# Final Report

# Asia Pacific Metrology Programme Supplementary Comparison (APMP.QM-S8)

Determination of Mass Fraction of Benzoic Acid, Methyl Paraben and n-Butyl Paraben in Soy Sauce

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# 1.0 INTRODUCTION

Preservatives such as benzoic acid and parabens are being tested on a regular basis by regulatory authorities and food testing laboratories as they are widely found in condiments and beverages such as fruit juices, sports drinks and soft drinks. The consumption of excessive amount of preservatives can lead to adverse health effects such as allergy and asthma. Also, the levels of these preservatives in food are strictly regulated and the maximum permitted concentrations in each type of food have been established in most countries.

The study was first proposed by the Health Sciences Authority (HSA), Singapore as a key comparison and was presented at the APMP TCQM meeting in Taipei in 2013. It was subsequently presented at the CCQM OAWG meeting in April 2014. The meeting recognised the need for a key comparison to underpin new and existing CMCs, and it also noted that the topic involved a condiment (see Objectives) which is commonly used in the Asia Pacific region. Hence, the meeting recommended that the study should proceed as a subsequent comparison under the APMP. CCQM OAWG members from other RMOs would also be invited to participate. An approval was subsequently obtained from the APMP TCQM Chair in April 2014 to organise this supplementary comparison, along with a parallel pilot study.

# 2.0 OBJECTIVES

The comparison aimed to enable participating NMIs/DIs to demonstrate their competence in the determination of three common preservatives; namely: benzoic acid, methyl paraben and n-butyl paraben. The chosen matrix material was soy sauce, a traditional condiment commonly used in the preparation of Asian cuisines.

# 3.0 MEASURANDS

The mass fractions (in mg/kg) of benzoic acid, methyl paraben and n-butyl paraben in soy sauce, were determined by the participating NMIs/DIs. The chemical structures and physical information of the analytes are as shown:



 Benzoic acid
 Methyl paraben
 n-Butyl paraben

 CAS No.: 65-85-0
 CAS No.: 99-76-3
 CAS No.: 94-26-8

 MW: 122.12
 MW: 152.15
 MW: 194.23

 logKow: 1.87<sup>1</sup>
 logKow: 1.96<sup>1</sup>
 logKow: 3.57<sup>1</sup>

The mass fractions of benzoic acid, methyl paraben and n-butyl paraben in the study material were in the range of 50 to 200 mg/kg. The study material was intentionally prepared to contain a lower concentration of the analytes than the regulatory limits in most countries in order to provide a higher level of analytical challenge for the participating NMIs/DIs.

# 4.0 STUDY MATERIAL

# 4.1 **PREPARATION**

The study material was prepared by HSA. The soy sauce was sourced from local supermarkets and screened for the presence of preservatives using liquid chromatographydiode array detection (LC-DAD). A brand of soy sauce with no preservatives found was then used to prepare the study material. The soy sauce was poured into a clean, dry drum and spiked with benzoic acid (NIST, in-house qNMR purity value:  $998 \pm 4 \text{ mg/g}$ ), methyl paraben (Sigma Aldrich, in-house qNMR purity value:  $994 \pm 4 \text{ mg/g}$ ) and n-butyl paraben (Sigma Aldrich, in-house qNMR purity value:  $1000 \pm 5 \text{ mg/g}$ ) in methanolic solution. The mixture was then stirred under an atmosphere of nitrogen for 2 hours at room temperature (about 18 to 25 °C). Thereafter, the soy sauce was aliquoted into amber glass bottles. A total of 89 bottles of study material were prepared.

<sup>&</sup>lt;sup>1</sup> S.G. Machatha, S.H. Yalkowsky, International J. Pharmaceutics, 294 (2005), 185-192.

# 4.2 HOMOGENEITY STUDIES

The homogeneity of the study material was established by employing gas chromatographyisotope dilution mass spectrometry (GC-IDMS). Ten bottles were randomly and stratifically selected. Two subsamples were taken from each bottle. The sample size was about 1 g. Using one-way ANOVA at 95 % level of confidence, the material was found to be sufficiently homogeneous for the study. Summaries of the ANOVA for the homogeneity study are given in Tables 1-3.

| Source of variance | SS    | DF | MS   | F    | p-Value | <b>F</b> <sub>critical</sub> |
|--------------------|-------|----|------|------|---------|------------------------------|
| Between bottles    | 14.23 | 9  | 1.58 | 1.05 | 0.46    | 3.02                         |
| Within bottles     | 15.01 | 10 | 1.50 |      |         |                              |
| Total              | 29.24 |    |      |      |         |                              |

## Table 1: Summary of ANOVA for benzoic acid

## Table 2: Summary of ANOVA for methyl paraben

| Source of variance | SS    | DF | MS   | F    | p-Value | F <sub>critical</sub> |
|--------------------|-------|----|------|------|---------|-----------------------|
| Between bottles    | 6.75  | 9  | 0.75 | 2.14 | 0.13    | 3.02                  |
| Within bottles     | 3.50  | 10 | 0.35 |      |         |                       |
| Total              | 10.25 |    |      |      |         |                       |

## Table 3: Summary of ANOVA for n-butyl paraben

| Source of variance | SS   | DF | MS   | F    | p-Value | F <sub>critical</sub> |
|--------------------|------|----|------|------|---------|-----------------------|
| Between bottles    | 4.75 | 9  | 0.53 | 1.56 | 0.25    | 3.02                  |
| Within bottles     | 3.38 | 10 | 0.34 |      |         |                       |
| Total              | 8.13 |    |      |      |         |                       |

The relative uncertainties of the between-bottle homogeneity,  $u_{bb}$ , for benzoic acid, methyl paraben and n-butyl paraben were found to be 0.37 %, 0.45 % and 0.32 %, respectively.

# 4.3 STABILITY STUDIES

The stability of the study material at about 40 °C and 25 °C was established by LC-DAD and GC-IDMS, respectively.

A short-term stability testing using isochronous design was carried out over a period of 28 days at a simulated transport temperature of about 40 °C. Two randomly selected bottles were stored at the reference temperature of about 4 °C. Two bottles were transferred from about 40 °C to the reference temperature each time, on two occasions over the study period. One subsample was taken from each bottle and analysed at least five times. The effect of exposure time on the stability of the analytes was determined by fitting linear regression lines to the data set. The slope (*b*) was tested for statistical significance using Student's *t* test at 95% confidence level, where the *t* value was calculating by dividing *b* by its standard deviation *s*(*b*) and compared against the critical *t* value. The statistical results given in Table 4 indicated that no significant trend at 95% confidence level was detected. Hence, no significant instability of the study material was observed at about 40 °C for a period of 28 days.

| Descriptions                             | Benzoic Acid | Methyl Paraben | n-Butyl Paraben |
|--|--------------|----------------|-----------------|
| Slope of the regression line (b)         | -0.149       | 0.056          | 0.048           |
| Intercept of the regression line $(b_0)$ | 161.128      | 106.189        | 102.552         |
| Variance of the points $(s^2)$           | 1.201        | 0.363          | 0.166           |
| Standard deviation of the points (s)     | 1.096        | 0.602          | 0.408           |
| Uncertainty of slope [s(b)]              | 0.0553       | 0.030          | 0.021           |
| Calculated $t(\frac{ b }{s(b)})$         | 2.69         | 1.85           | 2.32            |
| Critical t factor $(t_{0.95,n-2})$       | 12.71        | 12.71          | 12.71           |

Table 4: Summary of stability study for benzoic acid, methyl paraben and n-butyl at about 40 °C for 28 days

The same approach was used to determine the stability of the study material under the storage temperature of about 25 °C using a classical design. The study was carried out on

six occasions over a period of 454 days. The statistical results for the significance test of the regression coefficient are given in Table 5. The results showed that the slopes of the line were not significantly different from zero, using Student's *t*-test at 95 % confidence level, i.e. no significant instability was observed when the study sample was stored at about 25 °C. The uncertainties of the stability of the analytes were estimated by calculating  $s(b)^*454$ .

Table 5: Summary of stability study results for benzoic acid, methyl paraben and n-butyl paraben at about 25 °C for 454 days

| Descriptions                                | Benzoic Acid | Methyl Paraben | n-Butyl Paraben |
|---|--------------|----------------|-----------------|
| Slope of the regression line (b)            | -0.0041      | -0.0010        | -0.0022         |
| Intercept of the regression line $(b_0)$    | 156.606      | 100.768        | 98.386          |
| Variance of the points $(s^2)$              | 2.301        | 0.985          | 0.376           |
| Standard deviation of the points (s)        | 1.517        | 0.992          | 0.614           |
| Uncertainty of slope [s(b)]                 | 0.0038       | 0.0025         | 0.0016          |
| Calculated $t(\frac{ b }{s(b)})$            | 1.06         | 0.41           | 1.40            |
| Critical <i>t</i> factor ( $t_{0.95,n-2}$ ) | 2.78         | 2.78           | 2.78            |
| Relative standard uncertainty of stability  | 1.12 %       | 1.14 %         | 0.72 %          |

# 5.0 SCHEDULE

| Table 6: Schedule of the comparison |                                    |  |  |  |  |
|-------------------------------------|------------------------------------|--|--|--|--|
| Period                              | Event                              |  |  |  |  |
| September 2014                      | Call for participation             |  |  |  |  |
| January 2015                        | Distribution of study sample       |  |  |  |  |
| May 2015                            | Deadline for submission of results |  |  |  |  |

# 6.0 **REGISTRATION, SAMPLE RECEIPT AND REPORT SUBMISSION**

A total of ten NMIs/DIs participated in the comparison for benzoic acid, while six NMIs/DIs participated in the comparison for methyl paraben and n-butyl paraben.

One package comprising one bottle of study material and the Material Safety Data Sheet of soy sauce was sent to all participating NMIs/DIs on 26 Jan 2015. A temperature strip was pasted on the bottle to indicate any exposure of the study material to above 40 °C during transportation. Eight of the participating NMIs/DIs received the package within four days after dispatch. NIM, China received the package 16 days after dispatch due to customs issue. All the participating NMIs/DIs received the study sample intact and the study materials were not exposed to above 40 °C. The NMIs/DIs were informed to keep the study material at or below 25 °C, before analysis.

Due to the coincidence of the CCQM meeting and technical issues faced by a participating NMI/DI, the initial reporting deadline of 24 April 2015 was extended to 8 May 2015 and later re-extended to 22 May 2015, respectively, with the approval of the APMP TCQM Chair. All the participating NMIs/DIs submitted their results on or before the final deadline.

Information on participating NMIs/DIs, contacts, measurands registered, sample receipts and report submissions are summaried in Table 7.

Table 7: Information on participating NMIs/DIs, contacts, measurands registered, sample receipts and report submissions

| No. | Participating Institutes,<br>Department/Laboratory  | Economy                 | Contact Person      | Measurands Registered                                  | Sample<br>Receipt Date | Report<br>Submission<br>Date |
|-----|---|-------------------------|---------------------|--|------------------------|------------------------------|
| 1   | EXHM/GCSL-EIM<br>National Laboratory of Chemical<br>Metrology/General Chemistry State<br>Laboratories - Hellenic Institute of Metrology,<br>Chemical Metrology Laboratory | Greece                  | Dr Elias Kakoulides | Benzoic acid   | 29 Jan 2015            | 21 May 2015                  |
| 2   | GLHK<br>Government Laboratory, Hong Kong SAR,<br>Additives, Contaminants and Composition<br>Section   | Hong Kong<br>SAR, China | Dr Sin-Kam Hui      | Benzoic acid   | 28 Jan 2015            | 19 May 2015                  |
| 3   | HSA<br>Health Sciences Authority, Chemical<br>Metrology Division, Chemical Metrology<br>Laboratory  | Singapore               | Dr Teo Tang Lin     | Benzoic acid, methyl<br>paraben and n-butyl<br>paraben | N.A.                   | 6 May 2015                   |
| 4   | INMETRO<br>Instituto Nacional de Metrologia, Qualidade e<br>Tecnologia, Chemical Metrology Division<br>(DQUIM), Organic Analysis Laboratory<br>(LABOR)                    | Brazil                  | Eliane Rego         | Benzoic acid   | 28 Jan 2015            | 12 May 2015                  |
| 5   | KRISS<br>Korea Institute of Standards and Science,<br>Center of Analytical Chemistry, Division of<br>Metrology for Quality of Life, Center for<br>Organic Analysis        | Republic of<br>Korea    | Dr Seonghee Ahn     | Benzoic acid, methyl<br>paraben and n-butyl<br>paraben | 28 Jan 2015            | 21 May 2015                  |
| 6   | NIM<br>National Institute of Metrology, Food Safety<br>Laboratory   | PR. China               | Dr Xiuqin Li        | Benzoic acid, methyl<br>paraben and n-butyl<br>paraben | 11 Feb 2015            | 22 May 2015                  |

| 7  | RCChem-LIPI<br>Research Center for Chemistry – Indonesian<br>Institute of Sciences, Laboratory of Metrology<br>in Chemistry                                      | Indonesia   | Dyah Styarini              | Benzoic acid, methyl<br>paraben and n-butyl<br>paraben | 27 Jan 2015 | 22 May 2015 |
|----|--|-------------|----------------------------|--|-------------|-------------|
| 8  | STD-ITDI<br>Industrial Technology Development Institute,<br>Standards and Testing Division   | Philippines | Dr Benilda S. Ebarvia      | Benzoic acid   | 27 Jan 2015 | 21 May 2015 |
| 9  | TISTR<br>Thailand Institute of Scientific and<br>Technological Research, Industrial Metrology<br>and Testing Services Centre, Analytical<br>Chemistry Laboratory | Thailand    | Thippaya Junvee<br>Fortune | Benzoic acid, methyl<br>paraben and n-butyl<br>paraben | 27 Jan 2015 | 22 May 2015 |
| 10 | UME<br>TÜBİTAK Ulusal Metroloji Enstitüsü, Organic<br>Chemistry Laboratory   | Turkey      | Dr Ahmet Ceyhan<br>Gören   | Benzoic acid, methyl<br>paraben and n-butyl<br>paraben | 30 Jan 2015 | 22 May 2015 |

# 7.0 REPORTING OF RESULTS

A Report of Results Form was provided to the participating NMIs/DIs for completion. The participating NMIs/DIs were requested to report their results based on at least three subsamples, each with a recommended size of not less than 1 g. The results were reported in the unit of mg/kg, which included standard and expanded uncertainties (95 % level of confidence) for the mean of the replicate determinations. A complete description of the analytical procedure and the uncertainty estimation was also provided by the participating NMIs/DIs.

# 8.0 RESULTS SUBMITTED BY PARTICIPATING INSTITUTES

The reported results are summarised in Tables 8-10.

| NMI/DI            | Overall<br>mean<br>(mg/kg) | No. of<br>subsamples | Combined<br>standard<br>uncertainty<br>(mg/kg) | Coverage<br>factor, k (95 %<br>confidence<br>level) | Expanded<br>uncertainty to give<br>95 % confidence<br>level (mg/kg) |
|-------------------|----------------------------|----------------------|--|---|---|
| INMETRO           | 148.6                      | 9                    | 1.6  | 2.10  | 3.3   |
| TISTR             | 150.5                      | 14                   | 4.4  | 1.97  | 8.7   |
| EXHM/GCSL-<br>EIM | 150.66                     | 10                   | 1.75   | 2.31  | 4.03  |
| UME               | 152.38                     | 6                    | 3.46   | 2   | 6.92  |
| NIM               | 154.5                      | 6                    | 1.6  | 2   | 3.2   |
| GLHK              | 154.6                      | 18                   | 2.4  | 2   | 4.8   |
| KRISS             | 155.03                     | 5                    | 1.90   | 2.262   | 4.29  |
| HSA               | 155.5                      | 9                    | 2.4  | 2   | 4.8   |
| STD-ITDI          | 155.6                      | 4                    | 1.9  | 2   | 3.8   |
| RCChem-LIPI       | 157.5                      | 13                   | 3.9  | 2   | 7.8   |

Table 8: Summary of results for benzoic acid

| NMI/DI          | Overall<br>mean<br>(mg/kg) | No. of<br>subsamples | Combined<br>standard<br>uncertainty<br>(mg/kg) | Coverage factor, k (95 %<br>confidence level) | Expanded<br>uncertainty to<br>give 95 %<br>confidence<br>level (mg/kg) |
|-----------------|----------------------------|----------------------|--|---|--|
| RCChem-<br>LIPI | 96.6                       | 13                   | 3.2  | 2   | 6.4  |
| TISTR           | 98.9                       | 15                   | 3  | 1.99  | 5.9  |
| HSA             | 100.9                      | 9                    | 1.0  | 2   | 1.9  |
| NIM             | 101.0                      | 6                    | 1.1  | 2   | 2.2  |
| UME             | 101.46                     | 6                    | 2.68   | 2   | 5.35   |
| KRISS           | 101.49                     | 5                    | 1.10   | 2.228   | 2.44   |

## Table 9: Summary of results for methyl paraben

Table 10: Summary of results for n-butyl paraben

| NMI/DI      | Overall<br>mean<br>(mg/kg) | No. of<br>subsamples | Combined<br>standard<br>uncertainty<br>(mg/kg) | Coverage<br>factor, k (95 %<br>confidence<br>level) | Expanded<br>uncertainty to<br>give 95 %<br>confidence level<br>(mg/kg) |
|-------------|----------------------------|----------------------|--|---|--|
| HSA         | 97.2                       | 9                    | 1.2  | 2   | 2.3  |
| NIM         | 97.3                       | 6                    | 1.3  | 2   | 2.6  |
| RCChem-LIPI | 97.3                       | 12                   | 3.5  | 2   | 7.0  |
| TISTR       | 100.8                      | 15                   | 3.7  | 2.01  | 7.5  |
| KRISS       | 101.09                     | 5                    | 1.21   | 2.365   | 2.86   |
| UME         | 103.58                     | 6                    | 2.77   | 2   | 5.54   |

# 8.1 ADDITIONAL INVESTIGATION

When the Draft A report was first submitted, the then APMP TCQM Chair, Dr Euijin Hwang, commented that while the results of benzoic acid and methyl paraben seemed very good

and consistent, the results of n-butyl paraben would need more discussion. As there appeared to be two distinct groups of results for n-butyl paraben (one group from TISTR, KRISS and UME and the other from HSA, NIM and RCChem-LIPI), additional investigative work was carried out by HSA. In the investigation, a single analyst in HSA performed sample preparation and clean-up using the procedures of HSA (GC-IDMS), NIM (results similar to HSA's but using LC-IDMS/MS) and KRISS (results obtained using LC-IDMS/MS and with similar C18 columns as UME and TISTR; extensive sample preparation/clean-up). Three subsamples were taken and subjected to each sample preparation and clean-up procedure under repeatability conditions. All nine subsamples were analysed using LC columns used by NIM (BEH SHIELD RP18) and KRISS (BEH C18) on a LC-MS/MS. The measurements and associated standard deviation from the injections agreed within the reported results and associated combined standard uncertainty of the reported results made by HSA and KRISS.

