

Description

ANTI-CYANOBACTERIA RECOMBINANT ANTIBODY POLYPEPTIDE, GENE THEREOF, AND PREPARATION METHOD THEREOF

TECHNICAL FIELD

The invention relates to biotechnology in environment protection, especially a anti-cyanobacteria recombinant antibody polypeptide, gene thereof, and preparation method thereof.

BACKGROUND

Cyanobacteria proliferation and water pollution caused by water eutrophication are the biggest harm for the worldwide water environment that lead to great economic loss and irreparable damage towards the biosphere of the earth. Along with the accelerated industrialization and urbanization by the economic development of China, environment pollution and degeneration are severer and severer, and bionomic control on water environment has become a crucial problem to be faced and solved.

The present antibacterial agents are hardly effective against cyanobacteria, a prokaryocyte that belongs to Phylum X of cyanobacteria, Domain Bacteria. . Currently, there is only heavy metal chemical reagent, for example, copper sulfate and etc, which can control cyanobacteria. However, in practical use against cyanobacteria, the overdosed and repeated use of drugs due to their limited effects destroys against other beneficial algae, aquatic plants, organisms, and brings irreversible and perpetual wreck to the environment, as well as heavy metal agent residue with increasing amount in both the environment and agricultural products. Therefore, it is of great demand for the development of high-effective and safe anti-cyanobacterial drugs.

SUMMARY

In one aspect, the present invention provides genes, recombinant plasmids, and polypeptides of a novel anti-cyanobacteria recombinant antibody polypeptide; said polypeptides are capable of specifically killing cyanobacteria cells but not damaging other

algae cells, plants and animals. Another aspect of the present invention is to provide methods for preparing said anti-cyanobacteria recombinant antibody polypeptides.

Provided is a hybridoma with a depository number of CGMCC No.4783.

Also provided is a monoclonal antibody secreted by said hybridoma.

Also provided is an antibody mimics polypeptide with function of recognizing and binding cyanobacteria, which is an antigen-binding fragment of said monoclonal antibody, or a small molecular polypeptide that is designed on the basis of functional regions of said antigen-binding fragment of said monoclonal antibody.

Said antibody mimic polypeptide, its amino acid sequence sets forth in SEQ ID NO. 3.

Also provided is gene encoding any one of said antibody mimic polypeptides?

Said gene has a nucleotide sequence set forth in SEQ ID NO.2.

Also provided is an anti-cyanobacteria recombinant antibody polypeptide, constructed by operably and linearly connecting said antibody mimic polypeptide capable of recognizing and binding cyanobacteria to the C-terminus of colicin polypeptide, and said colicin is selected from colicin E1, Ia, Ib, A, B or N.

Said anti-cyanobacteria recombinant antibody polypeptide has an amino acid sequence set forth in SEQ ID NO.7.

Also provided is a gene encoding said any one of anti-cyanobacteria recombinant antibody polypeptide.

Said gene has a nucleotide sequence shown in SEQ ID NO.6.

Also provided is a recombinant expression vector, which comprises the gene encoding said anti-cyanobacteria recombinant antibody polypeptide.

Also provided is a method for preparing said anti-cyanobacteria recombinant polypeptide, with steps as follows: transducing the gene encoding the anti-cyanobacteria recombinant antibody polypeptide into an *E. coli* expression system, cultivating the transduced *E. coli*, and separating the polypeptide expressed by *E. coli*.

Also provided is the use of said anti-cyanobacteria recombinant antibody polypeptide on the control of water eutrophication.

The invention provides an anti-cyanobacteria recombinant antibody polypeptide, which is constructed from colicin polypeptide and an antibody mimic polypeptide against

cyanobacteria cell surface antigen. In said polypeptide molecule, the antibody mimetics against cyanobacteria cell surface antigen recognizes and binds to cyanobacteria, leading colicin of the molecule to attack the recognized cyanobacteria cell. With an antibody mimetic recognizing cyanobacteria cell surface antigen specifically, the polypeptide of the invention attacks cyanobacteria cells with no damage to other water organisms and no water pollution, thus being environment-protecting and safe on control of water cyanobacteria pollution. The antibody mimetic in the anti-cyanobacteria recombinant antibody polypeptide of the invention is an antibody mimetic capable of recognizing and binding to cyanobacteria, and is designed based on the amino acid sequences and nucleotide sequences of the light and heavy chain of the antigen-binding fragment (Fab) of the monoclonal antibody secreted by hybridoma CGMCC No.4783, such as mimic antibodies well known in the art, *e.g.*, single-chain antibody and small molecular antibody mimetics, *etc.* In the invention, the designed antibody mimetic just need to be capable of recognizing the antigen, as the only task of the antibody mimetics is to guide the recombinant polypeptide molecule to target cell surface, and build an ion channel on the cell membrane by colicin in the recombinant polypeptide, thereby resulting in the target cell death due to intracellular substance leak. Therefore, any recombinant polypeptides constructed by linking a colicin to an antibody mimetic that is designed on the basis of said monoclonal antibody and can recognize the target antigen, will fall into the claimed scope of the invention.

In an embodiment of the invention, the antibody mimetic is a 28-peptide antibody mimetics constructed by linking domains of V_H CDR1, V_H FR2 and V_L CDR3 of a monoclonal antibody against surface antigen of cyanobacteria protoplast (secreted by hybridoma CGMCC No.4783) in order of V_H CDR1- V_H FR2- V_L CDR3. Indoor and outdoor experiments proved that, the anti-cyanobacteria polypeptide constructed by said antibody mimetics has biological activity of moving around and automatically looking for the cyanobacteria cell surface antigen in water. The 28-peptide antibody mimetics has advantages that it is easily to be cloned, expressed, and has recognizing activity of the original antibody; it can kill cyanobacteria by guiding the colicin of the recombinant polypeptide to target cell membrane to build an ion channel. Said recombinant polypeptide has more ecological and environmental safety, compared with current drugs on controlling

water pollution (e.g., copper sulfate).

In the invention, the colicin for the construction of the anti-cyanobacteria recombinant antibody polypeptide may be selected from colicin E1, Ia, Ib, A, B or N that could form ion channel. The recombinant polypeptide constructed by anti-cyanobacteri antibody mimetics and colicin Ia is verified by examples of the invention. However, with the property of forming an ion channel on the cell membrane of cyanobacteria, any of these colicins according to conception of the invention may be linked linearly to anti-cyanobacteria antibody mimetics for constructing the recombinant polypeptide to carry out the invention, that is, to provide recombinant polypeptides that kill cyanobacteria.

The invention also provides preparation method of the recombinant plasmid comprising the gene of recombinant polypeptide, wherein the nucleotide sequence encoding antibody mimetics is inserted into the C-terminus of colicin by double strand oligonucleotides site-directed mutagenesis, to form the recombinant plasmid of the invention. Taking colicin Ia as an example, the recombinant plasmid of the invention is constructed by inserting the nucleotide sequence encoding antibody mimetics into the site after amino acid 626 of colicin Ia gene (the C-terminus). The original vector pSELECT-1 is purchased from Promega Corp., which contains genes of colicin Ia and Immunity protein. In Example 1 of the present invention, several pairs of primer sequences are designed for genes of antibody mimetics. The recombinant plasmid, e.g. pCHCcyanoMA1, is prepared according to the instruction of the kit from Strategene Corp. by double strand oligonucleotides site directed mutagenesis.

