

Growth, photosynthesis and some related metabolites as suitable selection criteria for the copper tolerance of *Ankistrodesmus falcatus*.

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Abstract:

This investigation was conducted to study the effects of copper on growth and metabolism of *Ankistrodesmus falcatus* isolated from sewage water at El-Dare Treatment Plant. The alga was grown in BG-11 medium containing different cu concentrations (0.0, 0.5, 5, 10, 50, 100, 150, 200 and 250 $\mu\text{m Cu}^{+2}$). The test organism was left to grow for 12 days under the various cu levels. The effect of copper on the growth, photosynthetic pigments, protein metabolism (soluble, insoluble), antioxidant compounds (phenolic, Proline) were determined. The cell number and dry weight were increased highly significantly up to 10 $\mu\text{m cu}^{+2}$, then the cell number and dry weight reduced gradually by increasing the doses of the copper. The concentration of photosynthetically active pigment peaked up to 10 $\mu\text{m cu}^{+2}$, Then they remained more or less unchanged up to 50 $\mu\text{m cu}^{+2}$, after this it was reduced. The content of soluble carbohydrates decreased dramatically up to 10 $\mu\text{m cu}^{+2}$, and then it was increased smoothly as the cu concentration increased in culture media. Cu Stress induced a significant changes in the soluble protein up to the level of 10 $\mu\text{m Cu}^{+2}$, then a high significant accumulation was obtained which peaked at the highest concentration of Cu^{+2} (250 μm). *The insoluble and total proteins remained more or less unchanged at the all Cu concentration used.* Aminoacids content accumulated progressively and irregularly by cu treatment. While proline content decreased by about 65.6% at the level of 10 $\mu\text{m cu}^{+2}$ concentration. It is on other hand increased by 261.7% at the level of 250 $\mu\text{m cu}^{+2}$ in relation to control value. Phenolic compounds seemed to be not affected by copper treatment.

Keyword: Cu toxicity- oxidative stress – *Ankistrodesmus falcatus*

Introduction:

Heavy metal contamination of soil is a serious problem for the environment. Some metals are required for plant growth and development, but some of these metals are very toxic at high concentration (Rout and Das, 2003; Broadley *et al.*, 2007). Various sources of Cu^{+2} including industrial and domestic wastes, agricultural practices, copper mine drainage, copper-based pesticides, have contributed to a progressive increase in copper concentrations in varied environments (Ma *et al.*, 2003 and Andrade *et al.*, 2004). Copper can be good and evil to algae as it is an essential micronutrient for algal growth, participating an important biological reactions as an enzymatic cofactor and electron carrier in the photosynthetic and respiratory processes (Andrade *et al.*, 2004). but at high concentration, it becomes highly toxic (Dewez *et al.*, 2005). Growth inhibition

and chlorosis are common symptoms of metal phyto-toxicity in several algae, in which photosynthesis is probably the most affected metabolic process. Copper reduces growth as well as photosynthetic and respiratory activities (Nale-wajko and Olaveson, 1995, Ali *et al.*, 2006). The photosynthetic apparatus is particularly susceptible to this cation, resulting in a decrease in the activity of photo-system II and electron transfer rate (Fernade and Henriques, 1991; Malick and Mohn, 2003 and Perales-Vela *et al.*, 2007). Because of its redox properties, copper induces oxidative stress by generating reactive oxygen species like superoxide and hydroxyl radicals via Haber-Weiss and Fenton reactions. Toxicity of copper might also result from the oxidation of sulphhydryl groups of enzymes leading to their inhibition (Teissire and Guy, 2000). Oxidative stress directly

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damages proteins, amino acids, nucleic acid, and membrane lipids often leading to alterations in cell structure and mutagenesis (Nagalakshmi and Prasad, 1988). Under acute conditions, however, the toxic effects of the pollutants may overwhelm the antioxidant defenses. This may result in cell death or the shut down of all cellular machinery. Toxicity may result in diverse effects, which depend on the type of algae, the nature and concentration of the metal, and the environmental conditions accompanying heavy metal stress (Satoh *et al.*, 2005). The aim of this investigation was to study the effects of different concentrations of Cu on growth and some related metabolites of *Ankistrodesmus falcatus* isolated from sewage water at El-Dare Treatment plant. This organism is unicellular green alga which can provide important information on the toxic effects of a pollutant on general metabolic processes and often used as an indicator of contamination.

