

COMPARATIVE PHYTOCHEMICAL ANALYSIS OF *PHLOGACANTHUS THYRSIFLORUS* NEES: IMPLICATIONS ON ATTENUATION OF PRO-OXIDANTS AND PATHOGEN VIRULENCE IN *CAENORHABDITIS ELEGANS* MODEL SYSTEM

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ABSTRACT

Objectives: *Phlogacanthus thyrsoiflorus* Nees of Acanthaceae family is endogenous to sub-tropical Himalayas. It has been reported to be used traditionally in Jaintia tribe of Meghalaya, India, for the treatment of many ailments. The aim was to detect the active compounds present in the leaves and evaluation of its *in vitro* free radicals scavenging potential. Leaves protective effects *in vivo* will be investigated using *Caenorhabditis elegans* model system utilizing wild type and mutant strains and the phenomena of host-pathogen interactions.

Methods: Gas chromatography-mass spectrometry was used for detection of different compounds present. The versatility of leaf extracts to scavenge different free radicals generated *in vitro* was assessed with different *in vitro* methods. Survival analysis of wild type and mutant strains *C. elegans* under enhanced pro-oxidants exposure was investigated *in vivo*. The fast killing assay was also performed to study the extracts modulatory activities on host *C. elegans* survival under pathogen *Pseudomonas aeruginosa* infection.

Results: About 40 compounds were detected in methanolic fraction of the extract with variable percentages. Both aqueous and methanol extract possessed remarkable, versatile free radical scavenging activity irrespective of the types of free radical generated. The *in vivo* experiments are in compliance with observable increased survival ability percentage of *C. elegans* under intense exogenous oxidative stress and pathogen infection.

Conclusion: Our findings enlightened the different bioactive compounds present with versatility of *P. thyrsoiflorus* in tackling different free radicals generated both *in vitro* and *in vivo* that highly support for its candidature as a good antioxidant source. Our findings may justify the historical relevance of this plant in herbal remedies that could form the basis for inquiry of new active principles.

Keywords: Free radicals, Oxidative stress, *Caenorhabditis elegans*, *Phlogacanthus thyrsoiflorus*, Phytochemicals.

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INTRODUCTION

Cells are inherently equipped with an impressive repertoire of integrated antioxidant and immune systems for a sustained protection of an organism. However, with an unavoidable consequence of aerobic metabolism, a stepwise unwanted reduction of molecular oxygen (O_2) under exposure or electron-transfer reactions leads to production of the highly reactive O_2 species (ROS) and subsequently reactive nitrogen species [1,2]. When the level of ROS exceeds the antioxidative defense mechanisms, a cell is said to be in a state of "oxidative stress." Under this state, an overwhelmed ROS reacts and propagates with atoms of diverse biomolecules to modulate their functions. At the molecular levels, deregulations with an unwanted inhibition and activation of proteins, mutagenesis of DNA and gene transcription disrupts cell homeostasis to enhanced cell death and/or proliferation. Oxidative stress has been associated with the cause and etiologies of over hundreds of human diseases such as cancer, cardiovascular disease, aging, and neurodegenerative diseases [3,4].

Compounds which can maintain the level of ROS and modulate integrity of the immune system are important players in the amelioration and prevention of various diseases [5]. Antioxidants are diverse compounds pivotal in maintaining different cell types in a reduced environment. Thus, antioxidant ultimately helps in preserving the integrity and adequate function of membrane lipids, cellular proteins, and nucleic acids in signal transductions and gene expressions [6]. It is noteworthy that the traditional system of medicine has been practiced since historical times for curing ailments. Compounds isolated and

derived from medicinal plants and other natural sources remain an important source of new drugs, new chemical entities and new drug leads [7-9].

Phlogacanthus thyrsoiflorus Nees of Acanthaceae family is endogenous to sub-tropical Himalayas, upper Gangetic plains, Bihar, North Bengal, and North Eastern regions of India in the continent of Asia. The plant is an evergreen shrub up to 2.4 m high, branchlets quadrangular with leaves up to 13-35 cm long. *P. thyrsoiflorus*, has been reported to be used traditionally for the treatment of many ailments [10,11]. Jaintia tribe of Meghalaya uses this plant for the historically significant antihyperglycemic, hypolipidemic, hepatoprotective activity they believed it possessed [12-14]. Considering its traditional medicinal values, this study is an attempt to evaluate on the different bioactive principles present, its antioxidative potentials both *in vitro* and *in vivo* using the genetically tractable model *Caenorhabditis elegans* which additionally can provide us with a useful tool to study plant potential on modulation of host immune system and the mechanism of host-pathogen interactions.

METHODS**Reagents and chemicals**

All chemicals and reagents used were of analytical grade. They were obtained from the following companies: Sigma Chemical Company, St. Louis, USA; Thermo Fisher Scientific Pvt. Ltd. Mumbai, India; Santa Cruz Biotech; Qualigens and Himedia Laboratories, India. Miscellaneous were purchased locally.

