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The Effect of the Aqueous Enzymatic Extraction Method towards *Momordica charantia* Seed Oil and Its Lignocellulosic Biomass

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Abstract

The development of a sustainable product using natural resources will gain much attention nowadays. This work aimed to study the effect of aqueous enzymatic extraction (AEE) using a different ratio of enzyme cocktails HEL1 and X7 towards the production of *Momordica charantia L. (M. charantia*) seed oil and to identify the composition of the seeds' lignocellulosic biomass (extracted-free materials, EFM). The *M. charantia* seed oil contained omega-3 fatty acids, such as stearidonic acid, that possess hypotriglyceridemic properties. The oil obtained was derivatized to fatty acid methyl esters (FAMEs) before biochemical quantification using gas chromatography (GC). The EFM was subjected to Soxhlet extraction and further analyzed to identify lignin, acid-soluble sugars, and hemicellulose composition. The results showed that the highest seed oil percentage, $(6.26\pm0.53 \%)$ was extracted from M2 sample using a combination of 5 % HEL1 to 1.25 % X7 enzyme cocktails, as compared to the extraction without enzyme ($3.32\pm0.38 \%$). The percentage of stearidonic acid in the seed oil increased from 13.55 % (without enzyme) to 19.43 % (M2), proving that there were some change in terms of fatty acid composition in the seed oil.

Keywords: Momordica charantia L., Seeds, Aqueous enzymatic extraction, Lignocellulosic biomass

Introduction

Momordica charantia L. (M. charantia), from the Cucurbitaceae family, is a thin climbing vine with long-stalked leaves that usually flowers from July to August and is cultivated as a vegetable in Asia, Africa, and the Caribbean. *M. charantia* plant can adapt to a wide range of climates, but grows best in warm weather. **Figure 1** shows a cross-section of *M. charantia* fruits. It is also known as bitter melon, balsam pear, bitter squash, or bitter gourd.



Figure 1 Cross-section of *M. charantia* fruit.

The local Malay name for this plant is "peria katak". The bitter taste of the fruit is attributed to the presence of cucurbitacin-like alkaloids, momordicin, and triterpene glycoside [1,2]. Bitter melon is also used in traditional medicine to treat a variety of ailments. The indigenous people of the Amazon drink the tea from the leaves to expel intestinal gas and to promote menstruation, as well as to reduce blood sugar levels for diabetic patients [3]. The use of bitter melon fruit as a natural remedy for treating diabetes is a common practice among the local population of Asia, Africa, and the Caribbean [4]. Many studies have reported that the hypoglycemic properties of *M. charantia* are induced by the presence of steroidal glycosides [5-7], insulin mimetic lectins [8], and triterpenoids [9,10]. However, most of the studies focused on the leaves, pulps, and stems of *M. charantia*. The seeds are discarded without any further uses. Some studies have shown that the seeds of *M. charantia* contain a high content of omega-3 polyunsaturated fatty acids, such as α -linolenic acid, eicosapentaenoic acid, stearidonic acid, and α -eleostearic acid [11-13]. Dietary intake of omega-3 polyunsaturated fatty acids is known to lower triglyceride content in humans, as the polyunsaturated fatty acids can attach to and clear out the unhealthy fats [14]. According to Chen *et al.*, the α -eleostearic acid presence in bitter melon seed oil could reduce hepatic triglyceride accumulation [15].

In Malaysia, approximately 122.80 hectares of farmland is planted with *M. charantia*, which can produce a total of 1372.46 metric tons of *M. charantia* fruits every year. In accordance, about 548.984 metric tons of *M. charantia* seed, that can be used to produce seed oil, are being discarded every year in Malaysia [16]. This is a waste of natural resources, as the extraction of oil from the seed can easily be carried out using many methods. With the increasing demand for oilseed from agricultural plants, oil extraction technology has become more advanced, such as the alternative solvent extraction methods including supercritical fluids [17], enzyme-assisted aqueous extraction (AEE) [18], and microwave-assisted extraction [19].

