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Original Article

Rapid investigation of α -glucosidase inhibitory activity of *Phaleria macrocarpa* extracts using FTIR-ATR based fingerprinting



Sabina Easmin^a, Md. Zaidul Islam Sarker^{a,*}, Kashif Ghaffoor^b,
Sahena Ferdosh^c, Juliana Jaffri^a, Md. Eaquib Ali^d, Hamed Mirhosseini^e,
Fahad Y. Al-Juhaimi^b, Vikneswari Perumal^a, Alfi Khatib^a

^a Faculty of Pharmacy, International Islamic University, Kuantan Campus, Pahang, Malaysia

^b Department of Food Science and Nutrition, King Saud University, Riyadh, Saudi Arabia

^c Faculty of Science, International Islamic University, Kuantan Campus, Pahang, Malaysia

^d Nanotechnology and Catalysis Research Centre, University of Malaya, Kuala Lumpur, Malaysia

^e Faculty of Food Science and Technology, Universiti Putra Malaysia, UPM Serdang, Selangor DE, Malaysia

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ABSTRACT

Phaleria macrocarpa, known as “Mahkota Dewa”, is a widely used medicinal plant in Malaysia. This study focused on the characterization of α -glucosidase inhibitory activity of *P. macrocarpa* extracts using Fourier transform infrared spectroscopy (FTIR)-based metabolomics. *P. macrocarpa* and its extracts contain thousands of compounds having synergistic effect. Generally, their variability exists, and there are many active components in meager amounts. Thus, the conventional measurement methods of a single component for the quality control are time consuming, laborious, expensive, and unreliable. It is of great interest to develop a rapid prediction method for herbal quality control to investigate the α -glucosidase inhibitory activity of *P. macrocarpa* by multicomponent analyses. In this study, a rapid and simple analytical method was developed using FTIR spectroscopy-based fingerprinting. A total of 36 extracts of different ethanol concentrations were prepared and tested on inhibitory potential and fingerprinted using FTIR spectroscopy, coupled with chemometrics of orthogonal partial least square (OPLS) at the 4000–400 cm^{-1} frequency region and resolution of 4 cm^{-1} . The OPLS model generated the highest regression coefficient with $R^2Y = 0.98$ and $Q^2Y = 0.70$, lowest root mean square error estimation = 17.17, and root mean square error of cross validation = 57.29. A five-component (1+4+0) predictive model was built up to correlate FTIR spectra with activity, and the responsible functional groups, such as $-\text{CH}$, $-\text{NH}$, $-\text{COOH}$, and $-\text{OH}$, were identified for the bioactivity. A successful multivariate model was constructed using FTIR-attenuated total reflection as a simple and rapid technique to predict the inhibitory activity.

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* Corresponding author. Faculty of Pharmacy, International Islamic University, Kuantan Campus, Jalan Sultan Ahmad Shah, Bandar Indera Mahkota, 25200 Kuantan, Pahang, Malaysia.

E-mail address: zaidul@iiu.edu.my (Md.Z.I. Sarker).

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1. Introduction

Phaleria macrocarpa (Scheff.) Boerl, also known as Mahkota dewa, belongs to Thymelaeaceae family and is one of the most familiar and widely used traditional medicines in South Asian countries. This plant is used for the prevention of liver disease, diabetes mellitus, gout, kidney disorders, cancer, heart disease, impotence, allergies, insect bites, hemorrhoids, strokes, blood-related disease, migraines, and acne and skin ailments [1]. In addition, *P. macrocarpa* also possess various types of metabolites, including carbohydrates, saponins, tannins, alkaloids, flavonoids, polyphenolic glycosides, palmitic acid, phenols, terpenoids, dodecanoic acid, lignans, ethyl stearate, and polyphenolic compounds. Especially in Malaysia and Indonesia, a large number of people use this plant in processed or unprocessed forms, such as tea and juice, as antidiabetic remedies. However, only a few compounds, such as phalerin [1], carbohydrates (structure is suspected analogous with myglitol or acarbose) [2], and magniferin [3], have been reported for the α -glucosidase inhibitory activity of *P. macrocarpa* based on bioassay-guided traditional method. There are several antidiabetic activities of *P. macrocarpa* that have been reported recently, such as the highest α -glucosidase inhibitory activity was observed in the methanol extract of the flesh and n-hexane extract of the stem of *P. macrocarpa* [4] and reduction of plasma glucose level in the Sprague Dawley rats after the treatment with methanolic extract of *P. macrocarpa* [5].

Metabolomics is a novel, holistic, and rapidly emerging field that provides a comprehensive chemical analysis of all existing metabolites, their variation, and metabolic pathway in a biological system ensuring the production of robust and high-quality data in an unbiased way [6].

Assuring the identity, safety, and quality of herbal products is a growing concern for the extravagantly complicated and unknown mechanism of disease treatment. In a herbal material, there are many active components present in meager amounts, and they cannot be isolated easily because of their variability. Consequently, the classical isolation and measurement methods are laborious, expensive, time consuming, unreliable, and involve toxic chemicals for the quality control of these herbs [7]. Therefore, there is a need to use more crude extracts (polytherapy). Moreover, only a few isolated pharmacologically active principles cannot represent the bioactivity of complex herbal extracts, and the activity of herbal preparation arises from combined interactions of thousands of constituents and their synergist/antagonist consequences [8,9]. Therefore, it is important to develop a rapid and accurate analytical method for ensuring bioactivity and safety of herbal materials. To overcome this limitation, metabolomics combined with several common analytical techniques have become an important tool for multicomponent analysis of herbal extracts where multiple components can be assessed in a single analysis [10].

