# AN IMPROVED AhR BIOASSAY FOR DETERMINING DIOXIN LEVELS IN SOILS, SEDIMENTS, AND FISHES

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## Introduction

Organochlorine compounds including chlorinated polyaromatic compounds, such polychlorinated dibenzo-pdioxins and dibenzofurans (PCDD/Fs), are lipophilic, bio-accumulative, and non-biodegradable and thus are recognized as environmental endocrine disruptors. The contamination of dioxin-like compounds in the environment and foodstuffs has raised public concerns due to its potential adverse effects on human health<sup>1</sup>. Invitro aryl-hydrocarbon-receptor (AhR) reporter gene assays, e.g., the dioxin-responsive element-driven chemical activated luciferase gene expression (DRE-CALUX) bioassay, are rapid, low-cost, and semi-quantitative methods for large-scale dioxin surveillance<sup>2</sup>. In the decade since the development of the DRE-CALUX bioassay, its sensitivity has been improved and its applicationshave been continuously expanded. Recently, two new invitro AhR reporter gene assays were developed using the mouse hepatoma cell line Hepa1c1c7, namely the DR-EcoScreen cells/DR-cell assay<sup>3</sup> and the H1L7.5c3 cells/CALUX bioassay<sup>4</sup>. The sensitivity of the *in-vitro* AhR reporter gene assay was improved by reducing the incubation temperature from 37 to 33°C and adding dexamethasone (DEX) and phorbol-12-myristate-13-acetate (PMA) to the cell culture medium<sup>4,5</sup>. Our *in-vitro* AhR reporter gene assay (Huh7-DRE-Luc cells/DRE-driven luciferase assay) combined with a fast cleanup system (a coupled carbon-acid silica column; Cape Technologies, South Portland, ME) was previously reported and data from analyses of dioxins in soil and sediment samples using high-resolution gas chromatograph/highresolution mass spectrometry (HRGC/HRMS) method were well correlated with those using the AhR reporter gene assay<sup>6</sup>. The present study aimed to improve the sensitivity of our AhR reporter gene assay by modification of some cell culture conditions.

### Materials and methods

Standard solutions of PCDD/Fs including labeled compound stock, internal standard spiking, cleanup standard spiking, and EPA Method 1613 calibration and verification solutions were obtained from Wellington Laboratories (Ontario, Canada). A liquid standard of seventeen 2,3,7,8-substituted PCDD/F congeners (Cat.# EDF-5416) known as Cerilliant Analytical Reference Standards was from Cambridge Isotope Laboratories, Inc. (Cambridge, MA, USA). CRM DX-1 is a certified dioxins- and furans-containing reference material (reference sediment DX-1 from the National Water Research Institute, Environment Canada, Quebec, Canada). All solvents were pesticide residue grade from Merck (Darmstadt, Germany), Tedia (Fairfield, OH, USA), and Sigma-Aldrich (St. Louis, MO, USA). PMA was purchased from Merck (Darmstadt, Germany). The rapid cleanup system was a coupled acid silica column-activated carbon mini-column (CAPE Technologies, South Portland, ME, USA). Silica gel (100–200 mesh) was purchased from Fisher (Leicestershire, England).

The environmental samples of soils (n=6), sediments (n=7), and fish (n=3) were collected from Taiwanese dioxin-contaminated areas in 2010. The samples (10 grams each) were extracted with 300 ml of toluene using a Soxhlet extractor. Extracted samples were evaporated to near dryness and then transferred to a coupled carbon-acid silica column for cleanup. The cleanup procedure was previously described<sup>6</sup>. For the HRGC/HRMS method, all samples were spiked with the different PCDD/F internal standards before extraction and the labeled cleanup standards for PCDD/Fs analysis were added before the cleanup procedure. The extract was analyzed using a HRGC/HRMS system (HP6890/JEOL JMS-700; Hewlett-Packard, Palo Alto, CA; JEOL, Tokyo, Japan) equipped with a DB-5MS 60 m column (J&W Scientific Folsom, CA, USA). The quality assurance (QA)/quality control (QC) procedures met the criteria of the Taiwanese Environmental Protection Administration (EPA).

