

***Carissa macrocarpa* leaves polar fraction ameliorates doxorubicin-induced neurotoxicity in rats via downregulating the oxidative stress and inflammatory markers**

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Supplementary Material

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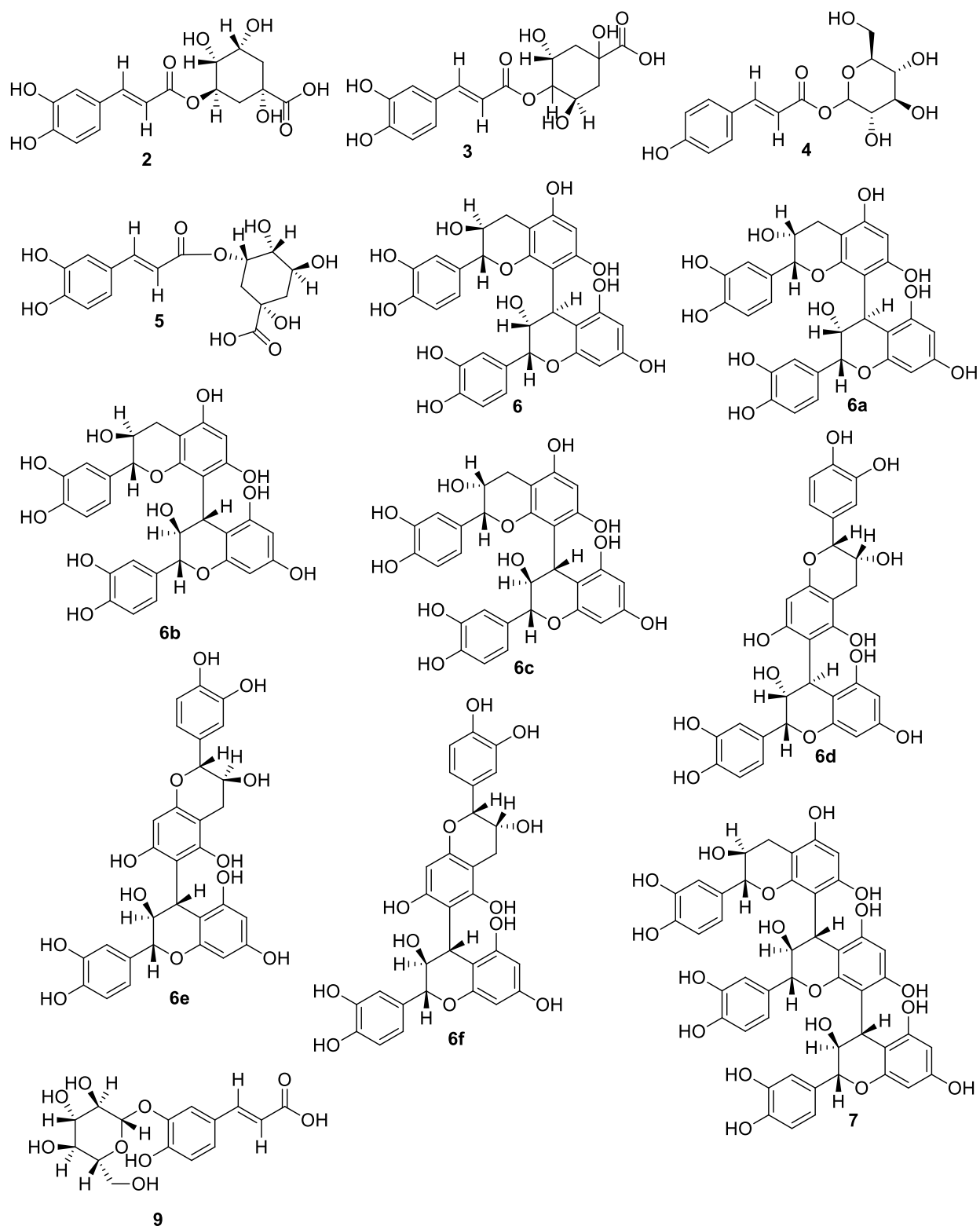


Figure S1. Structures of compounds evaluated by molecular docking simulation.

