Gluconeogenic fructose-1,6-bisphosphatase from the mature sporocarps of common aquatic ferns: partial purification and basic characterization of this enzyme from Marsilea minuta (Polypodiopsida)

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Abstract. The present communication reports substantial activity of gluconeogenic fructose-1,6-bisphosphatase (FBPase; EC 3.1.3.11) in three common heterosporous aquatic ferns (Marsilea minuta, Salvinia natans, and Azolla pinnata) and also describes a protocol for its partial purification from mature sporocarps of Marsilea minuta. The cytosolic FBPase, obtained from Marsilea minuta, Salvinia natans, and Azolla pinnata was recognized as gluconeogenic enzyme due to its drastic catabolic inactivation in presence of externally administered glucose and its insensitivity towards photosynthetic light illumination. Cytosolic gluconeogenic FBPase was partially purified from mature sporocarps of Marsilea minuta to about 22-fold over homogenate following low-speed centrifugation (11, 400 × g), 30–80% ammonium sulfate fractionation followed by subsequent chromatography using matrices like CM-Cellulose, Sephadex G-200, and Ultrogel AcA 34. The profile of partially purified FBPase in PAGE under non-denaturing condition was recorded. The enzyme activity increased linearly with respect to protein concentration to about 100 μg and with respect to time up to 75 minutes. Temperature optimum was found at 35 °C. The effect of substrate concentration and kinetic analyses for FBPase were carried out using D-fructose-1,6-bisphosphate (D-FBP, the substrate) in the range of 0.0 to 1.0 mM at an interval of 0.1 mM concentration. The \( K_m \) value for D-FBP of FBPase was 0.06129 mM and \( V_{max} \) was 4525 nmole P released (mg)\(^{-1}\) protein h\(^{-1}\) as determined by nonlinear regression kinetics using Prism 8 software (Graph Pad). The enzyme was functional in a constricted pH range of 7.0 to 8.0, giving maxima at pH 7.5. This cytosolic enzyme was significantly stimulated by Mg\(^{2+}\) and strongly inhibited by Hg\(^{2+}\), Cu\(^{2+}\) and Zn\(^{2+}\).

Keywords: aquatic fern, enzyme purification and characterization, FBPase, gluconeogenesis, sporocarp

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Aquatic ferns bear specialized and unique reproductive structures known as ‘sporocarps’. These are sclerified seed-like propagules whose morphological nature has been variously interpreted. It is now generally agreed that these specialized, stomaticiferous foliar units are modified leaflets with two reflexed and conenate segments, each bearing a number of elongate sori following the veins (Kramer, 1990). The young immature sporocarps are green, leathery and photosynthetic, while on maturation they become non-photosynthetic, blackish brown in colour and eventually break down as a result of disintegration of the indusial and the sporangial walls releasing the spores to the external environment. These unique desiccation tolerant structures have been extensively studied from various angles like morphology, anatomy, physiology, cytology, phytoremediation, environmental biology, phylogeny, antioxidant activity, etc. over a period of several decades (Bierhorst, 1971; Bold et al., 1987; Gifford, Foster, 1987; Raubeson et al., 1994; Rothwell, Stockey, 1994; Nagalingum et al., 2006; Sood et al., 2012; Arokiyaraj et al., 2018). However, a thorough biochemical analysis of this specialized structure during its development has not been conducted adequately. There is a dearth of information regarding the biochemical aspects of sporocarp maturation and sprouting. It has been reported earlier that adequate glucose concentration is the central parameter in living organisms for optimization and fulfilment of all metabolic events either directly or indirectly (Eckstein et al., 2012). In plants this sugar is available in all parts basically via photosynthesis in leaves followed by phloem translocation. It has also been observed that under certain physiological conditions, the requisite concentration of glucose is provided via non-photosynthetic pathway like gluconeogenesis. The conversion of immature fern sporocarps into mature and senescent ones involves a definite loss of photosynthetic activity which in turn indicates the possible occurrence of gluconeogenic pathway and the concomitant involvement of the gluconeogenic marker enzyme, fructose-1,6-bisphosphatase (FBPase).

