



# ISOLATION AND CHARACTERIZATION OF A HIGH SALT-TOLERANT AND GLYPHOSATE-DEGRADING STRAIN OF *Agrobacterium tumefaciens* BZ8

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**Abstract** – In this study, a high salt-tolerant and glyphosate-degrading strain named BZ8 was isolated from activated sludge. According to 16S rDNA sequencing methods, morphological, physiological and biochemical analysis, strain BZ8 was identified as *Agrobacterium tumefaciens*. The growth and glyphosate-degrading capability of *A. tumefaciens* BZ8 were investigated and the results showed that the optimum conditions for glyphosate degradation under 6% of NaCl concentration were found as follows: inoculation size of 10% (v/v), incubation temperature 37°C and initial pH of 5.0. Salt tolerance test showed that *A. tumefaciens* BZ8 grew well and could thoroughly degrade 2000 mg/L glyphosate in 36 h if the concentration of NaCl was lower than 6%, while the degradation rate decreased gradually with increasing NaCl concentration. But the glyphosate degradation rate could still reach 62% when the salt concentration was 8%. In addition, the kinetic parameters for *A. tumefaciens* BZ8 grown on 100-2800 mg/L glyphosate according to Haldane's model could predict the cell growth tendency successfully. These results showed that *A. tumefaciens* BZ8 could be used to control glyphosate wastewater with high salt content. Therefore, it has potential application.

**Keywords:** salt-tolerant; glyphosate-degrading strain; identification; growth kinetics.

## INTRODUCTION

Glyphosate is the most widely used herbicide in the world because it is a cheap, high efficient, low residue, broad-spectrum and foliar applied weed-killer. It can be used on non-cropland as well as on a great variety of crops. Nowadays, China is the largest exporter of glyphosate in the world. However, during the production of glyphosate, the discharge of wastewater can cause serious water pollution problems and environmental damage (Xu et al.,

2007). This is because the wastewater mainly contains glyphosate, NaCl and so on (Amoros et al., 2007). It was reported that producing one ton of glyphosate could cause production of 5-6 tons of glyphosate wastewater. Therefore, it is necessary to find safe and effective treatment methods for glyphosate wastewater.

Up to now, several technologies, which can be defined as physical methods, chemical techniques and biological treatments, have been developed for the treatment of glyphosate wastewater. Among these, physical and

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chemical methods have proven to be costly and always produce additional harmful pollution. Biological treatment is more economical than physical and chemical methods because it can be cost-effective and achieve the complete degradation of organic pollutants (Fan et al., 2012). Therefore, biological treatment of glyphosate wastewater has received increased attention for contamination prevention, and a number of different microorganisms, including yeast (Tang and Chen, 2010), *Pseudomonas* (Wu et al., 2010), *Flavobacterium* (Terry et al., 1986), *Arthrobacter* (Pipke and Amrhein, 1988) etc., have been isolated and tested for their ability to degrade glyphosate. However, there have been few studies on microorganisms that can function under the conditions of high NaCl concentration.

In this study, a salt-tolerant and glyphosate-biodegrading bacterium was screened from activated sludge and identified. In addition, the effects of environmental factors such as inoculation size, initial pH, temperature and salinity on the growth and glyphosate biodegradation were evaluated. The intrinsic kinetics of the growth on glyphosate using the Haldane model was also investigated. The aim of this study was to provide possible new resources for glyphosate wastewater treatment.

