

Spirulina: A Green Factory



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Spirulina (Arthrospira) :

Is a photosynthetic, filamentous, spiral-shaped, multicellular, blue-green microalga

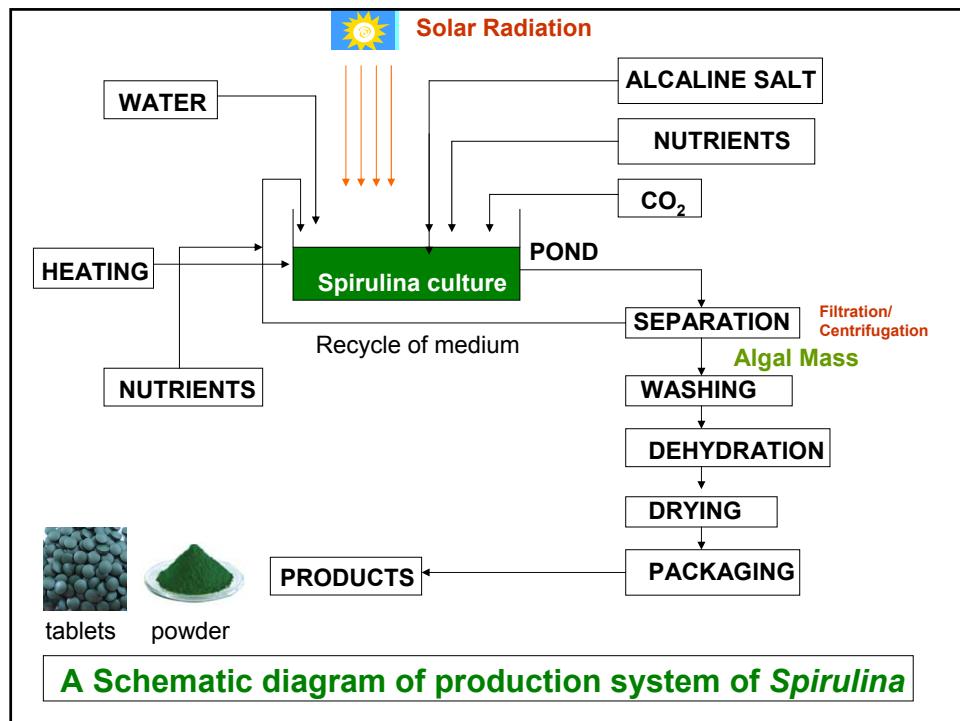
*It belongs to Cyanobacteria,
Oscillatoriaceae (family),
Arthrospira (genus)*

Most commercial production systems are based on large shallow raceways in which spirulina cultures are mixed by a paddle wheel under controlled conditions

It has a long history of use as a safe health food and as an important source of protein.



*Arthrospira sp. cultured
in our Lab*



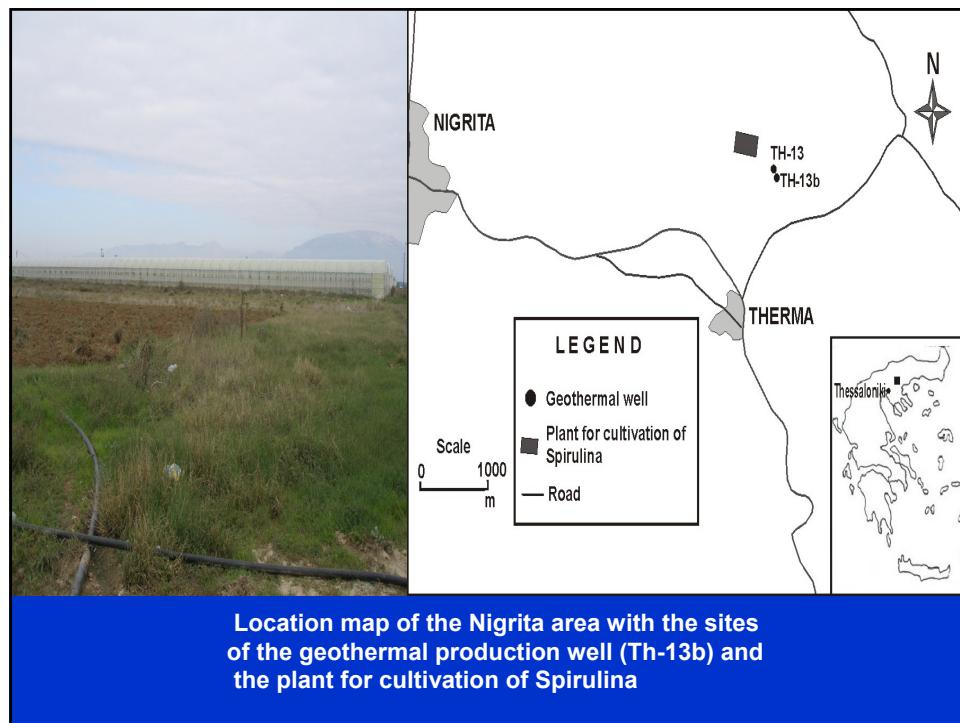
Geothermal Fluids used for Spirulina Production in Greece