AEE is getting some attention nowadays due to its efficient, sustainable, and eco-friendly characteristics. This extraction process depends on the specific properties of the enzyme itself. The enzyme, acting as a catalyst under optimum experimental conditions, disrupts the plant cell wall by hydrolyzation to release the intracellular components [20]. The plant cell walls are a complex structure, mainly made of cellulose, hemicellulose, lignin, and protein, in addition to pectin [21,22]. Rosenthal reported that the cell wall degradation enzyme for rapeseed consists of a combination of pectinase, cellulase, and hemicellulase [23]. In a previous study, Latif and Anwar reported that the active mechanism of enzyme cocktail (Alcalase 2.4L, Viscozyme L, Protex 7L, Natuzyme, and Kemzyme)

during AEE was the degradation of cellulose, pectin, and protein. Thus, the destruction of structural cell wall integrity led to higher sesame oil recovery; 16.5 - 24.8 % for AEE was observed, compared to only 12.3 % for the control (without enzyme treatment) [24].

The seed residue or lignocellulosic biomass obtained from the AEE method is known to have different hemicellulose compositions, as the enzymatic hydrolysis is influenced by both the structural features of cellulose and the mode of enzyme action. Given the advantages of this extraction technology, this research study aimed to highlight the essential characteristics of the *M. charantia* seed oil using a specific AEE method, and the seed biomass compositions, such as lignin, soluble sugar, uronic acid, and hemicellulose, after the AEE method.

Materials and methods

Seed sample preparation

Unripe *M. charantia* fruits were purchased from a vegetable farm in Ara Kuda, Seberang Perai Utara, Penang, Malaysia, between June and August 2015. The seeds were separated, cleaned, and sundried. The seed samples were stored in polyethylene bags under vacuum for further processing.

Aqueous enzyme extraction of seed samples

AEE was performed to extract oil from the *M. charantia* seed according to the method as described by Muniglia *et al.* [25]. The enzyme cocktails, HEL1 and X7, were blended in the Laboratoire D'ingénierie des Biomolécules (Nancy, France). The enzyme cocktail HEL1 contained the average mixture of cellulase (45.0 μ mol/min/mL), hemicellulase (7.0 μ mol/min/mL), betaglucosidase (4.5 μ mol/min/mL), endocellulase (27.0 μ mol/min/mL), exocellulase (9.0 μ mol/min/mL), arabinase (465.0 μ mol/min/mL), xylanase (300.0 μ mol/min/mL), galactanase (750.0 μ mol/min/mL), endopolygalacturonase (61.0 μ mol/min/mL), and pectin methylesteras (1.0 μ mol/min/mL) which aimed for cell wall degradation, while the enzyme cocktail X7 contained protease and pectinase, which aimed for protein degradation.

Initially, the freshly-ground dried *M. charantia* seeds were mixed with distilled water at a ratio of 1:4 into a pre-programmed bioreactor (Celligen Plus, New Brunswick Scientific, U.S.A.) with continuous stirring. Then, the enzyme cocktails were added consecutively into the bioreactor once the temperature of the bioreactor reached 50 °C, starting with HEL1, followed by X7, at their different ratios (**Table 1**). The reaction was left to carry on for 4 h in the bioreactor, followed by centrifugation (0.158 kilohertz, 20 min, 20 °C) to separate the products. The final products of this extraction process were: 1) oil, 2) seed residue, and 3) an aqueous part.

Seed samples	Enzyme (%)/Seed weight (g)				
	HEL1	X7			
M0	-	-			
M1	5.00	-			
M2	5.00	1.25			
M3	5.00	2.50			
M4	5.00	5.00			

 Table 1 Seed samples and their enzyme cocktail ratios.

The seed oil samples were initially derivatized to fatty acid metal esters (FAMEs) using the esterification process as described by Metcalfe and Schmitz before being further analyzed to identify their biochemical compositions using gas chromatography (GC). The polar properties of underivatized fatty acid usually lead to the formation of hydrogen bonding, which results in adsorption issue. Hence, reducing their polarity by derivatization makes them more amenable for analysis [26]. Simultaneously, the seed residue was further analyzed to identify the lignin, soluble sugar, uronic acid, and hemicellulose compositions using the Soxhlet extraction method, ionic chromatography, and size exclusion chromatography (SEC). The aqueous part was discarded.

