**Supporting information**

**Focus on Mascarenes endemic plants with specific phytochemical composition, potent antioxidant and antiproliferative properties.**

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**Experimental methods**

**GC-MS ANALYSIS**

The GC-MS analysis was carried out using the Agilent 7890A gas chromatography systems. The apparatus included an HP5-MS column coupled with Agilent MS model 5975C MSD with Triple axis detector (Agilent Technologies, US). The analysis began with the initial oven temperature set at 60°C for 2min, which increased at the rate of 10°C/min to a maximum of 300°C, and was maintained for another 4 min to yield a total run of 30 min under a constant helium pressure (10psi). Trimethylsilylimidazole (TMSI) derivatization of fractions was achieved by adding 10 µL of pyridine was added, followed by 50 µL of N, O-Bis(trimethylsilyl)trifluoroacetamide (BSTFA) to an Eppendorf tube containing 1 mg of lyophilized extract and incubated at 40°C – 50°C for 1 hour. Analysis was conducted by injecting 2 µL of the Trimethylsilylimidazole (TMSI) derivatized sample into the Agilent 7890A gas chromatography systems and the chromatogram analyzed by matching the MS spectra of the peaks with those stored in the NIST 2011 Mass Spectral Library (Agilent Technologies, USA) (**Figure S4**).

**LC-MS ANALYSIS**

LC-MS analysis was achieved using a HESI II electrospray ion source on a Thermo Scientific Q-Exactive Orbitrap mass spectrometer system coupled to a Waters BEH C18 (2.1x100 mm 1.7 µm particle size) reversed phase column via a Thermo Scientific Ultimate U3000 ultra-performance liquid chromatography system. Fractions or isolated compounds were separated using a linear gradient method utilising mobile phase A (0.1% formic acid in water) and mobile phase B (acetonitrile with 0.1% formic acid) at a flow rate of 0.4 mL/min The gradient consisted: 0 min (95% A), 1 min (95% A), 7 min (100% B), 7.01 min (95% A) and 10 min (95% A). The MS method utilised a full-MS experiment in negative ion mode with 140,000 resolutions and a scan range of 300-2500.

**NMR ANALYSIS**

1H NMR (500 MHz), 13C NMR (125 MHz), and 2D-NMR (COSY, HSQC and HMBC) spectra were obtained on a Bruker AVII 500 MHz instrument equipped with a dual 13C/1H cryoprobe. Chemical shifts were reported in δ (ppm) referenced to the solvents shifts (methanol-d4 or acetone-d6) and coupling constants (J) were measured in Hertz.

**HepG2 CELL VIABILITY PROFILES**

|  |  |
| --- | --- |
| Cell viability (% control)  **B** | **A** |
|  |
| **C** |
|  | *T. bentzoë* crude/ fractions cconcentration (µg/ml) |

**Figure S1: Cell viability profile of HepG2 cells treated with *T. bentzoë* extract and fractions.** HepG2 cells were incubated with varying concentrations of test extracts for 48 hours and cell viability was determined using MTT assay. For each concentration, the percentage of cell viability was calculated relative to the negative control (cells treated with 0.125 % DMSO).