In another aspect of the invention, provided is a preparation method of said anti-cyanobacteria polypeptide of the invention, wherein a recombinant plasmid comprising a gene encoding anti-cyanobacteria recombinant antibody polypeptide is obtained by operably linking a gene encoding antibody mimetics with a colicin gene, and the obtained recombinant plasmid is transfected into an *E. coli* engineering bacteria to obtain a engineering bacteria cell that may produce anti-cyanobacteria recombinant antibody polypeptide. Said anti-cyanobacteria polypeptide of the invention is then obtained from separation and purification with ion-exchange column.

Said anti-cyanobacteria polypeptide of the invention may be used in the control of

water pollution. Drug combinations suitable for environment may be prepared by adding vectors, excipients, or other alternative components into the polypeptide of the invention, wherein the mechanism is as follows: antibody mimetics within the polypeptide that recognize cyanobacteria cell surface antigen guide colicin to the cyanobacteria cell membrane with the hydrophobic fragment of ion channel domain in colicin inserted into, to form an ion channel, thereby cyanobacteria cell death is incurred by intracellular substance leak.

Said anti-cyanobacteria polypeptide of the invention has an advantage of target specificity compared with traditional anti-cyanobacterial drugs, thus it does not attack other algae, plants and animal cells; toxicity and untoward effect far less than traditional anti-cyanobacterial drugs. Furthermore, since the ion channel is directly formed on cyanobacteria cell membrane to kill cyanobacteria cells, it is difficult for cyanobacteria cells to repair the geometrical damages on cell membrane by way of gene mutation and protein expression, in the short time of intracellular substance leak through anti-cyanobacteria polypeptide ion channel to cyanobacteria cell death. Therefore, it is with low probability that cyanobacteria cells obtains drug resistance of the anti-cyanobacteria polypeptide of the invention, which indicates there is a promising application prospect to said anti-cyanobacteria polypeptide of the invention.

Deposit information of hybridoma:

Deposit Number: CGMCC No.4783

Deposit Name: Monoclonal antibody hybridoma against cyanobacteria cell surface antigen

Depository Date: April 4, 2011

Depository institution: China General Microbiological Culture Collection Center, CGMCC

Address: No.3, Courtyard No. 1, Beichen West Road, Chaoyang District, Beijing

DESCRIPTION OF DRAWINGS

FIG.1 schematic illustration of the structure of recombinant plasmid pCHCeyanoMA1 comprising genes of antibody mimetics and colicin Ia

FIG. 2 schematic illustrates the structure of the anti-cyanobacteria recombinant antibody of present invention.

FIG.3 schematic illustration of the structure of recombinant plasmid pCHCcyanoMA2 comprising genes of antibody mimetics and colicin Ia with reversed N-C order in part.

FIG.4 the experiment results about the inhibition of the anti-cyanobacteria

polypeptide of the invention on *Microcystis aeruginosa* cultivated in liquid medium,

Wherein, **A**, blank control; **B**, 50 µg/ml of ampicillin; **C**, broad 50 µg/ml of spectrum pheromone (Ph-NM); **D**, 50 µg/ml of anti-cyanobacteria polypeptide (1); **E**, 50 µg/ml of anti-cyanobacteria polypeptide (2).

FIG.5 the experiment results about the inhibition of the anti-cyanobacteria polypeptide (1) of the invention on *Microcystis aeruginosa* cultivated in liquid medium

Wherein, **A**, result of treating for 48 hours; **B**, result of treating for 72 hours; **C**, result of treating for 96 hours.

Wherein, there are 6 flasks in each column, from left to right, namely as follows: **1**, control; **2**, 0.1 µg/ml of anti-cyanobacteria polypeptide (1); **3**, 0.5 µg/ml of anti-cyanobacteria polypeptide (1); **4**, 1 µg/ml of anti-cyanobacteria polypeptide (1); **5**, 5 µg/ml of anti-cyanobacteria polypeptide (1); **6**, 10 µg/ml of anti-cyanobacteria polypeptide (1).

FIG.6 the experiment results about the inhibition of the anti-cyanobacteria polypeptide (1) of the invention on *Scenedesmus* cultivated in liquid medium.

Wherein, there are 6 flasks in each column which are treated for 10 days, from left to right, namely as follows: **1**, control; **2**, 1 µg/ml of anti-cyanobacteria polypeptide (1); **3**, 5 µg/ml of anti-cyanobacteria polypeptide (1); **4**, 10 µg/ml of anti-cyanobacteria polypeptide (1); **5**, 15 µg/ml of anti-cyanobacteria polypeptide (1); **6**, 20 µg/ml of anti-cyanobacteria polypeptide (1).

FIG. 7 the experiment results about the inhibition of the anti-cyanobacteria polypeptide (1) of the invention on *Microcystis aeruginosa* and *Scenedesmus* cultivated in liquid medium. Wherein, the result of *Microcystis aeruginosa* is shown in the upper part, and that of *Scenedesmus* is shown in the lower part.

FIG. 8 the experiment results about the inhibition of the anti-cyanobacteria polypeptide

of the invention on *Microcystis aeruginosa*, *Anabaena*, *Chlorella*, and *Scenedesmus* cultivated in liquid medium.

Wherein, the left flask is control, and the right one is 35 $\mu\text{g/ml}$ of anti-cyanobacteria polypeptide (1). Wherein A: *Microcystis aeruginosa*; B: *Anabaena*; C: *Chlorella*; D: *Scenedesmus*.

FIG.9 the experiment result about the inhibition of the anti-cyanobacteria polypeptide of the invention on the natural water polluted by cyanobacteria.

Wherein, A, blank control; B, 1 $\mu\text{g/ml}$ of anti-cyanobacteria polypeptide (1); C, 5 $\mu\text{g/ml}$ of anti-cyanobacteria polypeptide (1); D, 10 $\mu\text{g/ml}$ of anti-cyanobacteria polypeptide (1); E, 0.7 $\mu\text{g/ml}$ of copper sulfate.

FIG.10 some respects of results of the inhibition experiment in Fig. 9

Wherein, A, changes of optical density in water; B, changes of Chlorophyll a in water; C, pH changes in water; D, changes of dominant algae population in water.

Wherein, D-1, Control group; D-2, the group treated by 1 $\mu\text{g/ml}$ of anti-cyanobacteria polypeptide (1); D-3, the group treated by 0.7 $\mu\text{g/ml}$ of copper sulfate.

FIG.11 microscopic observation (magnified by 400-fold) of the inhibition of the anti-cyanobacteria polypeptide on natural water polluted by cyanobacteria in Fig. 9

Wherein, changes of algae (*Anabaena*) that is treated by control, 1 $\mu\text{g/ml}$ of anti-cyanobacteria polypeptide (1) and 0.7 $\mu\text{g/ml}$ of copper sulfate in water on the first Day 1 (A), the second Day (B), the third Day (C), the fourth Day(D), the fifth Day(E) are shown respectively.

DETAILED DESCRIPTION

Example 1 Construction of plasmid expressing anti-cyanobacteria polypeptide and preparation of anti-cyanobacteria recombinant polypeptide

Material: Hybridoma CGMCC No.4783.

