

Comparative Adaptability Assessment of Two Mangroves from Indian Sundarbans: Some Biochemical Appearances

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Abstract

Comparative adaptability against salinity was assessed between the two well-known mangroves (*Avicennia marina* and *Heritiera fomes*) from Indian Sundarbans *in vitro*. Occurrence of *H. fomes* is intermittent in and around of this mangrove swamp. A harmony has to maintain between ROS production and efficient scavenging of ROS by the plant itself for sustainability. In the present work, extent of salt tolerance was evaluated by mainly two ways: i) accumulation of free amino acids in the cytoplasm for proficient osmotic adjustment and ii) promoting elevated amount of antioxidants (both enzymes and secondary metabolites) with respect to substrate salinity. Occurrence of free amino acids (Alanine, Leucine and Proline) in *A. marina* are well correlated ($p \le 0.01$) with the increasing salinity and *H. fomes* (Alanine and Phenyl Alanine) correlation value showed $p \le 0.05$. ROS scavenging reflected through ABTS, DPPH and Fe^{2+} chelating activity and results indicating that *A. marina* have some advantage over the other investigated taxa. Amount of phenols and flavonoids also designated the same. Additional number of isoforms of two antioxidant enzymes (peroxidase and super oxide dismutase) occurred in *A. marina* as the salinity enhanced, but in case of *H. fomes*, which was lacking. The experimental results might be designated towards the comfortable adaptability to *A. marina*, rather to *H. fomes*.

Keywords

ABTS, DPPH, Free Amino Acids, Mangroves, Phenols

1. Introduction

Mangroves, being a productive and protective halophilic group of plant community dwell in along the tropical

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How to cite this paper: Dasgupta, N., Chowdhury, P. and Das, S. (2015) Comparative Adaptability Assessment of Two Mangroves from Indian Sundarbans: Some Biochemical Appearances. *Natural Science*, **7**, 519-534. http://dx.doi.org/10.4236/ns.2015.712053 and subtropical estuaries of the world. Being a protective buffer against the frequency and ferocity of episodic cyclones, hurricanes, Tsunami, storms and floods; the significance of mangrove restoration has claimed a priority research. According to FAO report [1], 198,000 km² of mangroves in 1980, and 157,630 km² in 1990 presently represents only 146,530 km² across the globe. Development and industrialization have triggered the existence crisis of this important habitat [2] [3]. Simultaneously, relative sea level rise due to global climatic changes also has great impact on mangrove vegetation [4]. Duke *et al.* [5] opined that the decline of the mangrove forest occur at a faster rate than inland tropical forest and coral reef and postulated that 30% - 40% of coastal wet lands and 100% of the mangrove forest could be lost in the next 100 years. Hence, an attentive responsiveness demands towards the potentiality of mangrove restoration programme.

Sundarbans, the largest single block of mangrove delta, includes two countries, India (West Bengal) and Bangladesh. Due to geomorphic features, an upliftment in the north-western part (India) and subsidence in the east (Bangladesh) has conquer an elevated salinity in the western part (India) which imparts a major impact on mangrove species distribution [6]. A cumulative effects of damming in the lower Gangetic plane, industrial pollution, geotectonic characteristics and unplanned reformation of coastal area have pave the way of massive siltation on the river bed which in turn, caused reduction of fresh water influx through the river system and its tributaries to the sea leading to hike salinity in the Indian Sundarbans [7]. These instigate disastrous (sometimes complete elimination) for some important plant species (such as *Aegialitisrotundifolia*, *Heritierafomes*, *Nypafruticans*, *Xylocarpusgranatum* and *X. mekongensis*) from the swamps [8]-[10]. *Heritiera fomes* prefers less than 5ppt salinity [11]. In India, this species is rapidly deteriorating (only 6% of 100 sampling sites) [12]. *H. fomes* is considered as red listed category taxa [13]. Although the species diversity is highest in this region, the conservation and sustainable management presently need serious attention. The most important criteria of mangroves, is their NaCl tolerance at least up to seawater level (~500 mM NaCl) [14] [15]. Moreover, mangroves are considered as potential models for studying the salt tolerance mechanisms in plants [16]. Hence, investigations on adaptive potential in mangroves might be in formative towards salt management.

Different mangrove species displays change in osmoregulation, which can be validated with excess accumulation of compatible solutes, like sugar, free amino acids and low molecular weight proteins [17] [18]. Dual effects of osmotic adjustment for cellular homoeostasis involve: uptake of inorganic ions and synthesis of compatible organic solutes within the cell sap [19]-[22]. To avoid ionic toxicity, absorbed inorganic ions are sequestered into the vacuole (especially Na⁺ and Cl⁻) [23]-[25]. Occurrence of excess inorganic ions in the vacuole, the cytoplasm has to confront bi-phasic stresses: internal stress caused by the accumulated ions and external stress [26]. To minimise this conflict, plants accumulate *de novo* some organic compatible solutes like amino acids (e.g. proline, alanine, tyrosine etc.), soluble sugars (e.g. sucrose), polyols (e.g. mannitol), and betaines (e.g. glycine betaine) to adjust the osmotic potential, and moreover these organic solutes do not interfere with normal metabolic processes [14] [27]-[29]. However, the degree of salt tolerance can be attributed to the accumulation of compatible solutes in the cell sap [30].

