



Polyphenolic profiles in edible *Annona* spp. using high-performance liquid chromatography (HPLC-MS/MS)

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Abstract

Polyphenolic compounds of fruits of various plant species play an important role in physiological functions related to human health. Polyphenols have important biological activities like antioxidant activity which helps in normal functioning of human body. The objective of this study was to investigate the profiles of polyphenolic compounds in different edible *Annona* spp. fruits. Total of eighteen identified individual phenolic compounds were found, among which *p*-coumaric acid (321.53-90.17 $\mu\text{g g}^{-1}\text{FW}$), *o*-coumaric acid (70.80-19.00 $\mu\text{g g}^{-1}\text{FW}$), 2,4-dihydroxybenzoic acid (39.49-10.43 $\mu\text{g g}^{-1}\text{FW}$), caffeic acid (35.26-3.43 $\mu\text{g g}^{-1}\text{FW}$), gentisic acid (24.69-10.46 $\mu\text{g g}^{-1}\text{FW}$), protocatechuic acid (17.04-4.23 $\mu\text{g g}^{-1}\text{FW}$), *t*-cinnamic acid (22.68-3.93 $\mu\text{g g}^{-1}\text{FW}$) and ferulic acid (21.78-3.43 $\mu\text{g g}^{-1}\text{FW}$) were abundant in annona fruits while benzoic acid (23.28-4.61 $\mu\text{g g}^{-1}\text{FW}$), *p*-hydroxybenzoic acid (1.79-0.31 $\mu\text{g g}^{-1}\text{FW}$), salicylic acid (6.00-2.40 $\mu\text{g g}^{-1}\text{FW}$), 3-hydroxybenzoic acid (6.05-0.88 $\mu\text{g g}^{-1}\text{FW}$), vanillic acid (19.13-2.16 $\mu\text{g g}^{-1}\text{FW}$), gallic acid (15.88-2.74 $\mu\text{g g}^{-1}\text{FW}$), ellagic acid (1.12-0.20 $\mu\text{g g}^{-1}\text{FW}$), syringic acid (0.78-0.34 $\mu\text{g g}^{-1}\text{FW}$) and sinapic acid (2.16-0.79 $\mu\text{g g}^{-1}\text{FW}$) were limited. However, chlorogenic acid was not detected. The results obtained in this study will furnish a better knowledge of the polyphenolic composition in annona fruits.

Key Words: *Annona* spp., HPLC-MS/MS, *o*-coumaric acid, *p*-coumaric acid, polyphenols

Introduction

Biological properties of polyphenolic compounds like antioxidant activities are essential to prevent many diseases including cancer, rheumatoid arthritis, cardio vascular and diseases related to degenerative process (Almeida *et al.*, 2011; Dembitsky *et al.*, 2011). Several studies have shown high correlation between the consumption of fruits in prevention and treatment of various diseases (Loizzo *et al.*, 2012). The genus *Annona* belongs to the annonaceous family and comprises of approximately 162 species of trees and shrubs (Chatrou *et al.*, 2012). Economically, this genus is the most important of the annonaceae family due to its edible fruits and medicinal properties. While, there has been no comparative study on the biochemical properties of these fruits (Sousa *et al.*, 2012). In this paper, a comprehensive study on the

polyphenolic profiles of eight different annona genotype using an HPLC method with a TQD (Triple Quadrupole) mass spectrometry is reported.

Material and Methods

Fresh annona fruits used for experimentation were procured from the orchards of Bioersity International, Indian Institute of Horticultural Research (IIHR) and Regional Horticulture Research and Extension Centre (RHREC), Bengaluru. Ripened fruits of uniform size, shape and maturity, free from visible damages were procured and carefully placed in corrugated fibre board boxes and brought to laboratory.

The individual polyphenolic acids for High performance liquid chromatography-tandem mass spectrometry (HPLC-MS/MS) (Acquity Class-H system, Waters, Burnsville, USA) analysis were isolated using 80 per cent methanol (Weidner *et al.*, 2000). 10 grams of *Annona* fruit pulp was homogenized in methanol (80%), centrifuged and volume was made up to 50 mL. 20 mL of extract was evaporated near to dryness under the vacuum at 45°C and then diluted to 5 mL with water. Later

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extracted thrice with petroleum ether and then in 40 mL of ethyl acetate using separating funnel. Aqueous layer formed was discarded; ethyl acetate extract was evaporated to dryness under vacuum at room temperature. To the dry residue, 4 mL of 2N NaOH was added and allowed overnight for hydrolyzing. Extract was acidified to pH 2 by using 5 mL 2N HCl, again re-extracted with 50 mL ethyl acetate. Ethyl acetate layer was again re-extracted twice with 25 mL of 0.1N NaHCO₃. The aqueous layer was further acidified to pH 2 with 5 mL 2N HCl and extracted thrice with 25 mL ethyl acetate, the ethyl acetate layer was dried completely in rotary evaporator and the residue was dissolved in 2 mL MS grade methanol, filtered through 0.2 µm nylon filter prior to injection into HPLC-MS/MS system (Acquity Class-H system, Waters, Burnsville, USA) for polyphenolic acid estimation.

