Simultaneous removal of harmful algal blooms and microcystins using microorganism and chitosan modified local soil

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ABSTRACT: Cyanobacterial harmful algal blooms (cyano-HAB) and microcystins (MCs) can cause potential threat to public health. Here, a method for simultaneous removal of cyano-HAB and MCs was developed using chitosan modified local soil (MLS) flocculation plus microorganism modified soil capping. The experiment was conducted in simulated columns containing algal water collected from Lake Taihu (China). More than 90% algal cells and intracellular MCs were flocculated and removed from water using chitosan-MLS and the sunk flocs were treated by different capping materials including Pseudomonas sp. An18 modified local soil. During 40 days’ incubation, dissolved MC-LR and MC-RR showed 10-fold increase in the flocculation only system. The increase of MC-LR and MC-RR in water was reduced by 30% and 70% in soil capping treatments, however, the total content of MCs in the sediment-water column remained similar with that in the control and flocculation only systems. In contrast, both dissolved MCs and total MCs were reduced by 90% in Pseudomonas sp. An18 modified soil capping treatment. The high performance of toxin decomposition was due to the combined effects of flocculation and MCs-degrading bacteria that embedded in the capping material, which prevents dilution of bacteria biomass, concentrates the algal cells, confines the released toxins, and enhances toxin biodegradation.
INTRODUCTION

Cyanobacteria harmful algal blooms (cyano-HABs) have increasingly occurred worldwide in various water sources, and one of the negative consequences is the production of cyanotoxins. Microcystins (MCs) are the frequently encountered cyanotoxins that contain over 90 variants. MCs are primarily retained in algal cells but released into waters in case of cells lysis, and can be accumulated in aquatic lives or transferred to higher trophic levels, which presents a potential threat to public health as tumor promoters. Therefore, rapid and sustainable degradation of MCs is imperative during the mitigation of cyano-HABs in natural waters.

Several methods including the application of algaecides, chemical and clay flocculation have been tested to remove cyano-HABs in natural waters,
whereas the methods often result in release of intracellular MCs into water.\textsuperscript{11} Removal of extracellular MCs using activated carbon, plant-mineral composite and phoslock\textsuperscript{®} are investigated,\textsuperscript{12-14} however, in addition to the high cost for large scale application, these methods only transfer MCs from water to other media,\textsuperscript{15} excessive amount of MCs could return back into water column. Only when MCs are degraded to less toxic products can the negative affects be diminished.\textsuperscript{16} Pan et al. propose a chitosan Modified Local Soil Induced Ecological Restoration (MLS-IER) technology to flocculate algal cells and then convert them into submerged vegetation in shallow waters.\textsuperscript{17} However, the released toxins could be diffused or mixed back to water column in deep water system where submerged macrophytes cannot be restored. MCs are degradable in natural waters by a wide range of microorganisms,\textsuperscript{5, 18} however, MCs may not be rapidly degraded by indigenous microorganisms if massive intracellular MCs are released due to the intensive lysis of the settled algal cells.\textsuperscript{19} Embedding exogenous MCs-degrading bacteria in the capping layer provide a potential alternative to enhance the biodegradation of MCs. However, challenges are presented due to the poor adaptability of MCs-degrading bacteria obtained through conventional artificial culture media (mostly in mineral salts), where the isolated bacteria often cannot form sustainable colonies and continuously degrade MCs in natural waters.\textsuperscript{20, 21}

Aiming at improving bacterial adaptability, we have previously proposed a method to isolate MCs-degrading bacteria using concentrated water extract of sediment as natural culture media.\textsuperscript{21} It was proved that the obtained strain
Pseudomonas sp. An18 displayed enhanced and continuous degradation of MCs in collected Taihu lake water under continuous addition of MCs. We anticipate that if Pseudomonas sp. An18 is embedded into the capping materials, it may greatly enhance the decomposition of MCs following MLS flocculation, since both the toxins and the bacteria can be well confined under the capping layer and the increased availability of MCs as well as the improved bacterial adaptability may work together to enhance the biodegradation of MCs. So far, there are no previous studies of embedding sustainable microorganisms into the capping materials and its combination to MLS flocculation-capping method for simultaneous removal of cyano-HABs and degradation of MCs.