## MATERIALS AND METHODS

### Isolation of the salt-resistant and glyphosate-degrading strain

Activated sludge samples collected from a glyphosate wastewater treatment plant located in Anhui Hong Sifang Co., Ltd. currently manufacturing glyphosate in Hefei, Anhui province, People's Republic of China were acclimated for one month in the selection medium (SM) supplemented with 500-2000 mg/L glyphosate and 60 g/L NaCl for selection of a salt-tolerant and glyphosate-degrading strain. The SM contained (g/L) KH<sub>2</sub>PO<sub>4</sub> (0.5), K<sub>2</sub>HPO<sub>4</sub> (0.5), MgSO<sub>4</sub>·7H<sub>2</sub>O (0.5), CaCl<sub>2</sub> (0.04). Initially, 10 g of the samples were inoculated into 90 mL SM with 500 mg/L glyphosate and incubated at 37°C, 200 rpm for one week. Then, 10% of the culture was sub-cultured in SM with a glyphosate concentration of 1000 mg/L under the conditions as described above. The process was repeated until 2000 mg/L glyphosate was biodegraded by the samples. Then the resulting culture was serially diluted with sterilized water and spread onto selection medium agar plates supplemented with 2000 mg/L glyphosate as the sole carbon source. The plates were incubated at 37°C. Colonies that appeared on the plates were picked out and streaked several times to obtain pure isolates.

The obtained strains were stored on slants at 4°C and their glyphosate-degrading potential tested by inoculating the pure cultures into replicated flasks. Each flask contained 100 mL SM supplemented with 2000 mg/L glyphosate. The

cultures were maintained at 37°C with a shaker speed of 200 rpm. The strain that completely metabolized glyphosate at the fastest rate was chosen for further research.

Identification of the salt-resistant and glyphosate-degrading strain

The screened strain was characterized and identified based on its morphological, biochemical properties and 16S rRNA gene sequence analysis. The obtained cells of exponential growth phase in SM were harvested by centrifugation at 12000 rpm, 4°C for 20 min and washed with sterile water. By using the method mentioned by Zhai et al. (Zhai et al., 2012), the genome of the strain was extracted.

Part of the 16S rRNA gene was amplified by PCR using the forward primer: Eu27F (5'-AGAGTTTGATCATGGCTCAG-3') and the reverse primer: 1492R (5'-TACGGCTACCTTGTTACGACTT-3'). DNA amplifications were performed in 50 µL reactions containing approximately 100 ng of total DNA, 4 µL of 10×buffer, 4 µL of 25 mmol/L MgCl<sub>2</sub>, 4 µL of 2.5 mmol/L dNTPs, 2 µL of each 10 mmol/L primer and 0.4 µL of 5 U/µL Taq polymerase. The PCR conditions were as follows: 94°C for 3 min followed by 30 cycles of denaturation at 94°C for 30 s, annealing at 55°C for 30 s and primer extension at 72°C for 2 min, followed by a final step at 72°C for 10 min. PCR products were purified using a commercial kit (Sangon, Shanghai, China), and then the obtained purified PCR products were sequenced by Sangon Biotech Company (Shanghai, China). The sequencing result was submitted to GenBank for BLAST analysis. A phylogenetic tree was constructed by the neighbor-joining method using the software MEGA 4.0 (Tamura et al., 2007).

Colony morphology was observed after cultivation on SM at 37°C for 24 h. Cell morphology was examined after 24 h of incubation at 37°C by optical microscopy (CX31, OLYMPUS, Japan). Gram staining was performed by a conventional method (Dong and Cai 2001). After heat treatment at 80°C for 10 min, the presence of spores was investigated by optical microscopy following crystal violet staining for 2 min. Activities of catalase and nitrate reductase, hydrolysis of starch and gluten, utilization of citrate, nitrate reduction and production of indole were carried out according to the conventional methods (Smibert et al., 1994; Dong and Cai, 2001). The utilization of various substrates (sucrose, maltose, glycerol, D-galactose, xylose, D-glucitol, mannitol) as the sole carbon source was conducted according to the methods reported by Liu et al. (2014).

### Biodegradation experiments

The stored culture of strain BZ8 maintained on a nutrient agar slant was inoculated into 100 mL SM supplemented with 2000 mg/L glyphosate. Then the bacterial sub-culture

at the late exponential phase was used as inoculum. The biodegradation experiments were performed in 500 mL flasks with a working volume of 100 mL at 200 rpm.