Results of the investigation were presented at the 2015 CCQM OAWG meeting in October and 2015 APMP TCQM meeting in November. NIM confirmed their result after the APMP meeting. HSA updated the 2016 CCQM OAWG meeting in April that there was no technical reason for the appearance of the two groups of results.

# 8.2 REFERENCE MATERIALS USED

Summaries of the reference standards and internal standards used are given in Tables 12 to 14.

Benzoic acid (acidimetric) certified reference material (CRM) from NIST (SRM 350b) was used as calibrant by INMETRO, KRISS, STD-ITDI and UME. Additional purity assessment on the chromatographic purity, water content and non-volatile and volatile residues of the CRM was carried out by KRISS. INMETRO verified the calibrant using NMR and mass spectrometry, STD-ITDI verified the calibrant using mass spectrometry, while UME did not report verification of the calibrant. EXHM/GCSL-EIM used benzoic acid (acidimetric) CRM from BAM. Benzoic acid CRM from HSA was used as calibrant by GLHK, HSA, RCChem-LIPI, and TISTR. NIM used its benzoic acid solution CRM.

When the Draft A report was first submitted, the then APMP TCQM Chair, Dr Euijin Hwang, suggested that the traceability issue related to the use of acidimetric CRMs for the determination of benzoic acid should be further discussed.

At the 2015 CCQM OAWG meeting in October, the results were presented and HSA sought the WG's view on results (INMETRO, STD-ITDI and UME), which were traceable to NIST SRM 350b. The WG acknowledged that in principle, the purity value of the SRM was assigned through the amount of H<sup>+</sup> and not benzoic acid. However, NIST vouched for the high purity of the SRM and also pointed out that the SRM had been widely used as the internal standard for qNMR measurements. KRISS had carried out purity assessment on the SRM using mass balance approach and also found the purity value to be high, except that its uncertainty is larger than that reported by NIST using a coulometric method. Despite the larger uncertainty, it would not change the overall uncertainty of the results due to its small contribution. In view of the high purity of the SRM, the WG considered this to be a special case.

Following the OAWG meeting, the results were presented at the APMP TCQM meeting in November 2015. The TCQM decided that it was necessary for INMETRO, STD-ITDI and UME to re-calculate the benzoic acid results with purity value (999.93±3.1mg/g, k=4.3 at approximately 95 % confidence level) from KRISS for traceability, even if it would not affect the result. EXHM/GCSL-EIM performed additional qNMR experiments on the benzoic acid (acidimetric) CRM from BAM using DMSO<sub>2</sub> as an internal standard (previously measured against 3,5-BTFMBA CRM 4601-a from NMIJ).

The re-calculated results of the four NMIs/DIs are given in Table 11. The results of STD-ITDI and INMETRO did not change, while those of EXHM/GCSL-EIM and UME did not deviate much from their original submitted results. Subsequently, the original submitted results of EXHM/GCSL-EIM and UME were used for evaluation.

| Participating | Original res | sult (mg/kg)   | Re-calculated | Re-calculated result (mg/kg) |  |  |
|---------------|--------------|----------------|---------------|------------------------------|--|--|
| institute     | Result       | U (at 95 % CI) | Result        | U (at 95 % CI)               |  |  |
| EXHM/GCSL     | 150.66       | 4.03           | 150.65        | 4.04                         |  |  |
| INMETRO       | 148.6        | 3.4            | 148.6         | 3.4                          |  |  |
| STD-ITDI      | 155.6        | 3.8            | 155.6         | 3.8                          |  |  |
| UME           | 152.38       | 6.92           | 152.37        | 7.02                         |  |  |

| Table 11: Original and re-calculated results for benzoi | c acid |
|---|--------|
| Table 11. Original and to baloalated results for benzo  | 5 uolu |

Labelled isotopes ( $D_5$ -benzoic acid or  ${}^{13}C_6$ -benzoic acid) were used as internal standard by all participating NMIs/DIs, except RCChem-LIPI which employed external calibration.

Methyl paraben CRM from HSA was used as calibrant by HSA, RCChem-LIPI and TISTR, while NIM used its methyl paraben CRM. Methyl paraben standards from commercial sources were used by KRISS and UME. The commercial standard was purity assessed inhouse using qNMR by UME. Purity assessment on the chromatographic purity, water contents and non-volatile and volatile residues of the commercial standard was carried out by KRISS.

Labelled isotopes ( $^{13}C_6$ -, D<sub>3</sub>- or D<sub>4</sub>-methyl paraben) were used as internal standards by HSA, KRISS, NIM, TISTR and UME. RCChem-LIPI did not use any internal standard as external calibration was employed.

n-Butyl paraben CRM from HSA was used as calibrant by HSA, RCChem-LIPI and TISTR, while NIM used its n-butyl paraben CRM. n-Butyl paraben standards from commercial sources were used by KRISS and UME. Purity assessment on the chromatographic purity, water contents and non-volatile and volatile residues of the commercial standard was carried out by KRISS. The commercial standard was purity assessed in-house using qNMR by UME.

Labelled isotopes ( ${}^{13}C_6$ - or D<sub>4</sub>-n-butyl paraben) were used as internal standards by HSA, KRISS, NIM, TISTR and UME. RCChem-LIPI did not use any internal standard as external calibration was employed.

|                   |                         |                            |   | Calibrant    |  |   |   |
|-------------------|-------------------------|----------------------------|---|--------------|--|---|---|
| NMI/DI            | Source                  | Purity value               | Expanded<br>uncertainty of<br>purity value    | Traceability | Verification<br>Technique  | Purity assay  | Internal<br>Standard                          |
| EXHM/GCSL-<br>EIM | BAM U1009               | 99.997%                    | 0.066%  | BAM          |  |   | D₅-Benzoic acid                               |
| GLHK              | HSA                     | 999.9 mg/g                 | 3.3 mg/g                                      | HSA          |  | Conducted by the<br>producer of the<br>calibrant (HSA)  | <sup>13</sup> C <sub>6</sub> -Benzoic<br>acid |
| HSA               | HSA (HRM-<br>1002A)     | 999.9 mg/g                 | 3.3 mg/g                                      | HSA          |  |   | <sup>13</sup> C <sub>6</sub> -Benzoic<br>acid |
| INMETRO           | NIST SRM<br>350b        | 999.78 mg/g<br>(99.9978 %) | 0.044 mg/g<br>(0.0044 %)                      | NIST         | NMR chemical<br>shifts and MS<br>spectrum<br>consistent with<br>benzoic acid | No, a CRM was<br>used instead   | D₅-Benzoic acid                               |
| KRISS             | NIST SRM                | 999.93 mg/g                | 3.1 mg/g                                      | KRISS        | LC/MS, LC-UV, K-<br>F titration, TGA   | Purity assayed by<br>KRISS with<br>chromatographic<br>purity, water<br>contents and non-<br>volatile & volatile<br>residues | <sup>13</sup> C <sub>6</sub> -Benzoic<br>acid |
| NIM               | NIM CRM<br>GBW(E)100006 | 1.00 mg/mL                 | 0.01 mg/mL<br>( <i>U</i> =1%,<br><i>k</i> =2) | NIM          |  |   | <sup>13</sup> C <sub>6</sub> -Benzoic<br>acid |
| RCChem-LIPI       | HSA,<br>Singapore       | 999.9 mg/g                 | 3.3 mg/g                                      | HSA          |  |   |   |

Table 12: Reference standards and internal standards used for the determination of benzoic acid

| STD-ITDI | NIST (SRM<br>350b)       | 999.978 mg/g | 0.044 mg/g | NIST | MS | $D_5$ -Benzoic acid                           |
|----------|--------------------------|--------------|------------|------|----|---|
| TISTR    | HSA (HRM-<br>1002A)      | 999.9 mg/g   | 3.3 mg/g   | HSA  |    | <sup>13</sup> C <sub>6</sub> -Benzoic<br>acid |
| UME      | Benzoic acid<br>SRM 350b | 0.999978     | 0.000044   | NIST |    | D <sub>5</sub> -Benzoic acid                  |

|                 |                         |              |  | Calibrant    |                                      |   |   |
|-----------------|-------------------------|--------------|--|--------------|--------------------------------------|---|---|
| NMI/DI          | Source                  | Purity value | Expanded<br>uncertainty of<br>purity value | Traceability | Verification<br>Technique            | Purity assay  | Internal<br>Standard                            |
| HSA             | HSA (HRM-<br>1003A)     | 999.5 mg/g   | 3.2 mg/g                                   | HSA          |                                      |   | <sup>13</sup> C <sub>6</sub> -Methyl<br>paraben |
| KRISS           | SIGMA                   | 997.90 mg/g  | 0.44 mg/g                                  | KRISS        | LC/MS, LC-UV, K-<br>F titration, TGA | Purity assayed by<br>KRISS with<br>chromatographic<br>purity, water<br>contents and non-<br>volatile & volatile<br>residues | <sup>13</sup> C <sub>6</sub> -Methyl<br>paraben |
| NIM             | NIM CRM<br>GBW(E)100074 | 996 mg/g     | 1.992 mg/g<br>(0.2% , <i>k</i> =2)         | NIM          |                                      |   | <sup>13</sup> C <sub>6</sub> -Methyl<br>paraben |
| RCChem-<br>LIPI | HSA,<br>Singapore       | 999.5 mg/g   | 3.2 mg/g                                   | HSA          |                                      |   |   |
| TISTR           | HSA (HRM-<br>1003A)     | 999.5 mg/g   | 3.2 mg/g                                   | HSA          |                                      |   | Methyl paraben-<br>d <sub>4</sub>               |
| UME             | Across<br>Chemicals     | 0.99998      | 0.003                                      | TÜBİTAK UME  | NMR and Mass<br>Spectrometry         | QNMR  | Methyl (CD <sub>3</sub> )<br>paraben            |

Table 13: Reference materials and internal standards used for the determination of methyl paraben

|                 |                         |              |  | Calibrant    |                                      |   |   |
|-----------------|-------------------------|--------------|--|--------------|--------------------------------------|---|---|
| NMI/DI          | Source                  | Purity value | Expanded<br>uncertainty of<br>purity value | Traceability | Verification<br>Technique            | Purity assay  | Internal<br>Standard                            |
| HSA             | HSA (HRM-<br>1004A)     | 999.2 mg/g   | 3.3 mg/g                                   | HSA          |                                      |   | <sup>13</sup> C <sub>6</sub> -n-Buty<br>paraben |
| KRISS           | Dr.<br>Ehrenstorfer     | 999.92 mg/g  | 0.7 mg/g                                   | KRISS        | LC/MS, LC-UV, K-<br>F titration, TGA | Purity assayed by<br>KRISS with<br>chromatographic<br>purity, water<br>contents and non-<br>volatile & volatile<br>residues | <sup>13</sup> C <sub>6</sub> -n-Buty<br>paraben |
| NIM             | NIM CRM<br>GBW(E)100077 | 997 mg/g     | 1.994 mg/g<br>(0.2% , <i>k</i> =2)         | NIM          |                                      |   | <sup>13</sup> C <sub>6</sub> -n-Buty<br>paraben |
| RCChem-<br>LIPI | HSA,<br>Singapore       | 999.2 mg/g   | 3.3 mg/g                                   | HSA          |                                      |   |   |
| TISTR           | HSA (HRM-<br>1004A)     | 999.2 mg/g   | 3.3 mg/g                                   | HSA          |                                      |   | n-Butyl<br>paraben-d,                           |
| UME             | Alfa Aesar              | 0.99990      | 0.00313                                    | TÜBİTAK UME  | NMR and Mass<br>Spectrometry         | QNMR  | Butyl (C <sub>4</sub> D <sub>9</sub><br>paraben |

Table 14: Reference materials and internal standards used for the determination of n-butyl paraben

# 8.3 EXPERIMENTAL DETAILS

The participating NMIs/DIs used their own methods for the determination of benzoic acid, methyl paraben and n-butyl paraben. Details of the sample treatment, instrumental methods, preparation of calibration solutions and blends, and calibration methods are summarised in Tables 15 and 16. Generally, the participating NMIs/DIs performed simple dilution, liquid-liquid extraction or filtration before analyses.

For the determination of benzoic acid, gas chromatography coupled with mass spectrometry (GC-MS) analyses were carried out by INMETRO, GLHK and HSA; liquid chromatography coupled with tandem mass spectrometry (LC-MS/MS) analyses were carried out by EXHM/GCSL-EIM, GLHK, KRISS, NIM, STD-ITDI, TISTR and UME; and liquid chromatography coupled with diode array detection (LC-DAD) was carried out by RCChem-LIPI.

For the determination of methyl paraben, GC-MS analyses were carried out by HSA, LC-MS/MS analyses were carried out by KRISS, NIM, TISTR and UME; and LC-DAD was carried out by RCChem-LIPI.

For the determination of n-butyl paraben, GC-MS analyses were carried out by HSA and RCChem-LIPI; and LC-MS/MS analyses were carried out by KRISS, NIM, TISTR and UME.

Various GC columns (VF-1MS, DB-5MS and CP-Wax52 CB) have been used. Also, various LC columns (BEH Shield RP18, 100 SIELC PrimeSep 100, Poroshell HPH-C18, Hypersil 5µ C18 Phenomenex, AQUITY UPLC BEH C18, Agilent Zorbax Eclipse Plus C18, Porpshell 120, EC-C18 and Phenomenex Luna 5µ C18) were used.

Isotope dilution mass spectrometry with single-point, bracketing and multi-level calibration methods for quantification has been employed. EXHM/GCSL-EIM and TISTR used matrix-matched calibration blends for quantification. Most of the participating NMIs/DIs reported the use of qualifying ions/MRM transitions for verification of measurements.