In this study, microorganisms modified capping materials were prepared by embedding Pseudomonas sp. An18 into Lake Taihu local soil. Simulated column experiment containing Lake Taihu water was conducted to evaluate the effect of chitosan-MLS flocculation followed by Pseudomonas sp. An18 modified soil capping on the removal of algal cells and degradation of MC-RR and MC-LR. The objective is to study the effect and the mechanism on cyano-HABs removal and MCs degradation using MLS flocculation and capping method.

MATERIALS AND METHODS

Toxin Standards and Reagents. MCs standards (MC-RR and MC-LR) were obtained from Taiwan Algal Science Inc (purity ≥95% by HPLC). Enkephalin (internal standard) was purchased from Sigma-Aldrich.

Soil and modifier. The soil used in flocculation experiment was collected from
the bank of Meiliang Bay of Lake Taihu at Wuxi, and was washed with distilled water and dried for 10 h at 90 °C. The soil was sieved through 180 meshes (74 μm). Chitosan (solid) was obtained from Qingdao Haisheng Bioengineering Co.Ltd. The molecular weight of chitosan was 500000, and the deacetylation degree was 83.6%. Deionized water was prepared using a Milli-Q filtration system (Millipore, Bedford, MA, USA).

**Preparation of Chitosan Modified Soil.** Chitosan was dissolved by adding 100 mg chitosan to 10 ml of 1% HCl and mixing until all chitosan was dissolved. This solution was diluted with deionized water to obtain a working solution of 1 mg/mL. To modify the soils, a certain volume of chitosan working solution was added to soil suspension. The mixture was well stirred and then ready for use in the flocculation experiment. The SEM picture of chitosan modified soil was presented in our previous study.22

**Preparation of Microorganism Modified Soil.** Local soil was collected from the shore of Lake Taihu (Meiliang Bay, Wuxi, China) and sieved through 40 meshes. *Pseudomonas sp.* An18 was isolated through water extract of sediment followed by concentration treatment according to the method described in our previous study.21 The bacteria were incubated in mineral salts medium containing MC-RR (50 μg/L) and MC-LR (6 μg/L) for 3 days, then harvested by centrifugation at 10,000 rpm for 5 min and washed twice with 50 mM phosphate buffer (pH 7.0). The local soil were dried in muffle furnace at 250 °C for 3 h, then the resultant soil (10g) was mixed with 150 ml *Pseudomonas sp.* An18 suspensions.
(5\times 10^{13} \text{ CFU/mL}) in 250 \text{ ml erlenmeyer flask} with shaken at 150 \text{ rpm for 4h. The suspension solution was centrifuged at 7,000 \text{ rpm} and the precipitate was washed 3 times with sterile water, finally, 1\text{M NaCl} was added to liberate the weakly retained bacterial cells. The modified soil was capable of embedding 10^{15} bacterial cells/g soil. The evaluation of \textit{Pseudomonas sp. An18} proliferation in the modified soil was conducted in mineral salt medium with addition of filter sterilized MCs every 2 days, the microorganism modified soil was vortexed to detach the embedded \textit{Pseudomonas sp An18} from soil and the bacteria number were counted using plate counting method in mineral salt medium containing MC-RR (50 \mu \text{g/L}) and MC-LR (6 \mu \text{g/L}).

\textbf{Incubation Experiment.} The experiment was conducted in 2000 \text{ mL plexiglass columns} with diameter of 8.4 cm. The Taihu local soil was autoclaved and used as sediment in the column. A volume of 1500 \text{ mL cyano-HABs water} were collected from Meiliang Bay, Lake Taihu and added into the columns (the initial algae concentration was 1.38\times 10^{10} \text{ cells/L}, concentration of MC-RR and MC-LR were 5.14 and 0.84 \mu \text{g/L}, respectively). Twelve columns containing the same amount of sediment and algal water were prepared. Three of them were used as controls where no flocculants or capping materials were added. Chitosan modified local soil (2 \text{ mg/L chitosan and 100 mg/L soil}, this optimized dosage was obtained according jar test presented in Figure S1 and S2) was added to all the rest of the columns and then followed by three different capping treatments. No capping materials were added in the flocculation only systems (three columns). Among the
rest of the columns, the flocculated algal flocs were capped with 1 cm local soil
(three columns) or 1 cm *Pseudomonas sp.* An18 modified soil (three columns).
The twelve columns were incubated at 25 °C with an alternating cycle of 12h light
and 12h darkness for 40 days.

