

## CONFIRMATION OF BARBITURATES BY GAS CHROMATOGRAPHY – MASS SPECTROMETRY

### 8.1 POLICY

This test method may be used to confirm the presence of amobarbital (AMB), butalbital (BTB), pentobarbital (PTB), phenobarbital (PHB) and secobarbital (SCB) in biological samples. Quantitative results obtained through the use of this method will only be reported within the validated dynamic range. Reporting of results following the application of this method will be contingent upon a thorough review and acceptance of quality control data and the qualification of individual results under the criteria for acceptance.

Any adjustments or deviations from the procedures below must be approved by either the State Toxicologist, a Manager, or a Supervisor, and appropriately documented in the batch file.

### 8.2 PURPOSE

The purpose of this standard operating procedure (SOP) is to provide technical direction for the identification and quantitation of select barbiturates present in biological specimens. This procedure will serve as the laboratory document describing sample preparation, instrumental analysis, data analysis, criteria for acceptance and reporting of the specified compounds.

### 8.3 PRINCIPLE

The targeted compounds and internal standard, hexobarbital (HXB), are isolated from whole blood, serum, plasma, urine or other submitted biological samples by the use of liquid-liquid extraction (LLE). Following LLE, the specimens, now termed extracts, are injected into a gas chromatograph (GC) where they are separated between a gaseous mobile and liquid stationary phase. Each compound exits the GC at a reproducible time which is termed its retention time.

The GC is coupled to a mass spectrometer (MS) detector equipped with an electron ionization source. As each compound is ionized in the source, selected-ion-monitoring is used to measure the mass-to-charge ratios of each compound and its related fragments. Multiple-point, internal standard calibration is used to generate a calibration curve. The concentration of any target compound identified in a sample is determined from its calibration curve.

### 8.4 SPECIMENS

8.4.1 The specimen volume is 0.5 mL.

8.4.2 Specimens include whole blood, serum, plasma, urine, and tissue homogenate.

8.4.3 Dilutions of specimens may be analyzed at the Forensic Scientist's discretion; however, this should be done in addition to testing the standard specimen volume, unless sample quantity dictates otherwise.

8.4.4 Analysis of larger specimen volumes must be approved and documented.

### 8.5 REAGENTS, MATERIALS AND EQUIPMENT

#### 8.5.1 REAGENTS

- 8.5.1.1 Acetonitrile
- 8.5.1.2 Certified blank blood
- 8.5.1.3 Ethyl acetate
- 8.5.1.4 Extraction solvent  
Add 20 mL ethyl acetate to a glass flask. Add 20 mL hexanes and mix. Use on date of preparation only.
- 8.5.1.5 Hexanes
- 8.5.1.6 Hydrochloric acid (HCl, concentrated)
- 8.5.1.7 0.1M Hydrochloric acid  
To 400 mL DI H<sub>2</sub>O, add 4.2 mL concentrated HCl. Dilute to 500 mL with DI H<sub>2</sub>O. Store this in a glass bottle at room temperature for up to 6 months.
- 8.5.1.8 Methanol
- 8.5.1.9 0.1M phosphate buffer (pH6)  
Dissolve 1.7 g Na<sub>2</sub>HPO<sub>4</sub> and 1.14 g NaH<sub>2</sub>PO<sub>4</sub> in 800 mL DI H<sub>2</sub>O. Dilute to 1 L with DI H<sub>2</sub>O and mix. Check the pH and, if necessary, adjust to 6 ±0.5. Store the buffer in a glass bottle at room temperature for up to one year.
- 8.5.1.10 0.1M phosphate buffer (pH5)  
Add 32 mL 0.1M HCl to 300 mL 0.1M phosphate buffer pH6 and mix. Adjust pH as necessary. Store in a glass bottle at room temperature for up to 1 year.
- 8.5.1.11 Sodium phosphate, dibasic anhydrous (Na<sub>2</sub>HPO<sub>4</sub>)
- 8.5.1.12 Sodium phosphate, monobasic monohydrate (NaH<sub>2</sub>PO<sub>4</sub> • H<sub>2</sub>O)
- 8.5.1.13 Trimethylanilinium hydroxide (TMPAH) in methanol

## 8.5.2 MATERIALS

- 8.5.2.1 Autosampler vials, inserts and caps
- 8.5.2.2 Disposable 16 x 100mm tubes
- 8.5.2.3 Disposable screw-cap tubes or centrifuge tubes with closures
- 8.5.2.4 Disposable pipette tips
- 8.5.2.5 Disposable safety closures for 16 x 100mm tubes
- 8.5.2.6 GC column (Agilent HP-5MS; 30 m x 0.250 mm i.d. x 0.250 µm film thickness, or equivalent)