Table 15: Participating NMIs/DIs' sample treatment and instrumental methods (BA denotes benzoic acid; MP denotes methyl paraben; and BP denotes n-butyl paraben)

| NMI/DI            | Sample                    | Extraction/Clean-up  | Instrument                       | Chromatographic                                      | Detection<br>(Ionisation               | Ions/MRM (Q: Quantify                     | ving; q: qualifying)                  |
|-------------------|---------------------------|--|----------------------------------|--|--|---|---------------------------------------|
|                   | size (g)                  | Extraction/Clean-up  | used                             | column   | mode for<br>MS)                        | Analyte                                   | Internal standard                     |
| EXHM/GC<br>SL-EIM | 1                         | Addition of internal standard,<br>dilution with mobile phase,<br>sonication then filtration.   | LC-MS/MS                         | 100 SIELC<br>PrimeSep 100<br>(250mm x 2.1mm,<br>5µm) | MS [ESI<br>(negative)]                 | BA: 121/77                                | BA: 126/82                            |
| GLHK              |                           | liquid-liquid extraction with LC-MS/MS DE<br>diethyl ether, drying then 0<br>reconstitution with CHCl <sub>3</sub> (for<br>GC-MS) or mobile phase (for<br>LC-MS/MS). Derivatisation LC-MS/MS (for GC-MS) | GC: Agilent J&W<br>DB-5MS (30m x | MS [EI for<br>GC-MS, ESI                             | BA (GC-MS): 179<br>(Q),105 (q)         | BA (GC-MS): 185<br>(Q), 111 (q)           |                                       |
|                   |                           |  |                                  | 0.25mmID x<br>0.25µm)                                | (negative) for<br>LC-MS/MS]            | BA (LC-MS/MS): 121/77<br>(Q), 121/121 (q) | BA (LC-MS/MS):<br>127/83 (Q)          |
|                   |                           |  | AS). Derivatisation              | LC: Poroshell<br>HPH-C18 2.1 x<br>50mm, 2.7µm        |  |   | ,co(u)                                |
| HSA               | 1                         | Addition of internal standard, GC-MS acidification with 5% HCl, dilution with $CH_3CN$ , liquid-liquid extraction with NaCl, centrifugation, exchange with   | GC-MS                            | CP-Wax 52 CB<br>(30m x                               | MS (EI)                                | BA: 105.0 (Q), 77.0 (q),<br>122.0 (q)     | BA: 111.0 (Q), 83.0<br>(q), 128.0 (q) |
|                   |                           |  |                                  | 0.25mmID,<br>0.25µm)                                 |  | MP: 152.0 (Q), 93.0 (q),<br>121.0 (q)     | MP: 158.0 (Q), 99.0<br>(q), 127.0 (q) |
|                   | $CH_3CN$ then filtration. |  |                                  |  | BP: 138.0 (Q), 194.0 (q),<br>121.0 (q) | BP: 144.0 (Q),<br>200.0 (q), 127.0 (q)    |                                       |

| INMETRO         | 1    | Addition of internal standard,<br>liquid-liquid extraction with<br>NaCl and ethyl ether,<br>centrifugation, drying of<br>organic extract under nitrogen<br>and vacuum at 50 °C for 40<br>min, followed by derivatisation<br>with MSTFA. | GC-MS                      | VF-1MS (30m x<br>0.25mmID,<br>0.25µm)                          | MS (EI, scan<br>mode: m/z<br>50 to 300) | BA: 179 (Q), 194 (q)                            | BA: 184 (Q), 199<br>(q)               |
|-----------------|------|---|----------------------------|--|---|---|---------------------------------------|
| KRISS           | 1    | Addition of internal standard,  | LC-MS/MS                   | AQUITY UPLC  | MS [ESI                                 | BA: 121/77                                      | BA: 127/83                            |
|                 |      | dilution with ammonium acetate, cleanup with SPE,   |                            | BEH C18, 2.1 x<br>10 mm, 1.7 μm                                | (negative)]                             | MP: 151/92                                      | MP: 157/98                            |
|                 |      | drying, reconstitution with mobile phase then filtration.   |                            | · • · · · · · · · · · ·  |   | BP: 193/92                                      | BP: 199/98                            |
| NIM             | ,    | BEH Shield RP18   | MS [ESI                    | BA: 121.1/77.1 (Q),  | BA: 127.1/83.1(Q)                       |   |                                       |
|                 |      | dilution with CH <sub>3</sub> CN,<br>ultrasonication, centrifugation,<br>dilution of supernatant with<br>water, then filtration.  |                            | (1.7 µm, 2.1   | (negative)]                             | 121.1/103.1 (q)                                 | 127.1/109.1 (q)                       |
|                 |      |   |                            | mm×100 mm)   |   | MP: 151.1/92.1 (Q),<br>151.1/136.1 (q)          | MP: 157.1/98.1 (Q)<br>157.1/142.1 (q) |
|                 |      |   |                            |  |   | BP: 193.2/92.1 (Q),<br>193.2/136.2 (q)          | BP: 199.2/98.1 (Q)<br>199.2/142.1 (q) |
| RCChem-<br>LIPI | 2    | Dilution, cleanup with SPE<br>then filtration (for HPLC-DAD)<br>and 5-fold dilution (for GC-  | BA and<br>MP: HPLC-<br>DAD | LC: Hypersil 5µ<br>C18 Phenomenex<br>4,6 x 150 mm              | DAD (BA:<br>230 nm; and<br>MP: 254 nm)  | BP (underivatised): 121<br>(Q), 138 (q), 93 (q) |                                       |
|                 | MS). |   | BP: GC-MS                  | GC: DB-5MS<br>(30m x 0.25mm x<br>0.25um)                       | BP: MS (EI)                             |   |                                       |
| STD-ITDI        | 0.2  | Dilution with mobile phase  | LC-MS                      | Agilent Zorbax<br>Eclipse Plus C18,<br>4.6 x 100 mm, 3.5<br>µm | MS [ESI<br>(negative)]                  | BA: 77.2  | BA: 82.1                              |

| TISTR | 1   | Extract with CH <sub>3</sub> OH, sonicate then filter | LC-MS/MS                       | Porpshell 120,<br>EC-C18 2.7<br>µm,2.1x100mm | MS [ESI<br>(negative)] | BA: 77<br>MP: 136 (Q), 92 (q)<br>BP: 92 (Q), 136 (q) | BA: 77<br>MP: 140 (Q), 96 (q)<br>BP: 96 (Q), 140 (q) |
|-------|-----|---|--------------------------------|--|------------------------|--|--|
| UME   | 1.2 | Liquid-liquid extraction with                         | LC-MS                          | Phenomenex                                   | MS [ESI                | BA: 120.7/77.0                                       | BA: 125.7/82.0                                       |
|       |     | methanol, filtration then dilution                    |                                | Luna 5µ C18<br>column (150 x 2               | (negative)]            | MP: 150.5/135.5                                      | MP: 153.7/135.5                                      |
|       |     |   | mm i.d., 5μm<br>particle size) |  | BP: 192.7/136.0        | BP: 202.0/136.0                                      |  |

Table 16: Participating NMIs/DIs' preparation of calibration solutions and blends, and calibration methods (continued)

|               | Calibration solutions and blends  |   | <b>.</b>   |  |
|---------------|---|---|--|--|
| NMI/DI        | Preparation   | Matrix/Solvent  | Calibration  |  |
| EXHM/GCSL-EIM | Dilution of multiple calibration solutions. Preparation of 10 different matrix-matched calibration blends.  | Matrix-matched  | IDMS (single-point, exact-matching)                              |  |
| GLHK          | Gravimetric preparation of multiple calibration solutions.<br>Preparation of multiple calibration blends.   | Methanol  | IDMS (bracketing)  |  |
| HSA           | Gravimetric preparation of multiple calibration solutions.<br>Preparation of multiple calibration blends.   | Methanol  | IDMS (single-point, exact-matching)                              |  |
| INMETRO       | Dilution of methanolic stock solutions of calibrants with water,<br>followed by addition of NaCl and internal standard to prepare<br>an eight-level calibration curve. Subjected solutions to same<br>treatment as the samples. | Water   | BA: IDMS (multi-level)<br>MP: Internal calibration (multi-level) |  |
| KRISS         | Gravimetric preparation of multiple calibration solutions.<br>Preparation of multiple calibration blends.   | 5 mmol/L ammonium<br>acetate buffer (pH4.1) /<br>methanol (v/v, 50/50)  | IDMS (single-point)  |  |
| NIM           | Dilution of multiple calibration solutions. Preparation of multiple calibration blends.   | CH <sub>3</sub> CN:H <sub>2</sub> O (1:1 v/v)                           | IDMS (single-point)  |  |
| RCChem-LIPI   | Dilution of multiple calibration solutions to prepare five-point calibration curves.  | Methanol  | External (both HPLC-DAD and GC-MS analyses                       |  |
| STD-ITDI      | Preparation of two calibration blends each from four mixed<br>standard solutions of natural BA, MP and BP and mixed<br>solutions of their isotopically-labelled standards.  | Mobile phase diluent<br>(50:50,<br>methanol:ammonium<br>acetate buffer) | IDMS (single-point)  |  |

| TISTR | Gravimetric preparation of multiple calibration blends. | Exact matrix-matching calibration. | IDMS (bracketing, exact matrix-matching) |
|-------|---|------------------------------------|--|
| UME   | Preparation of multiple calibration solutions.          | Matrix calibration solution        | IDMS (single-point)                      |

# 9.0 MEASUREMENT EQUATION AND ESTIMATION OF MEASUREMENT UNCERTAINTY

The measurement equations and contributors to the measurement uncertainty (MU) budgets are summarised in Table 17 below. The full uncertainty evaluation reported by the participating NMIs/DIs is given in Appendix I.

Table 17: Participating NMIs/DIs' measurement equations and contributors to measurement uncertainty budgets.

| NMI/DI            | Measurement Equation  | Contributions to measurement uncertainty budget   |
|-------------------|---|---|
| EXHM/GCSL<br>-EIM | $\begin{split} w_{B,S} &= w_{B,C} \; \frac{m_{D,S}}{m_{B,S}} \times \frac{m_{B,C}}{m_{D,C}} \times \frac{R_S}{R_B} \\ w_{BS} &= \text{mass fraction of benzoic acid in test material, } (mg/kg); w_{BC} &= \text{mass fraction of benzoic acid in calibration solution, } (mg/kg); \\ m_{D,S} &= \text{mass of benzoic acid-d5 solution added to sample blend, } (g); m_{B,S} &= \text{mass of test material in sample blend, } (g); m_{B,C} &= \text{mass of benzoic acid solution added to calibration blend, } (g); m_{D,C} &= \text{mass of benzoic acid-d5 solution added to calibration blend, } (g); R_S &= \text{mass of benzoic acid-d5 solution added to calibration blend, } (g); R_S &= \text{mass of benzoic acid-d5 solution added to calibration blend, } (g); R_S &= \text{measured peak area ratio of the selected ions in the sample blend; } R_C &= \text{measured peak area ratio of the selected ions in the sample blend; } R_C &= \text{measured peak area ratio of the selected ions in the sample blend}; } \end{split}$ | <ul> <li>Method precision</li> <li>Mass fraction of benzoic acid in<br/>calibration solution</li> <li>Mass fraction of benzoic acid-d5<br/>solution added to sample blend</li> <li>Mass of test sample in sample blend</li> <li>Mass of benzoic acid solution added<br/>to calibration blend</li> <li>Mass of benzoic acid-d5 solution<br/>added to calibration blend</li> <li>Measured peak area ratio of the<br/>selected ions in the sample blend</li> <li>Measured peak area ratio of the<br/>selected ions in the calibration blend</li> </ul> |
|                   | Equation used to estimate measurement<br>uncertainty:<br>${}^{u(w_{BS})} = {}^{w_{BS}} \sqrt{\left(\frac{SDr/\sqrt{n}}{w_{BS}}\right)^{2} + \sum \left(c_{I}\frac{u(m_{I})}{m_{I}}\right)^{2} + \sum \left(c_{I}\frac{u(R_{I})}{R_{I}}\right)^{2} + \left(c_{I}\frac{u(w_{BC})}{w_{BC}}\right)^{2}}$ where <i>SD</i> <sub>r</sub> is the standard deviation under<br>reproducibility conditions, <i>n</i> the number of<br>determinations and <i>C</i> <sub>i</sub> the sensitivity   |   |
|                   | determinations and $C_j$ the sensitivity coefficients associated with each uncertainty component. The uncertainty of  |   |

the peak area ratios was considered to have been included in the estimation of

method precision.

| GLHK | $c_{X} = c_{Z} \cdot \frac{m_{Y}}{m_{X}} \cdot \frac{m_{Zc}}{m_{Yc}} \cdot \frac{R_{B}}{R_{Bc}}$ $C_{X}: \text{ mass fraction of native analyte in sample; } C_{Z}: \text{ mass fraction of the native standard used to prepare the calibration blend; } m_{Y}: \text{ mass of labelled standard added to sample; } m_{X}: \text{ mass of sample used; } m_{Zc}: \text{ mass of the native standard added to the calibration blend; } m_{Yc}: \text{ mass of the native standard added to the calibration blend; } m_{Yc}: \text{ mass of the native standard added to the calibration blend; } m_{Yc}: \text{ mass of the labelled standard added to the calibration blend; } R_{B}: \text{ measured ion ratio (native/labelled) of the sample blend; } R_{Bc}: \text{ measured ion ratio (native/labelled) of the calibration blend}$  | <ul> <li>Precision</li> <li>Method bias</li> <li>Purity of calibrants</li> </ul>   |
|------|--|--|
| HSA  | $C_x = C_z \cdot \frac{m_y \cdot m_{zc}}{m_x \cdot m_{yc}} \cdot \frac{R_y - R_B}{R_B - R_x} \cdot \frac{R_{Bc} - R_Z}{R_y - R_{Bc}}$ $C_z : \text{mass fraction of benzoic acid/methyl paraben/n-butyl paraben in the calibration standard solution used to prepare the calibration blend; m_{y'}: mass of internal standards solution added to the sample blend; m_{yc}: mass of internal standard solution added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: mass of standard solutions added to the calibration blend; m_{zc}: observed isotope abundance ratio in the study material; R_y: observed isotope abundance ratio in the calibration standard; R_{B}: observed isotope abundance ratio in the calibration blend; R_{Bc}: observed isotope abundance ratio in the calibration blend; R_{Bc}: observed isotope abundance ratio in the calibration blend; R_{Bc}: observed isotope abundance ratio in the calibration blend; R_{Bc}: observed isotope abundance ratio in the calibration blend; R_{Bc}: observed isotope abundance ratio in the calibration blend; R_{Bc}: observed isotope abundance ratio in the calibration blend.$ | <ul> <li>Method precision</li> <li>Bias in the result due to choice of instrument (GC-MS and LC-MS)</li> <li>bias in the result due to choice of ion pair</li> <li>Method recovery</li> <li>Concentration of calibration standard solution (purity of calibration standard, weighing, different calibration blends)</li> <li>Mass of study material, standard solutions</li> </ul> |

## **INMETRO**

$$w_{a} = \left(\frac{A_{a}}{A_{IS}} - b0\right) \times \frac{m_{IS} \times m_{solIS} \times Pa}{b1 \times m_{sample} \times m_{totalsolIS}}$$

( 1

$$C_{X} = F_{p} \cdot F_{i} \cdot F_{ip} \cdot F_{r} \cdot C_{Z} \cdot \frac{m_{Y} \cdot m_{Zc}}{m_{X} \cdot m_{Yc}} \cdot \frac{R_{Y} - R_{B}}{R_{B} - R_{X}} \cdot \frac{R_{Bc} - R_{Z}}{R_{Y} - R_{Bc}}$$

 $w_a$  = Mass of anlyte;  $A_a$  = Area of analyte signal;  $A_{lS}$  = Area of internal standard signal; b0= Linear coefficient;  $m_{ls}$ = Mass of internal standard weighted in the stock solution:  $m_{solls}$  = Mass of internal standard solution added in the sample; Pa= Purity of analyte; b1= Angular coefficient;  $m_{sample}$  = Mass of sample weighted;  $m_{totalsollS}$  = Total mass of internal standard stock solution.

Subsequently,  $w_a$  was multiplied by 1000000 in order to express the mass fraction in mg/kg.

### Area ratio

- Mass of sample
- Mass of internal standard solution
- Angular coefficient (b1)
- Calibration Curve
- Linear coefficient (b0)
- Purity of calibrant

## **KRISS**

$$\mathbf{C}_{\text{sample}} = \frac{\mathbf{M}_{\text{is-solspiked}} \cdot \mathbf{AR}_{\text{sample}} \cdot \mathbf{M}_{\text{s-solstd.mix.}} \cdot \mathbf{C}_{\text{s-sol}}}{\mathbf{M}_{\text{sample}} \cdot \mathbf{AR}_{\text{std.mix.}} \cdot \mathbf{M}_{\text{is-solstd.mix.}}}}$$

 $C_{\text{sample:}}$  is the concentration of analytes in the sample;  $C_{s-sol}$  is the concentration of the analytes standard solution;  $M_{\text{sample}}$  is the mass of the sample taken for analysis;  $M_{is}$ . sol. spiked: is the mass of the isotope standard solution added to the sample aliquot;  $M_{is-sol}$ . std. mix .: is the mass of the isotope standard solution added to the isotope ratio standard solution; M<sub>s-sol, std. mix.:</sub> is the mass of the standard solution added to the isotope ratio standard solution; AR<sub>sample</sub>: is the area ratio of analyte/isotope for sample extract, observed by GC/MS; AR<sub>std. mix.</sub> is the area ratio of analyte/isotope for the isotope ratio standard solution, observed by GC/MS.

Combined standards uncertainties:

$$u_{total.} = \sqrt{u_{systematic}^2 + \frac{s^2}{n}}$$

Systematic (*u*<sub>systematic</sub>)

- Uncertainty of purity assay
- Uncertainty of gravimetric preparation for standard solutions
- Uncertainty of gravimetric mixing for calibration isotope standard mixtures
- · Uncertainty of observed area ratio of calibration isotope standard mixture by LC/MS (AR<sub>std. mix</sub>)

Random(*u*<sub>random</sub>)

- Uncertainty of observed area ratio of each subsample by LC/MS (AR<sub>sample</sub>). The uncertainty is included into s.
- · Uncertainties of weighing sample  $(M_{\text{sample}})$  and isotope standard solution spiked to the sample  $(M_{is-sol})$ spiked). The uncertainties are included into s.

