SIMPLE AND FAST DETERMINATION FOR GAMMA-HYDROXYBUTYRATE (GHB) IN URINE SAMPLE BY LLE AND GC-MS

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Received on 25/06/2020; accepted on 05/08/2020; published on the web on 21/09/2020

Gamma-hydroxybutyrate (GHB) is a depressant of the central nervous system able to promote disinhibition, anterograde amnesia and its use is related to Drug-Facilitated Crimes (DFC). Urine is the biological matrix of choice for toxicological analysis in victims of rape and/or theft in suspected victims of DFC. The aim of this work was the development of a simple and ultra-fast methodology for GHB determination in urine samples using GC-MS. Briefly, 250 µL of urine was pipetted in a vial and fortified with 10 µL (250 µg mL⁻¹) of internal standard GHB-d6. Then, the sample was acidified and submitted to extraction with ethyl acetate. After stirring, the material was taken to a microcentrifuge (1 minute at 5000 rpm). Then the dried residue was derivatized with BSTFA + 1% TCMS for 10 min at 70 ºC before injection. This method was successfully validated according UNODC and EMEA guidelines on bioanalytical method validation with limit of detection and limit of quantification were 0.25 and 0.50 µg mL⁻¹, respectively. Considering all steps for GHB determination, this analytical procedure allows the fastest detection of GHB in urine samples when compared to literature allowing the analysis of DFC suspect cases or clinical when endogenous GHB requires to be measured.

Keywords: GHB; GC-MS; urine; LLE; simple and fast.

INTRODUCTION

Gamma-hydroxybutyrate (GHB) is a relatively simple molecular structure, with a pKa = 4.72 and is derived from the neurotransmitter GABA naturally produced by mammals. Its mechanism of action is based on its binding to a specific GABA receptor, as well as altering dopaminergic reception.1,2 GHB absorption is fast when administered orally, reaching maximum plasma concentration between 25 and 45 minutes without binding plasma proteins. GHB distribution is fast and its effects last around 1 to 4 hours depending on the dose.3 Multiple doses of GHB does not promote accumulation in the body and endogenous metabolites are excreted after metabolization. A small amount of unchanged GHB (1-5% from administrated dose) is eliminated by urine, which it can be determined within 12 hours. After this period, GHB levels in urinary samples returns to endogenous baseline.1,3,4 Disinhibition and increased sociability, anterograde amnesia and even increased libido are described as GHB effects.3,5,6 Beginning in the 1990s, GHB and its precursors (gamma-butyrolactone and 1,4-butanediol) began to be used in situations involving sexual abuse and other crimes due to the relatively easy obtaining and manufacture in any laboratory. Also, the possibility of reducing time-consuming sample processing steps, decreasing the number of reagents, and increasing the analytic speed and efficiency is always welcome in any forensic toxicology laboratory. So, we purpose a simple and fast methodology for GHB determination in urine samples using GC-MS to be applied in any laboratory.

EXPERIMENTAL

Material and methods

Reagents and standards

HPLC grade acetonitrile (ACN), methanol (MeOH), ethyl acetate (EtOAc), chloroform (CHCl₃), dichloromethane (DCM) and isobutanol (i-BuOH) were obtained from Vetec® (Duque de Caxias, RJ, Brazil). Hydrochloric acid (HCl), sodium chloride (NaCl) and anhydrous sodium sulfate (Na₂SO₄) were purchased from Synth® (Diadema, SP, Brazil). GHB and GHB-D6 were purchased from Cerilliant® (Round Rock, TX, USA). N,O-Bis(trimethylsilyl) trifluoroacetamide with trimethylchlorosilane (TCMS) (BSTFA + 1% TCMS) was purchased from Sigma–Aldrich® (St.Louis, MO, USA). Blank urine was obtained from volunteers who did not use the test substance and were collected from volunteers admitted to the Vila Serena treatment center for chemical dependence (São Paulo, SP, Brazil). Stock solutions were prepared with MeOH and stored at -21 ºC. Calibration and control solutions were prepared by successive dilutions of the stock standard solutions in methanol. Aliquots of calibration solutions were stored at -20 ºC until use.