Table 1. Treatment details

SN	Species / Varieties used in the study
1.	<i>Annona squamosa</i> L.(Sweetsop)
a.	Balanagar
b.	Arka Sahan
c.	Red Sitaphal
2.	<i>Annona atemoya</i> Hort.
3.	<i>Annona reticulata</i> L. (Ramphal)
4.	<i>Annona muricata</i> L.(Soursop)
5.	<i>Annona cherimola</i> M. (Cherimoya)
6.	<i>Annona glabra</i> L. (Pond apple)

LC and MS-MS conditions

The polyphenolic acids were resolved on the analytical column BEH-C18 (2.1 x 50 mm, 1.7 µm) from waters India Ltd., protected by a Vanguard BEH C-18 (Waters, USA) with the gradient flow of organic and aqueous phase with the flow rate of 0.3 mL minute⁻¹. The column temperature was maintained at 25°C during analysis and the sample injection volume was 2 µL. The eluted polyphenolic acids were monitored by a PDA detector and the UPLC column effluent pumped directly without any split into the TQD-MS/MS (Waters, USA) system optimized for the polyphenolic acid analysis.

Mobile phase

Solvent - A: 0.1 per cent formic acid in water and Solvent - B: 0.2 per cent formic acid in methanol

Statistical analysis

Completely Randomized Design (CRD) and mean values were compared using Duncan's Multiple Range Test (DMRT).

Results and Discussion

Total seventeen individual polyphenolic acids were fractionated and quantified from the methanolic extracts of fruit pulp of *Annona* spp. by HPLC-MS/MS analysis. It is apparent from mean values that polyphenolic acids *viz.*, *p*-coumaric acid, *o*-coumaric acid, 2,4-dihydroxybenzoic acid, caffeic acid, gentisic acid, protocatechuic acid, *t*-cinnamic acid and ferulic acid were abundant in annona fruits while benzoic acid, *p*-hydroxybenzoic acid, salicylic acid, 3-hydroxybenzoic acid, vanillic acid, gallic acid, ellagic acid, syringic acid and sinapic acid were limited (Figure 1a and 1b). However, chlorogenic acid was not detected. The polyphenolic acid content was significantly different among the annona genotypes and is depicted in Table 2.

There was a significant difference for benzoic acid among the annona genotypes. A range of 3.63 to 23.28 µg g⁻¹FW of benzoic acid was recorded in different genotypes. As evident from the treatment means, maximum benzoic acid was found in *A. muricata* (23.28µg g⁻¹FW) which was followed by *A. reticulata* (13.69µg g⁻¹FW), Balanagar (10.02µg g⁻¹FW), Red Sitaphal (7.62µg g⁻¹FW) and *A. cherimola* (6.79µg g⁻¹FW). The minimum benzoic acid was found in *A. atemoya* (3.63µg g⁻¹FW), *A. glabra* (4.61µg g⁻¹FW) and Arka Sahan (6.09µg g⁻¹FW). The *p*-hydroxybenzoic acid was significantly different among the annona genotypes. *p*-hydroxybenzoic acid ranged from 0.31 to 1.97 µg g⁻¹FW in different genotypes. As evident from treatment means, the maximum *p*-hydroxybenzoic acid was noticed in Arka Sahan (1.97µg g⁻¹FW) which showed statistically (*p*<0.05) par with Red Sitaphal (1.79µg g⁻¹FW), was followed by *A. reticulata* (1.02µg g⁻¹FW), *A. glabra* (0.61 µg g⁻¹FW), *A. muricata* (0.67µg g⁻¹FW) and Balanagar (0.58µg g⁻¹FW). The minimum *p*-hydroxybenzoic acid was observed in *A. atemoya* (0.31µg g⁻¹FW) and *A. cherimola* (0.45µg g⁻¹FW). The salicylic acid was significantly different among the annona genotypes. Salicylic acid ranged from 2.40 to 6.00 µg g⁻¹FW in different genotypes. The maximum

Polyphenolic profiles in edible *Annona* spp.

