# EVALUATION OF INTERLABORATORY STUDY ON PCDD, PCDF AND DIOXIN LIKE PCB IN THE REFERENCE MATERIAL IN JAPAN (9TH ROUND FY 2011 RESEARCH GROUP ON ULTRA TRACE ANALYSES, JEMCA, JAPAN) 

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## Introduction

Inter-laboratory round robin is available for maintaining the quality/skills of dioxin analysis through testing bycertified laboratories. There are over 100 accredited Laboratories for dioxin by MLAP (Specified Measurement Laboratory Accreditation Program) system of Ministry of Economy Trade and Industry (METI) in Japan.Ministry of Environment (MOE) has another program for examining the order of competence also. But it is more important to maintain QA/QC system and evaluate quality of daily analysis data continuously. There are some official proficiency tests for dioxin analysis by JSAC (The Japan Society for Analytical Chemistry), MOE and METI in Japan.
Research Group on Ultra Trace Analyses (UTA) which is accompanying organization of Japan Environmental Measurement \& Chemical Analysis Association (JEMCA) established in 2003. The UTA consists of 63 private dioxin testing laboratories in 2011 and is responsible for developing the analytical potential of not only dioxins but also other trace level analysis of well known POPs in the environment. UTA carried out inter-laboratory round robin studies annually since 2003, R-1:flyash extract in 2003, R-2:soil in 2004, R-3:PUF fortified extract in 2005, R-4:soil in 2006, R-5:soil in 2007, R-6:flyash in 2008 ,R-7:sediment in 2009, R-8:sediment in 2010, and R-9:flyash in 2011 for polychlorinated dibenzo-p-dioxins (PCDDs), polychlorinated dibenzofurans (PCDFs) and dioxin-like polychlorinated biphenyls (DL-PCBs). This paper summarizes the recent inter-laboratory study (R-9, FY 2011) conducted by UTA group for PCDDs, PCDFs and DL- PCBs in sediment sample.

## Materials and methods

The flyash reference material the ninth round robin study (R-9) was sent to 63 laboratories. The flyash reference material was collected, dried, milled, homogenized and packed in 25 g portions.
All member laboratories were asked to report all 2,3,7,8-substituted PCDD/DFs congeners, homologues and 12 DL-PCBs. A special result form was sent to all members in which, the following details were requested; 1 . The analytical results obtained, including recovery percentage, 2 . Complete analytical procedure followed and 3. SIM Chromatograms of each sample. Results of these studies are evaluated for median, normalized interquartile range (NIQR), coefficient of variation by Robust method (CV \% rob) for each PCDDs, PCDFs and DL-PCBs. Furthermore Z-score was calculated and evaluated by ISO/IEC 17043(JIS Q 17043). Laboratories, which exceed > $\pm 3$ of Z-score were required cause analysis and report of corrective action.

## Results and discussion:

The results of the ninth round robin study on an isomer/congener specific basis with median, and their NIQR and CV \% rob are summarized in Table 1. Every data set was used to identify obvious outliers. Obvious outliers were defined as having each Z -score over 3.
As our earlier report, significant differences were observed between laboratories, in particular for 1,2,3,7,8-PeCDF and $1,2,3,4,7,8-\mathrm{HxCDF}$, depending upon the capillary column that was used for the analysis. The main causes of these
differences are due to co-eluting congeners in polar GC phase (SP-2331 or CP-Sil88) (ex. 1,2,3,7,8-PeCDF co-eluting $1,2,3,4,8-\mathrm{PeCDF}, 1,2,3,4,7,8-\mathrm{HxCDF}$ co-eluting 1,2,3,4,7,9-HxCDF). During combined data processing for these congeners, the CV \% rob value were clearly increased.
The CV \% rob in R-9 ranged from $4.9 \%$ to $9.8 \%$ for PCDDs/DFs congeners, $4.4 \%$ to $9.1 \%$ for DL-PCBs. and $3.83 \%$ for TEQ (not indicated in the table).

