INTRODUCTION

Breast cancer is one of the cancer types with a high percentage of mortality in Indonesia [1]. Most of the cases are caused by estrogen receptor (ER)-positive cancer, so anticancer from selective estrogen receptor modulators (SERMs) are usually used as the first line of the adjuvant therapy [2]. Tamoxifen (TAM) is an anticancer of SERMs and it is a prodrug that will be metabolized by CYP2D6 into active metabolites that have higher affinity to ER than TAM itself [3-5]. TAM is metabolized by CYP2D6 enzyme into endoxifen (END) and 4-hydroxytamoxifen (4-HT), both of which are metabolites with antiestrogenic activity 30 to 100-fold more potent than TAM itself [5]. Cytochrome 2D6 (CYP2D6) enzyme is polymorphic, so it might result in different metabolic activities for each patient [3-5]. TAM analysis has been done mostly using plasma samples, but plasma sampling of breast cancer patients by venipuncture is considered invasive and painful to patient [3]. Therefore, it is necessary to develop a method for analyzing TAM using sampling that considers patient comfort, namely dried blood spot (DBS), in which this method is more simple, requires less blood volume, and the analyte is considered non-invasive and painful to patient [3].

The previous study showed that the efficacy of TAM therapy depended on END threshold concentration. Patient who has END concentration above 5.9 ng/ml has less recurrence compared to patient with END level below the threshold concentration [4]. This concentration is extrapolated to DBS, resulted in threshold concentration of 3.3 ng/ml in DBS [3]. In the present study, the monitoring of TAM, END, and 4-HT through quantification in DBS breast cancer patients was developed and applied to determine the levels of TAM metabolites in the blood of breast cancer patients in Indonesia. This method can be used to evaluate the effectiveness of TAM therapy in breast cancer patients and can be considered by doctors for follow-up therapy.

MATERIALS AND METHODS

Reference standard samples and materials

TAM was purchased from Sigma-Aldrich (Singapore). N-desmethyl-4-HT (END/E/Z mixture) and 4-HT were purchased from Santa Cruz Biotechnology (USA). Clomiphene was purchased from Fabrica Italiana Sintetici (Italy). Formic acid, acetonitrile HPLC grade, and methanol for analysis were obtained from Merck (Darmstadt, Germany). Perkin Elmer 226 paper was obtained from Perkin Elmer (USA).

Preparation of solutions and standards

TAM, END, and 4-HT stock solutions were prepared in methanol to obtain a concentration of 1,000 ng/ml [5]. The stock solution was diluted to obtain intermediate solutions at 100 ng/ml. Working solutions were prepared by dilution of the intermediate solution with methanol. Calibration samples were prepared in whole blood diluting working solutions at 1:20 with whole blood free from the analytes. Clomiphene stock solution was prepared in methanol at 1,000 ng/ml concentration. Intermediate clomiphene solutions were prepared by dilution of the stock with methanol to obtain a concentration of 100 ng/ml.

Sample preparation

DBS samples were prepared from freshly drawn whole blood which spotted on the DBS paper and dried for 2 h. The method was referred to [5] with some modification. The blood spots were cut through the spot's diameter. For the extraction of the analytes from the DBS, a 100 μl methanol containing 0.1% clomiphene and 1.000 μl methanol as an extraction solvent were added to the DBS. Then
the tube was vortexed for 1 minute and sonicated for 25 minutes. 850 μl of the sample mixture was dried under a gentle stream of nitrogen (55 °C) for 15 minutes and the dried extract was dissolved in 100 μl 0.1% formic acid-0.1% formic acid in acetonitrile (35:65). The sample mixture was centrifuged for 10 minutes at 3,000 rpm, then 60 μl of the supernatant was transferred to an auto sampler vial and 10 μl was injected onto the LC-MS/MS system.

Liquid chromatography tandem mass spectrometry (LC-MS/MS)

An LC-MS/MS method was previously used for quantification of TAM and END and also 4-HT in plasma [3]. Positive electrospray ionization was applied and the transition from precursor into product was set at m/z 372.2>72.27 for TAM; 374.29>58.20 for END; 388.29>72.19 for 4-HT; and 406.28>100.17 for clomiphene.

Validation assay

Validation assay was performed based on Food and Drug Administration (2013) and European Medicines Agency (2011) guidelines for validation of bioanalysis [9, 10].