Gas chromatography analysis of FAMEs

Each derivatized oil (FAME) was analyzed to identify its biochemical composition using flame ionization detector gas chromatography (GC-FID), as it is the most suitable detector to detect organic compounds or any easily-flammable substances with higher carbon concentrations. The detector responds to the presence of ionic substances, and the number of ions formed can be calculated. The peak area of GC chromatogram can then be used to determine the qualitative measurement of the eluate, as the peak area of the signal is proportional to the concentration of the compound. In this study, the analysis of FAMEs was carried out by using Shimadzu GC-2010 plus an instrument equipped with Zebron capillary GC column ZB-Wax, 60 m \times 0.25 mm \times 0.25 mm. Pure helium gas was used as the carrier gas, pressurized and flowed at 141.2 kPA and 11.7 mL/min, respectively. Then, 1 μ L of each FAME sample was injected into the GC and was separated using temperature programming, starting with 120 °C and held for 2 min, followed by heating to 180 °C with a heating rate of 40 °C/min and held for another 2 min, before the temperature was increased to 220 °C with a heating rate of 3 °C/min, and held for another 25 min. The fatty acid standards were used for calibration (7 points per curve). The reagents and standards were provided by Sigma-Aldrich and Fisher Scientific.

Soxhlet extraction of seed residue

The dried seed residue was further extracted using the Soxhlet extraction method to ensure all extractive components were removed before hydrolysis to determine their lignin and acid-soluble sugar contents. The presence of extractive components hinders the hydrolysis process by cross-reacting with acid and condensing to acid-insoluble components, resulting in inaccurate analysis. To execute the Soxhlet extraction, an amount of 50 g of each dried seed residue sample was loaded into an extraction thimble, which was placed into a Soxhlet extractor (Pyrex, U.S.A) fitted with a condenser (Pyrex, U.S.A) on an isomantle heater (Fisherbrand, U.S.A). Then, 250 mL of *n*-hexane was added into a 500 mL round bottom flask fixed to the Soxhlet extractor and heated at *n*-hexane boiling point (68 °C). The condensate dripped into the reservoir containing the thimble. The solvent from the reservoir was poured into a round bottom flask once the solvent level reached the siphon. The process was continued for 5 h. Then, *n*-hexane was evaporated using a rotary evaporator (Buchi, Switzerland). The final products of the Soxhlet extractive materials (EFM) and 2) extractive materials. The extractive materials were discarded, and the EFM was weighed and kept for further analysis.

Determinations of lignin, soluble sugars, and uronic acid of EFM

An amount of 175 mg of each dried EFM sample was placed in a conical bottom centrifuge tube, and 1.5 mL of 72 % sulfuric acid was added dropwise. The mixture was placed into a water bath and heated at 30 °C for 1 h with periodical stirring using a glass rod. A volume of 42 mL of distilled water was added into the mixture to achieve a concentration of 4 % sulfuric acid by mass. The mixture was then autoclaved at 121 °C for 1 h. The mixture was allowed to cool down to room temperature before being filtered. The filtrate was then diluted with 56.5 mL of distilled water in a volumetric flask. The resulting solution was further analyzed by high-performance anion-exchange chromatography with pulsed amperometry detection (HPAEC-PAD) (PerkinElmer, U.S.A) to determine its soluble sugar and uronic acid contents. The solid residue (lignin) was dried and weighed. **Table 2** shows the seed sample codes for soluble sugar and uronic acid analysis.

Sample codes	Enzyme (%)/Sample Weight (g)					
1	HEL1	X7				
$M0_L$	-	-				
$M1_L$	5	-				
$M2_L$	5	1.25				
M3 _L	5	2.5				
$M4_L$	5	5				

Table 2 Seed sample codes for acid soluble sugar analysis and their descriptions.