Fructose-1,6-bisphosphatase is a ubiquitous enzyme which catalyzes the hydrolytic dephosphorylation of D-fructose-1,6-bisphosphate (Fru-1,6-P₂) to D-fructose-6-phosphate (Fru-6-P);

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\text{Fru-1,6-P}_2 \rightarrow \text{Fru-6-P} + \text{Pi}
\]

List of abbreviations

BSA – bovine serum albumin; CM – carboxymethyl; EDTA – ethylene diaminetetraacetate; F-6-P – fructose-6-phosphate; FBP – fructose-1,6-bisphosphate; FBPase – fructose-1,6-bisphosphatase; g – gravitational field; \( K_m \) – michaelis constant; OD – optical density; PAGE – polyacrylamide gel electrophoresis; \( P_i \) – inorganic phosphate; \( V_{max} \) – maximum velocity.
6-phosphate and inorganic phosphate. Two isoforms of FBPase has been recognized for more than five decades. The first one is associated with the photosynthetic plastids and is involved in the reductive carbon assimilation cycle of photosynthesis. The second is located in the cytoplasm, which is involved in both gluconeogenesis and sucrose synthesis (Zimmerman et al., 1978; Stitt, Heldt, 1985). Cytosolic fructose-1,6-bisphosphatase is primarily associated with the regulation of gluconeogenesis in non-photosynthetic tissues like mature fern sporocarps. The occurrence of gluconeogenic FBPase is widespread in living organisms vacillating from bacteria to highly structured plants and animals. Several workers have studied this ancestral gluconeogenic enzyme from a wide number of microbial, plant and animal genera with diverse phylogenetic significance (Kobr, Beevers, 1971; Yu, 1989; Botha, Turpin, 1990; Jardon et al., 2008; Dey et al., 2014). However, perusal of available literature clearly reveals a serious information gap regarding the existence of this enzyme in vascular cryptogams, particularly in non-green senescent sporocarps of common aquatic ferns like Marsilea sp., Salvinia sp., and Azolla sp. Taking all the above facts into consideration the present investigation was aimed towards detection, partial purification and characterization of this ancestral gluconeogenic enzyme from three common heterosporous water ferns belonging to Marsileaceae and Salviniaceae.

Materials and methods

Chemicals and reagents

D-fructose-1,6-bisphosphate (di-sodium salt), D-fructose-6-phosphate (di-sodium salt), D-glucose-6-phosphate (di-sodium salt), D-galactose-6-phosphate (Gal-6-P, di-sodium salt), D-fructose-6-phosphate (F-6-P, di-sodium salt), D-mannose-6-phosphate (M-6-P, di-sodium salt), bovine serum albumin (BSA), were obtained from SRL (Mumbai, India). 2-Mercaptoethanol (ME), ammonium molybdate, acetic acid, ammonium sulphate, ammonium chloride, copper chloride, potassium chloride, tris(hydroxymethyl)aminomethane, trichloroacetic acid (TCA), dipotassium hydrogen phosphate, magnesium chloride, hydrochloric acid, sodium chloride, sulphuric acid, zinc chloride were purchased from E Merck India Ltd. (Mumbai, India). 2-Glycerophosphate was obtained from BDH, England. Coomassie Brilliant Blue (R-250 and G-250), Ultragel Aca 34 (U8878) were obtained from Sigma Chemical Company, St. Louis, Missouri, USA. Sephadex G-200 was obtained from Amersham Pharmacia Biotech marketed by SRL, Mumbai, India. Ascorbic acid, sodium bicarbonate, ammonium sulfate and ammonium chloride were procured from SRL (Mumbai, India). CM-Cellulose column matrix was purchased from Genei, Bangalore, India. Manganese chloride, dialysis membrane, etc. were obtained from Hi Media Laboratories Ltd. (Mumbai, India). Mercuric chloride and calcium chloride were obtained from Qualigens Fine Chemicals (Mumbai, India). All other chemicals used were of analytical grade and acquired from renowned Indian companies.