Material and methods:

Algal species and culturing:

Ankistrodesmus falcatus was isolated from sewage water at El-Dare Treatment plant at Sohag district, Egypt. The alga was grown in Bold's basal medium according to (Bischoff and Bold, 1963). under the conditions of fluorescent illumination (2500 lux) and room temperature (25±2). Filtered dry air was allowed let to bubble in the culture vessels to provide carbon dioxide and to prevent settling of algal cells. In this experiment, the algae was grown in BG-11 medium containing different copper concentrations (0.0, 0.5, 5, 10, 50, 100, 150, 200 and 250 µm CU⁺².) For 12 days at the same conditions mentioned above. Each treatment was made in three replicates. At the end of incubation period, the algal cells were harvested and used for growth and metabolic determinations.

Determination of growth parameters:

While was used for isolation of eukaryotic and green algae.

Cell number:

The cell count of control and treated cultures was measured by Hemacytometer, 0.1 mm deep, having improved Naubauer ruling (A.O. Spencer "Bright fine"). The count was expressed as cells / ml algal suspension.

Dry weight:

Dry weight was determined according to (Utting, 1985). by filtering Culture aliquots (50 mL) through Whatman GF/C filters. The filters were dried and weighed.

Photosynthetic pigment extraction:

Chlorophyll a, b and Carotenoid were extracted in 100% methanol at 65°C and their contents were determined spectrophotometrically (SPEKOL 11, CARL ZEISS, JENA, GERMANY) according to (Metzner *et al.*, 1965).

Biochemical determinations:

Carbohydrate content estimation:

Carbohydrate content was determined in aqueous (soluble carbohydrate) and in HCl solutions (total carbohydrate) with anthrone sulphuric acid reagent according to (Fales, 1951). using glucose as a standard. The blue green color developed was measured at the 620 nm using spectrophotometer.

Estimation of total free amino acids:

Total free amino acids were determined according to (Moore *et al.*, 1958). The quantity of total free amino acids was calculated as µ gm./mg. dry weight.

Estimation of proline:

Free proline content of algal suspension was determined according to (Bates *et al.*, 1973). Briefly, 10 mL of algal suspension was centrifuged and the pellete was extracted in 5 mL of aqueous 3% sulfosalicylic acid for 3 h. The extract was centrifuged at 4000 rpm for 10 min. Two mL of the supernatant were mixed with 2 ml of fresh acid ninhydrin solution and 2 mL glacial acetic acid in a test tube for 1 h at 100°. The tubes were cooled, and the mixture was extracted with 4 mL toluene. The extract was vigorously stirred for 20 seconds. Therefore, the chromophore-containing toluene was aspirated from the aqueous phase, and its absorbance was measured at 520 nm. Proline was used as a standard.

Protein content estimation:

Protein content was determined according to (Lowry *et al.*, 1951). The alga of 10 mL of algal suspension was extracted in distilled-water (soluble protein) and in NaOH (Total protein) for 2 h at 90°C. The extract was centrifuged and the supernatants were pooled. The water-soluble protein was estimated by the Folin-phenol reagents and measured spectrophotometrically (SPEKOL

11, CARL ZEISS, JENA, GERMANY.) Bovine serum albumin was used as a standard.

Determination of phenolic compounds:

Phenolic compounds content was determined according to (Dai *et al.*, 1994). 0.1 gm of fresh tissue. Algal samples were homogenized with a plastic pestle in an Eppendorf tube containing 1 ml. phosphate buffer 0.1 M pH= 7.0. The homogenate was centrifuged in an Eppendorf microcentrifuge at 12800 for 10 min. Aliquots of 50 μ L. were added to a reaction mixture containing 3% of sodium carbonate and 0.3 M Folin reagent in a final volume of 1 ml. The reaction mixture was incubated for 2 hr. at room temperature and the absorbance at 765 nm. Total phenolic compounds were expressed as Nan equivalents of Gallic acid using a calibration curve prepared with 10-50 μ M of Gallic acid.