Step 1. Obtaining of the nucleotide sequence and amino acid sequence of antibody mimetics: the monoclonal antibody secreted by hybridoma was collected and sequenced by using conventional methods to obtain the sequence of heavy chain and light chain of

antigen-binding fragment Fab; the amino acid sequence of antibody mimetics was designed on the basis of the amino acid sequence of complementarity-determining regions, as shown in SEQ ID NO.3. The nucleotide sequence encoding said antibody mimetics is set forth in SEQ ID NO.2.

Step 2. The original vector is the plasmid pSELECT-1 (purchased from Promega Corp.), which carries genes of colicin Ia and Immunity protein (said genes are loaded in the lab where the experiments of this invention are conducted). The gene encoding anti-cyanobacteria antibody mimetics (nucleotide sequence shown in SEQ ID NO.2 in the sequences listing) was inserted after the site of I626 on the gene of colicin Ia by double-strand oligonucleotide Site-Directed Mutagenesis technology (QuickChange™ Kit, Stratagene Corp.), to obtain a recombinant plasmid pCHCcyanoMA1 (as shown in Fig. 1). The recombination plasmid was then transfected into the engineering bacteria of *E. coli* B834 (DE3) to prepare the polypeptide. Said anti-cyanobacteria polypeptide with amino acid sequence shown in SEQ ID NO.7 (hereinafter referred to as “anti-cyanobacteria polypeptide (1)”) and molecular weight about 70,000 was obtained.

The process of double-strand oligonucleotide site-directed mutagenesis was performed according to the manual of Stratagene QuickChange Site-Directed Mutagenesis Kit (catalog#200518).

1. Preparation of reactants for site-directed mutagenesis:

5 µl 10 x buffer

2 µl (10 ng) wild-type colicin Ia plasmid

1 µl (125ng) 5'-3' oligonucleotide primers as designed

1 µl (125ng) 5'-3' oligonucleotide primers as designed

1 µl dNTP

Double-distilled water 50 µl

1 µl pfu

(All is provided in the Kit except for the plasmid, primers and double-distilled water)

2. PCR amplification, amplification conditions: 20 cycles of denaturation at 95°C for 35 seconds, anneal at 53°C for 70 seconds, and extension at 68°C for 17 minutes;

3. 1 µl of endonuclease Dpn I was added to digest DNA chain (37°C, 1 hour); 1 µl of reactant and 50 µl of HMS174 competent cell were incubated together in ice for 30 minute, and then incubated in ice for 2 minute after a heat shock at 42°C for 45 seconds;

4. 0.5 ml of culture medium NZY was added, and cultivated by shaking at 37°C, 220 rpm for 1 hour. 50-100 µl of reactant was spread on LB medium plate plus 1% agar and 50 µg/ml of ampicillin for cultivating overnight at 37°C;

5. Colonies were picked up after cultivating for 18 hours. Plasmid was extracted by using extraction Kit of Qiagene Corp. or Gibco Corp., and then sequenced to confirm if the mutation is successful;

6. 50 ng of recombination plasmid was incubated with 50 µl of the prepared E. coli B834 (DE3) competent cells in ice for 30 minute, and heat shocked at 42°C for 30 second. 50-100 µl of the reactant was added with 0.5 ml of SOC culture medium, cultivated by shaking at 37°C, 220 rpm for 1 hour and spread on LB medium plate plus 1% agar and 50 µg/ml of ampicillin for incubating at 37°C for 12-16 hours. Afterwards, single colony was picked up;

7. Bacteria were amplified in 8-16 liters of LB medium at 250 rpm, 37°C for 6-8 hours. The bacterium was precipitated by centrifugation at 6000 g, 4°C for 20 minutes, resuspended with 50-80 ml of 50 mM Borate, 2mM EDTA (pH 9.0) at 4°C; 250 microliters of 0.2M PMSF was added and treated with ultrasonication at 4°C, 400W for 2 minutes. Bacterium debris was high-speed centrifugated at 4°C, 75,000 g for 1.5 hours. The supernatant was added with streptomycin sulphate to precipitate DNA, dialyzed overnight in dialysis bag with molecular weight of 15,000 as well as with 10 L of 50 mM Borate, 2mM EDTA, pH 9.0 buffer at 4°C, and then loaded on a CM column (Amersham Biosciences). The column was eluted by a buffer of 50 mM Borate, 0.3 M NaCl, 2mM EDTA, pH 9.0 to obtain the anti-cyanobacteria recombinant polypeptide (1) (See Fig. 2)

Sequences of double strand oligonucleotides designed for the preparation on genes of antibody mimetics of the plasmid pCHCcyanoMA1 are as follows:

pCHCcyanoMA1

1. 5'-3'

gcg aat aag ttc tgg ggt att TCT TAT TGG ATG CAG TGG GTT AAG CAA taa ata
aaa tat aag aca ggc

3'-5'

gcc tgt ctt ata ttt tat tta TTG CTT AAC CCA CTG CAT CCA ATA AGA aat acc cca
gaa ctt att cgc

2. 5'-3'

tat tgg atg cag tgg gtt aag caa AGA CCG GGG CAA GGG CTG GAG TGG ATT
GGC taa ata aaa tat aag aca ggc

3'-5'

gcc tgt ctt ata ttt tat tta GCC AAT CCA CTC CAG CCC TTG CCC CGG TCT ttg ctt
aac cca ctg cat cca ata

3. 5'-3'

ggg caa ggg ctg gag tgg att ggc CAG CAG TAT YGG TCT ACG CCC CCG TGG
ACG taa ata aaa tat aag aca ggc

3'-5'

gcc tgt ctt ata ttt tat tta CGT CCA CGG GGG CGT AGA CCA ATA CTG CTG gcc
aat cca ctc cag ccc ttg ccc

Example 2 Construction of a control plasmid expressing anti-cyanobacteria polypeptide and preparation of the control anti-cyanobacteria polypeptide

The original vector is the plasmid pSELECT-1 (purchased from Promega Corp.), which carries genes of colicin Ia, and Immunity protein (said genes are loaded in the lab where the experiments of this invention are conducted). The gene encoding anti-cyanobacteria antibody mimetics (nucleotide sequence shown in SEQ ID NO.4 in the sequence list) was inserted after the site of I626 on the gene of colicin Ia by double-strand oligonucleotide Site-Directed Mutagenesis technology (QuickChange™ Kit, Stratagene Corp.) to obtain recombinant plasmid pCHCcyanoMA2 (as shown in Fig. 3). The recombination plasmid was transfected into the engineering bacteria of *E. coli* B834 (DE3) to prepare the polypeptide. Said anti-cyanobacteria polypeptide whose sequence list is shown in SEQ ID NO.9 (hereinafter referred to as “anti-cyanobacteria polypeptide (2)”) and the molecular weight is about 70,000 was obtained.

Sequences of double strand oligonucleotides for the preparation on genes of antibody mimetics of the plasmid pCHCcyanoMA2 was designed and prepared according to the method in Example 1.

Example 3 Experiment results of inhibition of anti-cyanobacteria polypeptide on *Microcystis aeruginosa*

Experiment (1)

Step 1. Preparation of algae for experiment: *Microcystis aeruginosa* FACHB-978 is purchased from the Institute of Hydrobiology, Chinese Academy of Sciences.

