Mapping metabolome changes in *Luffa aegyptiaca* Mill fruits at different maturation stages via MS-based metabolomics and chemometrics

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**G A R A P H I C A L A B S T R A C T**

**A B S T R A C T**

**Introduction:** *Luffa aegyptiaca* Mill, sponge gourd or Egyptian cucumber, is grown worldwide for its edible fruit consumed as a vegetable like cucumber. Unlike young fruit (YF), the fully mature ripened fruit (MF) is strongly fibrous and is used as a cleanser to make scrubbing bath sponges. YF undergoes a complex series of physiological and biochemical changes during fruit ripening. However, the chemical compositional differences between YF and MF in *Luffa aegyptiaca* have not been distinguished to date.

**Objectives:** Comprehensively compare the metabolites profile of YF and MF to give insight on how maturation stage affects chemical composition.

**Methods:** Mass-based metabolomics comprising GC/MS and UHPLC/MS were adopted in this study targeting its volatile and non-volatile metabolites coupled with chemometrics to rationalize for the differences.

**Results:** A total of 53 volatile metabolites were identified via headspace solid phase microextraction (SPME) comprising 66.2% aldehydes/furans, 51.6% alcohols, 38.2% ketones, 15.1% acids and 10.1% aromatics of which aldehydes/furans were dominant at both fruit stages. Young fruit was in general more enriched in metabolites as revealed from UHPLC/MS and GC/MS analyses. The YF group encompassed higher levels of short chain alcohols (1-octen-3-ol) and aldehydes ((E)-2-hexenal and cucumber...
Introduction

*Luffa aegyptiaca* (Mill), a synonym of *Luffa cylindrica* (Linn) M. Roem, is a unique vegetable member of the Cucurbitaceae family, native to Asia and widely cultivated in several tropical and subtropical regions worldwide for its economical, medicinal and nutritional uses [1]. It is commonly named sponge gourd, vegetable sponge, bath sponge, dish cloth gourd and loofa [2,3].

The green young fruit (YF) is edible as raw like cucumber or after being cooked as in squash. A myriad of metabolites are reported in its YF: i.e., phenolic acids, flavonoids, vitamins, carotenoids, saponins and triterpenes which add to its health effects [4]. In contrast, the abrasive nature of the fibrous sponge gourd mature fruit (MF) allows for its use in skin exfoliation, body cleansing and or stimulation of blood circulation [5]. Traditionally, in Japan, the “Hechimasui” or the water extract of the plant’s vascular bundle, is used as skin lotion owing to its saponins content aside from amino acids and minerals which help maintain the skin healthy [6]. Industrially, *L. aegyptiaca* is used for water purification acting as adsorbent for heavy metals from waste water [3]. With regards to its health benefit effects, *L. aegyptiaca* fruits exhibit diverse traditional biological activities such as antimicrobial, stomachic, purgative, emollient, tonic, galactagogue and antipyretic, also useful treatment of syphilis, tumours, lung complaints, splenopathy and leprosy along with nephritis and jaundice [5,7]. While externally it is used for rheumatism, backache, as well as in hemorrhoids [5]. Chemical content mediating for luffa fruit uses include flavonoids, saponins and steroidal compounds [8]. Many phenolic and flavonoid glycosides were separated using antioxidation-guided assay suggesting that consumption of sponge gourds can mitigate against oxidative stress in human body [9]. Polyphenolic metabolites in luffa may also contribute to its anti-inflammatory action by inhibiting LPS-induced NO generation [9,10]. Another two compounds named, 3-hydroxy-1-methylene-tetrahydroxy-naphthlene-2-carbaldehyde and dihydroxy spinasterol, were isolated from *L. aegyptiaca* fruit petroleum ether extract and found to exhibit moderate antimicrobial action [11].

Seeds encompassed within *L. Aegyptiaca* fruit are also recognized with several effects. For example, luffin-a and luffin-b are two proteins isolated from *L. aegyptiaca* seeds to exhibit ribosome-inactivating cytotoxic action along with an abortifacient effect [12]. Luffacilin is another peptide isolated from sponge gourd seeds with anti-fungal activity against *Mycosphaerella arachidicola* and *Fusarium oxysporum* [13]. Two triterpenoids isolated from *L. aegyptiaca* seeds i.e., oleanolic acid and echninocystic acid exhibited immune-stimulatory effect [14]. An investigation by Muthumani, Meera [15] assessed *L. aegyptiaca* seeds anti-inflammatory, bronchodilator and antimicrobial activities against *S. aureus* and *Candida albicans*. In addition, the polypeptide “luffin P1” isolated from seeds of *L. aegyptiaca* exhibited anti-HIV-1 activity [16].

*Luffa* peel was also found to be enriched in phyttonutrients at higher levels relative to its pulp with more potent anti-inflammatory effect similar to citrus peel [17]. Considering that fruit undergoes major changes in its composition upon ripening to affect its effects and usage warrants for monitoring such changes using analytical tools. Indeed, no detailed study has uncovered compositional difference in metabolites of *L. aegyptiaca* fruits represented by young and mature stages.

Analytical techniques that aid in profiling a complex mixture of metabolites in biological systems are increasingly applied in food research and to employ chromatographic separation using either gas chromatography (GC) or liquid chromatography (LC) coupled to mass spectrometry (MS) targeting its volatile and non volatile metabolites, respectively.