Considerable scientific evidences suggested that under oxidative stress, reactive oxygen species (ROS) such as peroxyl, superoxide and hydroxyl radicals are generated and the ratio between oxidation and anti-oxidation is supposed to be a critical concept for maintaining a biological homeostasis [31]. Consequently, research on anti-oxidant principles of plants has been fast-tracked and halophytes have been recognized for marked antioxidant potential activities [32]. Polyphenols are antioxidants, having redox properties, act as reducing agents, hydrogen donators and singlet oxygen quenchers. A number of secondary metabolites were traced from mangroves time to time and have been recognized as pivotal bioactivities towards efficient slat management [33]-[35]. Despite of innumerable bioactivities, polyphenols perhaps are mostly attributed to antioxidant property [36]-[38]. Some essential phenols and flavonoids were traced from several mangroves [39]-[41].

Polyphenols in plants are well established as a defensive force against detrimental impact of abiotic stress which leads to generate free radicals [42]-[44]. The commonly determination methods of antioxidant activity are ABTS [2, 2'-azino-bis (3-ethylbenzothiazoline-6-sulphonicacid)] and DPPH (1, 1-diphenyl-2-picrylhydrazyl) assay. Both of them have intense reproducible characteristics under certain assay conditions. ABTS⁺ is dissolved in aqueous and organic media, where in the antioxidant activity can be measured [45]. In contrast, DPPH has been used extensively as a free radical evaluating reducing substances [46]. DPPH is only soluble in organic media, especially in ethanol, this have a limitation while deducing the role of hydrophilic antioxidants.

Removal of H_2O_2 is vital for cellular DNA homeostasis [47]. The antioxidant capacity of compounds has been

attributed to various factors such as prevention of chain reaction, chelating metals, radical scavenging ability and reductive capacity [48] [49]. Chelation of pro-oxidant metals is one of the most important mechanisms of action of secondary antioxidants. Iron and other transition metals (cobalt, copper, chromium, vanadium, arsenic, cadmiumand nickel) act as catalysts of free radical reactions promote oxidation. The transition metal ion, Fe²⁺ hold the capacity to shift single electrons by virtue of which it can allow the formation and proliferation of many radical reactions [50]. The main strategy to avoid ROS generation involves chelating of the metal ions that is associated with redox active metal catalysis.

Salt tolerance of plant is mainly attained by four mechanisms: 1) osmotic adjustment of the cytoplasm as a result of accumulation of compatible solutes; 2) exclusion of salt from the cell across the plasma membrane using ion transporters [51]-[53]; 3) accumulation of salt in the vacuoles using tonoplast transporters [54]-[57]; and 4) triggering on the elevated production of antioxidants (both enzymes and secondary metabolites) for scavenging reactive oxygen species [58]-[62]. A compulsory consequence of aerobic metabolism is generation of reactive oxygen species (ROS) which embrace free radicals such as superoxide anion (O2 -), hydroxylradical (•OH), as well as non-radical molecules like hydrogen peroxide (H₂O₂), singlet oxygen (¹O₂). Salt tolerance in plant primarily relies on minimizing the water loss through stomata and metabolic shifting through toxic ion homeostasis and osmotic adjustment. On the other hand, it has been established that salt tolerance can be correlated with enhanced ROS scavenging ability [63]-[65]. Enzymes like Superoxide dismutase (SOD), Peroxidase (PRX), Catalase (CAT) are potentially involved in H₂O₂ metabolism leading to photo protection. An increased accumulation of PRX and SOD were reported in water logging stress in Kandeliacandel and Bruguieragymnorrhiza [66]. Antioxidative enzymes in relation to salt tolerance in different plants have been assessed much, but remain focused because of enhance accumulation of these antioxidant enzymes not only concomitant with salt tolerance, but also with salt sensitivity too [67]. Experimental evidence revealed that, elevated synthesis of antioxidative enzymes and increased number of their isoforms can be attributed in selection of salt resistant [68]. In view of the above, this work aims to understand the extent of salt tolerance of two mangroves (one of which is presently in alarming existence in Indian Sundarbans and rest one shows predominant occurrence in the same regime) by assessing the leaf water potential, some major free amino acids accumulation and regulation of two antioxidant enzymes (peroxidase and superoxide dismutase) across in different salinity gradients (in vitro experiments). These findings might be enlightening towards the probable reasons of their extinction and priority areas of preservation strategy.

2. Materials and Methods

Healthy seeds/fruits of two true mangroves (*Avicennia marina* and *Heritiera fomes*) were collected (**Figure 1**) from Indian Sundarbans (of which, *H. fomes* is now in stress condition and the other one, *Avicennia marina* grown luxuriantly in the same habitat and were considered as natural control) and allowed to germinate in polythene bag in mesophytic soil. All the seedlings were grown up to a certain maturity with fresh water treatment for 30 days and thus maintained equable soil environments and age of the plants. After 30 days, the seedlings were treated with different concentrations of saline water (100, 200, 300 and 400 mM of NaCl), leaving one set of experimental pot (to be used as control). These salinity treatments (watering in every alternate day) were continued for another 30 days. At 60 days of maturity, young leaves were collected from three sets of seedlings from each treatment for further experiments.

