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Aerobic Biodegradation of 2-Picolinic Acid by a *Comamonas Sp.*

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Abstract

A strain was able to use 2-picolinic acid (2-PA) as a sole source of carbon, nitrogen and energy under aerobic conditions was isolated in this paper. Based on 16S rRNA gene sequence analysis, this strain was identified as *Comamonas sp.* and named as ZD3. The effects of pH and initial concentrations of 2-PA on the degrading capacity of ZD3 were investigated. It was found that ZD3 degraded 2-PA effectively under a pH range of 5.0 to 9.0, in which 7.0 was proved to be the optimal value. When the initial concentration of 2-PA was 100, 200, 400, 600 and 800 mg/L, ZD3 completely degraded 2-PA within 10, 18, 22, 78 and 114 h, respectively. Zero order kinetic model expressed the degradation behavior of 2-PA by ZD3 well. It was observed that under the concentration range of 100 to 400 mg/L, the degradation rate constant increased with the increase in concentration and reached the maximum at 400 mg/L. However, under the concentration range of 600 to 800 mg/L, the degradation rate constant began to decrease, indicating an inhibitory effect. Moreover, ultraviolet scanning and high performance liquid chromatography ion trap time-of-flight mass spectrometry showed that during the degradation of 2-PA the first reaction was α hydroxylation so that 6-hydroxy picolinic acid (6-HPA) was produced.

Keywords: *Comamonas sp.*; 2-Picolinic acid (2-PA); Aerobic; Biodegradation

Introduction

Pyridine and its derivatives belong to N-heterocyclic compounds. Pyridine-contained wastewater mainly stems from mining, coal and shale oil industry processing, wood preservative treatment, medicine and food industry, dye manufacture and agricultural production, etc. [1,2]. Due to its high toxicity and carcinogenicity, pyridines are listed as priority-controlled pollutants by United States Environmental Protection Agency [2]. 2-Picolinic acid, a kind of pyridine compounds, is intensively used for the manufacture of pharmacy, pesticides, daily chemical products, and additives of animal husbandry foods [3,4]. On account of high hydrophilicity (solubility=887 g/L, 20°C), 2-PA can steadily and persistently exist in water body and consequently endangers the environment. Recent studies have reported that the aerobic biological method is able to effectively remove 2-picolinic acid from wastewater [5], and the microbes mainly belong to the genera of Arthrobacter [6], Bacillus [7], and Streptomyces [8]. However, quantitatively describing the degradation process of 2-PA by microbes are still absent. Moreover, it is also unknown that whether 2-PA can be utilized by other species of microorganism or not [1,2]. In this paper, a strain was able to use 2-PA as a sole source of carbon, nitrogen and energy under aerobic conditions was isolated. Its cell morphology and homology analysis based on nucleic acid sequence were carried out. Environmental factors which influenced the degradation process of 2-PA were investigated, and consequently the degrading kinetic models were established in order to quantitatively describe the degradation capacity of this new strain. The purpose of this paper is to provide a scientific basis for the decontamination of wastewater loaded with 2-PA.

Materials and Methods

Chemicals

Both 2-PA (97%) and formic acid (95%) were of chromatographical purity and purchased from Sigma-Aldrich (American). Methanol (97%) was brought from the Fisher Chemical (American). Other reagents were of analytical purity. Ultrapure water was prepared by SPI-11-10T apparatus (resistivity=18 M Ω -cm).

Isolation and identification

The activated sludge used in this work was collected from a wastewater treatment plant (located in Xi'an, Shaanxi Province, China) that treated wastewater contaminated with 2-PA. A certain volume (10 mL) of the activated sludge was aseptically added to 90 mL of the sterilized MS medium (pH=7.0) [9] in a 250 mL flask. The 250 mL flask was supplemented with 2-PA with a concentration of 100 mg/L and placed in a rotary shaker (ZHWY-2102C, Zhicheng Analytical Instrument Manufacturing Co., Ltd., Shanghai, China) at 180 r/ min and 30°C. When the culture became obviously turbid, 10 mL of the culture was transferred to 90 mL fresh MS medium in a new 250 mL flask with 100 mg/L 2-PA. This cultivation was repeated until the degradation of 2-PA came to a stable level, and then the culture was diluted with sterilized NaH²PO⁴-K²HPO⁴ buffer (pH=7.0) and spread onto agar (1.5%) plates with MS medium and 100 mg/L 2-PA. The agar plates were incubated at 30°C for 2-3 d. Three colonies appeared on the agar plates and stain ZD3 was selected for the further study.