In order to investigate the influence of initial inoculation size, initial pH and temperature on glyphosate degradation, replicate flasks with SM supplemented with 2000 mg/L glyphosate and 60 g/L NaCl were conducted at initial inoculum sizes varying from 2% to 15%, initial pH ranges from 3 to 7 and temperatures from 20°C to 42°C, respectively. A similar experimental procedure was used for testing the capacity of the strain BZ8 to degrade glyphosate under the optimized conditions.

The effect of NaCl concentrations on glyphosate degradation was investigated by applying different NaCl concentration (20, 40, 60, 80 and 100 g/L) at a constant glyphosate concentration of 2000 mg/L.

For the estimation of intrinsic kinetic parameters, a pure culture of strain BZ8 was inoculated into 100 mL SM in 500 mL flasks with 60 g/L NaCl under the optimized conditions. The SM was supplemented with glyphosate concentrations varying from 100 to 2800 mg/L. All the experiments were performed in triplicate. Culture samples were regularly taken for measurement of biomass and glyphosate concentration, and results were shown as means  $\pm$  confidence intervals.

### Glyphosate degradation kinetics

The most commonly used kinetic model to describe microbial growth is the Monod kinetic model. However, this model does not represent the growth kinetics of inhibitory substrates. Haldane's model developed from the Monod kinetic model has been used widely to characterize the metabolic inhibition of substrate on the microbial growth (Li et al., 2010). Therefore, in this study, Haldane's equation was used to describe strain BZ8 growth kinetics. For each flask with a certain initial glyphosate concentration, the specific growth rate ( $\mu_x$ , 1/h) was calculated as:

$$\mu_x = \frac{r_x}{\rho_x} = \frac{d\rho_x}{dt} \frac{1}{\rho_x} \quad (1)$$

where  $r_x$  is the cell growth rate (mg/L/h) and  $\rho_x$  is the cell concentration (mg/L). Then  $\mu_x$  was modeled using Haldane's model described as follows:

$$\mu_x = \frac{\mu_{\max} \rho_S}{K_s + \rho_S + \rho_S^2 / K_{SI}} \quad (2)$$

where  $\mu_{\max}$  is the maximum specific cell growth rate (1/h),  $\rho_S$  is the glyphosate concentration (mg/L),  $K_S$  is the saturation constant (mg/L), and  $K_{SI}$  is the self-inhibition constant (mg/L).

The yield coefficient (mg dry cell mass/mg glyphosate)  $Y_m$  was calculated using the following equation:

$$Y_m = \frac{M_{\max} - M_i}{\rho_{S_i} - \rho_{S_r}} \quad (3)$$

where  $M_{\max}$  is the maximum biomass concentration (mg/L),  $M_i$  is the initial cell concentration (mg/L),  $\rho_{S_i}$  is the initial substrate concentration (mg/L), and  $\rho_{S_r}$  is the residual glyphosate concentration when the cell concentration reached the maximum (mg/L).

### Analytical procedures

The cell concentration was monitored spectrophotometrically at 600 nm. The uninoculated sterile medium was used as a control. The optical density (OD) value was then converted to dry cell mass (DCW) using a calibration curve (Vasiliadou et al., 2008). The relationship between DCW and OD was found to be  $DCW \text{ (mg/L)} = 287.3 \times OD_{600} - 3.27$ ;  $R^2 = 0.9997$ . The concentration of glyphosate in the culture was determined by a high performance liquid chromatography (HPLC) method on a 1100 series HPLC (Agilent, American) as described by Kawai et al. (1991).

## RESULTS AND DISCUSSION

### Characterization and identification of glyphosate-degrading strain BZ8

Several strains were isolated from activated sludge by the method mentioned above. The glyphosate degradation ability was confirmed by the glyphosate degradation rate. Among them, a bacterium, named BZ8, was found to exhibit the highest glyphosate degradation ability and thus was chosen for further study. Colonies of strain BZ8 were smooth, round, moderate humidity, neat edges and white opaque. This bacterium is non-spore-forming, aerobic and gram-negative rod-shaped. It was V.P experiment-negative, nitrate reductase-positive, catalase-negative, utilization of citrate, hydrolysis of starch, gluten-positive and indole production-negative. It also utilizes sucrose, maltose, glycerol, D-galactose, xylose, D-glucitol, mannitol as sole carbon sources for growth.