Free amino acids: Six free amino acids (*viz.* Aspartic acid, Alanine, Leucine, Phenyl-Alanine, Proline and Tryptophan) were assayed from three taxa grown in different salinity gradients. Two grams of freeze-dried leaves were homogenized in acetone and the extracts were centrifuged at 6000 rpm for 10 min. Supernatants were subjected to thin layer chromatography [69] and spots were allowed to run in running buffer (mixture of 4:1:1 n-butanol, acetic acid and distilled water, v/v) up to an equal distances. In the chromatogram, greyish purple, bluish purple and yellow spots appeared for different amino acids, which were identified from their respective standard R_{mf} values [69]. These spots were eluted by 80% ethanol and optical densities were measured in specific wavelengths (550 nm for proline and 400 nm for the other amino acids). Estimations were done based on standard curves plotted for individual amino acid. In case of tryptophan and tyrosine, where the R_{mf} values were very close, confirmatory tests (Hopkins-Cole test for tryptophan and Millon's reaction for tyrosine) were performed following [69]. The value for each amino acid is the average of nine observations, *i.e.* three replica from each plant and three plants of each species.

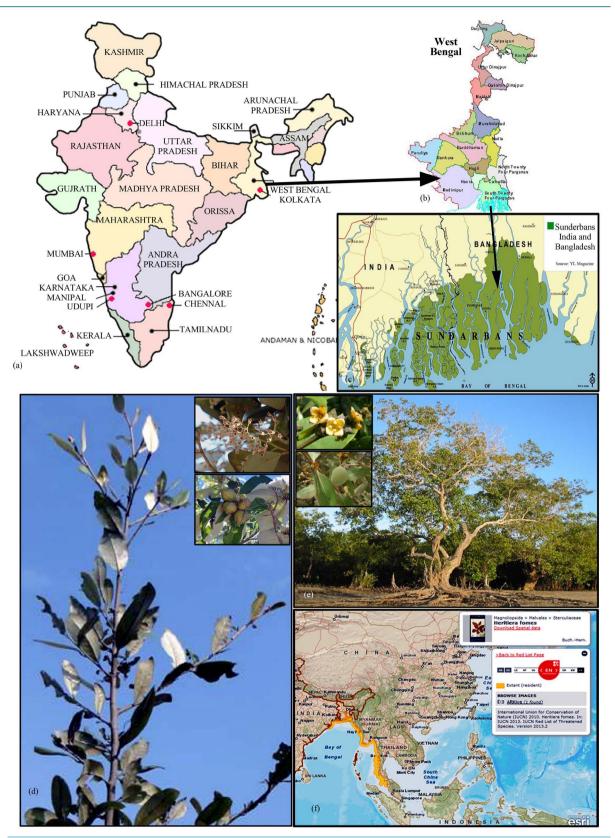


Figure 1. (a)-(c) Geographical Location of the study area; (d)-(e) Mature plants with flowering and fruiting twigs; (d) *Heritierafomes*; (e) *Avicennia marina*; (f) World distribution map of *H. fomes* according to ICUN.

2.1. Antioxidant Secondary Metabolites

Preparation of Extraction

Methanolic extracts were prepared by maceration of 10 gm of mature leaves (from the two taxa, each treatment) with 250 ml of 95% ethanol, at room temperature for 24 hours and then filtered. The procedure was repeated thrice. The extracts were filtered and centrifuged (4500 rpm, 15 min). The solvent was evaporated on rotary evaporator (Neocool Circulator, CF 300, Yamato). The concentrated methanolic leaf extracts were kept frozen (-20°C) until further use [70].

Total Phenols: Folin-Ciocalteu method [71] was followed to determine the total phenol contents in the extracts. 0.1 ml of extractant in methanol containing 0.05 - 0.3 mg/ml conc. of the leaf extract was mixed with 2 ml Folin-Ciocalteu reagent (previously diluted with water at 1:10 v/v) and 1.6 ml (7.5%) of Sodium carbonate (Na₂CO₃). The tubes were mixed and allowed to stand at room temperature for 30 min for color development. Spectrophotometer readings of absorbance were taken at 765 nm using. Gallic acid in the same concentration as the sample was used as positive control. The total phenolic content was expressed as Gallic acid equivalents (GAE) in milligram per gram of dry material using the calibration curve, where X was the absorbance and Y was the Gallic acid equivalent (mg·g⁻¹).

Total Flavonoids: Estimation of total flavonoids in the plant extracts were carried out following the method of Jia *et al.* [72]. In 0.1 ml of extractant in methanol containing 0.2 - 1.2 mg/ml conc. of the leaf extract, 1.2 ml distilled water, 0.120 ml of 5% Sodium nitrite (NaNO₂) were added and mixed vigorously. Keep the mixture at 25° C temperature for 5 mins. After that 0.120 ml of 10° AlCl₃ solution was added and mixed thoroughly. Then the tubes are allowed to stand at room temperature for 5 minutes. Then 0.8 ml of 1 mM Sodium hydroxide (NaOH) solution and 1.16 ml distilled water were added. The absorbances were measured at 510 nm. Quercetin in the same concentration as the sample was used as positive control. Total flavonoids content was calculated as Quercetin (mg·g⁻¹) using the calibration curve, where X was the absorbance and Y was the Quercetin equivalent (mg·g⁻¹).