8.5.2.7 Laboratory glassware (graduated cylinders, flasks)

8.5.2.8 Volumetric glassware (flasks)

### 8.5.3 EQUIPMENT

8.5.3.1 Agilent GC (6890 or equivalent)

8.5.3.2 Agilent MS (5973 or equivalent)

8.5.3.3 Calibrated, adjustable air-displacement pipettes

8.5.3.4 Centrifuge

8.5.3.5 Evaporator (Caliper LS, formerly Zymark, TurboVap)

8.5.3.6 pH Meter and/or indicating pH paper

8.5.3.7 Rotary mixer

8.5.3.8 Vortex mixer

## 8.6 STANDARDS, CALIBRATORS AND CONTROLS

### 8.6.1 STANDARDS

8.6.1.1 Reference materials (referred to interchangeably in this method as stock standards) are used for the preparation of working standards which in turn are used to produce calibrators, positive controls and the working internal standard.

8.6.1.2 Stock standards and stock internal standard are purchased from an approved reference material supplier and include the following:

|                       |           |
|-----------------------|-----------|
| a. Amobarbital:       | 1.0 mg/mL |
| b. Butalbital:        | 1.0 mg/mL |
| c. Hexobarbital (IS): | 1.0 mg/mL |
| d. Pentobarbital:     | 1.0 mg/mL |
| e. Phenobarbital:     | 1.0 mg/mL |
| f. Secobarbital:      | 1.0 mg/mL |

8.6.1.3 Working standard (0.1 mg/mL)

- Using a calibrated pipette, measure 1 mL each of AMB, BTB, PTB, PHB and SCB stock standards into a 10 mL class-A volumetric flask.
- Add methanol to the flask to the designated volume.
- The final concentration of the working standard is 0.1 mg/mL. The working standard is stored in the freezer in an amber bottle and expires one year from the date of preparation.

8.6.1.4 Working internal standard (0.05 mg/mL)

- Using a calibrated pipette, measure 0.5 mL HXB stock internal standard into a 10 mL class-A volumetric flask.
- Add methanol to the flask to the designated volume.

- c. The final concentration of the working internal standard is 0.05 mg/mL. The working internal standard is stored in the freezer in an amber bottle and expires one year from the date of preparation.

## 8.6.2 CALIBRATORS

- 8.6.2.1 Calibrators are prepared in certified blank blood at the time of analysis using the working standard. The preparation of the calibrators is detailed in 8.7 SAMPLE PREPARATION. If necessary, calibrators may be prepared in alternate matrices provided that the matrix has been previously determined to not contain any of the compounds tested for by this procedure.

## 8.6.3 CONTROLS

### 8.6.3.1 Negative Control

- a. At least one negative whole blood control is tested with every batch. The negative control is prepared using certified blank blood.
- b. When testing different sample types, wherever possible, include a negative control prepared from that matrix. (For example, when analyzing whole blood and urine samples the batch shall include at least one negative whole blood control and at least one negative urine control.)

### 8.6.3.2 Positive Controls

- a. Two positive whole blood controls are tested with every batch. The positive controls are prepared using certified blank blood to which the designated volume of control working standard has been added.
- b. Control stock standards are obtained from an approved reference material supplier.
- c. The control stock standards must be either a different lot number or from a different supplier to those used in producing the working standard.
- d. The control working standard (0.1 mg/mL) is prepared as described in 8.6.1.3.
- e. The preparation of the positive whole blood controls is detailed in 8.7 SAMPLE PREPARATION. Alternatively, quality control personnel may provide in-house positive controls.
- f. When testing different sample types, wherever possible, include at least one positive control prepared from that matrix.

## 8.7 SAMPLE PREPARATION

- 8.7.1 Label a clean 16 x 100mm tube for each member of the test batch. (i.e. Calibrator, control, case sample)
- 8.7.2 Place 1 mL of 0.1M phosphate buffer pH5 into each tube.
- 8.7.3 Add 0.5 mL of certified blank whole blood into each of the five calibrator tubes, the two positive control tubes and the negative control tube(s).
- 8.7.4 Prepare a 1:10 dilution of the working standard. (0.01 mg/mL)

- a. Using a calibrated pipette, combine 0.1 mL of the working standard with 0.9 mL of methanol in a labeled tube.
- b. Cap and vortex mix. This dilution shall be disposed of after calibrator preparation.