Instrumentation

The analyzes were performed using an Agilent 6850 Network GC System gas chromatograph coupled with an Agilent® 5975 Series quadrupole mass selective detector (MSD). The injections were done

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in splitless mode and the chromatographic separation was achieved with a capillary fused silica HP-5MS column (30 m x 0.25 mm x 0.1 film thickness) (Agilent, Santa Clara, CA, USA), using helium as a carrier gas at 1.0 mL min⁻¹, in a constant flow mode. The column oven temperature program was as follows: first hold at 50 °C for 1 minute; increase the temperature at a rate of 20 °C min⁻¹ up to 120 °C (hold 1 minute); then at a rate of 30 °C min⁻¹ up to 300 °C. The total analysis time was 11.50 min. The temperatures at the injection port and transfer line were 280 °C. The analysis of the analytes by the MS was through electronic ionization (EI) (70 eV) with selected ion monitoring (SIM) mode. The monitored ions were 233 (quantifier ion), 204 and 117 m/z; 239 (quantifier ion), 210 and 123 m/z for GHB and GHB-D6, respectively.

Method development for determining GHB in urine samples

Optimization of GHB derivatization

The GHB derivatization optimization study was carried out considering the incubation time and temperature using BSTFA + 1% TCMS as a derivatizing agent. The efficiency of the reaction was assessed through the total area under the conditions tested. The following parameters were studied: Derivatization in GC liner port (injection-port derivatization); Temperature (70, 80 and 90 °C); Time (5, 10, 20 and 30 minutes).

Optimization of liquid-liquid extraction (LLE) for GHB

Urine samples were fortified to a concentration of 10 µg mL⁻¹ of GHB and it was evaluated extractor solvent: ethyl acetate, acetonitrile, iso-butyl alcohol, chloroform and a mixture of methanol:dichloromethane (20:80); pH of the biological matrix: 0, 1, 2, 3, 4 and 5; Extraction time: 10, 30, 60 and 600 seconds; Agitation: 1200, 1600, 2000 and 2400 rpm; Salting out effect: 0, 0.5 and 10% NaCl (m v⁻¹) to the sample before the extraction procedure.

Sample preparation

A 250 µL aliquot of urine was placed in a vial and fortified with 10 µL (250 µg mL⁻¹) of internal standard GHB-D6. Sample was acidified with a solution of 3 mol L⁻¹ of HCl until pH 2 and then 500 µL of ethyl acetate (organic phase) was added. The microtube was closed and stirred for 10 seconds at 2400 rpm in a vortex. After stirring, the material was taken to a microcentrifuge (1 minute at 5000 rpm). Then 350 µL of organic phase were removed and transferred to bottles containing anhydrous sodium sulfate. The supernatant resulting from this step was placed in a road with silanol groups deactivated for drying at 40 °C under nitrogenous flow. Posteriorly, 35 µL of BSTFA + 1% TCMS was added following vortex agitation and posterior incubation during 10 minutes at 70 °C. Subsequently, the vials were cooled to room temperature and 2 µL aliquot was injected into the GC-MS. All steps for this procedure are showed in the Figure 1.

Method validation of GHB

The method was validated in urine samples according to Guidance for the Validation of Analytical Methodology and Calibration of Equipment used for Testing of Illicit Drugs in Seized Materials and Biological Specimens from United Nation Office on Drugs and Crime (UNODC) and Guideline on bioanalytical method validation from European Medicine Agency (EMEA).