2.2. ROS Scavenging Ability Assay

DPPH• quenching assay: The free radical scavenging activity of different plant extracts were determined using the stable radical DPPH (1,1-diphenyl-2-picrylhydrazyl) following Blois [73]. A freshly prepared DPPH solution (25 mg·L⁻¹) in methanol was prepared and 3.9 ml of this solution was mixed with 0.1 ml of extract in methanol containing 0.05 - 0.3 mg/ml conc. of the extract. 30 min later, the absorbances were measured at 517 nm using Spectrophotometer (Helios γ , Thermo Electron Corporation). Butylated Hydroxy Toluene (BHT) in the same concentration as the sample was used as positive control. The capability to scavenge the DPPH radical was calculated using the following equation:

DPPH radical scavenging activity (%) =
$$\left\{ \frac{Ac - At}{Ac} \right\} \times 100$$

where Ac is the absorbance of the blank reaction and At is the absorbance in presence of the sample of the extracts.

ABTS⁺ scavenging assay: The method of Re *et al.* [74] was adopted to determine ABTS [2, 2'-azino-bis (3-ethylbenzothiazoline-6-sulphonic acid)] radical scavenging assay. The stock solutions included 7 mM ABTS solution and 2.4 mM Potassium Per sulfate solution. The working solution was then prepared by mixing the two stock solutions in equal quantities and allowing them to react for 12 hour at room temperature in the dark. The solution was then diluted by mixing 1 ml ABTS solution with 60 ml methanol. Fresh ABTS solution was prepared for each assay. Plant extracts (0.1 ml) in methanol containing 0.005 - 0.03 mg/ml conc. of the extract were allowed to react with 3 ml of the ABTS solution and the absorbances were taken at 734 nm after 7 min using Spectrophotometer. Butylated Hydroxy Toluene (BHT) in the same concentration as the sample was used as positive control. The ABTS scavenging capacity of the extract was calculated as:

ABTS radical scavenging activity (%) =
$$\left\{ \frac{Ac - At}{Ac} \right\} \times 100$$

where Ac is the absorbance of the blank reaction and At is the absorbance in presence of the sample of the

extracts.

Fe²⁺ **chelation activity**: The method described by Haro-Vicente *et al.* [75] was followed for assessing the chelating capacity of the plant extracts for ferrous ion. The reaction was carried out in 20 mM HEPES buffer, pH 7.2. Different aliquots of the plant extract, ranging from 0.05 - 0.3 mg/ml were added to 12.5 mM ferrous sulfate solution and the reaction was initiated by the addition of ferrozine (75 mM). The reaction mixture was shaken vigorously which followed incubation for 20 min at room temperature. The absorbances were measured at 562 nm. EDTA in the same concentration as the sample was used as positive control. The Fe²⁺ Chelation capacity of the extract was calculated as:

Fe²⁺ Chelationability (%) =
$$\left\{ \frac{Ac - At}{Ac} \right\} \times 100$$

where Ac is the absorbance of the blank reaction and At is the absorbance in presence of the sample of the extracts.

In all scavenging/chelating ability are expressed in terms of IC_{50} . It is defined as the concentration of the plant extract that's needed to scavenge/chelate 50% of the scavengers/chelating ions present were calculated by the following equation:

$$IC_{50} = \left\{ \frac{Perccent inhibition}{Concentration of the Sample} \right\} \times 50$$

2.3. Estimation of Enzymes: Gel Electrophoresis

Extraction of enzymes: Two grams of young leaf buds were macerated to powder with liquid Nitrogen in a mortar-pestle and then added with 0.1 gm PVP and 5 ml of extraction buffer (1 M Sucrose, 0.2 M Tris-HCl and 0.056 M β -Marcaptoethanol; pH was adjusted at 8.5 after volume make up by double distilled water). The homogenates were centrifuged at 1500 rpm for 20 min at 4°C and use the supernatants in PAGE analysis.

PAGE analysis: Equimolar amounts of proteins (leaf extracts) were loaded in each well. Different isoforms of enzymes were separated by native gel electrophoresis. Gels were stained for definite enzyme following Das and Mukherjee [76]. Gels were documented with a Gel-Doc System (Biostep GmbH – Germany) and analysis for band intensity and Relative Mobility Factor (R_{mf}) were performed with Kodak-MI software. This experiment was repeated five times with the same molecular weight markers. In each case, the number of isoforms and respective OD values were obtained at more or less same R_{mf} .

2.4. Enzyme Assay

Peroxidase (PRX, E.C.1.11.1.7): From 200 mg fresh leaf, extraction was prepared in 1 - 1.5 ml of 0.9% KCl and centrifuged at 12,000 rpm for 15 min at 4°C; the supernatants used as enzyme sample. Absorbance's were taken spectrophotometrically (Helios γ , Thermo electron Corporation, USA) at 460 nm with respect to the standard curve prepared following Shannon *et al.* [77] with minute modification.