There are a variety of spectroscopic techniques for herbal fingerprint analysis, such as gas chromatography mass spectrometry, LC–MS, and nuclear magnetic resonance (NMR) spectroscopy [11]. Although these are the principal analytical methods applied in plant metabolomics, they have some

advantages, such as comprehensiveness, sample throughput, and precision. Additionally, these systems are not constantly available in developing countries where natural products are generally well known. The application of most available, simple, and rapid methods like Fourier transform infrared spectroscopy (FTIR) could be an efficient alternative for herb quality control [12]. FTIR spectroscopy, used to determine functional groups, is reagentless, cost effective, rapid, nondestructive, and suitable for investigations that ensure high reproducibility, specificity, easy, and minimal sample preparation [13]. In addition, the functional groups absorb the incoming FTIR radiation at specific wavelength and chemical bonds vibrate in characteristic ways, such as bending or stretching, leading to a spectrum or “fingerprint” [14,15]. Recently, researches have inserted this idea to manage big data sets to characterize samples. For instance, metabolomics together with projection-based multivariate data analysis was successfully applied for obtaining reliable results [16,17].

The objective of this study was to develop a rapid multivariate predictive model based on orthogonal partial least square (OPLS) method combined with FTIR technique by analyzing a metabolic fingerprint and chemical changes in the different ratios of ethanol–water extracts of *P. macrocarpa*. The model has been established from a correlation between the α -glucosidase inhibitory activity and FTIR spectra of *P. macrocarpa*. To the best of our knowledge, this is the first attempt to develop a rapid predictive model by fingerprinting the compositions of this plant under different solvent treatments using FTIR and metabolomics approach.

2. Methods

2.1. Plant materials

The fruits of *P. macrocarpa* were purchased from the farm at Kota Bharu, Kelantan, Malaysia. The fruits were identified by voucher specimens (PIUM 230-1) prior to being deposited in the herbarium of the Pharmacy Department, International Islamic University, Pahang, Malaysia.

2.2. Chemicals

The organic solvent ethanol (analytical grade), which was supplied by R & M chemicals (Essex, UK), was used for the extraction of the plant material. For enzymatic assay, enzyme (α -glucosidase) was purchased from Megazyme (Wicklow, Ireland). The substrate 4-nitrophenyl- α -D-glucopyranoside and quercetin were supplied by Sigma-Aldrich (St. Louis, MO, USA).

2.3. Sample preparation

First, the collected fruits were washed to avoid any debris. The seeds were separated, and the fruits were cut into small pieces and dried at room temperature for 1 week. The dried sample was ground to fine powder using grinding machine (Fritsch Universal Cutting Mill-115 PULVERISSETTE 19; Fritsch, Germany) and stored in refrigerator at -10°C for FTIR-ATR and enzyme analyses. A total of 36 samples, resulted from six

different concentrations of ethanol with six replications, were prepared for analysis. About 5.0 g of each grinded sample was immersed in 150 mL of ethanol and water at varying concentrations (100%, 80%, 60%, 40%, 20%, and 0%) and sonicated for 30 minutes. The solids were separated from the extracts by centrifugation (D-78532 tuttlingen, Universal 32R; Hettich Zentrifugen, Germany) at 20°C temperature, 4000 rpm, and 15 minutes, and the extraction solvent was evaporated using a rotary vacuum evaporator (Buchi, B-490, IL, USA) at 40°C. For further analyses, the sample was placed in a freeze dryer (Crist, Alpha 1-4 LD) to remove the remaining solvent and stored at –80°C.

2.4. α -glucosidase inhibition assay

The inhibitory activity of *P. macrocarpa* extract was measured using Collins et al [18] method with some modifications. To prepare the stock solution, 1 mg of extract was immersed in 1 mL of dimethyl sulfoxide. In addition, 1 mg quercetin was dissolved in 1 mL of dimethyl sulfoxide to make a positive control [19]. To dilute the prepared stock solution, 30 mM phosphate buffer (pH 6.5) was added to get a final concentration of 40 μ g/mL [20]. α -glucosidase enzyme (α -glucosidase type 1 from *Saccharomyces cerevisiae*; Megazyme 130301) was dissolved in 50 mM potassium phosphate buffer (pH 6.5) and used as an enzyme solution. The highest concentration (0.02 U/well) was used and incubated for 5 minutes at room temperature (20°C). Next, 20 mL of 50 mM phosphate buffer (pH 6.5) was added in 6.0 mg of substrate (p-nitrophenyl- α -D-glucopyranose; Sigma-Aldrich, N1377-5G) and allowed to incubate at room temperature for 15 minutes. After incubation, the reaction was ceased by adding 50 μ L of 2M glycine (pH 10). The yellow color was produced for p-nitrophenol (released from the substrate) after addition of glycine. According to the aforesaid procedure, the 30 mM phosphate buffer, fruit extract, quercetin (positive control), α -glucosidase (enzyme), substrate, and stopping reagent (glycine) were put in a 96-well microplate, with a final volume of 250 μ L. The absorbance (Abs) was analyzed at 405 nm using a spectrophotometer (Nanoquant/Tecan, Infinite M 200) [20]. The α -glucosidase inhibition was measured by the following equation (Eq. 1) of Wang et al [21].

$$\text{Percentage inhibition} = (\text{Abs}_{\text{control}} - \text{Abs}_{\text{sample}}) / \text{Abs}_{\text{control}} \times 100 \quad (1)$$

All experiments were conducted in triplicates. Metabolites with the highest inhibitory activity were further characterized by analyzing the concentration needed to inhibit 50% of the α -glucosidase activity (IC_{50}) under the assay conditions, which were ascertained graphically. Different parts of the plant were compared on the basis of their IC_{50} values calculated from the dose response curves.