The 16S rRNA gene is a highly conserved gene and was used for the phylogenetic analysis of taxa at higher levels. The 16S rRNA gene sequences of strain BZ8 were obtained (comprising 1385 nucleotides) and submitted to GenBank (<http://www.ncbi.nlm.nih.gov>). The sequence displayed the highest similarity (99%) to that of *A. tumefaciens* (GenBank accession AB535688.1 and FR828338.1). A phylogenetic tree was constructed based on the 16S

rRNA coding gene sequences of the isolate and the nearest relatives (Fig. 1). Combined with the morphological, physiological and biochemical analysis, the strain can be identified as *A. tumefaciens*. So, this strain was named *A. tumefaciens* BZ8.

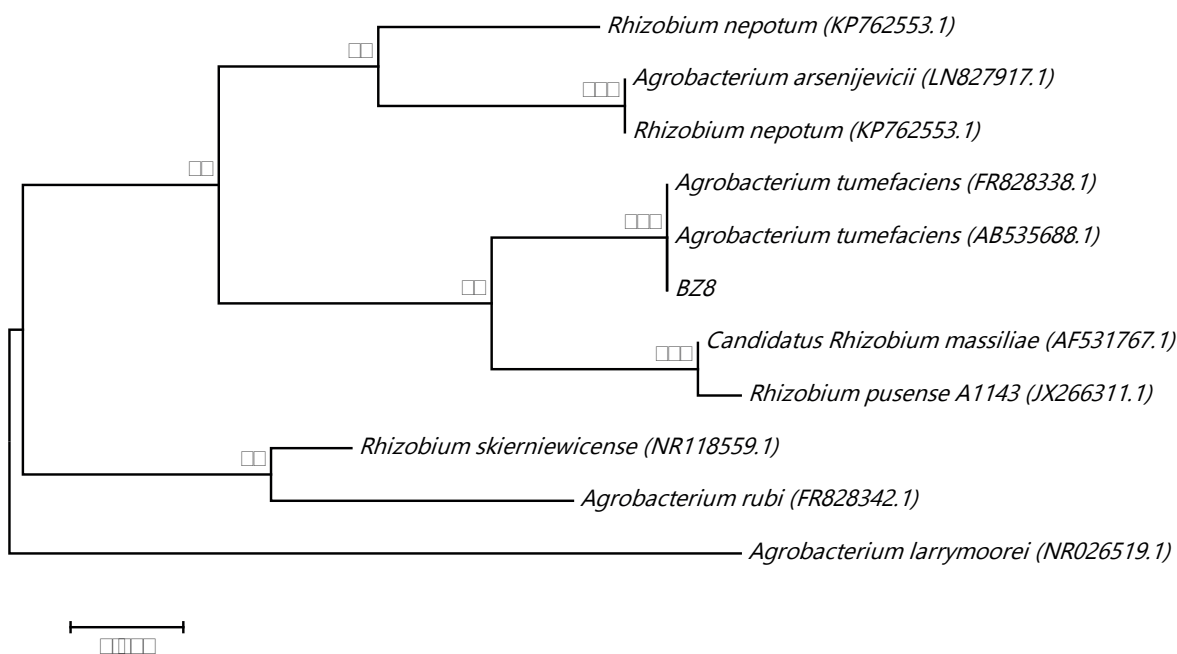
Effects of inoculation size, initial pH, temperature and salinity on glyphosate degradation