Note: M refers to the EFM, which had undergone AEE, while, $_{L}$ refers to the soluble sugar and uronic acid sample of the EFM.

Determination of soluble sugar and uronic acid contents of EFM

The separation and quantification of soluble sugars and uronic acids of the EFM were carried out using HPAEC-PAD operating on Dionex ICS-3000 system, which consisted of a gradient pump SP, a passer AS, an electrochemical detector ED operating with a gold electrode, a reference electrode Ag/AgCl, and Chromeleon version 6.8 (Dionex Corp, USA). A Carbopac PA20 column (3×150 mm², Dionex) with a guard column (3×30 mm², Dionex) was used in the stationary phase. The mobile phase was made of water, 250 mM NaOH in water, and 1 M NaOAc/20 mM NaOH in water. A diluted 46 - 48 % sodium hydroxide solution (Fisher Scientific) and anhydrous sodium acetate (Sigma Aldrich) in ultrapure water were used as eluents. The eluents were degassed with helium gas for 20 min before being injected. After each analysis, the column was flushed for 10 min with 250 mM NaOH solution and restored for 10 min to the initial conditions of the analysis. The samples were injected with an entire loop of 25 µm, and the separations were carried out at 35 °C at a flow rate of 0.4 mL/min. The pulse sequence for pulsed amperometric detection consisted of potentials of +100 mV (0 - 200 ms), +100 mV integration (200 - 400 ms), +100 mV (400 - 410 ms), -2000 mV (410 - 420 ms), +600mV (430ms), and -100mV (440 - 500ms) [27]. Soluble sugar and uronic acid standards were used for calibration (7 points per curve). The reagents and standards were provided by Sigma-Aldrich and Fisher Scientific.

Hemicellulose extraction of EFM

Initially, the EFM was subjected to the bleaching process to produce holocellulose. Sodium chlorite was used as the bleaching agent due to its moderately-active oxidation properties. The bleaching process was conducted by milling 1.5 g of each dried treated EFM sample into 0.05 mm size before being dispersed slowly into 125 mL deionized water in a 250 mL conical flask. Acetic acid glacial, 1 mL, was added into the mixture, with an additional 1 g of sodium chlorite. The mixture was sealed and placed in a Julabo SW 22 shaking water bath at 70 °C for 2 h. Then, another 1 mL of acetic acid glacial and 1 g of sodium chlorite were added consecutively into the reaction mixture with continuous shaking in the same conditions for 2 h. The process was repeated until the color of the residue turned to white. The reaction mixture was filtered and washed thoroughly with deionized water. The holocellulose obtained was collected and dried overnight in a fume hood, which was then further reprecipitated into hemicellulose. An amount of 200 mL of 2 % w/v sodium hydroxide aqueous solution was added into 10 g of each holocellulose sample in a round bottom flask and stirred for 2 h at 80 °C. Then, an additional 200 mL of 2 % w/v sodium hydroxide aqueous solution was added into the reaction mixture and stirred for another 2 h. The pH of the mixture was adjusted to 5 - 6 by adding 20 % w/v acetic acid solution. The reprecipitation process of the mixture was repeated using 800 mL of ethanol. The precipitate formed was recovered by centrifugation. The percentage yield of hemicellulose was calculated and analyzed using Fourier-transform infrared (FTIR) spectroscopy (PerkinElmer, England), as well as SEC.

Determination of molecular weight distribution of hemicellulose by SEC

The molecular weight distribution of hemicellulose was determined by SEC. The analysis was performed using a Hewlett-Packard 1090 series HPLC system consisting of an autosampler, a UV detector, and 3 columns of Styragel HR1, HR3, and HR4 (Waters Inc.) linked in series using tetrahydrofuran (THF) as the eluent. The hemicellulose was dissolved in 1 mg/mL of THF, and the solution was filtered through a 0.45 mm filter. Then, 20 mL of the filtered solution was injected into the system and was detected using a UV detector at 280 nm. Data were collected using Agilent ChemStation rev. A.10.01 and analyzed using Agilent GPC Addon rev. A.02.02 software.