Cultivation method of the algae: algae seed was inoculated with an initial cell density of 5×10^4 /mL into a 500 ml conical flask containing 250 ml of BG11 medium and placed into an illumination incubator, wherein the conical flask was covered with breathable film to prevent pollutions. Algae were cultivated at 25°C with illumination intensity of (2500±10%) lux and light/dark ratio of 12h/12h. Conical flasks were shaken and position-exchanged once every 2-3 hours. Samples were taken and tested once every 24 hours since the 5th day after the inoculation. The next step could be performed till the cell density reached about 10^6 /mL.

Step 2. 1 ml of prepared cyanobacteria liquid is added respectively into 5 tubes, namely, **A, B, C, D, and E**, and then inhibitors were added as follows: **A**, blank control; **B**, 50µg/ml of ampicillin; **C**, 50µg/ml of broad spectrum pheromone (Ph-AM1 disclosed in Patent Application No. CN200910092128.4); **D**, 50µg/ml of anti-cyanobacteria polypeptide (1); **E**, 50µg/ml of anti-cyanobacteria polypeptide(2). After shaking cultivation at low speed for 12hours in a incubator, the 5 tubes were respectively added with acridine orange (50nM) and propidium iodide (600nM) for incubating for 30 min; several microliters were sampled onto a slide to observe the alive stained cyanobacteria by a fluorescence microscope (Nikon 90i with filter of DM505 and DM565).

Results are shown in Fig. 4. It's manifested by results that only anti-cyanobacteria polypeptide (1) killed the cultivated *Microcystis aeruginosa*, whereas none of anti-cyanobacteria polypeptide (2), ampicillin and Ph-NM showed effects on the cultivated *Microcystis aeruginosa*.

Experiment (2)

Step 1. It was same as in Step 1 in Experiment (1).

Step 2. Anti-cyanobacteria polypeptides were added by concentration gradient into 6 of 500 ml conical flasks filled with algae. The anti-cyanobacteria polypeptide was added to *Microcystis aeruginosa* in doses of 0, 0.1, 0.5, 1, 5, and 10 µg/ml. The inhibition on cyanobacteria was observed every day.

Results were shown in Fig. 5. Because the concentration of anti-cyanobacteria polypeptide (1) was different in each group, the time of colour changing from green to light yellow of *Microcystis aeruginosa* liquid in each conical flask was proportional to the concentration of anti-cyanobacteria polypeptide (1). However, it was observed visually that *Microcystis aeruginosa* in all treatment could be killed effectively in 72 hours.

The above result showed that anti-cyanobacteria polypeptide (1) is capable of target killing the cultivated *Microcystis aeruginosa*, whereas anti-cyanobacteria polypeptide (2) hardly shows any effect.

Experiment (3)

Material: *Scenedesmus obliquus* FACHB-417 is purchased from the Institute of Hydrobiology, Chinese Academy of Sciences, and cultivated according to the method in Step 1 of Experiment (1).

Step 2. Anti-cyanobacteria polypeptide (1) was added into 6 of 500 ml conical flasks with algae by concentration gradient 0, 1, 5, 10, 15, and 20 µg/ml respectively. The inhibition on *Scenedesmus* was observed every day.

Results are shown in Fig. 6. Although the dose of anti-cyanobacteria polypeptide (1) was 10 times higher than that added in *Microcystis aeruginosa*, no killing effect of anti-cyanobacteria polypeptide (1) on *Scenedesmus* was observed visually during the experiment (190 hours).

Experiment (4)

0.1 mL of culture liquid is taken from each flask of the Experiment (1) - (3) and placed

in a blood counting chamber for observing and counting under a microscope. It's shown by the algae cell number in Fig. 7 that, compared to the control, 1 µg /ml of anti-cyanobacteria polypeptide (1) can kill *Microcystis aeruginosa* in 72 hours, whereas 1 µg/ml of anti-cyanobacteria polypeptide (1) hardly showed any effect against *Scenedesmus*.

Example 4. Experiment of anti-cyanobacteria polypeptide against *Microcystis aeruginosa*, *Anabaena*, *Chlorella* and *Scenedesmus* in liquid culture medium.

Microcystis aeruginosa, *Anabaena*, *Chlorella* and *Scenedesmus* are purchased from the Institute of Hydrobiology, Chinese Academy of Sciences, cultivated and treated according to the methods in Step 1, Experiment (1) of Example 3.

40 ml of the algae liquid with cell density of 10^6 /mL was distributed into two 300ml conical flasks, and added respectively with A: blank control liquid with same amount, B: 35 µg/ml of anti-cyanobacteria polypeptide (1), and cultivated for 4 days. The growth was observed.

Results were shown in Fig. 8. After 24 hours, the color of algae liquid of *Microcystis aeruginosa* and *Anabaena* turned light and became clear on the fourth Day, whereas the colour and turbidity of *Chlorella* and *Scenedesmus* liquid stayed constant. It's illustrated by the results that, anti-cyanobacteria polypeptide 1 inhibited the growth of the *Microcystis aeruginosa* and *Anabaena*, but did not inhibit the growth of *Chlorella* and *Scenedesmus*.

The above results proved that anti-cyanobacteria polypeptide (1) is capable of target killing cyanobacteria.

Example 5 Experiment of killing cyanobacteria in natural water by anti-cyanobacteria polypeptide

Natural water seriously polluted by cyanobacteria was taken from a shrimp breeding pool of Suzhou Yangcheng Lake, 25~35 L of the water was distributed into plastic breeding basins of 60 x 40 x 30 cm, with 10 min of air supply by bumping in every 20 min, temperature automatically adjusted to 27 °C, illumination intensity of (2500±10%) lux and light/dark ratio of 12h/12h. Water was sampled every day to detect

the pH, absorbance at 650 nm, and Chlorophyll a absorbance (See Fig. 9).

The absorbance at 650 nm was detected by placing the sample liquid into a visible spectrophotometer (See Fig. 10).

Chlorophyll a content was detected by using hot ethanol- freeze thawing- spectrophotometry method (See Fig. 10). The absorbances at 650 and 700 nm were detected by a spectrophotometer to calculate concentration of Chlorophyll a, with a formula of:

$$[Chla]=[12.12 (D664-D750)-1.58(D647-D750)-0.08(D630 - D750)]VE/VS \cdot d.$$

pH, absorbance at 650 nm, and Chlorophyll a absorbance in water of the control group remained high during the experiment, whereas those of the anti-cyanobacteria polypeptide (1) group and copper sulfate group decreased dramatically since the second day.

30 ml of water was sampled every day, fixed with formalin, and then concentrated for 10 times by centrifugation. 0.1 mL was taken into a blood counting chamber to be observed and counted under a microscope (See Fig. 11). It's displayed by microscopic examination that, on the first day, plenty of blossoming *Anabaena* could be seen in control group, anti-cyanobacteria polypeptide (1) group and copper sulfate group; whereas since the second day, *Anabaena* in anti-cyanobacteria polypeptide (1) group and copper sulfate group decreased and fragmented greatly in water; and no *Anabaena* could be seen in anti-cyanobacteria polypeptide (1) group and copper sulfate group on the fifth day, but plenty of *Anabaena* in control group were still alive.