World Health Organization 1998 (WHO<sub>1998</sub>) toxic equivalency factors (TEFs) and international (I)-TEFs were used in this study<sup>1</sup>. For the bioassay, the labeled standards were not added in the extraction and cleanup procedure. After the extraction and cleanup procedure, the final extracts were evaporated to dryness and dissolved in dimethyl sufoxide (DMSO) (100 µl) for the DRE-driven luciferase bioassay<sup>6</sup>. In brief, Huh7-DRE-Luc cells in DMEM medium were seeded at a density of  $1 \times 10^4$  cells/well on a 96-well white plate (cat. 36101, Nalge Nunc, Roskilde, Denmark). Following incubation for 24 h, the cells were treated with extracts for 24 h. In parallel, other cells were also treated with different concentrations of TCDD (0.1, 0.3, 1, 3, 10, 30, 100, 300, 1000, 3000, and 10,000 pM) for 24 h to generate a calibration curve for the assay of TCDD-dependent DREdriven luciferase activity. If PMA was needed in the extracts or the standards, 10 µL of 8 µM PMA was added to the recombinant hepatoma cells at the 16<sup>th</sup> hour of the incubation period. After treatments, 50 µl of 1×lysis buffer (Promega, Madison, WI, USA) was added per well. Cells were freeze-thawed three times using liquid nitrogen, vortexed (90 rpm at 37°C for 10 min to complete the cell lysis, and assayed using the standard protocol of the Luciferase Assay System (GLOMAX, Promega) to determine luciferase activity. The luciferase activity was expressed as relative light units (RLU)/well. The values of the sigmoid semi-logarithmic TCDD dose-response calibration curve were fitted using the Hill equation<sup>6</sup>. The luciferase response was converted to a bioanalytical equivalent (BEQ) by comparison of the extract induction responses to a standard curve of TCDD concentrations. BEQs of each sample were independently measured three times. Exploring the correlation between the HRGC/HRMS method and the modified DRE-driven Luciferase bioassay, dioxin I-TEQ levels in soils, sediments, and fish were predicted by BEQ values from this bioassay. The statistical analyses in this study were performed using SPSS 12.0 version.

## **Results and discussion**

This is the first report to examine whether both reducing the incubation temperature (i.e., to 35°C) and adding non-AhR ligand (i.e., PMA) affect TCDD-induced luciferase activation in mammalian recombinant hepatoma cells. Our AhR reporter gene assay cell line, Huh7-DRE-Luc, when treated with TCDD  $\leq$  1000 pM, showed the highest fold induction of luciferase activity by TCDD at incubation temperature 35°C (Fig, 1A); two previous studies reported a similar finding<sup>4,5</sup>. In the report by Zhao et al., luciferase activity of a recombinant Hepa1c1c7 cell, i.e., H1L6.1c2, treated with sediment extracts at 33°C was significantly increased (5–15 fold) above the luciferase activity of the same cells with the same treatment at 37°C<sup>5</sup>. Using Hepa1c1c7 cells stably transfected with a new CALUX plasmid with an increased DRE number, namely H1L7.5c3 or third generation (G3) CALUX cells, the same research team showed that TCDD treatment at 33°C caused a 4- to 10-fold increase in luciferase activity above that caused by TCDD treatment at 37°C<sup>4</sup>. Fig. 1B also shows that addition of PMA significantly enhanced the TCDD-induced luciferase activation by 1.8–3.2 fold, consistent with previous findings<sup>5</sup>. Results in Fig. 1 clearly show that incubation of Huh7-DRE-Luc cells with PMA at 35°C markedly improved the sensitivity of the cells used in TCDD detection.

Huh7-DRE-Luc cells were treated with TCDD at 35°C in the presence of 0.8  $\mu$ M PMA and a dose–response induction of luciferase activity by TCDD was demonstrated in Figure 2A, showing a semi-logarithmic dose-response calibration curve with a sigmoid appearance (R<sup>2</sup>>0.995, *p*<0.001). Moreover, Huh7-DRE-Luc cells were treated with a liquid standard (EDF-5416) or a certified reference material (DX-1) at 35°C in the presence of PMA. As shown in Table 1, the resultant BEQ levels by EDF-5416 (10X dilution) or DX-1 (1X, 10X, and 100X dilution) were 4.22–5.71 times higher than WHO<sub>1998</sub>-TEQ levels. As compared with the previous assay using Huh7-DRE-Luc cells treated at 37°C<sup>6</sup>, stability and sensitivity of the present AhR reporter gene assay using the cells treated at 35°C in the presence of PMA were obviously improved (data not shown).