Nematode strains

The *C. elegans* strains used includes the wild type 'Bristol N2', which was kindly supplied by National Centre for Biological Sciences (NCBS), Bengaluru, India. Mutant strains which include GA480 [SOD-2 (gk257) I, SOD-3 (tm760) X.] and TJ1052 [age-1 (hx546) II] strains were obtained from *Caenorhabditis* Genetics Center (CGC), University of Minnesota, St. Paul, MN, USA.

Plant material

Healthy disease free leaves of *P. thyrsoiflorus* were collected from Dawki, West Jaintia Hills District, Meghalaya, India, and the submitted herbarium was identified by Dr. N. Parthasarathy, Professor, Department of Ecology and Environmental Sciences, Pondicherry University, India (Voucher no KSU101#16). The collected sample was thoroughly washed and shade dried for 20-25 days, followed by pulverized with an electrical blender to obtain a powder form of the dried leaves. The powdered samples were stored in an air tight container at room temperature without any exposure to sunlight for further use.

Preparation of plant extracts

Plant material was extracted using water (aqueous) and methanol solvents (1:10 w/v). For aqueous extract, powdered leaves were extracted by macerated in water for 24 hrs in a shaker. Then, the solution was first filtered through muslin cloth and Whatman filter paper No. 1, the filtrate thus obtained was concentrated and dried at 40°C using rotavapor. For methanol extract (ME), the dried leaf powdered was placed in the thimble of Soxhlet apparatus using 500 ml of methanol as a solvent. The extraction was continued until the clear solvent was seen in the thimble. Finally, the solvent was concentrated in vacuum under reduced pressure using a rotary flask evaporator, allowed to evaporate until the dry powdered was obtained. All the collected, dried powdered samples were stored at 4°C for further use, and the percentage yield of the extract was calculated using the following formula

$$\text{Percentage yield} = \frac{\text{Final weight of the dried extract}}{\text{Initial weight of the powder}} \times 100$$

Qualitative and quantitative phytochemical analysis

Preliminary qualitative phytochemical analysis of extracts was performed as per the standard protocol [15]. Quantitatively, the total phenolic content of leaves was determined by Folin-Ciocalteu method [16]. The total alkaloid and flavonoid contents were determined according to methods described previously [17,18]. The mean of four readings repeated thrice was used and the total phenolic content was expressed in mg of gallic acid equivalents (GAE)/100 g dry weight of extract and flavonoids mg of quercetin equivalents (QE)/g compound.

Gas chromatography-mass spectrometry (GC-MS) analysis of the extract

GC-MS analysis was performed using JEOL GCMATE II GC-MS spectrometer. This instrument was set to an initial temperature of 110°C, and maintained at this temperature for 2 minutes. At the end of this period, the oven temperature was risen to 280°C, at the rate of 5°C/minute, and maintained for 9 minutes. The injection port temperature was ensured as 250°C and helium flow rate as 1 ml/minute. The ionization voltage was 70 eV. The samples were injected in split mode as 10:1. Mass spectral scan range was set at 45-450 (m/z). Using computer searches on a NIST Ver. 2.1 MS data library, comparing of the spectrum obtained through GC-MS compounds present in the plants sample were identified.

In vitro antioxidant assay

2, 2-diphenyl-1-picrylhydrazyl (DPPH) radical scavenging was determined by following standard protocol [19]. Hydrogen peroxide scavenging activity of plant extract was assessed by method as described [20]. Nitric oxide was generated from sodium nitroprusside and measured by Gries's reaction [21]. The chelating of ferrous ions

by extracts was estimated by the method of Dinis [22], wherein the Fe²⁺ chelating ability was monitored by the absorbance of the ferrous iron-ferrozine complex at 562 nm. The total antioxidant capacity of the extracts was evaluated by phosphomolybdenum method [23]. Different concentrations of AE and ME were used and compared with ascorbic acid as positive control in all the assays.

In vitro DNA damage protection assay

Plasmid DNA nicking assay was performed using pUC 18 plasmid DNA as per the method described previously [24]. Briefly, 10 µl of AE and ME (final concentration of 0.25 µg/µl and 0.5 µg/µl) kept separately in Eppendorf tubes was mixed with plasmid DNA and incubated at room temperature followed by the addition of 10µl of Fenton's reagent (30 mM H₂O₂, 50 µM ascorbic acid and 80 µM FeCl₃). The final volume of the mixture was made up to 20 µl and incubated for additional 30 minutes at 37°C. Finally, 20 µl of the sample was loaded in each well of 1.5% agarose gel, electrophoresed and the bands were visualized and photograph using Biorad gel documentation system.

In vivo screening on C. elegans

Synchronization and culture conditions of C. elegans

Worms GA480 [SOD-2 (gk257) I, SOD-3 (tm760) X.] and TJ1052 [age-1 (hx546) II] strains and wild type N2 were grown at 20°C on NGM medium spread lawn, fed with *Escherichia coli* OP50 strain (henceforth termed OP50) as food and maintained at 20°C. Synchronized culture of gravid worms was obtained by lysis in an alkaline hypochlorite solution. The eggs were allowed to hatch in NGM plates at 20°C and young L1 were grown on NGM plates seeded with OP50.

