



TECHNOLOGY DEVELOPMENT FOR MASS PRODUCTION OF *SPIRULINA PLATENSIS* IBSD-174 AND ITS UTILIZATION AS NUTRIENT RICH FORMULATIONS

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Spirulina platensis IBSD-174 was isolated from freshwater Loktak lake of Manipur, India during 2005-06 and partial characterization with special attention on morphological and biochemical has been done. The soluble protein content stabilized and enhanced up to 82.7 ± 0.08 $\mu\text{g/ml}$ with the treatment of different level of pH, nitrogen and phosphorus for consecutive three years in laboratory conditions. The strain was deposited and obtained accession number from the fresh water cyanobacterial and microalgal repository of IBSD-DBT, Imphal, Manipur, INDIA a creation of Department of Biotechnology, Govt. of India (Ref. No.: BT/PR11323/PBD/26/171/2008 dated 31st March 2009) for ready reference and commercial exploitation in future.

Keywords: *Spirulina platensis*, Loktak lake, Manipur, Accession number

Microalgae are the organisms capable of producing valuable metabolites such as pigments, proteins and vitamins for feed additive, pharmaceutical and nutraceutical purposes (Gladue and Maxey 1994, Cartens *et al.* 1996, Guerin *et al.* 2003, Hu 2004). *Spirulina platensis* is a planktonic photosynthetic filamentous cyanobacterium that forms massive populations in tropical and subtropical bodies of water which have high levels of carbonate and bicarbonate and alkaline pH of up to 11. *Spirulina platensis* is recognized by one main morphological feature *i.e.* the arrangement of multicellular cylindrical trichomes in an open left-hand helix along the entire length of the filament (Vonshak 1997). *Spirulina platensis* is the most popular in microalgal biotechnology with respect to its being an easy to grow species and having a simple harvest and drying process. *Spirulina platensis* has been produced commercially all over the world due to its high content of protein (up to 70%), pigments (especially the blue pigment, phycocyanin), essential fatty acid, vitamin B₁₂ and minerals (Cohen 1997, Boussiba and Richmond 1980, Mahajan and Kamat 1995, Becker 1981). It is also successfully used in aquaculture and poultry industry as well (Belay *et al.* 1996, Wikdors and Ohno 2001).

High pH and temperature are the key factors for the large scale cultivation of *Spirulina* under outdoor cultures. The optimal temperature for the cultivation

of *Spirulina* has been reported in the range of 35-38°C. This microalga also requires relatively high pH in the range of 9.5-9.8 (Hu 2004) which inhibits the contamination of most algae in the culture. High amounts of sodium bicarbonate must always be present in the medium to sustain high pH and prevent fluctuations (Zarrouk 1966). Different types of culture systems *e.g.* open ponds (Lee 1997), tubular photobioreactors (Torzillo *et al.* 1986), inclined glass panels (Hu *et al.* 1996) etc. are used in the mass production of *Spirulina*. On a commercial scale, *Spirulina* can be grown in raceway type open ponds which involve less investment and sun light can be utilized for photosynthesis (Torzillo *et al.* 1986). The objective of the study was to evaluate the influence of pH on the biomass and total soluble proteins of *Spirulina*. The study was also undertaken regarding its mass production under net house out door conditions and possibility of further utilization in the form of value added formulations.

MATERIALS AND METHODS

The experimental organism *Spirulina platensis* IBSD-174 was obtained from the freshwater cyanobacterial and microalgal repository (FWCMR) of IBSD-DBT, Imphal, Manipur, INDIA a creation of the Department of Biotechnology, Govt. of India with reference number BT/PR11323/PBD/26/171/2008

Table 1: Influence of pH on total soluble proteins ($\mu\text{g}/\text{ml}$) in *Spirulina platensis* - IBSD 174

pH	Total soluble proteins ($\mu\text{g}/\text{ml}$)
7.0	54.0 \pm 0.10
7.5	59.0 \pm 0.15
8.0	62.0 \pm 0.10
8.5	66.0 \pm 0.11
9.0	70.9 \pm 0.09
9.5	79.3 \pm 0.07
10.0	82.7\pm0.08
10.5	77.9 \pm 0.09