2.5. Instrumentation and FTIR-ATR measurements

The FTIR spectra of the samples were read at room temperature (20°C) using attenuated total reflectance (ATR) and an internal reflection element made of diamond using Spectrum Two instrument (Perkin Elmer Inc., USA). The spectral region

was 4000–400 cm^{-1} at a resolution of 4 cm^{-1} . The spectra were measured by Spectrum 10.03.09.0139 software (Perkin Elmer Inc., USA). ATR scan technique was used directly for all the samples. The freeze-dried extracts were analyzed, without any preparation, directly on the diamond ATR crystal. The ATR crystal was carefully cleansed between the measurement using solvent and acetone.

2.6. Data preprocessing and calibration model development

Each FTIR spectrum was baseline corrected using Spectrum software (Perkin Elmer, Inc.) to minimize the differences between spectra because of baseline shifts. The spectra were then exported to ASCII file that reduced the FTIR spectra automatically, and excel files were prepared for chemometric analysis. The multivariate calibration (OPLS model) for the data of FTIR spectra was produced using the SIMCA-P software version 13 (Umetrics, Umeå, Sweden). Investigation was based on a 36×3600 data matrix gathered so that each column corresponded to the spectra data and each row represented the sample at a given wavelength. The calibration of functioning of OPLS was evaluated using the coefficient of determination values (R^2Y) and root mean square error of estimation (RMSEE). In addition, Q^2Y and root mean square error of cross-validation (RMSE_{CV}) were applied for the assessment of validation capability of OPLS.

The full frequency region (4000–400 cm^{-1}) of the FTIR spectra was applied at a resolution of 4 cm^{-1} ; consequently, a total of 7189 variables were acquired. The model was constructed and validated by the 36 independent observations. The model validity was evaluated by internal cross validation according to the value of R^2Y and Q^2Y cumulative.

2.7. Statistical analysis

The results were assessed using Minitab 16 (Minitab Inc., State College, PA, USA) by one-way analysis of variance with Tukey's comparison test. A 95% confidence interval was used, and the differences were considered significant at $p < 0.05$.

3. Results

3.1. Extraction yield

Extracts of different solvent concentrations were obtained from fruit flesh of *P. macrocarpa* under investigation. The extraction yields of different solvent ratios of *P. macrocarpa* fruit flesh are illustrated in Table 1. The percentage yield of aqueous and ethanolic extracts of *P. macrocarpa* were significantly ($p > 0.05$) different from one another. The extract yield mainly depends on the solvent properties. Depending on the extraction solvent, the range of the percentage yields of the fruit extracts was 10–25%, as presented by the results. Comparing all extracts in this study, the highest extraction yield (25%) was obtained from the water extracts of *P. macrocarpa*. The percentage yield of the ethanol extract was found to be 10%, which was the lowest yield. This result indicates that the extraction yield depends on the solvent polarity. A more

Table 1 – Comparison of the extraction yield and α -glucosidase inhibitory activity of the *Phaleria macrocarpa* at different concentrations of ethanol.

Extraction solvent (%)	Extraction yield (%)	α -glucosidase inhibitory activity IC ₅₀ (μ g/mL)
Water	25.40 $\pm$ 0.60 ^a	299.24 $\pm$ 29.40 ^a
20% Ethanol	23.56 $\pm$ 1.15 ^b	137.99 $\pm$ 31.89 ^b
40% Ethanol	22.83 $\pm$ 1.09 ^{bc}	65.24 $\pm$ 16.37 ^c
60% Ethanol	21.47 $\pm$ 1.11 ^c	32.82 $\pm$ 5.89 ^d
80% Ethanol	19.28 $\pm$ 1.09 ^d	13.15 $\pm$ 2.96 ^d
100% Ethanol	10.26 $\pm$ 1.08 ^e	7.42 $\pm$ 1.70 ^d
Quercetin	ND	4.34 $\pm$ 1.08 ^d

Values are presented as the mean $\pm$ standard deviation (SD), $n = 6$. Values in each column with different subscript letters are significantly different ($p < 0.05$). ND = not determined.

polar solvent gives a higher yield. The results demonstrated that water was the most suitable extraction solvent according to the extract's yield. The yields of different extracts exhibited the following trend: water > 20% > 40% > 60% > 80 > 100%, which was in agreement with the previous studies [22,23]. Javadi et al [22] reported that the water extract of *Cosmos caudatus* leaves showed maximum yield compared with the various organic solvents. Sultana et al [23] also found that the aqueous methanol extract of medicinal plant leaves was higher than the various absolute organic solvents.