Oxidative stress resistance assay

Pilot experiments were performed to ascertain the maximum time and dose used before the experiment [25]. The assay was performed in 96 well microtiter plates with 150 µl of media/well in triplicates per condition (P10 worms/well) and the same experiment was repeated thrice at different times independently. Briefly, L3-L4 worms were transferred and allowed to grow for 30 hrs in liquid S-medium with OP50 containing dissolved AE and ME at 1 mg/ml concentration. Control wells consist of worms that were incubated without the extract. After 30 hrs, worms were transferred to a freshly prepared liquid S-medium (supplemented with 200 µM 5-Fluoro-2'-deoxyuridine; FuDR). 10 mM tert-butyl hydroperoxide was added to the medium to induce oxidative stress at zero time and the survival rate of the worms was evaluated every 4 hrs in a time-dependent manner. A worm was scored as dead when it did not respond to light mechanical stimulus.

Fast killing assay

Pilot experiments were performed to ascertain the different time and doses used in the experiment. Fast killing assay plates were prepared using peptone, glucose, and sorbitol agar containing 1% bacto-peptone, 1% glucose, 1% NaCl, 150 mM sorbitol, and 1.7% bacto-agar [26]. To the plates, 50 µl of the overnight grown culture of virulence strain PA14, *Pseudomonas aeruginosa* (PA) was added to the plates and allowed to grow for 24 hrs at 37°C, followed by transferring to 23°C for 24 hrs. Later, the solid plates were seeded with a synchronized population of L3-L4 worms previously incubated for 24 hrs in liquid S-medium containing OP50 with dissolved AE and ME (final concentration of 0.5 mg/ml and 1 mg/ml). Control plates consist of worms that were incubated without the extract. All the assays were performed in triplicate containing 20 worms/plate and the same experiment was repeated thrice independently. Finally, the plates were scored for dead worms every 8 hrs. A worm was considered dead and removed if it failed to move after gentle touch with a platinum wire worm picker.

Statistical analysis

All values reported are mean±SD of four to five replicates in each group except for survival analysis. Data obtained from different sets were analyzed using Student's t-test for paired data. The level of significance

($p < 0.05$) was considered to be statistically significant and reported as percentage or absorbance. The Kaplan–Meier analysis method was used for processing and plotting survival data against time [27].

RESULTS

Recovery percent, quantitative analysis, and total antioxidant capacity

The extract of *P. thyrsoiflorus* was prepared as described and the percent yield was found to be 12.6% for AE and 15% for ME. Quantitative analysis results showed that total phenolic content present in AE and ME at 18 and 21 mg GAE/g of phenolic compounds. The total flavonoid content of AE and ME was found to be 9 and 31.8 mg QE/g of flavonoids. Finally, the total alkaloid content present for AE is approximately 5.5 and ME at 16.5 mg/g of extract. Preliminary qualitative phytochemical analysis of extracts showed the presence of metabolites such as carbohydrates, proteins, flavonoids, phenols, steroids, terpenoids, cardiac glycosides, tannins, quinones, and saponins (Fig. 1).

GC-MS analysis

To elucidate the chemical profile of the plant, we performed the GC-MS analysis of ME. Our result revealed the presence of 40 phytochemical constituent compounds that could contribute to the medicinal quality

of the plant. The major bioactive principles with their retention time (RT), molecular formula, molecular weight, and peak area in percentage are presented in Fig. 2. The first compound identified with less RT (10.545) was phenol 2, 6-dimethoxy whereas 2-ethylacridine was the last compound which took longest RT (33.839 minutes). The GC-MS analysis of extract revealed the abundance of phytol with the highest percentage followed by hexadecanoic acid, methyl ester, stigmasterol, allooromadendrene, squalene, and 1, 8-diamino-3, 6-dioxaoctane compounds. The detailed reports of some of the major compounds present are listed in Table 1.

In vitro antioxidant assays

DPPH and hydroxyl radical scavenging assay

Percentage of DPPH scavenging activity of AE and ME increases with increased in extract concentration. In comparison with a 96.70% inhibition of ascorbic acid, AE and ME exhibited an inhibitory percentage of 42.95% and 72.13% at 300 $\mu\text{g/ml}$, respectively (Fig. 3a). From Fig. 3b it can be inferred that the $\text{OH}\cdot$ radical scavenging activity of AE and ME also increased with increases in extract concentration. AE with 68.9% inhibitory activity possessed a comparatively better $\text{OH}\cdot$ scavenging activity than ME with 61.2% at 500 $\mu\text{g/ml}$.