Plant material

Sporocarps of Marsilea minuta L., Salvinia natans L., and Azolla pinnata R.Br. were collected in different seasons from different ponds and paddy fields of Bongaon (23°4′ N, 88°49′ E) and Hasnabad (22.5°7′ N, 88.9°1′ E), 24-Parganas (N), WB, India in fresh condition and kept under –20 °C in the laboratory until use. Specimens were identified using standard taxonomic literature and herbarium specimens available at Central National Herbarium (CAL). Voucher specimens were deposited in the Departmental herbarium of Barasat Govt. College.

Isolation of fructose-1,6-bisphosphatase

In order to assay the enzyme (FBPase), and for detecting its activity in the sporocarps of the experimental samples separately at the initial phase of our work, the enzyme was isolated according to the following steps. The experimental material (about 10 gms) were collected and washed thrice with chilled distilled water and homogenized in a pre-cooled mortar and pestle in two volumes of the extraction buffer [50 mM tris-acetate (pH 7.5) containing 0.2 mM β-mercaptoethanol (ME)]. Neutral sea-sand was used in the mortar during the process of extraction. This homogenate was filtered to eliminate sand, unbroken cells and cellular debris. The resultant residue was discarded and the filtrate (homogenate) was collected. Half of the total volume of the filtrate was centrifuged at 11, 400 × g for 30 minutes in a Remi C 24 Plus cold centrifuge and the pellet was discarded. The supernatant (low-speed supernatant fraction) thus obtained was dialyzed overnight against the same buffer solution used for homogenization. On completion of dialysis, the low-speed supernatant was recovered from the dialysis bag and used for the preliminary assay.
Assay of fructose-1,6-bisphosphatase

A simple assay method of fructose-1,6-bisphosphatase was designed during the course of the present investigation in our laboratory, based on the method of Biswas et al. (1981). The typical incubation mixture restricted 100 mM tris-acetate buffer (pH 7.5), 5 mM MgCl₂, 0.2 mM fructose-1,6-bisphosphate (FBP), 0.1 mM EDTA, and an appropriate protein aliquot (50–200 μg) in a total volume of 1.0 mL. 0.2 mM fructose-6-phosphate (F-6-P) substituted FBP in each set of assay which served as control against non-specific phosphatases. Zero minute controls for both FBP and F-6-P, where the protein of the incubation mixture was destroyed by 0.25 mL of 20% chilled trichloroacetic acid (TCA) prior to the addition of substrate, served as double check against prying inorganic phosphate (if any). In totalling to these sets, an appropriate blank (minus enzyme) was also maintained. The enzyme incubation was carried out at 37 °C for 1 hour. After 1 hour of first (enzyme) incubation, the reaction was finished by the procedure applied for zero minute control. Therefore, in an assay set, there were six tubes, i.e., one blank, one zero minute control and one experimental tubes for each set (i.e., FBP set and F-6-P set). Afterwards, the amount of inorganic phosphate released on hydrolysis from the substrate(s) by the enzyme action was estimated according to the procedure of Chen et al. (1956) with slight modifications (Adhikari et al., 1987). 3.0 mL of a reagent (Pᵢ-reagent) and 3.0 mL H₂O were added to the reaction mixture (total volume thus became 7.25 mL) and incubated at 37 °C for 1 hour. The Pᵢ-reagent was prepared immediately before use by adding H₂SO₄ (6N), ascorbic acid (10 % w/v), ice-cold ammonium molybdate (2.5 % w/v) and H₂O in 1:1:1:2 ratios in that order. After incubation, the blue colour developed was measured at 820 nm in a Systronics-106 spectrophotometer. The free inorganic phosphate was estimated with the help of a standard curve prepared from different known quantities (0–100 μg) of phosphorous (K₂HPO₄ was used as the source). The enzyme-dependent release of inorganic phosphate was considered by subtracting the second corrected value (experimental tube – zero minute tube of ‘FBP set’) from the first corrected value (experimental tube – zero minute tube of ‘F-6-P set’). Protein was estimated by the method of Bradford (1976) with minor modifications using BSA as standard. A standard curve could thus be obtained and using that, the protein content of the experimental samples were estimated. As 1 mole of inorganic phosphate was hydrolyzed from one mole of FBP, mole number of inorganic phosphate quantified was equal to the number of FBP hydrolyzed at C-1 position.