3.2. Inhibition of α -glucosidase activity

The inhibition activities (values of IC₅₀) of *P. macrocarpa* at different ethanolic concentrations are presented in Table 1. A lower IC₅₀ value is desirable for higher inhibitory potential. The IC₅₀ values of six different extracts of *P. macrocarpa* ranged from 7.4 μ g/mL to 299.2 μ g/mL. The inhibition potential of different samples was compared on the basis of their resulting IC₅₀ values. At 100%, 80%, and 60% ethanolic extracts, there were no significant differences ($p > 0.05$) among the IC₅₀ values generated for these extracts. However, the inhibition potential (IC₅₀ value) of water and 20% and 40% ethanolic extracts showed a significant difference ($p < 0.05$). When the ethanol concentration of the extract was increased to 100%, it showed a very high inhibitory activity against α -glucosidase, which was significantly different ($p < 0.05$) when compared with aqueous extract. Absolute ethanol and 80% ethanol exhibited the highest inhibition of α -glucosidase activity with an IC₅₀ of 7.4 μ g/mL and 13.1 μ g/mL, respectively, compared with an IC₅₀ of 4.3 μ g/mL estimated for quercetin (positive control). At high ethanol concentrations, there was a high recovery of the more lipophilic compounds, but a low recovery of the more hydrophilic compounds. The water extract exhibited the lowest activity (299.2 μ g/mL). Ethanol is less polar than water. Therefore, ethanol recovers the more lipophilic compounds, and water recovers the more hydrophilic compounds. In this study, 100% ethanol extract having the highest activity indicates that more lipophilic compounds in *P. macrocarpa* may be responsible for α -glucosidase activity. This finding is in agreement with a previous study, which reported that 100%

ethanol extract of air dried *Phyllanthus niruri* showed the highest inhibitory activity of α -glucosidase compared with water and 50% and 70% of ethanol extract of air dried *Phyllanthus niruri* [24]. Moreover, this is further supported by a recent study which reported that 100% of ethanol extract of *Ipomoea aquatic* showed the highest inhibitory activity of α -glucosidase compared with water and 20%, 50%, and 80% ethanol extract of *Ipomoea aquatic* [25]. This activity showed the following trend: 100% > 80% > 60% > 40% > 20% > water (Table 1). The inhibitory potential of *P. macrocarpa* decreased significantly with increasing the solvent polarity (water ratio), resulting in a lower potential of water solvent than organic solvents for extracting metabolites responsible for inhibitory activity [22,26]. This finding might be also for the concentrations of the extracted biologically active constituents and their variability at different solvent ratios. This result is in agreement with the other studies reporting that absolute organic solvents showed a strong biological activity compared with a mixture solvent (aqueous–organic) [22,23]. Besides that, the α -glucosidase inhibitory activity of *P. macrocarpa* was in accordance with a previous study as potential inhibiting carbohydrate hydrolyzing enzymes in both *in vitro* and *in vivo* effectively [3].

3.3. Analysis of FTIR spectra

FTIR spectroscopy has been widely used and is one of the advanced techniques that could be used for analysis of bioactive compounds of natural products. The importance of IR spectroscopy for the qualitative analysis arises from its properties, especially as fingerprint technique, meaning that there are no two bioactive compounds having the same FTIR spectra. Figure 1 shows typical spectra of six different extracts of *P. macrocarpa*, and the functional group assignment is summarized in Table 2. The whole range of FTIR raw spectra looks very similar by observation because of the similarity in chemical composition. That is, significant relationship within the FTIR spectra and α -glucosidase inhibition was not detected by raw evaluation. The obvious differences between spectra of the two samples, spectra 1 and 6, were due to the intensity at wave numbers of 2852 cm⁻¹, 1734.3 cm⁻¹, 1368.6 cm⁻¹, 1162 cm⁻¹, and 572.72 cm⁻¹. The polarity of pure water and ethanol are 1.000 and 0.654, respectively. The dissimilarities observed between spectra 1 and 6 at 2852 cm⁻¹, 1734.3 cm⁻¹, 1368.6 cm⁻¹, 1162 cm⁻¹, and 572.72 cm⁻¹ are due to the slightly different solvating properties of ethanol and water. The peak at 3292.2 cm⁻¹ is for O–H bonds of alcohols and phenols (H–bonded hydroxyl group) and carboxylic acids. The peaks at 2929.8 cm⁻¹ and 2852 cm⁻¹ are assigned to the sp³ and sp² stretching of C–H bonds of methyl and aldehyde groups [27]. The peak at 1734.3 cm⁻¹ is for C=O stretching due to the presence of aldehyde group. The peak at 1612.4 cm⁻¹ indicates the presence of C=C stretching of alkenes and N–H bending of amines and amides. The peak at 1514.2 cm⁻¹ is assigned to the vibration of N=O in nitro (R–NO₂) and stretching of C=C in aromatics. The presence of N=O vibration in nitro group and C–H bending in alkanes (–CH₃) are indicated by the peak at 1416 cm⁻¹ [28]. The bands at 1368.6 cm⁻¹, 1273.8 cm⁻¹, 1199.2 cm⁻¹, 1162 cm⁻¹, and 1043.5 cm⁻¹ represent S=O