dated 31st March 2009. This strain was originally isolated from Loktak lake of Bishnupur, Manipur, India during 2005-06 through enrichment culture technique. The cyanobacterial strain *Spirulina platensis* IBSD-174 was grown and maintained in chemically defined modified CFTRI medium (Venkataraman *et al.* 1995) under diffused light intensity of 54-67.5 $\mu\text{mole m}^{-2}\text{s}^{-1}$ at 28 \pm 2 $^{\circ}\text{C}$ temperature with 14:10 hrs light and dark phase. The purity and identification was authenticated based upon the keys given by Desikachary (1959). The effect of pH (6.0-10.5) was examined on total soluble proteins (Herbert *et al.* 1971). All the ingredients of CFTRI medium was added accordingly for 1L of medium and the contents were mixed properly with magnetic stirrer and the pH of the medium was adjusted (Thermo Scientific, ORION 2 STAR) using 1N NaOH and 1N HCl. The pure culture of *Spirulina platensis* IBSD-174 was inoculated in 5 litres of modified CFTRI medium (Venkataraman and Becker 1985) in plastic bucket and / or tubs of 15 litres capacity. The containers were kept in a net house covered with polythene sheets. The buckets and tubs were covered with white muslin cloth to prevent the growing cultures from contamination with dust particles and also to protect them from laying of eggs by mosquitoes and other insects. Mass cultivation was carried out under such conditions without providing any additional illumination or temperature regulation for fifteen days. The

microalgal suspension was regularly mixed with the help of sterilized stainless steel rods. The algal biomass was harvested on 15th day of growth by filtering through a dense net. The harvested biomass was washed twice with distilled water to remove the adhering salts and dried in an oven at 30-50 $^{\circ}\text{C}$ for 30-45 mins.

Isolation of genomic DNA and 16S rRNA analysis

Isolation of genomic DNA was done as per the method described by Tillett and Neilan, 2000 and 16S rRNA was carried out by using PCR mix for 25 μl reaction mixture containing 1X PCR (Tris-EDTA with MgCl_2) buffer, dNTP mix (25 mM each of dNTPs), Taq DNA polymerase, 0.3 μM forward primer (536f 5' -GTGCCAGCAGCCGCGGTRATA-3'), 0.3 μM reverse primer (1488R 5' -CGGTTACCTTGTTACGACTTCACC-3'), template DNA (50 ng) and sterile double distilled water.

The PCR conditions were set for 28 cycles with initial denaturation at 95 $^{\circ}\text{C}$ for 5 mins then 95 $^{\circ}\text{C}$ for 1 min, annealing at 55 $^{\circ}\text{C}$ for 1 min and final extension at 72 $^{\circ}\text{C}$ for 2 mins using thermal cycler (Mastercycler gradient, eppendorf).

The amplified PCR product was loaded with loading buffer in the ratio 5:1 in 2% agarose (Sigma) gel and ran in electrophoresis unit (Elchrom Scientific) using 1X TAE buffer (Promega) for 30 mins at 90V. The size of the PCR product was compared with 200 bp DNA markers (Himedia). The product bands were observed in Gel documentation system (Vilber Lourmet). The quality and quantity of amplified PCR product was checked by nanodrop (Biospectrometer, eppendorf).

Sequencing and analysis of the 16S rRNA gene

The amplified PCR products of the 16S rRNA were sequenced at NCCS-DBT, Pune using the same forward and reverse primers and nucleotide sequences obtained from DNA sequencing were compared with sequence information available in the National Center for Biotechnology Information (NCBI) database using BLAST (Altschul *et al.* 1990, 1997) with nucleotide option by logging onto <http://>

www.ncbi.nlm.nih.gov. The sequences retrieved were converted to FASTA format.

RESULTS AND DISCUSSION

The identification of the *Spirulina platensis* IBSD-174 was authenticated based upon microscopic examination and molecular tools. The microalga *Spirulina platensis* IBSD-174 is a multicellular filamentous cyanobacterium, whose cells (diameter from

1-12 μm) are organized in helicoidal trichomes without ramifications (Richmond 1990; plate 1b). The influence of growth conditions on the chemical composition of *Spirulina* has been studied by many researchers with the purpose of optimizing the production of economically and nutritionally interesting compounds (Cohen *et al.* 1993). The extraction of nutritionally active compounds in pure form is expensive but direct consumption of *Spirulina* as a nutraceutical food is a viable alternative. The pH of

Table 2. Sequences producing significant alignments:

Accession	Description	Max score	Total score	Query coverage	E value	Max ident
DQ279773.1	<i>Arthrospira platensis</i> strain Sp-5 16S ribosomal RNA, tRNA-Ile, and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene, partial sequence	1478	1478	90%	0.0	90%
EF432318.1	<i>Arthrospira platensis</i> Sp-16 16S ribosomal RNA, tRNA-Ile, and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
EF432314.1	<i>Arthrospira platensis</i> Sp-8 16S ribosomal RNA, tRNA-Ile, and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
EF432313.1	<i>Arthrospira platensis</i> Sp-7 16S ribosomal RNA, tRNA-Ile, and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
EU427543.1	<i>Arthrospira platensis</i> HZ01 16S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
AY672713.1	<i>Arthrospira platensis</i> strain AICB 49 16S ribosomal RNA gene, partial sequence; tRNA-Ile and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
DQ393282.1	<i>Arthrospira platensis</i> strain SAG 257.80 16S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
DQ393279.1	<i>Arthrospira platensis</i> strain NIES 39 16S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
AF329392.1	<i>Arthrospira</i> sp. FACHB438 16S ribosomal RNA gene, complete sequence; tRNA-Ile and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene, partial sequence	1476	1476	90%	0.0	90%
EF432315.1	<i>Arthrospira platensis</i> Sp-12 16S ribosomal RNA, tRNA-Ile, and tRNA-Ala genes, complete sequence; and 23S ribosomal RNA gene	1474	1474	85%	0.0	91%