Changes of algae shape was observed under a microscope, and various algae were counted to detect the dynamic changes about dominant population of floating plants in water (See Fig. 10). In control group, *Anabaena* and *Microcystis aeruginosa* were dominant algae all the time; in anti-cyanobacteria polypeptide (1) group, *Anabaena* and *Microcystis aeruginosa* decreased gradually while *Chlorella* became dominant; in copper sulfate group, *Anabaena* and *Microcystis aeruginosa* decreased gradually, and another *Cyanobacteria*, the *Oscillatoria*, became dominant.

Table 1 Number count and shape change of floating plants treated by anti-cyanobacteria polypeptide (/L)

	1 st Day of	2 nd Day	3 rd Day	4 th Day	5 th Day

	treating				
Control group	32745	26363 No change in algae shape	33194 No change in algae shape	41068 No change in algae shape	17954 No change in algae shape
1 mg/L of polypeptide	50394	43981 Most <i>Anabaena</i> became single cells or short chains of 2-5 cells	10624 Most of <i>Anabaena</i> become short chains of 1-5 cells, without long chain	5370 Most of <i>Anabaena</i> become short chains of 1-5 cells with little amount	3432 Almost no <i>Anabaena</i> cells
5 mg/L of polypeptide	43987	40828 Same as the change treated by 1 mg/L polypeptide	20744 Same as the change treated by 1 mg/L polypeptide	9890 Same as the change treated by 1 mg/L polypeptide	3680 Almost no <i>Anabaena</i> cells
10 mg/L of polypeptide	48101	41787 Same as the change treated by 1 mg/L polypeptide	23443 Same as the change treated by 1 mg/L polypeptide	14214 Same as the change treated by 1 mg/L polypeptide	4902 Almost no <i>Anabaena</i> cells
20 mg/L of polypeptide	45588	39194 Same as the change treated by 1	33453 Same as the change treated by 1	4800 Almost no <i>Anabaena</i> cells	3508 Almost no <i>Anabaena</i> cells

		mg/L polypeptide	mg/L polypeptide		
0.7 mg/L of copper sulfate	22930	1266 all <i>Anabaena</i> became single cells	499 Almost all the algae became organic detritus	305 Few algae	379 Few algae

Results are shown in Figs. 9-11 and Table 1. It's manifested by results that 0.7 ppm copper sulfate killed all kinds of algae without selection. However, anti-cyanobacteria polypeptide (1) has targeting inhibition on the growth of cyanobacteria (*Anabaena*, *Microcystis aeruginosa*) in water, but has little influence on the growth of other phytoplankton.

CLAIMS

1. A hybridoma with a depository number of CGMCC No.4783.
2. A monoclonal antibody secreted by said hybridoma of claim 1.
3. An antibody mimetic polypeptide with function of recognizing and binding to cyanobacteria, said antibody mimetic polypeptide being an antigen-binding fragment of said monoclonal antibody of claim 2, or a small molecular polypeptide that is designed on the basis of functional regions of said antigen-binding fragment and artificially expressed and isolated.
4. The antibody mimetic polypeptide according to claim 3, its amino acid sequence sets forth in SEQ ID NO. 3.
5. A gene encoding the antibody mimetic polypeptide of claim 3 or 4.
6. The gene according to claim 5, its nucleotide sequence sets forth in SEQ ID NO.2.
7. A anti-cyanobacteria recombinant antibody polypeptide, constructed by operably and linearly connecting said antibody mimetic polypeptide with function of recognizing and binding to cyanobacteria of claim 3 or 4 to the C-terminus of colicin polypeptide, said colicin is selected from colicin E1, Ia, Ib, A, B or N.
8. The anti-cyanobacteria recombinant antibody polypeptide according to claim 7, its amino acid sequence sets forth in SEQ ID NO.7.
9. A gene encoding the anti-cyanobacteria recombinant antibody polypeptide.
10. The gene according to claim 9, its nucleotide sequence sets forth in SEQ ID NO.6.
11. A recombinant expression vector comprising the gene encoding the anti-cyanobacteria recombinant antibody polypeptide.
12. A method of preparing the anti-cyanobacteria recombinant polypeptide of claim 7 or 8, comprising steps of: transforming the gene encoding the anti-cyanobacteria recombinant antibody polypeptide into an *E. coli* expression system, cultivating the transformed *E. coli*, and separating the polypeptide expressed by *E. coli*.
13. Application of said anti-cyanobacteria recombinant antibody polypeptide of claim 7 or 8 on control water eutrophication.

SEQUENCE LISTING

<110> Protein Design Lab, Ltd.

<120> ANTI-CYANOBACTERIA RECOMBINANT ANTIBODY POLYPEPTIDE,
GENE THEREOF, AND PREPARATION METHOD THEREOF

<130> PC11853/JJQ

<160> 9

<170> PatentIn version 3.3

<210> 1

<211> 1878

<212> DNA

<213> Nucleotide sequence encoding colicin Ia

<400> 1

```

atgtctgacc ctgtacgtat tacaaatccc ggtgcagaat cgctggggta tgattcagat      60
ggccatgaaa ttatggccgt tgatatttat gtaaacccctc cacgtgtcga tgtcttcat      120
ggtaccccg ctcgatggag ttcttcggg aacaaaacca tctggggcgg aaacgagtgg      180
gttgatgatt cccaacccg aagtgatatc gaaaaaaggg acaaggaaat cacagcgtac      240
aaaaacacgc tcagcgcgca gcagaaagag aatgagaata agcgtactga agccggaaaa      300
cgctctctg cggcgattgc tgcaaggga aaagatgaaa acacactgaa aacactccgt      360
gccggaaacg cagatgccgc tgatattaca cgacaggagt tcagactcct gcaggcagag      420
ctgagagaat acggattccg tactgaaatc gccggatatg acgccctccg gctgcataca      480
gagagccgga tgctgtttgc tgatgtgat tctcttcgta tatctcccc ggaggccagg      540
tcgttaatcg aacaggctga aaaacggcag aaggatgcgc agaacgcaga caagaaggcc      600
gctgatatgc ttgctgaata cgagcgcaga aaaggtattc tggacacccg gttgtcagag      660
ctggaaaaaa atggcggggc agcccttgcc gttcttgatg cacaacaggc ccgtctgctc      720
gggcagcaga cacggaatga cagggccatt tcagaggccc ggaataaact cagttcagtg      780
acggaatcgc ttaacacggc ccgtaatgca ttaaccagag ctgaacaaca gctgacgcaa      840
cagaaaaaca cgctgacgg caaacgata gttcccctg aaaaattccc ggggcgttca      900
tcaacaaatc attctattgt tgtgagcgg gatccgagat ttgccgtac gataaaaatc      960
acaaccagcg cagtcacga taaccgtgca aacctgaatt atctctgag ccattccggt     1020
ctggactata aacgcaatat tctgaatgac cggaatccgg tggtagacaga ggatgtggaa     1080