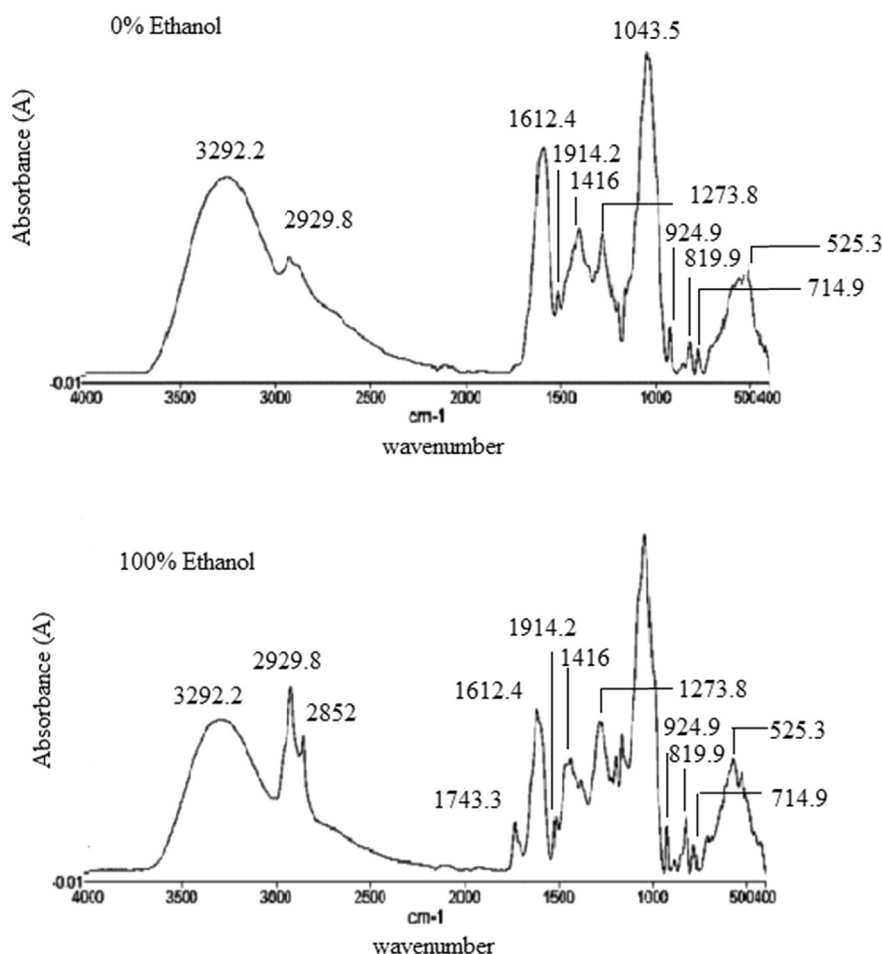


Figure 1 – Typical FTIR spectra of extracts of 100% ethanol extract (active) and 0% ethanol extract (nonactive).

vibration of sulfoxide, C–O stretching of alcohols and carboxylic acids, and C–N vibration of amines [29]. The band at 924.93 cm^{-1} is associated with C–H out of plane bending vibration of mono-substituted aliphatic alkenes, where

819.94 cm^{-1} , 779.3 cm^{-1} , and 714.96 cm^{-1} are for C–H out of plane bending vibrations of meta and para-substituted of aromatics. The absorbance at about 714.96 cm^{-1} is assigned to the C–H bond of long chain alkanes [28].

Table 2 – Functional groups and modes of vibration in ethanolic and aqueous extracts.

Wavenumber (cm^{-1})	Functional group	Vibration mode
3292	O–H	Stretching (sym)
2929 and 2852	–C–H (CH_2)	Asymmetric & symmetric stretching vibration of methylene (– CH_2) group
2852	–C–H (CHO)	Stretching (sym)
1734	C=O (CHO, COOCH_3)	Carbonyl (C=O) functional group from the ester linkage & aldehyde
1612	N–H (NH_2), C=C (alkene)	Bending, stretching
1514	N=O (NO_2), C=C (aromatic)	Stretching
1416	–C–H (CH_3)	Bending
1368	S=O (SO_2), C–X, N=O (NO_2)	Stretching
1273–1043	C–X, S=O (SO_2), C–N (NH_2), C–O (OH, COOH)	Stretching
924	–C–H (alkene)	Bending & out of plane vibration
819	–C–H (aromatic)	bending & out of plane vibration
779–540	C–X (halogen)	Vibration of halogen & out of plane vibration of disubstituted aromatics

3.4. OPLS and statistical validation

The OPLS model was established from 36 observations (samples) and 7189 ($X = 7188$, $Y = 1$) variables (spectra) using one predictive and four orthogonal (1+4+0) components with $R^2Y = 0.98$ and $Q^2Y = 0.70$.

The OPLS, a new analog of partial least square (PLS), is able to rotate the projection so that the model focuses on the effect of interest that is often masked by other unwanted variation, and the first OPLS component shows the difference between the classes. In addition, PLS considers only component 1 and component 2, but OPLS considers orthogonal component also and multiple data set are incorporated by OPLS [30]. This phenomenon was successfully used to obtain new information of a particular combination of metabolite, protein, and transcript correlation [29]. Therefore, OPLS can incorporate a large data set for better identification and interpretation of relevant information.

To select the most beneficial model, some parameters such as regression coefficient (R^2Y), RMSEE, and $RMSE_{CV}$ were evaluated. A value of R^2Y presents the goodness of fit for the components, whereas Q^2Y indicates the predictive ability. R^2Y is the percentage of variation of the response explained by the model; Q^2Y is the percentage of the variation of the response predicted by the model according to cross validation. The model fitness and predictive ability are regarded as effective when the cumulative values of R^2Y and Q^2Y are 0.5 or above [31,32]. RMSEE is a measure of average deviation of the model from the data [33]. For this model, the RMSEE value was 17.14, which is comparable to other predictive model developed by other studies [34]. Additionally, $RMSE_{CV}$ is a measure of the predictive quality of the model for the new samples. The lower the $RMSE_{CV}$ value, the higher the predictive accuracy for new sample. The $RMSE_{CV}$ value was 57.29 for this developed OPLS model.

Table 3 shows the results obtained from the OPLS model in terms of R^2Y , RMSEE, and $RMSE_{CV}$ either for normal or other spectra filters such as its derivatives, multiplicative signal correction and standard normal variant. As presented in Figure 2 and Table 3, OPLS calibration in the second derivatives spectra showed the highest R^2Y (0.98) and the lowest RMSEE (17.17) and $RMSE_{CV}$ (57.29) values compared with other spectra filter.