Linearity

Aliquots of 950 μl blank blood were spiked with 50 μl stock solutions to obtain eight calibration levels of samples containing TAM (5, 10, 25, 50, 100, 150, 200 ng/ml), END (1, 2, 5, 10, 15, 20, 30 and 40 ng/ml), and 4-HT (0.5, 1; 2; 5; 7.5; 10; 15 and 20 ng/ml). Blank, zero, and calibration samples were applied to Perkin Elmer 226 paper (3 x 20 μl), dried at room temperature for 2 h. Replicates at each concentration were analyzed within 2–4 h. Replicates for each concentration were analyzed as described above. Calibration curves were calculated relating the area ratios from TAM, END, and 4-HT peaks to the clomiphene peak and with the nominal concentrations of the calibration samples.

Lower limit of quantification (LLOQ)

An independent DBS quality control sample at the lowest point of the calibration curve at concentrations of 5.0; 1.0; and 0.5 ng/ml for TAM, END, and 4-HT, respectively was included in the accuracy and precision experiments. This assay was performed with five replicates. The acceptance criteria established for the LLOQ is maximum 20% for the % coefficient variation (CV) value.

Accuracy and precision

Aliquots of blank blood were spiked with standard solutions to obtain 4 types of concentration: LLOQ, Quality control sample in low, medium, high concentration (QCL, QCM, and QCH) containing TAM, END, and 4-HT. Samples were extracted with the extraction method as described above. This assay was divided into 2 types: within run and between run. Both assays were performed with five replicates. The acceptance criteria established for the accuracy and precision is maximum 20% for the %CV value of LLOQ concentration and 15% for the %CV value of QCL, QCM, and QCH concentrations.

Recovery

Aliquots of blank blood were spiked with standard solutions into obtain QCL, QCM, and QCH concentrations, then samples were extracted and injected to the LC-MS/MS system. Standards containing TAM, END, and 4-HT on QCL, QCM, and QCH concentrations were also injected to the LC-MS/MS system. Both assays were performed with three replicates for each concentration. The acceptance criteria is established for maximum 15% for the CV value.

Matrix effect

Standard solutions of TAM, END, and 4-HT were diluted in the solvent to obtain QCL and QCH concentrations, then injected into the LC-MS/MS system. Furthermore, whole blood from six different sources was extracted with the extraction method as described above, then the analytes were inserted into the samples after the extraction process to obtain QCL and QCH concentrations. The assay was performed with two replicates for each concentration. Matrix effect is showed by calculating the matrix factor. The acceptance criteria established for the matrix effect is maximum 15% for the %CV value in the calculation of standardized normalized matrix factor.

Application of the method

A total of 29 breast cancer patients who were taking adjuvant hormonal treatment with TAM (20 mg/day) were enrolled in the study. They were given a full explanation of the procedures during sampling and signed the informed consent prior to participating in this study. The study inclusion criteria were patient that has been diagnosed with breast cancer (ER+) and confirmed by histopathology report, had received TAM in their therapy regimen (20 mg/day) for at least 2 mo and aged 25-55 y old during the blood collection. The exclusion criteria were the patient who declared they were unwilling to participate in the study by not signing the informed consent sheet.

Finger prick blood samples were collected from 29 breast cancer patients of Dharmais Cancer Hospital, Jakarta, Indonesia. Around 200 μl blood samples were collected from the fingertips. Blood was drawn by finger prick technique using lancet and the first drop of blood from fingertip was thrown away by rubbing it with alcohol swab. The blood drops were collected in a 0.5 ml K3-EDTA micro tube. After that, 20 μl blood was immediately transferred to DBS paper using a calibrated micropipette and the paper was allowed to dry for 2 h at room temperature. After the drying process, the DBS paper was stored in plastic seal bag containing desiccant until the time of analysis.

This research has passed the ethical review from the Health Research Ethics Committee of the Faculty of Medicine, Universitas Indonesia with number 0076/UN2. F1/ETIK/2018 and notification of approval from the Health Research Ethics Committee of Dharmais Cancer Hospital with number 038/KEPK/III/2018.