Table 2. Phenolic acid ($\mu\text{g g}^{-1}$ FW) profiling of the selected edible *Annona* spp. fruits by HPLC-MS/MS

Species / Varieties	Benzoic acid	<i>p</i> -hydroxy benzoic acid	Salicylic acid	3-hydroxy benzoic acid	<i>t</i> -cinnamic acid	2,4-dihydroxy benzoic acid	Gentisic acid	Protocatechuic acid	<i>p</i> -coumaric acid	<i>o</i> -coumaric acid	Vanillic acid	Galic acid	Caffeic acid	Ferulic acid	Syringic acid	Sinapic acid	Ellagic acid	Chlorogenic acid
<i>Annona squamosa</i> a) Balanagar	10.02 ^c	0.58 ^{cd}	2.42 ^d	2.89 ^b	18.29 ^b	12.39 ^{cd}	18.16 ^b	4.23 ^b	115.38 ^{de}	48.11 ^b	4.96 ^{bc}	10.25 ^b	6.31 ^f	5.93 ^d	0.77 ^a	2.16 ^a	0.28 ^{cd}	ND
b) Arka Sahar	6.09 ^{de}	1.97 ^a	4.69 ^b	1.35 ^{cd}	22.68 ^a	16.31 ^{cd}	14.22 ^{bc}	14.33 ^a	90.17 ^{fg}	19.00 ^{de}	3.11 ^{cd}	14.06 ^a	23.88 ^c	6.22 ^d	0.34 ^b	1.11 ^{cd}	0.27 ^{cd}	ND
c) Red Sitaphal	7.62 ^d	1.79 ^a	2.69 ^{cd}	2.92 ^b	17.54 ^b	10.43 ^d	14.84 ^{bc}	4.82 ^b	103.76 ^{ef}	36.23 ^{bc}	4.10 ^{cd}	6.57 ^c	3.43 ^f	5.91 ^d	0.78 ^a	0.87 ^{cd}	0.20 ^d	ND
<i>Annona atemoya</i>	3.63 ^f	0.31 ^d	4.73 ^{ab}	1.32 ^{cd}	14.07 ^c	14.83 ^{cd}	9.77 ^c	15.82 ^a	78.54 ^g	7.91 ^e	2.59 ^{cd}	8.34 ^{bc}	10.07 ^e	7.24 ^d	0.75 ^a	0.86 ^{cd}	0.27 ^{cd}	ND
<i>Annona reticulata</i>	13.69 ^b	1.02 ^b	6.00 ^a	6.06 ^a	10.20 ^d	39.49 ^a	17.38 ^b	6.99 ^b	172.65 ^c	48.64 ^b	3.71 ^{cd}	15.88 ^a	25.59 ^{bc}	6.13 ^d	0.38 ^b	0.79 ^d	0.40 ^c	ND
<i>Annona muricata</i>	23.28 ^a	0.67 ^c	5.12 ^{ab}	3.13 ^b	4.47 ^c	27.84 ^b	24.69 ^a	17.04 ^a	321.53 ^a	70.80 ^a	2.16 ^d	6.03 ^c	16.76 ^d	15.94 ^b	0.36 ^b	1.67 ^b	0.69 ^b	ND
<i>Annona cherimola</i>	6.79 ^d	0.45 ^{cd}	2.40 ^d	0.89 ^d	3.93 ^c	18.38 ^c	11.82 ^c	16.47 ^a	260.66 ^b	23.46 ^{cd}	6.87 ^b	6.35 ^c	35.26 ^a	21.78 ^a	0.67 ^a	1.26 ^{bc}	1.12 ^a	ND
<i>Annona glabra</i>	4.61 ^{ef}	0.61 ^c	3.97 ^{bc}	1.81 ^c	3.95 ^c	30.87 ^b	10.46 ^c	16.56 ^a	124.05 ^d	50.31 ^b	19.13 ^a	2.74 ^d	27.78 ^b	12.28 ^c	0.63 ^a	0.99 ^{cd}	0.26 ^{cd}	ND
Mean	9.47	0.92	4.00	2.54	11.89	21.32	15.17	12.03	158.34	38.06	5.83	8.78	18.63	10.18	0.58	1.21	0.42	--
S. Em (\pm)	0.518	0.096	0.448	0.301	1.068	2.228	1.778	1.362	5.485	4.793	0.839	0.777	1.110	0.839	0.055	0.157	0.0560	--
CD (0.05)	1.553	0.289	1.345	0.903	3.203	6.681	5.331	4.085	16.445	14.371	2.515	2.331	3.329	2.518	0.166	0.470	0.167	--