Fig. 2 illustrates the effect of inoculation size on the glyphosate degradation of *A. tumefaciens* BZ8. Experiments were carried out with the same initial glyphosate concentration of 2000 mg/L. It was obvious that the glyphosate concentration decreased with time for different inoculation sizes. Glyphosate was thoroughly degraded after 54 h, 48 h, 34 h and 36 h at inoculation sizes of 2%, 5%, 10% or 15%, respectively (Fig. 2). The effect of inoculum size on cell growth of strain BZ8 is also shown in Fig. 2. As shown in Fig. 2, with the increase of inoculum size, the lag period of the cell growth shortened gradually, while the cell growth rate and the substrate utilization rate were accelerated. Table 1 presents the values of glyphosate degradation rate with respect to inoculation size. The glyphosate degradation rate was greatly increased with inoculation size up to 10% (v/v), but it kept almost constant as the inoculation size further increased up to 15%. Taking into account the growth of strain BZ8 and glyphosate degradation, 10% (v/v) was chosen as the optimum inoculum size and used for all following experiments.

Glyphosate degradation and the cell growth were also tested at various initial pH values (Fig. 3). As expected, the pH greatly influenced glyphosate degradation as well as cell growth in the strain BZ8 culture system. Fig. 3 illustrates

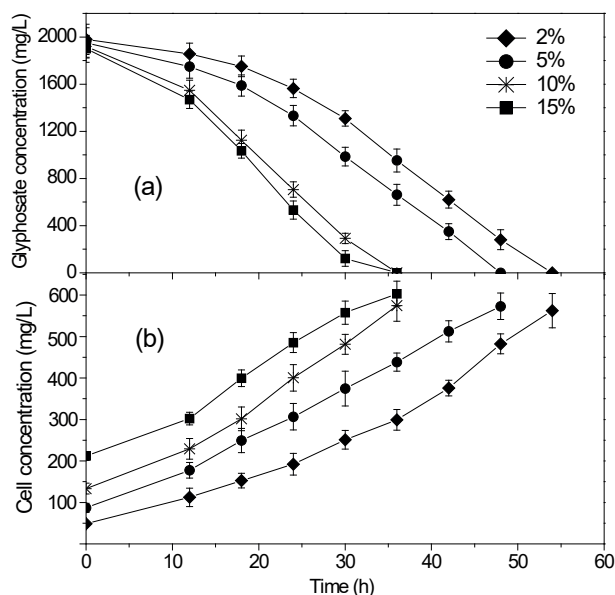
that the growth of strain BZ8 was positively related with the glyphosate degradation. The optimal pH for *A. tumefaciens* BZ8 growth occurred at pH 5.0. The shortest time to degrade glyphosate thoroughly occurred at pH 5.0. Apparently, a pH value of 5.0 would be the optimum for degradation of glyphosate by this bacterial strain. However, there was no distinct difference in glyphosate degradation rate as pH increased from 4.0 to 6.0 (Table 1). Although the glyphosate biodegradation efficiencies of the other two systems (pH 3.0 and pH 7.0) were significantly delayed, the strain BZ8 was still capable of entirely consuming glyphosate. This special pH adaptability indicates that the strain BZ8 could be applied to acidic or neutral conditions without altering the pH.

The influence of temperature on the glyphosate degradation and growth of strain BZ8 is illustrated in Fig. 4. The results show that strain BZ8 could effectively degrade glyphosate in the temperature range from 30°C to 42°C. Especially under the temperature of 37°C, BZ8 showed higher efficiency of glyphosate degradation as compared to the other three temperatures. The glyphosate degradation rate was seen to decrease with increasing or falling temperature (Table 1). However, at a low temperature of 20°C and a high temperature of 42°C the strain BZ8 could still degrade glyphosate thoroughly within 70 h and 40 h, respectively. Therefore, strain BZ8 is a potentially useful microorganism that could function at both comparatively low and high temperatures. This broad temperature adaptability may be important for biodegradation in glyphosate-contaminated environments that undergo daily and seasonal temperature changes.