- They represent a cheap and renewable energy form that is sequestered in underground aquifers
- They come from high depths
- They have high temperatures
- They usually contain dissolved CO₂

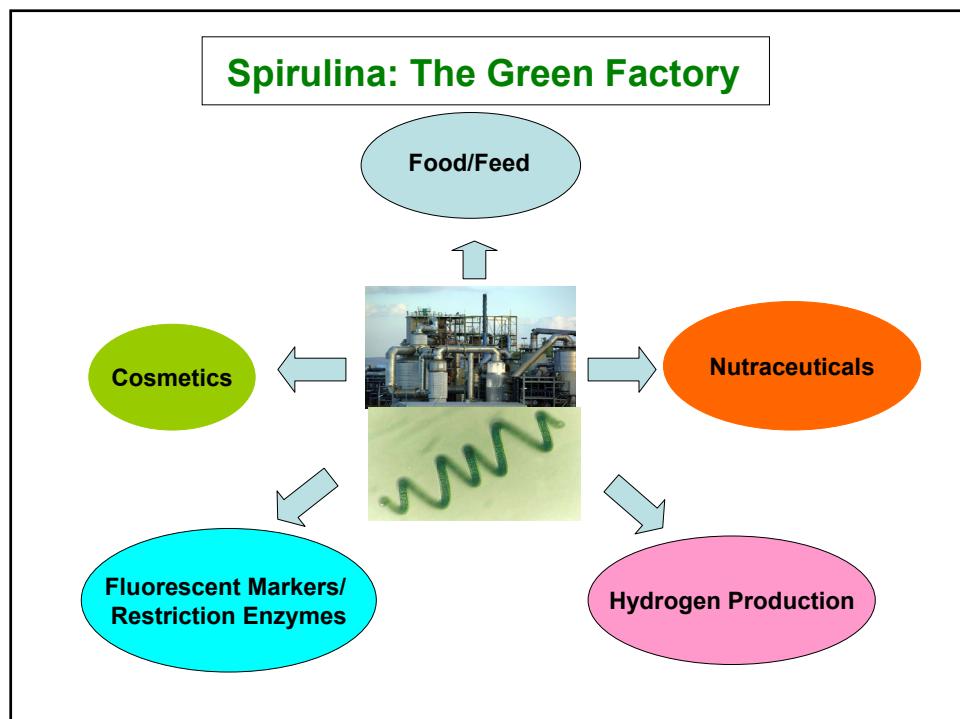
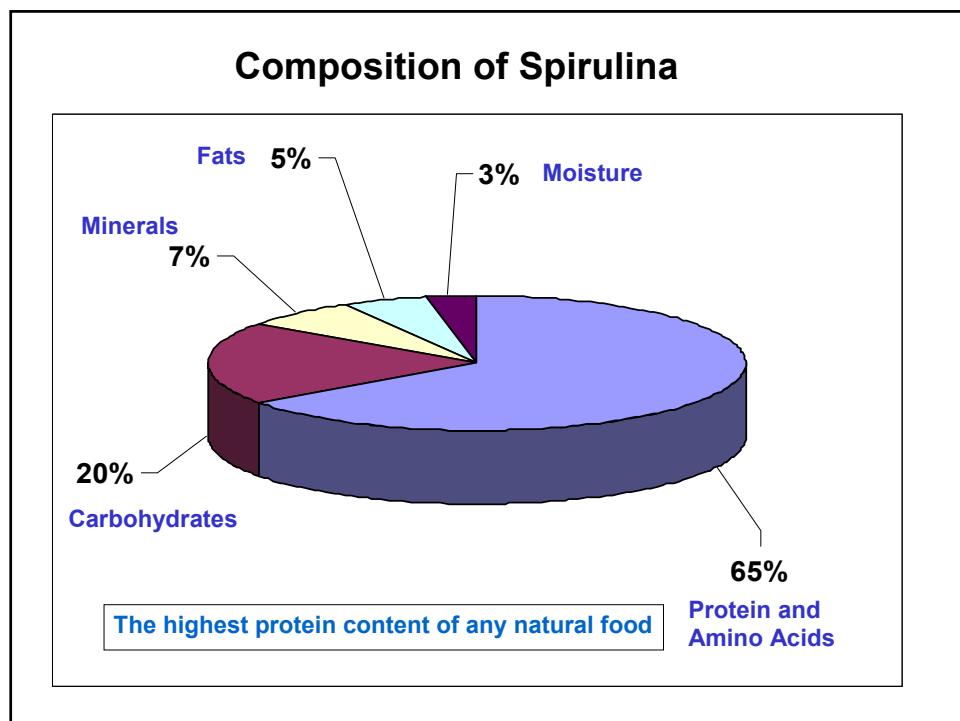
- ✓ The geothermal energy used in this way in greenhouse cultivation ponds contribute to increasing the daily yield by 20-30%