**Sample Collection and Preparation.** Water samples were collected at 1 cm
below the water surface every 3 days. After filtration through 0.22 µm membrane,
the water was spiked with 100 µl enkephalin internal standard (10 µg/L) to reduce
the matrix effects during measurement of MCs. The algal cells were enumerated
every 3 days. After 40 days’ incubation, in order to assess the distribution of
MC-RR and MC-LR in different parts of the sediment-water systems, the
overlaying water was carefully taken off from each column and filtered through
0.45 µm membrane to separate the suspended algal cells. The sediment in the
columns were sliced into three sections: 0-1 cm (capping layer), 1-2 cm (flocs
layer), and 2-6 cm (sediment layer). For the control system, there was no capping
and flocs layer. The capping layer (local soil or *Pseudomonas sp.* An18 modified
soil) and sediment layer were freeze-dried and extracted in 0.1 M EDTA-Na$_4$P$_2$O$_7$
for three times. After centrifugation at 4000 rpm, the aqueous extractions were
passed through SPE cartridge (300 mg, 3 ml, Waters), and the eluted solution was
evaporated under nitrogen and dissolved in 1 ml distilled water for UPLC-MS/MS
analysis. Suspended algal cells and algae flocs were freeze-dried and extracted
using 40% (v/v) methanol solution with ultrasonication (300W, 5 mins) three
times, followed by the evaporation of the elution containing MCs from SPE
cartridge, as was described previously.\textsuperscript{26} The total content (µg) of MC-LR and MC-RR in the whole systems after 40 days’ incubation were calculated according to the toxin concentration in filtered water, suspended algae cells and that in sediment, flocs layer, and capping layer (dry weight).

**MCs Determination.** Quantitative analysis of MCs was performed using a LC-MS system equipped with electrospray ionization tandem mass spectrometry (UPLC-ESI-MS/MS) (Acquity UPLC, Quattro Premier XE, Waters, USA). Sample separation was carried out with UPLC system and Acquity UPLC BEH C\textsubscript{18} column (i.d. 2.1 mm×50 mm, particle size 1.7 µm, pore size 130 Å, Waters). The column oven was kept at 40 \textdegree C and the injection volume was 10 µL. The flow rate used was 0.2 ml/min. Water and acetonitrile were used as mobile phases, the organic phase was linearly increased from 45 % to 60 % in 2.5 min, then increased to 100% in 1.5 min and held for 0.5 min, and finally brought back to 45% and held for 3 min until the next injection. The mass spectrometer was operated in positive mode electrospray ionization in multiple-reaction monitoring mode. Source temperature and desolvation gas temperature were held at 120 and 350 \textdegree C, respectively. The measured recoveries of the method ranged between 91.0-103.2 %. The detection limit was 6.0 and 3.0 ng/L for MC-RR and MC-LR, respectively.

**Analysis of MCs Biodegradation Products.** Biodegradation products of the toxins were analyzed using matrix-assisted laser desorption/ionization-time of flight mass spectrometry (MALDI-TOF MS, BIFLEX III, Bruker Inc., USA).
After 40 days incubation, MCs from the sediment of local soil capped system and *Pseudomonas sp.* An18 modified soil capped system were extracted. The extracted MCs solution (containing MCs and its degradation products) was dried with nitrogen, then dissolved in distilled water. MALDI-TOF-MS instrument was equipped with a nitrogen laser operating at 337 nm, and a two-state ion source operating in the delayed extraction mode. Briefly, 2 μL of a saturated solution of a-cyano-4-hydroxycinnamic acid (in 3:2 v/v acetonitrile-0.1% trifluoroacetic acid) were premixed with 2 μL of the MCs solution. Afterward, 2 μL of this mixture were applied to the sample plate, and air-dried at 24 °C. Measurements were performed at an acceleration voltage of 20 kV using reflector mode, allowing the determination of monoisotopic mass values. Each spectrum refers to the sum of 100-200 individual laser shots.