Note: Values represent means of triplicate readings, values with different superscripts in the same column are significantly different at 95% level of confidence based on Duncan's multiple-range test



salicylic acid was found in *A. reticulata* ($6.00\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. muricata* ($5.12\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* ($4.73\mu\text{g g}^{-1}\text{FW}$), Arka Sahan ($4.69\mu\text{g g}^{-1}\text{FW}$) and *A. glabra* ($3.97\mu\text{g g}^{-1}\text{FW}$). The minimum salicylic acid was found in *A. cherimola* ($2.40\mu\text{g g}^{-1}\text{FW}$), Balanagar ($2.42\mu\text{g g}^{-1}\text{FW}$) and Red Sitaphal ($2.69\mu\text{g g}^{-1}\text{FW}$). Annona genotypes exhibited significant difference for 3-hydroxybenzoic acid and it ranged from 0.89 to $6.06\mu\text{g g}^{-1}\text{FW}$ in different genotypes. As evident from treatment means, the maximum 3-hydroxybenzoic acid was found in *A. reticulata* ($6.06\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. muricata* ($3.13\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal ($2.92\mu\text{g g}^{-1}\text{FW}$), Balanagar ($2.89\mu\text{g g}^{-1}\text{FW}$) and *A. glabra* ($1.81\mu\text{g g}^{-1}\text{FW}$). The minimum 3-hydroxybenzoic acid was found in *A. cherimola* ($0.89\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* ($1.32\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan ($1.35\mu\text{g g}^{-1}\text{FW}$). The *t*-cinnamic acid was significantly different among the annona genotypes. A range of 3.93 to $22.68\mu\text{g g}^{-1}\text{FW}$ of *t*-cinnamic acid was recorded in different genotypes. As evident from the treatment means, maximum *t*-cinnamic acid was found in Arka Sahan ($22.68\mu\text{g g}^{-1}\text{FW}$) which was followed by Balanagar ($18.29\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal ($17.54\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* ($14.07\mu\text{g g}^{-1}\text{FW}$) and *A. reticulata* ($10.20\mu\text{g g}^{-1}\text{FW}$). The minimum *t*-cinnamic acid was observed in *A. cherimola* ($3.93\mu\text{g g}^{-1}\text{FW}$), *A. glabra* ($3.95\mu\text{g g}^{-1}\text{FW}$) and *A. muricata* ($4.47\mu\text{g g}^{-1}\text{FW}$). The 2,4-dihydroxybenzoic acid was significantly different among the annona genotypes. A range of 39.49 to $10.43\mu\text{g g}^{-1}\text{FW}$ of 2,4-dihydroxybenzoic acid was recorded in different genotypes. As evident from the treatment means, maximum 2,4-dihydroxybenzoic acid was found in *A. reticulata* ($39.49\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. glabra* ($30.87\mu\text{g g}^{-1}\text{FW}$), *A. muricata* ($27.84\mu\text{g g}^{-1}\text{FW}$), *A. cherimola* ($18.38\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan ($16.31\mu\text{g g}^{-1}\text{FW}$). The minimum 2,4-dihydroxybenzoic acid was noticed in *A. atemoya* ($14.83\mu\text{g g}^{-1}\text{FW}$), Balanagar ($12.39\mu\text{g g}^{-1}\text{FW}$) and Red Sitaphal ($10.43\mu\text{g g}^{-1}\text{FW}$).

Annona genotypes exhibited significant difference for gentisic acid. Gentisic acid ranged from 9.77 to $24.69\mu\text{g g}^{-1}\text{FW}$ in different genotypes. The maximum gentisic acid was found in *A. muricata* ($24.69\mu\text{g g}^{-1}\text{FW}$) which was followed by Balanagar ($18.16\mu\text{g g}^{-1}\text{FW}$), *A. reticulata* ($17.38\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal ($14.84\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan

($14.22\mu\text{g g}^{-1}\text{FW}$). The minimum gentisic acid was found in *A. atemoya* ($9.77\mu\text{g g}^{-1}\text{FW}$), *A. glabra* ($10.46\mu\text{g g}^{-1}\text{FW}$) and *A. cherimola* ($11.82\mu\text{g g}^{-1}\text{FW}$). The protocatechuic acid was significantly different among the annona genotypes. A range of 4.23 to $17.04\mu\text{g g}^{-1}\text{FW}$ of protocatechuic acid was recorded in different genotypes. As evident from the treatment means, maximum protocatechuic acid was found in *A. muricata* ($17.04\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. glabra* ($16.56\mu\text{g g}^{-1}\text{FW}$), *A. cherimola* ($16.47\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* ($15.82\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan ($14.33\mu\text{g g}^{-1}\text{FW}$). The minimum protocatechuic acid was observed in Balanagar ($4.23\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal ($4.82\mu\text{g g}^{-1}\text{FW}$) and *A. reticulata* ($6.99\mu\text{g g}^{-1}\text{FW}$). There was a significant difference for *p*-coumaric acid among the annona genotypes. A range of 78.54 to $321.53\mu\text{g g}^{-1}\text{FW}$ of *p*-coumaric acid was recorded in different genotypes. The maximum *p*-coumaric acid was found in *A. muricata* ($321.53\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. cherimola* ($260.66\mu\text{g g}^{-1}\text{FW}$), *A. reticulata* ($172.65\mu\text{g g}^{-1}\text{FW}$), *A. glabra* ($124.05\mu\text{g g}^{-1}\text{FW}$) and Balanagar ($115.38\mu\text{g g}^{-1}\text{FW}$). The minimum *p*-coumaric acid was found in *A. atemoya* ($78.54\mu\text{g g}^{-1}\text{FW}$), Arka Sahan ($78.54\mu\text{g g}^{-1}\text{FW}$) and Red Sitaphal ($103.76\mu\text{g g}^{-1}\text{FW}$).