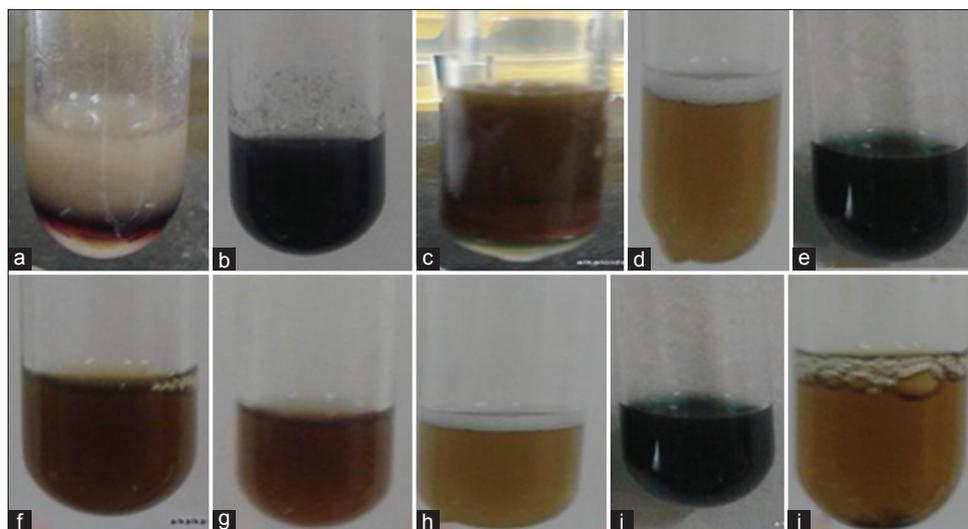


Fig. 1: Preliminary qualitative phytochemical analysis of aqueous extract showed the presence (a) carbohydrates (b) proteins (c) flavonoids (d) phenols, (e) steroids (f) terpenoids (g) cardiac glycosides (h) tannins (i) quinones (j) Saponins

Table 1: Major compounds identified in the methanol extract of *P. thyrsoiflorus* Nees by GC-MS

RT	Compound name	Formula	Mol wt.	PA%
10.545	Phenol, 2,6-dimethoxy-	$\text{C}_8\text{H}_{10}\text{O}_3$	154.1	2.32
13.917	Dispiro [2.2.2.2] deca-4,9-diene	$\text{C}_{10}\text{H}_{12}$	132	3.19
	N, N-dimethyltrimethylsilamine	$\text{C}_5\text{H}_{15}\text{NSi}$	117	3.19
14.393	1, 8-diamino-3,6-dioxaoctane	$\text{C}_6\text{H}_{16}\text{N}_2\text{O}_2$	148	2.24
15.629	Benzoic acid, 4-hydroxy-3,5-dimethoxy-, hydrazide	$\text{C}_9\text{H}_{12}\text{N}_2\text{O}_4$	212	2.21
15.930	Orcinol	$\text{C}_7\text{H}_8\text{O}_2$	124	1.84
16.190	2,6,6-trimethyl-bicyclo[3.1.1]heptane, trans	$\text{C}_{10}\text{H}_{18}$	138	1.36
17.118	Hexadecanoic acid, methyl ester	$\text{C}_{17}\text{H}_{34}\text{O}_2$	270	5.68
18.805	9-Octadecenoic acid (Z)-, methyl ester	$\text{C}_{19}\text{H}_{36}\text{O}_2$	296.4	1.69
18.902	Phytol	$\text{C}_{20}\text{H}_{40}\text{O}$	296.5	8.55
19.043	Methyl stearate	$\text{C}_{19}\text{H}_{38}\text{O}_2$	298.5	0.92
21.026	Cyclopropane, isothiocyanate-	$\text{C}_4\text{H}_5\text{NS}$	99.1	0.67
22.482	7H-Purin-6-amine, 7-methyl-	$\text{C}_6\text{H}_7\text{N}_5$	149.1	1.78
24.525	Squalene	$\text{C}_{30}\text{H}_{50}$	410.7	2.61
25.127	Allooromadendrene	$\text{C}_{15}\text{H}_{24}$	204.3	3.02
26.583	Stigmasterol	$\text{C}_{29}\text{H}_{48}\text{O}$	412.69	4.11
29.577	5-methyl-2-phenyl-1H-indole	$\text{C}_{12}\text{H}_{13}\text{N}$	207.2	0.96
33.989	2-Ethylacridine	$\text{C}_{15}\text{H}_{13}\text{N}$	207.2	0.96

RT: Retention time, Mol wt: Molecular weight, PA%: Peak area percentage, *P. thyrsoiflorus*: *Phlogacanthus thyrsoiflorus*

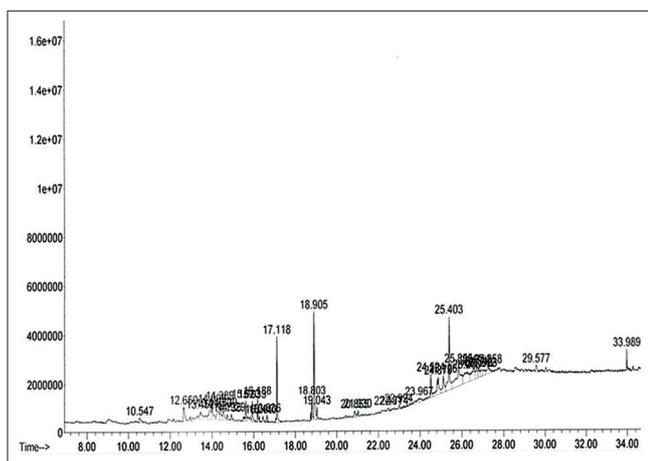


Fig. 2: Gas chromatography-mass spectrometry chromatogram of methanolic extract of *Phlogacanthus thyriflorus*