Table 1. Statistical analysis of the 9th round robin (R-9, 2011) study results of PCDDs/PCDFs and DL-PCBs.

| PCDDs/DFs, DL-PCB | $\begin{gathered} \text { MEDIAN } \\ (\mathrm{pg} / \mathrm{g}) \end{gathered}$ | NIQR | $\begin{gathered} \mathrm{CV} \% \\ \text { rob } \end{gathered}$ | $\begin{aligned} & \mathrm{MIN} \\ & (\mathrm{pg} / \mathrm{g}) \end{aligned}$ | $\begin{aligned} & \text { MAX } \\ & (\mathrm{pg} / \mathrm{g}) \end{aligned}$ | MEAN (pg/g) | SD | N |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 2378-TeCDD | 41 | 2.93 | 7.1 | 28 | 49 | 41 | 3.85 | 63 |
| 12378-PeCDD | 211 | 14.1 | 6.7 | 177 | 261 | 210 | 16.2 | 63 |
| 123478-HxCDD | 233 | 17.4 | 7.5 | 191 | 304 | 232 | 19.5 | 63 |
| 123678-HxCDD | 836 | 47.8 | 5.7 | 652 | 1030 | 830 | 60.5 | 63 |
| 123789-HxCDD | 726 | 41.9 | 5.8 | 577 | 897 | 727 | 54.4 | 63 |
| 1234678-HpCDD | 2438 | 166 | 6.8 | 2035 | 3110 | 2436 | 174 | 63 |
| OCDD | 977 | 87.7 | 9.0 | 785 | 1120 | 968 | 83.0 | 63 |
| 2378-TeCDF | 455 | 36.0 | 7.9 | 355 | 582 | 456 | 43.7 | 63 |
| 12378-PeCDF | 345 | 121 | 35.1 | 279 | 581 | 400 | 93.3 | 63 |
| 12378-PeCDF(*) | 337 | 16.5 | 4.9 | 279 | 459 | 336 | 27.3 | 41 |
| 12378-PeCDF(*) | 529 | 36.0 | 6.8 | 445 | 581 | 520 | 35.9 | 22 |
| 23478-PeCDF | 442 | 24.8 | 5.6 | 330 | 492 | 438 | 28.1 | 63 |
| 123478-HxCDF | 261 | 40.4 | 15.5 | 199 | 363 | 271 | 39.0 | 63 |
| 123478-HxCDF(*a) | 253 | 17.0 | 6.7 | 199 | 363 | 249 | 26.6 | 41 |
| 123478-HxCDF(*b) | 311 | 21.5 | 6.9 | 278 | 354 | 313 | 20.2 | 22 |
| 123678-HxCDF | 262 | 13.0 | 5.0 | 217 | 299 | 260 | 16.4 | 63 |
| 123789-HxCDF | 39 | 3.8 | 9.8 | 26 | 58 | 40 | 6.0 | 62 |
| 234678-HxCDF | 265 | 24.1 | 9.1 | 215 | 320 | 268 | 25.3 | 63 |
| 1234678-HpCDF | 481 | 25.2 | 5.2 | 409 | 578 | 481 | 32.5 | 63 |
| 1234789-HpCDF | 104 | 8.75 | 8.4 | 83 | 126 | 105 | 9.69 | 63 |
| OCDF | 268 | 23.8 | 8.9 | 215 | 327 | 267 | 24.2 | 63 |
| 344'5-TeCB(\#81) | 119 | 5.19 | 4.4 | 101 | 137 | 118 | 7.27 | 63 |
| 33'44'-TeCB(\#77) | 489 | 25.6 | 5.2 | 380 | 558 | 484 | 34.4 | 63 |
| 33'44'5-PeCB(\#126) | 183 | 14.5 | 7.9 | 139 | 240 | 184 | 19.4 | 63 |
| 33'44'55'-НxCB(\#169) | 40 | 3.0 | 7.6 | 32 | 119 | 41 | 10.6 | 63 |
| 2'344'5-PeCB(\#123) | 75 | 5.2 | 6.9 | 60 | 104 | 76 | 7.11 | 63 |
| 23'44'5-PeCB(\#118) | 1625 | 125 | 7.7 | 1188 | 2050 | 1636 | 147 | 63 |
| 233'44'-PeCB(\#105) | 733 | 61.5 | 8.4 | 612 | 877 | 744 | 58.5 | 63 |
| 2344'5-PeCB(\#114) | 85 | 7.75 | 9.1 | 66 | 111 | 85 | 8.23 | 63 |
| 23'44'55'-НxCB(\#167) | 123 | 8.25 | 6.7 | 98 | 152 | 124 | 10.8 | 63 |
| 233'44'5-HxCB(\#156) | 319 | 24.1 | 7.6 | 250 | 369 | 319 | 25.5 | 63 |
| 233'44'5'-НxCB(\#157) | 99 | 8.01 | 8.1 | 82 | 111 | 99 | 7.15 | 63 |
| 233'44'55'-HpCB(\#189) | 78 | 4.15 | 5.3 | 61 | 91 | 78 | 5.06 | 63 |