For the limit of detection (LOD) and limit of quantification (LOQ), a pool of urine samples from 20 individuals was used as blank sample. From this pool, different concentrations of GHB standard were evaluated in triplicates by fortification with known amounts of analytical standard, which chromatographic peaks were discounted from baseline endogenous GHB present. After obtaining the result of the triplicate, the same procedure was performed in sextuplicate to guarantee the LOD value. The response equivalent to three times signal:noise defined this parameter. The acceptance criteria for LOQ was the response within ±20% of the expected value at the lower concentration. Linearity was assessed in six replicates with addition of GHB standard in blank urine samples, which the first point was LOQ (0.5; 1.0; 2.5; 5; 10 and 20 µg mL⁻¹).

Selectivity/specificity was assessed using 13 potential interfering compounds (11-nor-9-carboxy-tetrahydrocannabinol, ethanol, paracetamol, acetylsalicylic acid, methylmalonic acid, succinic acid, beta-estradiol, caffeine, lorazepam, phenobarbital, benzocaine,

Figure 1. Step-by-step of purpose methodology for GHB determination in urine samples
phenylpropanolamine, carisoprodol) at a concentration of 1000 ng mL\(^{-1}\), evaluating if these molecules can interfere with the GHB analysis following the extraction procedure developed in the present work. The acceptance criteria for this assay was the absence of interfering substances in the retention time of the analyte of interest and its respective internal standard.

Precision and accuracy of the studies were carried out by analyzing urine samples containing three known concentrations (1.5, 9.0 and 16 µg mL\(^{-1}\)) on three consecutive days. The acceptance criteria for these assays was coefficient variation for the lower concentration controls must be lower than 20% and for the other control levels better than 15%. Absolute recovery studies were performed by preparing two sets of urine samples. A set of samples (set A), which consists of three concentrations, where the first point with a concentration 3 times greater than the LOQ of the analyte (1.5, 9 and 16 µg mL\(^{-1}\)). In the second set (set B), analytes were added to the samples with the same concentration used in the set A after the LLE procedure.

Absolute recovery was assessed by comparing the average response obtained for set A and the response for set B (100%). UNODC Analytical Methods Validation Guide does not consider this factor as indicative for failing the test, if the other validation parameters achieve the desired purposes.\(^{10}\)

| Dilution integrity | The addition of NaCl (salting out effect) and the frequency of agitation (rpm) do not influence the GHB extraction by LLE in urine samples, however the agitation time is essential. Although 1 and 10 minutes showed better results, we opted for a quick procedure (10 seconds) for the validation of the method, considering that the data indicated a low coefficient of variation with absolute area. The work focuses on the determination of GHB in urine quickly and simply.

| Application to real cases | All parameters of the method validation are in accordance with the UNODC guide.\(^{10}\) Linearity showed an equation of \(y = 0.1144X - 0.1247\) (\(R^2 = 0.9932\)) and the heteroscedasticity phenomenon was not observed through the F test (Snedecor F distribution). No chromatographic interference was verified with the tested substances, demonstrating the specificity/selectivity of the proposed methodology. However, these results were only possible when SIM mode was applied, since countless molecules with similar characteristics to GHB are excreted in the urine samples and

RESULTS AND DISCUSSION

GHB determination is considered a forensic challenge because it is an endogenous compound of low molecular weight, fast excretion and which has a structural similarity with other organic acids eliminated in the urine. In addition, the analysis of GHB by GC-MS is difficult due to the high polarity of this molecule. So, there is a need for the derivatization step to determine GHB by GC-MS.\(^{12}\) Previously published papers opted for the conversion of GHB to GBL for analysis in GC-MS due to the possible interconversion in vitro,\(^{13-15}\) but GBL is still small sized and can be confused with other similar structures.