8.7.5 Using the working standard and the prepared dilution, spike the calibrators according to the following table.

| Calibrator Description  | Volume ( $\mu$ L) Added | Working Standard |
|-------------------------|-------------------------|------------------|
| Calibrator 1 (0.5 mg/L) | 25                      | 0.01 mg/mL       |
| Calibrator 2 (2.0 mg/L) | 100                     | 0.01 mg/mL       |
| Calibrator 3 (5.0 mg/L) | 25                      | 0.1 mg/mL        |
| Calibrator 4 (10 mg/L)  | 50                      | 0.1 mg/mL        |
| Calibrator 5 (20 mg/L)  | 100                     | 0.1 mg/mL        |

8.7.6 Prepare a 1:10 dilution of the control working standard. (0.01 mg/mL)

- a. Using a calibrated pipette, combine 0.1 mL of the control working standard with 0.9 mL of methanol in a labeled tube.
- b. Cap and vortex mix. This dilution shall be disposed of after control preparation.

8.7.7 Using the control working standard and the prepared dilution, spike the positive controls according to the following table.

| Control Description  | Volume ( $\mu$ L) Added | Control Working Standard |
|----------------------|-------------------------|--------------------------|
| Control 1 (3.0 mg/L) | 100                     | 0.01 mg/mL               |
| Control 2 (15 mg/L)  | 100                     | 0.1 mg/mL                |

8.7.8 If in-house positive controls are being used, transfer 0.5 mL of each into their labeled tubes.

8.7.9 Sample 0.5 mL of each case sample into its respective tube.

8.7.10 Add 50  $\mu$ L of the working internal standard solution to each tube. Final concentration of the internal standard is 5 mg/L.

8.7.11 Cap the tubes and briefly vortex mix.

8.7.12 Add 3 mL extraction solvent to each tube.

8.7.13 Cap the tubes and place on a rotary mixer for 20 minutes.

8.7.14 Centrifuge the tubes for 10 minutes at 3500rpm.

8.7.15 Transfer the organic layer to clean, labeled 10 mL centrifuge or screw cap tubes.

8.7.16 Transfer the tubes to the evaporator and evaporate the extracts to dryness at 50°C. Make sure the extracts are evaporated to dryness before reconstitution.

8.7.17 Reconstitute the extracts by the addition of 100  $\mu$ L TMPAH in methanol to each tube and cap. Briefly vortex mix the tubes. If necessary, centrifuge the tubes for 2 minutes at 2000 rpm to collect the extracts at the bottom of the tubes.

8.7.18 Transfer the extracts to labeled autosampler vials with inserts and cap.

## 8.8 INSTRUMENTAL PARAMETERS

The instrumental parameters can be found in Appendix A. Prepare a sequence table by first setting the data path in ChemStation to the date of the test. After entering all vial locations, sample descriptions, comments and/or lot numbers in the sequence table ensure that the method listing in the table is BARB.M for each line.

## 8.9 SPECIAL INSTRUCTIONS

Methanol is to be used when solvent blanks are included in the test sequence.

## 8.10 DATA ANALYSIS

8.10.1 Analysis of the batch data is conducted using the instrumental data analysis software in ChemStation.

8.10.2 Quantitative calculations are generated by internal standard, multi-point, linear regression with a 1/a (inverse of concentration) weighting factor. The calibration curves are updated using the calibrator results for the batch; no historical calibration curves are permitted.

8.10.3 Printed reports for each vial in the batch are generated for review along with the updated calibration curves.

8.10.4 Technical review of the batch is conducted according to the criteria listed below.

## 8.11 CRITERIA FOR BATCH ACCEPTANCE

If the analysis of the batch meets the criteria listed below, the results for the specimens are accepted.

### 8.11.1 Calibrators and calibration curves

8.11.1.1 Chromatographic peaks for AMB, BTB, PTB, PHB, SCB and internal standard shall appear symmetrical (i.e. no co-elution, split peaks, or shoulders).

8.11.1.2 Retention times shall be within  $\pm 2\%$  and ion ratios shall be within  $\pm 20\%$  of those in calibrator 3. These are inclusive ranges.

8.11.1.3 Quantitative results for target compounds in each calibrator shall be within  $\pm 20\%$  of their target values with the exception of calibrator 1 which shall be within  $\pm 25\%$  of their targets. These are inclusive ranges. Result comparisons will use values truncated after the first decimal place in units of mg/L.

8.11.1.4 The calibration curves for AMB, BTB, PTB, PHB and SCB shall have correlation coefficients  $\geq 0.99$ .

8.11.1.5 The failure to meet any of these criteria for one compound does not invalidate the acceptability of another compound.

### 8.11.2 Controls

8.11.2.1 The negative control(s) shall not identify AMB, BTB, PTB, PHB or SCB above its limit of detection. Identification is based on a)

acceptable retention time matching, b) distinct peaks present for all selected ions, and c) acceptable ion ratios.