**Soil Characterization.** Micrographs of soil and microorganism modified soil were characterized using a XL-30 scanning electron microscope (Philips Corp., The Netherlands).

**Statistical Analysis.** The t-test was used to determine significant differences among different measurements, a significance was assumed when P < 0.05.

**RESULTS**

**Soil Characterization.** Scanning electron microscope (SEM) images illustrated a smooth surface and discrete particles of local soil particles (Figure 1A), while modification by *Pseudomonas sp.* An18 substantially roughened the surface of soil with widely distributed ridges and large density of *Pseudomonas sp.* An18
Evaluation of *Pseudomonas sp.* An18 carrying capability revealed the modified soil was capable of embedding $10^{15}$ bacterial cells/g soil (Figure S3). Proliferation tests showed that the biomass of *Pseudomonas sp.* An18 in the microorganism modified soil dramatically increased after 1 days’ lag phase and yielded stable CFU counts (nearly $10^{19}$ CFU/mL) after 3 days (Figure 2), suggesting the soil was favorable for *Pseudomonas sp.* An18 growth when MCs were supplied.

**Figure 1.** SEM photographs of local soil and *Pseudomonas sp.* An18 modified soil. (A) local soil (200×magnification), (B) *Pseudomonas sp.* An18 modified soil (200×magnification), (C) *Pseudomonas sp.* An18 modified soil (3000×magnification). The arrow in 1C pointed to *Pseudomonas sp.* An18.

![Graph](image)
Figure 2. Proliferation of *Pseudomonas sp.* An18 on the microorganism modified soil in mineral salts medium containing MCs. Data represent the mean values of triplicate ± SD.

**Algae Removal.** In the chitosan-MLS flocculation only systems, the algae removal rate reached 99% within 1 day, however, re-suspension of algal cells occurred in the following days and the removal rate reduced to about 90% at day 40 (Figure 3). The re-suspension was effectively avoided when local soil or *Pseudomonas sp.* An18 modified soil was used to cap the sunk algae flocs, where the removal rate remained at 99% throughout the 40 days’ period. A small algae removal (maximum of 13% at day 40) was also observed in the control systems (Figure 3).

**Figure 3.** Changes in algae concentration in water columns. Data represent the mean values of triplicate ± SD.

**MCs in Water Column.** The initial concentration of dissolved MC-RR in the Taihu water sample was 5.14 µg/L. In the control systems, an obvious increase of dissolved MC-RR was observed from day 9, which reached the maximum level of
108.14 μg/L between day 24 to 40 that is 20-fold higher than the initial level (Figure 4A-a). Compared to the control systems, the increase of dissolved MC-RR was reduced by 50% in chitosan-MLS flocculation only systems (Figure 4A-b), 70% in flocculation-local soil capping systems (Figure 4A-c), and 96% in flocculation plus *Pseudomonas* sp. An18 modified soil capping systems (Figure 4A-d), respectively. The dissolved MC-LR showed a similar trend with that of MC-RR among different treatments. The initial concentration of dissolved MC-LR in Taihu water sample was 0.84 μg/L. In the control systems, dissolved MC-LR was quickly increased from day 9 to day 24 and reached 15.45 μg/L between day 24 to 40 (Figure 4B-a). The MC-LR release was reduced, at day 40, to 10.64 μg/L by chitosan-MLS flocculation only treatment (Figure 4B-b), 6.65 μg/L by flocculation plus local soil capping treatment (Figure 4B-c), and 0.49 μg/L by flocculation plus *Pseudomonas* sp. An18 modified soil capping treatment (Figure 4B-d), respectively. The diffusion of released MCs from sediment to water were largely reduced in soil capping treatment (Figure 4A-c, Figure 4B-c) and entirely disappeared in microorganism modified soil treated systems (Figure 4A-d, Figure 4B-d).
**Figure 4.** Changes of dissolved MC-RR (A) and MC-LR (B) concentration in the surface water of different systems. a, b, c, d indicate different treatment. Data represent the mean values of triplicate ± SD.