In this respect, derivatization with BSTFA + 1% TCMS is considered the best option due to its low cost, description of the derivatized in mass spectrometry libraries and handling facilities when compared to other derivatives.\(^{16,17}\) Derivatization is often carried out off-line after extraction. However, the possibility of performing this derivatization on-line could be a way of optimizing the time-consumption of the method. To develop an ultra-fast methodology by GC-MS, derivatization in GC liner port (injection-port derivatization) was carried out after drying the extract.\(^{18}\)

In this procedure, the derivatization occurred in the gas phase, where the sample and the derivatization reagent were injected together into the liner port. So, before the splitless injection 35 µL of BSTFA + 1% TCMS were added in the dried extract followed by vortexing for 30 seconds. After this 2 µL were injected into the liner port at 280 °C, where the derivatization step occurred. Initial results were promising with this approach. However, in a sequence of chromatographic injections, matrix interferers became more evident in the chromatograms, difficulting toxicological analysis and decreasing the useful life of the liner and the chromatographic column. So, incubation time and temperature were evaluated. The results indicated that 70°C for 10 minutes is the best derivatization process for this silylant agent, indicating a shorter time than established by the literature.\(^{16,17,19-27}\)

Selected ion monitoring mode (SIM mode) was chosen because of an increase of selectivity and decrease of the instrument’s response to anything that interferes with the measurement, allowing the determination of a component in a complex chromatogram where the compounds do not resolve.\(^{24}\) Thus, SIM mode was chosen, considering there are many compounds with similar retention time as well as fragmentation pattern compared to GHB in urine samples. In fact, it is impractical to perform a GHB analysis on urinary matrix using GC-MS in SCAN mode.

For optimization of the GHB in urine samples by LLE, we evaluated polar solvents considering partition coefficient (Log P = -0.51) and ethyl acetate (Log P = 0.71) obtained better results, as already demonstrated in literature.\(^{25}\) Isobutyl alcohol (Log P = 0.8) and especially acetonitrile (Log P = -33) can form 2 phases in the urine samples. However, addition of GHB standard diluted with methanol in urinary matrices, even in very low volumes, promotes a system homogeneous, not allowing the separation of phases and consequently the extraction. An alternative procedure was the use of the mixture methanol: dichloromethane (20:80), however the results showed inefficiency of this mixture.

Similarly, different mixtures of ethyl acetate and acetonitrile were evaluated and there were no differences between the combinations 9:1 and 8:2 when compared with extracts exclusively from ethyl acetate. The composition ethyl acetate: acetonitrile (7:3) showed greater extractive capacity. However, the appearance of interfering matrix did not allow the continuation of this procedure. So, the chosen extractor solvent was ethyl acetate.

GHB has acidic characteristics (pKa = 4.72) indicating that its extraction must be carried out in an acidic environment. The pH range used (0-5) demonstrated that pH 2 was the best for the procedure, indicating that the molecule will be in the non-ionized form. When pH values are less than 2 conversion of GHB to GBL may occur through dehydration and consequently closing the aliphatic structure in cyclic, even in the absence of temperature.\(^{30,31}\) At a pH greater than 2, this molecule has greater ionized fractions,\(^{32}\) decreasing the extractive capacity of the solvent by LLE.

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can interfere in the analysis. The other results of the validation are described in Table 1.

Table 1. Confidence parameters of the validated method for the determination of GHB in urine samples (sextuplicate for each point)

| Confidence parameters   | GHB            |
|------------------------|----------------|
| Absolute recovery (%)  |                |
| C1                     | 23.9           |
| C2                     | 20.3           |
| C3                     | 20.0           |
| LOD (µg mL⁻¹)          | 0.25           |
| LOQ (µg mL⁻¹)          | 0.50           |
| Intra-day precision (CV%) |            |
| C1                     | 13.7           |
| C2                     | 8.60           |
| C3                     | 9.80           |
| Inter-day precision (CV%) |            |
| C1                     | 9.90           |
| C2                     | 11.6           |
| C3                     | 9.40           |
| Accuracy (%)           |                |
| C1                     | 93.2           |
| C2                     | 108.3          |
| C3                     | 111.9          |
| Dilution integrity     |                |
| Precision (CV%)        |                |
| 10 times               | 6.00           |
| Accuracy (%)           |                |
| 10 times               | 101.1          |

Due to the possibility of obtaining urine samples with high concentrations of GHB, dilution integrity was validated according to Guideline on bioanalytical method validation from European Medicine Agency.¹¹ GHB standard was added in urine samples to a concentration of 50 µg mL⁻¹ and the matrix was diluted 1/10 with distilled water to obtain a final concentration of 5 µg mL⁻¹. Subsequently, the extraction process was carried out with a calibration curve. The result obtained was 5.05 µg mL⁻¹ and the CV% was less than 15%. Figure 2 shows a typical chromatogram using SCAN and SIM mode obtained by the purpose methodology.