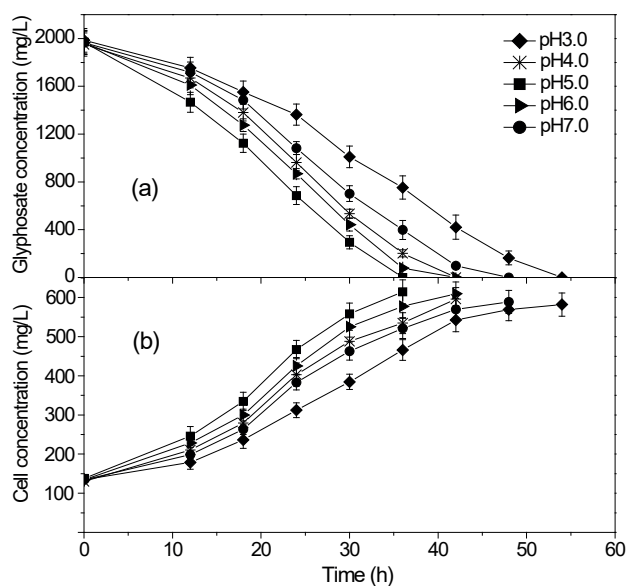


**Figure 1.** Phylogenetic tree of strain BZ8 based on 16S rRNA gene sequence analysis. Bootstrap values obtained with 1000 repetitions are indicated as percentages at all branches. The scale bar indicates 0.002 substitutions per nucleotide position.





**Figure 2.** Effect of inoculum size on biodegradation of glyphosate (a) and cell growth (b) by strain BZ8



**Figure 3.** Effect of pH on biodegradation of glyphosate (a) and cell growth (b) by strain BZ8

**Table 1.** Glyphosate degradation rates of *A. tumefaciens* BZ8 under different conditions

Initial inoculum size (%)	Glyphosate degradation rate (mg/L/h)	Initial pH	Glyphosate degradation rate (mg/L/h)	Culture temperature (°C)	Glyphosate degradation rate (mg/L/h)
2	36.67	3	36.76	20	26.5
5	40.63	4	46.43	30	47.12
10	54.16	5	54.78	37	55.18
15	53.33	6	46.79	42	46.15
—	—	7	40.93	—	—

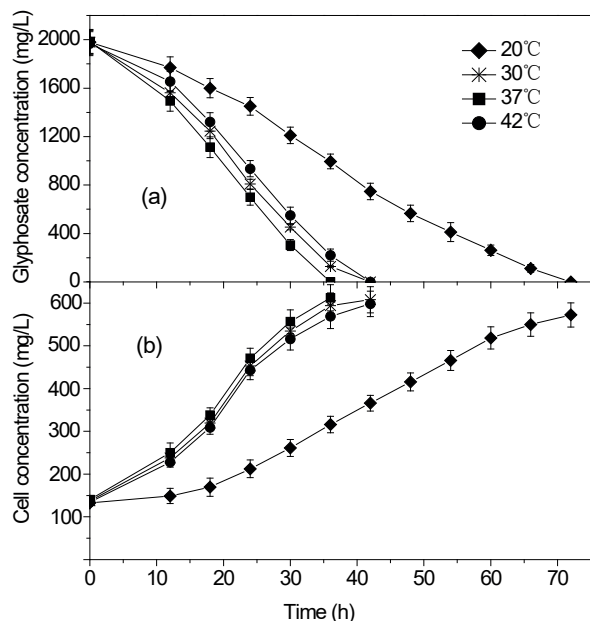
*A. tumefaciens* BZ8 grew well over the temperature ranges from 30°C to 42°C, which was significantly different at 20°C (Fig. 4). It reached the stable phase after 36 h at 30–42°C. At 20°C, the stable phase was recorded after 66 h. The fastest cell growth speed of BZ8 occurred at 37°C. According to these results, the temperature of 37°C was applied in the following set of experiments.