```

ggtgacaaga aaattataa tgctgaagtt gctgaatggg ataagttacg gcaaagattg	1140
cttgatgcc aaaaataaat cacctctgct gaatctgcgg taaattcggc gagaaataac	1200
ctcagtgcc gaacaaatga gcaaaagcat gcaaatgacg ctcttaatgc cctgttgaag	1260
gaaaaagaga atatacgtaa ccagcttcc ggcatcaatc agaagatagc ggaagagaaa	1320
agaaaacagg atgaactgaa ggcaacgaaa gacgcaatta attcacaac agagttcctg	1380
aatcagttt cagaaaaata tggcgcaaaa gctgagcagt tagccagaga gatggccggg	1440
caggctaaag ggaagaaaat acgtaatgtt gaagaggcat taaaacgta tgaaaagtac	1500
cgggctgaca ttaacaaaaa aattaatgca aaagatcgtg cagcgattgc cgcagccctt	1560
gagtctgtga agctgtctga tatatctct aatctgaaca gattcagtcg gggactggga	1620
tatgcaggaa aatttacaag tcttctgac tggatcactg agtttggtta ggctgtccgg	1680
acagagaact ggcgtcctct tttgttaa acagaaacca tcatagcagg caatgccgca	1740
acggctcttg tggcactggt cttcagtatt ctaccggaa gcgcttagg cattatcggg	1800
tatggtttac tgatggctgt caccggtgcg ctgattgatg aatcgcttgt ggaaaaagcg	1860
aataagttctggggtatt	1878

<210> 2

<211> 87

<212> DNA

<213> Gene of antibody mimetics against *Microcystis aeruginosa* surface antigen

<400> 2

tcttattgga tgcagtgggt taagcaaaga ccggggcaag ggctggagtg gattggccag	60
cagtattggt ctacgccgcc gtggacg	87

<210> 3

<211> 29

<212> PRT

<213> Amino acid sequence of antibody mimetics against *Microcystis aeruginosa* surface antigen

<400> 3

Ser Trp Trp Met Gln Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu
 1 5 10 15
 Trp Ile Gly Gln Gln Tyr Trp Ser Thr Pro Pro Trp Thr
 20 25
 <210> 4
 <211> 87
 <212> DNA
 <213> VHCDR1 N-C sequence in reversed order of gene of antibody mimetics
 against *Microcystis aeruginosa* surface antigen
 <400> 4
 cagatgtggt attcttgggt taagcaaaga ccggggcaag ggctggagtg gattggccag 60
 cagtattggt ctacgccgcc gtggacg 87
 <210> 5
 <211> 29
 <212> PRT
 <213> VHCDR1 N-C sequence in reversed order of amino acid sequence of
 antibody mimetics against *Microcystis aeruginosa* surface antigen
 <400> 5
 Gln Met Trp Tyr Ser Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu
 1 5 10 15
 Trp Ile Gly Gln Gln Tyr Trp Ser Thr Pro Pro Trp Thr
 20 25
 <210> 6
 <211> 1965
 <212> DNA
 <213> Nucleotide sequence encoding anti-cyanobacteria polypeptide I
 <400> 6
 atgtctgacc ctgtacgtat tacaatccc ggtgcagaat cgctggggta tgattcagat 60

ggccatgaaa ttatggccgt tgatatttat gtaaaccctc cacgtgtcga tgtctttcat 120
 ggtaccccg ctcgatggag ttcttcggg aacaaaacca tctggggcgg aaacgagtgg 180
 gttgatgatt ccccaacccg aagtgatatc gaaaaaaggg acaaggaaat cacagcgtag 240
 aaaaacacgc tcagcgcgca gcagaaagag aatgagaata agcgtactga agccggaaaa 300
 cgctctctcg cggcgattgc tgcaaggga aaagatgaaa acacactgaa aacactccgt 360
 gccggaaacg cagatgccgc tgatattaca cgacaggagt tcagactcct gcaggcagag 420
 ctgagagaat acggattccg tactgaaatc gccggatatg acgccctccg gctgcataca 480
 gagagccgga tgctgtttgc tgatgctgat tctcttcgta tatctccccg ggaggccagg 540
 tcgttaatcg aacaggctga aaaacggcag aaggatgcgc agaacgcaga caagaaggcc 600
 gctgatatgc ttgctgaata cgagcgcaga aaaggatttc tggacaccg gttgtcagag 660
 ctggaaaaaa atggcggggc agcccttgcc gttcttgatg cacaacaggc cgtctgctc 720
 gggcagcaga cacggaatga cagggccatt tcagaggccc ggaataaact cagttcagtg 780
 acggaatgc ttaacaggc ccgtaatgca ttaaccagag ctgaacaaca gctgacgcaa 840
 cagaaaaaca cgctgacgg caaacgata gttccctg aaaaattccc ggggcgttca 900
 tcaacaaatc attctattgt tctgagcggg gatccgagat ttgccgtac gataaaaatc 960
 acaaccagcg cagtcacga taaccgtgca aacctgaatt atctctgag ccattccggt 1020
 ctggactata aacgcaatat tctgaatgac cggaatccgg tggtagacaga ggatgtggaa 1080
 ggtgacaaga aaatttataa tgctgaagtt gctgaatggg ataagttacg gcaaagattg 1140
 ctgtatgcca gaaataaaat cacctctgct gaatctgcgg taaattcggc gagaaataac 1200
 ctcatgcca gaacaaatga gaaaagcat gcaaatgacg ctctaatgc cctgtgaag 1260
 gaaaaagaga atatacgtaa ccagcttcc ggcatcaatc agaagatagc ggaagagaaa 1320
 agaaaacagg atgaactgaa ggcaacgaaa gacgcaatta attcacaac agagttcctg 1380
 aatcagttt cagaaaaata tggcgcaaaa gctgagcagt tagccagaga gatggccggg 1440
 caggctaaag ggaagaaaat acgtaatgtt gaagaggcat taaaaacgta tgaaaagtac 1500
 cgggctgaca ttaacaaaaa aattaatgca aaagatcgtg cagcgattgc cgcagccctt 1560
 gagtctgtga agctgtctga tatatcgtct aatctgaaca gattcagtcg gggactggga 1620
 tatgcaggaa aatttacaag tcttgctgac tggatcactg agtttggtta ggctgtccgg 1680
 acagagaact ggcgtcctct tttgttaaa acagaaacca tcatagcagg caatgccgca 1740
 acggctcttg tggcactggg cttcagtatt cttaccggaa gcgcttagg cattatcggg 1800
 tatggtttac tgatggctgt caccgggtgc ctgattgatg aatcgcttgt ggaaaaagcg 1860

aataagttct ggggtatttc ttattggatg cagtgggtta agcaaagacc ggggcaaggg 1920
 ctggagtgga ttggccagca gtattggtct acgccgccgt ggacg 1965

<210> 7

<211> 654

<212> PRT

<213> Amino acid sequence of anti-cyanobacteria polypeptide 1

<400> 7

Ser Asp Pro Val Arg Ile Thr Asn Pro Gly Ala Glu Ser Leu Gly Tyr
 1 5 10 15
 Asp Ser Asp Gly His Glu Ile Met Ala Val Asp Ile Tyr Val Asn Pro
 20 25 30
 Pro Arg Val Asp Val Phe His Gly Thr Pro Pro Ala Trp Ser Ser Phe
 35 40 45