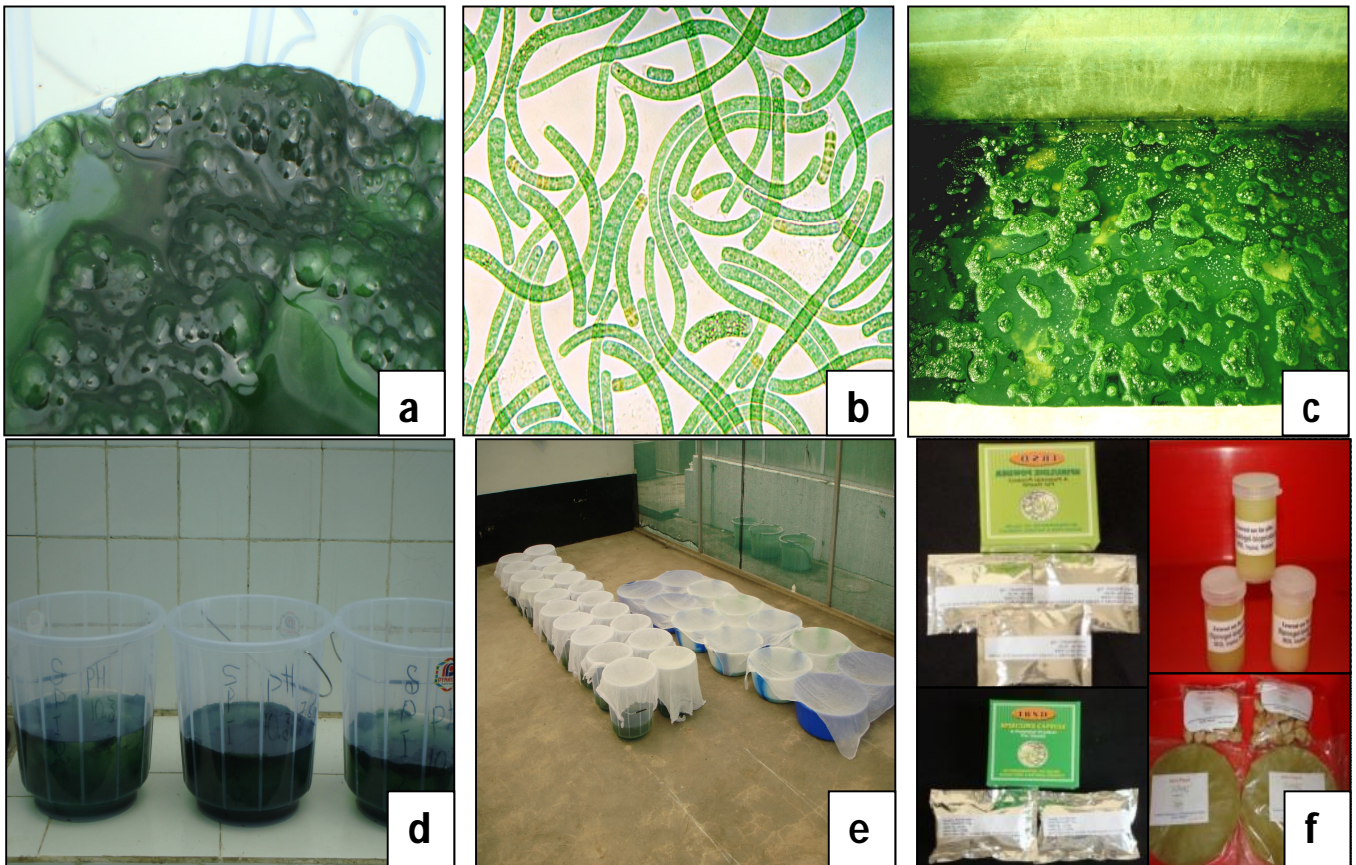


Plate 1(a-f):Growth of *Spirulina platensis* and its mass production for possible utilization as nutritionally rich formulations (a) *Spirulina* biomass in liquid culture (b) Photomicrograph showing *Spirulina* filaments (c) Growing *Spirulina* in tanks (d) Buckets with *Spirulina* (e) Growth units of *Spirulina* covered with muslin cloth (f) *Spirulina* formulations: Spiro-capsule, Spiro-powder, Spiro-gel, Spiro-bori and Spiro-papad developed with dried *Spirulina* biomass