The actual glyphosate wastewater normally contains various concentrations of NaCl, which directly influences the activity of glyphosate-degrading microorganisms. In this study, strain BZ8 was inoculated into SM medium with 2000 mg/L glyphosate at different NaCl concentrations and incubated at 37°C for 36 h. As shown in Fig. 5, the strain grew well and glyphosate could be completely degraded in 36 h when the concentration of NaCl was lower than 60 g/L. With increasing salt concentration, the cell growth was inhibited gradually and the glyphosate degradation rate decreased greatly. The possible reason is that high salinity could cause osmotic stress or inhibit the reaction

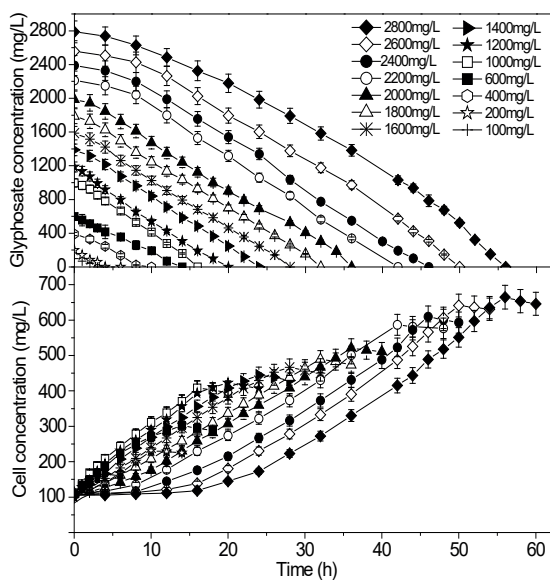
pathways in the organic degradation process. This results in a significant decrease in biological treatment efficiency or biodegradation kinetics. In addition, high salt content induces cell lysis, which causes increased effluent solids. However, the glyphosate degrading rate of strain BZ8 could still achieve 62% when the salt concentration reached 80 g/L. Compared with the other salinities, strain BZ8 grew difficultly at the salinity of 100 g/L, and the total glyphosate removed was only 35%. Therefore, strain BZ8 could be applied for comparatively high salinity glyphosate wastewater treatment.

#### Growth kinetics of *A. tumefaciens* BZ8

The consumption of glyphosate and growth of *A. tumefaciens* BZ8 at various initial glyphosate concentrations were investigated with the same NaCl concentration of 60 g/L (Fig. 6). As shown in Fig. 6,

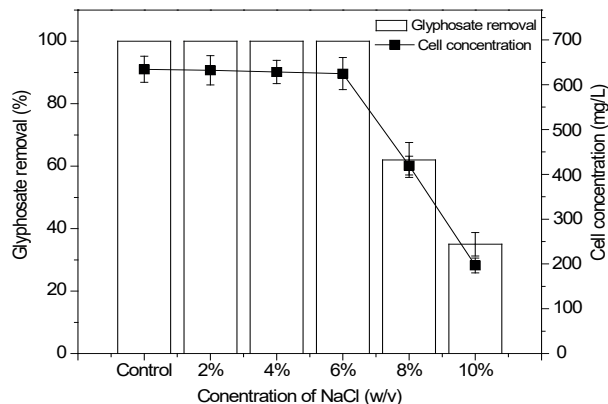


**Figure 4.** Effect of temperature on biodegradation of glyphosate (a) and cell growth (b) by strain BZ8

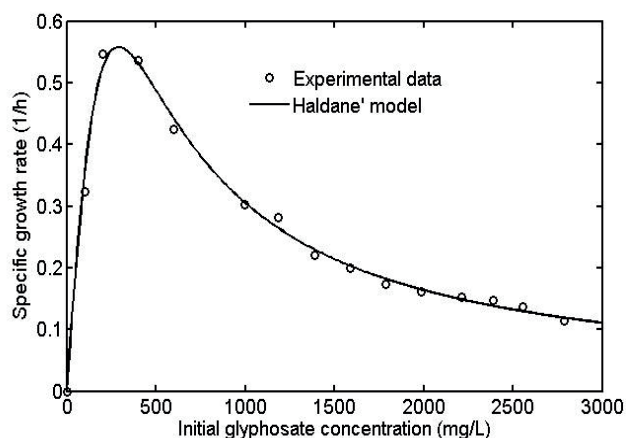


**Figure 6.** Biodegradation of glyphosate (a) and growth of strain BZ8 (b) at different glyphosate concentration

with the increase of glyphosate concentration, the time to degrade glyphosate thoroughly gradually shortened. It was obvious that the strain BZ8 could completely degrade 2800 mg/L glyphosate within 56 h (Fig. 6a). This maximum biodegradable glyphosate concentration of BZ8 was higher than *Bacillus cereus* CB4 (Fan et al. 2012) and *Arthrobacter* sp. N4 (Bazot and Lebeau, 2008), and it is inferred that strain BZ8 could tolerate a relatively