**Distribution of MCs.** Measurements of MCs in water, algae cells, sediment, flocs, and capping layer illustrated the distribution of MCs in different parts of the sediment-water column systems. For the control systems, among the total MC-RR (2360 µg), 51% of the intracellular MC-RR (1209 µg) was released into the water, 48% was still left in suspended algal cells (1135 µg) and 1% in the sediment at day 40 (Figure 5A-a). Chitosan-MLS flocculation transferred 85% of the MC-RR into the sunk algal flocs, and left 10% in suspended algae cells and 4% in water (Fig. 5A-b). In the flocculation and soil capping treatment, MC-RR was almost entirely removed from suspended algae cells and water (2% left), 78% and 19% were transferred into the flocs layer and capping layer (Fig. 5A-c), respectively. However, the total content of MC-RR in the flocculation only and flocculation-soil capping systems remained similar to that of the control. In
contrast, nearly 90% MC-RR in the whole sediment-water system was degraded in chitosan-MLS flocculation plus *Pseudomonas sp*. An18 modified soil capping systems (Fig. 5A-d), and the dissolved MC-RR left in the water (6 µg) was reduced by 201, 13 and 5-fold compared to the control, flocculation only and flocculation-soil capping systems, where the original total content of toxins were similar. The distribution of MC-LR followed a similar pattern with that of MC-RR. The dissolved MC-LR in the control was 356 µg (Figure 5B-a) and was reduced to 23 and 10 µg in flocculation (Figure 5B-b) and flocculation-soil capping systems (Figure 5B-c), respectively, but the total MC-LR content (ranged from 620 to 650 µg) remained similar among these three systems (Figure 5B). In contrast, the total MC-LR content was reduced by 90% and dissolved MC-LR by 99% in flocculation plus *Pseudomonas sp*. An18 modified soil capping treatment (Figure 5B-d) compared to the control systems.

**Figure 5.** Distribution of total content of MC-RR (A) and MC-LR (B) in different parts of the sediment-water system of each treatment after 40 days’ incubation.
Data represent the mean values of triplicate.

**Biodegradation Products of MCs.** The MS spectrum of MC-RR revealed a major ion at m/z 1038.27 after 40 days incubation in soil capped sediment (Fig. 6a), in contrast, two major degradation products of MC-RR (MW=332.92 and 707.27) were observed in *Pseudomonas sp*. An18 modified soil capped layer (Fig. 6b), corresponding to the reduced intensity of MC-RR (MW=1038.27). Similarly, in comparison to the great intensity of MC-LR in soil capped sediment (Fig. 6c), the predominate ion in *Pseudomonas sp*. An18 modified soil capped layer converted to MW=332.73 and 301.65 (Fig. 6d), of which the MC-LR ion at m/z 995.83 almost disappeared.

![Mass spectra](image)

**Figure 6.** Mass spectra of MC-RR in the sediment of soil capping layer after 40 days incubation (a), MC-RR and its biodegradation products after 40 days in the sediment of *Pseudomonas sp*. An18 modified soil capped layer (b), MC-LR in the sediment of soil capping layer after 40 days incubation (c), MC-LR and its
biodegradation products after 40 days in the sediment of Pseudomonas sp. An18 modified soil capped layer (d).