Catabolic inactivation by externally administered glucose

In set I, sporocarps of Marsilea minuta, Salvinia natans, and Azolla pinnata were washed with distilled water and then two similar batches were designed, each containing 2 gms of fresh tissue for each individual genus. These experimental materials belonging to each batch were then kept in two different petri plates over moist filter paper (soaked thoroughly with distilled water). The experimental sets were maintained in presence of 5% glucose (at 24–26 °C) and control sets were also maintained parallel without glucose under similar temperature regimes. This treatment was carried out for 7 days and rotational light / dark condition was maintained identically for both the sets (experimental & control). After completion of the treatment, respective homogenate (s) of the tissues were prepared independently from both the sets (materials placed separately in presence of 0% and 5% glucose). FBPase assay was carried out following the procedure described in the previous section.

Influence under photosynthetic light

In set II, Marsilea minuta, Salvinia natans, and Azolla pinnata were washed with distilled water and then two similar batches were designed, each containing 2 gms of fresh tissue. These materials of each batch and for each genera were then maintained in two different petri plates over moist filter paper (soaked thoroughly with distilled water). One of these investigational sets were placed under alternate light/dark condition (8 hours light, 16 hours dark) for 7 days and the control set was maintained under complete darkness for 7 days. Subsequent to 7 days of treatment, respective homogenate (s) of the tissues were prepared independently from both the batches. FBPase assay was carried out following the procedure described earlier.

Partial purification of FBPase (from the sporocarps of Marsilea minuta)

10 gms of mature Marsilea sporocarps were washed with chilled distilled water twice and the partial purification of the cytosolic fructose-1,6-bisphosphatase from this sample was carried out following the steps as mentioned below (all the steps were carried out under 0 to 4 °C).
Homogenate
Water-washed sporocarps of Marsilea minuta was homogenized in a mortar and pestle with 2-volumes of 50 mM tris-acetate (pH 7.5) buffer containing 0.2 mM ME in presence of a little amount of washed sand. This homogenate was filtered to eliminate the sand, unbroken cells and cellular debris. The resultant residue was discarded and the filtrate (homogenate) collected.

Low speed supernatant
The resultant homogenate was centrifuged at 11, 400 × g for 30 minutes in a Remi C-24 Plus cold centrifuge. On completion of centrifugation, the pellet was discarded and the supernatant fraction was collected from the centrifuge tubes and pooled together.

Ammonium sulphate fractionation
The low-speed supernatant fraction obtained from the preceding step was then made 30–80% saturated with (NH₄)₂SO₄ by adding requisite amount of solid salt slowly with constant rousing (by using a Remi magnetic-stirrer). Then it was kept at 0 °C for 30 minutes and then centrifuged at 11, 400 × g for 30 minutes using Remi C-24 Plus cold centrifuge. The resultant pellet thus obtained after this centrifugation was dissolved in minimal volume of 50 mM tris-acetate buffer having 0.2 mM ME and dialyzed overnight against the same buffer (at least 1000 volumes) for complete removal of (NH₄)₂SO₄. On completion of dialysis, the 30–80% (NH₄)₂SO₄ fraction was recovered from the dialysis bag.

Cation exchange column chromatography through CM-cellulose
The 30–80% ammonium sulphate (dialyzed) fraction obtained from the earlier step were loaded onto a column (0.8 × 10.9 cm) of CM-cellulose [before this experimentation, commercially available CM-cellulose was flooded with 300–400 volumes of glass-distilled water over-night at room-temperature for complete swelling of gel beads. The excess water was then poured out and the column material equilibrated earlier with 50 mM tris-acetate buffer (pH 7.5) having 0.2 mM ME and dialyzed overnight against the same buffer (at least 1000 volumes) for complete removal of (NH₄)₂SO₄. On completion of dialysis, the 30–80% (NH₄)₂SO₄ fraction was recovered from the dialysis bag.