Results and discussion

Two different types of enzyme cocktails, namely HEL1 and X7, were used in this study to extract oil, soluble sugars, and uronic acid of *M. charantia* seeds. **Table 3** shows that the introduction of enzyme cocktails into the batch reaction increased the percentage of extracted seed oil. In the control experiment, without the addition of enzyme cocktail (M0), only 3.32 ± 0.38 % of seed oil was extracted. The highest percentage seed oil (6.26 ± 0.53 %) was extracted from M2 using a combination of 5 % HEL1 to 1.25 % X7 enzyme cocktails, followed by M4 (5.03 ± 0.28 %) using a combination of an equal ratio of HEL1 and X7 enzyme cocktails.

The results indicated that cellulase, proteinase, and pectinase present in the enzyme cocktails HEL1 and X7 could facilitate the breakdown of cellular and subcellular structures of the seed to promote oil release. The enzymes could have weakened the vegetal cells in the seed to help release the oil [28]. In a previous study, Latif and Anwar [29] reported that the active mechanism of a selected enzyme cocktail during enzymatic aqueous extraction was the degradation of cellulose, pectin, and protein, thus destructing the structural cell wall integrity of the seed cells [29]. Additionally, the incorporation of enzymes may have cleaved the protein networks of the cotyledon cells and oleosin-based membranes that surround the lipid bodies of the seed cell wall, resulting in oil release [30,31]. The results also showed that the increment of enzyme cocktail X7 to HEL1 ratio was not proportional to the percentage of oil being extracted.

Sample	Oil extracted (%)				
M0	3.32±0.38				
M1	4.67±0.11				
M2	6.26±0.53				
M3	4.10 ± 0.44				
M4	5.03 ± 0.28				

Table 3 Percentages of seed oil recovery by aqueous enzymatic extraction method using combination of different enzyme cocktail ratios.

Note: The extraction process was triplicated.

The percentage of seed oil extracted = $\frac{Weight of oil obtained (mg)}{Weight of dry seed used (mg)} \times 100 \%$

The extracted seed oil samples (M0, M1, M2, M3, and M4) were subjected to derivatization into FAMEs before identification of their biochemical compositions by GC-FID since FAMEs are more volatile and more easily analyzed. GC-FID has been widely used in the separation and detection of

FAMEs. The GC chromatogram revealed the presence of some common and non-essential FAMEs, as shown in **Table 4**. Seven known fatty acids were identified. Three unknown fatty acids were also detected. However, 3 unknown peaks of free fatty acids were observed from the chromatogram that were co-eluting almost at the same time. It is suggested that some compounds may co-elute more significantly in size and elute together at the end during separation [32,33]. Hence, 1 dimensional GC may not be the best option for FAME separation.

Table 4 FAME com	positions of M.	charantia seed	oils of M0.	M1, M2, M3, and M4.
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Fatty and	Compositions (%)							
Fatty acid	M0	M1	M2	M3	M4			
Palmitic acid (C16)	8.84±0.12	7.21±0.09	17.97±0.11	17.26 ± 0.10	11.23 ± 0.09			
Stearic acid (C18)	-	-	3.28 ± 0.13	-	-			
Elaidic acid (C18:1n9t)	0.47 ± 0.11	8.06 ± 0.08	20.42 ± 0.09	28.06 ± 0.11	29.63 ± 0.08			
Oleic acid (C18:1n9c)	35.80 ± 0.08	17.31 ± 0.11	-	-	-			
Vaccenic acid (C18:1n7)	$2.84{\pm}0.10$	5.02 ± 0.09	2.52 ± 0.09	5.02 ± 0.11	7.58 ± 0.10			
Stearidonic acid (C18:4n3)	13.55 ± 0.11	13.42 ± 0.11	19.43 ± 0.08	16.42 ± 0.08	12.83 ± 0.09			
Paullinic acid (C20.1)	6.50±0.12	6.19±0.12	13.51±0.11	5.19 ± 0.09	2.41 ± 0.10			
Unknown 1	10.66 ± 0.09	10.14 ± 0.09	7.81 ± 0.10	15.14 ± 0.09	15.31 ± 0.10			
Unknown 2	9.55±0.11	9.96 ± 0.10	$6.00{\pm}0.10$	4.96 ± 0.10	8.69±0.11			
Unknown 3	11.79 ± 0.12	8.83±0.11	$9.07 {\pm} 0.09$	7.98 ± 0.11	12.32 ± 0.11			

Note: Values are % of total fatty acid expressed as mean±SD of 3 separate determinations.