the growing medium has been reported to be a very important parameter which influences the growth of the organism. The enhancement in the pH from 7.0 to 10.0 enhanced the level of total soluble proteins and further increase in pH decreased its content at 15th day of incubation (Table 1). These results are in accordance with the reports of other laboratories where an increase in pH above 10.5 was also shown to be harmful for the growth of this alkaline pH loving organism (Richmond *et al.* 1980, Goksan *et al.* 2007). The optimal growth temperature for *Spirulina platensis* was shown to be between 30°C to 35°C and higher temperature than this may certainly be deleterious for its growth (Danesi *et al.* 2001, Vonshak 1997). The *Spirulina* cultures grew efficiently as liquid culture and in plastic tubs or buckets (Plate-1 a, c, d) and muslin cloth cover prevented the contamination of *Spirulina* biomass (Plate-1 e). Detailed chemical com-

position of the harvested biomass indicated that the product had immense biotechnological potential (crude protein- 67.00%; crude lipids-3.1%; carbohydrate- 14.5%; nucleic acid- 4.5%; essential minerals- 7.6% and vitamins-1.2%). Growth conditions optimized for biomass production and productivity were usually used in the commercial production of *Spirulina* without considering chemical composition, but higher concentrations of potentially useful compounds such as polyunsaturated fatty acids, proteins and phenolics can be obtained by manipulating growth conditions. According to Torzillo and Bernardini (1991), for outdoor cultures of *Spirulina*, upto 34% of the biomass produced during the daylight period may be lost through respiration at night such chemically analyzed biomass was utilized for the development of different formulations (Plate-1 f). Our studied strain showed maximum similarity with *Ar-*

throssira platensis strain Sp-5 16S ribosomal RNA, partial sequence bearing accession no. DQ279773.1 having 90% similarity revealing that morphological identification is similar with the molecular characterization.

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>          i          b          s          d
1246bpCTTCCGGGGGCGTGGGTAAC TCCCCC
GTGTGACGGGGGTGTGTACAAGGCCGGGA
ACGGATTCACCGCGGTATGCTGACCCGCGAT
TACTAGCGATTCCACCTTCATGCACGCGATTT
GCAGACTGCAATCTAAACTGAAGAAAGTTTT
AGGGAATGGCTCCAGACCCCGTCTTGGCTC
CCCTTGTACCCCCCATTTGAAAAGGGGTGA
ACCCAGGACAAAGGGCGCATGATACTTTGA
CTTCTCCCCCCTTTCCCCCGGTTTGTCCCCG
GCATTCCCCCTAAAGTGCCCAATTGAATGATG
GCAACTAAGGACGAGGGTTGCGCTCGTTGCG
GGACTTAACCAACATCTCACGACACGAGCT
GACGACAGCCATGCACCACCTGTGTTCCCGC
TCCCGAAGGCAATCACGGCTTTCACCAAGAT
TCCAGACATGTCAAGCCCTGGTAAGGTTCTTC
CCGTTGCATCCAATTAACCACATACTCCACC
GATTGGGCGGGCCCCCGTCAATTCTTTGAGT
TTCAGACTTGCGTGCCTACTCCCCAGGGGGG
AACTTAACTCGTTAGTTTCGGCACAGCCCGG
GTCGATACAGGATACACCTAGTTTCCATTGTT
TACGGCTAGGACTACTGGGGTATCTAATCCC
ATTCGCTCCCCTAGCTTTCGTCCTCAGTGTC
AGTTACGGCCCAGCAGAGCGCTTTCGCCACA
CGGTGTTATTCCCGATATCTACGCATTTACC
GCTACACCGGGAATTCCCTCCGCCCCACCCT
ACTCTAGGGGTTTCAGTTTCCACTGCCTTTCAG
TAGTTGAGCTACTGTCTTTGACAGCAGATTTG
AACTACCACCTACACCCTCTGTACGCCAATC
ATTCTGGCAAACAGTTACCTGGTGCGTATTAC
CGCGGTGCCGGGCACGGAGTTAGCCGAGGCT
TATTCCTCAAGTACCGTCATTGTGTTCTTCCTT
GAGAAAAGGGGTTTACAACCCAAGAGCCTTC
TTCCCCCTCGCGGTCTTGCTCCGTCAGGCTTG
CGCCCATTGCGGAAAATTCCCCGGAGTGCCT
CCCGTAGTCAGTCCCCGTGGTGTGTCAGTCATCA
GTGCAGACCTCATCCTGTGTCAGTCCCCTTGTA
TCGTGCGCTCCCCTTGTATATCTCGCACACTC
ATCAAATAGGAGAGGGCCTCCAACATCTTCT
CCCGTAGGGCTCA

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