RESULTS AND DISCUSSION

Chromatography system and preparation of samples

The LC-MS/MS system analyzed TAM, END, 4-HT, and clomiphene simultaneously with Multiple Reaction Monitoring (MRM) mass spectrometry mode. The capillary pipe voltage was 3.5 kW with 350 °C for desolvation temperature and 650 L/hour for the gas flow rate. The detection condition is shown in table 1.

| Compounds | Parent ion (m/z) | Daughter ion (m/z) | Cone (V) | Collision energy (V) |
|-----------|-----------------|-------------------|----------|---------------------|
| TAM       | 372.2           | 72.27             | 50       | 27                  |
| END       | 374.29          | 58.20             | 45       | 30                  |
| 4-HT      | 388.29          | 72.19             | 50       | 27                  |
| Clomiphene| 406.28          | 100.17            | 45       | 25                  |

Optimal conditions for the analysis can be obtained by optimizing the analysis and sample preparation method. Optimization of the mobile phase combination, composition, and flow rate was done to obtain a good separation. The mobile phase consisted of formic acid 0.1%-formic acid 0.1% in acetonitrile (35:65) using an isocratic system with a flow rate of 0.25 ml/min. The analysis was performed in 4 min and peaks were separated in a short retention time; TAM (1.86 min), END (1.06 min), 4-HT (1.1 min), and clomiphene (1.8 min).

Sample preparation is an important factor in bioanalysis because biological samples contain endogenous materials that can interfere with the analysis of compounds. Appropriate sample preparation is
necessary to minimize the disturbances in the analyses, so the accuracy of the analysis results can be obtained. Protein preparation was chosen because of its simplicity and efficiency [11, 12].

The optimum separation conditions for TAM, END, and 4-HT using the protein precipitation method with 1,000 μl methanol as the extraction solution and sonication was applied for 25 min. The samples were injected 10 μl into the LC-MS/MS system using a 0.1% formic acid-0.1% formic acid in acetonitrile (35:65) phase with 0.25 ml/min resulted in retention time for TAM at 1.86 min, END at 1.11 min, 4-HT at 1.15 min, and clomiphene at 1.41 min.

The method proposed in the present study has several advantages over the methods reported in the literature for TAM, such as short analysis time (4 min per sample), simple sample preparation, and more convenient for the patient. Moreover, the method was used to measure the concentration of TAM and its metabolites (END and 4-HT) for a clinical study of breast cancer patients.

Validation assay

Validation assay has been conducted in the previous study and in the same laboratory. As shown in table 2, the intra and inter-assay precision and accuracy experiments performed on LLOQ and QC samples (QCL, QCM, and QCH) were fulfilled the requirement of EMA and FDA. Within run and between run imprecisions were in the 3.56 to 8.67% and 4.2 to 10.82% ranges, respectively. Accuracy within run and between run was estimated at % bias -9.47 to 14.88% and 19.63 (LLOQ) to 11.4%. LC with MS/MS methods should exclude any matrix effects. The effect of matrix was showed by analysis between the analyte area added after the blank extraction process compared to the area of standard solution in the solvent. As shown in table 2, this method showed acceptable matrix effect ranging from 85.78 to 99.71% with variability (CV) under 10%.

| Analyte  | QC     | Conc. (ng/ml) | Precision (% CV) | Accuracy (% bias) | Matrix effect | Internal standard-normalized matrix factor |
|----------|--------|---------------|------------------|-------------------|--------------|------------------------------------------|
| TAM      | LLOQ   | 5.00          | 4.76 ± 2.94     | -16.95            | 85.78±2.32   | 0.89±0.02                                |
|          | QCL    | 25.00         | 4.71 ± 2.94     | -9.90             | 98.55±0.46   | 1.02±0.01                                |
|          | QCM    | 75.00         | 5.24 ± 2.94     | -10.16            | 93.71±2.74   | 0.97±0.03                                |
| END      | LLOQ   | 5.00          | 6.27 ± 3.42     | -19.63            | 92.93±6.35   | 0.96±0.08                                |
|          | QCL    | 150.00        | 4.02 ± 2.94     | -6.15             | 14.88±7.36   | 0.91±0.07                                |
|          | QCM    | 30.00         | 5.20 ± 3.42     | -12.29            | 87.86±7.36   | 0.93±0.08                                |
| 4-HT     | LLOQ   | 0.50          | 8.67 ± 3.42     | -18.11            | 99.61±7.71   | 1.03±0.08                                |
|          | QCL    | 2.50          | 5.72 ± 3.42     | -11.40            | 80.79±1.32   | 0.96±0.13                                |
|          | QCM    | 7.50          | 7.66 ± 3.42     | -4.99             | 81.87±7.80   | 0.97±0.01                                |
|          | QCH    | 15.00         | 4.65 ± 3.42     | -12.22            | 79.61±7.71   | 0.91±0.07                                |