Used GC column for 1,2,3,7,8-PeCDF and 1,2,3,4,7,8-HxCDF analysis
(※a) BPX-DXN, DB-5, BPX-5, RH-12ms etc.: separate single peak
(※b) SP-2331, CP-Sil88 etc.: including co-elute congeners

Table 2 describes the trends of CV \% rob from the 1st to 9 th round robin study. It seems better CV\% rob result for 9th round robin study as compared with the result of the past. These results indicate appreciable improvement of the analytical techniques and systems of UTA individual laboratories every year. They have gradually increased number of laboratories to use GC columns that can separate other congeners in the analysis of $1,2,3,7,8-\mathrm{PeCDF}$ and $1,2,3,4,7,8-$ HxCDF. (e.g. during R-9 study the use of such columns is $65 \%$ while it was only $24 \%$ during R-1).

Table 2. Trends of the round robin study results (CV \% rob)

| PCDDs/DFs, DL-PCB | $\begin{gathered} \hline 2003 \\ \text { (1st) } \\ \text { flyash } \\ \text { ext. } \end{gathered}$ | $\begin{gathered} \hline 2004 \\ \text { (2nd) } \\ \text { soil } \end{gathered}$ | $\begin{gathered} \hline 2005 \\ \text { (3rd) } \\ \text { PUF } \\ \text { fortified } \end{gathered}$ | $\begin{gathered} \hline 2006 \\ (4 \mathrm{th}) \\ \text { soil } \end{gathered}$ | $\begin{gathered} \hline 2007 \\ (5 \text { th) } \\ \text { soil } \end{gathered}$ | $\begin{gathered} \hline 2008 \\ \text { (6th) } \\ \text { flyash } \end{gathered}$ | $2009$ <br> (7th) sediment | $\begin{gathered} \hline 2010 \\ \text { (8th) } \\ \text { sediment } \end{gathered}$ | $\begin{gathered} \hline 2011 \\ \text { (9th) } \\ \text { flyash } \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 2378-TeCDD | 8.5\% | 15.8\% | 7.5\% | 24.8\% | 15.4\% | 11.4\% | 15.0\% | 15.6\% | 7.1\% |
| 12378-PeCDD | 6.3\% | 16.7\% | 5.9\% | 16.6\% | 10.3\% | 9.6\% | 9.3\% | 7.2\% | 6.7\% |
| 123478-HxCDD | 10.3\% | 15.8\% | 8.5\% | 16.2\% | 10.7\% | 10.0\% | 7.6\% | 6.9\% | 7.5\% |
| 123678-HxCDD | 8.7\% | 16.6\% | 9.7\% | 13.3\% | 10.4\% | 9.9\% | 6.9\% | 6.7\% | 5.7\% |
| 123789-HxCDD | 8.3\% | 17.5\% | 6.6\% | 13.4\% | 6.8\% | 10.8\% | 8.0\% | 9.2\% | 5.8\% |
| 1234678-HpCDD | 6.3\% | 16.5\% | 9.0\% | 15.4\% | 10.2\% | 8.9\% | 6.3\% | 6.4\% | 6.8\% |
| OCDD | 8.5\% | 11.5\% | 6.4\% | 13.9\% | 7.5\% | 8.3\% | 6.7\% | 8.5\% | 9.0\% |
| 2378-TeCDF | 9.1\% | 17.9\% | 9.1\% | 15.6\% | 8.8\% | 13.8\% | 11.0\% | 10.7\% | 7.9\% |
| 12378-PeCDF | 10.0\% | 23.4\% | 11.1\% | 22.2\% | 23.6\% | 38.8\% | 32.2\% | 27.1\% | 35.1\% |