**DISCUSSION**

**Effect of Flocculation and Capping on Algae and Toxin Removal.** Algal cells tend to suspend and float in water because of the negatively charged cell surface, low specific gravity and specific buoyant structure. Chitosan, a cellulose-like polyelectrolyte biopolymer, is derived from the alkaline deacetylation of crustacean chitin, which possesses several intrinsic characteristics of coagulants and flocculants such as high cationic charge density, long polymer chains, bridging of aggregates and precipitation. The high content of positively charged amine groups in the chitosan structure facilitates electrostatic interaction between polymer chains and negatively charged algal cells,\(^{27, 28}\) which caused rapid mitigation of cyano-HABs (Figure 3). For toxin producing cyano-HABs, most toxins are contained inside the cells during the bloom, which will be released and contributed to the dissolved MCs when the cells lysis occur.\(^{29}\) In the chitosan-MLS flocculation only systems, although 99% algal cells were removed from the water column at day 1 (Figure 3), dissolved MC-RR (Figure 4A-b) and MC-LR (Figure 4B-b) began to increase in the water column at day 12 and reached the maximum concentration during day 24 to 40. This result suggests that flocculation alone can only remove algae cells from water column, however, dissolved MCs toxins may be significantly released if the algae flocs are not further treated. The algae flocs may be resuspended back to the water and
continue to grow under the field conditions especially in shallow waters.\textsuperscript{30} By using soil capping after the flocculation, the increase of dissolved MC-RR (Figure 4A-c) and MC-LR (Figure 4B-c) from the sediment was postponed about a week and reduced by 74\% and 40\% compared to the flocculation only treatment at day 40. Soil capping not only physically sealed the algal cells in sediment and depress the diffusion of dissolved MCs from the sediment to the water, but also significantly promoted the cells lysis (Figure S4). Algal cell lysis is crucial for the reduction of algae recruitment,\textsuperscript{30} however, it will result in the release of intracellular MCs.\textsuperscript{29} If the released toxin is only buried under the capping layer but not degraded, it will be diffused into the water in a postponed period (Figure 4A-c, Figure 4B-c). Previous reports found that once the capping layer was saturated, the excessive toxins could eventually be released into water.\textsuperscript{31} The problems mentioned above can be solved if the capping layer is highly efficient for MCs degradation. Indeed, in the flocculation plus microorganism modified capping systems (1 cm), both MC-RR and MC-LR were almost entirely decomposed in the water (Figure 4A-d, 4B-d) and sediment (Figure 5A-d, 5B-d) throughout the tested period. By increasing the thickness of the \textit{Pseudomonas sp. An18} modified capping layer, the microorganism may obtain more time to degrade the upcoming MCs molecules under the capping layer.

\textbf{Adaptability of Incubated Microorganisms in Natural Environment.} In cyano-HABs waters, degradation of MCs by the indigenous MCs-degrading microorganism is one of the main pathways for toxins removal.\textsuperscript{32, 33} In our
simulated systems, the maximum dissolved MC-RR and MC-LR reached 108.14 µg/L (Figure 4A-a) and 15.45 µg/L (Figure 4B-a) during 40 days’ incubation, suggesting long time was required for removal of the toxins by the native microorganism only. This agreed well with the report that MCs can not be rapidly removed relying on the native MCs-degrading microorganisms after abrupt release of MCs under collapse of severe cyano-HABs.  

Although exogenous MCs-degrading bacteria have been widely studied under laboratory conditions, one of the problems is that most of the incubated bacteria can hardly be sustainable in natural waters due to the poor bacterial adaptability.  

The conventional approach of obtaining MCs-degrading bacteria is collection of microorganism communities from MCs contaminated sediment or water, followed by enrichment and isolation in artificial medium (mostly mineral salts medium) where MCs were supplied as the sole carbon and nitrogen source.  

The discrepancy of chemical property between artificial media and natural environment, in addition to the competition with indigenous microorganism communities in natural environment often result in poor bacterial adaptability of the incubated bacteria and limited toxins removal efficiency.  