Table 2: Summary of full validation results

| Analyte  | Linear regression | QC     | Conc. (ng/ml) | Precision (% CV) | Accuracy (% bias) | Recovery %±SD | Mean ±SD | % CV |
|----------|-------------------|--------|---------------|------------------|-------------------|---------------|----------|-------|
| TAM      | y=0.0143+0.0115x  | LLOQ   | 5.00          | 4.90 ± 2.94      | -7.70             | 71.27±4.34   | 6.09     |
|          | r=0.9996          | QCL    | 25.00         | 6.34 ± 2.94      | -12.90            | 81.15±4.33   | 5.34     |
|          | END               | QCM    | 75.00         | 5.42 ± 2.94      | -8.25             | 85.78±0.46   | 1.02±0.01|
|          | y=0.0003+0.0017x  | QCH    | 150.00        | 5.46 ± 2.94      | 7.27              | 93.71±2.74   | 0.97±0.03|
|          | r=0.9977          | LLOQ   | 1.00          | 4.37 ± 2.94      | 18.69             | 92.93±6.35   | 0.96±0.08|
|          | END               | QCL    | 5.00          | 5.47 ± 2.94      | -10.88            | 87.86±7.36   | 0.91±0.07|
|          | y=0.0029+0.0034x  | QCM    | 15.00         | 2.18 ± 2.94      | -13.11            | 81.87±7.80   | 9.52     |
|          | r=0.9958          | QCH    | 30.00         | 3.14 ± 2.94      | 6.29              | 85.78±0.46   | 1.02±0.01|
|          | 4-HT              | LLOQ   | 0.50          | 14.44 ± 2.94     | 17.60             | 85.78±0.46   | 1.02±0.01|
|          | y=0.0029+0.0034x  | QCL    | 2.50          | 10.25 ± 2.94     | 11.01             | 81.87±7.80   | 9.52     |
|          | r=0.9958          | QCM    | 7.50          | 7.36 ± 2.94      | -9.50             | 85.78±0.46   | 1.02±0.01|
|          |                   | QCH    | 15.00         | 5.13 ± 2.94      | 6.73              | 81.87±7.80   | 9.52     |

Table 3: Summary of partial validation results

The partial validation was performed in this study as shown in table 3. The linearity of each calibration curve was determined by plotting the peak area ratio (y) of the analyte to the internal standard versus the nominal concentration (x) of TAM, END, and 4-HT. The calibration curves were linear over the concentration range of 5.0-200.0 ng/ml for TAM; 1.0-40.0 ng/ml for END; and 0.5-20.0 ng/ml for 4-HT with a correlation coefficient of 0.99 for those three compounds.

Accuracy and precision were calculated using within run and between run test at LLOQ, QCL, QCM, and QCH concentration levels. The accuracy (%bias value) TAM, END, and 4-HT were less than 20%. The precision (%CV value) of TAM 4.90-6.34%; 2.18-4.37% for END; and 5.13-14.44% for 4-HT. Recovery was calculated using three replicates at three concentration levels: QCL, QCM, and QCH. The mean recovery (%recovery value) for TAM, END, and 4-HT was 71.27±4.34 %; 81.15±4.33%; and 81.87±7.80 with %CV value less than 15% for all compounds. The results showed that the method is reliable within the analytical range.

This method was applied for the determination of DBS concentration of TAM, END, and 4-HT in 29 patients with a daily dose of 20 mg TAM. Sampling was conducted in outpatient subject that...
already reach steady state level of TAM. The results of the analysis on 29 samples showed that all samples contained TAM, END and 4-HT. Our data showed a wide range of drug DBS concentrations among patients.