Superoxide dismutase (**SOD**, **E.C.1.15.1.1**): Cell sap extracted by macerating 200 mg of leaf in 1 - 1.5 ml of 50 mM Phosphate buffer, pH adjusted to 7.0 and centrifuged at 12,000 rpm for 15 min at 4°C. The supernatants used as enzyme samples. Standard curves were prepared with pure enzyme samples at different aliquots (50, 100, 150, 200, 250 µg·ml⁻¹). Absorbances were measured at 550 nm following the protocol described by Keith *et al.* [78] with slight modification.

The value for each species is the average of nine observations, *i.e.* three replica from each plant and three plants of each species. SPSS 12.0 version was employed for all the statistical calculations.

3. Results

Estimation of free amino acids: Among the six amino acids investigated in leaves of the two species (*A. marina* and *H. fomes*) grown at different NaCl treatment (control, 100, 200, 300 and 400 mM) resulted that not all six amino acids occur in both the species except Aspartic acid (Asp) and Alanine (Ala). Apart from these two, *A. marina* possesses Leucine (Leu) and Proline (Pro) and instead of Leucine and Proline, *H. fomes* have Phenyl-Alanine (Phe) and Tryptophan (Trp). In *Avicennia*, Asp, Ala, Leu and Pro were recorded as 11.37, 3.48, 3.74

and $9.73~\mu g \cdot g^{-1}$ leaf weight respectively in control plants; sharp increment up to 400 mM substratum observed in all the amino acids were detected (19.15, 29.95, 26.36 and 47.39 $\mu g \cdot g^{-1}$ respectively) except Asp, where amount increased was less than the other three. In *Avicennia*, the trend line of proline increment according to salinity rise was most prominent (9.73 in control to 47.39, in 400 mM saline treatment). In *Heritiera*, Asp increment occurred over control up to 200 mM salinity (12.54 to 22.21 $\mu g \cdot g^{-1}$); beyond this, it was decreased. Similarly, Ala, Phe and Trp were observed in the approach of augmentation up to 300 mM NaCl concentration (23.03, 11.96 and 11.93 $\mu g \cdot g^{-1}$ respectively) and drop down at 400 mM (20.63, 9.51 and 7.44 $\mu g \cdot g^{-1}$ respectively) (**Figure 2(a)**).

Incidence of total phenols: Estimated total phenol (TP) in different salinity level of the two studied mangroves is much higher over the control one. It is expressed as Gallic acid equivalent (GAE) in all cases. In *Avicennia*, TP is observed as gradual increment according to the gradual increment of substrate salinity (19.28 mg·g⁻¹ in control and 39.41 mg·g⁻¹ in 400 mM saline concentration). But this trend did not occur in *Heritiera*, where the augmentation of TP over control treatment is recorded up to 200 mM of NaCl treatment (12.65 mg·g⁻¹ and 31.54 mg·g⁻¹

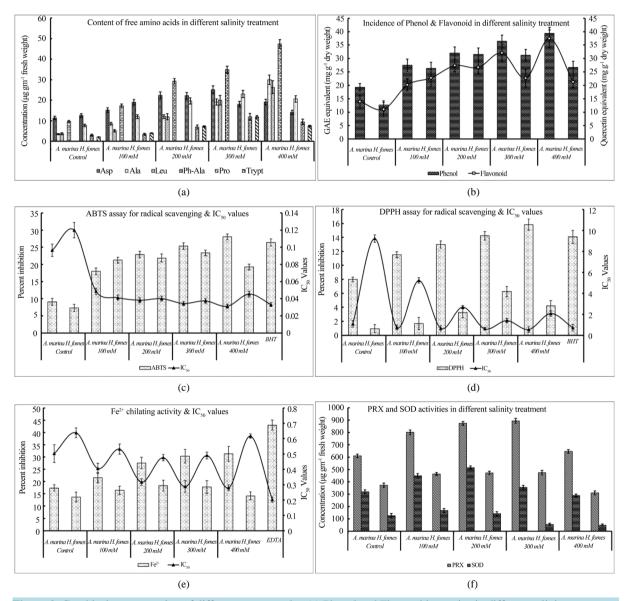


Figure 2. Graphical representation of different assay results. (a) Phenol and Flavonoid contains in different salinity treatment; (b) Amino acids contains in different salinity treatments; (c) ABTS assay and IC_{50} values; (d) DPPH assay and IC_{50} values; (e) Fe^{2+} chelating activity and IC_{50} values; (f) PRX and SOD activities in different salinity treatment.

respectively); the amount was more or less unchanged in 300 mM (31.2 mg·g⁻¹) and instead of increment in 400 mM, it was depleted (26.7 mg·g⁻¹) (**Figure 2(b)**).

Incidence of total flavonoids: The incidence of total flavonoids (TF) was expressed in Quercetin equivalent in the present work. Like TP, total flavonoid was assessed as a considerable increment as salinity increased from 0 to 400 mM of NaCl treatment in *Avicennia* (13.96 mg·g⁻¹ and 37.4 mg·g⁻¹), but the trend is quite dissimilar in case of *Heritiera*, where the escalation recorded up to 200 mM of NaCl treatment over the control one (11.06 and 26.71 mg·g⁻¹ respectively) and it considerably dropped down in 300 and 400 mM of NaCl treatment (22.68 and 21.28 mg·g⁻¹ respectively) (**Figure 2(b)**).