Nitrate reductase assay in vivo:

For in vivo assay of nitrate reductase, the method of (Jaworski, 1971). was used. Algal cells of 10 ml algal suspension of diuron-treated alga and untreated were precipitated and incubated in anaerobic dark conditions for 1 h in 5 ml of 0.1 M K-phosphate (ph=7.5) containing 50 mm KNO₃ and 1% (v/v) n-propanol at 28 °c. The reaction was stopped by boiling in water bath for 5 min and then centrifuged. The supernatant of one ml sample mixed well with two ml 1% w/v sulphonilamide in 1N hcl and two ml 0.1 % w/v N- (1- naphthyl) ethylenediaminedihydrochloride in distilled water. The absorbance was measured by using spectrophotometer (SPEKOL 11, CARL ZEISS, JENA, GERMANY) at 540 nm. Nitrate reductase activity was expressed as μ g NO₂/ml algal suspension h-1.

Results:

The preliminary data showed that the growth of *Ankistrodesmus falcatus* completely died beyond 250 μ M Cu⁺². The data in Fig (1) revealed that, The cell number and dry weight were increased highly significantly up to 10 μ M Cu⁺², Then the growth criteria decreased gradually by the further increase in the Cu. The harmful effect of Cu was much more pronounced at the higher doses (200 μ M Cu⁺² and 250 μ M Cu⁺²). In a conformity the photosynthetic pigments also stimulated up to the level 10

μ M Cu⁺², Then they remained more or less unchanged up to 50 μ M Cu⁺², then a gradual reduction in photosynthetic pigments was obtained which was much more obvious at the highest level of the Cu (200 μ M Cu⁺² and 250 μ M Cu⁺²).

the content of soluble carbohydrates decreased dramatically up to 10 μ M Cu⁺², then it increased smoothly as the Cu concentration increased in culture media. The highest reduction was recorded at the level of 10 μ M Cu⁺² (about 55.3%) while the highest increase was observed at the higher concentration of Cu (9% over the control value), the insoluble fraction increased progressively up to 10 μ M Cu⁺², then it remained more or less unchanged, with a general tendency to decrease especially at the higher level of Cu. The reduction was not exceeded than 9% at the level of 250 μ M Cu⁺². Consequently the total carbohydrate remained around those of control even at the highest concentration of Cu.

Cu stress induced insignificant changes in the soluble protein up to the level of 10 μ M Cu⁺², then a high significant accumulation was obtained which peaked at the highest concentration of Cu (250 μ M Cu⁺²) it was increased by 32.97% over the control at the level of 250 μ M Cu⁺². The insoluble and total proteins remained more or less unchanged at the all Cu concentration level used.

Aminoacids content accumulated progressively and irregularly by Cu treatment. The highest accumulation was recorded at the level of 50 μ M Cu⁺² (149.24% over the control), and the lowest accumulation was reported at the lower levels (0.5 and 5 μ M Cu⁺²).

proline content decreased dramatically up to 10 μ M Cu⁺² then a sharp and quick accumulation was exhibited beyond these level and continue to be increased as the Cu increased in the culture media. Thus while proline content decreased by about 65.6% at the level of 10 μ M Cu⁺² concentration. It is on other hand increased by 261.7% at the level of 250 μ M Cu⁺² in the relation to control. The activity of nitrate reductase increased slightly up to 5 μ M Cu⁺², then a sharp increase in the activity was recorded at the level of 10 μ M Cu⁺². (174, 03% of control), there after it decreased highly significantly as the Cu concentration increased in the culture media.

This reduction was much more obvious at the higher doses of cu. At the level of 250 μM Cu^{+2} , the percent reduction was about 56% in relation to the control value.

Cu stress did not affect the concentration of phenolic compounds even at the highest concentration of Cu. The content of phenolic compound remained mostly around those of control.