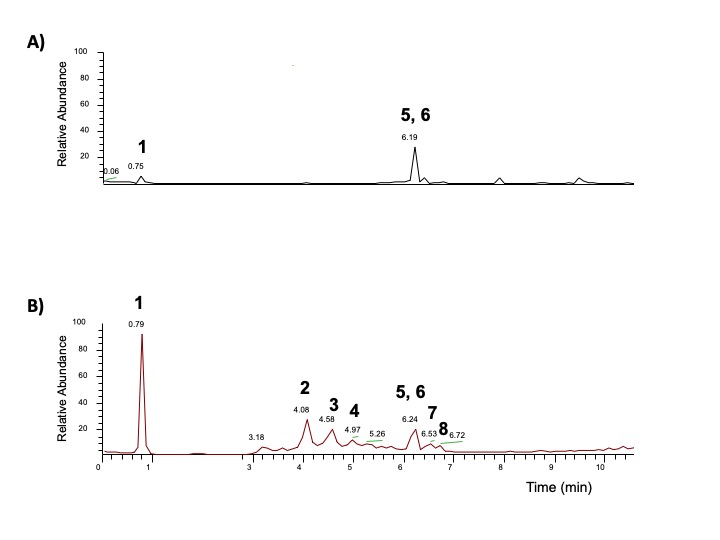
**HepG2 FLOW CYTOMETRIC PROFILES.**

|  |  |
| --- | --- |
| 2.  A |  |
| B | C |
| D | E |
| F | J |

**Figure S2: Representatives annexin V-FITC/PI flow cytometric profile of HepG2 cells, 48 hours post extract/control treatment.** Cells were treated for 48 hours with (A) negative control (0.025% DMSO); (B, C, D) 10 µg/ml, 20 µg/ml and 40 µg/ml of *T. bentzoë*, respectively; (E, F, G) 1 µg/ml, 2 µg/ml and 4 µg/ml of etoposide, respectively.

|  |  |
| --- | --- |
| A |  |
| B | C |
| E |  |
| D  F | G |

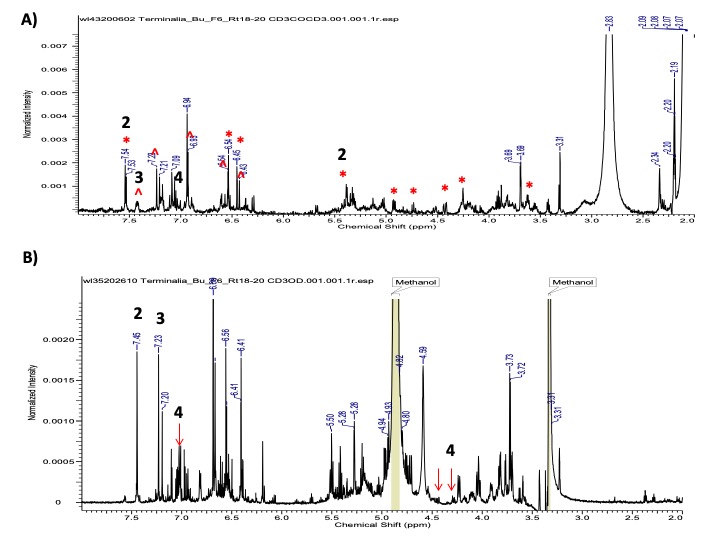
**Figure S3: Representatives cell cycle histogramof HepG2 cells, 48 hours post extract/control treatment.** Cells were treated for 48 hours with (A) negative control (0.025% DMSO); (B, C, D) 10 µg/ml, 20 µg/ml and 40 µg/ml of *T. bentzoë*, respectively; (E, F, G) 1 µg/ml, 2 µg/ml and 4 µg/ml of etoposide, respectively.



**Figure S4: UPLC ESI MS of the butanol fraction 6 at the positive (A) and negative (B) mode.** The identified compounds are indicated in Table 6.