In this study, headspace solid phase micro-extraction (SPME) was used to collect volatile constituents of *Luffa aegyptiaca* prior to its analysis using GC–MS. SPME is a fast, non-solvent technique that allows volatile sampling emitted from the plants at low levels and is superior to hydro-distillation in being sensitive due to thermal heating without the generation of artifacts [18–20].

For analysis of non-volatile high molecular weight secondary metabolites, ultra high performance liquid chromatography (UHPLC) coupled to ESI-mass detector is a well developed technology that provides monitoring of a large number of metabolite peaks at high sensitivity [21,22]. Considering the complexity of metabolites dataset and to assess for differences in *L. aegyptiaca* fruits at two maturation stages in an untargeted manner, multivariate data analyses were employed. Principal component analysis (PCA) as an unsupervised clustering way without dataset knowledge alongside supervised OPLS were employed for analysing GC–MS and UHPLC-MS datasets [23,24].

To the best of our knowledge, detailed metabolites profile of *L. aegyptiaca* and impact of maturity stage upon its chemical composition have never been addressed before, issues of value to rationalize for its different uses and or biological effects.

Materials & methods

Plant material

Luffa aegyptiaca fruits were collected fresh from the field at Behira Governorate, Egypt at the two different ripening stages, young (YF) and old mature (MF) during the month of May 2016. The description regarding old versus mature is based on the fruit pulp (interior part) in case of young is a soft vegetable part whereas in case of old mature the pulp is almost completely fibrous. Voucher specimens are deposited at the Department of Pharmacognosy, College of Pharmacy, Cairo University.

Chemicals and fibers

SPME fiber of stableflex coated with divinylbenzene/carboxen/polydimethylsiloxane (DVB/CAR/PDMS, 50/30 μm) was purchased by Supelco (Oakville, ON, Canada). All chemicals and standards were purchased from Sigma Aldrich (St. Louis, MO, USA). Acetonitrile and formic acid (LC–MS grade) were obtained from J. T. Baker (Netherlands), milliQ water was used for UHPLC/PDA/ESI-Orbitrap HRMS analysis.
**Headspace volatiles analysis of L. aegyptiaca**

The HS-SPME volatile analysis was carried out as stated in [20] with slight modifications. Dried finely ground fruits (100 mg) were placed in SPME screw cap vials (1.5 ml) and spiked with (Z)-3-hexenyl acetate dissolved in water at a final concentration of 2 μg per vial. The SPME fiber was inserted manually into vial containing samples placed in an oven kept at 50 °C for 30 min. The fiber was subsequently withdrawn into the needle and then injected into the injection port of the gas chromatography-mass spectrometer (GC–MS). GC–MS analysis was performed on a Schimadzu GC-17A gas chromatograph equipped with DB-5 column (30 m × 0.25 mm i.d. × 0.25 μm film thickness; Supelco) and coupled to Schimadzu QP5050A mass spectrometer. The interface and the injector temperatures were both set at 220 °C. The following gradient temperature program was used for volatiles analysis. The oven temperature was kept first at 40 °C for 3 min, then increased to 180 °C at a rate of 12 °C min⁻¹, kept at 180 °C for 5 min, and finally ramped at a rate of 40 °C min⁻¹ to 240 °C and kept at this temperature for 5 min. The carrier gas Helium was used at a total flow rate of 0.9 ml/min. Splitless injection mode was used for volatiles analysis. In samples with lower levels of volatiles, the SPME fiber was inserted to the next analysis by placing it in the injection port for 2 min at 220 °C to ensure complete elution of volatiles. Blank runs were made during sample analyses. The HP quadruple mass spectrometer was operated in El mode at 70 eV. A scan range was set at 40–500. Volatile components were identified by comparing their retention indices (RI) relative to n-alkanes (C6-C20), mass matching to NIST, WILEY library data and with standards whenever available. Peaks were first deconvoluted using AMDIS software (www.amdis.net) prior to mass spectral matching. GCMS files shall be made available upon request.

**GC–MS data processing for multivariate analysis**

Volatile metabolites abundance data were prepared for multivariate data extraction by using MET-IDEA software [19] for data extraction. The mass peak abundance was first normalized to the amount of spiked (Z)-3-hexenyl acetate then subjected to principal component analysis (PCA) and partial least squares-discriminant analysis (OPLS-DA) using SIMCA-P version 13.0 software package (Umetrics, Umeå, Sweden). All variables were mean centered and scaled to Pareto variance.

**Metabolites extraction for UHPLC- Orbitrap HRMS analysis**

Extraction of luffa fruits for UHPLC/MS was made by homogenizing freeze dried powder (120 mg) with 5 ml 100% MeOH containing umbelliferone (an internal standard for relative quantification using UPLC-MS) present at a concentration of 10 μg/mL using a Turrax mixer (11,000 RPM). Extracts were then vortexed vigorously and centrifuged at 3000g for 30 min to remove plant debris. For UHPLC-MS analyses, 500 μl was aliquoted and filtered through 20μm size filter. Three microliters were used for UPLC-MS analysis. For each specimen, three biological replicates from each fruit stage were provided and extracted in parallel under identical conditions.