- ✓ The use of freely released geothermal CO₂ reduces the production costs of *Spirulina* biomass by over 25%.





ALGAE A.C. Spirulina production pond

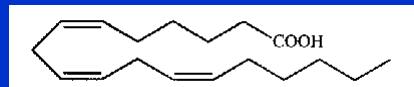


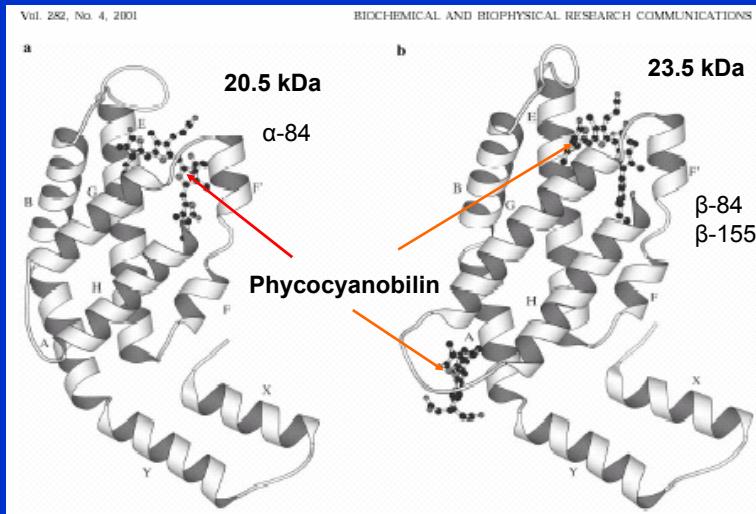
Potential Applications of *Spirulina* as Nutritional and Therapeutic Supplement in Health Management

- Antioxidant
- Anticancer effects
- Immunomodulation effects
- Anti-viral effects

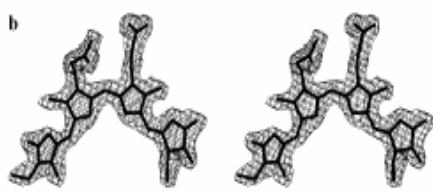
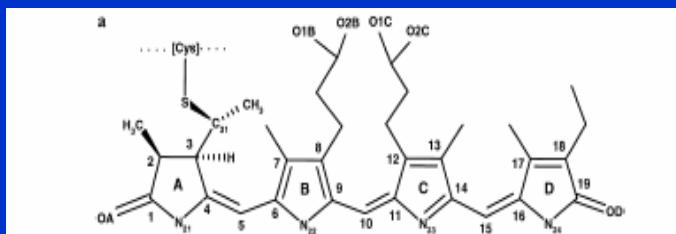
Health benefits from Spirulina are mainly due to some major bioactive components:

- ***Phycocyanin (Phycocyanin C/Allophycocyanin proteins)***
- ***Spirulan (sulphated polysaccharide : MW~220,000 ; rhamnose (52%), uronic acids (16%))/***
Immulina (High MW polysaccharide(~10⁷) ; rhamnose (35.4%)
- ***Gamma Linolenic acid (GLA) (ω -6 PUFA)***





Ribbon representation of *spirulina platensis* Phycocyanin C subunits (Phycocyanin C: Trimer of α/β heterodimers)

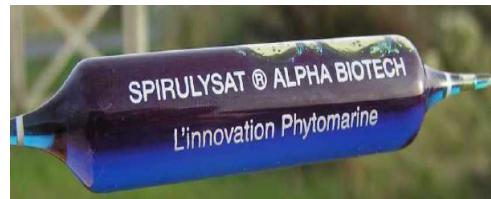
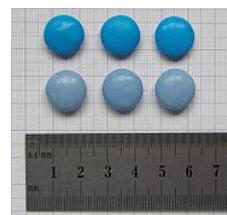


PCB is covalently attached to cysteines of the apoprotein via a thioether bond to C-31 on ring A

The final step in phycobiliprotein biosynthesis is the covalent chromophore attachment to the apoprotein by heterodimeric lyases

Commercial products containing Spirulina bioactive components

Chocolate confectionary



Phycocyanin extract

Old and new
Blue smarties (Nestlé)
with phycocyanin (2008)
(artificial coloring was removed)



A Spirulina high Mr Polysaccharide
Nutraceutical

Commercial GLA preparations
contain plant-based oil GLA

Present Research activities

Studies of *Spirulina* bioactive components in order to evaluate their potential use in biomedicine and cosmetics

- Lipid components
- Sulphated polysaccharides
- Phycocyanin and phycocyanobilin (PCB)
- Enzyme activities present in dried microalgal biomass

Lipid Components

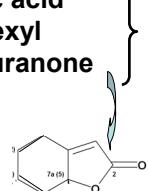
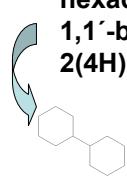
What is the chemical composition and Bioactivity
of Volatile Lipid Components in Commercial
Spirulina Dry Biomass ?

Major Volatile Lipid Components of *Spirulina* Dry Biomass Identified by GC-MS

heptadecane (77.88%)
6,10,14-trimethyl-2-pentadecanone (6.09%) } Volatiles by
Hydrodistillation

L-linalool (10.5%)
9(Z)-octadecenoic acid (10.19%)
hexadecanoic acid (5.0%)
cis-linalool oxide (4.08%) } Pentane Extract

hexadecenoic acid
1,1'-bicyclohexyl
2(4H)-benzofuranone } Fractions of
Si gel Chromatography of
Dichloromethane extract



Antimicrobial Action of Spirulina extracts and Si-Gel Chromatography Fractions

Antimicrobial activity was assayed against six human pathogenic bacteria and three fungi through both the disc diffusion (inhibition zones (mm)) as well as the dilution methods.

Tested samples	<i>S. aureus</i>	<i>S. epidermidis</i>	<i>P. aeruginosa</i>	<i>E. cloacae</i>	<i>K. pneumoniae</i>	<i>E. coli</i>	<i>C. albicans</i>	<i>C. tropicalis</i>	<i>C. glabrata</i>
Cyclohexane extract	8	10	NA	NA	NA	10	NA	8	NA
Dichloromethane extr.	11	11	NA	NA	NA	9	NA	9	8
Methanol extract	15	14	10	11	10	10	9	10	9
¹Fraction 70-74	12	10	9	10	10	9	8	9	9
¹Fraction 50-56	11	12	10	10	9	9	8	9	9
¹Fraction 148-151	12	12	10	10	9	10	8	10	10
¹Fraction 169-170	11	12	9	10	9	11	7	10	11
¹Fraction 223-224	12	11	10	11	10	11	7	10	10
¹Fraction 268	12	13	10	11	11	10	8	10	10
¹Bicyclohexane	12	13	10	10	11	10	8	9	10
Netilmicin	17(0.004)	19(0.004)	16(0.088)	17(0.008)	18(0.008)	16(0.010)	NT	NT	NT
Itraconazole	NT	NT	NT	NT	NT	NT	20	22	23
5-flucytocine	NT	NT	NT	NT	NT	NT	20(0.01)	21(0.001)	22(0.0001)

