-0.66	
		FI at <i>m/z</i> 299.0762	5	299.0767	C ₁₃ H ₁₅ O ₈	-1.66	
		FI at <i>m/z</i> 324.0997	12	324.0997	C ₁₆ H ₁₉ O ₄ NCl	0	
C21 G	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) glucuronide isomer 3						5.1
	MS¹	PM at <i>m/z</i> 500.1321 (M+H)	7	500.1318	C ₂₂ H ₂₇ O ₁₀ NCl	0.59	
	MS²	FI at <i>m/z</i> 107.0495	52	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 201.0313	100	201.0318	C ₉ H ₁₀ O ₃ Cl	-2.73	
		FI at <i>m/z</i> 218.0578	36	218.0578	C ₉ H ₁₃ O ₃ NCl	0	
		FI at <i>m/z</i> 324.0997	31	324.0997	C ₁₆ H ₁₉ O ₄ NCl	0	
		FI at <i>m/z</i> 394.0906	4	394.0899	C ₁₅ H ₂₁ O ₉ NCl	1.68	
C22 G	25C-NBOMe-M (<i>O,O</i>-bis-demethyl-hydroxy-) glucuronide isomer 4						5.3
	MS¹	PM at <i>m/z</i> 500.1315 (M+H)	7	500.1318	C ₂₂ H ₂₇ O ₁₀ NCl	-0.61	
	MS²	FI at <i>m/z</i> 123.0443	100	123.0446	C ₇ H ₇ O ₂	-2.48	
		FI at <i>m/z</i> 185.0365	25	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
		FI at <i>m/z</i> 202.0628	11	202.0629	C ₉ H ₁₃ O ₂ NCl	-0.66	
		FI at <i>m/z</i> 324.0996	9	324.0997	C ₁₆ H ₁₉ O ₄ NCl	-0.35	
		FI at <i>m/z</i> 378.0945	1	378.0950	C ₁₅ H ₂₁ O ₈ NCl	-1.38	
C27 G	25C-NBOMe-M (<i>O</i>-demethyl-hydroxy-) glucuronide isomer 1						5.5
	MS¹	PM at <i>m/z</i> 514.1483 (M+H)	6	514.1475	C ₂₃ H ₂₉ O ₁₀ NCl	1.65	
	MS²	FI at <i>m/z</i> 107.0495	33	107.0497	C ₇ H ₇ O	-1.77	
		FI at <i>m/z</i> 137.0597	100	137.0603	C ₈ H ₉ O ₂	-4.05	
		FI at <i>m/z</i> 185.0365	1	185.0369	C ₉ H ₁₀ O ₂ Cl	-2.34	
		FI at <i>m/z</i> 313.0918	4	313.0923	C ₁₄ H ₁₇ O ₈	-1.74	
		FI at <i>m/z</i> 338.1152	17	338.1154	C ₁₇ H ₂₁ O ₄ NCl	-0.48	
C28/29 G	25C-NBOMe-M (<i>O</i>-demethyl-hydroxy-) glucuronide isomer 2						6.5
	MS¹	PM at <i>m/z</i> 514.1479 (M+H)	8	514.1475	C ₂₃ H ₂₉ O ₁₀ NCl	0.87	
	MS²	FI at <i>m/z</i> 123.0443	100	123.0446	C ₇ H ₇ O ₂	-2.48	
		FI at <i>m/z</i> 199.0521	26	199.0526	C ₁₀ H ₁₂ O ₂ Cl	-2.43	
		FI at <i>m/z</i> 216.0787	11	216.0786	C ₁₀ H ₁₅ O ₂ NCl	0.54	
		FI at <i>m/z</i> 338.1152	10	338.1154	C ₁₇ H ₂₁ O ₄ NCl	-0.48	
C31/32 G	25C-NBOMe-M (hydroxy-) glucuronide isomer 1						5.2
	MS¹	PM at <i>m/z</i> 528.1640 (M+H)	7	528.1631	C ₂₄ H ₃₁ O ₁₀ NCl	1.70	
	MS²	FI at <i>m/z</i> 91.0548	46	91.0548	C ₇ H ₇	0	
		FI at <i>m/z</i> 121.0651	100	121.0653	C ₈ H ₉ O	-1.98	
		FI at <i>m/z</i> 352.1313	18	352.1310	C ₁₈ H ₂₃ O ₄ NCl	0.81	
C33 G	25C-NBOMe-M (hydroxy-) glucuronide isomer 2						6.3
	MS¹	PM at <i>m/z</i> 528.1638 (M+H)	7	528.1631	C ₂₄ H ₃₁ O ₁₀ NCl	1.32	
	MS²	FI at <i>m/z</i> 109.0652	73	109.0653	C ₇ H ₉ O	-1.28	
		FI at <i>m/z</i> 137.0598	100	137.0603	C ₈ H ₉ O ₂	-3.32	
		FI at <i>m/z</i> 313.0921	17	313.0923	C ₁₄ H ₁₇ O ₈	-0.78	
		FI at <i>m/z</i> 352.1324	3	352.1310	C ₁₈ H ₂₃ O ₄ NCl	3.94	