Unfortunately, it was not possible to apply this method in real samples from individuals until they had contact with GHB because no suspected cases reached our group as well as the São Paulo Scientific Police at the period which the methodology was developed. Therefore, it must be considered that GHB showed pharmacological effects like other central depressants drugs and rapid excretion (less than 12 hours in urine samples). These aspects could be responsible for the absence of real cases during the development of this method.³³

According to the literature, the methodology described by Busardò et al.³³ is considered the fastest to analyze this molecule in biological fluids by GC-MS. This method is considered ultra-fast by the author, with time previously stipulated for agitation and centrifugation (5 minutes), derivatization (5 minutes) and posterior chromatographic analysis (16 minutes). However, the developed procedure in the present work used only 10 seconds of agitation followed by 1 minute of centrifugation and 10 minutes of reaction with BSTFA + 1% TMCS. The complete chromatographic runtime was 13 minutes, which is faster when compared to others in the literature with a full validation. LOQ values (0.50 µg mL⁻¹) is noteworthy, which is 20 times less than the

![](image_url)

Figure 2. Chromatogram obtained through the analysis of GHB in urine samples according the purposed methodology. (A) Blank urine with addition of 10 µg mL⁻¹ GHB-D6 in SCAN mode; (B) Blank urine with addition of 10 µg mL⁻¹ GHB-D6 in SIM mode; (C) Blank urine with addition of 10 µg mL⁻¹ GHB-D6 and GHB respectively. Blank urine contains endogenous (basal) GHB.
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Linear Range (µg mL⁻¹)

| Extraction procedure | Derivatization | Analysis time (minutes) | Linear Range (µg mL⁻¹) | Reference |
|----------------------|----------------|-------------------------|------------------------|-----------|
| LLE                  | BSTFA + 1% TMCS| 23                      | 0.5-20*                | purposed method |
| LLE                  | BSTFA + 1% TMCS| 26                      | 0.5-40                 | Busardo et al., 2014³³ |
| LLE                  | BSTFA + 1% TMCS| 32                      | 1-100                  | Kütting et al., 2019³⁵ |
| SPME                 | Hexyl-chloroformate/pyridine | 33          | 0.05-20                | Brown et al., 2007⁴⁴ |
| LLE                  | BSTFA + 1% TMCS| 35                      | 5-500                  | Hassan & Cooper, 2015⁴¹ |
| LLE                  | BSTFA + 1% TMCS| 38                      | 2-12                   | Andreeen et al., 2010⁴² |
| Headspace-Trap       | GBL conversion | 40                      | 5-1000                 | LeBeau et al., 2000⁴³ |
| LLE                  | GBL conversion | 46                      | 2-200                  | Ferrara et al., 1993⁴⁴ |
| LLE                  | BSTFA + 1% TMCS| 46                      | 20-500                 | Aromatario et al., 2012⁴⁵ |
| LLE                  | BSTFA + 1% TMCS| 49                      | 0.01-30                | Shima et al., 2009⁴⁶ |
| Headspace-SPME       | GBL conversion | 50                      | 1-100                  | Frison et al., 2000⁴⁷ |
| LLE                  | BSTFA + 1% TMCS| 51                      | 1-200                  | Kintz et al., 2005⁴⁸ |
| LLE                  | BSFTA + 1% TMCS| 51                      | 1-200                  | Jamey et al., 2016⁴⁹ |
| SPE                  | BSFTA + 1% TMCS| 51                      | 1-100                  | Kim et al., 2019⁵⁰ |
| LLE                  | BSFTA + 1% TMCS| 52                      | 1-100                  | Kietzerow et al., 2020³⁷ |
| LLE                  | BSFTA + 1% TMCS| 55                      | 0.2-20                 | Elian, 2000⁵⁵ |
| Headspace-Trap       | DMS/TBA-HSO₄   | 56                      | 5-150                  | Ingels et al., 2013⁵¹ |
| LLE                  | BSFTA + 1% TMCS| 60                      | 1-200                  | Villain et al., 2003⁵⁵ |
| LLE                  | BSFTA + 1% TMCS| 76                      | 0.1-5                  | Crookes et al., 2004⁵⁶ |
| LLE                  | GBL conversion | 83                      | 0.01-100               | Braisford et al., 2010⁵⁷ |
| SPE                  | BSFTA          | 111                     | 1.4-55                 | Arnold et al., 2018⁴⁴ |
| SPE                  | BSFTA + 1% TMCS| 121                     | 5-50                   | Adamowicz e Kala, 2010³³ |