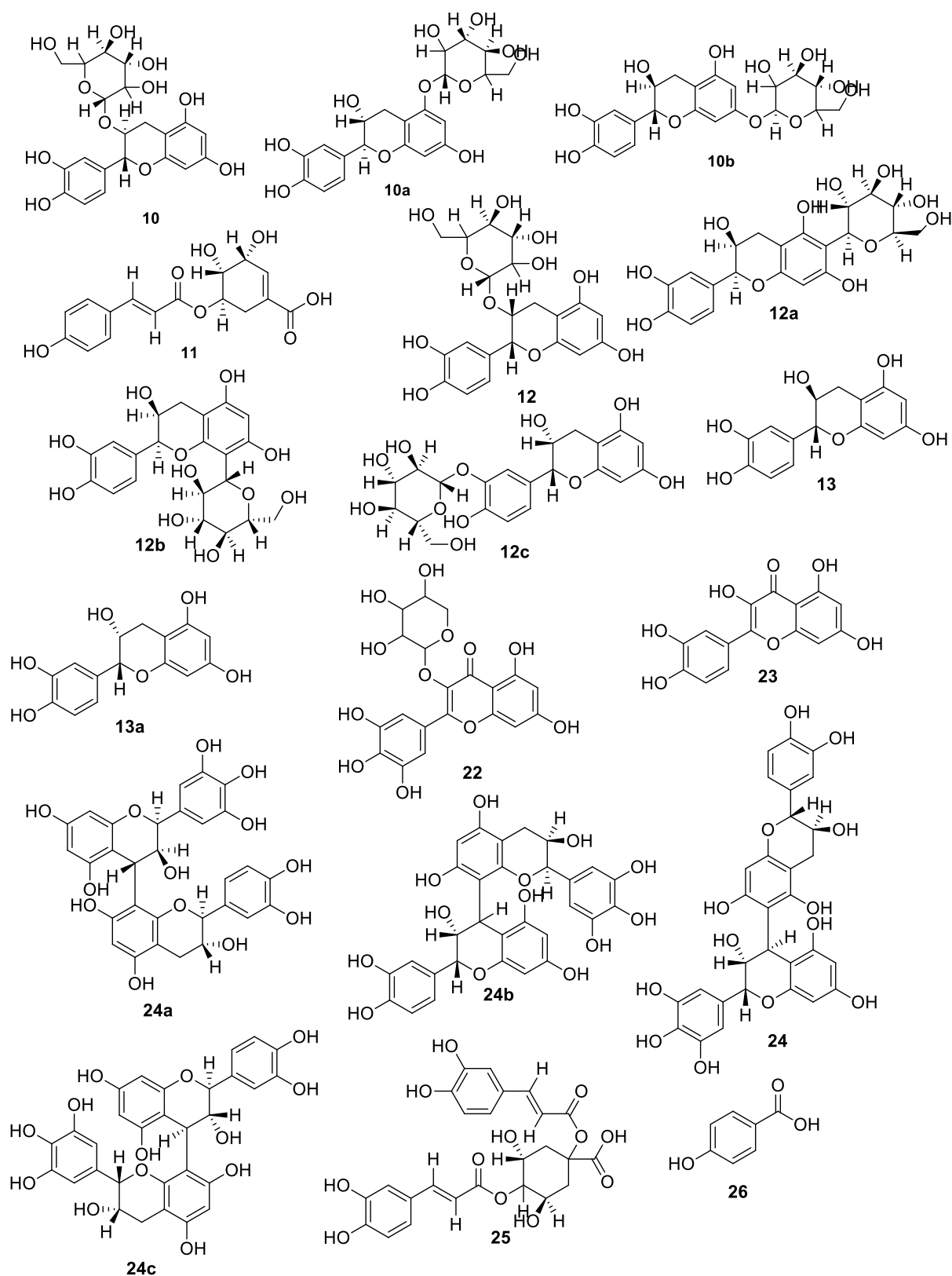


Figure S2. Structures of compounds evaluated by molecular docking simulation: Continued