Table 3 – Multivariate calibration for determining of α -glucosidase inhibitory metabolites using FTIR based on OPLS technique.

Multivariate calibration	Data filter	R^2Y	RMSEE	$RMSE_{CV}$
OPLS	Normal	0.344523	87.6634	100.576
	1 st derivative	0.973757	18.7113	47.2206
	2 nd derivative	0.977128	17.1745	57.2975
	3 rd derivative	0.937399	27.9516	66.2902
	MSC	0.898656	37.421	50.6524
	SNV	0.859918	42.5036	53.5465

MSC = multiplicative signal correction; OPLS = orthogonal partial least square; R^2Y = regression coefficient; $RMSE_{CV}$ = root mean square error of cross-validation; RMSEE = root mean square error of estimation; SNV = standard normal variant.

The first two OPLS components showed the maximum variation with a value of 35.5% and 36.9% for components 1 and 2, respectively. Figure 2 illustrates the OPLS score scatter plot produced by multivariate data analysis. The processed data obtained from this plot indicated that the *P. macrocarpa* extracts were well separated based on activity. A gradient shifting of the sample along the OPLS component 1 to the left side represents an increase in the activity. Based on the activity, the active extracts (100%, 80%, 60%, and 40% ethanol) were significantly separated by predictive component t_1 [1] from the non-active extracts (20% ethanol and water). By contrast, the most active extracts (100% ethanol) were at the left of the ellipse separated significantly by the orthogonal component t_0 [1] from their less active samples (water extracts) which were at the right side.

Figure 3 illustrates the coefficient of determination (R^2Y) value of the regression line between the predicted and observed IC_{50} value based on FTIR spectra. The value of R^2Y in this research was 0.98, which exhibited the very good fitness and predictive ability of the regression model.

3.5. Bioactivity of spectral characteristic and representing functional groups

Figure 4 illustrates the line loading plot obtained from multivariate data analysis of the extracts of *P. macrocarpa*. This plot can make the recognition of the possible biological activity of peaks from the FTIR spectrum easier. This plot assesses the spectral characteristics that accelerate or decelerate the α -glucosidase inhibition. The inhibitory activity with a positive pq [1] value has been presented in the rightmost point of the loading plot. The wave number with increased pq [1] values accelerate the inhibitory activity. The reported α -glucosidase inhibitory activity containing compounds (Figure 5) of *P. macrocarpa* are phalerin [1], carbohydrate (structure is suspected analogous with myglitol or acarbose) [2], and magniferin [3]. The peaks at 2912 cm^{-1} , 2848 cm^{-1} , 1720 cm^{-1} , 1179 cm^{-1} , and 827 cm^{-1} have the highest pq [1] value and 3367 cm^{-1} , 1454 cm^{-1} , 1291 cm^{-1} , $1083\text{--}998\text{ cm}^{-1}$, $756\text{--}549\text{ cm}^{-1}$ also have a positive pq [1] value, representing the signal from the α -glucosidase inhibitory compound. The peaks at 2912 cm^{-1} and 2848 cm^{-1} are assigned to the stretching of C–H of methyl group in phalerin and aldehyde groups, whereas 1720 cm^{-1} is for the vibration of C=O in ketone (magniferin and phalerin), aldehyde, and carboxylic acid group. The peak at 1179 cm^{-1} may be assigned to the stretching of C–O of alcoholic group present in phalerin, magniferin, myglitol, and acarbose and carboxylic acid, whereas 827 cm^{-1} was for C–H out of plane bending in alkenes and aromatics. The absorption at 3315 cm^{-1} also leads to the inhibition potential because of the –OH (hydrogen bonded) of alcoholic and phenolic group present in all reported compounds and –NH of amines group present in acarbose, 1291 cm^{-1} is for the stretching of C–N of amines group observed in myglitol, whereas the absorbance at $756\text{--}549\text{ cm}^{-1}$ is for C–Cl bond which was observed in acarbose. The reported compounds such as carbohydrate, phalerin, and magniferin in *P. macrocarpa* have –CH, –CO, –N=, –NH, and –OH stretching. However, aldehyde and carboxylic acid group containing compound has not yet been

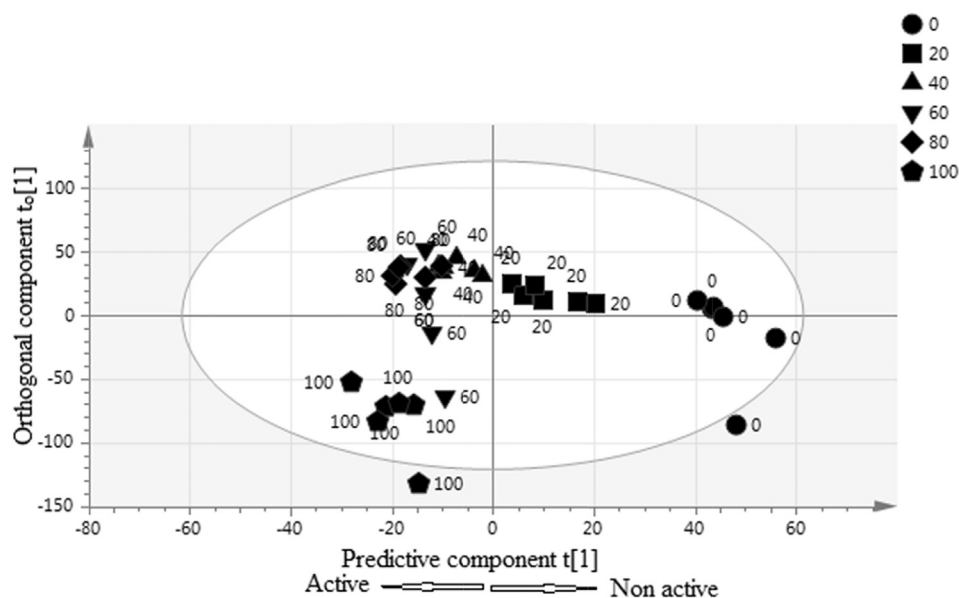


Figure 2 – Orthogonal partial least squares score scatter plot of different ratio of *Phaleria macrocarpa* extracts.