**UHPLC-Orbitrap HRMS analysis**

The negative ion high-resolution ESI and collision-induced dissociation (CID) MSn spectra were obtained from an Orbitrap Elite mass spectrometer (Thermo Fisher Scientific, Germany) equipped with a heated electrospray ion source (negative spray voltage of 3 kV, capillary temperature of 300 °C, source heater temperature of 250 °C, FTMS resolution of 30,000). Nitrogen was used as sheath and auxiliary gas. The MS system was coupled to an UHPLC system ( Dionex UltiMate 3000, Thermo Fisher Scientific), equipped with a RP-18 column (particle size 1.8 μm, pore size 100 Å, 150×1 mm ID,Acquity HSS T3, Waters; column temperature of 40 °C) and a photodiode array detector (220–600 nm, Thermo Fisher Scientific). The mobile phases were H2O and CH3CN with 0.1% formic acid by using the following binary gradient at a flow rate of 150 μl/min: 0–1 min, isocratic 95% A, 5% B; 1–11 min, linear from 5 to 100% B; 11–19 min, isocratic 100% B; and 19–30 min, isocratic 5% B. The injection volume was 2 μL. The CID mass spectra (buffer gas: helium) were recorded using normalized collision energy (NCE) of 35%. Metabolites were characterized by their UV–VIS spectra (220–600 nm), retention times relative to external standards, mass spectra and comparison to phytochemical dictionary of natural products database (CRC) and reference literature. UHPLC-MS files are uploaded in study number MTBLS1305 in Metabolights https://www.ebi.ac.uk/metabolights/database. Relative Luffa metabolites analyzed after UHPLC/MS was performed using XCMS data analysis software, which can be downloaded freely from https://bioconductor.org/packages/release/bioc/html/xcms.html [22,25]. Data were subjected to PCA and OPLS-DA, using the SIMCA-P 13.0 software package (Umetrics, Umeå, Sweden). Markers were subsequently identified by analyzing the S-plot, which was declared with covariance (p) and correlation (pcor). All variables were mean centered and scaled to Pareto variance.

**Results & discussion**

*L. aegyptiaca* volatiles profiling using SPME coupled to GC–MS.

In this study, the metabolic profiles of *L. aegyptiaca* fruits were characterized at two different maturity stages represented by young and old mature fruits. Three independent biological replicates from different fruits representing each stage were extracted and analyzed under identical conditions. The volatile analysis of *L. aegyptiaca* (Fig. 1) revealed for the presence of 53 volatiles listed in Table 1, categorized into acids, alcohols, aldehydes/furan, aromatics, esters, ketones, aliphatic hydrocarbons and monosesqui-terpene hydrocarbons. Both fruit ripening stages showed relatively a comparable aroma composition as shown in their gas chromatograms (Fig. 2). This is the first report on aroma compositional differences of *L. aegyptiaca* fruits at two different stages of growth worldwide.

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**Fig. 1.** A photo of *L. aegyptiaca* fruit collected at the 2 different ripening stages (YF) and (MF).
Table 1
Relative percentile levels of volatile components detected in young and old mature L. egyptiaca fruit using SPME-GC–MS measurements (n = 3).