The 16S rRNA gene sequence of strain ZD3 was amplified by PCR with primers 27F (5'-AGTTTGATCMTGGCTCAG-3') and 1492R (5'-GGTTACCTTGTTACGACTT-3'). PCR mixtures contained template (genomic DNA 20-50 ng/L) (0.5 μ L), 2.5 mM dNTP (1 μ L), 10 × buffer (with Mg²⁺) (2.5 μ L), 27 F (10 μ M, 0.5 μ L), 1492R (10 μ M, 0.5 μ L), d H²O (up to 25 μ L), and Dream Taq-TM DNA Polymerase (0.2 μ L). The PCR instrument used was a 2720 thermal cycler (Applied Biosystems, ABI, USA), the electrophoresis instrument used was a DYCP-31DN (Beijing Liuyi Instrument Factory, China), and the electrophoresis image-forming instrument used was a FR980 (Shanghai

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Furi Science and Technology Co., Ltd., China). The amplification settings were a starting temperature of 94°C for 4 min; 94°C for 45 sec, 55°C for 45 sec, 72°C for 1 min, 30 cycles; 72°C for 10 min; and finally 4°C for termination. A partial sequence of the PCR products was determined for the isolate using an ABI PRISM[™] 3730XL DNA Sequencer (Applied Biosystems). BLAST analysis (NCBI) of the forward and reverse sequences obtained for strain ZD3 was performed. The phylogenetic tree was generated using the aligned sequences through the Neighbo-Joining algorithm using the MEGA 6 software.

Degradation of 2-PA

One loop of strain ZD3 on the agar plates was transferred aseptically to 100 mL sterilized MS medium (pH=7.0) in a 250 mL flask with 100 mg/L 2-PA. After 12 h of cultivation ten milliliters of the culture was aseptically inoculated into 90 mL sterilized MS medium in a 250 mL flask: (1) the pH of MS medium varied from 4.0 to 12.0 and then supplemented with 2-PA of 100 mg/L; (2) the pH of MS medium was fixed at 7.0 and then supplemented with 2-PA ranged from 100 to 800 mg/L. These cultures were incubated on a rotary shaker (30°C, 180 r/min). All cultivations were repeated three times. A flask with the same amount of the autoclaved cells (20 min, 120°C) was used as abiotic degradation. Samples were withdrawn periodically for the analysis of 2-PA concentration and total organic carbon (TOC). Prior to the analysis of 2-PA concentration and TOC, samples were centrifuged under 15°C at 12,000 r/min for 20 min to collect the supernatant which then was filtered through a 0.22-µm membrane to remove the biomass.

Analysis

Residual 2-PA concentration was determined at 265 nm with high performance liquid chromatography (S600, Sykam, Germany) using a Reprospher 100 C8 (5 μ m, 150 mm × 4.6 mm) (Sykam, Germany). The mobile phase was a mixture of water to methanol (v/v, 90: 10) containing 0.2% of formic acid. The flow rate was 1 mL/min and the column temperature was 30°C. Mass spectra of 6-hydroxy picolinic acid were acquired using a high performance liquid chromatography ion trap time-of-flight mass spectrometry (HPLC-IT-TOF-MS) (Shimadzu, Japan). The reaction mixtures were precipitated with ethanol and centrifuged (20000 r/min, 20 min, 4°C). The supernatants were analyzed by direct insertion mass spectra in the negative mode. Scanning electron microscopy (SEM) was used for the morphology observation and the method was the same as described in Ref. [9].