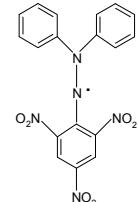
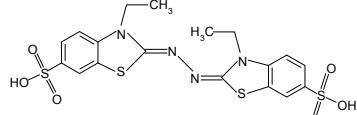
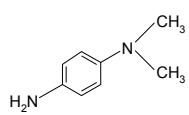
NA, Not Tested; NA, Not Active; Chromatography ¹Fractions of Dichloromethane extract

Anionic Polysaccharides

**Spirulina anionic polysaccharides express
Immunomodulating and Antiviral activities**

Free radical Scavenging Activity ?

Scavenging of stable free radicals by Spirulan (spectrophotometric determination)



$EC_{50} = 0.2 \text{ mg/ml}$

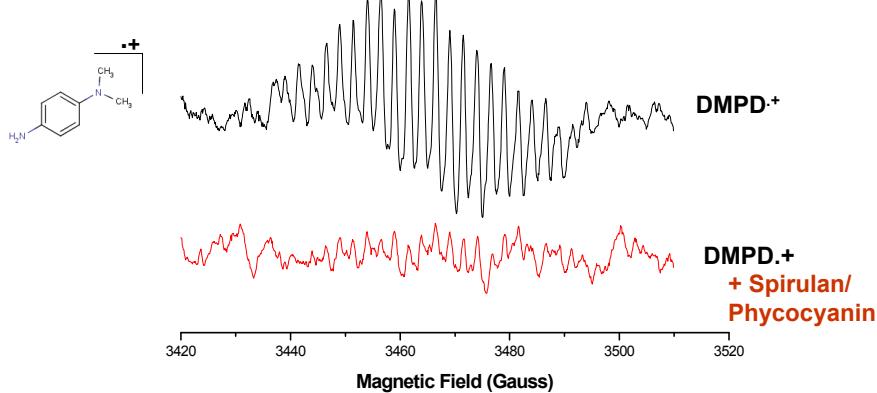
$EC_{50} = 0.3 \text{ mg/ml}$

No

Polysaccharide Specificity

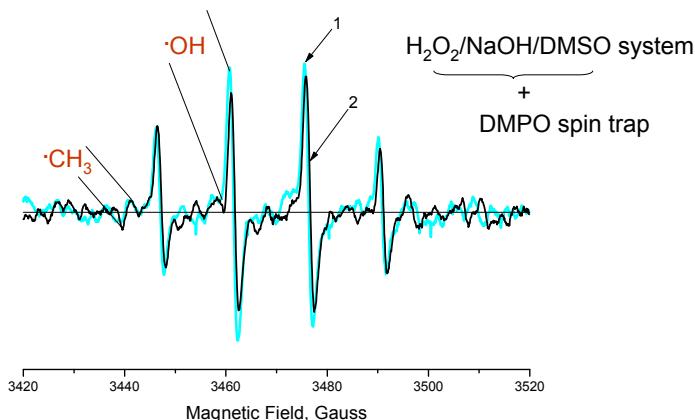
**Chondroitin sulfate, Hyaluronic acid and Heparin
express very low scavenging activities**

Scavenging of DMPD⁺ by Spirulan or Phycocyanin followed by EPR Spectroscopy



Tsapra et al. submitted

Spin-Trapping of Free Radicals in absence (1) and presence (2) of a *Spirulina platensis* Polysaccharide Fraction



Analysis of the EPR spectrum indicates that the polysaccharide scavenges hydroxyl radicals

Phycocyanobilin (PCB)