| Peak | Rt (min) | KI Identification | Class | Mature average ± Std. Dev. | Young average ± Std. Dev. |
|------|----------|-------------------|-------|---------------------------|---------------------------|
| 1    | 8.51     | 951 Pentanoic acid | Fatty acid | 2.67 ± 2.01 | 0.46 ± 0.37 |
| 2    | 10.01    | 1071 Heptanoic acid |       | 0.85 ± 0.55 | 0.24 ± 0.20 |
| 3    | 11.42    | 1180 Octanoic acid |       | 3.93 ± 2.30 | 1.46 ± 0.77 |
| 4    | 12.71    | 1270 Nonanoic acid |       | 3.86 ± 2.02 | 1.60 ± 0.84 |
| 5    | 8.49     | 949 1-Octen-3-ol Alcohol |       | 6.85 ± 9.55 | 15.94 ± 1.01 |
| 6    | 8.78     | 971 3-Octanolo |       | 0.74 ± 0.73 | 2.15 ± 0.16 |
| 7    | 9.29     | 1010 6,6-Dimethyl-1,3-heptadien-5-ol |       | 3.45 ± 2.77 | 0.91 ± 0.52 |
| 8    | 9.43     | 1045 Benzyl alcohol |       | 0.49 ± 0.19 | 0.27 ± 0.09 |
| 9    | 9.98     | 1080 Dihydromyrcenol |       | 2.23 ± 1.30 | 0.10 ± 0.05 |
| 10   | 10.39    | 1107 β-Linalool |       | 1.62 ± 0.09 | 9.86 ± 3.38 |
| 11   | 10.67    | 1127 Phenylethyl Alcohol |       | 0.34 ± 0.12 | 0.29 ± 0.21 |
| 12   | 11.19    | 1164 β-Terpineol |       | 0.54 ± 0.30 | 0.40 ± 0.12 |
| 13   | 11.57    | 1191 Camphol |       | 0.52 ± 0.25 | 1.88 ± 0.61 |
| 14   | 11.85    | 1211 α-Terpineol |       | 1.99 ± 1.45 | 0.16 ± 0.08 |
| 15   | 12.23    | 1317 2,2-Dodecadiol |       | 0.81 ± 0.41 | 0.13 ± 0.06 |
| 16   | 6.19     | 773 2-Hexenal Aldehyde/furan |       | 0.17 ± 0.50 | 10.58 ± 1.26 |
| 17   | 6.56     | 879 3-Furanmethanol |       | 0.19 ± 0.32 | 1.69 ± 0.19 |
| 18   | 7.08     | 906 Heptanal |       | 0.68 ± 0.41 | 0.29 ± 0.07 |
| 19   | 8.22     | 929 Benzaldehyde |       | 0.06 ± 0.40 | 4.09 ± 1.38 |
| 20   | 8.63     | 960 2-n-Pentylfuran |       | 0.02 ± 0.01 | 0.01 ± 0.01 |
| 21   | 8.86     | 1009 Octanal |       | 0.50 ± 0.21 | 0.36 ± 0.08 |
| 22   | 9.57     | 1054 Benzenecetaldehyde |       | 0.83 ± 0.67 | 2.57 ± 0.59 |
| 23   | 9.65     | 1038 Cumarine aldehyde |       | 0.60 ± 0.34 | 0.17 ± 0.18 |
| 24   | 9.78     | 1047 2-Octenal, (E)- |       | 0.85 ± 0.21 | 1.06 ± 0.44 |
| 25   | 10.45    | 1111 Nonanal |       | 8.76 ± 1.84 | 1.67 ± 0.38 |
| 26   | 11.18    | 1163 Nonadienal(cucumberaldehyde) |       | 1.03 ± 0.99 | 5.39 ± 2.83 |
| 27   | 11.28    | 1170 2-Nonenal, (E)- |       | 1.97 ± 0.30 | 2.37 ± 0.66 |
| 28   | 11.90    | 1214 Decanal |       | 8.98 ± 3.08 | 0.77 ± 0.05 |
| 29   | 12.16    | 1234 p-Phthalaldehyde |       | 0.27 ± 0.24 | 0.03 ± 0.01 |
| 30   | 6.49     | 878 p-Xylene |       | 1.93 ± 0.88 | 0.76 ± 0.54 |
| 31   | 8.73     | 967 Mesitylene (Benzene, 1,3,5-trimethyl-) |       | 2.22 ± 1.07 | 0.70 ± 0.25 |
| 32   | 9.25     | 1033 m-Cymene |       | 0.26 ± 0.07 | 0.38 ± 0.06 |
| 33   | 10.09    | 1087 Dimethylethylbenzene |       | 0.61 ± 0.30 | 0.13 ± 0.03 |
| 34   | 10.50    | 1115 p Cymene |       | 0.14 ± 0.09 | 0 ± 0 |
| 35   | 11.21    | 1165 Unknown aromatic |       | 0.47 ± 0.24 | 0.07 ± 0.04 |
| 36   | 12.17    | 1320 β-Methylnaphthalene |       | 1.66 ± 0.52 | 0.18 ± 0.25 |
| 37   | 15.71    | 1511 β-Vinylnaphthalene |       | 0.54 ± 0.27 | 0.06 ± 0.10 |
| 38   | 13.08    | 1304 Bornyl acetate |       | 0.16 ± 0.04 | 0.59 ± 0.22 |
| 39   | 14.09    | 1376 3-Hydroxy-2,4,4-trimethylpentyl 2-methylpropanoate |       | 1.90 ± 1.54 | 0.22 ± 0.05 |
| 40   | 14.50    | 1420 Cyclopentanecarboxylic acid, isopropyl ester |       | 0.09 ± 0.04 | 0.01 ± 0.02 |
| 41   | 8.57     | 955 3-Octanone |       | 4.74 ± 3.73 | 26.95 ± 2.02 |
| 42   | 9.20     | 1003 Acetophenone |       | 0.74 ± 0.32 | 0.21 ± 0.06 |
| 43   | 11.78    | 1205 2,4-Dimethyl-3-hexanone |       | 5.02 ± 0.66 | 0.50 ± 0.56 |
| 44   | 10.36    | 1104 n-Udecan |       | 0.33 ± 0.19 | 0.18 ± 0.07 |
| 45   | 13.09    | 1305 Tridecan |       | 1.65 ± 1.11 | 0.22 ± 0.06 |
| 46   | 14.32    | 1393 Tetradecan |       | 4.49 ± 2.72 | 0.60 ± 0.27 |
| 47   | 15.54    | 1505 Pentadecan |       | 3.17 ± 2.10 | 0.40 ± 0.25 |
| 48   | 17.01    | 1601 Hexadecan |       | 2.96 ± 2.09 | 0.42 ± 0.41 |
| 49   | 9.33     | 1014 Limonene |       | 0.40 ± 0.15 | 0.28 ± 0.07 |
| 50   | 10.77    | 1134 Isoyocorene |       | 0.19 ± 0.15 | 0.04 ± 0.03 |
| 51   | 14.74    | 1440 Unknown |       | 0.06 ± 0.04 | 0 ± 0 |
| 52   | 11.07    | 1155 Allyl dithioacetate |       | 0.20 ± 0.08 | 0.06 ± 0.05 |
| 53   | 13.87    | 1369 Eugenol |       | 0.03 ± 0.04 | 0.13 ± 0.04 |