Copper stress resulted in the appearance of new polypeptides bands, consequently while the number of polypeptides in control was 15 polypeptides, they were 35 at 250 μM Cu^{+2} (more than 2 folded which fluctuated between the low molecular weight polypeptides to the high molecular weight polypeptides. For example 37,34 K Da and 22 K Da polypeptide appeared only under copper stress also the higher molecular weight polypeptides (208 K Da to 71 K Da) appeared only in copper treated *Ankistrodesmus falcatus*.

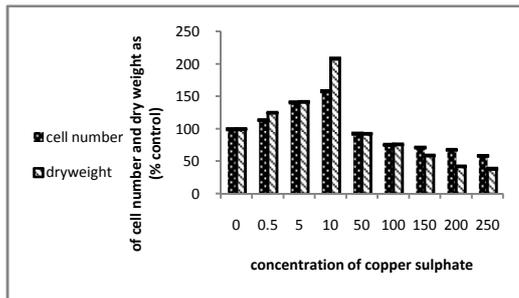


Fig. (1): Effect of different concentrations of copper ion on the percentage control of cell number and dry weight as (%control) of *Ankistrodesmus falcatus*.

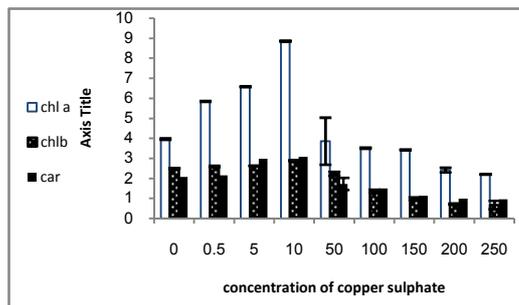


Fig. (2): Effect of different concentration of copper ion on photosynthetic pigments (chlorophyll a, chlorophyll b and carotenoids) (mg/g fresh weight) of *Ankistrodesmus falcatus*.

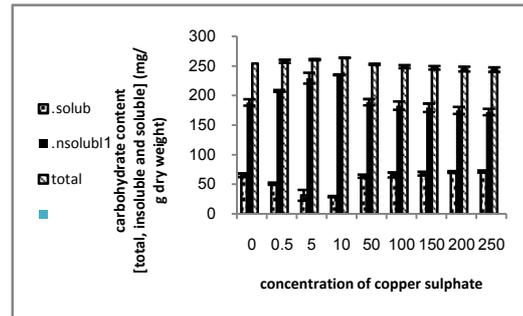


Fig. (3): Effect of different concentrations of copper ion on the percentage control of carbohydrate contents of *Ankistrodesmus falcatus*.

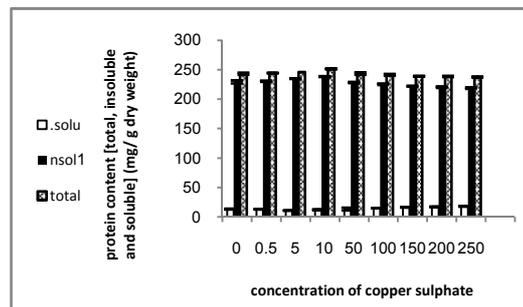


Fig. (4): Effect of different concentrations of copper ion on protein content [total, insoluble and soluble] (mg/g dry weight) of *Ankistrodesmus falcatus*.

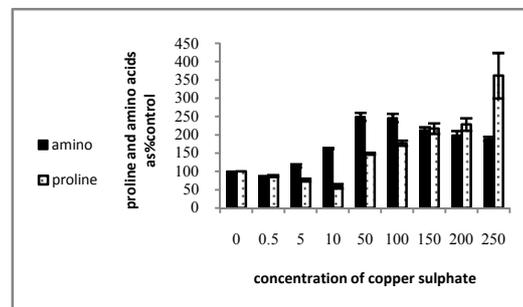


Fig. (5): Effect of different concentrations of copper ion on the percentage control of proline and amino acids of *Ankistrodesmus falcatus*.