H₂O₂ and nitric oxide radical scavenging activity

The percentage of H_2O_2 scavenging activity of AE and ME was found to be 87.5% and 90.7%, at par with that of ascorbic acid which exhibited a 91.3% inhibitory percentage at 200 $\mu\text{g/ml}$ (Fig. 3c). Fig. 3d depicts the scavenging activity of extracts for nitric oxide radical. The percentage inhibition was in the order of ascorbic acid>ME>AE with a corresponding inhibitory percentage of 65.6%, 60%, and 41.5%, respectively.

Total antioxidant capacity and ferrous (Fe^{2+}) ion chelating assay

The total antioxidant capacity of the *P. thyriflorus* leaf extracts was determined by phosphomolybdate method and the increased absorbance at 695 nm indicated higher total antioxidant capacity. Our results depicted uniform increased in absorbance from 100 to 500 $\mu\text{g/ml}$ in all the three samples studied, with total antioxidant capacity in the order of ascorbic acid>ME>AE (Fig. 3e). Fig. 3f represented the ferrous ions (Fe^{2+}) chelating activities of *P. thyriflorus* extracts which were found to be comparatively higher than ascorbic acid at 200 $\mu\text{g/ml}$. This percentage chelation activity was in the order of ME>AE>ascorbic acid corresponding with percentage of 64.3%, 62.7%, and 58.5%, respectively.

In vitro DNA damage protective assay

Hydroxyl ($\cdot\text{OH}$) radicals generated by the Fenton's reaction are known to induced breaks in DNA strands. Fig. 4 of agarose gel migration image showed that when plasmid DNA was dissolved in Fenton's reagent, electrophoretically two visible bands were observed due to the formation of open circular DNA (OC-Form) in lane 2 besides the intact supercoil DNA (SC-Form) in lane 1. Addition of extracts resulted in the partial or full recovery of super coiled DNA (Lane 3-6). In comparison with AE, ME loaded in lane 5 and 6 exhibited maximum recovery of SC-Form irrespective of the concentrations used (Fig. 4).

In vivo screening on *C. elegans*

Oxidative stress resistance assay

The effect of tert-butyl hydroperoxide (10 mM) treatment on the survival time of untreated wild type control and mutated *C. elegans* strains was investigated until all the worms were dead (zero survival). Our results showed that TJ1052 mutants survive the longest with 44 hrs survival time, followed by N2 wild type with 32 hrs (Fig. 5a and c). Oxidative stress sensitive strain GA480 has the shortest survival time with zero survival corresponding at 24 hrs (Fig. 5b). Taking into account the time of zero survival in individual strain as mentioned, treatment with AE and ME showed increased survival percentage in all the strains studied. Our data revealed that N2 worms previously treated with AE and ME exhibited a 90.47% and 88.37% survival percentage at 32 hrs whereas, GA480 strain exhibited a 75.0% and 71.45% survival percentage at

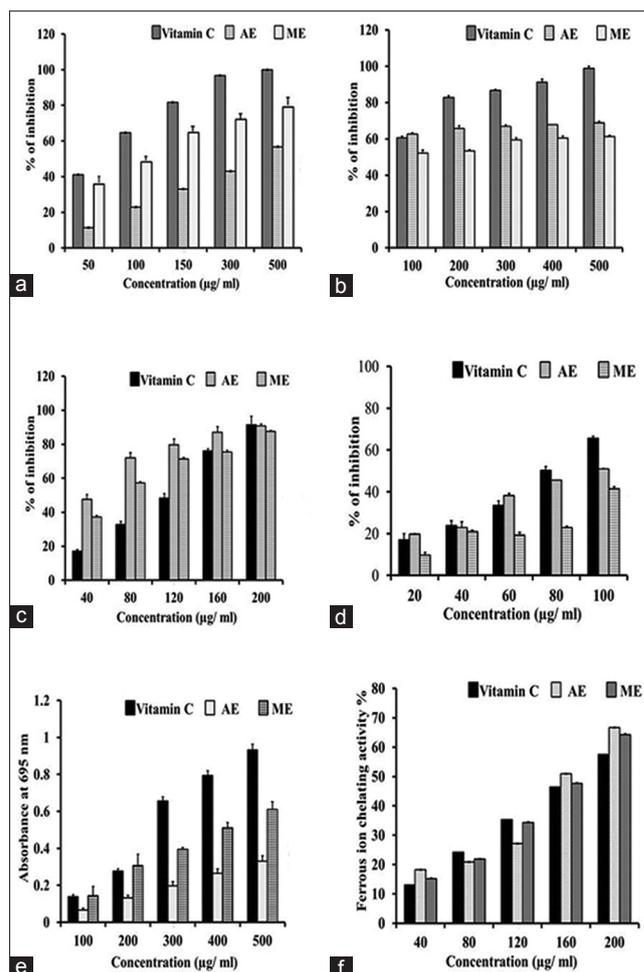


Fig. 3: *In vitro* antioxidant activities of AE and ME: (a) DPPH radical scavenging activity, (b) hydroxyl radical scavenging activity, (c) hydrogen peroxide scavenging activity, (d) nitric oxide radical scavenging activity, (e) total antioxidant activity, (f) ferrous ion (Fe^{2+}) chelating activity