Based on the compositions of *M. charantia* seed oil (**Table 4**), the percentage of elaidic and palmitic acids increased with the introduction of the enzyme cocktails. In the control experiment (M0), only 0.47 % of elaidic acid was present in the extracted oil. With the introduction of the enzyme cocktails, the percentage of elaidic acid increased to 8.06, 20.42, 28.06, and 29.63 % for M1, M2, M3, and M4, respectively. The presence of the enzyme cocktails showed a similar trend for palmitic acid, albeit with lesser increment. The oil release could be due to the presence of cellulase in the enzyme cocktail, which disrupted other polysaccharide constituents in the seed cell wall. The cellulase presence in HEL1 and X7 promoted an efficient cellulose degradation. The cellulase mechanism, described as the cleaving of internal glucosidic bonds within an unbroken glucan chain by endoglucanase enzyme, could result in the creation of a non-reducing chain of cellobiohydrolase substrate, cleaving the cellobiose dimers from the glucan chain that was released in the solution. Furthermore, β -glucosidase split the cellobiose dimer into glucose monomer, concluding the cellulose hydrolysis process [34]. The combination of enzymes played a dominant effect in disrupting the seed cell walls, which was also reported in other findings [35].

These results also showed that the oil composition differed with different X7 enzyme ratios. In M4, the percentage distribution of elaidic acid, vaccenic acid, palmitic acid, stearidonic acid, and paullinic acid increased with increasing ratio of X7. The results suggested that the enzyme cocktail X7 supported the reaction of enzyme cocktail HEL1 hydrolyzed the seed cellular structures as each enzyme cocktail consisted of different types of an enzyme mixture. It can be concluded that the *M. charantia* seed oil obtained from AEE exhibited slight changes in the integrity and percentage distribution of the active constituents. This finding was in agreement with the findings of earlier researchers, who showed that enzyme assisted extraction can lead to an increase in oil yield as well as slight modification in the biochemical properties of the extracted oil [36].

The biochemical composition of extracted oil also differed from the previous studies. Ali *et al.* extracted seed oil with the compositions of palmatic acid (4.84 ± 0.13 %), stearic acid (20.21 ± 0.40 %), linoleic acid (6.37 ± 0.46 %), and eleostearic acid (53.22 ± 0.36 %) [37]. Meanwhile, the seed oil being extracted using AEE had a higher amount of palmitic acid. In addition, the saturated fatty acid, stearic

acid, was only detected in M2. Additionally, the omega-3 fatty acid detected by Ali *et al.* was eleostearic acid, whilst in the oil extracted using AEE, the highly unsaturated stearidonic acid was detected.

The lignin, soluble sugar, and uronic acid contents of the EFM were determined. Lignin consists of a very complex structure which is difficult to cleave [38]. The lignin content of the EFM; M0, M1 (15.91 %) > M2 (15.06 %) > M4 (14.71 %) > M3 (14.54 %) did not show any significant changes with the presence of enzymes in AEE. Different types of soluble sugars and uronic acid were released and further analyzed with HPAEC-PAD. The results are presented in Table 5. The main soluble sugar composition of EFM was glucose (rt. 15.90 min) and rhamnose (rt. 10.42 min), respectively. No definite trend was shown between the percentage of enzyme ratios used with the percentage distribution of soluble sugars and uronic acid obtained. M0L showed the lowest amount of acid galacturonic (rt. 31.87 min, 8.42 %). The percentage of acid galacturonic increased with the presence of enzyme cocktails in $M_{3_{L}}$ (16.12 %) < $M_{4_{L}}$ $(16.37 \%) < M1_{L} (22.00 \%) < M2_{L} (22.25 \%)$. These results suggested that the presence of pectinase in the enzyme cocktail X7 facilitated the breakdown of cell wall pectic polysaccharides into simpler molecules like acid galacturonic [39-41]. The concentration of glucose in M0 was only 57 %. The percentage of glucose increased with the introduction of enzyme cocktails; M3 (65.42 %) \leq M4 (66.77 %) < M1 (71.34 %) < M2 (76.20 %). These results indicated that more cellulose from the cell wall was cleaved into different types of sugars (glucose, xylose, galactose, arabinose, rhamnose, mannose, and fucose) with the help of the enzyme cocktails [42-44].