As shown in Table 2, levels of PCDD/Fs range from 49.0 to 2760 ng BEQ/kg dry weight (d.w.) for 6 soil samples, from 25.6 to 6570 ng BEQ/kg d.w. for 7 sediment samples, and from 13.1 to 239 ng BEQ/kg wet weight (w.w.) for 3 fish samples. The relative standard deviations (RSD) of the tested samples were between 4.01 and 16.8%. PCDD/F measurements with RSD value  $\leq 15\%$  meet the Taiwanese EPA criteria of the *in-vitro* AhR bioassay method (NIEA S901.60B). Notably, fish viscera have a significantly higher BEQ value than whole fish and fish meat. Because levels of I-TEQs and BEQs of 16 samples were non-normally distributed by the Kolmogorov-Smirnov test, the data were logarithmically transformed to approximate normal distribution. The relationship between I-TEQs and BEQs was determined by a linear regression model with the equation: log (PCDD/Fs I-TEQs) =  $1.19 \times \log$  (BEQs)–1.15 (R<sup>2</sup>=0.95) (Fig. 2B).

Several strategies have been used in the *in-vitro* AhR reporter gene assay to improve the sensitivity of dioxin analysis. The strategies include modification of the original reporter plasmid<sup>7</sup>, development of a new recombinant reporter plasmid<sup>8</sup>, increase in the copy number of DRE in the promoter<sup>4,5</sup>, addition of PMA or DEX to the cell culture medium<sup>5</sup>, and reduction in incubation temperature from 37°C to 33°C<sup>4,5</sup>. Among these strategies, PMA or DEX addition to the cell culture medium and reduction in incubation temperature from 37°C to 33°C<sup>4,5</sup>. Among these strategies, PMA or DEX addition to the cell culture medium and reduction in incubation temperature from 37°C to 33 or 35°C are the easier ways to improve detection sensitivity of already established *in-vitro* AhR reporter gene assays, such as our Huh7-DRE-Luc cells/DRE-driven luciferase assay. We have previously demonstrated that the addition of a coupled carbon-acid silica column (CAPE) to the Huh7-DRE-Luc cells/DRE-driven luciferase assay is a rapid and useful semi-quantitative green technique for large-scale dioxin surveys<sup>6</sup>. Here, we further show that altering the incubation conditions (adding PMA at 35°C) is able to markedly enhance the sensitivity of the Huh7-DRE-Luc cells/DRE-driven luciferase assay of dioxin, suggesting a potential environmentally friendly technique for high-throughput screening of dioxins. This high-throughput screening technique in conjunction with the HRGC/HRMS method may significantly facilitate large-scale survey for dioxin, particularly for environmental dioxin in heavily contaminated areas and dioxin contamination in foodstuffs.

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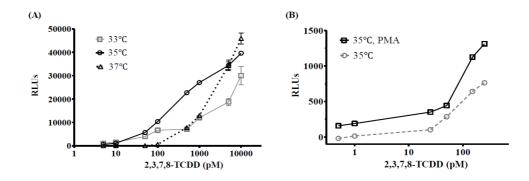


Figure 1. Effects of temperature and PMA on the Huh7-DRE-Luc cells/DRE-driven luciferase assay. (A) Huh7-DRE-Luc cells were treated with TCDD for 24 hours at 33, 35, and 37°C. (B) The cells were also treated with TCDD as indicated for 24 hours at 35°C in the presence of 0.8  $\mu$ M PMA. After treatments, luciferase activity was determined. Data are presented as means  $\pm$  SD (n = 3) and are expressed as relative light units (RLUs).

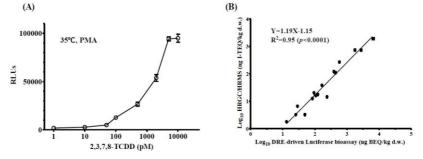


Figure 2. (A) Dose–response induction of luciferase activity by TCDD. Huh7-DRE-Luc cells were treated with TCDD for 24 hours at 35°C in the presence of 0.8  $\mu$ M PMA. After treatments, luciferase activity was determined. Data are presented as means  $\pm$  SD (n=3) and are expressed as relative light units (RLUs). (B) A linear regression was determined between the HRGC/HRMS assay values and the bioassay values (R<sup>2</sup>=0.95, p<0.001, n=16). Thirteen environmental samples (6 from soil and 7 from sediment) and 3 biological samples (from fish) were included. The mathematic equation was log (PCDD/Fs I-TEQs) = 1.19 × log (BEQs) – 1.15.