ABTS⁺ scavenging assay: The trend observed in this assay is quite similar to DPPH scavenging assay. The magnitude of percent reduction of ROS increase in case of *A. marina* is tremendous from Control to 100 mM treatment (9.06% - 18.01%) following which the increment is gradual with maximum at 400 mM of NaCl treatment (28.02%). Similarly in case of *H. fomes*, relative increase in value of ABTS activity is evident from Control to 300 mM (7.29% - 23.35%), while a decrease is apparent with 400 mM treatment (19.24%). Also a trace amount of increment is observed from 100 mM to 200 mM (21.25% - 21.85%). Reference BHT value was observed to be 26.36%. Highest IC₅₀ value for *H. fomes*was found at control treatment (0.11%) and the lowest at 300 mM (0.037%) and in *A. marina* maximum at control (0.096%) and lowest in 400 mM NaCl treatment (0.031%) (**Figure 2(c**)).

DPPH assay for radical scavenging: The DPPH assay in different salinity level of the two studied taxaon an average showed a continuous increase in percent reduction of evolved ROS according to the gradual increment of substrate salinity. In *A. marina* the DPPH assay value is maximum in 400 mM NaCl treated plant (15.54%) which is much higher than the control value (7.98%). Whereas in case of *H. fomes*, the trend was dissimilar with an increment from control (0.94%) to 300 mM NaCl treated plant (6.24%) and a decrease at 400 mM NaCl (4.21%) respectively. The value of DPPH activity in case of the positive control (BHT) was found to be 14.08%. IC₅₀ values were calculated in all the cases which reflected the usual reverse diagrams. IC₅₀ is inversely proportional to percent inhibition. Experimental results showed that the maximum IC₅₀ value in *H. fomes* at control treatment (9.21%) and minimum in *A. marina* at 400 mM treatment (0.55%) (**Figure 2(d)**).

Fe⁺² **chelation activity**: It is evident from the experiment that the Fe²⁺ chelation activity in case of *A. marina* is found to be constant hike from control (17.42%) to 400 mM of NaCl treatment (31.29%) but the rate of change is not same in all treatments (*e.g.* increment from 300 mM to 400 mM is. 30.37% - 31.29%). *H. fomes* showed a different form of chelation activity with an increment from Control to 200 mM (13.71% - 18.39%) and a gradual decrease in 300 mM (17.86%) and 400 mM (14.2%) with the lowest value at control (13.71%). The positive control, EDTA showed a value of 43.02% which is higher as compared to *A. marina* and *H. fomes*. IC₅₀ value showed similar trends as DPPH and ABTS with the maximum in *H. fomes* at control (0.63%) and lowest for *A. marina* at 400 mM treatment (0.27%) (**Figure 2(e**)).

3.1. Quantitative Assay of PRX and SOD

Peroxidase: Experimental results showed increase in the amount of PRX enzyme in case of *A. marina* from control to 300 mM ($610.56 - 894.56 \, \mu g \cdot g^{-1}$) NaCl treatment while a depletion occurs at 400 mM treatment ($647.08 \, \mu g \cdot g^{-1}$). The results obtained from *H. fomes*, where maximum value is observed at 300 mM treatment ($474.78 \, \mu g \cdot g^{-1}$) over the control following which the value decreased to 309.86 $\mu g \cdot g^{-1}$ at 400 mM NaCl treatment, though the increment from 200 mM to 300 mM is nominal but in 400 mM saline treated plants giving a value less than control plant (372.08%) (**Figure 2(f)**).

Superoxide Dismutase: Increase in SOD level is evident in case of A. marina from control to 200 mM treatment (317.86 $\mu g \cdot g^{-1}$), where after it dropped down at 300 mM (354.04 $\mu g \cdot g^{-1}$) and 400 mM (289.06 $\mu g \cdot g^{-1}$) NaCl treatment. Unlikely H. fomes showed increase till 100 mM treatment (167.07 $\mu g \cdot g^{-1}$) but when treated with 200 mM NaCl treatment the amount of enzyme depleted to 141.83 $\mu g \cdot g^{-1}$ and lowest at 400 mM (49.07 $\mu g \cdot g^{-1}$) (**Figure 2(f)**).

3.2. PAGE Analysis

Peroxidase: Experimental data shows that *A. marina* expresses comparatively more bands than *H. fomes*, two bands appear in Control treatment with (138.2, 151.74) ODs at their respective R_{mf} values are (0.31, 0.47); in 100 mM treatment of NaCl number of bands increased to three (147.93, 153.00 188.97; R_{mf} 's are 0.31, 0.48 and

0.66 respectively), further an increment in band number is observed in 200 mM and 300 mM NaCl treatment where four (OD 127.0, 131.0, 122.44, 138.6; R_{mf} s are 0.31, 0.42, 0.66, 0.73 respectively) and five (OD 18.4, 26.02, 16.76, 18.96, 53.34; R_{mf} s are 0.09, 0.18, 0.47, 0.59, 0.68 respectively) bands are expressed respectively, where after the count decreased to three at 400 mM NaCl treatment. In *H. fomes*, unlike *A. marina* three bands occur under Control treatment (O.D. 28.53, 2.49, 1.99; R_{mf} values are 0.25, 0.4, 0.57 respectively), which remained same at 100 mM (O.D. 158.45, 128.77, 90.09; R_{mf} 's values are 0.24, 0.44, 0.58 respectively), further at 200 mM treatment of NaCl, bands increased to four (O.D. 72.64, 39.3, 74.1, 82.65; R_{mf} s are 0.24, 0.34, 0.42, 0.6 respectively). A considerable decrease in the band number is noticed at 300 (OD 160,131; R_{mf} 's are 0.06 and 0.31) and 400 mM with two bands per concentration of NaCl treatment (**Figure 3(a)**, **Figure 3(b)**).