24 hrs (Fig. 5a and b). Furthermore, TJ1052 treated with AE and ME showed a survival percentage of 65.7% and 62.85% at 44 hrs (Fig. 5c). In addition, preliminary assays were done before this study and no observable effect of the added plant extracts on worm survival, lifespan or brood size on OP50 (data not shown), indicating a lack of toxicity of the compounds present.

Fast Killing assay

C. elegans is commonly used as an infection model for pathogenesis studies in *P. aeruginosa*. The standard virulence assays rely on the slow and fast killing or paralysis of nematodes. The ability of plant extract to protect *C. elegans* from PA infection under fast killing assay was demonstrated in Fig. 6. After 72 hrs, the survival of the AE treated nematode was found to be 48% and 63%, respectively, at 0.5 mg/ml and 74% at 1 mg/ml (Fig. 6a). Similarly, ME showed a similar protective ability with the worms survival percentage of 62% at 0.5 mg/ml and 74% at 1 mg/ml (Fig. 6b). In addition, both AE and ME exhibited no inhibition sign on PA growth and/or perturbation of *C. elegans* feeding rate. This directly indicated that the observable effect of extracts on the survival of host worms is by modulation of toxins generated and/or enhance its various defences against the pathogen (Fig. 7a, b and c).

DISCUSSION

Our *in vitro* experiments indicated that both AE and ME possessed remarkable, versatile, free radicals scavenging activities irrespective

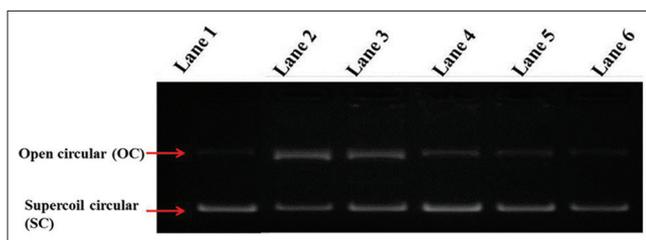


Fig. 4: DNA protective assay using pUC18 plasmid. Representative lane includes: Lane 1: DNA (pUC18 plasmid alone), Lane 2: DNA + Fenton reagent, Lane 3: DNA + Fenton reagent + 0.25 µg/µl AE, Lane 4: DNA + Fenton reagent + 0.50 µg/µl AE, Lane 5: DNA + Fenton reagent + 0.25 µg/µl ME, and Lane 6: DNA + Fenton reagent + 0.50 µg/µl ME

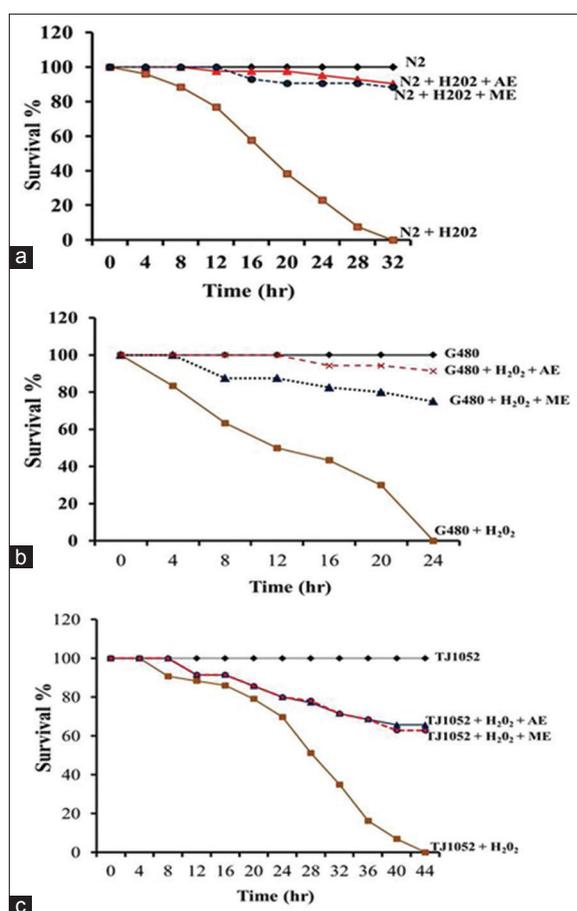


Fig. 5: *In vivo* effects of aqueous and methanol leaf extract against tert-butyl hydroperoxide induced oxidative stress on the survival of (a) N2 wild type (b) G480 and (c) TJ1052 mutant strains. Results are presented as percentage in comparison with respective control group