Analysis time consider all steps for GHB determination per sample. BSTFA, N-O-Bis(trimethylsilyl)trifluoroacetamide; TMCS, trimethylchlorosilane; DMS, Dimethyl sulfate; TBA, tributylamine; LLE, liquid-liquid extraction; SPE, solid phase extraction; SPME, solid phase microextraction. *Dilution integrity was successfully performed considering the concentration of 50 µg mL⁻¹.

cut-off established by UNODC and SOFT.⁵­³⁴ So, the purpose method allows endogenous GHB determination when necessary. Several studies have evaluated the endogenous levels of GHB in a series of volunteers to determine a cut-off, differentiating endogenous from exogenous GHB. Cut-off value from 10 µg mL⁻¹ has been reported in the literature⁵⁰,⁵³-⁵⁹ and DFC guidelines.⁴³,⁴⁵ However, cut-off value from 5 µg mL⁻¹⁴,⁴⁰,⁴¹ and 6 µg mL⁻¹¹⁴,⁴² has already been reported as the best index to be used in DFC situations based on observations of endogenous values that are excreted in the urine.

However, the urinary concentration of GHB after ingesting a dose of this drug (50 mg kg⁻¹) was determined previously.⁴ It was verified an extensive concentration range whose average was 3.8 µg mL⁻¹ from 6 hours after GHB consumption while urinary levels lower than 5 µg mL⁻¹ from 12 hours posterior GHB administration. So, this is an indication the detection period could be less than 12 hours after GHB administration.⁴ These results were corroborated by other findings, where the authors indicate GHB analysis in urine specimens within 3 days of sampling with the specimens stored at -20 °C or 4 °C or add sodium fluoride (1-2%) into the sample, which acts as an enzyme inhibitor. These procedures prevent in vitro GHB formation.⁴³ Despite the discussions about the cut-off values to differentiate endogenous from exogenous GHB and its stability, the proposed method was able to unequivocally determine concentrations lower than the lowest value proposed (LOQ = 0.5 µg mL⁻¹).

Table 2 shows the scientific literature studies that deals with the analysis of GHB by GC-MS in urine samples and summarizes the total analysis time considering all steps, demonstrating that the purposed methodology adopted by the authors is more efficient in terms of time and handling. In this regard, the procedure developed to determine this analyte in this biological matrix has a working range compatible with lower concentrations proposed by other studies without the need for changes.

CONCLUSIONS

The purposed procedure to determine GHB in urine samples by GC-MS is simple and ultra fast, compatible with any toxicology laboratory with immediate application, being able to unequivocally identify the analyte, even in the lower concentrations proposed by other studies as well as the cut-off established by UNODC and SOFT (10 µg mL⁻¹). Therefore, this methodology can be used for clinical analysis when endogenous GHB requires to be measured.

ACKNOWLEDGEMENTS

We thank CNPq for the doctoral scholarship for carrying out this study.

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