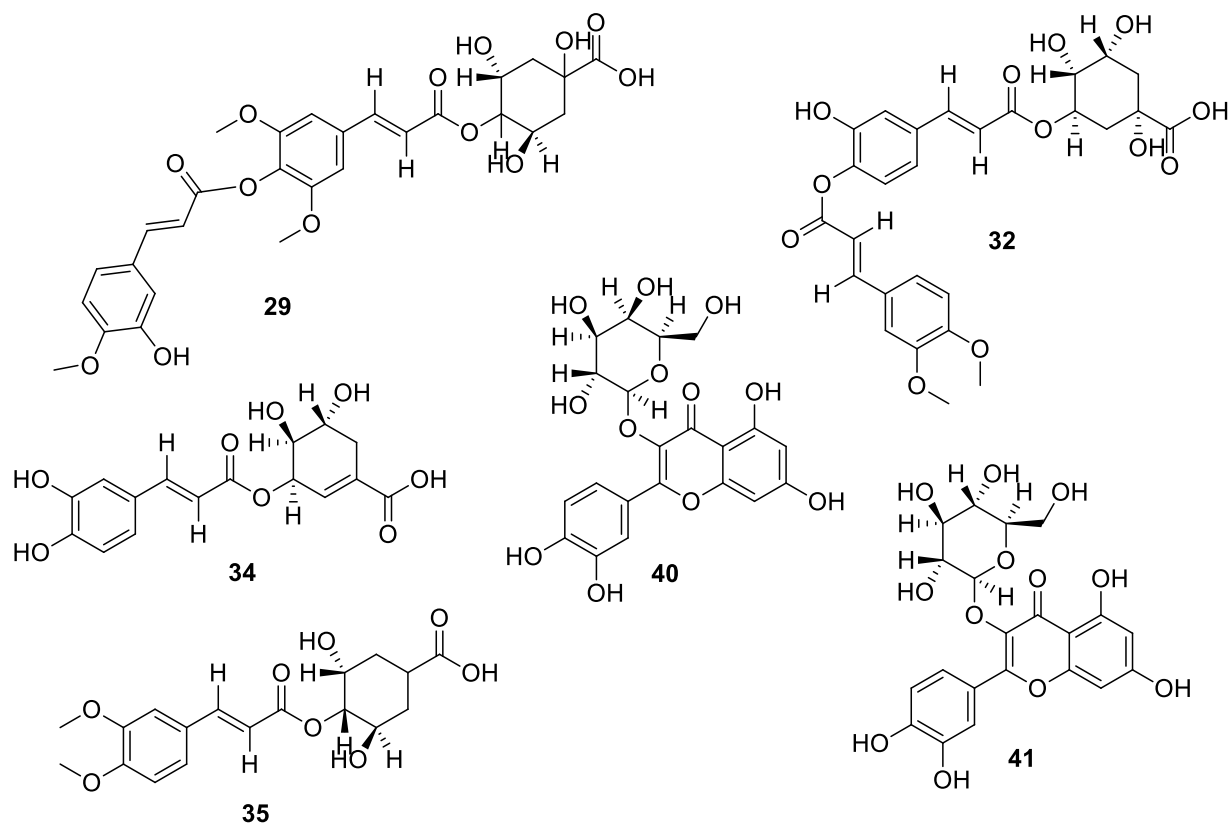


Figure S3. Structures of compounds evaluated by molecular docking simulation: Continued