Superoxide Dismutase: In case of *A. marina*, plants showed similar band expressions as peroxidase. Plants under control treatment showed two bands at OD 189.0 and 245.0 with respective R_{mf} values are 0.19 and 0.37. An increase in band number is seen when plant is treated with 100 mM NaCl (OD 127.4, 146.6, 168.4; R_{mf} s are 0.17, 0.36 and 0.49 respectively) which continued to increase till 200 mM NaCl (OD 178.0, 186.0, 222.0, and 26.1, R_{mf} s are 0.1, 0.37, 0.48 and 0.53 respectively) and then the band number dropped at 300 mM NaCl (OD 63.0, 126.5, 103.45, 65.31; R_{mf} s are 0.06, 0.2, 0.31 and 0.42) and more at 400 mM NaCl treatment, the number of band depleted (OD 44, 104; R_{mf} s are 0.14, 0.42). SOD graph for *H. fomes* showed three bands under control, 100 mM and 200 mM NaCl treatment (OD 5.43, 6.48, 5.42; R_{mf} s are 0.33, 0.63, 0.75 and 110.51, 53.32, 67.72; R_{mf} s are 0.04, 0.26, 0.54 and 42.98, 162.2, 70.59; R_{mf} s are 0.08, 0.21, 0.34). Bands number dropped down to two at 300 mM and 400 mM NaCl (OD 48.59, 29.58 at 0.54, 0.1 and 48.06, 67.4 at 0.34, 0.58 R_{mf} s) (**Figure 3(c)**, **Figure 3(d)**).

A two tailed bivariate correlation results shows that in *A. marina*, the available free amino acids, studied polyphenols, two antioxidant enzymes and ROS scavenging activity are positively correlated with the salinity imposed during experiments (either $P \le 0.01$ or $P \le 0.05$ level). To the contrary, in *H. fomes*, except leucine and phenyl alanine, all parameters are not correlated with the increment of salinity level (Table 1).

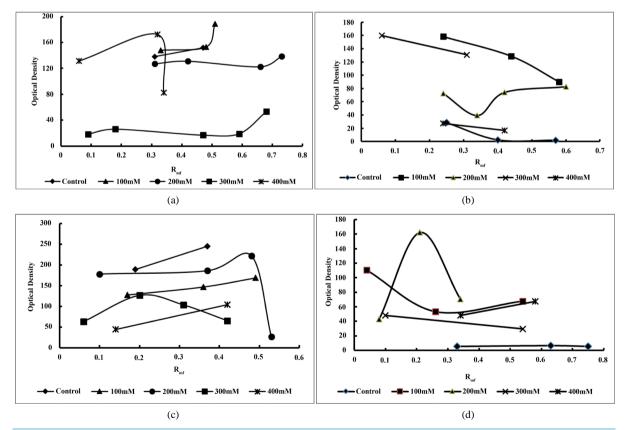


Figure 3. Graphical expression of the results obtained from Gel electrophoresis showing Optical Density and R_{mf} values of two enzymes. (a) *A. marina* (PRX); (b) *H. fomes* (PRX); (c) *A. marina* (SOD); (d) *H. fomes* (SOD).

Table 1. Correlation of different parameters with salini	Table 1.	of different parameters with salinity.
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Parameter	Avicennia marina	Heritiera fomes	Parameter
PRX	0.202	0.241	
SOD	0.256	0.790	
Asp	0.731	0.088	
Ala	0.976^{**}	0.902^{*}	
Leu	0.981**	-	
PhAla	-	0.880^*	
Pro	0.994**	-	Salinity
Trypt	-	0.783	
Phenol	0.982**	0.678	
Flavonoid	0.997**	0.548	
ABTS	0.959^{*}	0.633	
DPPH	0.974**	0.835	
Fe^{2+}	0.969*	0.178	

^{**}Correlation is significant at the $P \le 0.01$ (2-tailed). *Correlation is significant at the $P \le 0.05$ (2-tailed).

4. Discussion

Reactive oxygen species (ROS) are shaped from usual bi-product of plant metabolic activity. Any environmental disaster leads to enhanced fabrication of ROS which instigate progressive oxidative damage and ultimately triggering apoptosis. Even their destructive activity, they are considered as additional messengers towards conferment of tolerance to ecological hassles. A harmony between ROS assembly and their scavenging is maintained for proper signal transduction and defense against premature oxidative destruction to the plant cells [34] [35]. Competent scavenging of unwanted ROS created during any environmental stresses involves several non-enzymatic as well as enzymatic antioxidants activities in the tissues. In this work, we describe a comparative analysis of incidence of ROS scavenging enzymes and secondary metabolites in two true mangroves from Sundarbans mangrove forest in different salinity treatment (*in vitro*) in relation to their sustainability. The fact that the occurrence non-enzymatic factors and enzymatic disparities play an important role towards their sustainability has been discussed in detail.