Phycocyanin : ~ 14% of Spirulina dry biomass

PCB: ~ 0.66% of Spirulina dry biomass
(15 g of Spirulina (a heaping tablespoon) contains about 100 mg PCB)

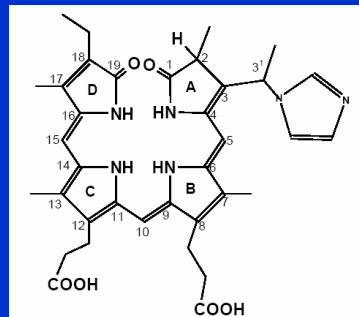
Phycocyanin exhibits a variety of pharmacological properties

Phycocyanobilin ?

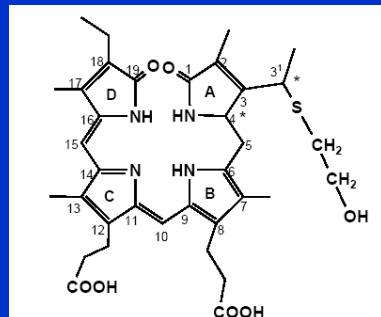
PCB Preparation

Spirulina → Phycocyanin → Methanolysis → PCB → Chromatography

Since a Spontaneous formation of PCB adducts has been observed
 (Tu et al (2009) JACS, 131, 5399)



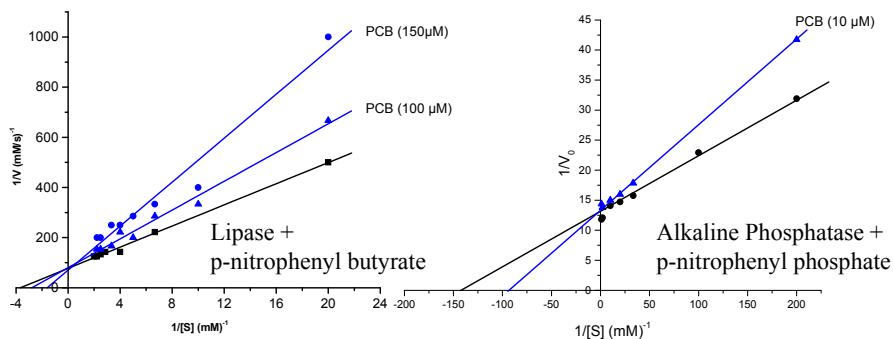
PCB-imidazol adduct



PCB-mercaptopropanol adduct

Therefore: Our aim was to examine the interaction of PCB with various biomolecules

Inhibition of Lipase and Alkaline Phosphatase by Phycocyanobilin (PCB)



$$K_i = 320 \mu\text{M}$$

$$K_i = 18.9 \mu\text{M}$$

PCB irreversibly modified:

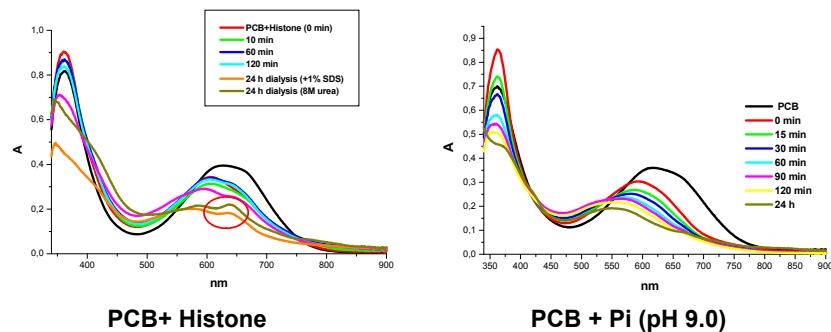
BSA, Histone H2B, Phosvitin, certain anionic polysaccharides and DNA