Sam	HEL1	N7	Percentage of yield (%)								
Sam	ΠELI	X7	Glu	Xyl	Galac	Ara	Rham	Man	Fuc	Ac. Galac	Ac. Gluc
M0	_		57.00	19.02	4.21	3.95	1.70	1.09	0.42	11.77	0.83
$M0_L$		-	0.61	1.84	7.60	6.98	62.98	6.41	2.40	8.42	2.77
M1	5		71.34	13.85	2.57	2.28	1.07	0.91	0.25	7.31	0.42
$M1_L$	3	-	1.29	2.67	9.17	10.37	35.36	12.64	5.24	22.00	1.26
M2	5	1.25	76.20	10.36	2.60	1.98	1.06	0.79	0.17	6.54	0.30
M2 _L			1.10	2.32	9.16	9.40	37.26	12.62	4.54	22.25	1.34
M3	5	2.5	65.42	22.34	2.72	3.37	1.41	0.78	0.18	2.94	0.83
$M3_L$	5	2.5	0.57	1.19	8.92	7.13	48.00	13.16	3.94	16.12	0.96
M4	5	5	66.77	22.06	2.48	2.77	1.21	1.00	0.17	2.69	0.85
$M4_L$			0.61	1.66	7.96	5.95	48.34	14.40	2.93	16.37	1.79

Table 5 Sugar analysis of EFM by HPAEC-PAD.

Note: M refers to the EFM, which had undergone AEE. $_{\rm L}$ refers to the soluble sugar and uronic acid sample of the EFM.

The hemicellulose content of the seed biomass was determined. M0 produced the highest percentage of hemicellulose (16.98 %). The percentage of hemicellulose decreased slightly with the presence of enzyme cocktails; M1 (14.23 %), M2 (13.44 %), M3 (13.41 %), and M4 (12.98 %). Despite significant knowledge in the synergism among cellulase and enzyme cocktail components, a full mechanistic understanding of lignocellulose hydrolysis by cellulolytic enzymes is still limited and needs further refinement [45,46]. The variation in the hemicellulose composition showed that, as a

polysaccharide of simpler structure than cellulose, hemicellulose was also prone to breakdown with the presence of hemicellulase [47].

The results of further analysis of hemicellulase obtained using FTIR spectroscopy are shown in **Figure 2**. The FTIR spectrum showed that there were differences in the peak intensity and between the absorption bands obtained. According to the Beer-Lambert law, the higher is the concentration; the stronger is the peak intensity [48]. The absorption of 3272, 2927, 1628, 1416, 1239, and 1038 cm⁻¹ peaks represented the characteristic absorption of hemicellulose [49]. The broad absorption band at ~3272 cm⁻¹ usually represents the O-H or the absorption of a large number of individual molecules, such that hydrogen is bonded to a slightly different compound.

Meanwhile, the absorption band ~2927 cm⁻¹ indicated the presence of saturated carbon band, C-H. The band ~1737 cm⁻¹ referred to the stretching band of a carbonyl group. The absorption that appeared in ~1628 cm⁻¹ is usually associated with the absorbed water [50]. A small peak at ~1737 cm⁻¹ could be attributed to the acetyl groups of the hemicellulose. A strong band that appeared at ~1038 cm⁻¹ could be assigned to C-OH bending in xylan, attributed to the dominant presence of xylan, one of the 3 main sub-groups of hemicellulose with the backbone of 1 - 4 linked β -D-pyranose attributed to the presence of xylan-type hemicellulose in *M. charantia* seed [51].