| Table 1. Huh7-DRE-Luc cells treated with a liquid stan | ndard (EDF-5416) or a certified reference |
|--|---|
|--|---|

| Certified standards   |   | DRE-driven luciferase bioassay |      |                          |      |      |       |
|-----------------------|---|--------------------------------|------|--------------------------|------|------|-------|
| Standard              | Concentration                                 | Na                             | Mean | Range                    | SD   | RSD  | Ratio |
| EDF-5416              |   |                                |      |                          |      |      |       |
| 10X °                 | 7.50 <sup>d</sup>                             | 9                              | 42.9 | 37.4-46.4                | 4.29 | 10.0 | 5.71  |
| (ng WHO1998-TEQ/mL)   |   |                                | (1   | ng BEQ/mL <sup>e</sup> ) |      | (%)  |       |
| CRM DX-1 <sup>g</sup> |   |                                |      |                          |      |      |       |
| 1X                    | 466   | 6                              | 1970 | 1894-2039                | 121  | 6.15 | 4.22  |
| 10X                   | 46.6  | 12                             | 221  | 113-279                  | 28.7 | 13.0 | 4.75  |
| 100X                  | 4.66  | 9                              | 23.6 | 15.9-28.6                | 2.74 | 11.6 | 5.06  |
| (ng WH                | O <sub>1998</sub> -TEQ/kg d.w. <sup>f</sup> ) |                                | (ng  | g BEQ/kg d.w.)           | )    | (%)  |       |

<sup>a</sup> N is the number of measurements.

<sup>b</sup>Ratio=(the BEQ value of the bioassay)/(the certified TEQ value of the standard)

<sup>c</sup> 10X means that the standard was diluted to 1/10.

 $^{\rm d}$  The original concentration was 75.0 ng WHO  $_{\rm 1998}\text{-}TEQ/mL.$ 

<sup>e</sup>ng BEQ/mL.

<sup>f</sup>ng WHO<sub>1998</sub>-TEQ/kg dry weight (d.w.).

 Table 2. Analysis of PCDD/F levels in environmental and biological samples with Huh7-DRE-Luc cells/DRE-driven luciferase assay

| Samples                   | Ν  | Mean | Range     | SD    | RSD (in %) |
|---------------------------|----|------|-----------|-------|------------|
| Soil <sup>a</sup>         |    |      |           |       |            |
| Soil-1                    | 15 | 2760 | 1050-4520 | 204   | 7.37       |
| Soil-2                    | 18 | 394  | 182-789   | 49.1  | 12.5       |
| Soil-3                    | 6  | 166  | 115-216   | 17.66 | 4.27       |
| Soil-4                    | 3  | 84.0 | 55.4-144  | 10.08 | 12.0       |
| Soil-5                    | 3  | 107  | 81.0-124  | 4.28  | 4.01       |
| Soil-6                    | 3  | 49.0 | 32.3-76.4 | 4.40  | 8.98       |
| Sediment a                |    |      |           |       |            |
| Sediment-1                | 18 | 6570 | 3270-9940 | 952   | 14.5       |
| Sediment-2                | 12 | 584  | 473-753   | 61.3  | 10.5       |
| Sediment-3                | 9  | 1760 | 1650-2510 | 203   | 11.6       |
| Sediment-4                | 6  | 124  | 119-128   | 10.6  | 8.58       |
| Sediment-5                | 6  | 94.0 | 81.9-106  | 6.20  | 6.59       |
| Sediment-6                | 3  | 440  | 343-555   | 44.1  | 10.0       |
| Sediment-7                | 3  | 25.6 | 12.7-52.7 | 3.05  | 11.9       |
| Fish                      |    |      |           |       |            |
| Whole fish b              | 24 | 13.1 | 2.31-19.3 | 2.10  | 16.1       |
| Fish meat <sup>c</sup>    | 24 | 28.8 | 20.2-36.6 | 3.46  | 12.0       |
| Fish viscera <sup>d</sup> | 27 | 239  | 96.1-314  | 40.0  | 16.8       |

<sup>a</sup> The unit was ng BEQ/kg d.w.

<sup>b</sup> The whole fish of Perth herring (Nematalosa come).

<sup>c</sup> The meat of Tilapia (Oreochromis niloticus niloticus).

<sup>d</sup> The viscera of Tilapia (*Oreochromis niloticus niloticus*); the unit was ng BEQ/kg w.w.