Method precision

 Standard deviations (s) of multiple measurement results from five subsamples (n=5). It includes u<sub>random</sub>. Therefore, *u<sub>random</sub>* is not combined into utotal.

| NIM             | $X = \left(\frac{A_s}{A_i}\right)_s \times \left(\frac{A_i}{A_s}\right)_r \times \left(\frac{M_s}{M_i}\right)_r \times \frac{(M_i)_s}{W}$<br>X: Content of measurands in soy sample (mg/kg); $\left(\frac{A_s}{A_i}\right)_s$ : Peak area ratio of measurand and internal standard in sample solution; $\left(\frac{A_i}{A_s}\right)_r$ : Peak area ratio of internal standard and measurand in standard solution; $\left(\frac{M_s}{M_i}\right)_r$ : Mass ratio of measurand and internal standard solution; $\left(\frac{M_s}{M_i}\right)_r$ : Mass ratio of measurand and internal standard solution; $\left(\frac{M_s}{M_i}\right)_r$ : Mass of internal standard added in soy sample (µg); <i>w</i> : Mass of soy sample (g).  | <ul> <li>Method precision (between batch precision for the method as a whole)</li> <li>Standard solution (purity of standard solution, weight of standard solution, weight of internal standard solution, weight of solvent)</li> <li>Sample pretreatment (weight of sample, weight of internal standard solution, weight of solvent)</li> </ul>   |
|-----------------|---|--|
| RCChem-<br>LIPI | HPLC:<br>$X = C_{HPLC} * \frac{m_{centrifuge}}{m_{sample}} * \frac{m_{hplc}}{m_{spe}} * \frac{1}{\text{Re}c}$ GC-MS:<br>$X = C_{GCMS} * \frac{m_{centrifuge}}{m_{sample}} * \frac{m_{hplc}}{m_{spe}} * \frac{m_{final}}{m_{200ul}} * \frac{1}{\text{Re}c}$ X: mass fraction of the analyte (mg/kg);<br>C <sub>HPLC</sub> : Concentration analyte from<br>calibration curve in the HPLC system;<br>C <sub>GCMS</sub> : Concentration analyte from<br>calibration curve in the GCMS system;<br>M <sub>centrifuge</sub> : mass of solution of sample in<br>water; M <sub>sample</sub> : mass of 2 ml of sample;<br>M <sub>hplc</sub> : mass of final methanol solution after<br>celan up with SPE; M <sub>SPE</sub> :mass of 1 ml<br>aliquot of sample in water solution that is<br>pass through in to the SPE cartridge; M <sub>final</sub> :<br>mass of the final solution for analysis with<br>GCMS<br>M <sub>200ul</sub> : mass of the 200 uL of final methanol<br>solution aliquot to be diluted<br>Rec : recovery factor | <ul> <li>HPLC:</li> <li>Calibration curve</li> <li>Mass of final methanol solution</li> <li>Mass of 1 ml aliquot of sample in water solution that is pass through in to the SPE cartridge</li> <li>Mass of sample</li> <li>Mass of solution of sample in water</li> <li>Recovery</li> <li>Repeatability sample</li> <li>Middle standard solution</li> <li>GC-MS</li> <li>Mass of solution of sample in water</li> <li>Mass of solution of sample in water</li> <li>Mass of solution of sample in water</li> <li>Mass of solution for sample in water</li> <li>Mass of solution of sample in water</li> <li>Mass of solution that is pass through in to the SPE</li> <li>Mass of final methanol solution after celan up with SPE</li> <li>Mass of the 200 uL of final methanol solution to be diluted</li> <li>Mass of the final solution for analysis with GCMS</li> <li>Presision</li> <li>Calibration curve</li> <li>Pacewory</li> </ul> |

- Recovery
- Middle standard solution

## STD-ITDI

TISTR

$$C_{Measurand} = \frac{m_{is-sp} \ x \ m_{std} \ x \ AR_{sample} \ x \ C_{stock,std}}{m_{sample} \ x \ AR_{std} \ x \ m_{is-sp,std}}$$

 $C_{Measurand}$ :concentration of measurand;  $m_{is}$ .  $_{sp,sample}$ : mass of isotope solution added to the sample solution;  $m_{std}$ : mass of standard stock solution in the standard-isotope calibration blend;  $AR_{sample}$ : observed response ratio of the standard/isotope in the sample;  $c_{stock,std}$ : concentration of the standard stock solution;  $m_{sample}$ : mass of the sample taken for analysis;  $AR_{std}$ : observed response ratio of standard/isotope in the standard-isotope calibration blend;  $m_{is-sp,std}$ : mass of isotope stock solution in the standard-isotope blend

$$C_{X} = C_{Z} \cdot \frac{M_{Y} \cdot M_{Zc}}{M_{X} \cdot M_{Yc}} \cdot \frac{R'_{B}}{R'_{Bc}}$$

For uncertainty estimation:

$$w_x = F_P \cdot F_E \cdot w_{z,c} \cdot \frac{m_y \cdot m_{z,c}}{m_x \cdot m_{y,c}} \cdot \frac{R'_b}{R'_{b,c}}$$

 $w_x =$  mass fraction of analyte in sample;  $w_{Z,C} =$  mass fraction of analyte in the calibration solution used to prepare the calibration blend;  $m_Y =$  mass of spike solution added to sample blend;  $m_{Y,C} =$ mass of spike solution added to calibration blend;  $m_x =$ mass of sample added to sample blend;  $m_{Z,C} =$  mass of solution added to calibration blend;  $R_B, R_{B,C} =$ observed isotope amount ration in the sample blend and the calibration blend, respectively;  $F_P =$  method precision factor;  $F_E =$  Extraction efficiency factor

- Concentration of standard stock solution purity and gravimetric preparation)
- Calibration standard mixture (gravimetric mixing of blends)
- · Mass of sample for analysis
- Mass of internal standard in the sample
- Peak area ratio of analyte and internal standard in calibration blend
- Peak area ratio of analyte and internal standard in the sample

Benzoic acid

- Method precision
- Weighing
- Mass fraction of analyte in the calibration solution
- R'b and R'bc
- Extraction effects/matrix effects

Methyl paraben and n-butyl paraben

- Method precision
- Weighing
- Mass fraction of analyte in the calibration solution
- R'b and R'bc
- Matrix effects
- Interference from two different ion pairs

| UME | C <sub>sample</sub>  | <ul> <li>Weighing of sample</li> </ul>            |
|-----|--|---|
|     | $=\frac{Area_{sample}}{C_{Std}}\frac{Area_{STD-IS}}{C_{STD-IS}}C_{STD-IS}$     | <ul> <li>Weighing of internal standard</li> </ul> |
|     | $= \frac{1}{Area_{IS}} \frac{1}{C_{IS}} \frac{1}{Area_{Sample}} C_{Sample-IS}$ | <ul> <li>Native stock solution</li> </ul>         |
|     | 15 16 Sumple   | <ul> <li>Labelled stock solution</li> </ul>       |
|     |  | <ul> <li>Intermediate precision</li> </ul>        |
|     |  | Recovery  |
|     |  | Repeatability                                     |

# 10.0 SUPPLEMENTARY COMPARISON REFERENCE VALUES

Table 18 summarises the proposed supplementary comparison reference values (SCRVs) and their associated standard uncertainties for the three measurands. Two statistical approaches were undertaken, using all results submitted in the supplementary comparison.

The proposed SCRVs were presented at the 2016 CCQM OAWG meeting in April and 2016 APMP TCQM meeting in November. The meetings agreed to the use of the median approach for calculation of the SCRVs for all three measurands as there was no significant difference between the two approaches.

|   | Measurands   |                |                 |
|---|--------------|----------------|-----------------|
| ·   | Benzoic acid | Methyl paraben | n-Butyl paraben |
| No. of results ( <i>N</i> )                         | 10           | 6              | 6               |
| Approach 1:   |              |                |                 |
| Arithmetic mean                                     | 153.49       | 100.06         | 99.55           |
| Standard deviation (SD)                             | 2.81         | 1.94           | 2.68            |
| Standard uncertainty (SD/ $\sqrt{N}$ )              | 0.89         | 0.79           | 1.09            |
| Approach 2:   |              |                |                 |
| Median  | 154.55       | 100.95         | 99.05           |
| Median absolute deviation (MAD)                     | 1.61         | 0.53           | 1.80            |
| 1.483×MAD (MAD <sub>e</sub> )                       | 2.39         | 0.78           | 2.67            |
| Standard uncertainty<br>(1.25×MAD <sub>e</sub> /√N) | 0.94         | 0.40           | 1.36            |

Table 18: Proposed SCRVs and associated uncertainties calculated by arithmetic mean and median approaches

The participating NMIs/DIs' results with associated reported standard uncertainties, as well as the SCRV (median) and their associated standard uncertainty  $(1.25 \times MAD_e/\sqrt{N})$  for





Figure 1: SCRV (solid red line) and its standard uncertainty (red dotted line) for benzoic acid, with participating NMIs/DIs' results and the associated reported standard uncertainties



Figure 2: SCRV (solid red line) and its standard uncertainty (red dotted line) for methyl paraben, with participating NMIs/DIs' results and the associated reported standard uncertainties



Figure 3: SCRV (solid red line) and its standard uncertainty (red dotted line) for n-butyl paraben, with participating NMIs/DIs' results and the associated reported standard uncertainties

# 11.0 DEGREES OF EQUIVALENCE (DOE) CALCULATION

The degree of equivalence ( $D_i$ ) for each participating NMI/DI's result was calculated using the SCRV for each measurand. The  $D_i$  was calculated as follows:

$$D_i = X_i - X_{SCRV}$$

where  $X_i$  is the reported result of a participating NMI/DI; and  $X_{SCRV}$  is the SCRV.

The uncertainty associated with D<sub>i</sub> for each participating NMI/DI was estimated as follows:

$$u(D_i) = \sqrt{u^2(X_i) + u^2(X_{SCRV})}$$

The expanded uncertainty of the  $D_i$ , i.e.  $U(D_i)$  was calculated as:

$$U(D_i) = \sqrt{k_i^2 \times u^2(X_i) + k_{SCRV}^2 \times u^2(X_{SCRV})}$$

where  $k_i$  is the coverage factor reported by the participating NMI/DI; and  $k_{SCRV}$  is the coverage factor of the SCRV calculated from t(0.05; degrees of freedom) at 95% confidence level.

Tables 19 to 21 summarise the participating NMIs/DIs'  $D_i$  (mg/kg),  $U(D_i)$  (mg/kg) and  $D_i/U(D_i)$  for the three measurands. Figures 4 to 6 present the plots of  $D_i$  and their expanded uncertainties  $[U(D_i)]$  with the respective  $k_{SCRV}$  at 95% confidence level for the three measurands.

| NMI/DI        | Benzoic acid |          |              |
|---------------|--------------|----------|--------------|
|               | $D_i$        | $U(D_i)$ | $D_i/U(D_i)$ |
| INMETRO       | -5.95        | 3.98     | -1.49        |
| TISTR         | -4.05        | 8.93     | -0.45        |
| EXHM/GCSL-EIM | -3.89        | 4.57     | -0.85        |
| UME           | -2.17        | 7.24     | -0.30        |
| NIM           | -0.05        | 3.85     | -0.01        |
| GLHK          | 0.05         | 5.25     | 0.01         |
| KRISS         | 0.48         | 4.80     | 0.10         |
| HSA           | 0.95         | 5.25     | 0.18         |
| STD-ITDI      | 1.05         | 4.36     | 0.24         |
| RCChem-LIPI   | 2.95         | 8.09     | 0.36         |

Table 19: Participating NMIs/DIs'  $D_i$  (mg/kg), U( $D_i$ ) (mg/kg) and  $D_i$  / U( $D_i$ ) for benzoic acid



Figure 4: Plot of degrees of equivalence ( $D_i$ ) and their expanded uncertainties [ $U(D_i)$ ] with k=2.26 at 95% confidence level for benzoic acid.

| NMI/DI      | Methyl paraben |          |              |
|-------------|----------------|----------|--------------|
|             | $D_i$          | $U(D_i)$ | $D_i/U(D_i)$ |
| RCChem-LIPI | -4.35          | 6.48     | -0.67        |
| TISTR       | -2.05          | 6.06     | -0.34        |
| HSA         | -0.05          | 2.25     | -0.02        |
| NIM         | 0.05           | 2.43     | 0.02         |
| UME         | 0.51           | 5.46     | 0.09         |
| KRISS       | 0.54           | 2.66     | 0.20         |

Table 20: Participating NMIs/DIs'  $D_i$  (mg/kg), U( $D_i$ ) (mg/kg) and  $D_i$  / U( $D_i$ ) for methyl paraben



Figure 5: Plot of degrees of equivalence ( $D_i$ ) and their expanded uncertainties [ $U(D_i)$ ] with k=2.57 at 95% confidence level for methyl paraben.
| NMI/DI |       | n-Butyl paraben |              |  |  |  |  |  |
|--------|-------|-----------------|--------------|--|--|--|--|--|
|        | $D_i$ | $U(D_i)$        | $D_i/U(D_i)$ |  |  |  |  |  |
| HSA    | -1.85 | 4.25            | -0.44        |  |  |  |  |  |
| NIM    | -1.75 | 4.36            | -0.40        |  |  |  |  |  |
| LIPI   | -1.75 | 7.83            | -0.22        |  |  |  |  |  |
| TISTR  | 1.75  | 8.22            | 0.21         |  |  |  |  |  |
| KRISS  | 2.04  | 4.52            | 0.45         |  |  |  |  |  |
| UME    | 4.53  | 6.55            | 0.69         |  |  |  |  |  |



Figure 6: Plot of degrees of equivalence ( $D_i$ ) and their expanded uncertainties [ $U(D_i)$ ] with k=2.57 at 95% confidence level for n-butyl paraben.

## 12.0 CORE COMPETENCY AND HOW FAR DOES THE LIGHT SHINE?

This comparison covered a subject which is of wide interest and importance. It enabled participating NMIs/DIs to demonstrate their measurement capabilities in the determination of common preservatives in soy sauce, using procedure(s) that required simple sample preparation and selective detection in the mass fraction range from 50 to 1,000 mg/kg. The APMP TCQM has earlier agreed that the study can be extended to include other polar food preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous-based beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based condiments (e.g. vinegar, fish sauce, etc).

The Core Competency Tables of the participating NMIs/DIs for the measurands reported in APMP.QM-S8 are presented in APPENDIX II.

## 13.0 USE OF REPORT

This report is intended to be used as an internal reference for the participating NMIs/DIs and APMP TCQM. Its content shall not be disclosed to other parties or used for other purposes.

#### ACKNOWLEDGEMENT

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## APPENDIX I

## Full Uncertainty Evaluation Reported By the Participating NMIs/DIs

### EXHM/GCSL-EIM

The equation used to estimate standard uncertainty is :

$$u(w_{BS}) = w_{BS} \sqrt{\left(\frac{SDr}{\sqrt{n}} \frac{1}{w_{BS}}\right)^{2} + \sum \left(C_{j} \frac{u(m_{i})}{m_{i}}\right)^{2} + \sum \left(C_{j} \frac{u(R_{i})}{R_{i}}\right)^{2} + \left(C_{j} \frac{u(w_{BC})}{w_{BC}}\right)^{2}}$$

where  $SD_r$  is the standard deviation under reproducibility conditions, *n* the number of determinations and  $C_j$  the sensitivity coefficients associated with each uncertainty component. The uncertainty of the peak area ratios was considered to have been included in the estimation of method precision.

Uncertainty estimation was carried out according to JCGM 100: 2008. The standard uncertainties were combined as the sum of the squares of the product of the sensitivity coefficient (obtained by partial differentiation of the measurement equation) and standard uncertainty to give the square of the combined uncertainty. The square root of this value was multiplied by a coverage factor (95% confidence interval) from the t-distribution at the total effective degrees of freedom obtained from the Welch-Satterthwaite equation to give the expanded uncertainty.

| uncertainty component  | value   | sensitivity<br>coefficient | standrard<br>uncertainty | relative<br>uncertainty | $C_i \times u_i$ | $(C_i \times u_i)^2$ |
|--|---------|----------------------------|--------------------------|-------------------------|------------------|----------------------|
| method precision   | 1,00    | 150,660                    | 0,0042                   | 0,0042                  | 0,637            | 0,406273             |
| mass fraction of benzoic acid in calibration solution, $(mg/kg)$       | 106,71  | 1,412                      | 0,2832                   | 0,0027                  | 0,3998           | 0,159826             |
| mass of benzoic acid-d5 solution added to sample blend, $(g)$          | 1,53614 | 98,077                     | 6,52E-05                 | 4,25E-05                | 0,0064           | 0,000041             |
| mass of test material in sample blend, (g )                            | 1,01911 | -147,835                   | 6,21E-05                 | 6,10E-05                | 0,0092           | 0,000084             |
| mass of benzoic acid solution added to calibration blend, (g)          | 1,49937 | 100,482                    | 6,50E-05                 | 4,34E-05                | 0,0065           | 0,000043             |
| mass of benzoic acid-d5 solution added to calibration blend, ( $g$ )   | 1,54226 | -97,688                    | 6,53E-05                 | 4,23E-05                | 0,0064           | 0,00004              |
| measured peak area ratio of the selected ions in the sample blend      | 0,9075  | 166,024                    | 0,0048                   | 0,0053                  | 0,7950           | 0,632054             |
| measured peak area ratio of the selected ions in the calibration blend | 0,9419  | -159,950                   | 0,0085                   | 0,0090                  | 1,3596           | 1,848609             |
| mass fraction of benzoic acid in soy sauce sample (mg/kg)              | 150,66  |                            |                          |                         |                  |                      |
| combined standard uncertainty (mg/kg)                                  | 1,75    |                            |                          |                         |                  |                      |
| effective degrees of freedom   | 7,65    |                            |                          |                         |                  |                      |
| coverage factor (k, p=0,95)  | 2,31    |                            |                          |                         |                  |                      |
| expanded uncertainty (mg/kg)   | 4,03    |                            |                          |                         |                  |                      |

# GLHK

|                               | Benzoic acid |
|-------------------------------|--------------|
| Precision                     | 0.81%        |
| Method Bias                   | 1.26%        |
| Purity of calibrants          | 0.33%        |
| Combined standard uncertainty | 1.53%        |
| Coverage factor, k            | 2            |
| Expanded uncertainty          | 3.07%        |

# HSA

$$C_{X} = F_{p} \cdot F_{i} \cdot F_{ip} \cdot F_{r} \cdot C_{Z} \cdot \frac{m_{Y} \cdot m_{ZC}}{m_{X} \cdot m_{YC}} \cdot \frac{R_{Y} - R_{B}}{R_{B} - R_{X}} \cdot \frac{R_{BC} - R_{Z}}{R_{Y} - R_{BC}}$$

#### where

additional factors (F) contributing to biases in the result value of benzoic acid or methyl paraben or nbutyl paraben were included by assigning a mean value of 1, with an associated uncertainty value to this mean value.