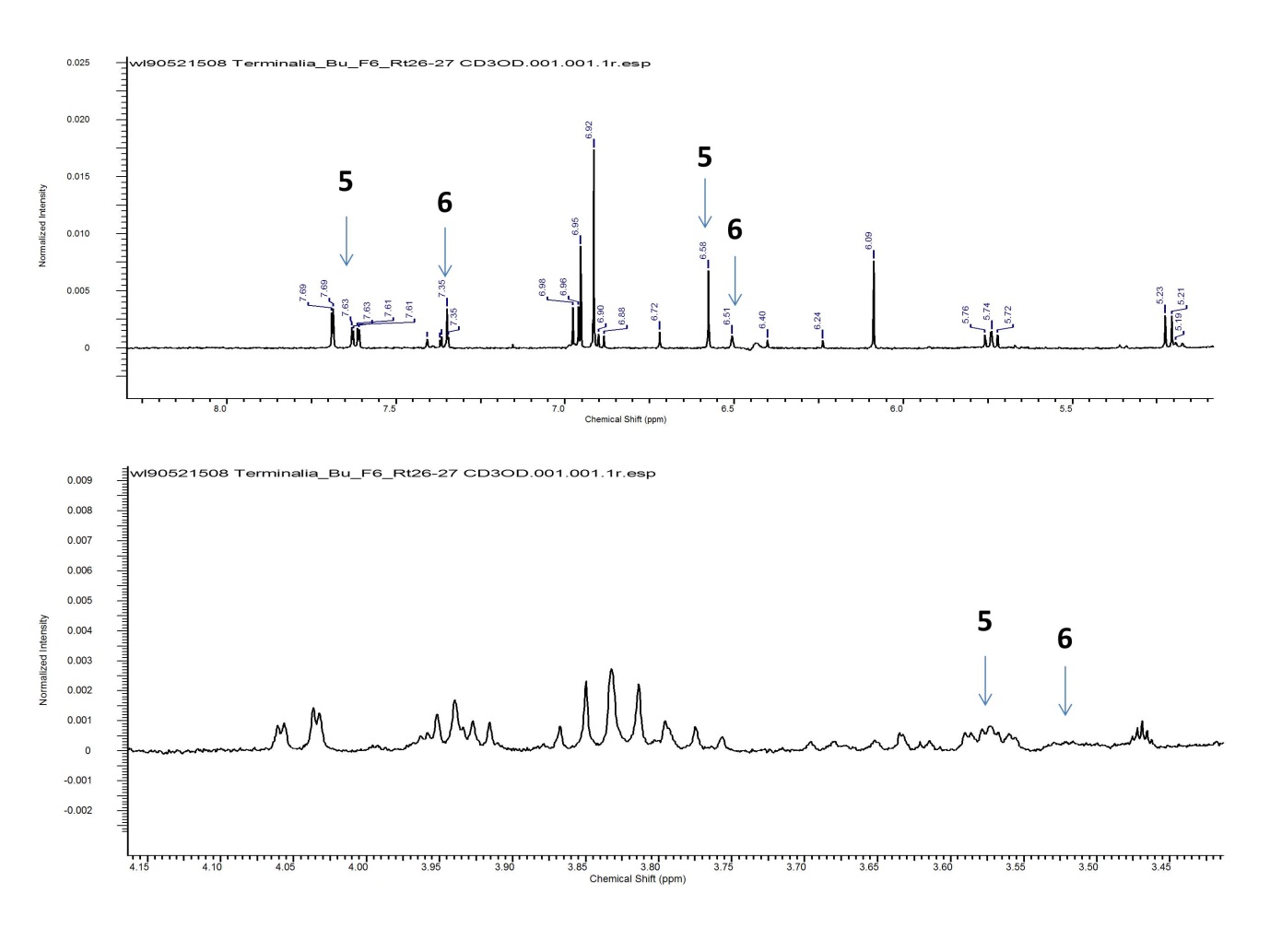


**Figure S5: 1H NMR spectrum (A) and HSQC spectrum (B) for punicalagin (1) present in a mixture in an HPLC fraction at RT 2-3 min in methanol-d4.** The nmr peaks marked as \* are consistent with those of punicalagin [1–4].