Results and Discussion

Isolation and identification

The morphological characteristics of strain ZD3 on agar plates was shown in Figure 1. It was creamy white and non-transparent with an irregular shape. Its surface was rough and flat. The edge was ragged. The colony closely attached to the surface of agar plates. As presented in Figure 2, the SEM image under 10000 times magnification showed that the cells of strain ZD3 were short rod with a length between 1 and 2 μ m. Based on the phylogenetic tree in Figure 3, it was found that the homology of strain ZD3 (Genbank No. KP900021) between *Comamonas sp.* VT3 reached 100%. Therefore, strain ZD3 was identified as a *Comamonas sp.* Recent researches have reported that *Comamonas sp.* can be applied for puring the wastewater contained polycyclic aromatic hydrocarbon (PAH), quinoline, cholesterin, steroids, and nitrobenzene, etc [10]. However, studies on the degradation of 2-PA by *Comamonas sp.* are still limited.



Figure 1: Morphology of strain ZD3 on agar plates.



Figure 2: SEM image of ZD3.

Degradtion of 2-PA

Effect of pH: Evironmental factors which affect the performance of microbial degradation mainly inculde temperature, pH, dissolved oygen (can be expressed by shaking speed), salinity, and initial concentration of substrates [11]. Experimental results suggested that the optimal temperature for the degradation of 2-PA by strain ZD3 was 30°C, shaking speed 150 rpm, and salinity 0% (w/v, NaCl) (data not shown). In this study, the influences of pH and initial concentrations of 2-PA were emphasized. As for pH, this paper adopted initial pH adjustment method and buffer method, respectively. Initial pH adjustment method was used HCl or NaOH for adjusting the pH (4.0-10.0) of MS medium with 2-PA (100 mg/L) as a sole carbon, nitrogen and energy source prior to the inoculation of strain ZD3.

As presented in Table 1, it can be concluded that the optimal pH value was 7.0, in which the degrading rate of 2-PA almost reached 100% after 12 h. TOC analysis showed that 2-PA was completely mineralized into carbon dioxide and water at 12 h. When the initial pH was 5.0 or 9.0, inhibitory effect emerged and the completely degrading time was prologed to 48h. When the initial pH was 4.0 or 10.0, 2-PA was not degraded at all. Moreover, when the initial pH ranged from 5.0 to 9.0, the value of pH fluctuated and finally showed a downward trend during the whole degradation process. The possible reason is 2-PA was biotransformed into intermediates with acidic or alkaline so that the value of pH changed. When the initial pH was 4.0 or 10.0, a downward

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Initial pH	<i>T</i> (h)	pН	Removal rate of 2-PA (%)
4.0	12	3.57	1.4
	24	3.41	1.3
	36	3.35	1.4
	48	3.40	1.5
5.0	12	4.71	1.7
	24	4.46	17.8
	36	4.87	34.5
	48	4.90	97.7
7.0	12	6.86	99.1
9.0	12	8.66	2.0
	24	8.69	2.3
	36	8.91	11.5
	48	8.66	98.3
10.0	12	9.88	0.1
	24	9.76	0.1
	36	9.63	0.5
	48	9.65	2.0

Table 1: pH changes and removal rates of 2-PA during the degradation process based on initial pH adjustment method.

trend of pH was also observed even no 2-PA was degraded. pKa^1 of 2-PA is 1.0 [12]. According to Henderson-Hasselbalch equation, when the initial pH of MS medium was 4.0 or 10.0, 2-PA was dissociated and in turn proton was released from the carboxyl group of 2-PA so the value of pH decreased.

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As for buffer method, experimental results showed that it was not suitable to use organic buffer solutions since some organic chemicals such as potassium acid phthalate served as a carbon source which in turn disturbed the utilization of 2-PA by strain ZD3. Toxic chemicals such as six methylene tetramine inhibited the activity of ZD3. Thus, inorganic buffer solutions were selected in this paper. From Figure 4a, it was apparent that the value of pH maintained constantly during the whole degradation process. As shown in Figure 4b, the degradation rate of 2-PA almost reached 100% within 10 h under pH of 7.0. When the pH value equaled to 6.0, 5.0 and 9.0, the completely degrading time was prolonged to 50 h, 60 h and 60 h, respectively. With pH of 4.0, 10.0, 12.0, all the degradation rates were less than 10% after 120 h, and the degradation rate was lowest at pH 4.0 (1.8%). It was concluded that the degradation performance of 2-PA by strain ZD3 at different pH values was in the following order: 7.0>6.0>5.0>9.0>10.0>12.0>4.0. In addition, the results obtained from initial pH adjustment method were nearly consistent with them from buffer method.