- $F_p$  = Factor representing method precision
- $F_i$  = Factor representing any bias in the result due to choice of instrument
- $F_{ip}$  = Factor representing any bias in the result due to choice of ion pair
- $F_r$  = Factor representing method recovery

The full uncertainty budget for the determination of benzoic acid is given in the Table below:



| Parameter             | X <sub>i</sub> | U <sub>xi</sub> | u <sub>xi</sub> /x | Contribution | Sources of uncertainty   |
|-----------------------|----------------|-----------------|--------------------|--------------|--|
| Fp                    | 1              | 0.004540        | 0.4543<br>%        | 8.5851%      | Standard deviation of the mean of 9<br>independent determinations on the study<br>material   |
| Fi                    | 1              | 0.006882        | 0.6882<br>%        | 19.6977%     | Comparison of results obtained using<br>GC-MS and LC-MS  |
| F <sub>ip</sub>       | 1              | 0.002984        | 0.2984<br>%        | 3.7039%      | Comparison of results obtained using different ion pairs   |
| Fr                    | 1              | 0.011866        | 1.1866             | 58.5625%     | Uncertainty in method recovery   |
| Cz                    | 100.93<br>71   | 0.480540        | 0.4761<br>%        | 9.4275%      | <ul> <li>Uncertainty in the purity value of<br/>benzoic acid certified reference<br/>material (HRM-1002A)</li> <li>Uncertainty in weighing based on value<br/>from the in-house balance calibration<br/>certificate</li> </ul> |
| m <sub>Y</sub>        | 0.5517         | 0.000071        | 0.0128             | 0.0068%      |  |
| m <sub>Yc</sub>       | 0.5371         | 0.000071        | 0.0132             | 0.0072%      | Uncertainty based on value from the  |
| m <sub>Zc</sub>       | 0.5366         | 0.000071        | 0.0132             | 0.0072%      | in-house balance calibration certificate   |
| m <sub>X</sub>        | 1.0304         | 0.000071        | 00069              | 0.0020%      |  |
| $R_{X_i} R_{Y_i} R_Z$ | Uncertair      | nty included i  | n method p         | precision    |  |
| $R_{B_{c}}R_{Bc}$     | Negligible     | e               |                    |              |  |

|                     |             |                 |                    | Precisio<br>0.7531 |   |
|---------------------|-------------|-----------------|--------------------|--------------------|---|
|                     |             |                 | overy<br>615%      |                    | 0.0193%<br>Mx<br>0.0052%<br>Cz<br>24.3074%<br>Interference<br>& Matrix<br>Instrument 8.1697%<br>9.9464%   |
| Parameter           | Xi          | U <sub>xi</sub> | u <sub>xi</sub> /x | Contribution       | Sources of uncertainty  |
| $F_{ ho}$           | 1           | 0.000824        | 0.0824<br>%        | 0.7531%            | Standard deviation of the mean of 9<br>independent determinations on the study<br>material  |
| Fi                  | 1           | 0.002993        | 0.2993<br>%        | 9.9464%            | Comparison of results obtained using<br>GC-MS and LC-MS   |
| F <sub>ip</sub>     | 1           | 0.002712        | 0.2712<br>%        | 8.1697%            | Comparison of results obtained using different ion pairs  |
| Fr                  | 1           | 0.007150        | 0.7150             | 56.7615%           | Uncertainty in method recovery  |
| Cz                  | 63.689<br>3 | 0.297979        | 0.4679<br>%        | 24.3074%           | <ul> <li>Uncertainty in the purity value of<br/>methyl paraben certified reference<br/>material (HRM-1003A)</li> <li>Uncertainty in weighing based on value<br/>from the in-house balance calibration<br/>certificate</li> <li>Comparison of results obtained from<br/>different calibration blends bracketing<br/>the same sample blend</li> </ul> |
| m <sub>Y</sub>      | 0.5517      | 0.000071        | 0.0128             | 0.0182%            |   |
| m <sub>Yc</sub>     | 0.5371      | 0.000071        | 0.0132             | 0.0192%            | Uncertainty based on value from the   |
| m <sub>Zc</sub>     | 0.5366      | 0.000071        | 0.0132             | 0.0193%            | in-house balance calibration certificate  |
| m <sub>X</sub>      | 1.0304      | 0.000071        | 0.0069             | 0.0052%            |   |
| $R_{X,}R_{Y,}R_{Z}$ | Uncertair   | nty included i  | n method p         | precision          |   |

The full uncertainty budget for the determination of n-butyl paraben is given in the Table below:



| Fi                  | 1           | 0.004118       | 0.4118<br>% | 11.7743%  | Comparison of results obtained using<br>GC-MS and LC-MS   |
|---------------------|-------------|----------------|-------------|-----------|---|
| F <sub>ip</sub>     | 1           | 0.002413       | 0.2413<br>% | 4.0426%   | Comparison of results obtained using<br>different ion pairs   |
| Fr                  | 1           | 0.009861       | 0.9861      | 67.5253%  | Uncertainty in method recovery  |
| Cz                  | 67.082<br>5 | 0.285183       | 0.4251<br>% | 12.5515%  | <ul> <li>Uncertainty in the purity value of n-<br/>butyl paraben certified reference<br/>material (HRM-1004A)</li> <li>Uncertainty in weighing based on value<br/>from the in-house balance calibration<br/>certificate</li> <li>Comparison of results obtained from<br/>different calibration blends bracketing<br/>the same sample blend</li> </ul> |
| my                  | 0.5517      | 0.000071       | 0.0128      | 0.0114%   |   |
| m <sub>Yc</sub>     | 0.5371      | 0.000071       | 0.0132      | 0.0120%   | Uncertainty based on value from the   |
| mzc                 | 0.5366      | 0.000071       | 0.0132      | 0.0121%   | in-house balance calibration certificate  |
| m <sub>X</sub>      | 1.0304      | 0.000071       | 0.0069      | 0.0033%   |   |
| $R_{X_i}R_{Y_i}R_Z$ | Uncertair   | nty included i | n method p  | precision |   |
| $R_{B_i}R_{Bc}$     | Negligible  | e              |             |           |   |

## INMETRO

Uncertainty was firstly estimated during the analysis of each subsample. Tables 1 and 2 show examples of the determination of benzoic acid and methyl paraben, respectively, for one of the subsamples. The same is shown in Figures 1 and 2 in order to highlight the major contributions. Combined standard uncertainty was then calculated by the square root of the sum of the squares of the standard uncertainties calculated for each subsample. This resulted in 1.6 mg/kg for benzoic acid and for methyl paraben, as reported in item 1. Coverage factors were calculated considering the effective degrees of freedom.

| Sources of uncertainty                                     | Туре   | Distribution | Standard<br>uncertainty | Sensitivity<br>coefficient | Uncertainty<br>component |
|--|--------|--------------|-------------------------|----------------------------|--------------------------|
| Area ratio   | А      | Normal       | 0.00562311              | 0.00011228                 | 6.314E-07                |
| Mass of sample   | В      | Normal       | 0.000015                | -0.00014586                | 2.188E-09                |
| Mass of IS solution  | В      | Normal       | 0.000015                | 0.00014895                 | 2.234E-09                |
| Angular coefficient (b1)                                   | А      | Normal       | 0.00459928              | -0.00013970                | -                        |
| Linear coefficient (b0)                                    | А      | Normal       | 0.00478592              | -0.00011228                | -                        |
| Calibration Curve  | А      | Normal       | -                       | -                          | 2.327E-07                |
| Purity of calibrant  | В      | Normal       | 0.00072093              | 0.00014921                 | 1.076E-07                |
| Standard uncertainty                                       |        |              |                         |                            | 6.814E-07                |
| Standard uncertainty (mg/kg)                               |        |              |                         |                            | 0.6814                   |
| calibration curve<br>mass of IS solution<br>mass of sample |        |              |                         |                            |                          |
| area ratio   |        |              |                         |                            |                          |
| 0 E+0  | 0 1 E- | -07 2 E-07   | 3 E-07 4 E-0            | 75E-0761                   | E-07 7 E-07              |
|  |        | unc          | ertainty compo          | onent                      |                          |

# KRISS

Combined standards uncertainties were obtained by combining systematic uncertainties and random uncertainties, which can be experimentally estimated by the standard deviation (s) of measurement results of multiple subsamples, as shown below equation

$$u_{total.} = \sqrt{u_{systematic}^2 + \frac{s^2}{n}}$$

Detailed for the full uncertainty budget is provided the below table.

|   | -   |
|---|---|
|   | Sources   |
|   | Uncertainty of purity assay   |
|   | Uncertainty of gravimetric preparation for standard solutions   |
| Systematic ( <i>u</i> <sub>systematic</sub> ) | Uncertainty of gravimetric mixing for calibration isotope standard mixtures   |
|   | Uncertainty of observed area ratio of calibration isotope standard mixture by LC/MS ( $AR_{std. mix}$ )                 |
|   | Uncertainty of observed area ratio of each subsample by   |
|   | LC/MS ( <i>AR</i> <sub>sample</sub> ). The uncertainty is included into s.  |
| Random( <i>u<sub>random</sub></i> )           | Uncertainties of weighing sample $(M_{sample})$ and isotope standard  |
|   | solution spiked to the sample ( $M_{is-sol, spiked}$ ). The uncertainties are included into s.                          |
|   | Standard deviations (s) of multiple measurement results from  |
| Method precision                              | five subsamples ( <i>n</i> =5). It includes $u_{random}$ . Therefore, $u_{random}$ . is not combined into $u_{total}$ . |

# NIM

|                        | Uncertainty b                                     | udget of benzoic acid                                 |             |          |  |
|------------------------|---|---|-------------|----------|--|
| Parameter              | Source of uncertainty                             | Source of data  | Xi          | Urel(Xi) |  |
| Method<br>precision    | Between batch precision for the method as a whole | Replicate analysis of<br>samples among six<br>batches | 154.5 mg/kg | 0.82%    |  |
|                        | Purity of standard solution                       | RM specification                                      | 1.00 mg/mL  | 0.5%     |  |
| Standard               | Weight of standard solution                       |   | 1 g         | 0.019%   |  |
| solution               | Weight of internal standard<br>solution           |   | 0.23g       | 0.083%   |  |
|                        | Weight of solvent                                 | Balance calibration                                   | 10 g        | 0.0019%  |  |
|                        | Weight of sample                                  | certificate   | 1 g         | 0.019%   |  |
| Sample<br>pretreatment | Weight of internal standard<br>solution           |   | 0.1 g       | 0.19%    |  |
|                        | Weight of solvent                                 |   | 8 g         | 0.0024%  |  |
| Combin                 | ed standard uncertainty                           | 0.98 %  |             |          |  |
| Relative ex            | xpanded uncertainty (k=2)                         | 2.0 %   |             |          |  |
|                        | Uncertainty Analysis Results                      |   |             |          |  |
|                        | Cx=   |   |             |          |  |
|                        | u(x) =  | 1.6 mg/kg   |             |          |  |
|                        | k=  | 2   |             |          |  |
|                        | U(x) =  | 3.1   | 2 mg/kg     |          |  |

|                                     | Uncertainty but                                   | dget of methyl paraber                                | า           |          |  |
|-------------------------------------|---|---|-------------|----------|--|
| Parameter                           | Source of uncertainty                             | Source of data  | Xi          | Urel(Xi) |  |
| Method<br>precision                 | Between batch precision for the method as a whole | Replicate analysis of<br>samples among six<br>batches | 101.0 mg/kg | 0.95%    |  |
|                                     | Purity of standard                                | RM specification                                      | 996 mg/g    | 0.1%     |  |
|                                     | Weight of standard                                |   | 0.01 g      | 0.19%    |  |
| Standard                            | Weight of standard solution                       |   | 1 g         | 0.019%   |  |
| solution                            | Weight of internal standard<br>solution           |   | 0.23g       | 0.083%   |  |
|                                     | Weight of solvent                                 | Balance calibration                                   | 10 g        | 0.0019%  |  |
|                                     | Weight of sample                                  | certificate   | 1 g         | 0.019%   |  |
| Sample<br>pretreatment              | Weight of internal standard<br>solution           |   | 0.1 g       | 0.19%    |  |
|                                     | Weight of solvent                                 |   | 8 g         | 0.0024%  |  |
| Combin                              | ed standard uncertainty                           | 1.0%  |             |          |  |
| Relative expanded uncertainty (k=2) |   | 2.0%  |             |          |  |
| Uncertainty Analysis Results        |   |   |             |          |  |
| Cx=                                 |   | 101.0 mg/kg   |             |          |  |
|                                     | u(x) =  | 1.1mg/kg  |             |          |  |
|                                     | k=  | 2   |             |          |  |
|                                     | U(x) =  | 2.1   | 2 mg/kg     |          |  |

|                              | Uncertainty but                                   | dget of n-Butyl parabe                                | n          |          |  |
|------------------------------|---|---|------------|----------|--|
| Parameter                    | Source of uncertainty                             | Source of data  | Xi         | Urel(Xi) |  |
| Method<br>precision          | Between batch precision for the method as a whole | Replicate analysis of<br>samples among six<br>batches | 97.3 mg/kg | 1.21%    |  |
|                              | Purity of standard                                | RM specification                                      | 997 mg/g   | 0.1%     |  |
|                              | Weight of standard                                |   | 0.01 g     | 0.19%    |  |
| Standard                     | Weight of standard solution                       |   | 1 g        | 0.019%   |  |
| solution                     | Weight of internal standard<br>solution           |   | 0.23g      | 0.083%   |  |
|                              | Weight of solvent                                 | Balance calibration                                   | 10 g       | 0.0019%  |  |
|                              | Weight of sample                                  | certificate   | 1 g        | 0.019%   |  |
| Sample<br>pretreatment       | Weight of internal standard<br>solution           |   | 0.1 g      | 0.19%    |  |
|                              | Weight of solvent                                 |   | 8 g        | 0.0024%  |  |
| Combin                       | ed standard uncertainty                           | 1.3%  |            |          |  |
| Relative ex                  | xpanded uncertainty (k=2)                         | 2.6%  |            |          |  |
| Uncertainty Analysis Results |   |   |            |          |  |
|                              | Cx=   | 97.3 mg/kg  |            |          |  |
|                              | u(x) =  | 1.3mg/kg  |            |          |  |
|                              | k=  |   | 2          |          |  |
|                              | U(x) =  | 2.  | 6 mg/kg    |          |  |

# **RCChem-LIPI**

| HPLC-DAD   | $C = \frac{C_{hplc} \times M_{hplc} x M_{centrifuge}}{M_{SPE} x M_{sample}} x \frac{1}{\text{Re } c}$  |
|--|--|
| $u_{combine} = C_{\sqrt{\left(\frac{u_{Chplc}}{C_{hplc}}\right)^2} + $ | $\left(\frac{u_{Mhplc}}{M_{hplc}}\right)^2 + \left(\frac{u_{MSPE}}{M_{SPE}}\right)^2 + \left(\frac{u_{Mcentrifug}}{M_{centrifuge}}\right)^2 + \left(\frac{u_{Msample}}{M_{sample}}\right)^2 + \left(\frac{u_{Rec}}{Rec}\right)^2 + \left(\frac{u_{rep}}{1}\right)^2 + \left(\frac{u_{Cstandard}}{C_{standard}}\right)^2$ |

#### **Benzoic Acid**

| Mass fraction (mg/kg) | 157.527 |
|-----------------------|---------|
| Replication           | 13      |
| sd                    | 4.942   |
| %RSD                  | 3.137   |

The uncertainty budget is summarized in the following table

| No     | Source of<br>uncertainty   | Symbol              | Value    | u           | Unit  | Relative<br>standard<br>uncertainty (%) |
|--------|--|---------------------|----------|-------------|-------|---|
| 1      | Calibration curve  | Co <sub>hplc</sub>  | 11.1374  | 0.193886065 | mg/kg | 1.741                                   |
| 2      | mass of final methanol solution  | M <sub>hplc</sub>   | 2.9817   | 2.12132E-05 | g     | 0.000711                                |
| 3      | mass of 1 ml<br>aliquot of sample<br>in water solution<br>that is pass<br>through in to the<br>SPE cartridge | M <sub>SPE</sub>    | 1.0200   | 2.12132E-05 | g     | 0.00208                                 |
| 4      | mass of sample   | M <sub>sample</sub> | 2.1023   | 0.000141421 | g     | 0.00673                                 |
| 5      | mass of solution of sample in water  | $M_{sentrifuge}$    | 10.1237  | 0.000141421 | g     | 0.00140                                 |
| 6      | Recovery   | Recovery            | 103.9089 | 1.580259091 | %     | 1.521                                   |
| 7      | Repeatability<br>sample  | Repeatability       | 1        | 0.008701578 | -     | 0.870                                   |
| 8      | Middle standard solution   |                     | 10.5472  | 0.030070407 | mg/kg | 0.285                                   |
|        | X (C)  | 157.5271755         |          | mg/kg       |       |   |
|        | u <sub>c(X)</sub>  | 3.91668857          |          | mg/kg       |       | 2.5                                     |
|        | ge factor at<br>nce level of 95%   | 2                   |          |             |       |   |
| Expand | ed uncertainty   | 7.83337714          |          | mg/kg       | ]     |   |
| Result |  | 157.5 ± 7.8         |          | mg/kg       |       |   |