In order to improve bacterial adaptability in natural waters, an isolation strategy using concentrated natural medium was proposed in our previous study, by which the obtained *Pseudomonas sp*. An18 showed enhanced adaptability and sustainability for MCs degradation in sampled Taihu Lake water when continuous addition of MCs were degraded.
In flocculation only systems, dramatic increase of dissolved MC-RR and MC-LR occurred after nearly 14 days incubation (Figure 4A-b, Figure 4B-b), despite the diffusion of released MCs can be largely reduced in soil capping treatment (Figure 4A-c, Figure 4B-c), the total residual toxins in the experimental systems were not significantly reduced compared to control (Figure 5), in contrast, the increase of dissolved MCs had entirely disappeared (Figure 4A-d, Figure 4B-d), the residual MC-RR (Figure 5A-d) and MC-LR (Figure 5B-d) in the whole sediment-water systems displayed 90% reduction in *Pseudomonas sp*. An18 microorganism modified soil treated systems, suggesting the adaptability of the incubated *Pseudomonas sp*. An18 strains and biodegradation of MCs in the experimental systems. In this study, *Pseudomonas sp*. An18 was enriched from water extracts of sediment thus experienced selective pressure and subsequent isolation in concentrated water extracts of sediment, which showed similar nutrient condition, component and PH with the real environment and led to adaptation of the re-incubated microbes in the simulated natural water.

**Concentrated Microorganisms and MCs by in-situ flocculation-capping.** Biodegradation of MCs by exogenous MCs-degrading bacteria under natural conditions requires not only the adaptation of bacteria, but also the reaction opportunity between the incubated bacteria and toxins. Direct addition of isolated bacteria into natural waters may encounter serious dilution problem. Embedding bacteria into soil can keep relative high bacterial biomass on the soil particles therefore enhance its reactivity with the toxins. Moreover, in natural
cyano-HABs waters, the algal cells together with MCs are predominant in the surface of water column, which may also impair the degradation reaction between toxins and bacteria. In comparison to 51% MC-RR (Figure 5A-a) and 56% MC-LR (Figure 5B-a) that were released and diluted in the water column in the control systems, 96% MC-RR (Figure 5A-c) and 98% MC-LR (Figure 5B-c) were confined under the capping layer in flocculation-soil capping systems, hence provided concentrated MCs under the capping layer, suggesting that if MCs-degrading bacteria can be embedded in the capping material, the biodegradation reaction may be enhanced due to the simultaneously raised toxins concentration and bacteria biomass. In this study, the bacteria modified soil loaded nearly $10^{15}$ *Pseudomonas sp.* An18 cells/g modified soil (Figure S3), indicating 1 cm thickness capping layer (approximately 140 g) can embed $1.4 \times 10^{17}$ $Pseudomonas sp.$ An18 cells, therefore created a micro-environment with higher bacterial biomass. Furthermore, proliferation of *Pseudomonas sp.* An18 in the soil led to $10^5$ increases of bacteria number during 6 days’ incubation (Figure 2), suggesting the soil favored bacteria inhabitation and proliferation.

**Degradation of MCs by flocculation-microorganism Modified Soil Capping.**

In addition to providing adequate time for bacteria to establish biodegradation ability before they were mixed back to water column, flocculation-microorganism modified soil capping treatment also retained the released MCs within the confined sediment layer. The concentrated toxins and bacteria in-situ hence increased the availability of toxins to *Pseudomonas sp.* An18, as the
biodegradation efficiency was positively proportional to the initial toxins concentrations within a certain range,\(^{10, 19}\) therefore, 90% degradation of MC-RR (Figure 5A-d) and MC-LR (Figure 5B-d) from the whole sediment-water systems were triggered. This is likely due to that when MCs are released and diffused into the microorganisms modified soil, the toxins are metabolized as carbon and nitrogen source by Pseudomonas sp. An18. This was confirmed by the analysis of MCs biodegradation products by MALDI-TOF mass spectra, which revealed that in Pseudomonas sp. An18 modified soil capped systems, the degradation products of MC-RR (MW=332.92 and 707.27, Fig 6b) and MC-LR (MW=332.73 and 301.65, Fig 6d) were evident as the concentration of MC-RR and MC-LR decreased in the experimental systems (Figure 5A-d and Figure 5B-d). Moreover, the appearance of MW=332, which was recognized as the Adda residues,\(^{38}\) in both the biodegradation products of MC-RR and MC-LR, indicating that the cleavage of Adda-Arg peptide bond, by which the toxicity of MCs can be substantially reduced.\(^{39}\) Zhang et al.(2010) studied the biodegradation pathway of MC-RR bacterium Sphingopyxis sp. USTB-05, the results suggested that Adda-Arg peptide bond of MC-RR was cleaved, then a hydrogen and a hydroxyl were combined onto the NH\(_2\) group of Adda and the carboxyl group of arginine,\(^{35}\) during which the MW=332 were indicated as the Adda residues.\(^{35, 38}\) The examined MW=332.92 in the present test implying the similar biodegradation pathway of MCs by Pseudomonas sp. An18, which need further investigations.