### Table 4: Result of analysis in breast cancer patients

| Patient | Measured concentration (ng/ml) | TAM | END | 4-HT |
|---------|-------------------------------|-----|-----|------|
| SN 01   | 164.61                        | 25.30 | 5.74 |
| SN 02   | 144.44                        | 24.10 | 4.88 |
| SN 03   | 114.12                        | 28.77 | 4.86 |
| SN 04   | 105.33                        | 24.84 | 7.36 |
| SN 05   | 139.95                        | 18.42 | 6.64 |
| SN 06   | 104.97                        | 6.62  | 4.32 |
| SN 07   | 183.52                        | 26.63 | 0.72 |
| SN 08   | 58.27                         | 6.72  | 6.58 |
| SN 09   | 98.21                         | 16.97 | 6.08 |
| SN 10   | 94.32                         | 17.10 | 6.73 |
| SN 11   | 67.15                         | 9.29  | 8.19 |
| SN 12   | 140.28                        | 10.88 | 1.00 |
| SN 13   | 127.88                        | 9.50  | 2.13 |
| SN 14   | 160.31                        | 4.55  | 3.21 |
| SN 15   | 178.72                        | 19.43 | 6.46 |
| SN 16   | 61.57                         | 13.95 | 11.28|
| SN 17   | 100.97                        | 21.56 | 3.11 |
| SN 18   | 154.82                        | 9.42  | 4.59 |
| SN 19   | 138.81                        | 10.72 | 3.36 |
| SN 20   | 120.02                        | 12.79 | 5.43 |
| SN 21   | 188.63                        | 18.18 | 7.26 |
| SN 22   | 13.38                         | 11.07 | 2.56 |
| SN 23   | 96.03                         | 3.75  | 1.87 |
| SN 24   | 39.68                         | 5.36  | 0.96 |
| SN 25   | 30.29                         | 2.22  | 0.21 |
| SN 26   | 54.46                         | 1.45  | 1.45 |
| SN 27   | 39.68                         | 5.96  | 0.44 |
| SN 28   | 35.23                         | 8.70  | 1.21 |
| SN 29   | 39.20                         | 3.98  | 1.45 |

The lowest TAM level was found in patient SN 25 of 30.29 ng/ml and the highest level of TAM which is 188.63 ng/ml was present in patient SN 21. The mean value of TAM levels in DBS was 110.02 ng/ml with standard deviation of 47.68 and the coefficient of variation of 43.33%.

The variation between patients was also found in the results of END analysis. The lowest END levels were present in patient SN 26 of 1.45 ng/ml and the highest levels of 28.77 ng/ml were present in patient SN 03. END levels in DBS had an average value of 13.04 ng/ml with standard deviation of 8.02 and the variation coefficient of 61.48%. The results of 4-HT analysis also showed variation between patients. The lowest 4-HT level of 0.21 ng/ml was found in patient SN 25 and the highest level of 11.28 ng/ml was found in patient SN 16. 4-HT level in DBS had an average value of 4.13 ng/ml with standard deviation of 2.79 and the variation coefficient of 67.75%. Based on the data obtained, it can be concluded that TAM, END, and 4-HT levels in the DBS between patients vary widely. The graph of TAM, END, 4-HT analysis in breast cancer patients is shown on fig. 1. The analytical result data is shown on table 4. Concentration range of END and 4-HT found in this study showed high variability. This shows that on the same dosage, different metabolism in every patient lead to different level of metabolites formation, thus affecting TAM efficacy.
The efficacy of TAM therapy in breast cancer patients is correlated to the END clinical threshold in blood. Based on the previous study, 96% patient with END serum concentration above 5.9 ng/ml had a 26% reduction of recurrences comparing with patients with END concentrations below the threshold [4]. This concentration is extrapolated to DBS, resulted in END threshold concentration of 3.3 ng/ml in DBS. Based on the END threshold concentration, 2 patients have END concentration below the clinical threshold. The variation of concentrations may be attributed to the inter-individual pharmacokinetic variability, including polymorphisms of CYP2D6 and CYP3A4 and drugs interaction [5, 13].

CONCLUSION

The developed method for determination TAM, END, and 4-HT using LC-MS/MS has a simple sample preparation and rapid analysis run time. This method met the 2011 European Medicines Agency (EMA) and 2013 Food and Drug Administration (FDA) bioanalytical guideline for validations. This method was successfully applied to 29 breast cancer patients, including 2 patients with END level below the threshold concentration.

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AUTHORS CONTRIBUTIONS

All authors have contributed equally

CONFLICTS OF INTERESTS

Declared none

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