The *o*-coumaric acid was significantly different among the annona genotypes. A range of 7.91 to $70.80\mu\text{g g}^{-1}\text{FW}$ of *o*-coumaric acid was recorded in different genotypes. The maximum *o*-coumaric acid was observed in *A. muricata* ($70.80\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. glabra* ($50.31\mu\text{g g}^{-1}\text{FW}$), *A. reticulata* ($48.64\mu\text{g g}^{-1}\text{FW}$), Balanagar ($48.11\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal ($36.23\mu\text{g g}^{-1}\text{FW}$), *A. cherimola* ($23.46\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan ($19.00\mu\text{g g}^{-1}\text{FW}$). The minimum *o*-coumaric acid was found in *A. atemoya* ($7.91\mu\text{g g}^{-1}\text{FW}$). The perusal of data on vanillic acid indicated significant difference among the annona genotypes. A range of 2.16 to $19.13\mu\text{g g}^{-1}\text{FW}$ of vanillic acid was recorded in different genotypes. The maximum vanillic acid was found in *A. glabra* ($19.13\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. cherimola* ($6.87\mu\text{g g}^{-1}\text{FW}$), Balanagar ($4.96\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal ($4.10\mu\text{g g}^{-1}\text{FW}$) and *A. reticulata* ($3.71\mu\text{g g}^{-1}\text{FW}$). The minimum vanillic acid was noticed in *A. muricata* ($2.16\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* ($2.59\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan ($3.11\mu\text{g g}^{-1}\text{FW}$). There was a significant difference obtained for gallic acid among the annona genotypes and it ranged from

Polyphenolic profiles in edible *Annona* spp.