#### 8.11.2.2 Positive controls

- a. Chromatographic peaks for AMB, BTB, PTB, PHB and SCB and internal standard shall appear symmetrical.
- b. Retention times shall be within  $\pm 2\%$  and ion ratios shall be within  $\pm 20\%$  of those in calibrator 3 for each compound in the positive control. These are inclusive ranges.
- c. Quantitative results for AMB, BTB, PTB, PHB and SCB in each control shall be within  $\pm 20\%$  of their target values. These are inclusive ranges. Result comparisons will use values truncated after the first decimal place in units of mg/L.
- d. The failure to meet any of these criteria for one compound does not invalidate the acceptability of another compound.
- e. At least one positive control must meet these criteria for all compounds for the batch to be accepted.

#### 8.12 CRITERIA FOR CASE SAMPLE ACCEPTANCE

If the criteria for batch acceptance have been satisfied, the results of individual case samples are deemed suitable for reporting if the following criteria are met.

- 8.12.1 Any chromatographic peak for AMB, BTB, PTB, PHB or SCB shall appear symmetrical.
- 8.12.2 The retention times for any reportable compounds are  $\pm 2\%$  and the ion ratios are within  $\pm 20\%$  of those in calibrator 3. These are inclusive ranges.
- 8.12.3 The quantitative results for each identified compound must be within the dynamic range of the test method.
- 8.12.4 When dilutions of case samples are tested, the quantitative result(s) before multiplication shall be within the dynamic range of the test method.

#### 8.13 REPORTING

- 8.13.1 Results are reported in units of milligrams per liter (mg/L).
- 8.13.2 Results are truncated to no more than two significant figures for reporting.
  - a. Example 1: amobarbital is measured as 6.38 mg/L.
  - b. The result is truncated to 6.3 mg/L (two significant figures) and reported.
  - c. Example 2: phenobarbital is measured at 15.92 mg/L.
  - d. The result is truncated to 15.9 mg/L, but reported as 15 mg/L (two significant figures).
  - e. Example 3: secobarbital is measured at 10.02 mg/L.
  - f. The result is truncated to 10.0 mg/L (three significant figures), but reported as 10 mg/L (one significant figure).
- 8.13.3 When multiple dilutions are analyzed, the smallest dilution within the dynamic range is reported.



8.14 METHOD PERFORMANCE

- 8.14.1 Limit of detection: 0.05 mg/L
- 8.14.2 Lower limit of quantification: 0.5 mg/L
- 8.14.3 Dynamic range: 0.5 mg/L – 20 mg/L
- 8.14.4 Upper limit of quantitation: 20 mg/L

8.15 TRACEABILITY

- 8.15.1 Traceability of the reference materials to SI units is provided through the certificate of analysis provided by the approved reference material supplier.

APPENDIX A  
 INSTRUMENTAL PARAMETERS

GAS CHROMATOGRAPH

| Split/Splitless Inlet |                                 | Oven/Column         |               |
|-----------------------|---------------------------------|---------------------|---------------|
| Mode                  | Split                           | Carrier Gas Mode    | Constant Flow |
| Inlet Liner           | 4mm splitless w/glass wool plug | Carrier Gas Flow    | 2.0 mL/min    |
| Temperature           | 280°C                           | Initial Temperature | 110° C        |
| Split Ratio           | 40:1                            | Initial Time        | 1.00 min      |
| Gas Type              | Helium                          | Ramp Rate           | 15° C/min     |
| Gas Saver             | On                              | Final Temperature   | 300° C        |
| Gas Saver Flow        | 15.0 mL/min                     | Final Time          | 0.33 min      |
| Gas Saver Time        | 2.00 min                        |                     |               |
| Autosampler           |                                 |                     |               |
| Injection Volume      | 2.0 µL                          |                     |               |
| Solvent Wash A        | 4 (Acetonitrile)                |                     |               |
| Solvent Wash B        | 4 (Methanol)                    |                     |               |
| Sample Pumps          | 2                               |                     |               |

MASS SPECTROMETER

| Solvent Delay     | 5.00 min      | MS Quad Temperature   | 150°C   |
|-------------------|---------------|-----------------------|---------|
| EM Offset         | Set in tune   | MS Source Temperature | 230°C   |
| Resolution        | Low           | Dwell Time            | 50 msec |
| Signals           | Ions          | Ion Ratios            |         |
| Amobarbital       | 169, 184, 185 | 184/169, 185/169      |         |
| Butalbital        | 196, 195, 209 | 195/196, 209/196      |         |
| Hexobarbital (IS) | 235, 169      | 169/235               |         |
| Pentobarbital     | 169, 184, 112 | 184/169, 112/169      |         |
| Phenobarbital     | 232, 175, 146 | 175/232, 146/232      |         |
| Secobarbital      | 196, 195, 181 | 195/196, 181/196      |         |