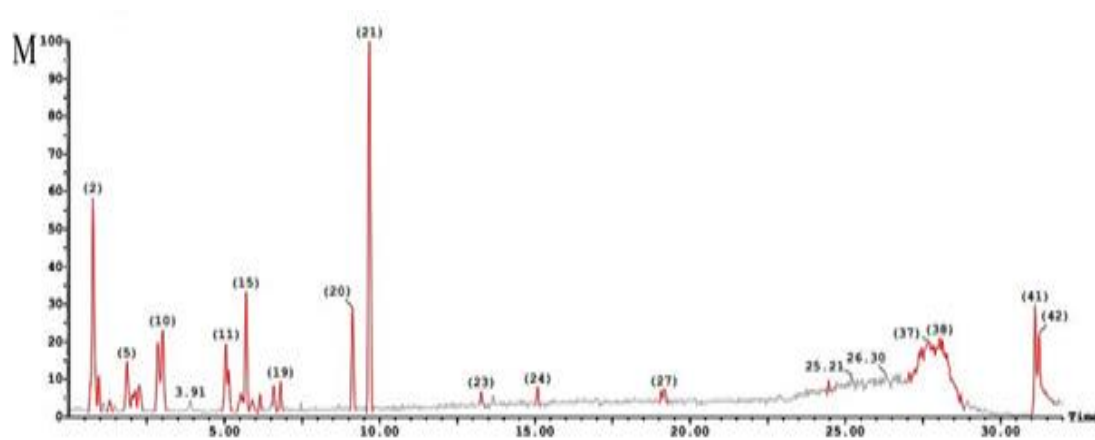


Figure S4. TIC of *C. macrocarpa* polar fraction of the leaves using UPLC-ESI-MS/MS in negative ionization mode.

Table S1. Docking score of detected polyphenolics and their isomers against TACE enzyme

No.	Compound Name	TNF-activating converting enzyme (TACE)	
		Score (kcal/mol)	RMSD refine (Å)
2	Chlorogenic acid	-12.6813	0.91
3	Cryptochlorogenic acid	-13.3260	0.93
4	coumaroyl-5- β -glucose	-11.9755	1.07
5	5- <i>O</i> -Caffeoylquinic acid	-13.6537	0.96
6	Procyanidin B1	-16.2139	2.66
6a	Procyanidin B2	-17.5405	1.82
6b	Procyanidin B3	-13.9250	1.60
6c	Procyanidin B4	-17.5085	0.84
6d	Procyanidin B5	-19.9566	1.77
6e	Procyanidin B6	-21.9152	1.25
6f	Procyanidin B8	-17.0187	1.92
7	Procyanidin C2	-13.2742	1.80
9	Caffeic acid 3-glucoside	-12.8895	1.44
10	Catechin 3- <i>O</i> - β -D-glucopyranoside	-18.5367	1.69
10a	Catechin 5- <i>O</i> - β -D-glucopyranoside	-17.1693	1.31
10b	Catechin 7- <i>O</i> - β -D-glucopyranoside	-15.3866	1.57
11	4- <i>O</i> - <i>p</i> -Coumaroylshikimic acid	-11.6970	1.24
12	Epicatechin 3- <i>O</i> - β -D-glucopyranoside	-20.6389	1.34
12a	Epicatechin 6-C-glucoside	-19.2057	1.30
12b	Epicatechin 8-C-glucoside	-19.1597	0.93
12c	Epicatechin-3'- <i>O</i> -glucoside	-14.7443	1.74
13	Catechin	-14.2871	1.02
13b	Epicatechin	-14.1969	0.84
22	Myricetin-3- <i>O</i> -xyloside	-18.2396	1.15
23	Quercetin	-13.1148	1.19
24a	Epigallocatechin-(4 β -8)-catechin	-18.0768	1.27
24b	Epicatechin (4 β .8) epigallocatechin	-18.3569	1.97
24c	Catechin-(4 α -8)-(-)-epigallocatechin	-18.0083	1.44
24	Epigallocatechin-(4 β -6)-(+)-catechin	-14.8778	1.82
25	1,4-Dicaffeoylquinic acid	-15.0462	1.10
29	Feruloyl- <i>O</i> -sinapoylquinic acid	-15.8150	1.08
32	Dimethoxycinnamoyl- <i>O</i> -caffeoylquinic acid	-13.6193	1.00
34	3-Caffeoylshikimic acid	-15.6133	0.89
35	Dimethoxycinnamoylquinic acid	-13.1423	1.48
40	Hyperoside	-19.9102	1.47
41	Isoquercetin	-16.0728	1.10