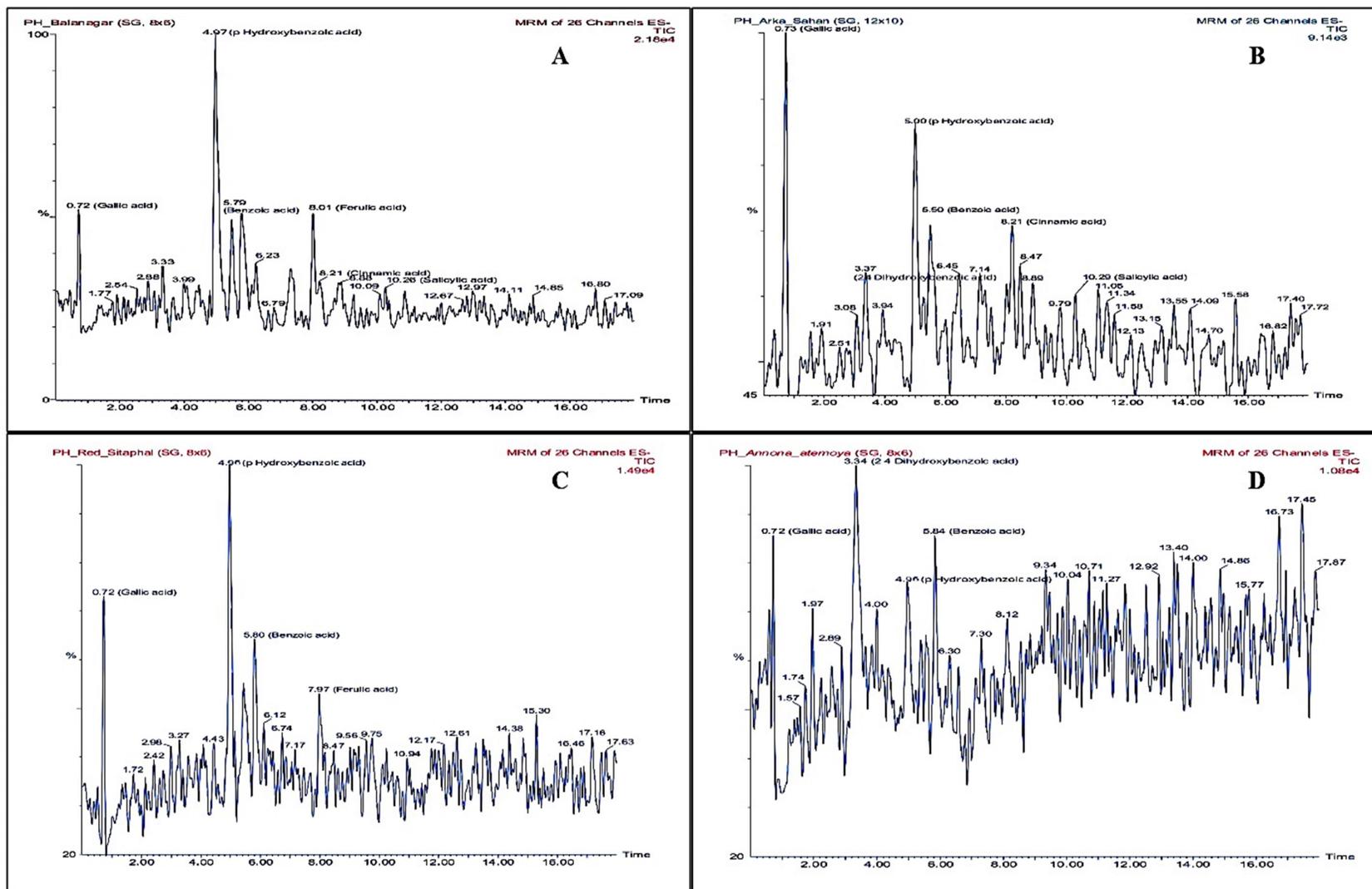


Figure 1a. Chromatograms of phenolic acid profiling in *A. squamosa* [Balanagar (A), Arka Sahan (B), Red Sitaphal (C)] and *A. atemoya* (D)