| 12378-PeCDF( ${ }^{( } \mathrm{a}$ ) |  |  | 7.8\% | 14.1\% | 9.8\% | 12.6\% | 5.7\% | 9.7\% | 4.9\% |
| 12378-PeCDF(*) |  |  | 9.0\% | 16.2\% | 7.1\% | 9.8\% | 9.2\% | 9.4\% | 6.8\% |
| 23478-PeCDF | 6.1\% | 13.7\% | 6.2\% | 9.7\% | 10.3\% | 9.2\% | 7.5\% | 11.6\% | 5.6\% |
| 123478-HxCDF | 6.3\% | 15.2\% | 8.9\% | 11.5\% | 13.0\% | 10.8\% | 9.9\% | 8.8\% | 15.5\% |
| 123478-HxCDF(*a) |  |  | 9.1\% | 9.9\% | 8.6\% | 9.2\% | 9.0\% | 8.1\% | 6.7\% |
| 123478-HxCDF(* ${ }^{\text {b }}$ ) |  |  | 7.2\% | 13.2\% | 8.0\% | 10.8\% | 5.6\% | 9.9\% | 6.9\% |
| 123678-HxCDF | 6.3\% | 10.7\% | 9.3\% | 14.5\% | 6.5\% | 10.6\% | 8.2\% | 9.4\% | 5.0\% |
| 123789-HxCDF | 12.9\% | 17.4\% | 10.3\% | 16.9\% | 14.7\% | 15.8\% | 13.7\% | 12.3\% | 9.8\% |
| 234678-HxCDF | 6.6\% | 10.4\% | 10.0\% | 21.8\% | 10.1\% | 10.3\% | 10.4\% | 10.0\% | 9.1\% |
| 1234678-HpCDF | 7.4\% | 10.5\% | 7.9\% | 16.2\% | 10.1\% | 9.9\% | 9.1\% | 7.7\% | 5.2\% |
| 1234789-HpCDF | 7.5\% | 11.5\% | 10.2\% | 15.2\% | 9.5\% | 9.8\% | 7.9\% | 7.3\% | 8.4\% |
| OCDF | 7.0\% | 10.7\% | 11.6\% | 11.5\% | 10.1\% | 10.2\% | 9.0\% | 6.6\% | 8.9\% |
| 344'5-TeCB(\#81) | 11.5\% | 12.9\% | 9.0\% | 16.1\% | 10.2\% | 10.0\% | 5.6\% | 11.9\% | 4.4\% |
| 33'44'-TeCB(\#77) | 8.2\% | 8.3\% | 9.3\% | 11.6\% | 5.7\% | 10.4\% | 6.2\% | 10.6\% | 5.2\% |
| 33'44'5-PeCB(\#126) | 8.6\% | 8.9\% | 9.4\% | 15.9\% | 9.6\% | 9.7\% | 8.9\% | 10.9\% | 7.9\% |
| 33'44'55'-НxCB(\#169) | 8.4\% | 23.1\% | 18.7\% | 18.5\% | 16.0\% | 7.6\% | 10.5\% | 15.0\% | 7.6\% |
| 2'344'5-PeCB(\#123) | 11.2\% | 10.7\% | 10.0\% | 14.8\% | 8.8\% | 12.9\% | 9.0\% | 7.3\% | 6.9\% |
| 23'44'5-PeCB(\#118) | 6.9\% | 8.3\% | 18.1\% | 13.0\% | 6.9\% | 20.1\% | 5.7\% | 7.1\% | 7.7\% |
| 233'44'-PeCB(\#105) | 8.3\% | 8.4\% | 18.5\% | 12.3\% | 8.7\% | 12.2\% | 6.2\% | 5.8\% | 8.4\% |
| 2344'5-PeCB(\#114) | 9.8\% | 12.2\% | 20.6\% | 19.0\% | 11.0\% | 12.6\% | 5.9\% | 10.9\% | 9.1\% |
| 23'44'55'-НxCB(\#167) | 8.1\% | 8.3\% | 9.8\% | 10.0\% | 10.8\% | 9.3\% | 7.5\% | 7.1\% | 6.7\% |
| 233'44'5-HxCB(\#156) | 8.3\% | 8.2\% | 8.9\% | 9.3\% | 7.2\% | 9.4\% | 6.2\% | 7.6\% | 7.6\% |
| 233'44'5'-НxCB(\#157) | 8.8\% | 11.5\% | 9.9\% | 9.4\% | 11.3\% | 7.6\% | 6.0\% | 5.0\% | 8.1\% |
| 233'44'55'-HpCB(\#189) | 7.2\% | 9.8\% | 7.7\% | 8.2\% | 9.0\% | 8.3\% | 7.1\% | 8.9\% | 5.3\% |
| PCDDs/DFs |  |  | 8.5\% | 15.4\% | 9.9\% | 10.6\% | 8.9\% | 9.1\% | 7.1\% |
| DL-PCB |  |  | 12.5\% | 13.2\% | 9.6\% | 10.9\% | 7.1\% | 9.0\% | 7.1\% |
| PCDDs/DFs, DL-PCB |  |  | 10.2\% | 14.5\% | 9.8\% | 10.7\% | 8.1\% | 9.1\% | 7.1\% |