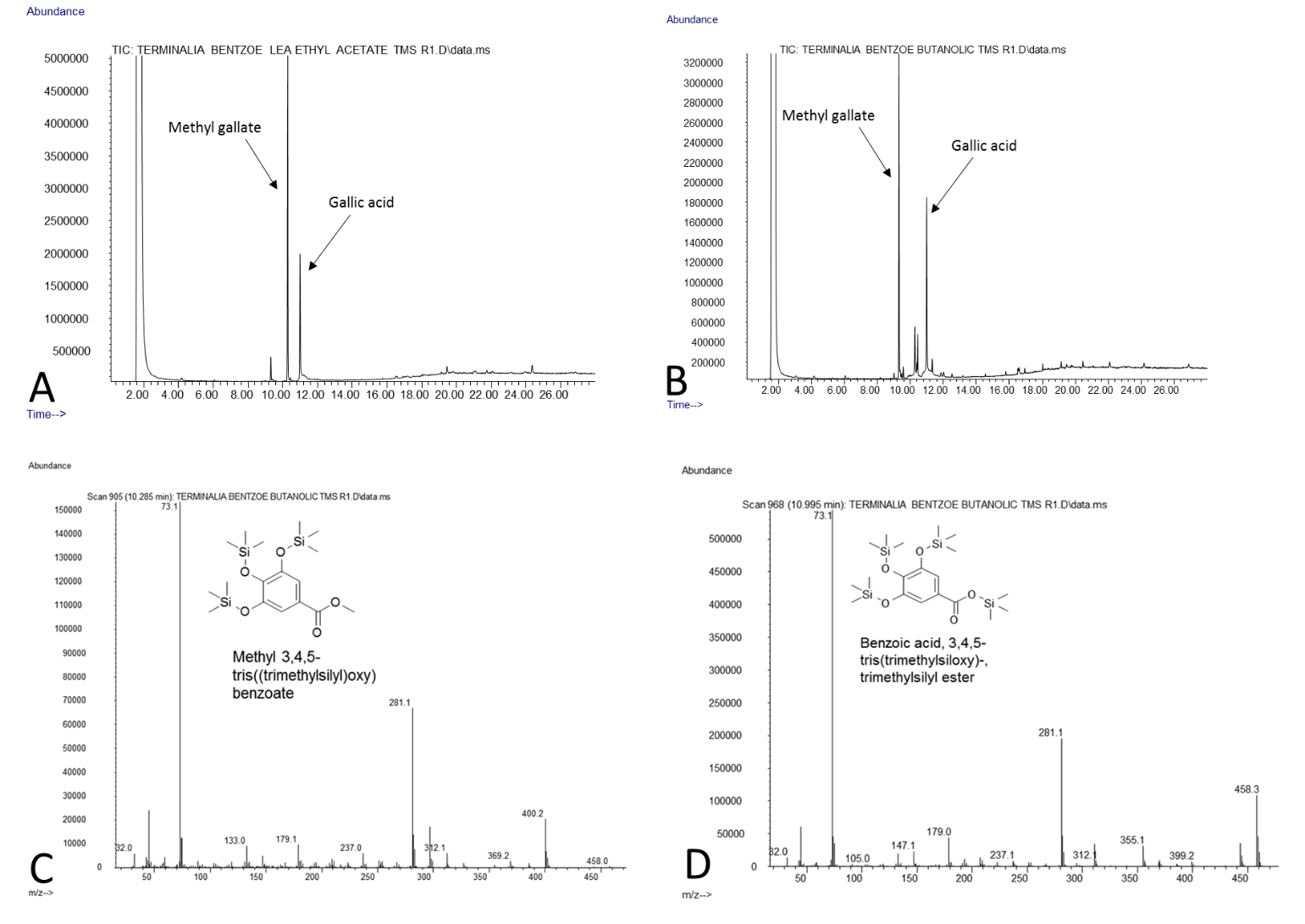


**Figure S6: 1H NMR spectra for a mixture of isoterchebulin (2), terflavin A (3), and**

**3,4,6-trigalloyl-beta-D-glucopyranose (4) in a HPLC fraction at RT18-20 min in acetone-d6 (A) and methanol-d4 (B).**The NMR peaks masked as \* and ^ in acetone-d6 are consistent with those of isoterchebulin (**2**) [1] and terflavin A (**3**) [5], respectively. The 1H NMR peaks measured in methanol-d4 (B) are consistent with those of 3,4,6-trigalloyl-beta-D-glucopyranose (**4**) [6].