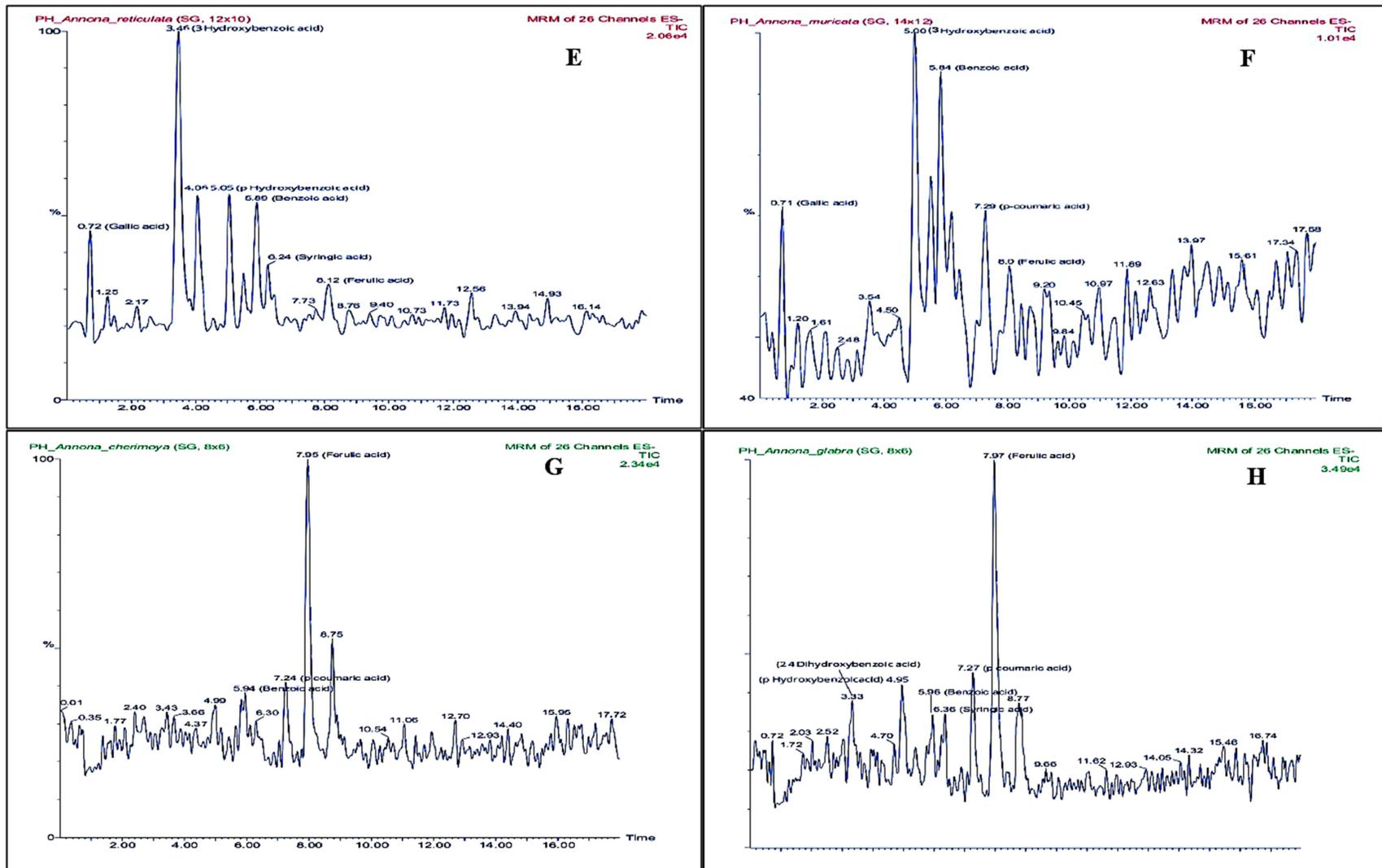


Figure 1b. Chromatograms of phenolic acid profiling in *A. reticulata* (E), *A. muricata* (F), *A. cherimoya* (G) and *A. glabra* (H).



2.74 to 15.88 $\mu\text{g g}^{-1}\text{FW}$. As evident from the treatment means, maximum gallic acid was observed in *A. reticulata* (15.88 $\mu\text{g g}^{-1}\text{FW}$) which was *on par* with Arka Sahan (14.06 $\mu\text{g g}^{-1}\text{FW}$) and was followed by Balanagar (10.25 $\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* (8.34 $\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal (6.57 $\mu\text{g g}^{-1}\text{FW}$) and *A. cherimola* (6.35 $\mu\text{g g}^{-1}\text{FW}$). The minimum gallic acid was found in *A. glabra* (2.74 $\mu\text{g g}^{-1}\text{FW}$) and *A. muricata* (6.03 $\mu\text{g g}^{-1}\text{FW}$).

The caffeic acid differed significantly among the annona genotypes and it ranged from 3.43 to 35.26 $\mu\text{g g}^{-1}\text{FW}$. The maximum caffeic acid was found in *A. cherimola* (35.26 $\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. glabra* (27.78 $\mu\text{g g}^{-1}\text{FW}$), *A. reticulata* (25.59 $\mu\text{g g}^{-1}\text{FW}$), Arka Sahan (23.88 $\mu\text{g g}^{-1}\text{FW}$) and *A. muricata* (16.76 $\mu\text{g g}^{-1}\text{FW}$). The minimum caffeic acid was found in *A. atemoya* (10.07 $\mu\text{g g}^{-1}\text{FW}$), Balanagar (6.31 $\mu\text{g g}^{-1}\text{FW}$) and Red Sitaphal (3.43 $\mu\text{g g}^{-1}\text{FW}$). There was a significant difference for ferulic acid among the annona genotypes and it ranged from 5.91 to 21.78 $\mu\text{g g}^{-1}\text{FW}$. The maximum ferulic acid was found in *A. cherimola* (21.78 $\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. muricata* (15.94 $\mu\text{g g}^{-1}\text{FW}$), *A. glabra* (12.28 $\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* (7.24 $\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan (6.22 $\mu\text{g g}^{-1}\text{FW}$). The minimum ferulic acid was observed in Red Sitaphal (5.91 $\mu\text{g g}^{-1}\text{FW}$), Balanagar (5.93 $\mu\text{g g}^{-1}\text{FW}$) and *A. reticulata* (6.13 $\mu\text{g g}^{-1}\text{FW}$). The perusal of data on syringic acid indicated significant difference among the annona genotypes and it ranged from 0.78 to 0.34 $\mu\text{g g}^{-1}\text{FW}$. As evident from treatment means, the maximum syringic acid was noticed in Red Sitaphal (0.78 $\mu\text{g g}^{-1}\text{FW}$) which was statistically ($p < 0.05$) *on par* with Balanagar (0.77 $\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* (0.75 $\mu\text{g g}^{-1}\text{FW}$), *A. cherimola* (0.67 $\mu\text{g g}^{-1}\text{FW}$) and *A. glabra* (0.63 $\mu\text{g g}^{-1}\text{FW}$) and was followed by Arka Sahan (0.34 $\mu\text{g g}^{-1}\text{FW}$), *A. muricata* (0.36 $\mu\text{g g}^{-1}\text{FW}$) and *A. reticulata* (0.38 $\mu\text{g g}^{-1}\text{FW}$). The sinapic acid differed significantly among the annona genotypes and it ranged from 0.87 to 2.16 $\mu\text{g g}^{-1}\text{FW}$. The maximum sinapic acid was found in Balanagar (2.16 $\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. muricata* (1.67 $\mu\text{g g}^{-1}\text{FW}$), *A. cherimola* (1.26 $\mu\text{g g}^{-1}\text{FW}$) and Arka Sahan (1.11 $\mu\text{g g}^{-1}\text{FW}$). The minimum sinapic acid was found in *A. reticulata* (0.79 $\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* (0.86 $\mu\text{g g}^{-1}\text{FW}$), Red Sitaphal (0.87 $\mu\text{g g}^{-1}\text{FW}$) and *A. glabra* (0.99 $\mu\text{g g}^{-1}\text{FW}$). There was a significant difference for ellagic acid among the annona genotypes and it ranged