**Total Esters** | 2.16 | 0.83 |

**Total ketones** | 10.50 | 27.66 |

**Total Aliphatic hydrocarbons** | 12.60 | 1.83 |

**Total Terpene hydrocarbons** | 0.66 | 0.32 |

**Total Miscellaneous** | 0.23 | 0.18 |
Aldehydes

Aldehydes/furans amounted for the major class present in mature fruit at ca 32–35%. Decanal (28) & nonanal (25) were the higher volatiles signal intensity percentile detected in MF (8.9% & 8.7%), respectively, and to reach (0.7 & 1.6%) in YF, respectively. Both aldehydes were reported to exhibit antimicrobial/antifungal activity and to possess a citrus but waxy/fatty odor\[26,27\]. In contrast, hexenal (16), with an odor of freshly cut grass\[28\] was the major aldehyde component in YF represents 10.6%. Other aldehydes included nonadienal (cucumberaldehyde) that predominated in young luffa aroma at 5.4% versus 1% in mature fruit\[29\]. Cucumberaldehyde was reported for it potential antimicrobial activities\[30\] and is used as commercial food flavoring agent, as well as in cosmetics and perfumes fragrance\[31\]. Benzaldehyde (19) with its characteristic almond like odour\[32\], represents 4.1% in YF rather than MF. Previous volatile analysis in the closely related vegetable ‘‘cucumber’’ suggested for aliphatic aldehydes i.e., hexenal, nonenal and nonadienal (cucumber aldehyde), to account for its characteristic pleasant aroma and to change upon ripening as observed in luffa\[33\]. Generation of these compounds is believed to be via enzymatic activity after mechanical rupture or cutting of the fruit ‘‘as in chewing process’’ to yield the characteristic flavour of fresh young cucumber fruit i.e., nonadienal\[29\]. Extending such aroma profile to also encompass L. aegyptiaca fruit being detected a higher levels at the edible stage with notable decrease upon ripening and to contribute majorly for young fruits aroma.

Alcohols

Next to aldehydes/furans class, alcohols presented the second major volatile class and also showing same pattern being higher in young fruits at 32.1% versus 19.6% at the mature stage. Octenol (5), β-linalool (10) and camphol (13) were the major components in YF present at (15.9, 9.9 & 1.9%) versus (6.8, 1.6 & 0.5%) in MF, respectively. Octenol is one of the major volatiles detected in YF volatiles (15.9%) characterized by mushroom like yet sweet aroma and flavor\[34\]. Whereas, linalool abundance in young fruit (10%) might contribute for the fruit antimicrobial effect\[35,36\], in addition to its potent fungitoxic action against Fusicladium effusum\[37\]. Other volatiles identified in L. aegyptiaca volatile blend of potential antimicrobial activity included cymene, napthalene, benzyl alcohol, phenylethyl alcohol and eugenol\[38,39\] and likely to contribute for its MF use as bath cleanser. In terms of aroma perception, β-linalool contributes strongly to orange oil aroma\[40\] and might also contribute to luffa overall aroma.

Ketones

The highest % ketone peak intensity in YF was for 3-octanone (41), detected at 27% and dropping to 5% upon fruit maturity. 3-octanone alongside octenol are involved in attracting insects for pollination due to its fruity sweet aroma\[34\]. In contrast, 2,4-dimethyl-3-hexanone (43) was detected in MF at 5% while almost diminished in the YF to 0.5%.

Aromatics

Aromatics were present at 7.8 and 2.3% in MF vs. YF, respectively with xylene (30), mesitylene (31) and methylnapthalene (36) as major components. Napthalene exhibits a strong unpleasant odour and insecticidal action\[41\]. Interestingly, napthalene showed higher % peak intensity in mature fibrous inedible stage of fruit rather than its edible fruit.

Terpenoids

Terpenoids accounted for minor levels as volatile class as both stages with limonene as the predominant monoterpenic present in both fruit stages ca 0.35%.

Fatty acids/hydrocarbons

Two major classes found extensively in MF (11.3 & 12.6%) and to decline in YF (3.7 & 1.8%) were short fatty acids and hydrocarbons, respectively and contrary to most other volatile classes showed higher peak intensity level in YF and suggesting that lipophilic VOCs i.e. fatty acids/hydrocarbons contribute more to MF aroma.
Multivariate data analysis of volatiles dataset in the different fruit ripening stages

Although difference in volatile patterns could be observed, we attempted to employ multivariate data analyses to help identify markers for each fruit ripening stage via principle component analysis (PCA) and supervised orthogonal partial least squares (OPLS). PCA score plot was prescribed by two orthogonal PCs with PC1 accounting for 79% of the total variance versus PC2 to account for 12% (Fig. 3A). Most notably, replicates representing volatiles analysis of each of the two ripening stages showed segregation of specimens to some extent based on ripening stage with YF exhibiting negative score values along PC1 versus positioning of mature fruits on the other side.

PCA loading plot examination (Fig. 3B) revealed that oxygenated volatiles i.e., aldehydes/ketones were the most variant volatiles and contributing for specimens segregation. 2-Hexenal (16), cucumber aldehyde (26), octen-3-ol (5), 3-octanone (41) and the terpenoid alcohol β-linalool (10) in YF and to contribute for its characteristic aroma compared to the abundance of nonanal (25), decanal (28), dimethylhexanone (43) and β-methylnaphthalene (36) in mature fruits (Fig. 3B). Such segregation pattern suggest that qualitative difference in aldehydes and ketones that accounted for the different fruits stages aroma composition, structures of the major key volatiles are illustrated in Suppl. Fig. S1.

Further, supervised multivariate data analysis, i.e., OPLS, was employed for samples classification. OPLS-DA outscores PCA in the separation of the predictive variation from orthogonal variation and enhances data interpretation [42]. The OPLS-DA score plot (Fig. 3C) showed better discrimination between the sample groups with the young group clustering separately from the mature one with variance coverage (R2 = 0.80) and prediction power of (Q2 = 0.73).