Gel filtration through Sephadex G-200
Active CM-cellulose fractions were pooled together and column chromatography was carried out through Sephadex G-200 (0.8 × 14.0 cm). Proteins were eluted with the 50 mM tris-acetate buffer (pH 7.5) having 0.2 mM ME in fractions of 3.0 mL at a flow rate of 25 minutes per tube. Eighteen such fractions were collected. The FBPase active Sephadex G-200 fractions were pooled together and molecular sieve chromatogram was made through UltrogelAcA 34 column.

Molecular sieve chromatography through UltrogelAcA 34
Active Sephadex G-200 fractions were pooled together and column chromatography was carried out through UltrogelAcA 34 column (0.8 × 9.5 cm). Proteins were eluted with the same buffer in fractions of 2.5 mL at a flow rate of 15 minutes per tube. Fifteen such fractions were collected. The FBPase active fractions obtained from UltrogelAcA 34 were pooled together and used as the enzyme preparation for biochemical characterization of FBPase.

Characterization of FBPase
Polyacrylamide gel electrophoresis was performed under native conditions following the method of Bollag et al. (1996) with minor modifications. All other experiments were carried out following conventional methods of enzymology.

Kinetic analysis
The $K_m$ value for D-FBP of FBPase was 0.2215 mM and $V_{max}$ was 7663 nmole P$_i$ released (mg$^{-1}$ protein h$^{-1}$) as determined by nonlinear regression kinetics using Prism 8 software (Graph Pad). All experiments were repeated thrice and results were expressed as means ± SE of three replicates.

Results
Screening of FBPase
Both young (green) and mature (brownish black) sporocarps of Marsilea minuta, Salvinia natans, and Azolla pinnata were screened separately for the presence of FBPase. In the young sporocarps of all the experimental samples, the specific activity of FBPase was relatively lower than the mature sporocarps obtained from the same source (Table 1).
Fig. 1. A–C: uninfluenced activity of FBPase under photosynthetic light illumination for 7 days in Marsilea minuta (A), Salvinia natans (B), and Azolla pinnata (C); D–F: catabolic inactivation of FBPase activity by externally administered glucose (5%) for 7 days in Marsilea minuta (D), Salvinia natans (E), and Azolla pinnata (F)
Fig. 2. A–C: elution profiles of FBPase and protein contents through CM-cellulose column (A), Sephadex G-200 column (B), and Ultrogel AcA34 column (C); D: Native PAGE profile with corresponding FBPase activity; E: effect of protein concentration on FBPase activity; F: effect of incubation time on FBPase activity
Identification of gluconeogenic FBPase

The activity of such FBPase in mature sporocarps remained unaltered under alternative light/dark conditions (Fig. 1, A–C) but, it was subdued under the presence of externally administered glucose for all the investigated genera (Fig. 1, D–F).

Partial purification of FBPase from the mature sporocarps of Marsilea minuta

This enzyme was partially purified from the mature sporocarps of Marsilea minuta to about 22-fold and with about 71% yield over homogenate following low-speed centrifugation (11, 400 × g), 30–80 % ammonium sulfate fractionation followed by subsequent chromatography using matrices like CM-cellulose, Sephadex G-200 and Ultrogel AcA 34 as detailed earlier in the methods section (Table 2). The general blueprint of elution profile of cytosolic FBPase and proteins are presented in Fig. 2, A–C.