In this column experiments, under the co-effects of prevented bacterial
dilution (embedding *Pseudomonas sp. An18* in the microorganism modified soil),
concentrated MCs concentration in-situ (flocculation and microorganism modified
soil capping), as well as enhanced bacteria adaptability, the reduction of dissolved
MCs (Fig. 4) in water and the decomposition of MCs in the sediment can be
achieved (Fig. 5).

In order to understand unambiguously the role of incubated *Pseudomonas sp. An18* and to exclude the biodegradation of MCs by the indigenous microorganism,
autoclaved local soil instead of in-situ sediment was used as sediment layer in the
column. Under field conditions, natural aquatic sediments have a wide range of
MCs-degrading microorganisms, it is possible that the combined effects of
indigenous bacteria in the natural sediment and *Pseudomonas sp. An18* in the
capping layer may influence the efficiency of MCs degradation. In addition, this
study was carried out in columns under stationery condition, therefore the
re-suspension of sunk algal cells (Figure 3) and diffusion of the released MCs
(Figure 4) were depressed through 1cm thickness soil or microorganism modifies
soil capping. However, capping as a measure for lake restoration is inevitably
influenced by the complicated conditions in the field, such as vertical transport
and re-suspension generated by bioturbation and horizontal transport induced by
the wind and current. Although re-suspension of algal flocs has been studied
under simulated laboratory conditions, the complicated impact of field condition
needs further studies at various scales including lab, mesocosm, whole water
ponds, and open waters.
Environmental Implications. The cyano-HABs in eutrophic lakes are usually followed by toxins release due to the decomposition of algal cells. The dissolved MCs in many cyano-HABs lakes can remain a few µg/L, but the toxins can be accumulated in zooplankton, aquatic organisms and eventually bioaccumulated in human bodies through food chain. Only when MCs are degraded to less toxic products, can the negative affects be diminished. In this study, the flocculation-capping technology converged algal cells coupled with intracellular toxins under the capping layer, accelerated algae decomposition which reduced the recruitment of algae cell, and potentially promote the transfer of released nutrients from algae cells to vegetation if submerged vegetation seeds were included in the capping material in shallow waters. Different methods should be jointly used to achieve both short term and long term effects. For instance, flocculation is important to remove toxic algae from water and create the light conditions for the vegetation restoration in shallow water systems. The problems of floc resuspension and algal toxin release associated with flocculation method can be compromised by using microorganism modified soil/sand capping method. The capping method is also important for maintaining the water clarity for an extended period so that it may create a window period for the restoration of submerged vegetation, where the remediation processes can be relayed into a longer term. Once the excess nutrients are redistributed from water to sediment and from algae to vegetation, the restoration of a healthy food chain and the aquatic ecology may become possible by using the nutrients as a resource. As the
released toxins were biodegraded, it is possible to diminish the risk of toxin accumulation in aquatic vegetation, organisms and animals, which may further impact on food web and public health, which needs further multidisciplinary long time studies.

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Supporting Information Available

Jar test for cyano-HABs removal using chitosan modified soil (Figure S1 and S2)

Assessment of *Pseudomonas* sp. An18 carrying capacity of microorganism modified soil. Embedding capability of *Pseudomonas* sp. An18 on the microorganism modified soil (Figure S3). Morphology characterization of algal cells (Figure S4). This information is available free of charge via the Internet at http://pubs.acs.org.

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