PCB interacts with:

the anions phosphates and borates as well as with the amine Tris

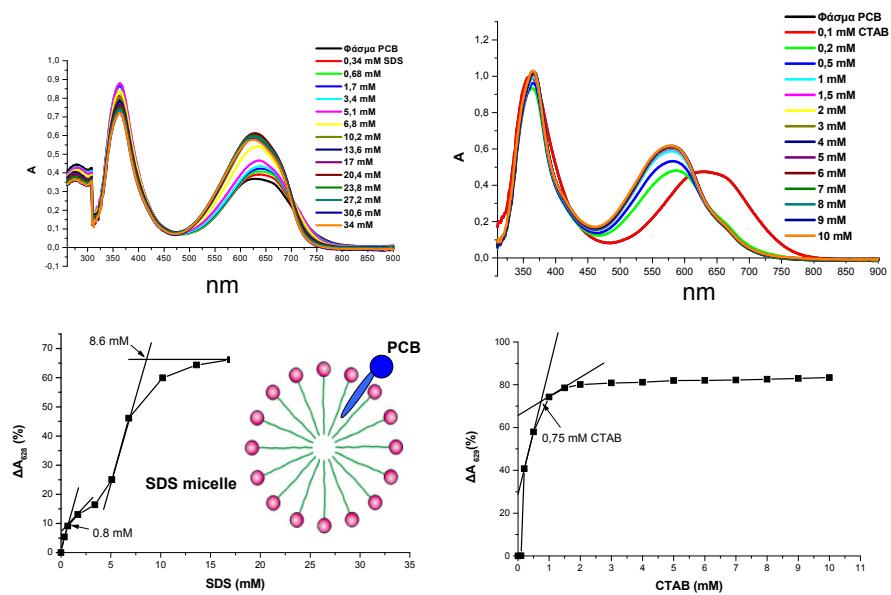
PCB interacts with surfactants:

SDS, Triton X100, CTAB



All the above interactions may cause autocatalytic PCB covalent modifications

Titration of PCB by SDS and CTAB



Peroxidase of *Spirulina* dry Biomass

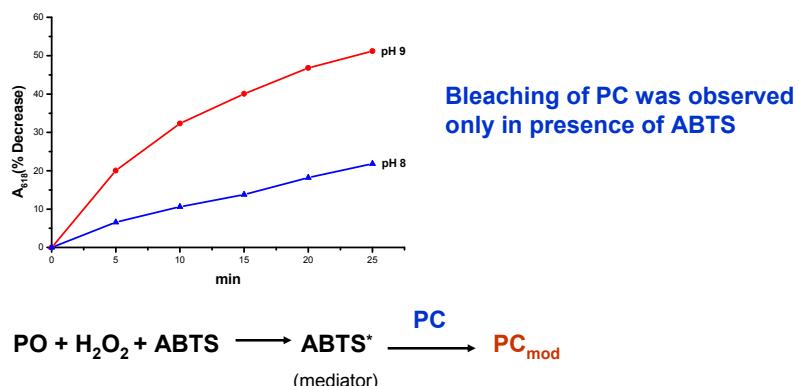
ROS-scavenging enzymes are increasingly used in skin health care products

Is Peroxidase Activity present
in Dry Commercial Spirulina Biomass ?

Peroxidase from Spirulina Dried Biomass

- ✓ Partial purification by ammonium sulfate fractionation and gel filtration
- ✓ Molecular mass: 43.500 Da
- ✓ Two pH optima: pH 4.0 and pH 8.0
- ✓ Km for ABTS: 46 µM (best substrate)
- ✓ Km for H₂O₂ : 21.2 mM

Spirulina Peroxidase (PO)-induced Spectral Change of Phycocyanin (PC) in presence of H₂O₂ and ABTS



Since Peroxidase activity survives the manufacturing conditions for *Spirulina* powder production, it may play significant role in cosmetic products in blocking hydrogen peroxide toxicity

Future Research activities

Metabolic Engineering, Metabolic Flux Analysis and determination of key metabolites using HPLC-MS/NMR techniques will be used for designing and modifying *Spirulina* pathways for the overproduction of important constituents of this microorganism

Synthesis of hydrophobic derivatives of PCB as membrane-targetted photosensitizers for photodynamic therapy