 Gly Asn Lys Thr Ile Trp Gly Gly Asn Glu Trp Val Asp Asp Ser Pro
 50 55 60
 Thr Arg Ser Asp Ile Glu Lys Arg Asp Lys Glu Ile Thr Ala Tyr Lys
 65 70 75 80
 Asn Thr Leu Ser Ala Gln Gln Lys Glu Asn Glu Asn Lys Arg Thr Glu
 85 90 95
 Ala Gly Lys Arg Leu Ser Ala Ala Ile Ala Ala Arg Glu Lys Asp Glu
 100 105 110
 Asn Thr Leu Lys Thr Leu Arg Ala Gly Asn Ala Asp Ala Ala Asp Ile
 115 120 125
 Thr Arg Gln Glu Phe Arg Leu Leu Gln Ala Glu Leu Arg Glu Tyr Gly
 130 135 140
 Phe Arg Thr Glu Ile Ala Gly Tyr Asp Ala Leu Arg Leu His Thr Glu
 145 150 155 160
 Ser Arg Met Leu Phe Ala Asp Ala Asp Ser Leu Arg Ile Ser Pro Arg
 165 170 175

Glu Ala Arg Ser Leu Ile Glu Gln Ala Glu Lys Arg Gln Lys Asp Ala			
180	185	190	
Gln Asn Ala Asp Lys Lys Ala Ala Asp Met Leu Ala Glu Tyr Glu Arg			
195	200	205	
Arg Lys Gly Ile Leu Asp Thr Arg Leu Ser Glu Leu Glu Lys Asn Gly			
210	215	220	
Gly Ala Ala Leu Ala Val Leu Asp Ala Gln Gln Ala Arg Leu Leu Gly			
225	230	235	240
Gln Gln Thr Arg Asn Asp Arg Ala Ile Ser Glu Ala Arg Asn Lys Leu			
245	250	255	
Ser Ser Val Thr Glu Ser Leu Asn Thr Ala Arg Asn Ala Leu Thr Arg			
260	265	270	
Ala Glu Gln Gln Leu Thr Gln Gln Lys Asn Thr Pro Asp Gly Lys Thr			
275	280	285	
Ile Val Ser Pro Glu Lys Phe Pro Gly Arg Ser Ser Thr Asn His Ser			
290	295	300	
Ile Val Val Ser Gly Asp Pro Arg Phe Ala Gly Thr Ile Lys Ile Thr			
305	310	315	320
Thr Ser Ala Val Ile Asp Asn Arg Ala Asn Leu Asn Tyr Leu Leu Ser			
325	330	335	
His Ser Gly Leu Asp Tyr Lys Arg Asn Ile Leu Asn Asp Arg Asn Pro			
340	345	350	
Val Val Thr Glu Asp Val Glu Gly Asp Lys Lys Ile Tyr Asn Ala Glu			
355	360	365	
Val Ala Glu Trp Asp Lys Leu Arg Gln Arg Leu Leu Asp Ala Arg Asn			
370	375	380	
Lys Ile Thr Ser Ala Glu Ser Ala Val Asn Ser Ala Arg Asn Asn Leu			
385	390	395	400
Ser Ala Arg Thr Asn Glu Gln Lys His Ala Asn Asp Ala Leu Asn Ala			
405	410	415	

Leu Leu Lys Glu Lys Glu Asn Ile Arg Asn Gln Leu Ser Gly Ile Asn
 420 425 430
 Gln Lys Ile Ala Glu Glu Lys Arg Lys Gln Asp Glu Leu Lys Ala Thr
 435 440 445
 Lys Asp Ala Ile Asn Phe Thr Thr Glu Phe Leu Lys Ser Val Ser Glu
 450 455 460
 Lys Tyr Gly Ala Lys Ala Glu Gln Leu Ala Arg Glu Met Ala Gly Gln
 465 470 475 480
 Ala Lys Gly Lys Lys Ile Arg Asn Val Glu Glu Ala Leu Lys Thr Tyr
 485 490 495
 Glu Lys Tyr Arg Ala Asp Ile Asn Lys Lys Ile Asn Ala Lys Asp Arg
 500 505 510
 Ala Ala Ile Ala Ala Ala Leu Glu Ser Val Lys Leu Ser Asp Ile Ser
 515 520 525
 Ser Asn Leu Asn Arg Phe Ser Arg Gly Leu Gly Tyr Ala Gly Lys Phe
 530 535 540
 Thr Ser Leu Ala Asp Trp Ile Thr Glu Phe Gly Lys Ala Val Arg Thr
 545 550 555 560
 Glu Asn Trp Arg Pro Leu Phe Val Lys Thr Glu Thr Ile Ile Ala Gly
 565 570 575
 Asn Ala Ala Thr Ala Leu Val Ala Leu Val Phe Ser Ile Leu Thr Gly
 580 585 590
 Ser Ala Leu Gly Ile Ile Gly Tyr Gly Leu Leu Met Ala Val Thr Gly
 595 600 605
 Ala Leu Ile Asp Glu Ser Leu Val Glu Lys Ala Asn Lys Phe Trp Gly
 610 615 620
 Ile Ser Tyr Trp Met Gln Trp Val Lys Gln Arg Pro Gly Gln Gly Leu
 625 630 635 640
 Glu Trp Ile Gly Gln Gln Tyr Trp Ser Thr Pro Pro Trp Thr
 645 650

<210> 8
 <211> 1965
 <212> DNA
 <213> Nucleotide sequence encoding antifungal polypeptide 2

<400> 8
 atgtctgacc ctgtacgtat tacaaatccc ggtgcagaat cgctggggta tgattcagat 60
 ggccatgaaa ttatggccgt tgatattat gtaaaccctc cacgtgtcga tgtcttcat 120
 ggtaccccg ctcgatggag ttcttcggg aacaaaacca tctggggcgg aaacgagtgg 180
 gttgatgatt ccccaaccg aagtgatat gaaaaagg acaaggaaat cacagcgatc 240
 aaaaacacgc tcagcgcgca gcagaaagag aatgagaata agcgtactga agccggaaaa 300
 cgcctctctg cggcgattgc tgcaaggga aaagatgaaa acacactgaa aacactccgt 360
 gccggaaacg cagatgccgc tgatattaca cgacaggagt tcagactcct gcaggcagag 420
 ctgagagaat acggattccg tactgaaatc gccggatatg acgccctccg gctgcataca 480
 gagagccgga tgctgttgc tgatgctgat tctcttcgta tatctccccg ggaggccagg 540
 tcgttaatcg aacaggctga aaaacggcag aaggatgcgc agaacgcaga caagaaggcc 600
 gctgatatgc ttgtgaata cgagcgcaga aaaggtattc tggacaccg gttgtcagag 660
 ctggaaaaaa atggcggggc agcccttgcc gttcttgatg cacaacaggc ccgtctgctc 720
 gggcagcaga cacggaatga cagggccatt tcagaggccc ggaataaact cagttcagt 780
 acggaatcgc ttaacacggc ccgtaatgca ttaaccagag ctgaacaaca gctgacgcaa 840
 cagaaaaaca cgctgacgg caaacgata gttccctg aaaaattccc ggggcgttca 900
 tcaacaaatc attctattgt tgtgagcgg gatccgagat ttgccgtac gataaaaatc 960
 acaaccagcg cagtcacga taaccgtgca aacctgaatt atcttctgag ccattccggt 1020
 ctggactata aacgcaatat tctgaatgac cggaatccgg tggtagaca ggatgtggaa 1080
 ggtgacaaga aaattataa tgctgaagt gctgaatggg ataagttacg gcaaagattg 1140
 ctgatgccca gaaataaaat cacctctgct gaatctgcgg taaattcggc gagaaataac 1200
 ctcatgccca gaacaaatga gaaaagcat gcaaatgacg ctcttaatgc cctgtgaag 1260
 gaaaaagaga atatacgtaa ccagcttcc ggcatcaatc agaagatagc ggaagagaaa 1320
 agaaaacagg atgaactgaa ggcaacgaaa gacgcaatta attcacaac agagttcctg 1380
 aaatcagttt cagaaaaata tggtgcaaaa gctgagcagt tagccagaga gatggccggg 1440