The S-loading plot derived from the OPLS-DA model (Fig. 3D) revealed that linalool and cucumber aldehyde could be identified as the most discriminatory aroma compounds for YF. To assess for validity of the GC–MS based OPLS models, Q2, R2X and R2Y values were close to. Permutation diagnostic analysis of 20 iterations provided reference distribution of R2/Q2 values and hence indicated statistical significance of these parameters, with most models showing a regression line crossing zero, which signifies the models validation. Also, the p-value for OPLS-DA model was calculated using CV-ANOVA was estimated at 0.03 below p-value of 0.05 (Suppl. Fig. S2).

Secondary metabolites profiling of L. aegyptiaca via UHPLC/PDA/Orbitrap HRMS

Different studies investigated the chemical composition of L. aegyptiaca organs [5,43–46]. However, this study provides the first comprehensive secondary metabolites profile in luffa fruits at the different ripening stages via high resolution UHPLC/PDA/Orbitrap HRMS (Fig. 4) and coupled to multivariate data analyses (Fig. 5).

Chemical constituents of young and mature fruits were analysed using gradient elution mobile phase (H2O: ACN) buffered with formic acid allowing for peaks elution. 40 Metabolites were detected belonging to 17 saponins, 9 fatty acids, 6 flavonoids and other miscellaneous compounds (Table 2, Fig. 6). Identification was based on molecular formulae, fragmentation patterns, UV absorbance, and comparison with compounds reported in the literature, mass bank and phytochemical dictionary of natural product database. The nature of the attached sugars in glycosides i.e. saponins and flavonoids was revealed from the lost fragment moieties that is /C0 162 amu; C6H10O5 for hexose (galactose or glucose), −146 amu; C6H10O4 for deoxyhexose (rhamnose), −132 amu; C6H9O3 for pentose and −176 amu; C6H8O6 for hexauronic acid [47,48].
Saponins

As a rich source of saponins, 17 compounds originating from 5 triterpene sapogenins were identified (Fig. 6). Saponins were previously reported in luffa [43] and revealed to exhibit biological actions as immunostimulatory, anti-inflammatory and fibrinolysis [5, 7, 14]. Saponins of the triterpene oleanene type identified in the current study included trihydroxy-12-oleanen-28-oic acid (hydroxyhederagenin/arjunolic acid) at \( m/z \) 487; C\(_{30}\)H\(_{47}\)O\(_5\) in peaks (11 and 27), dihydroxy-12-oleanen-28-oic acid (maslinic/machaerinic acid) at \( m/z \) 471; C\(_{30}\)H\(_{48}\)O\(_4\) in peaks (17, 20, 23, 31 and 32), dihydroxy-23-oxo-12-oleanen-28-oic acid (hydroxygypsogenin/quillaic acid) at \( m/z \) 485; C\(_{30}\)H\(_{45}\)O\(_5\) in peaks (9, 10, 12, 13, 16, 24 and 30), hydroxy-23-oxo-12-oleanen-28-oic acid (gypsogenin) at \( m/z \) 469; C\(_{30}\)H\(_{46}\)O\(_4\) in peaks (15 and 26) [5, 7, 43, 49, 50] and hydroxy-8-oleanen-29-oic acid (bryonolic acid) at \( m/z \) 455; C\(_{30}\)H\(_{48}\)O\(_3\) as in peak (25) that was reported before from L. cylindrica [51].

A fibrinolytic saponin lucyoside Nt (peak 9) was detected in MF [5], with a molecular ion (M–H\(^–\)) at \( m/z \) 1395.6017, C\(_{64}\)H\(_{99}\)O\(_{33}\). Tandem MS/MS revealed sequential loss of its attached sugars from fragments appearing at \( m/z \) 1233 (–162 amu, hexose); \( m/z \) 1101 (–132 amu, pentose); \( m/z \) 969 (–132 amu, pentose); \( m/z \) 823; C\(_{42}\)H\(_{63}\)O\(_{16}\) (–146 amu, rhamnose); \( m/z \) 643; C\(_{36}\)H\(_{51}\)O\(_{10}\) (–162 amu, hexose and 18 amu, H\(_2\)O) and quillaic acid at \( m/z \) 485; C\(_{30}\)H\(_{45}\)O\(_2\) (–176 amu, glucuronic acid) as showed in (Suppl. Fig. S3).

Flavonoids

6 Flavones were detected herein as confirmed from its UV absorbance spectra in both fruit stages as glucuronoide conjugates at \( m/z \) 461.0710 (\( \lambda_{max} \) 268, 346 nm), 445.0758 (\( \lambda_{max} \) 269, 338 nm) and 475.0867 (\( \lambda_{max} \) 270, 344 nm) in peaks (5, 6 and 7) along with their corresponding aglycones at \( m/z \) 285.0399, 269.0447 and 299.0554 for luteolin, apigenin and diosmetin, respectively in peaks (14, 18 and 19), illustrated in Suppl. Fig. S (4, 5 & 6). L. aegyptiaca is reported previously for the presence of flavone glycosides either as glucuronides or as methyl ester of glucuronic acid conjugates and to possess antioxidant, anti-emic and anti-inflammatory activities [9, 45].