Used GC column for 1,2,3,7,8-PeCDF and 1,2,3,4,7,8-HxCDF analysis
( $※$ a) BPX-DXN, DB-5, BPX-5, RH-12ms etc.: separate single peak
(※b) SP-2331, CP-Sil88 etc.: including co-elute congeners

Figure 1 shows Z-sore exceed $> \pm 3$ laboratory numbers in individual congeners in R-9 in 2011. Generally results from around $90 \%$ of the laboratories showed $< \pm 2$ Z-score in individual congeners data. Furthermore, reproducibility data on extraction procedure ( $\leqq 30 \%$ ) and injection ( $\leqq 10 \%$ ) showed appreciable results from many laboratories.
The trends number of laboratories whose results exceeded $> \pm 3$ of Z -score of at least one data in individual congeners, were 20 / 77 (total) for R-1, 27 / 83 (total) for R-2, 33 / 78 (total) for R-3, 23 / 75 (total) for R-4, 32 / 77 (total) for R-5, 20 / 77 (total) for R-6, 11 / 70 (total) for R-7 ,32 / 66 (total) for R-8 and 25/ 63(total) for R-9. These trends indicate that individual laboratories maintain QA/QC systems for Z-score in inter-laboratory round robin study.


Fig. 1. Z-score exceed $> \pm 3$ laboratory numbers in individual congeners (total 63 laboratories R-9 in 2011). Used GC column for 1,2,3,7,8-PeCDF and 1,2,3,4,7,8-HxCDF analysis
(※a) BPX-DXN, DB-5, BPX-5, RH-12ms etc.: separate single peak
(※b) SP-2331, CP-Sil88 etc.: including co-elute congeners

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