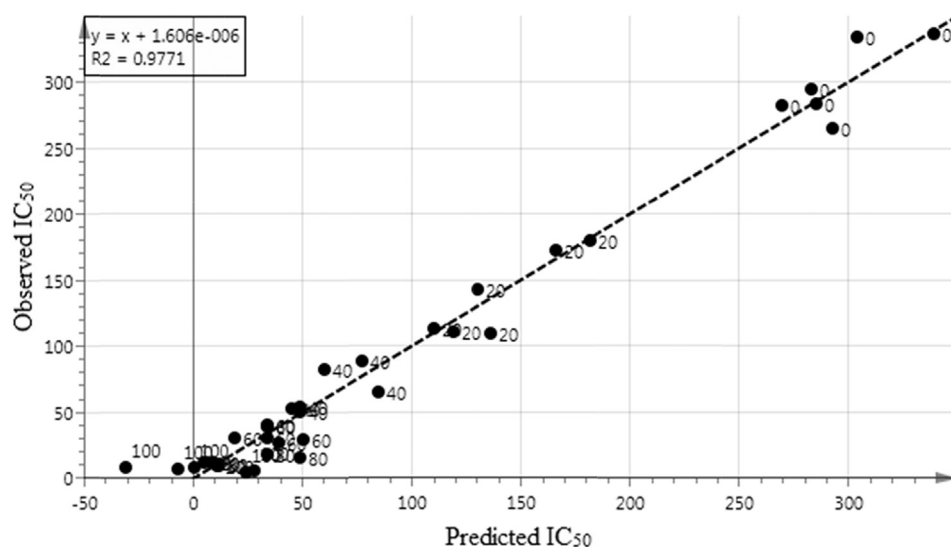


Figure 3 – Prediction versus observation IC_{50} values from all samples. The R^2 of the regression line indicates the goodness of fit between experimental observations and predicted model.

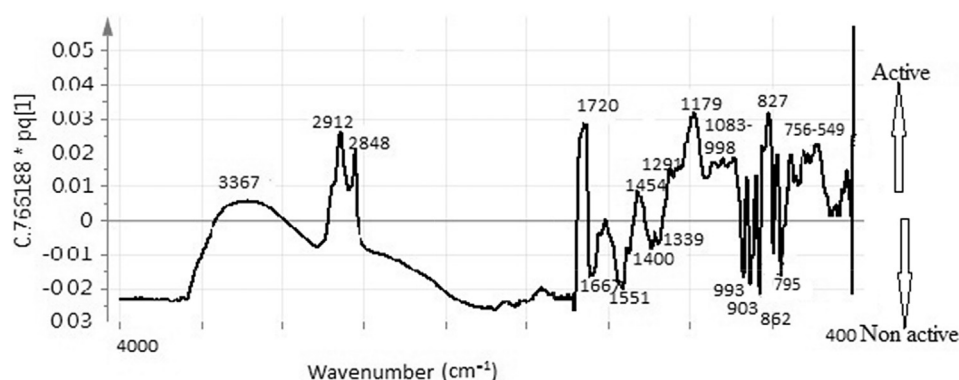


Figure 4 – Orthogonal partial least squares line loading plots of different ratio of *Phaleria macrocarpa* extracts.