#### **Methyl Paraben**

| Mass fraction (mg/kg)<br>Replication | 96.579<br>13 |
|--------------------------------------|--------------|
| sd                                   | 2.410        |
| %RSD                                 | 2.495        |

The uncertainty budget is summarized in the following table

| No  | Source of<br>uncertainty   | Symbol             | Value    | u           | Unit  | Relative<br>Standard<br>Uncertainty (%) |
|-----|--|--------------------|----------|-------------|-------|---|
| 1   | Calibration curve  | Co <sub>hplc</sub> | 6.9015   | 0.095503547 | mg/kg | 1.384                                   |
| 2   | mass of final<br>methanol solution   | M <sub>hplc</sub>  | 2.9817   | 2.12132E-05 | g     | 0.000711                                |
| 3   | mass of 1 ml<br>aliquot of sample<br>in water solution<br>that is pass<br>through in to the<br>SPE cartridge | M <sub>SPE</sub>   | 1.0200   | 2.12132E-05 | g     | 0.00208                                 |
| 4   | mass of sample   | $M_{sample}$       | 2.1023   | 0.000141421 | g     | 0.00677                                 |
| 5   | mass of solution of sample in water  | $M_{centrifuge}$   | 10.1237  | 0.00014142  | g     | 0.00140                                 |
| 6   | Recovery   | Recovery           | 103.2608 | 3.04839692  | %     | 2.952                                   |
| 7   | Repeatability<br>sample  | Repeatability      | 1        | 0.00692142  | -     | 0.692                                   |
| 8   | Middle standard solution   |                    | 10.4444  | 0.027436596 | mg/kg | 0.263                                   |
|     | X (C)  | 96.57882833        | mg/kg    |             |       |   |
|     | u <sub>c(x)</sub>  | 3.228987879        | mg/kg    |             |       | 3.3                                     |
|     | overage factor at<br>idence level of 95%   | 2                  |          |             |       | ·                                       |
| Exp | anded uncertainty  | 6.457975           | mg/kg    |             |       |   |
|     | Result   | 96.6 ± 6.5         | mg/kg    |             |       |   |

| G      | CMS   |                         |               |             |       |   |  |  |  |  |
|--------|---|-------------------------|---------------|-------------|-------|---|--|--|--|--|
| N      | Without derivatization technique  |                         |               |             |       |   |  |  |  |  |
| λ      | $X = C_{GCMS} * \frac{m_{centrifuge}}{m_{sample}} * \frac{m_{hplc}}{m_{spe}} * \frac{m_{final}}{m_{200ul}} * \frac{1}{\text{Re}c}$  |                         |               |             |       |   |  |  |  |  |
|        | ${}_{ombine} = C_{\sqrt{\left(\frac{u_{CGC}}{C_{GC}}\right)^2 + \left(\frac{u_{Mhplc}}{M_{hplc}}\right)^2 + \left(\frac{u_{MSPE}}{M_{SPE}}\right)^2 + \left(\frac{u_{Mcentrifug}}{M_{centrifuge}}\right)^2 + \left(\frac{u_{Msample}}{M_{sample}}\right)^2 + \left(\frac{u_{Rec}}{Rec}\right)^2 + \left(\frac{u_{rep}}{1}\right)^2 + \left(\frac{u_{Cs}\tan dard}{C_{s}\tan dard}\right)^2 + \left(\frac{u_{Mfinal}}{M_{final}}\right)^2 + \left(\frac{u_{200ul}}{M_{200ul}}\right)^2 + \left(\frac{u_{200ul}}{M_$ |                         |               |             |       |   |  |  |  |  |
|        | paraben   | 07 208                  |               |             |       |   |  |  |  |  |
| Replic | fraction (mg/kg)<br>ation   | 97.298<br>12            |               |             |       |   |  |  |  |  |
| sd     |   | 4.801                   |               |             |       |   |  |  |  |  |
| %RSD   |   | 4.934                   |               |             |       |   |  |  |  |  |
| The un | certainty budget is sun   | nmarized in the fo      | llowing table |             |       |   |  |  |  |  |
| No     | Source of uncertainty   | Symbol                  | Value         | u           | Unit  | Relative<br>Standard<br>Uncertainty (%) |  |  |  |  |
| 1      | mass of sample  | M <sub>sample</sub>     | 2.361723      | 0.000141421 | g     | 0.0059791                               |  |  |  |  |
| 2      | mass of solution of<br>sample in water  | M <sub>centrifuge</sub> | 10.37422      | 0.000141421 | g     | 0.001363196                             |  |  |  |  |
| 3      | mass of 1 ml aliquot<br>of sample in water<br>solution that is pass<br>through in to the SPE  | M <sub>spe</sub>        | 1.0403        | 2.12132E-05 | g     | 0.00203914                              |  |  |  |  |
| 4      | Contriduo     Mass of final       mass of final     Mhplc       after celan up with     Mhplc       SPE     2.5990  |                         |               |             |       |   |  |  |  |  |
| 5      | mass of the 200 uL of<br>final methanol<br>solution to be diluted   | M <sub>200ul</sub>      | 0.1589        | 2.12132E-05 | g     | 0.013350                                |  |  |  |  |
| 6      | mass of the final solution for analysis with GCMS   | M <sub>final</sub>      | 0.8008        | 2.12132E-05 | g     | 0.002649                                |  |  |  |  |
| 7      | Presision   | Repeatability           | 1             | 0.013685199 |       | 1.3685199                               |  |  |  |  |
| 8      | Calibration curve   | C <sub>GC</sub>         | 1.8120        | 0.0113      | mg/kg | 0.62362                                 |  |  |  |  |
| 9      | Recovery  | Recovery                | 1.0169        | 0.032558472 |       | 3.2017378                               |  |  |  |  |
| 10     | Middle standard solution  | Standard<br>solution    | 3.1895        | 0.011289444 | mg/kg | 0.353956                                |  |  |  |  |
|        | X (C)   | 97.29764                | mg/kg         |             |       |   |  |  |  |  |
|        | u <sub>c(x)</sub>   | 3,458882568             | mg/kg         |             |       | 3.6                                     |  |  |  |  |

## STD-ITDI

#### Benzoic Acid

| Components of Uncertainty  | Uncertainty<br>contribution, % |
|--|--------------------------------|
| <ol> <li>Concentration of Benzoic Acid Standard Stock Solution<br/>(purity and gravimetric preparation)</li> </ol> | 10.1                           |
| 2) Calibration standard mixture (gravimetric mixing of blends)   | 28.0                           |
| 3) Mass of Sample for analysis   | 4.0                            |
| 4) Mass of Benzoic Acid ring-D5 in the sample  | 0.9                            |
| 5) Peak Area Ratio of Benzoic-Benzoic acid ring-d5 in<br>calibration blend   | 46.1                           |
| 6) Peak Area Ratio of Benzoic acid-Benzoic acid ring-d5 in the<br>sample   | 10.9                           |

#### TISTR

#### Benzoic acid

| Factor                    | Values    | Uncertainties |          |
|---------------------------|-----------|---------------|----------|
|                           | X         | u(x)          | u(x)/(x) |
| Measurement equation      |           |               |          |
| factors                   |           |               |          |
| Method Precision          | 1         | 0.014331      | 1.433%   |
| mzc                       | 0.25537   | 0.000035      | 0.014%   |
| my                        | 0.15314   | 0.000035      | 0.023%   |
| тус                       | 0.15165   | 0.000035      | 0.023%   |
| mx                        | 1.00878   | 0.000035      | 0.004%   |
| wz                        | 599.61102 | 2.290295      | 0.382%   |
| R'b                       | 0.94256   | 0.018312      | 1.943%   |
| R'bc                      | 0.95785   | 0.011681      | 1.219%   |
| Extraction effects/Matrix |           |               |          |
| effect                    | 1         | 0.01          | 1.000%   |

| n-butyl paraben<br>Factor                    | Values           |                           | 1                       |
|--|------------------|---------------------------|-------------------------|
| Factor                                       |                  | Uncertainties             |                         |
|  | X                | u(x)                      | u(x)/(x)                |
| Measurement equation factors                 |                  |                           |                         |
| Method Precision                             | 1.0000           | 0.02598                   | 2.598%                  |
| m <sub>zc</sub>                              | 0.18838          | 0.000035                  | 0.0188%                 |
| m <sub>y</sub>                               | 0.10174          | 0.000035                  | 0.0348%                 |
| m <sub>yc</sub>                              | 0.10618          | 0.000035                  | 0.0333%                 |
| m <sub>x</sub>                               | 0.99996          | 0.000035                  | 0.0035%                 |
| Wz   | 533.5706         | 3.466894                  | 0.6498%                 |
| R'b  | 1.0847           | 0.015236                  | 1.4045%                 |
| R'bc   | 1.1295           | 0.019938                  | 1.7651%                 |
| Additional Factors                           |                  | Enter u(x) = 0 and veff = | = 1 for unused factors. |
| matrix effects                               | 1.000            | 0.0100                    | 1.000%                  |
| Interference from two different<br>ion pairs | 1.000            | 0.0053                    | 0.534%                  |
| Methyl paraben                               |                  |                           | 1                       |
| Factor                                       | Values           | Uncertainties             |                         |
|  | x                | u(x)                      | u(x)/(x)                |
| Measurement equation<br>factors              |                  |                           |                         |
| Method Precision                             | 1.0000           | 0.01897                   | 1.897%                  |
| m <sub>zc</sub>                              | 0.18838          | 0.000035                  | 0.0188%                 |
| m <sub>y</sub>                               | 0.10174          | 0.000035                  | 0.0348%                 |
| m <sub>yc</sub>                              | 0.10618          | 0.000035                  | 0.0333%                 |
| m <sub>x</sub>                               | 0.99996          | 0.000035                  | 0.0035%                 |
| W  | 532.9885         | 2.348845                  | 0.4407%                 |
| Wz   |                  |                           | 1.3397%                 |
| w <sub>z</sub><br>R'b                        | 1.1473           | 0.015370                  | 1.000170                |
| -  | 1.1473<br>1.1462 | 0.015370                  | 1.4712%                 |
| R'b<br>R'bc                                  |                  |                           | 1.4712%                 |
| R'b  |                  | 0.016863                  | 1.4712%                 |

# UME

| Uncertainty budget of Benzoic Acid |        |       |          |          |  |  |
|------------------------------------|--------|-------|----------|----------|--|--|
|                                    |        | Value | u(x)     | u(x)/x   |  |  |
| Weighing of sample (mg)            |        | 1250  | 2.56E-04 | 2.05E-07 |  |  |
| Weighing of IS (mg)                |        | 140   | 2.02E-04 | 1.44E-06 |  |  |
| Native stock solution (mg/kg)      |        | 1200  | 6.60E-03 | 5.50E-06 |  |  |
| Labelled stock solution (mg/kg)    |        | 1200  | 1.10     | 9.18E-04 |  |  |
| Intermediate precision             |        | 100   | 8.68E-01 | 8.68E-03 |  |  |
| Recovery                           |        | 1     | 1.32E-02 | 1.32E-02 |  |  |
| Repeatability                      |        | 100   | 1.62     | 1.62E-02 |  |  |
|                                    |        |       |          | 2.27E-02 |  |  |
| Result (mg/kg)                     | 152.38 |       |          |          |  |  |
|                                    |        |       |          |          |  |  |
| Combined uncertainty               |        | 3.46  |          |          |  |  |
| Expanded uncertainty               |        | 6.92  |          |          |  |  |
| % Relative uncertainty             |        | 4.54  |          |          |  |  |
| % Relative standard uncertainty    |        | 2.27  |          |          |  |  |

| Uncertainty budget of methyl paraben |        |       |          |          |  |  |
|--------------------------------------|--------|-------|----------|----------|--|--|
|                                      |        | Value | u(x)     | u(x)/x   |  |  |
| Weighing of sample (mg)              |        | 1250  | 2.02E-04 | 1.62E-07 |  |  |
| Weighing of IS (mg)                  |        | 140   | 2.56E-04 | 1.83E-06 |  |  |
| Native stock solution (mg/kg)        |        | 800   | 1.50E-02 | 1.88E-05 |  |  |
| Labelled stock solution (mg/kg)      |        | 800   | 2.94E-02 | 3.68E-05 |  |  |
| Intermediate precision               |        | 100   | 6.36E-01 | 6.36E-03 |  |  |
| Recovery                             |        | 1     | 2.40E-02 | 2.40E-02 |  |  |
| Repeatability                        |        | 100   | 8.78E-01 | 8.78E-03 |  |  |
|                                      |        |       |          | 2.64E-02 |  |  |
| Result (mg/kg)                       | 101.46 |       |          |          |  |  |
|                                      |        |       |          |          |  |  |
| Combined uncertainty                 |        | 2.68  |          |          |  |  |
| Expanded uncertainty                 |        | 5.35  |          |          |  |  |
| % Relative uncertainty               |        | 5.28  |          |          |  |  |
| % Relative standard uncertainty      |        | 2.64  |          |          |  |  |

| Uncertainty budget of butyl paraben |          |       |          |          |  |  |
|-------------------------------------|----------|-------|----------|----------|--|--|
|                                     | <u> </u> | Value | u(x)     | u(x)/x   |  |  |
| Weighing of sample (mg)             |          | 1250  | 1.01E-05 | 8.11E-09 |  |  |
| Weighing of IS (mg)                 |          | 140   | 1.48E-04 | 1.06E-06 |  |  |
| Native stock solution (mg/kg)       |          | 800   | 1.57E-02 | 1.96E-05 |  |  |
| Labelled stock solution (mg/kg)     |          | 800   | 1.51E-02 | 1.88E-05 |  |  |
| Intermediate precision              |          | 100   | 4.60E-01 | 4.60E-03 |  |  |
| Recovery                            |          | 1     | 2.58E-02 | 2.58E-02 |  |  |
| Repeatability                       |          | 100   | 5.47E-01 | 5.47E-03 |  |  |
|                                     |          |       |          | 2.68E-02 |  |  |
| Result (mg/kg)                      | 103.58   |       |          |          |  |  |
|                                     |          |       |          |          |  |  |
| Combined uncertainty                |          | 2.77  |          |          |  |  |
| Expanded uncertainty                |          | 5.54  |          |          |  |  |
| % Relative uncertainty              |          | 5.35  |          |          |  |  |
| % Relative standard uncertainty     |          | 2.68  |          |          |  |  |