from 0.27 to 1.12 $\mu\text{g g}^{-1}\text{FW}$. The maximum ellagic acid was found in *A. cherimola* (1.12 $\mu\text{g g}^{-1}\text{FW}$) which was followed by *A. muricata* (0.69 $\mu\text{g g}^{-1}\text{FW}$), *A. reticulata* (0.40 $\mu\text{g g}^{-1}\text{FW}$), Balanagar (0.28 $\mu\text{g g}^{-1}\text{FW}$), Arka Sahan (0.27 $\mu\text{g g}^{-1}\text{FW}$), *A. atemoya* (0.27 $\mu\text{g g}^{-1}\text{FW}$) and *A. glabra* (0.26 $\mu\text{g g}^{-1}\text{FW}$). The minimum ellagic acid was observed in Red Sitaphal (0.20 $\mu\text{g g}^{-1}\text{FW}$). Polyphenolic compounds are the secondary metabolites which are widely found in plant and plant derived foods. Several analytical methods are available for detection of the polyphenolics. Most of the times, these polyphenolic compounds are analyzed by High Performance Liquid Chromatography (HPLC) (Giusti *et al.*, 1999; Vagiri *et al.*, 2012), coupled with diode array detector and mass spectrometer (Revilla *et al.*, 1999). In the present investigation, seventeen polyphenolic compounds were identified by comparing their retention times and mass spectra with respective standards. The perusal of mean values indicated that polyphenolic acids *viz.*, *p*-coumaric acid, *o*-coumaric acid, 2,4-dihydroxybenzoic acid, caffeic acid, gentisic acid, protocatechuic acid, *t*-cinnamic acid and ferulic acid were abundant in annona fruits. However, chlorogenic acid was not identified in any of the methanolic extracts of fruits. The custard apple juice was found to contain polyphenolic acids like caffeic, ferulic, *p*-coumaric and sinapic acid in free form (Lee *et al.*, 2003). Cinnamic acid and *p*-coumaric acid were prevalent polyphenolic compounds in *A. muricata* pulp (Jimenez *et al.*, 2014). *p*-coumaric acid, caffeic acid, protocatechuic acid, gentisic acid and gallic acid were reported in custard apple wine (Jagtap and Bapat, 2015). It is clear from the HPLC-MS/MS analysis that individual polyphenolics were positively correlated with total polyphenolic contents of annona fruits. *A. muricata* was registered with elevated level of polyphenolic compounds, thus have wide health benefits due to their antioxidant characteristics. Reactive oxygen species (ROS) generated during metabolic processes in the human body are eliminated by enzymatic and non-enzymatic antioxidant systems that exist in the body. However, antioxidant system abnormalities, or excess generation of ROS due to various physical and chemical factors, induce oxidative stress, which causes tissue damage and gene mutation, resulting in various chronic diseases, such as diabetes,

Alzheimer's disease, and aging (Seifried *et al.*, 2007).

Conclusion

From the above study it may be concluded that research on the development of functional materials from edible natural resources with antioxidant effects is a need of the hour. In the present study, *Annona muricata* L. has been reported with highest concentration of polyphenolic compounds among the eight *Annona* spp. fruits. Total eighteen individual compounds were studied in the fruits, among which the *p*-coumaric acid, *o*-coumaric acid, 2,4-dihydroxybenzoic acid, caffeic acid, gentisic acid, protocatechuic acid, *t*-cinnamic acid and

ferulic acid. The results obtained in this study will help further the understanding of the polyphenolic composition of *Annona* fruits and the roles of these compounds in health-promoting physiological functions by nullifying reactive oxygen species (ROS).

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