of free radical generated. This overall significantly higher inhibition percentage can be positively correlates with the total phenolic, flavonoid, and alkaloid content present and the abundance of active and lead compounds as depicted in GC-MS analysis. By its capability to reduce a stable free radical DPPH, extracts provide us with a first clue that there were compounds present to donate an electron or hydrogen to DPPH, can react with it and thereby bleach the DPPH absorption. The significant OH[•] radical scavenging activity observed provides an additional and direct evidences that protection from the most common and single unpaired electron encountered in aerobic organism is possible thereby, its reactivity with all biological molecules

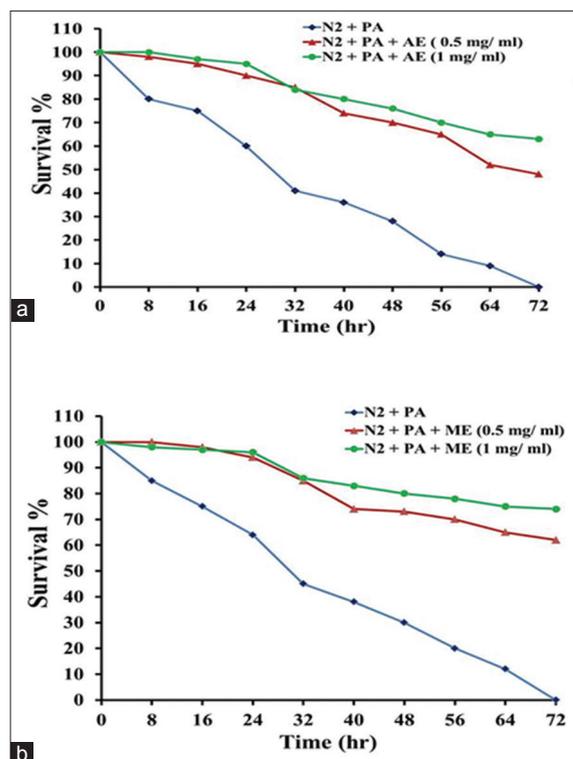


Fig. 6: Survival of N2 wild type *Caenorhabditis elegans* under *Pseudomonas aeruginosa* fast killing assay. (a) Represents worms treated previously with aqueous and (b) worms treated previously with methanol extract. Results are presented as percentage in comparison with respective control group

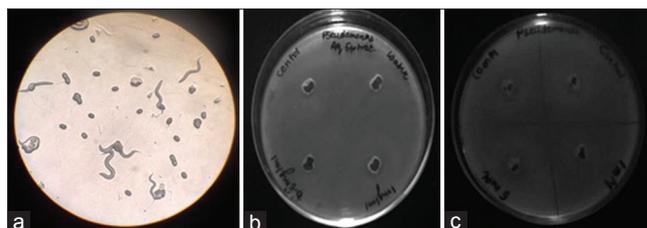


Fig. 7: (a) Population of *Caenorhabditis elegans* L-3 and L-4 worms with eggs. Disc diffusion method for *Pseudomonas aeruginosa* growth inhibition in the presence of (b) aqueous extract, and (c) methanol extract at 0.5 and 1 mg/ml concentration. No visible inhibition zones were observed

and subsequent declined in cellular damages such as lipid peroxidation, protein damage, and membrane destruction can be ameliorated [5,6]. In addition, although not so common, NO[•] radical toxicity increases greatly when it reacts with superoxide radical, forming the highly reactive peroxynitrite anion (ONOO⁻), an effective chain terminating agent in free radical mediated lipid oxidation with well-established role in carcinoma and inflammatory processes [28]. Thus, by scavenging NO[•] radical, compounds present in extracts helps in breaking of the chain reactions with an ultimate reduced toxicity that could play a role in protecting the physicochemical properties of membrane bilayers from free-radical induced severe cellular dysfunction. Finally, by having a ferrous ions (Fe²⁺) chelation activity, extracts renders an important effect via. Inhibiting metal-catalyzed oxidation reaction, halting directly at the main source of ROS production as applicable within animal and human systems [29].

At the nucleic acids level, oxidative modification of DNA has been suggested to contribute to aging and various diseases including

cancer and chronic inflammation. Our results clearly showed that both AE and ME rescued pUC18 plasmid DNA against damage caused by hydroxyl radical. The DNA cleavage analysis provides evidence that compounds present in *P. thyrsoflorus* extracts can directly or indirectly have a role in mutual antagonism relevance with genotoxicity and mutagenicity. It is also interesting to note that recent studies suggest that phytol, the most abundant compound present in this plant can indeed reduce both production of free radicals besides scavenging activity, a link directly correlated to its unique structural feature of branched-chain unsaturated alcohol [30,31]. Similarly, phytosterols such as stigmasterol, squalene, and isothiocyanates detected are reported to be responsible, at least in part, for preventive effects on the development of diseases due to ROS against lipid peroxidation [32-34]. With this study limitations and the rich active principles present in a single plant observed, we suggests that the synergism like effect of different compounds present provide a combinational advantages of synergy in efficacy thereby, cumulative with pronounced versatility against different free radicals and nuclei acid protection were observed.