Characterization of FBPase

The Partially purified FBPase profile was identified in PAGE under non-dissociating condition. Two major bands were marked in the native PAGE profile, out of which the first major protein band was found to coincide with the enzyme activity (Fig. 2, D). This partially purified enzyme exhibited variable activity in absence of different assay components (Table 3). The enzyme activity increased linearly with respect to protein concentration to about 100 μg (Fig. 2, E) and with respect to time up to 75 min (Fig. 2, F). Temperature optimum was found at 35 °C (Fig. 3, A). The effect of substrate concentration and kinetic analyses for FBPase were carried out using D-fructose-1,6-phosphate (D-FBP, the substrate) in the range of 0.0 to 1.0 mM at an interval of 0.1 mM. The \( K_m \) value for D-FBP of FBPase was 0.06129 mM and \( V_{max} \) was 4525 nmole P\(_i\) released (mg\(^{-1}\) Protein h\(^{-1}\)) as determined by nonlinear regression kinetics using Prism 8 software (Graph Pad) (Fig. 3, C). This FBPase specifically utilized fructose-1,6-bisphosphatase as its exclusive substrate. No other phosphorylated sugar could replace it (Table 4). The enzyme was functional in a confined pH 7–8 giving maxima at pH 7.5 using tris-HCl buffer in experiments (Fig. 3, B). This cytosolic enzyme was significantly stimulated by Mg\(^2+\) and highly subdued by Hg\(^2+\), Cu\(^2+\), and Zn\(^2+\) (Fig. 3, D).

Discussion

The reproductive structures of heterosporous ferns although referred collectively under a common term ‘sporocarp’, each structure is unique to their respective genera and are strikingly different from each other. The difference lies in terms of gross morphology, anatomy, time of maturation and mechanism of dispersal. Previous

Table 1. Activity of fructose-1,6-bisphosphatase from young and mature sporocarps of common aquatic ferns from low-speed supernatant fraction (11, 400 × g) [Results are mean ± SE, n = 3]

| Plant          | Plant part | Growth stage of sporocarps and collection period             | Specific activity of FBPase (nmole P\(_i\) released (mg\(^{-1}\) h\(^{-1}\)) | Total activity [nmole P\(_i\) released h\(^{-1}\)] | Yield (%) | Fold purification (Fold) |
|----------------|------------|--------------------------------------------------------------|-----------------------------------------------------------------|-----------------------------------------------|-----------|------------------------|
| Azolla pinnata | Sporocarps | Young (June 2019)                                           | 28.16 ± 4.21                                                    | 28.16 ± 4.21                                  | 100.00 ± 2.14 | 1.00 ± 0.03            |
|                |            | Mature (September 2019)                                      | 159.45 ± 25.77                                                 | 159.45 ± 25.77                                | 95.94 ± 3.06 | 1.13 ± 0.04            |
| Marsilea minuta| Sporocarps | Young (November 2019)                                        | 122.36 ± 18.05                                                 | 122.36 ± 18.05                                | 92.78 ± 7.16 | 1.34 ± 0.11            |
|                |            | Mature (February 2019)                                       | 210.16 ± 31.44                                                 | 210.16 ± 31.44                                | 73.37 ± 4.88 | 6.25 ± 0.43            |
| Salvina natans | Sporocarps | Young (December 2019)                                        | 71.22 ± 8.46                                                   | 71.22 ± 8.46                                  | 95.94 ± 3.06 | 1.13 ± 0.04            |
|                |            | Mature (March 2019)                                          | 176.31 ± 22.52                                                 | 176.31 ± 22.52                                | 92.78 ± 7.16 | 1.34 ± 0.11            |

Table 2. Partial purification of gluconeogenic FBPase from non-photosynthetic sporocarps of Marsilea minuta [Results are mean ± SE, n = 3]