Fatty acids

Several fatty acids (FA) were detected and found most abundant in YF (peaks 8, 21, 22, 28, 29, 34, 36, 37 and 39). Their order of elution was consistent with their degree of oxygenation starting with trihydroxylated and dihydroxylated unsaturated FA then finally saturated FA. Fatty acids have been reported as bioactive metabolites to lower diabetes incidence [52]. In contrast, biological actions reported for hydroxylated fatty acids (oxylipids) included cytotoxic, anti-inflammatory and antimicrobial effects [53], previously reported in luffa fruits and seeds [17, 54].
**Fig. 5.** Principal component analysis (PCA) and orthogonal projection to latent structures-discriminant analysis (OPLS) supervised data analysis of modelling young versus mature fruit specimens analysed via UHPLC-MS for their secondary metabolites. PCA score (A) and loading plot (B) \((n = 3)\); OPLS-DA score plot (C) and loading S-plot (D). Segregation in both score plots showed marker metabolites for YF as Mol.ion/Rt, namely: Lucyoside I \((649.39/11.9)\), Dihydroxy-23-oxo-12-oleanen-28-oic acid-O-dipentosyl deoxyhexosyl-O-hexosyl glucuronoside \((1233.54/11.21)\); Dihydroxy-23-oxo-12-oleanen-28-oic acid-O-deoxyhexosyl-hexoside \((793.43/11.78)\) and Dihydroxy-23-oxo-12-oleanen-28-oic acid-O-pentosyl dihexoside \((941.5/11.35)\).

**Table 2**

| No. | Rt (min.) | Identification | \((M-H)^{-}\) | Element composition | Error (ppm) | MSn ions m/z (-) ppm | Fruit \(\text{Young}^*\) | Fruit \(\text{Mature}^*\) |
|-----|-----------|----------------|----------------|---------------------|-------------|---------------------|----------------|----------------|
| 1   | 0.66      | Unknown        | \(\text{C}_1\text{H}_7\text{O}_4\) | 215.0324           | 6.8         | 179, 161, 89         | (+)            | (+)            |
| 2   | 0.87      | (Iso)citric acid | \(\text{C}_6\text{H}_7\text{O}_7\) | 191.0193           | 3.7         | 173, 111            | (+)            | (+)            |
| 3   | 9.9       | Gibberelin A8  | \(\text{C}_6\text{H}_7\text{O}_7\) | 363.1437           | 0.3         | 345, 319, 301, 275, 257, 239, 119 | (+)            | (+)            |
| 4   | 10.08     | Umbelliferone (internal standard) | \(\text{C}_6\text{H}_7\text{O}_7\) | 161.0241           | 4.8         | 133, 117            | (+)            | (+)            |
| 5   | 10.32     | Luteolin-O-glucuronide | \(\text{C}_6\text{H}_7\text{O}_7\) | 461.0709           | 1.1         | 285, 153            | (+)            | (+)            |
| 6   | 10.55     | Apigenin-O-glucuronide | \(\text{C}_6\text{H}_7\text{O}_7\) | 445.0757           | 1.7         | 269                | (+)            | (+)            |
| 7   | 10.7      | Diosmetin-O-glucuronide | \(\text{C}_6\text{H}_7\text{O}_7\) | 475.0867           | 0.78        | 299, 285, 270       | (+)            | (+)            |
| 8   | 10.8      | Trihydroxy-oxo-octadecenoic acid | \(\text{C}_6\text{H}_7\text{O}_7\) | 343.2116           | 0.2         | 325, 307, 289, 209, 201, 171, 153, 135 | (+)            | (+)            |
| 9   | 11.04     | Lucyoside N  | \(\text{C}_6\text{H}_7\text{O}_7\) | 1395.6016          | 2.8         | 1377, 1233, 1101, 969, 823, 643, 485 | (+)            | (+)            |
| 10  | 11.12     | Dihydroxy-23-oxo-12-oleanen-28-oic acid-O-tripentosyl-deoxyhexosyl-O-hexosyl glucuronide | \(\text{C}_6\text{H}_7\text{O}_7\) | 1365.5899          | 3.3         | 1347, 1233, 1101, 969, 823, 661, 643, 485 | (+)            | (+)            |
| 11  | 11.14     | Lucyoside A  | \(\text{C}_6\text{H}_7\text{O}_7\) | 811.4470           | 0           | 691, 649, 487       | (+)            | (+)            |
| 12  | 11.21     | Dihydroxy-23-oxo-12-oleanen-28-oic acid-O-dipentosyl deoxyhexosyl-O-hexosyl glucuronoside | \(\text{C}_6\text{H}_7\text{O}_7\) | 1233.5491          | 2.7         | 1101, 955, 823, 661, 643, 485 | (+)            | (+)            |
| 13  | 11.28     | Lucyoside J  | \(\text{C}_6\text{H}_7\text{O}_7\) | 809.4301           | 2           | 647, 485            | (+)            | (+)            |
| 14  | 11.29     | Luteolin     | \(\text{C}_6\text{H}_7\text{O}_7\) | 285.03998          | 1.5         | 1341, 1217, 807, 645, 627, 469 | (+)            | (+)            |
| 15  | 11.35     | Hydroxy-23-oxo-12-oleanen-28-oic acid-O-tripentosyl-deoxyhexosyl-O-hexosyl glucuronoside | \(\text{C}_6\text{H}_7\text{O}_7\) | 1349.9697          | 2           | 1331, 1217, 807, 645, 627, 469 | (+)            | (+)            |
| 16  | 11.35     | Dihydroxy-23-oxo-12-oleanen-28-oic acid-O-pentosyl dihexoside | \(\text{C}_6\text{H}_7\text{O}_7\) | 941.5097           | 0.7         | 779, 485            | (+)            | (+)            |
| 17  | 11.43     | Lucyoside C  | \(\text{C}_6\text{H}_7\text{O}_7\) | 795.4522           | 0.3         | 633, 471, 385       | (+)            | (+)            |
| 18  | 11.48     | Apigenin     | \(\text{C}_6\text{H}_7\text{O}_7\) | 269.04477          | 1.1         | 289                | (+)            | (+)            |
| 19  | 11.53     | Diosmetin    | \(\text{C}_6\text{H}_7\text{O}_7\) | 299.05548          | 1.1         | 289                | (+)            | (+)            |
| 20  | 11.55     | Lucyoside G  | \(\text{C}_6\text{H}_7\text{O}_7\) | 795.4516           | 1.09        | 633, 471, 405       | (+)            | (+)            |
| 21  | 11.6      | Trihydroxy-octadecenoic acid | \(\text{C}_6\text{H}_7\text{O}_7\) | 327.2166           | 3.3         | 299                | (+)            | (+)            |
| 22  | 11.66     | Trihydroxy-octadecenoic acid | \(\text{C}_6\text{H}_7\text{O}_7\) | 329.2323           | 3           | 311, 293, 229, 211, 171 | (+)            | (+)            |
| 23  | 11.7      | Dihydroxy-12-oleanen-28-oic acid-O-deoxyhexoside-O-dihexoside | \(\text{C}_6\text{H}_7\text{O}_7\) | 925.5141           | 1.4         | 779, 763, 617, 471  | (+)            | (+)            |
Miscellaneous