The role of oxidative stress in *C. elegans* lifespan has been investigated in several different ways. Correlations between life span and various aspects of oxidative metabolism have been examined together; with studies typically either examines age changes in wild type nematodes or differences between wild type nematodes and mutants with altered rates of aging. Our studies conceptualized on the idea that if ROS can accelerate and modulate survival and aging, then manipulating ROS levels should affect mortality and survival rate. Superoxide dismutase (SOD), the first line of antioxidant enzyme acts by converting superoxide to hydrogen peroxide, which can subsequently be converted to water by catalase or peroxiredoxin. In this study, we found that GA480 (mutants for SOD-2 and SOD-3) worms were indeed sensitive to induced oxidative stress by displaying the shortest survival time and/or zero survival within 24 hrs after treatment. We speculated that although under normoxic conditions, additional SOD's, chemical and enzymatic antioxidants may be useful for normal organismal defenses against oxidative damage, but under conditions of extreme oxidative burst like the prevailing experimentally induced elevation, the compensation for the loss of crucial function due to mutation can be easily overwhelmed and this may contribute to the shortest survival as seen in the case of G480 in comparison to N2 and TJ1052 with normal and overexpress SODs.

Furthermore, our findings clearly showed that groups treated with extracts exhibited significantly higher survival percentage in all the strains of *C. elegans* studied. We suggest that during this period of rapid oxidative burst, different compounds present as mentioned earlier can individually and/or synergistically modulate the loads of free radicals generated by donating an electron for reducing and converting them into stable unreactive metabolites. These benefits are thus manifested manifold either directly in the form of less macromolecular oxidative damage, besides safeguarding the normal functioning of active integrated antioxidant systems from the overwhelming effects of oxidative stress. In totality, the abundance of different antioxidative compounds as seen in the extracts can give a kind of compensation required for the altered SOD of G480 besides an enhanced additional protection to N2 and TJ1052 strains with an intact system for their better survival.

It is also noteworthy to mention that there are numerous aging genes that have been found to cause at least a 20% increase in the lifespan of *C. elegans*. TJ1052, an age-1 mutants are one such example with long life, normal fertility, and stress tolerant. Recent findings have revealed that long-lived age-1 mutants have increased SOD and catalase activity levels, which led to their resistance to oxidative stress in comparison to the wild type [35,36]. Other studies conducted by Honda *et al.* have shown that age-1 mutation leads to oxidative stress resistance phenotype in *C. elegans* by regulating through the insulin signaling pathway [37]. With an unclear common mechanism till date, our present findings of TJ1052 worms exceptional resistant to oxidative and electrophilic stresses,

with prolonged survival time will further contribute substantially to the notion that indeed *age-1* mutation strain has some favorable alterations in molecular expression/signaling events corresponding to the beneficial oxidative metabolism which contributed significantly to its longer lifespan. Furthermore, our findings also lend support to the hypothesis that oxidative stresses comprise key contributors to the process of aging [38], and that longevity depends at least in part on the balance between the generation of such stresses and resistance to them [39,40].

In fast killing assay, *C. elegans* death occurred within a period of hours and it was reported that the damage was caused due to the phenazines, low molecular-weight diffusible toxins excreted by PA [41,42]. Our results clearly demonstrated the ability of *P. thyrsoflorus* extracts in rescuing *C. elegans* from fatal PA infection. This allows us to speculate that besides enhancing the integrated antioxidative system of the host, compounds present in the plant can lead to the chemical modification of toxic phenazines to a lesser toxic intermediates thereby providing, with an ultimate protective effects to the host. Recently, forward genetic approach to identify components of a presumptive *C. elegans* innate immune response pathway upstream of induced defense responses has led to the discovery that a *C. elegans* homolog of the mammalian p38 mitogen-activated protein kinase is an important component that is upregulated in *C. elegans* defense response to bacterial pathogens [43-45]. Natural extracts or compounds present in plants are also reported to have immunomodulatory activities [46,47]. Alternatively, the presence of compounds in extracts capable of modulating the immune system of the host to fight against the penetration of invading pathogen cannot be ruled out and this speculation needs further evaluation.

CONCLUSION

It is worth mentioning that the data obtained from this study enlightened the different versatility of *P. thyrsoflorus* in tackling different free radicals generated both *in vitro* and *in vivo* in *C. elegans* that highly support for its candidature as a good antioxidant source. This work also showed the presence of many pharmacologically bioactive principles, which needs to be elucidated for their potential modulation on different molecules and signaling involved. This may help to develop a herbal medicinal product or an effective therapeutic agent involving immune modulation and strategy based biomedical usefulness in controlling different oxidative stress and age-related pathophysiological conditions.

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