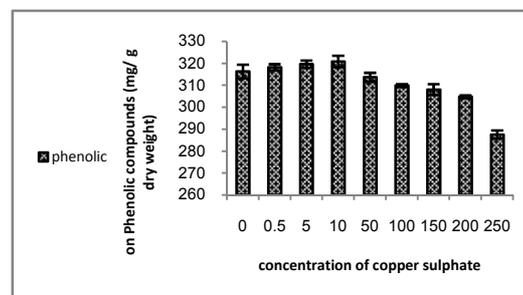


Fig. (6): Effect of different concentrations of copper ion on phenolic compounds ($\mu\text{g/mg}$ dry weight) of *Ankistrodesmus falcatus*.

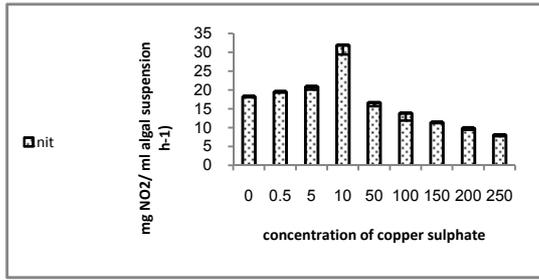


Fig. (7): Effect of different concentrations of copper ion on nitrate reductase activity in *Ankistrodesmus falcatus*.

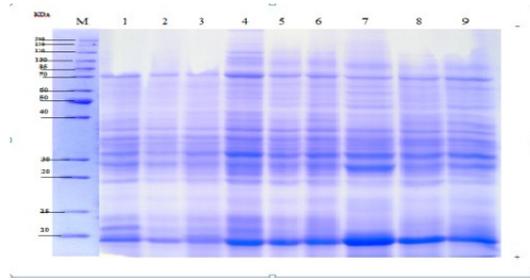


Fig. 8: Coomassie-stained SDS-10% polyacrylamide gel of polypeptides of *Ankistrodesmus falcatus* treated with different concentrations of copper (0.0 µM, lane 1; 0.5 µM, lane 2; 5 µM, lane 3; 10 µM, lane 4 and 50 µM, lane 5; 100 µM, lane 6; 150 µM, lane 7; 200 µM, lane 8 and 250 µM, lane 9).

No. Of band	M. Wt.	Cu conc. (µM)									
		Contr.	0.5	5	10	50	100	150	200	250	
1	208					+	+	+	+	+	+
2	165					+	+	+	+	+	+
3	141							+	+	+	+
4	118					+	+	+	+	+	
5	117									+	+
6	100						+	+	+	+	+
7	96			+	+		+	+	+	+	+
8	92									+	+
9	89										
10	85			+	+	+	+	+	+	+	+
11	74					+	+	+	+	+	+
12	73		+	+	+	+	+	+	+	+	+
13	71						+	+	+	+	+
14	70	+	+		+					+	+
15	69					+				+	
16	67			+	+	+	+	+	+	+	+
17	66					+	+			+	
18	64	+	+	+	+	+	+	+	+	+	+
19	63								+		+
20	62										+
21	60	+		+	+				+		
22	59						+	+	+	+	+
23	58	+	+	+	+					+	+
24	53								+	+	
25	50						+	+			
26	49		+	+		+	+	+	+	+	+
27	46	+	+	+	+	+	+	+	+	+	
28	45	+	+	+	+	+	+	+	+	+	+
29	42	+	+	+	+	+	+	+	+	+	+
30	41										
31	40							+		+	+
32	39	+									
33	38								+	+	+
34	37	+	+	+	+	+	+	+	+	+	+
35	36	+	+	+	+	+	+	+	+	+	+
36	35										
37	34					+	+	+	+	+	+
38	33	+	+			+	+	+	+	+	+
39	32			+	+						
40	31					+	+	+	+	+	+
41	28	+	+	+	+	+	+	+	+	+	+
42	26									+	+
43	24										+
44	22						+	+	+	+	+
45	21										
46	19	+	+	+	+	+	+	+	+	+	+
47	18	+									+
48	16										
49	15	+	+	+	+	+	+	+	+	+	+
50	Total	15	14	17	22	24	26	29	35	35	

Table (1): Molecular weights of protein bands detected in *Ankistrodesmus falcatus* treated with heavy metals. Data were obtained by Total Lab version 1.10 electrophoresis data system program

Discussion:

The most important feature of this work that *Ankistrodesmus falcatus* can survive up to 250 $\mu\text{m cu}^{+2}$. More ever growth criteria (cell number, dry weight and photosynthetic pigments estimated progressively up to 10 $\mu\text{m cu}^{+2}$, the dry matter content nearly doubled at 10 $\mu\text{m cu}^{+2}$, then gradually reduction was obtained in this growth parameter at copper concentration increased in culture media. This mean that:

- I- *Ankistrodesmus falcatus* tolerate the relatively high concentration of copper in nature.
- II- It posses high ability to picked up a lot amount of cu under the polluted conditions.
- III- consequently, it possesses a high capacity to clean up the copper polluted places (ponds and canals), accordingly *Ankistrodesmus falcatus* can used as an excellent biological cleaner in polluted water.

Such biphasic responses to copper were also revealed by other investigators using some plants and algae. Fageria (2002). reported that copper application significantly increased dry matter yield of upland rice and common bean. However, dry matter yields of the two species were decreased at the highest Cu concentrations.

Gao *et al.*, (2008). working on *Jatropha curcas* L. Seedlings found that the biomass in leaves increased slightly at lower Cu concentrations. There was a correlation between increasing Cu concentration and reduced seedling stems and roots mass. Lara and Luca (2005). reported that the peach root stock *prunus cerasifera* Mr.S. 2/5 plantlets grown in vitro on media containing either 10 or 50 μm of cuSO_4 did not show any visible signs of copper toxicity. The negative effect of Cu^{+2} on photosynthetic pigments in algae had been reported by (Dewez *et al.*, 2005; Perales-Vela *et al.*, 2007).

Copper tolerance of *Ankistrodesmus falcatus* was founded to be linked with the stable levels of carbohydrates and proteins even at the highest copper levels used. This mean that copper made *Ankistrodesmus falcatus* able to unregulate the interaction between carbon and nitrogen which might used as suitable marker for the copper

tolerance of *Ankistrodesmus falcatus*. this was linked with the marked and progressive accumulation of soluble protein which detoxify the copper toxicity (phytochelation) along with this amino acids and proline increased unexpectedly by copper. Amino acids approach to fold as the severe doses of copper. More ever proline content was fold at the highest doses of copper which indicated that this huge accumulation of amino acids and proline could play the major role in detoxification of copper and might responsible for increasing the copper tolerance of *ankistrodesmus falcatus*.

Such enhancement of amino acids and including proline content by metal stress is a common metabolic response of algae (El-Naggar, 1993). The mechanisms of proline action are not fully understood, but it has been suggested that increased accumulation permits osmotic adjustment, as well as provides protection for enzymes (Sharma *et al.*, 1998, Basake *et al.*, 2001). Biological membranes and polyribosome. Proline is capable of detoxifying free radicals by forming a stable complex with them, thus maintaining NAD (P)⁺/ NAD(P)H ratios during stress at values similar to normal conditions (Hare and Cress, 1997; Floyd and Nagy, 1984).

The data also reveal that Phenolic compound in *Ankistrodesmus falcatus* did not affected at any copper levels. They remained around the control value. Phenolic compound might used as antioxidant compound for scavenging the reactive oxygen species in alga and higher plant (Burritt *et al.*, 2002; Contreas *et al.*, 2005).

The activity of nitrate reductase stimulated markedly up to 10 mm, there is quick increase in the activity of nitrate reductase as the level of 10 mm (174.3%) (the same level which nearly doubled the dry matter content) then this activity began to decrease gradually which was more pronounced only at the severe doses of copper. This criteria in the activity of nitrate reductase could help this alga to kept its protein content even at the severe doses of copper.