Other detected compounds included citric acid (peak 2) at m/z 191.0193; C₆H₇O₇ found at both fruit stages and a lactone diterpenoid plant growth hormone; Gibberellin A₈ (peak 3) at m/z 363.1437; C₁₉H₂₃O₇ with fragment ions at m/z 319 (-44 amu, M-H-COO), m/z 275 (-88 amu, M-H-COO-COO) and m/z 257 (-106 amu, M-H-C₂O₄-H₂O) [55] confirming its structure detected exclusively in MF. In contrast, 3 phosphatidyl-hexanoic acid derivatives were detected only in YF samples (peaks 35, 38 and 40) that are involved in plant cell membrane regulation [56,57]. In addition, phosphatidyl-ethanolamine derivatives are incorporated with leukemic antiproliferative agent (methotrexate) to provide effective transport system of cytoxic drug [58].

Multivariate PCA and OPLS data analyses of the UPLC/MS dataset:

In an attempt to distinguish between young and mature L. aegyptiaca fruits, PCA and OPLS were employed as in GC/MS data-set as analysed via UHPLC/MS (Fig. 5). PCA score plot showed
samples segregation along PC1 and PC2 both accounting for more than 85% of the variance, whereas OPLS model exhibited good prediction power of (Q2 = 0.96) and (R2 = 0.98). Both PCA loading plot and OPLS S-plot revealed for secondary metabolites markers for YF as dihydroxy-23-oxo-12-olean-28-oic-0-dipentsyl deoxyhexosyl-O- hexosyl glucuronide, dihydroxy-23-oxo-12-olean-28-oic acid-0-pentosyl dihexoside, dihydroxy-23-oxo-12-olean-28-oic acid-0-deoxyhexosyl-hexoside and trihydroxy-12-olean-28-oic acid hexoside (lucyoside I) (peaks 12, 16, 24 and 27). It is obvious that all YF markers contribute to polyoxygenated sapogenins i.e., (dihydroxyalted-oxo-sapogenins along with trihydroxylated sapogenin for lucyoside I), while no markers were detected for MF. Such metabolites abundance is likely attributed to metabolic changes that occur upon fruit lignification for fibrous vascular bundles hardening and channeling metabolic pathways towards production of higher molecular weight metabolites not detected by UHPLC/MS. Cross validation of UHPLC/MS derived OPLS-DA is presented in Suppl. Fig. S8.

Conclusion

In this study, detailed compositional variation in sensory and secondary metabolitess presented herein for the first time in L. aegyptiaca fruits at different developmental stages via an untargeted metabolic approaches based on GC/MS and UPLC–PDA/ES I–Orbitrap HRMS MS Datasets were further analyzed using chemometric tools to determine metabolites heterogenity for both fruit stages and to account for its different morphology or biological effects. A total of 53 volatiles were detected and to account for this fruit aroma categorized into diverse chemical classes, edible YF showed richness in sensory metabolites to mediate for its aroma viz., hexenal, nonadinal (cucumber aldehydyl) along with terpenoidal alcohol β-linalool compared to the inedible mature fruit. It could be speculated that enzymes catalyzing these metabolites biosynthetic pathways are down-regulating in luffa fruit upon ripening. UHPLC/MS revealed 40 metabolites with saponins of oleanene triterpene glycosides as major bioactive secondary metabolites in both fruit stages, and with polyoxygenated sapogenins glycosides; dihydroxy-23-oxo-12-olean-28-oic acid derivatives and lucyoside I as metabolic markers in YF versus no marker to contribute for unedible lignified MF. Such chemical differences between the two Luffa aegyptiaca stages could lead to variation in biological effects, yet though to be determined in future work to exploit uses specific for each fruit maturity stage.

Compliance with Ethics requirements

This article does not contain any studies with human or animal subjects.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary material

Supplementary data to this article can be found online at https://doi.org/10.1016/j.jare.2019.10.009.

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