New cosmetic products containing *Spirulina* bioactive components

NHRE

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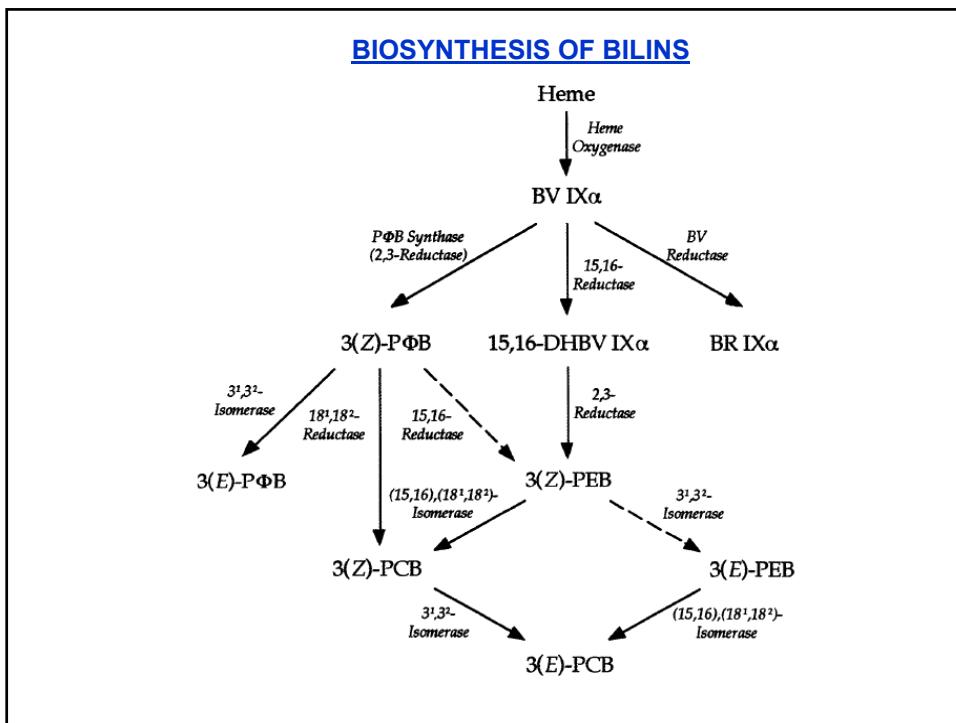
Prof. E. Nerantzis

ALGAE AC, Nigrita Serres

Bilateral Greece-Spain GSRT 006-γ Programme

01 PRAXE 11 GSRT Programme

Thank you for your attention



Arthrosphaera platensis str. Paraca, whole genome shotgun sequencing project

Genome Info:	Features:	BLAST homologs:	Links:	Review Info:
Refseq: NZ_ACSP00000000	Genes: 0	COG	Genome Project	Publications: None
GenBank: ACSK00000000	Protein coding: None	TaxMap	Refseq FTP	Refseq Status: WGS
Length: 4,997,563 nt	Structural RNAs: None	TaxPlot	GenBank FTP	Seq.Status: Draft
GC Content: 44%	Pseudo genes: None	GenePlot	BLAST	Sequencing center: University of Applied Sciences of Western Switzerland
% Coding: 0%	Others: None	gMap	TraceAssembly	Completed: 2009/07/20
Topology: other	Contigs: 1820	CDD	Organism Group	
Molecule: DNA	Other genomes for species:			

<http://www.ncbi.nlm.nih.gov/sites>

Arthrosira maxima CS-328, whole genome shotgun sequencing project

Genome Info:	Features:	BLAST homologs	Links:	Review Info:
Refseq: NZ_ABVK00000000	Genes: 5728	COG	Genome Project	Publications: None
GenBank: ABVK00000000	Protein coding: 5690	TaxMap	Refseq FTP	Refseq Status: WGS
Length: 6,003,314 nt	Structural RNAs: 38	TaxPlot	GenBank FTP	Seq.Status: Draft
GC Content: 44%	Pseudo genes: None	GenePlot	BLAST	Sequencing center: US DOE Joint Genome Institute (JGI-PGF)
% Coding: 82%	Others: None	gMap	TraceAssembly	Completed: 2008/10/15
Topology: other	Contigs: 129	CDD	Organism Group	
Molecule: DNA	Other genomes for species:			

<http://www.ncbi.nlm.nih.gov/sites/>