**Figure 5.** Effect of NaCl on biodegradation of glyphosate (a) and cell growth (b) by strain BZ8



**Figure 7.** Experimental and predicted specific growth rates of the culture due to Haldane's model

higher concentration of glyphosate than most glyphosate-degrading microorganisms in previous reports.

The biomass and glyphosate concentrations were linearly related for most of the active growth phase, so the yield coefficient ( $Y_m$ , mg/mg) was calculated by linearizing the decline of glyphosate with the increment of cell mass using Equation (3).

All coefficients of correlation ( $R^2$ ) were found to be above 0.99. The growth yields observed in this study varied between 0.201 mg/mg and 0.353 mg/mg as the initial glyphosate concentration was varied from 100 to 2800 mg/L (Fig. 6b). The highest value (0.353 mg/mg) was found at a glyphosate concentration of 200 mg/L, where the maximum specific growth rate was also obtained (Fig. 7).

When the concentrations were between 0 and 200 mg/L, the values of specific growth rate increased gradually with increase of glyphosate. There was a slight decrease in specific growth rate as the glyphosate concentrations increased from 200 mg/L to 400 mg/L; however, beyond 400 mg/L, with an increase of glyphosate concentration, a remarkable decline in specific growth rate occurred (Fig. 7).

The phenomenon could be explained by the fact that the sole carbon source of glyphosate was consumed mainly for assimilation into biomass and energy for cell growth and maintenance. When the inhibition effect of glyphosate becomes predominant above the glyphosate concentration of 400 mg/L, the proportion of the total glyphosate converted to energy for cell growth and maintenance increased as the specific growth rate decreased. More energy is required to overcome the effect of substrate inhibition at high glyphosate concentrations, while the proportion of the total substrate assimilated into biomass decreases as specific growth rates decrease. Therefore, substrate inhibition is known to reduce both specific growth rate and the yield coefficient.

The specific growth rate ( $\mu_x$ ) was estimated using Equation (1) by performing a linear least squares regression on the semi-logarithmic plot of the biomass concentration over cultivation time in the exponential growth phase. Then, the  $\mu_x$  data were employed to determine Haldane's parameters by nonlinear least-squares regression analysis using Matlab 7.0. The Haldane parameters for strain BZ8 grown on glyphosate were obtained as  $\mu_{max} = 1.28$  1/h,  $K_S = 84.82$  mg/L, and  $K_i = 227.59$  mg/L ( $R^2 = 0.992$ ). It is clear that Haldane's equation was strongly correlated with the experimental data (Fig. 7). This indicates that Haldane equation was suitable to describe the process of strain BZ8 glyphosate degradation in terms of cell growth behavior.

## CONCLUSIONS

In present study, a high salt-tolerant and glyphosate-degrading strain BZ8 was isolated and identified as *A. tumefaciens*. The growth and glyphosate-degrading capability of the strain BZ8 were investigated and the optimum degradation conditions were obtained. Salt tolerance tests showed that the strain grew well and could thoroughly degrade 2000 mg/L glyphosate in 36 h when the concentration of NaCl was lower than 6%, and the glyphosate degradation rate could still reach 62% when the salt concentration was 8%. The kinetic parameters for strain BZ8 grown on glyphosate according to Haldane's model were  $\mu_{max} = 1.28$  1/h,  $K_S = 84.82$  mg/L, and  $K_i = 227.59$  mg/L. The results demonstrate that strain BZ8 has a remarkable potential for application in the disposal of industrial glyphosate wastewater.

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