| Fraction          | Total protein (mg) | Specific activity [nmole P\(_i\) released (mg\(^{-1}\) protein h\(^{-1}\)) | Total activity [nmole P\(_i\) released h\(^{-1}\)] | Yield (%) | Fold purification (Fold) |
|-------------------|--------------------|---------------------------------------------------------------------|--------------------------------------------------|-----------|------------------------|
| Homogenate fraction | 27.08 ± 2.46       | 176.90 ± 16.11                                                       | 4790.45 ± 6.48                                   | 100.00 ± 2.14 | 1.00 ± 0.03            |
| 11, 400 × g supernatant fraction | 22.83 ± 2.13       | 201.32 ± 30.34                                                      | 4596.13 ± 4.11                                   | 95.94 ± 3.06 | 1.13 ± 0.04            |
| 30–80% (NH\(_4\))SO\(_4\) fraction | 18.75 ± 3.86       | 237.05 ± 26.71                                                      | 4444.68 ± 12.34                                  | 92.78 ± 7.16 | 1.34 ± 0.11            |
| CM-cellulose fraction | 3.93 ± 6.42        | 1106.15 ± 5.43                                                     | 4347.16 ± 6.73                                   | 90.74 ± 5.49 | 6.25 ± 0.43            |
| Sephadex G-200 fraction | 2.06 ± 5.09        | 1706.42 ± 2.11                                                      | 3515.22 ± 1.26                                   | 73.37 ± 4.88 | 9.64 ± 1.31            |
| UltrogelAcA 34 fraction | 0.85±0.13         | 4035.16±1.50                                                        | 3429.88±32.14                                   | 71.59±2.73  | 22.18±1.14             |

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Fig. 3. A: effect of incubation temperature on FBPase activity; B: pH sensitivity of partially purified cytosolic FBPase; C: effect of substrate concentration on FBPase activity; D: effect of some monovalent and divalent cations on cytosolic FBPase activity.
Table 3. Effect of different assay components on FBPase activity [Results are mean ± SE, n = 3]

| Sl. No. | Condition          | Specific activity [nmole P released (mg)⁻¹ protein h⁻¹] | An average activity (%) |
|---------|--------------------|----------------------------------------------------------|-------------------------|
| 1       | Complete set       | 3853.46 ± 234.11                                         | 100.00                  |
| 2       | Minus substrate (FBP) | 0.00                                                      | 0.00                    |
| 3       | Minus buffer       | 3326.05 ± 187.19                                         | 86.31                   |
| 4       | Minus MgCl₂        | 3642.85 ± 206.15                                         | 94.53                   |
| 5       | Minus EDTA         | 3516.31 ± 188.32                                         | 91.25                   |
| 6       | Heat killed enzyme | 0.00                                                      | 0.00                    |

Table 4. Substrate specificity on FBPase activity [Results are mean ± SE, n = 3]

| Substrate                  | Concentration (mM) | Specific activity [nmole P released (mg)⁻¹ protein h⁻¹] | An average activity (%) |
|----------------------------|--------------------|----------------------------------------------------------|-------------------------|
| D-fructose-1,6-bisphosphate| 0.2               | 3942.17 ± 187.76                                         | 100.00                  |
| D-glucose-6-P              | 0.2               | 0.00                                                      | 0.00                    |
| D-fructose-6-P             | 0.2               | 0.00                                                      | 0.00                    |
| D-galactose-6-P            | 0.2               | 0.00                                                      | 0.00                    |
| D-Manose-6-P               | 0.2               | 0.00                                                      | 0.00                    |

reports are also indicative of the fact that because of the striking overall morphological differences among the spore-bearing structures, the two heterosporous fern families Marsileaceae and Salviniaaceae are not considered to be related closely (Bower, 1928; Eames, 1936; Tryon, Tryon, 1982). In the present investigation the activity of cytosolic FBPase was found to vary considerably among the three aquatic ferns screened. The activity varied considerably according to the growth stage of the collected sporocarps. Young and immature sporocarps exhibited a low enzyme titre value while maximum activity was recorded in mature sporocarps which were about to dehisce. Out of the three species scanned, highest activity was recorded in mature sporocarps of Marsilea followed by Salvinia and Azolla, respectively (Table 1) and this may be due to genotypic and physiological differences and other accompanying environmental factors.