The magnitude of salt adaptation is determined by a number of biochemical pathways that include retention and/or acquisition of water (water potential of cells), protecting chloroplast functions and conserving ion homeostasis. The synthesis and accumulation of low molecular weight metabolites, known (compatible solutes), is an omnipresent device for osmotic adjustment in plants. Their main role is to increase water potential (ψ) without hindering the normal metabolism [79]. In the present study, in A. marina, Aspartic acid, Alanine, Leucine and Proline play an essential role towards higher water potential and offer a better adaptability in high substrate salinity. Whereas, in H. fomes, among the studied free amino acid only Alanine and Phenyl alanine sowing correlation (at $P \le 0.05$ level) with salinity, but the other two (Aspertic acid and Tryptophan) are not significantly correlated and complete absent of Proline. These indicate relatively less adaptive ability in higher salt concentration. Munns [80] opined that high salinity hinder important physiological processes, such as photosynthesis and nutrient uptake, which are due to the reduction of leaf water potential. The increase of alanine in stressed leaves leading to excess glycolysis for enhanced respiration for the requirement of higher energy for salt sequestration in the vacuoles and endow with carbon skeletons for the photorespiration [81]. Enhanced production of Leucine and tryptophan signify the change occurs in protein synthesis in excess salt substratum [82]. Increment of Proline was also reported in salt stressed plants, as it helps to maintain required Ψ during stress [83]. Increments of Asp, Ala, Leu and Pro with enhanced salinity indicate the better adaptability of A. marina over H. fomes. Synthesis and accumulation of free amino acids in plant cells are considered as adaptive stress response of the plant [84] and regarded as compatible solute that adjust osmotic potential in cytoplasm [85] [86]. Parida et al. [18]

experimentally showed that in the cell sap of *Bruguiera parviflora*, the increment of osmotic solutes, like total sugar and total free amino acids occurred in various degrees under NaCl treatment and those osmoregulatory solutes help to restore the water potential more negative and might be consider as marker of extent of salt tolerance.

Oxidative damage of the plant cell relies broadly on the delicate equipoise between ROS production, and their efficient scavenging. Competent scavenging of ROS produced during various abiotic stresses involves the exploit of several non-enzymatic as well as enzymatic antioxidants present in the cells. In the present work, among the two taxa, *A. marina* shows a steady state increment of phenol and flavonoid along the salinity gradient but this trend do not occur in *H. fomes*, phenol and flavonoid both show an increment up to 200 mM of salinity and drop down beyond this. Radical scavenge (per cent inhibition) estimation with ABTS, DPPH and Fe²⁺ chelating assays experiments reveal a considerable augmentation along salinity ascent in case of *A. marine* which are not found in the other taxa. Ebrahimzadeh *et al.* [87] experimentally proved that the highest chelating activity and the largest amount of phenolic compounds were found in *Mellilotus arvensis*. Mishra and Das [88] indicated that the non-secretor mangrove *Bruguiera parviflora* grows as alt tolerance appliance through elevated level of antioxidative enzymes and escalation in osmolytes like proline to acclimatize under salt stress condition.

Antioxidative enzymes mediated toxic ion homeostasis is an essential progression of a plant in abiotic stress. The present work reveals that both antioxidative enzymes PRX and SOD produced either added number of isoforms or enhanced activity (qualitative and quantitative estimation) along the increased salinity treatment in case of *A. marina* over the control treatment than that of *H. fomes*, in which, the increment in both estimations hiked up to certain level, beyond which it dropped down. Jetesh *et al.* [89] concluded that the antioxidant enzymes protect halophytes from deleterious ROS production and accumulation in cytoplasm during salt stress. Hence, extent of salt tolerant differ in the investigated taxa and salt tolerant can be attributed to the enhance accumulation of compatible osmolytes in the cell sap (free amino acids) as well as increased production of non-enzymatic ROS foragers like phenols and flavonoids and antioxidant enzymes (PRX and SOD).

Abogadallah [67] commented that elevated antioxidant activity could be inferred as better tolerance to oxidative stress; the plant suffers less oxidative stress due to higher antioxidant activity. Mallik *et al.* [90] experimentally proved the activity of both CAT and SOD increased significantly and also resulted in the synthesis of new isoforms of both CAT and SOD in response to the NaCl treatment among a group of plants. Turkan *et al.* [91] worked on antioxidant activity on two strains of *Plantago* (*P. maritima*, salt-tolerant and *P. media*, salt-sensitive) in water logging and salinity stressed condition and opined that a salt-tolerant species showed an amplified antioxidant activity against lipid peroxidation towards salt-tolerance than a salt-sensitive one. In plants, ROS are being generated through lickage during electron transport systems of chloroplasts, mitochondria, and plasma membranes [92]-[94]. Abiotic stresses such as drought, salinity, chilling, metal toxicity, and UV-B radiation disrupt the cellular homeostasis and lead to boosted generation of ROS [95] [96]. The present observation in respect of well correlated with the incidence of free amino acids and antioxidants (both enzymatic and non-enzymatic components) with salinity treatment might be specifying towards the advantage of *A. marina* than *H. fo-mes* for sustainable existence in present days' elevated substrate salinity.

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