The inhibition of Nitrate reductase activity in *Ankistrodesmus* at severe copper stress is likely due to impaired NO_3^- uptake in the presence of elevated levels of the test

metals. The other possibility could be direct inhibition of N-R activity by the test metals (De Filippis and Pallaghy, 1994). Proteins did not affected with the pronounced drop in the activity of Nitrate reductase especially at the higher doses of the inhibitor. This might indicated that, the two processes (the activity of Nitrate reductase and the machinery of protein synthesis) did not necessary linked (Abdel Rahman *et al.*, 2004).

the data of protein pattern reveal that:

- 1- the total number of polypeptides increased progressively as the copper concentration increased in the culture media. For example the number Of polypeptides at the higher doses of cu (200 $\mu\text{m Cu}^{+2}$ and 250 $\mu\text{m Cu}^{+2}$) were 35 compared to only 15 in the control.
- 2- thus a huge number of polypeptides appeared due to cu stress.
- 3- the new polypeptides which induced by cu stress scattered from a very high molecular weight polypeptides to very low molecular weight polypeptides. Interesting the highest molecular weight 208 K.Da appeared when the growth began to be reduced (the level of 50 $\mu\text{m Cu}^{+2}$ and higher).while the 165 and 118 K.Da appeared from 10 to 200 $\mu\text{m Cu}^{+2}$.cumulatively it is worth to mention that, the more higher molecular weight of protein(208-73 K.Da) appeared in the copper treated media. Thus one can say that all of those polypeptides considered a copper polypeptides . also the lower molecular weight polypeptides (22 K.Da) which considered a heat chock protein appeared only at the level from 50 to 250 $\mu\text{m Cu}^{+2}$.

Conclusion

According to the above result and discussion it can be concluded that *Ankistrodesmus falcatus* exerted surprising copper tolerance:

- I- The growth criteria was stimulated unexpectedly up to 10 $\mu\text{m Cu}^{+2}$.
- II- it can survived up to the relative higher doses of copper (250 $\mu\text{m Cu}^{+2}$).
- III- which intern induce the ability of this alga to picked the a lot amount of copper under the polluted condition. consequently it posses a high capacity to clean up the copper polluted places(ponds and canal), accordingly

Ankistrodesmus falcatus can used as an excellent biological cleaner in polluted water.

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الملخص العربي

أجريت هذه الدراسة لمعرفة تأثير النحاس على نمو وعلى بعض التغيرات الفسيولوجية لطحلب الانكستروديسمس فالكاتس المعزول من محطة معالجة مياه الصرف الصحي بالدير. نمت هذا الطحلب في المعمل على وسط غذائي صناعي تحت تأثير تركيزات مختلفة من النحاس (0، 5، 10، 50، 100، 150، 200، 250 ميكرومول). وترك هذا الطحلب لينمو تحت تأثير هذه التركيزات لمدة 12 يوم ثم تم تقدير بعض التغيرات الفسيولوجية مثل عدد الخلايا، الوزن الجاف، المحتوى الصيغي، المحتوى البروتيني الذائب والغير ذائب والكلية، المحتوى السكري الذائب والغير ذائب والكلية، محتوى الأحماض الامينية والبرولين، محتوى المركبات الفينولية. يزداد عدد الخلايا والوزن الجاف زياد كبيره بزيادة تركيز النحاس حتى تركيز 10 ميكرومول من النحاس ثم يتناقص تدريجيا بزيادة التركيز. المحتوى الصيغي (كلوروفيل ا، كلوروفيل ب، كاروتين) ازدادت زيادة ملحوظة حتى تركيز 10 ميكرومول من النحاس ثم ظلت ثابتة حتى تركيز 50 ميكرومول ثم تناقصت. المحتوى السكري الذائب تناقص تدريجيا حتى تركيز 10 ميكرومول ثم ازداد زيادة بسيطة بزيادة تركيز النحاس في الوسط. يسبب النحاس تغيرات ملحوظة في المحتوى البروتيني حيث يحدث تراكم للمحتوى البروتيني عند تركيز 250 ميكرومول. المحتوى البروتيني الكلية ومحتوى البروتيني الغير ذائب يظل تقريبا ثابت عند كل تركيزات النحاس المستخدمة. الأحماض الامينية الحرة تتراكم بطريقة غير منتظمة مع زيادة تركيز النحاس في الوسط الغذائي. المحتوى البرولين يتناقص حوالي 65.6% عند تركيز 10 ميكرومول من النحاس ثم يزداد إلى 61.7% عند تركيز 250 ميكرومول. محتوى المواد الفينولية لا يتأثر بزيادة تركيز النحاس.