The elevated activity of this enzyme in senescent sporocarps in comparison with their juvenile counterparts (Table 1) raises a very pertinent question regarding its actual physiological significance, which is yet to be conclusively resolved. A possible explanation of this phenomenon can be given from a biochemical point of view. Fern sporocarps are unusual modified reproductive structures that are reported to be highly desiccation tolerant [some are known to be viable 40 years after being collected (Bierhorst, 1971; Johnson, 1985)]. These structures have thick, sclerenchymatous walls that provide adequate protection against dryness and mechanical damage by insect herbivores. During maturation and before dispersal the sporocarp has to cope up with internal stress from the turgor developed. Just prior to spore dispersal the shielding coat of the sporocarp breaks down due to its prolonged acquaintance with water or a scratchy surface. Hydration changes the internal components and a gelatinous frame of pectin (i.e., the sorophore) comes out from the sporocarp.

Sprouting of sporocarps is dependent largely on the heterosachharide pectin, which besides providing mechanical strength and protection also helps the cells to cope up with the stress. Unequivocally, it is known that adequate glucose concentration is the central parameter of all living organisms for optimization of biological balance and for successful fulfillment of all metabolic events either directly or indirectly (Eckstein et al., 2012). It is also well established that under certain physiological conditions, the required concentration of glucose can be provided by non-photosynthetic pathway like gluconeogenesis. Since very little is known regarding the principal pathways of carbohydrate metabolism in fern sporocarps, the possibility of a gluconeogenic pathway in senescent fern sporocarps cannot be ruled out. Enzymatic studies on aquatic ferns have been conducted partially and discretely by Soltis and Soltis (1986), but their findings did not throw any light regarding the source of carbohydrates or sugars that are essentially required for growth and development of fern sporocarps, particularly at that time when the concerned vegetative mother plant becomes senescent and its photosynthetic efficiency and solute translocation remain on a low-level.

The present work accords with the pilot indication for the unique presence of the gluconeogenic fructose-1, 6-bisphosphatase for the first time in sporocarps of three heterosporous water ferns followed by its partial purification from mature sporocarps of Marsilea minuta. The results obtained in the present investigation reinvents the primordial nature of this protein in aquatic fern system. The persistence of this enzyme in vascular cryptogams and its alliance with the process of gluconeogenesis can clearly be proved through its catabolic suppression by glucose and simultaneously its lack of response under any light activation which is strongly evident in photosynthetic form of this enzyme (Fig. 1, A–F).

Previous reports dealing with the characterization of cytosolic FBPase have elucidated that the enzyme...
has a high affinity for its substrate (FBP), neutral pH optima, is inhibited by AMP and glucose and is also strongly stimulated by Mg^{2+} (Kobr, Beevers, 1971; Dey et al., 2014). The rudimentary characters of this enzyme illustrated in the present study are also comparable to other genera (Kobr, Beevers, 1971; Botha, Turpin, 1990; Jardon et al., 2008; Say, Fuchs, 2010; Yu, 1989). The preeminent activity of this FBPase in mature sporocarps in comparison with the young developing sporocarps (Table 1) nurtures a fundamental question regarding its physiological connotation, which is yet to be conclusively resolved. Is it a ‘safe-guard’ metabolic strategy adopted by the aquatic ferns for maintenance of the reserve carbohydrate pool during the stressful sprouting phase? In order to provide a satisfactory answer, further work may be carried out with the aid of radio isotope study, gene expression and proteomic analysis.

Conclusion

The present communication thus reports substantial activity of gluconeogenic FBPase in senescent spore bearing structures of three common heterosporous aquatic ferns. This ancient cytosolic enzyme (purified partially from Marsilea minuta sporocarps) was recognized as a gluconeogenic enzyme, due to its drastic inactivation in presence of glucose and its insensitivity towards photosynthetic light illumination. The results obtained in the present study also reveal the fact that a gluconeogenic pathway is operative in senescent sporocarps of Marsilea minuta, which may be regarded as a metabolic protective strategy adopted by the aquatic ferns for maintenance of the reserve carbohydrate pool during the stressful sprouting phase.

Declaration of Competing Interest

The authors declare that they have no conflict of interest.

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