caggctaaag ggaagaaaat acgtaatgtt gaagaggcat taaaaacgta tgaaaagtac 1500
 cgggctgaca ttaacaaaaa aattaatgca aaagatcgtg cagcgattgc cgcagccctt 1560
 gagtctgtga agctgtctga tatatcgtct aatctgaaca gattcagtcg gggactggga 1620
 tatgcaggaa aatttacaag tcttctgtac tggatcactg agtttggtta ggctgtccgg 1680
 acagagaact ggcgtcctct tttgttaaa acagaaacca tcatagcagg caatgccgca 1740
 acggctcttg tggcactggt ctacagtatt ctaccggaa gcgcttagg cattatcggg 1800
 tatggtttac tgatggctgt caccggtgcg ctgattgatg aatcgcttgt ggaaaaagcg 1860
 aataagttct ggggtattca gatgtggtat tcttgggtta agcaaagacc ggggcaaggg 1920
 ctggagtga ttggccagca gtattgtct acgccgccgt ggacg 1965

<210> 9

<211> 654

<212> PRT

<213> Amino acid sequence of anti-cyanobacteria polypeptide 2

<400> 9

Ser Asp Pro Val Arg Ile Thr Asn Pro Gly Ala Glu Ser Leu Gly Tyr

1 5 10 15

Asp Ser Asp Gly His Glu Ile Met Ala Val Asp Ile Tyr Val Asn Pro

20 25 30

Pro Arg Val Asp Val Phe His Gly Thr Pro Pro Ala Trp Ser Ser Phe

35 40 45

Gly Asn Lys Thr Ile Trp Gly Gly Asn Glu Trp Val Asp Asp Ser Pro

50 55 60

Thr Arg Ser Asp Ile Glu Lys Arg Asp Lys Glu Ile Thr Ala Tyr Lys

65 70 75 80

Asn Thr Leu Ser Ala Gln Gln Lys Glu Asn Glu Asn Lys Arg Thr Glu

85 90 95

Ala Gly Lys Arg Leu Ser Ala Ala Ile Ala Ala Arg Glu Lys Asp Glu
 100 105 110
 Asn Thr Leu Lys Thr Leu Arg Ala Gly Asn Ala Asp Ala Ala Asp Ile
 115 120 125
 Thr Arg Gln Glu Phe Arg Leu Leu Gln Ala Glu Leu Arg Glu Tyr Gly
 130 135 140
 Phe Arg Thr Glu Ile Ala Gly Tyr Asp Ala Leu Arg Leu His Thr Glu
 145 150 155 160
 Ser Arg Met Leu Phe Ala Asp Ala Asp Ser Leu Arg Ile Ser Pro Arg
 165 170 175
 Glu Ala Arg Ser Leu Ile Glu Gln Ala Glu Lys Arg Gln Lys Asp Ala
 180 185 190
 Gln Asn Ala Asp Lys Lys Ala Ala Asp Met Leu Ala Glu Tyr Glu Arg
 195 200 205
 Arg Lys Gly Ile Leu Asp Thr Arg Leu Ser Glu Leu Glu Lys Asn Gly
 210 215 220
 Gly Ala Ala Leu Ala Val Leu Asp Ala Gln Gln Ala Arg Leu Leu Gly
 225 230 235 240
 Gln Gln Thr Arg Asn Asp Arg Ala Ile Ser Glu Ala Arg Asn Lys Leu
 245 250 255
 Ser Ser Val Thr Glu Ser Leu Asn Thr Ala Arg Asn Ala Leu Thr Arg
 260 265 270
 Ala Glu Gln Gln Leu Thr Gln Gln Lys Asn Thr Pro Asp Gly Lys Thr
 275 280 285
 Ile Val Ser Pro Glu Lys Phe Pro Gly Arg Ser Ser Thr Asn His Ser
 290 295 300
 Ile Val Val Ser Gly Asp Pro Arg Phe Ala Gly Thr Ile Lys Ile Thr
 305 310 315 320
 Thr Ser Ala Val Ile Asp Asn Arg Ala Asn Leu Asn Tyr Leu Leu Ser
 325 330 335

His Ser Gly Leu Asp Tyr Lys Arg Asn Ile Leu Asn Asp Arg Asn Pro
340 345 350

Val Val Thr Glu Asp Val Glu Gly Asp Lys Lys Ile Tyr Asn Ala Glu
355 360 365

Val Ala Glu Trp Asp Lys Leu Arg Gln Arg Leu Leu Asp Ala Arg Asn
370 375 380

Lys Ile Thr Ser Ala Glu Ser Ala Val Asn Ser Ala Arg Asn Asn Leu
385 390 395 400

Ser Ala Arg Thr Asn Glu Gln Lys His Ala Asn Asp Ala Leu Asn Ala
405 410 415

Leu Leu Lys Glu Lys Glu Asn Ile Arg Asn Gln Leu Ser Gly Ile Asn
420 425 430

Gln Lys Ile Ala Glu Glu Lys Arg Lys Gln Asp Glu Leu Lys Ala Thr
435 440 445

Lys Asp Ala Ile Asn Phe Thr Thr Glu Phe Leu Lys Ser Val Ser Glu
450 455 460

Lys Tyr Gly Ala Lys Ala Glu Gln Leu Ala Arg Glu Met Ala Gly Gln
465 470 475 480

Ala Lys Gly Lys Lys Ile Arg Asn Val Glu Glu Ala Leu Lys Thr Tyr
485 490 495

Glu Lys Tyr Arg Ala Asp Ile Asn Lys Lys Ile Asn Ala Lys Asp Arg
500 505 510

Ala Ala Ile Ala Ala Ala Leu Glu Ser Val Lys Leu Ser Asp Ile Ser
515 520 525

Ser Asn Leu Asn Arg Phe Ser Arg Gly Leu Gly Tyr Ala Gly Lys Phe
530 535 540

Thr Ser Leu Ala Asp Trp Ile Thr Glu Phe Gly Lys Ala Val Arg Thr
545 550 555 560

Glu Asn Trp Arg Pro Leu Phe Val Lys Thr Glu Thr Ile Ile Ala Gly

	565	570	575
Asn	Ala	Ala	Thr
Ala	Leu	Val	Ala
Leu	Val	Phe	Ser
Ile	Leu	Thr	Gly
	580	585	590
Ser	Ala	Leu	Gly
Ile	Ile	Gly	Tyr
Gly	Leu	Leu	Met
Ala	Val	Thr	Gly
	595	600	605
Ala	Leu	Ile	Asp
Glu	Ser	Leu	Val
Glu	Lys	Ala	Asn
Lys	Phe	Trp	Gly
	610	615	620
Ile	Gln	Met	Trp
Tyr	Ser	Trp	Val
Lys	Gln	Arg	Pro
Gly	Gln	Gly	Leu
	625	630	635
Glu	Trp	Ile	Gly
Gln	Gln	Tyr	Trp
Ser	Thr	Pro	Pro
Trp	Thr		
	645	650	