UPLC-ESI-MS-MS metabolites characterization

Organic phenolic acids and their derivatives

Hydroxybenzoic acid (**26**) and syringic acid (**28**) were detected at different retention times as reported in Table 1. Compounds **2**, **3**, **5**, and **34** that showed a molecular ion at m/z 353 were detected at retention times 1.87, 2.03, 2.16, 15.06, and 16.52, respectively. Compounds **2** and **5** produced MS² fragments at m/z 191 ([quinic acid-H]), 179 ([Caffeic acid-H]), 161 ([caffeic acid-H₂O-H]), hence, assigned as 3-O-caffeolyquinic acid and 5-O-caffeolyquinic acid, respectively. Where 4-O-caffeolyquinic acid was designated for compound **3** with characteristic MS² fragments at m/z 191, 173 [191-H₂O], and 161 [1].

Additionally, 3-O-caffeoylshikimic acids has been proposed for **34**, depending upon MS² fragments (at m/z 179, 161) and published data [2]. Another caffeolyquinic acid derivative, diacetoxy-5-methoxyphenyl) acroyl-O-*p*-coumaroyl-O-caffeoylquinic acid (**30**) was identified by the aid of MS, MS² data (Table 1), and reported literature [3].

Other hydroxycinnamate derivatives such as 5-O-*p*-coumaroylshikimic and 4-O-*p*-coumaroylshikimic acids have been suggested for peaks **8** and **11**. MS² fragmentation exhibited a base peak at m/z 163 due to [*p*-coumaric acid-H₂O], as well as other characteristic fragments [3]. Moreover, peak **25**, which was detected at retention time 6.58 showed deprotonated molecular ion at m/z 515 and distinctive fragments at m/z 353 [M-162] and 179 that are matched with the reported literature of dicaffeoylquinic acid (**25**). Feruloyl-O-sinapoylquinic acid (**29**) was designated for peak detected at m/z 573 [M-H]⁻, based on MS² fragments (397 [M-ferulic acid-H], 223 [sinapic acid-H], 173 [quinic acid-H₂O-H]) as compared with the literature [2]. Besides, compounds, dimethoxycinnamoyl-O-caffeoylquinic acid (**32**) and its isomer (**33**) were observed at different retention times 13.30 and 15.66 but exhibited similar deprotonated molecular ions at m/z 543 and characteristic fragment ions at m/z 353 [M-dimethoxycinnamoyl-H₂O] and 173 that attributed to dimethoxycinnamoyl moiety. In addition, dimethoxycinnamoylquinic acid was suggested for peak **35** that was detected in positive ionization mode at m/z 383 (R_t 16.59) [2]. Moreover, hydroxycinnamate glycosides were detected in the current study at retention times 2.13 and 2.87, that are corresponding to peaks **4** and **9**, respectively. Where peaks **4** was suggested to be coumaric acid hexoside, due to molecular ions at m/z 325 [M-H]⁻ which produced fragment ions at m/z 187, 163, and 145 in accordance with the literature [4]. Also, caffeic acid hexoside was assigned for peak **9** depending upon its molecular ion at m/z 343 [M+H]⁺ with distinctive fragments at 326, 311, and 285, which in agreement with the reported data [5].

Flavonoids aglycone and glycosides

Flavonoids are considered one of the major polyphenolics detected in the polar fractions of the species under investigation. Peak **14** and **15** provided the same deprotonated molecular ion at m/z 755 [M-H]⁻ with different MS² fragments, which were 593 [M-162, sugar unit] and 285 [M-308, rutinoside unit] for peak **14** that assigned as kaempferol-O-hexoside-O-rutinoside. However, quercetin-O-deoxyhexoside-O-deoxyhexosyl-hexoside was proposed for peak **15** that produced fragment ions at m/z 609 [M-146, deoxyhexosyl unit] and 301 [M-146-308]⁻, deoxyhexosyl and rutinoside units]. Other kaempferol derivatives were kaempferol-O-

deoxyhexoside-O-deoxyhexosyl-hexoside isomers, which were designated for peaks **17** and **18**, respectively. Both compounds provided the same molecular ions (m/z 739 $[M-H]^-$) and MS² fragmentation patterns (m/z 593 and 285). Moreover, peaks **19** and **20** were tentatively identified as quercetin-O-deoxyhexosyl-hexoside isomers based on their molecular ion at m/z 609 $[M-H]^-$ and MS² fragment at m/z 301 due to loss of rutinoside unit (−308 amu). Also, the other isomer exhibited fragment ion at m/z 465 in positive ionization mode due to loss of deoxyhexosyl unit (−146 amu). All the previous results were confirmed upon comparison with the literature [1].