Influence of the initial concentration of 2-PA: When the initial concentrations of 2-PA were 100, 200, 400, 600 and 800 mg/L, the required time for complete degradation was 10, 18, 22, 78 and 114 h, respectively (Figure 5). The inhibitory effects became stronger with increase in the initial concentrations. Moreover, the lag phase was also prolonged. Figure 6 showed the ultraviolet spectrum under initial concentration of 200 mg/L. From 0 to 11 h, 2-picolinic acid (A²⁶⁵) kept unchanged, and then the peak intensity of 2-PA began to reduce at 12 h, simultaneously, an absorption peak at A³¹⁰ emerged indicating that 2-PA was bio-transferred into a new intermediate. From 13 to 18 h, the peak intensities of A²⁶⁵ and A³¹⁰ weakened until disappeared. TOC







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analysis showed that the final products at 18 h were carbon dioxide and water. When the initial concentrations of 2-PA were 100, 400, 600 and 400 mg/L, the same phenomenon were also observed.

Figure 7 presented HPLC-IT-TOF-MS of A³¹⁰ at 13 h corresponded to Figure 6, in which m/z=138.0233 was classified as the [M-H]⁻ adductive product of 6-hydroxy picolinic acid (6-HPA). In general, the first step for biodegradation of 2-PA is introduced a hydroxyl group on a carbon adjacent to N atom so that 6-HPA is generated [1,2]. In this paper, the experimental findings were consistent with these literatures [1,2]. The aerobic biodegradation pathway of 2-PA by strain ZD3 will be further studied.

Degradation kinetics: The models of zero-order kinetic and firstorder kinetic were applied to express the degradation process, which based on the following equations, respectively [13,14]:

$$\ln c = a + k_1 t \tag{1}$$

 $c = b + k_0 t \tag{2}$

where *a* and *b* are constants of degradation kinetics, respectively. *t* is time, k^1 and k^2 are rate constants of zero-order kinetic and first-order kinetic, respectively.

As seen in Figure 8, the zero-order kinetic and first-order kinetic models were used to fit in the linear range of each degrading profile in Figure 5. It was found that the correlation index (R^2) of zero-order kinetic was apparently higher than the first-order kinetic (Tables 2 and 3). k^0 increased with the increase in initial concentration (100 to 400



Figure 6: Ultraviolet spectra (UV) recorded the degradation and biotransformation of 2-PA.





Figure 8: Degradation kinetic models: (a) zero order kinetics model; (b) first order kinetics model.

Concentrations (mg/L)	The zero order kinetics equations	k _o	R ²
100	c=-10.867 t+139.01	10.87	0.9518
200	c=-24.956 t+433.13	24.96	0.9543
400	c=-47.390 <i>t</i> +1108.7	47.39	0.9466
600	c=-22.296 t+1725.8	22.29	0.9418
800	c=-15.941 <i>t</i> +1858.4	15.94	0.9302

Table 2: The parameters of zero order degradation dynamics model.

Concentrations (mg/L)	The first order kinetics equation	<i>k</i> ,	R ²
100	c=-0.0898 t+4.8224	0.09	0.8722
200	c=-0.7561 t+13.167	0.76	0.8642
400	c=-0.6481 t+15.780	0.65	0.8888
600	c=-0.1607 t+14.945	0.16	0.7920
800	c=-0.1175 t+16.037	0.12	0.8834

Table 3: The parameters of first order degradation dynamics model.

mg/L). It might be that under such a concentration range, the carbon, nitrogen and energy for the growth of strain ZD3 was not enough so that the higher initial concentrations, the faster the degradation rate [13]. When the initial concentrations of 2-PA increased from 600 to 800 mg/L, k^0 showed a trend of decline and the substrate inhibition effect emerged.

Conclusion

A strain utilized 2-PA as a sole carbon, nitrogen, energy sources under aerobic conditions were isolated and based on 16S rRNA gene sequence analysis it was identified as a *Comamonas sp.* The influence of pH and initial concentrations on 2-PA degradation were investigated. It was found that the optimal pH value was 7.0 and the required time for completely degradation was extended with the increase in initial concentration. The degradation kinetics model was established. Compared to the first order kinetics model, zero order kinetics model better expressed the degradation behavior of 2-PA.

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