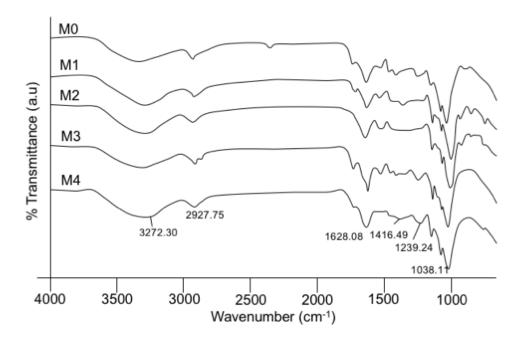


Figure 2 FTIR spectra of hemicellulose.

The extracted hemicellulose was then subjected to SEC analysis to evaluate the effects of the enzymes on their molecular mass distribution. Figure 3 shows the size exclusion chromatogram of the extracted samples.

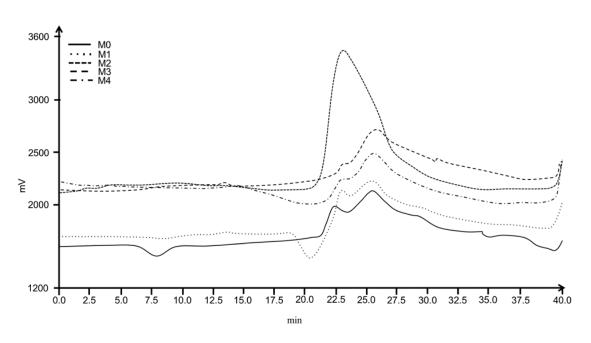


Figure 3 Size exclusion chromatogram of hemicellulose from different *Momordica charantia* seed extracted free materials.

Each extracted sample was observed to have a slightly different molecular mass distribution trend. The difference in the molecular mass distribution trend could be due to the effects of different enzymes towards the induction of different transformations in cellulose size. M0 had the highest weight average (M_w) of 447,148 - 457,148 gmol⁻¹, followed by M4 $(M_w: 36,186 - 378,116 \text{ gmol}^{-1})$, M3 $(M_w: 120,339 \text{ gmol}^{-1})$, M1 $(M_w: 87,865 \text{ gmol}^{-1})$, and lastly M2 $(M_w: 55,915 \text{ gmol}^{-1})$. These results indicated that the treatment degraded the macromolecular structure of hemicellulose to a noticeable extent with the incorporation of enzyme [47] as the hemicellulose obtained displayed higher molecular mass than that conventionally extracted, of ~38,000 and 56,000 gmol⁻¹ in neutral and basic medium, respectively [52]. Sun *et al.* suggested that the extraction of high M_w hemicelluloses was obtained with the coextraction of lignin-carbohydrates complexes [53]. In addition, the high hemicellulose polydispersity (M_w/M_n) from M0 M_w/M_n of 1.15 - 7.17, to M2 M_w/M_n of 6.06, to M4 M_w/M_n of 1.9 - 21.72, to M1 M_w/M_n of 107.71, and lastly to M3 M_w/M_n of 284.30 indicated a wide molecular weight distribution, which might be caused by hemicellulose chain aggregation [54,55].

Conclusions

The adaptation of the AEE method successfully aided the extraction of *M. charantia* seed oil utilizing the combination of HEL1 and X7 enzyme cocktails. The presence of enzymes disrupted the cell wall into simpler molecules (i.e., glucose, xylose, etc.) and facilitated the oil released. The introduction of the enzyme in the extraction process slightly changed the molecular mass distribution of the lignocellulosic biomass. Improved analytical methods are needed to characterize further, analyze, and study the biomass composition and enzyme mechanisms. This study proved that *M. charantia* seed contain significant contents of fatty acids, sugars, and hemicellulose, which can be further utilized to develop sustainable products such as biofuel, membrane, and biodegrade plastic.

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12 of 13

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