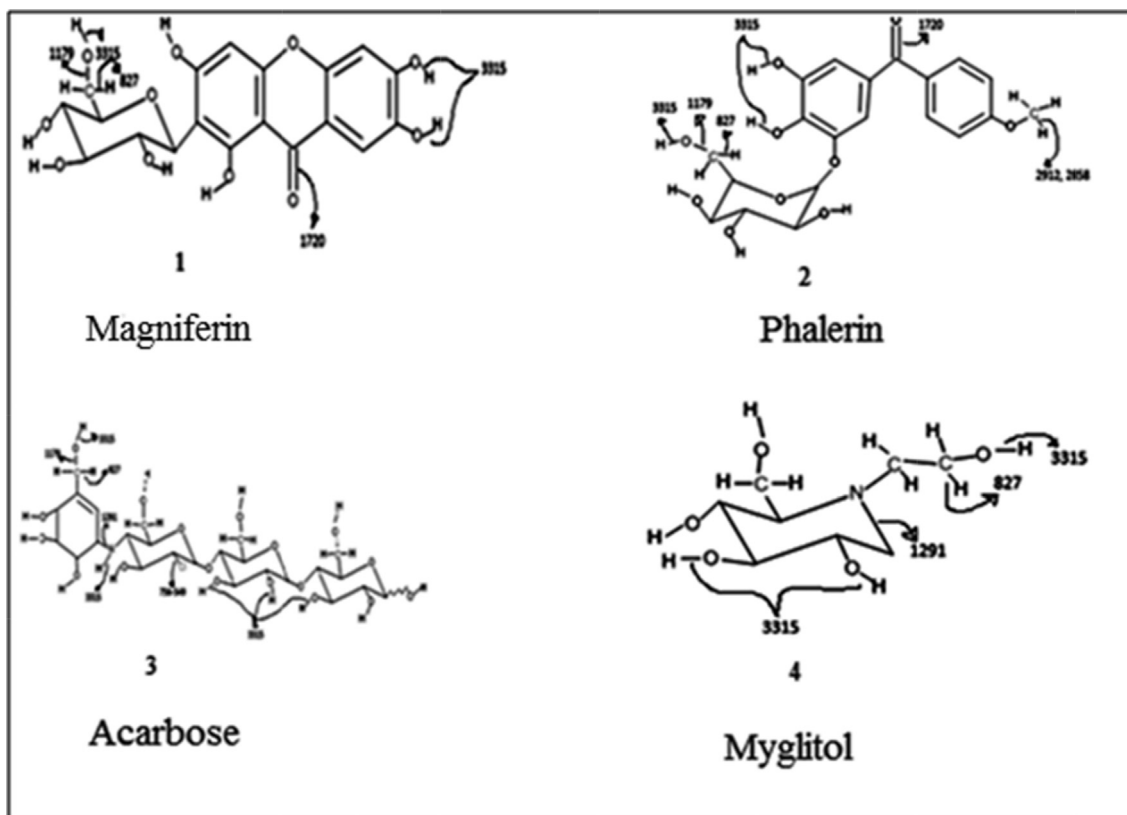


Figure 5 – Reported α -glucosidase inhibitory compounds from *Phaleria macrocarpa*.

reported in *P. macrocarpa*. However, in this study, it has been seen that the aldehyde and carboxylic acid groups also induce the α -glucosidase inhibitory activity. The findings are in agreement with the previous study reported by Javadi et al [22], indicating that carboxylic acid group (α -linolenic acid) might induce inhibitory activity. The contribution of aldehyde group to inhibitory activity has not yet been reported. This group may accelerate the α -glucosidase inhibitory activity. Flavonoids can inhibit the activity of α -glucosidase, the unsaturated C ring, 3–OH, 4–CO, the linkage of the B ring at the position 3, and the hydroxyl substitution of the B ring [35]. Meanwhile, the polyphenol shows its α -glucosidase inhibitory activity by the presence of free hydroxyl group at the 3'-position of theaflavin as well as the esterification of theaflavin with a mono-galloyl group. Thus, the existence of polyphenols, flavonoids, carbohydrates, sterols, amines, and terpenes might contribute to the α -glucosidase inhibition potential of the *P. macrocarpa* extract. Nevertheless, other peaks obtained from this OPLS line loading plot do not have the positive contributions to the α -glucosidase inhibition, and this plot can also provide the negative contributing peaks. For instance, the spectra at 1667 cm^{-1} , 1551 cm^{-1} , 1400 cm^{-1} , 1339 cm^{-1} , 933 cm^{-1} , 903 cm^{-1} , 862 cm^{-1} , 795 cm^{-1} reduced the α -glucosidase inhibitory activity. The band at 1667 cm^{-1} is for the stretching of C=N of imines and oximes groups, whereas the peaks at 1551 cm^{-1} and 1400 cm^{-1} are assigned to the asymmetric and symmetric stretch of N=O in $-\text{NO}_2$,

indicating that nitro, imine, and oxime groups containing compounds might contribute in the reduction of the inhibitory activity. The outcome of this study disagrees with the report by Damazio et al [36]. These results suggest that the effect of chalcones on serum glucose lowering seems to be a consequence of insulin secretion, and these chalcones represent potential compounds with strong α -glucosidase inhibitory properties. However, it has been also reported that the presence of nitro group and their position in the phenyl rings are responsible for the α -glucosidase inhibitory activity of chalcones. Therefore, the position of nitro group may be responsible for reducing the activity in *P. macrocarpa*. The peaks at 1339 cm^{-1} and 933 cm^{-1} are assigned to the $-\text{SO}$, $-\text{PH}$, or $-\text{PO}$ functional groups containing compounds that may also reduce the inhibitory activity. Another phase in this research is the insertion of the entire frequency region of the FTIR spectrum in model development instead of using a part of spectrum, as reported previously [27]. The advantage of incorporating the full wave length region with that a portion of spectrum might lead to over-fitting the model. By contrast, the selection of a part of spectrum is arbitrary that might lead to ambiguity in the interpretation. Therefore, this research demonstrates the success of containing the whole FTIR spectral bands to produce a predictive multivariate calibration model (OPLS) for α -glucosidase inhibition of the extract of *P. macrocarpa* as a rapid alternative model for the investigation of biological activity.

4. Conclusion

A successful multivariate model was developed using FTIR-ATR together with OPLS algorithm as a simple and rapid method for the analysis of α -glucosidase inhibitory activity of *P. macrocarpa* extracts. Further, the model has been used to identify the contribution to the bioactivity of the functional groups. This study reported that the functional groups, such as $-\text{CHO}$, $-\text{COOH}$, $-\text{NO}_2$, $-\text{NH}$, and $-\text{OH}$, induced the inhibitory activity, whereas $-\text{SH}$, $-\text{PH}$, and $-\text{PO}$ reduced the bioactivity. The constructed model having highly significant correlation with R^2Y of 0.98 could be applied for quality control process in pharmaceutical industry.

Conflicts of interest

There is no conflicts among the authors and institutes.

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