Additionally, compounds **40** and **41** belong to quercetin derivatives and have been identified as quercetin-3-O-galactoside, and quercetin-3-O-glucoside, respectively, as depicted from their MS, MS² data and comparison with literature [6]. Where, compounds **40** and **41** provided the same molecular ion at m/z 465 $[M+H]^+$ and MS² fragments at m/z 301, 300, 257, 255, 229, 179, and 151.

Furthermore, compound **22** is belonging to flavonoid glycoside [3] that was tentatively identified as myricetin pentoside [7] based on its molecular ions at m/z 449 $[M-H]^-$. Myricetin pentoside showed characteristic MS² fragment at 317 $[M-132]$. Finally, flavonoid aglycone such as quercetin has been proposed for peaks **23**.

Flavan-3-ol derivatives

C. macrocarpa is considered a main source of flavan-3-ol bioactive metabolites. From these interesting molecules, compound **13** has been identified as (epi) catechin based on a molecular ion at m/z 289 $[M-H]^-$, MS² fragmentation (m/z 245, 205, 203, 187, 179, and 161) and comparison with literature [8]. Furthermore, compounds **6**, **7**, **21**, and **31** belong to (epi)catechin derivatives of B-type; thus, (epi)catechin dimer was assigned for peaks **6**, **21**, and **31** based on deprotonated molecular ion at m/z 577 $[M-H]^-$, MS² fragmentation (m/z 425, 289) and comparison with the literature [1]. However, peak **7** that showed pseudo-molecular ion at m/z 865 $[M-H]^-$, and fragmentation patterns at m/z 451, 425, 407, and 289, were tentatively identified as (epi)catechin trimer [1].

Moreover, compounds **10**, **12**, and **16** provided the same deprotonated molecular ion at m/z 451 $[M-H]^-$ and different MS² fragments at m/z 408, 393, 351, 337, 301, 273, and 245 for compound **10** that suggested to be catechin 3-O-glucoside. However, epicatechin 3-O-glucoside and its isomer were designated for compounds **12** and **16** with MS² fragments at m/z 391, 343, 301, 287, 273, and 247 [9].

Compounds **24** and **27** were detected at the same molecular ion at m/z 593 $[M-H]^-$ and showed MS² fragments at m/z 557, 467, 441, 425, 407, and 289, hence proposed to be (epi)gallocatechin-(epi)catechin [7].

Sterols and triterpenes

Stigmasterol (**39**) was detected methanol soluble fraction [10,11]. Furthermore, the current study detected several triterpenoid aglycones at different retention times, such as compounds **36**, **37**, and **38**. Where compounds **36**, **37**, and **38** provided the same molecular ion at m/z 455 $[M-H]^-$,

MS² fragment ions (m/z 439, 419, 411, 410, 407, and 397) and designated as ursolic acid, carissic acid, and oleanolic acid upon comparison with literatures [12–15].

Miscellaneous

Lignans such as carinol (m/z 377 [M–H][–] and MS² at 333, 271, 257, 163, and 119) was assigned for compound **(1)** depending on a comparison of its data with that previously reported [13]. Compound **42** with m/z 621 [M–H][–] and distinctive fragment ion at 501 was assigned as 2(R)-26-([(2E)-3-(4-hydroxy-3-methoxyphenyl)-1-oxo-2-propen-1-yl]oxy)-2,3-dihydroxypropylester [16]. Finally, compound **43** that detected at retention time at 31.25 possessed molecular ion at m/z 429 [M–H][–] and 431 [M+H]⁺ with characteristic MS² fragments at m/z 205, 191, 177, 149, 121, hence assigned as α -tocopherol [17].

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