# APPENDIX II

# Core Competency Tables for Measurands Reported in APMP.QM-S8

| APMP.QM-S8   | EXHM/<br>GCSL-<br>EIM       | Determination of Mass Fraction of<br>Benzoic Acid, Methyl Paraben and n-<br>Butyl Paraben in Soy Sauce  |  |
|--|-----------------------------|---|--|
| <b>Scope of Measurement:</b><br>The supplementary comparison enables participating NMIs/DIs to demonstrate their<br>measurement capabilities in the determination of common preservatives in soy sauce, using<br>procedure(s) that required simple sample preparation and selective detection in the mass<br>fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food<br>preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous-<br>based beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based<br>condiments (e.g. vinegar, fish sauce, etc). |                             |   |  |
| Competency   | Tick,<br>cross,<br>or "N/A" | Specific Information as Provided by NMI/DI  |  |
| <b>Competencies for Value-Assignmer</b>  | nt of Calibra               | ant   |  |
| Calibrant: Did you use a "highly-pure substance" or calibration solution?  | ~                           | Pure material<br>Benzoic acid: BAM, U1009   |  |
| Identity verification of analyte(s) in calibration material.#  | N/A                         |   |  |
| For calibrants which are a highly-<br>pure substance: Value-Assignment /<br>Purity Assessment method(s). <sup>#</sup>  | ~                           | Performed additional qNMR experiment<br>on the benzoic acid (acidimetric) CRM<br>using DMSO <sub>2</sub> as internal standard<br>(previously measured against 3,5-<br>BTFMBA CRM 4601-a from NMIJ). The<br>re-calculated result (150.65 mg/kg $\pm$ 4.04<br>mg/kg) did not deviate much from the<br>original submitted result (150.66 mg/kg $\pm$<br>4.03 mg/kg). |  |
| For calibrants which are a calibration solution: Value-assignment method(s). <sup>#</sup>  | N/A                         |   |  |
| Sample Ánalysis Competencies   |                             |   |  |
| Identification of analyte(s) in sample   | ✓                           | Retention time, MRM mode with one ion pair, MS ion ratios   |  |
| Extraction of analyte(s) of interest from matrix   | ✓                           | Liquid-liquid extraction, sonication  |  |
| Cleanup - separation of analyte(s) of<br>interest from other interfering matrix<br>components (if used)  | ~                           | Filtration  |  |
| Transformation - conversion of<br>analyte(s) of interest to<br>detectable/measurable form (if used)  | N/A                         |   |  |
| Analytical system  | ✓                           | LC-MS/MS  |  |
| Calibration approach for value-<br>assignment of analyte(s) in matrix  | ~                           | IDMS<br>Single-point calibration, matrix-<br>matched, exact-matching  |  |

| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used) | ~   | Standard additions |
|--|-----|--------------------|
| Other  | N/A |                    |

| APMP.QM-S8                              | GLHK     | Determination of Mass Fraction of           |
|---|----------|---|
|   | OLIIK    |   |
|   |          | Benzoic Acid, Methyl Paraben and n-         |
|   |          | Butyl Paraben in Soy Sauce                  |
| Scope of Measurement:                   |          |   |
| The supplementary comparison enabl      |          |   |
|   |          | common preservatives in soy sauce, using    |
|   |          | tion and selective detection in the mass    |
|   |          | can be extended to include other polar food |
|   |          | other alkyl benzoates) in water, aqueous-   |
| condiments (e.g. vinegar, fish sauce, e |          | das, sports drinks, etc) and aqueous-based  |
| Competency                              | Tick,    | Specific Information as Provided by         |
| competency                              | cross,   | NMI/DI                                      |
|   | or "N/A" |   |
| Competencies for Value-Assignmer        |          | ant   |
| Calibrant: Did you use a "highly-pure   |          | Benzoic acid: HSA (HRM-1002A)               |
| substance" or calibration solution?     |          | $(999.9 \pm 3.3 \text{ mg/g})$              |
|   |          |   |
|   | ✓        | Methyl paraben: HSA (HRM-1003A)             |
|   |          | (999.5 ± 3.2 mg/g)                          |
|   |          |   |
| Identity verification of analyte(s) in  | N/A      |   |
| calibration material. <sup>#</sup>      |          |   |
| For calibrants which are a highly-      |          |   |
| pure substance: Value-Assignment /      | N/A      |   |
| Purity Assessment method(s).#           |          |   |
| For calibrants which are a calibration  |          |   |
| solution: Value-assignment              | N/A      |   |
| method(s).#                             |          |   |
| Sample Analysis Competencies            | I        | GCMS: Retention time and the ion ratio      |
| Identification of analyte(s) in sample  | ✓        | LCMSMS: Retention time and the ion ratio    |
|   | •        | transition ratio                            |
| Extraction of analyte(s) of interest    |          | Added 6N HCl to 1g of sample                |
| from matrix                             | ✓        | Added on the to the of sample               |
| Cleanup - separation of analyte(s) of   |          | Liquid-liquid extraction by diethyl ether.  |
| interest from other interfering matrix  |          | Dried and reconstituted in chloroform (for  |
| components (if used)                    | ✓        | GC-MS) or in mobile phase : MeOH (95:5)     |
|   |          | (for LC-MS/MS)                              |
| Transformation - conversion of          |          | For ID-GC-MS, the extraction solution was   |
| analyte(s) of interest to               | ✓        | derivatized by N-Methyl-N-trimethylsilyl-   |
| detectable/measurable form (if used)    |          | trifluoracetamide (MSTFA) for detection.    |
| Analytical system                       | ✓        | ID-GC-MS                                    |

|  |     | ID-LC-MS/MS   |
|--|-----|---|
| Calibration approach for value-<br>assignment of analyte(s) in matrix                | ~   | a) IDMS<br>b) Bracketing<br>(The sample was calibrated by isotope<br>dilution of ${}^{13}C_6$ -Benzoic acid (for benzoic<br>acid) and ${}^{13}C_6$ -Methyl paraben (for methyl<br>paraben). The calibration blends were<br>prepared by gravimetric method. The<br>value-assignment of analytes in matrix<br>was calculated by bracketing method.) |
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used) | ~   | Matrix CRM from HSA (HRM – 1005A)<br>(Benzoic Acid, Methyl Paraben and n-<br>Butyl Paraben in Soy Sauce) was used to<br>verify the method for value-assignments.  |
| Other  | N/A |   |

| APMP.QM-S8  | HSA           | Determination of Mass Fraction of<br>Benzoic Acid, Methyl Paraben and n-<br>Butyl Paraben in Soy Sauce              |  |
|---|---------------|---|--|
| Scope of Measurement:   |               |   |  |
| The supplementary comparison enables participating NMIs/DIs to demonstrate their measurement capabilities in the determination of common preservatives in soy sauce, using procedure(s) that required simple sample preparation and selective detection in the mass fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous-based beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based condiments (e.g. vinegar, fish sauce, etc). |               |   |  |
| Competency  | Tick,         | Specific Information as Provided by   |  |
|   | cross,        | NMI/DI  |  |
| Or man a familie a fam Malera Arabimmera  | or "N/A"      |   |  |
| Competencies for Value-Assignmen  | nt of Calibra |   |  |
| Calibrant: Did you use a "highly-pure substance" or calibration solution?   | ~             | Pure materials<br>Benzoic acid: HSA, HRM-1002A<br>Methyl paraben: HSA, HRM-1003A<br>n-Butyl paraben: HSA, HRM-1004A |  |
| Identity verification of analyte(s) in calibration material. <sup>#</sup>   | N/A           |   |  |
| For calibrants which are a highly-<br>pure substance: Value-Assignment /<br>Purity Assessment method(s). <sup>#</sup>   | N/A           |   |  |
| For calibrants which are a calibration solution: Value-assignment method(s). <sup>#</sup>   | N/A           |   |  |
| Sample Analysis Competencies  |               |   |  |
| Identification of analyte(s) in sample  | ✓             | Retention times, SIM mode with three ions on GC-MS, MS ion ratios   |  |
| Extraction of analyte(s) of interest from matrix  | ✓             | Liquid-liquid extraction, shaking, centrifugation   |  |
| Cleanup - separation of analyte(s) of   | ✓             | Filtration  |  |

| interest from other interfering matrix components (if used)   |     |  |
|---|-----|--|
| Transformation - conversion of<br>analyte(s) of interest to<br>detectable/measurable form (if used) | N/A |  |
| Analytical system   | ✓   | GC-MS  |
| Calibration approach for value-<br>assignment of analyte(s) in matrix                               | ✓   | IDMS<br>Single-point calibration, exact-matching |
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used)                | ~   | Two qualifying ions per measurand                |
| Other   | N/A |  |

| APMP.QM-S8  | INMETRO                     | Determination of Mass Fraction of<br>Benzoic Acid, Methyl Paraben and n-<br>Butyl Paraben in Soy Sauce         |
|---|-----------------------------|--|
| Scope of Measurement:<br>The supplementary comparison enables participating NMIs/DIs to demonstrate their<br>measurement capabilities in the determination of common preservatives in soy sauce, using<br>procedure(s) that required simple sample preparation and selective detection in the mass<br>fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food<br>preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous-<br>based beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based<br>condiments (e.g. vinegar, fish sauce, etc). |                             |  |
| Competency  | Tick,<br>cross, or<br>"N/A" | Specific Information as Provided by NMI/DI   |
| Competencies for Value-Assignmer  | nt of Calibrar              | nt   |
| Calibrant: Did you use a "highly-pure substance" or calibration solution?   | ~                           | NIST SRM 350b (highly-pure benzoic<br>acid). 999.93 mg/g $\pm$ 3.1 mg/g, k=4.3,<br>95% (balance mass approach) |
| Identity verification of analyte(s) in calibration material. <sup>#</sup>   | ~                           | GC-MS and NMR (NMR chemical shifts<br>and MS spectrum consistent with<br>benzoic acid)                         |
| For calibrants which are a highly-<br>pure substance: Value-Assignment /<br>Purity Assessment method(s). <sup>#</sup>   | N/A                         |  |
| For calibrants which are a calibration solution: Value-assignment method(s). <sup>#</sup>   | N/A                         |  |
| Sample Analysis Competencies  |                             |  |
| Identification of analyte(s) in sample  | ✓                           | GC-MS  |
| Extraction of analyte(s) of interest from matrix  | ~                           | Liquid-liquid extraction with ethyl ether  |
| Cleanup - separation of analyte(s) of<br>interest from other interfering matrix<br>components (if used)   | N/A                         |  |
| Transformation - conversion of  | ✓                           | Derivatisation with MSTFA, resulting in  |

| analyte(s) of interest to detectable/measurable form (if used)                       |              | Benzoic acid trimethylsilyl ester   |
|--|--------------|---|
| Analytical system  | ✓            | GC-MS   |
| Calibration approach for value-<br>assignment of analyte(s) in matrix                | $\checkmark$ | GC-IDMS using calibration curve<br>approach with isotope labeled internal<br>standard   |
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used) | $\checkmark$ | Recovery studies of fortified blank matrix  |
| Other  | ✓            | The purity value (999.93 mg/g ± 3.1 mg/g,<br>k=4.3, 95% CI) of the benzoic acid CRM<br>determined by KRISS (based on mass<br>balance approach) was used in place of<br>NIST's certified value (based on coulometric<br>assay). Using this purity value, INMETRO's<br>re-calculated result did not change.<br>The uncertainty associated with the result for<br>benzoic acid would be increased by 4.0% to<br>cross the DoE line (Di = -5.95 mg/kg for<br>reported result of 148.6 mg/kg) for the<br>purpose of CMC claim in future. |

DoE result for benzoic acid does not cross zero. The reported value was not consistent with the KCRV. No specific competency in the Table above was identified as the reason. However, INMETRO would increase the uncertainty of their measurement in future.

| APMP.QM-S8  | KRISS  | Determination of Mass Fraction of<br>Benzoic Acid, Methyl Paraben and n-<br>Butyl Paraben in Soy Sauce   |  |
|---|--|--|--|
| <b>Scope of Measurement:</b><br>The supplementary comparison enables participating NMIs/DIs to demonstrate their measurement capabilities in the determination of common preservatives in soy sauce, using procedure(s) that required simple sample preparation and selective detection in the mass fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous-based beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based condiments (e.g. vinegar, fish sauce, etc). |  |  |  |
| Competency  | Tick,<br>cross,<br>or "N/A"                    | Specific Information as Provided by NMI/DI   |  |
| <b>Competencies for Value-Assignmer</b>   | Competencies for Value-Assignment of Calibrant |  |  |
| Calibrant: Did you use a "highly-pure substance" or calibration solution?   | ~  | Benzoic acid was purchased from NIST,<br>methyl paraben from SIGMA, and n-butyl<br>from Dr. Ehrenstorfer. Purities of three<br>compounds were assayed by KRISS |  |
| Identity verification of analyte(s) in calibration material. <sup>#</sup>   | ✓  | LC/MS/MS   |  |
| For calibrants which are a highly-  |  | Mass balance: LC/UV (related organic   |  |

| pure substance: Value-Assignment /<br>Purity Assessment method(s). <sup>#</sup><br>For calibrants which are a calibration |              | impurities), TGA (non-volatile residue)<br>Headspace GC/MSD (residual solvent)<br>and Karl-Fisher titmetry (water contents)<br>Gravimetrically prepared four solutions |
|---|--------------|--|
| solution: Value-assignment method(s). <sup>#</sup>  | ✓            | were analysed and cross checked by ID-<br>LC/MS/MS.  |
| Sample Analysis Competencies  |              |  |
| Identification of analyte(s) in sample  | ✓            | LC/MS/MS   |
| Extraction of analyte(s) of interest from matrix  | N/A          | No extraction  |
| Cleanup - separation of analyte(s) of<br>interest from other interfering matrix<br>components (if used)                   | ~            | C18 SPE  |
| Transformation - conversion of<br>analyte(s) of interest to<br>detectable/measurable form (if used)                       | N/A          | No transformation  |
| Analytical system   | $\checkmark$ | ID-LC/MS/MS  |
| Calibration approach for value-<br>assignment of analyte(s) in matrix   | ✓            | One-point exact-matching calibration with an isotope ratio standard solution.  |
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used)                                      | ~            | KRISS CRM 108-06-001   |
| Other   | N/A          |  |

| APMP.QM-S8  | NIM                | Determination of Mass Fraction of                |
|---|--------------------|--|
|   |                    | Benzoic Acid, Methyl Paraben and n-              |
|   |                    | Butyl Paraben in Soy Sauce                       |
| Scope of Measurement:   |                    |  |
| The supplementary comparison enable   | es participat      | ing NMIs/DIs to demonstrate their                |
| measurement capabilities in the deterr  | nination of c      | common preservatives in soy sauce, using         |
| procedure(s) that required simple sam   | ple prepara        | tion and selective detection in the mass         |
| fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food   |                    |  |
| preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous- |                    |  |
|   |                    | das, sports drinks, etc) and aqueous-based       |
| condiments (e.g. vinegar, fish sauce, e   |                    |  |
|   | ,                  | Creation of Dravided by                          |
| Competency  | Tick,              | Specific Information as Provided by<br>NMI/DI    |
|   | cross,<br>or "N/A" |  |
|   |                    |  |
| Competencies for Value-Assignmen  | t of Calibra       |  |
| Calibrant: Did you use a "highly-pure   |                    | Calibration solution:                            |
| substance" or calibration solution?   |                    | NIM-CRM GBW(E)100006 Benzoic acid                |
|   |                    | in water 1.00 mg/mL ( <i>U</i> =1%, <i>k</i> =2) |
|   | 1                  | Highly-pure substance:                           |
|   | ×                  | NIM-CRM GBW(E)100074 Methyl                      |
|   |                    | Paraben 996 mg/g ( <i>U</i> =0.2%, <i>k</i> =2)  |
|   |                    | NIM-CRM GBW(E)100077 n-Butyl                     |
|   |                    | Paraben 997 mg/g ( $U$ =0.2%, $k$ =2)            |

| Identity verification of analyte(s) in calibration material. <sup>#</sup>   | N/A | /   |
|---|-----|---|
| For calibrants which are a highly-<br>pure substance: Value-Assignment /<br>Purity Assessment method(s). <sup>#</sup> | N/A | /   |
| For calibrants which are a calibration solution: Value-assignment method(s). <sup>#</sup>                             | N/A | /   |
| Sample Analysis Competencies  |     |   |
| Identification of analyte(s) in sample  | ~   | LC-MS/MS quantifying ions, qualifying ions and retention time |
| Extraction of analyte(s) of interest from matrix  | ~   | Liquid –liquid extraction and ultrasonic extraction           |
| Cleanup - separation of analyte(s) of<br>interest from other interfering matrix<br>components (if used)               | N/A | /   |
| Transformation - conversion of<br>analyte(s) of interest to<br>detectable/measurable form (if used)                   | N/A | /   |
| Analytical system   | ✓   | LC-MS/MS (ACQUITY UPLC <sup>™</sup> Xevo TQ-<br>S)            |
| Calibration approach for value-<br>assignment of analyte(s) in matrix   | ~   | Single-point calibration (IDMS)                               |
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used)                                  | ~   | HPLC-DAD external calibration                                 |
| Other   | N/A | /   |

| APMP.QM-S8   | RCChem-   | Determination of Mass Fraction of                |  |  |  |  |
|--|---|--|--|--|--|--|
| AFINIF.QINI-30   | LIPI  |  |  |  |  |  |
|  | LIFI  | Benzoic Acid, Methyl Paraben and n-              |  |  |  |  |
|  |   | Butyl Paraben in Soy Sauce                       |  |  |  |  |
| Scope of Measurement:  |   |  |  |  |  |  |
| The supplementary comparison enable  | oles participat   | ing NMIs/DIs to demonstrate their                |  |  |  |  |
| measurement capabilities in the dete   | measurement capabilities in the determination of common preservatives in soy sauce, using |  |  |  |  |  |
| procedure(s) that required simple sample preparation and selective detection in the mass |   |  |  |  |  |  |
| fraction range from 50 to 1000 mg/kg   | . The study c   | an be extended to include other polar food       |  |  |  |  |
| preservatives (e.g. sorbic acid, propie  | onic acid and   | other alkyl benzoates) in water, aqueous-        |  |  |  |  |
|  |   | das, sports drinks, etc) and aqueous-based       |  |  |  |  |
| condiments (e.g. vinegar, fish sauce,  | etc).   |  |  |  |  |  |
| Competency   | Tick,   | Specific Information as Provided by              |  |  |  |  |
|  | cross, or   | NMI/DI   |  |  |  |  |
|  | "N/A"   |  |  |  |  |  |
| Competencies for Value-Assignment of Calibrant   |   |  |  |  |  |  |
| Calibrant: Did you use a "highly-  |   | We used "highly-pure substance" from             |  |  |  |  |
| pure substance" or calibration   |   | Health Science Authority(HSA), Singapore         |  |  |  |  |
| solution?  | $\checkmark$  | <ul> <li>HSA HRM-1002A (Benzoic acid,</li> </ul> |  |  |  |  |
|  |   | Purity 999.9±3.3 mg/g)                           |  |  |  |  |
|  |   | HSA HRM-1003A(Methyl paraben,                    |  |  |  |  |

|   |                       | Purity 999.5±3.2 mg/g )                            |
|---|-----------------------|--|
|   |                       | <ul> <li>HSA HRM-1004A(n-Butyl paraben,</li> </ul> |
|   |                       | Purity 999.2±3.3 mg/g)                             |
| Identity verification of analyte(s) in      | N/A                   |  |
| calibration material. <sup>#</sup>          |                       |  |
| For calibrants which are a highly-          |                       |  |
| pure substance: Value-Assignment            | N/A                   |  |
| / Purity Assessment method(s). <sup>#</sup> |                       |  |
| For calibrants which are a                  |                       |  |
| calibration solution: Value-                | N/A                   |  |
| assignment method(s).#                      |                       |  |
| Sample Analysis Competencies                |                       |  |
| Identification of analyte(s) in             | ✓                     | Identification of analyte(s) in sample was         |
| sample                                      | •                     | done by using GC-MS.                               |
| Extraction of analyte(s) of interest        |                       | Extraction and Clean up was done by                |
| from matrix                                 |                       | employing conditioned C18 Solid Phase              |
|   |                       | Extraction cartridge. The cartridge                |
|   | <ul> <li>✓</li> </ul> | containing sample was then washed with             |
|   |                       | 4 ml of 10% methanol in 1% phosphoric              |
|   |                       | acid and then the analytes were eluted             |
|   |                       | with 3 ml of methanol.                             |
| Cleanup - separation of analyte(s)          |                       |  |
| of interest from other interfering          | ✓                     |  |
| matrix components (if used)                 |                       |  |
| Transformation - conversion of              |                       |  |
| analyte(s) of interest to                   | N/A                   |  |
| detectable/measurable form (if              |                       |  |
| used)<br>Analytical system                  |                       | HPLC-DAD and GC-MS were used for                   |
| Analytical System                           | ✓                     |  |
|   | <b>v</b>              | quantitative analysis of analyte(s) in matrix.     |
| Calibration approach for value-             |                       | External calibration approach was used             |
| assignment of analyte(s) in matrix          | ✓                     | for value-assignment of analyte(s) in              |
|   |                       | matrix.  |
| Verification method(s) for value-           |                       | The validated analytical method was used           |
| assignment of analyte(s) in sample          | ✓                     | for value-assignment of analyte(s) in              |
| (if used)                                   |                       | sample.  |
| Other                                       |                       | CRM matrix (HSA HRM-1005A) was used                |
|   | ✓                     | as quality control material for evaluating         |
|   |                       | the performance of the analytical method.          |

| APMP.QM-S8            | STD-ITDI* | Determination of Mass Fraction of   |
|-----------------------|-----------|-------------------------------------|
|                       |           | Benzoic Acid, Methyl Paraben and n- |
|                       |           | Butyl Paraben in Soy Sauce          |
| Scope of Measurement: |           |                                     |