**Figure S7: 1H NMR spectra for a mixture of 2"-O-galloyl-orientin (3) and 2"-O-galloyl-isoorientin (6) with a ratio of 5:2 based on the integration of single hydrogen in each compound.** The assignment of each peak is listed in **Table S1**.



**Figure S8: GC Chromatogram of TMSi derivatives of ethyl acetate (A) and butanol (B) fraction of *T. bentzoë*.** Mass spectrum of the peak for TMSi derivative of methyl gallate (C) and gallic acid (D) present in both organic fractions. Methyl gallate, GC-MS of TMSi derivative of methyl gallate, Rt, 11.030 min, *m/z* 400.2 [M]+, 369.2, 312.2, 281.1,179.1, 133.0, 73.1. The data are consistent with those stored in the NIST 2011 Mass Spectral Library (Agilent Technologies, USA). Gallic acid, GC-MS of TMSi derivative of gallate, Rt, 10.320 min, *m/z* 458.1 [M]+, 355.1, 314.9, 281.1, 73.1. The *m/z* pattern is consistent with those previously reported [7].

**Table S1 1H NMR spectral data (500 MHz, CD3OD) of the O-galloyl-C-glycosylflavones 7 and 8 (5:2) [" in ppm, multiplicities and J values (Hz) are given in parentheses].**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| proton | 7 | 2”-O-Galloylorientin  [2] | 8 | 2”-O-Galloylisoorientin  [2] |
| 3 | 6.58, 6.40a (s) (10:1) | 6.55, 6.37a (s) | 6.51 (s) | 6.45 (s) |
| 6 | 6.09, 6.24a (s) (10:1) | 6.06, 6.21a (s) | 7.35 (d; 2.0) | 7.34 (d; 2.0) |
| 8 |  |  | 6.43 (s) | 6.39 (s) |
| 2’ | 7.69, 7.37a (d; 2.2) | 7.68, 7.38a (d; 1.9) |  |  |
| 5’ | 6.97 (d; 8.5) | 6.97 (d; 8.4) | 6.89 (d; 8.0) | 6.89 (d; 8.9) |
| 6’ | 7.62, 7.40a (dd; 8.4, 2.2) | 7.61, 7.40a (dd; 8.4, 1.9) | 7.36 (dd; 8.0, 2.0) | 7.35 (dd; 8.9, 2.0) |
| *Glucosyl* |  |  |  |  |
| 1’’ | 5.22 (d; 10.1) | 5.22, 5.35a (d; 10.1) | 5.18 (d; 10.1) | 5.16 (d; J=10.0) |
| 2’’ | 5.74 (t; 10.1) | 5.74, 5.66a (t; 9.7) | 5.74 (t; 9.1) | 5.72 (br t) |
| 3” | 3.85 (t; 9.0) | 3.83 (t; 9.0) | 3.78 (t; 9.3) | 3.76 (t; 9.6) |
| 4” | 3.81 (t; 9.0) | 3.79 (t; 9.0) | 3.63 (t; 9.4) | 3.61 (t; 9.4, 9.1) |
| 5” | 3.57 (m) | 3.55 (m) | 3.52 (m) | 3.50 (m) |
| 6a” | 4.05 (dd; 2.2, 12.1) | 4.02 (dd; 1.9, 12.1) | 3.93 (overlapped) | 3.93 (dd; 12.0, 1.8) |
| 6b” | 3.93 (dd; 6.0, 12.1) | 3.91 (dd; 5.8, 12.1) | 3.80 (overlapped) | 3.79 (dd; 5.5, 12.0) |
| *Galloyl* |  |  |  |  |
| 2’’’ | 6.92, 6.72a (s) | 6.89, 6.69a (s) | 6.95 (s) | 6.93 (s) |
| 6” | 6.92, 6.72a (s) (10:1) | 6.89, 6.69a (s) | 6.95 (s) | 6.93 (s) |

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