The supplementary comparison enables participating NMIs/DIs to demonstrate their measurement capabilities in the determination of common preservatives in soy sauce, using procedure(s) that required simple sample preparation and selective detection in the mass

fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueousbased beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based condiments (e.g. vinegar, fish sauce, etc).

| Competency  | Tick,<br>cross, or<br>"N/A" | Specific Information as Provided by NMI/DI  |  |
|---|-----------------------------|---|--|
| Competencies for Value-Assignme   | ent of Calibr               | ant   |  |
| Calibrant: Did you use a "highly-<br>pure substance" or calibration<br>solution?                                      | ~                           | The benzoic acid standard (NIST SRM-<br>350b) was purchased from NIST.  |  |
| Identity verification of analyte(s) in calibration material. <sup>#</sup>   | ~                           | The identity of benzoic acid was confirmed by LCMSMS  |  |
| For calibrants which are a highly-<br>pure substance: Value-Assignment<br>/ Purity Assessment method(s). <sup>#</sup> | N/A                         |   |  |
| For calibrants which are a calibration solution: Value-assignment method(s).#   | N/A                         |   |  |
| Sample Analysis Competencies  |                             |   |  |
| Identification of analyte(s) in sample  | ~                           | The identity of benzoic acid was confirmed by LCMSMS  |  |
| Extraction of analyte(s) of interest from matrix  | ~                           | Extraction with mobile phase<br>(methanol:ammonium acetate buffer<br>50:50)   |  |
| Cleanup - separation of analyte(s)<br>of interest from other interfering<br>matrix components (if used)               | N/A                         |   |  |
| Transformation - conversion of<br>analyte(s) of interest to<br>detectable/measurable form (if<br>used)                | N/A                         |   |  |
| Analytical system   | ✓                           | LCMSMS  |  |
| Calibration approach for value-<br>assignment of analyte(s) in matrix   | ~                           | Exact matching IDMS   |  |
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used)                                  | ~                           | HPLC-PDA (gravimetric).<br>Use of CRM from HSA (HRM-1001A) as<br>quality control material.  |  |
| Other   | ~                           | The purity value (999.93 mg/g ± 3.1 mg/g,<br>k=4.3, 95% CI) of the benzoic acid CRM<br>determined by KRISS using mass balance<br>approach was used in place of NIST's<br>certified value (based on coulometric<br>assay). Using this purity value, STD-ITDI's<br>re-calculated result did not change. |  |

\*Important Note: the metrology in chemistry activities under STD-ITDI will eventually be part of the National Metrology Laboratory of ITDI. However, there will be no major changes to the manpower and infrastructure supporting the capability demonstrated in this supplementary comparison following the change in organisational structure.

| APMP.QM-S8  | TISTR         | Determination of Mass Fraction of<br>Benzoic Acid, Methyl Paraben and n-<br>Butyl Paraben in Soy Sauce |
|---|---------------|--|
| Scope of Measurement:   |               |  |
| The supplementary comparison enable                                       | • •           | •  |
| •   |               | common preservatives in soy sauce, using   |
|   |               | tion and selective detection in the mass   |
| fraction range from 50 to 1000 mg/kg.                                     | The study c   | can be extended to include other polar food  |
| preservatives (e.g. sorbic acid, propior                                  | nic acid and  | other alkyl benzoates) in water, aqueous-  |
| based beverages (e.g. fruit juices, tea                                   | extracts, so  | das, sports drinks, etc) and aqueous-based   |
| condiments (e.g. vinegar, fish sauce, e                                   | etc).         |  |
| Competency  | Tick,         | Specific Information as Provided by  |
|   | cross,        | NMI/DI   |
|   | or "N/A"      |  |
| Competencies for Value-Assignmer  | nt of Calibra |  |
| Calibrant: Did you use a "highly-pure substance" or calibration solution? |               | Highly-pure substance purchased from<br>HSA  |
| substance of calibration solution?  |               | ПЗА  |
|   |               | HSA (HRM-1002A) Benzoic acid   |
|   | ✓             | HSA (HRM-1003A) Methyl paraben   |
|   |               | calibrant  |
|   |               | HSA (HRM-1004A) n-Butyl paraben  |
|   |               | calibrant  |
| Identity verification of analyte(s) in                                    |               | Identity verification of analytes were done  |
| calibration material.#  | ✓             | by LC-MS/MS, qualifying ions and   |
| For collegate which are a bight.  |               | retention time   |
| For calibrants which are a highly-<br>pure substance: Value-Assignment /  | N/A           |  |
| Purity Assessment method(s). <sup>#</sup>                                 | IN/A          |  |
| For calibrants which are a calibration                                    |               |  |
| solution: Value-assignment  | N/A           |  |
| method(s). <sup>#</sup>   |               |  |
| Sample Analysis Competencies  | •             |  |
| Identification of analyte(s) in sample                                    |               | Identity verification of analytes were done  |
|   | ✓             | by LC-MS/MS, qualifying ions and   |
| Entre ation of each (c/c) of interest                                     |               | retention time   |
| Extraction of analyte(s) of interest<br>from matrix                       | ✓             | Extraction with methanol followed by   |
| Cleanup - separation of analyte(s) of                                     |               | sonication<br>Filtration   |
| interest from other interfering matrix                                    | ✓             |  |
| components (if used)  |               |  |
| Transformation - conversion of  |               |  |
| analyte(s) of interest to   | N/A           |  |
| detectable/measurable form (if used)                                      |               |  |
| Analytical system   | ✓             | LC-MS/MS (Agilent 6400 Series Triple   |
|   |               | Quad LC/MS)  |

| Calibration approach for value-<br>assignment of analyte(s) in matrix                | ~   | Bracketing, exact-matching double IDMS.<br>Matrix-matched calibration standards. |
|--|-----|--|
| Verification method(s) for value-<br>assignment of analyte(s) in sample<br>(if used) | N/A | Recovery studies by spiking in blank matrix                                      |
| Other  | N/A |  |

| APMP.QM-S8 | UME | Determination of Mass Fraction of   |
|------------|-----|-------------------------------------|
|            |     |                                     |
|            |     | Benzoic Acid, Methyl Paraben and n- |
|            |     | Butul Develop in Cov Course         |
|            |     | Butyl Paraben in Soy Sauce          |

#### Scope of Measurement:

The supplementary comparison enables participating NMIs/DIs to demonstrate their measurement capabilities in the determination of common preservatives in soy sauce, using procedure(s) that required simple sample preparation and selective detection in the mass fraction range from 50 to 1000 mg/kg. The study can be extended to include other polar food preservatives (e.g. sorbic acid, propionic acid and other alkyl benzoates) in water, aqueous-based beverages (e.g. fruit juices, tea extracts, sodas, sports drinks, etc) and aqueous-based condiments (e.g. vinegar, fish sauce, etc).

| Competency  | Tick,<br>cross,<br>or "N/A" | Specific Information as Provided by NMI/DI  |  |  |
|---|-----------------------------|---|--|--|
| Competencies for Value-Assignment of Calibrant  |                             |   |  |  |
| Calibrant: Did you use a "highly-pure<br>substance" or calibration solution?  | ~                           | -Benzoic Acid SRM350b (NIST) (KRISS<br>and UME (qNMR) re-evaluated the value)<br>-Methyl paraben calibrant(Across<br>Chemicals) 0.99998 ±0.003 mg/g*<br>-n-Butyl paraben calibrant (AlfaAesar)<br>0.9999±0.00313*<br>*q NMR was used for value assignment |  |  |
| Identity verification of analyte(s) in calibration material. <sup>#</sup>   | ✓                           | LC-MS and qNMR  |  |  |
| For calibrants which are a highly-<br>pure substance: Value-Assignment /<br>Purity Assessment method(s). <sup>#</sup> | ~                           | qNMR  |  |  |
| For calibrants which are a calibration solution: Value-assignment method(s). <sup>#</sup>                             | N/A                         |   |  |  |
| Sample Analysis Competencies  |                             |   |  |  |
| Identification of analyte(s) in sample  | ✓                           | LC-MS   |  |  |
| Extraction of analyte(s) of interest from matrix  | ✓                           | Liquid-liquid extraction with methanol  |  |  |
| Cleanup - separation of analyte(s) of<br>interest from other interfering matrix<br>components (if used)               | ~                           | Filtration<br>Sample Dilution   |  |  |
| Transformation - conversion of<br>analyte(s) of interest to<br>detectable/measurable form (if used)                   | N/A                         |   |  |  |

| Analytical system                  | ✓            | LC-MS/MS |
|------------------------------------|--------------|----------|
| Calibration approach for value-    |              | IDMS     |
| assignment of analyte(s) in matrix | •            |          |
| Verification method(s) for value-  |              | IDMS     |
| assignment of analyte(s) in sample | $\checkmark$ |          |
| (if used)                          |              |          |
| Other                              | N/A          |          |

# APPENDIX III OTHER INFORMATION PROVIDED IN REPORT OF RESULTS

## UME

## Synthesis of Calibrants

# Methyl (CD<sub>3</sub>) 4-hydroxybenzoate

4-hydroxybenzoic acid (1g, 7.24 mmol) was dissolved in methanol (CD<sub>3</sub>OD, 6.66 mL, 0.164 mol). After adding concentrated sulphuric acid (1 mL, 10 M) the mixture was refluxed for 5 hours. Saturated sodium hydroxide ( $\approx$ 15 mL) was added to the cooled solution for neutralization. The resulting mixture was allowed to stand for 15 min, before being poured into a cool beaker which was made up to 500 mL with water. White crystalline solid (1.1 g, 98.2 %) was obtained after filtering and drying the precipitate.

<sup>1</sup>H NMR, δ (600MHz, CDCl<sub>3</sub>) 7.96 (2H, d, *J* 8.8 Hz, ArH); 6.89 (2H, d, *J* 8.8 Hz, ArH) <sup>13</sup>C NMR, δ (150 MHz, CDCl<sub>3</sub>) 167.43; 160.18; 131.95; 122.32; 115.26. [M<sup>-</sup>] m/z 153.7.

# Butyl (C<sub>4</sub>D<sub>9</sub>) 4-hydroxybenzoate

4-hydroxybenzoic acid (0.5 g, 3.62 mmol) was refluxed overnight in n-butanol(d10) (1.32 mL, 14.48 mmol) and toluene (0.765 mL, 7.24 mmol) using sulphuric acid as a catalyst (0.15 mL). The solvent was distilled off and the residual oily liquid was poured into a cool beaker with ice-water. After filtration and drying the precipitates, the white solid of butyl paraben was obtained (88 %).

<sup>1</sup>H NMR, δ (600MHz, CDCl<sub>3</sub>) 7.96 (2H, d, *J* 8.8 Hz, ArH); 6.88 (2H, d, *J* 8.8 Hz, ArH) <sup>13</sup>C NMR, δ (150 MHz, CDCl<sub>3</sub>) 166.87; 159.93; 131.87; 122.82; 115.17. [M<sup>-</sup>] m/z 202.0.

## Instrumunets and chromatographic conditions

Experiments were performed by a Zivak® HPLC and Zivak® Tandem Gold Triple quadrupole Mass Spectrometry equipped with a Phenomenex Luna 5u C18 column (150 x 2 mm i.d., 5µm particle size). The mobile phase was composed of (A, acetonitrile) in water (B, 5 mM ammonium acetate, pH:4.2), the gradient programme of which was 0-1.00 minute 20 % A and

80 % B, 1.01-8.00 minutes 80 % A and 20 % B, 8.01-8.06 minutes 20 % A and 80 % B and finally 08.06-12.00 20 % A and 80 % B. The flow rate of the mobile phase was 0.30 mL/min, and the column temperature was set to 30  $^{\circ}$ C. The injection volume was 10  $\mu$ L (table 1, figure 1).

# Table 1. LC-MS/MS parameters of selected compounds

|   | Compounds         | Parent ion | Daughter<br>ion | Collision energy<br>(V) |
|---|-------------------|------------|-----------------|-------------------------|
| 1 | Benzoic acid      | 120.70     | 77.00           | 10                      |
| 2 | Benzoic acid-IS   | 125.70     | 82.00           | 10                      |
| 3 | Methyl paraben    | 150.50     | 135.50          | 15                      |
| 4 | Methyl paraben-IS | 153.70     | 135.50          | 15                      |
| 5 | Butyl paraben     | 192.70     | 136.00          | 15                      |
| 6 | Butyl paraben-IS  | 202.00     | 136.00          | 15                      |



Figure 1. LC-IDMS chromatogram of methyl paraben, butyl paraben and benzoic acid. Traceability of calibrants

The purity assessment of native and synthesized labeled parabens was done by quantitative nuclear magnetic resonance (qNMR). 1,3,5-trimethoxybenzene was used as internal standard (IS) and methanol-D<sub>4</sub> (Merck, 99.8%) was used as solvent. 1,3,5-trimethoxybenzene purity (99.798 %  $\pm$  0.144) was determined by Nist 350b benzoic acid CRM (NIST, Gaithersburg,

US). Three different samples were prepared from each paraben and each sample was analyzed with three repetitions.

The sample solution of parabens was prepared by following steps: Paraben (20-30 mg) and 1,3,5-Trimethoxybenzene (20-30 mg) were accurately weighed, dissolved in CD<sub>3</sub>OD (2.1 mL), stirred with vortex for 30 seconds and 0.7 mL solution transferred to an NMR tube. All NMR measurements were carried out on a Varian 600 spectrometer operating at 599.90 MHz. The probe used was a Varian's One NMR. The following parameters were employed for acquisition of spectra: spectral width, 16 ppm; acquisition time, 3.4 s; relaxation delay, 40 s; 90° pulse width, 7.2  $\mu$ s; time domain, 64K data points; 32 scans; temperature, 298.15 K.

All NMR spectra were processed with the software Mestrenova 8.1.0. An exponential line broadening window function of 0.3 Hz was used in the data processing. After Fourier transformation of the free induction decays, the spectra were baseline corrected, phased, and integrated in the appropriate region. The peaks for the analyte and the internal standard were integrated inside, that is, excluding, the 13C satellites.

The calculation equation of qNMR for the purity is as follows:

$$P_x = \frac{I_x}{I_{Std}} \frac{N_{Std}}{N_x} \frac{M_x}{M_{Std}} \frac{m_{Std}}{m_x} P_{Std}$$

I<sub>Std</sub>, N<sub>Std</sub>, M<sub>Std</sub>, m<sub>Std</sub> and P<sub>Std</sub> are the peak area, number of proton, molecular weight, mass and purity of the internal standard, respectively. Ix, Nx, Mx, mx and Px are the peak area, number of proton, molecular weight, mass and purity of the sample, respectively.

The calculation equation of the relative standard uncertainty is as follows:

$$\frac{u(Px)}{Px} = \sqrt{\left(\frac{u(Ix/Istd)}{Ix/Istd}\right)^2 + \left(\frac{u(Mx)}{Mx}\right)^2 + \left(\frac{u(Mstd)}{Mstd}\right)^2 + \left(\frac{u(mx)}{mx}\right)^2 + \left(\frac{u(ms)}{ms}\right)^2 + \left(\frac{u(Ps)}{Ps}\right)}$$
The

uncertainty from balance is the most important component for the total uncertainty value so all of the samples were weighed with an advanced balance. The uncertainty from molecular weight is often pretty small than integration repeatability. All of the free induction decays of



