

Accepted Manuscript

Promoting the treatment of crude oil alkane pollution through the study of enzyme activity

Meng Long, Li Wen, Bao Mutai, Sun Peiyan



PII: S0141-8130(18)32936-2

DOI: [doi:10.1016/j.ijbiomac.2018.07.160](https://doi.org/10.1016/j.ijbiomac.2018.07.160)

Reference: BIOMAC 10204

To appear in: *International Journal of Biological Macromolecules*

Received date: 14 June 2018

Revised date: 25 July 2018

Accepted date: 25 July 2018

Please cite this article as: Meng Long, Li Wen, Bao Mutai, Sun Peiyan , Promoting the treatment of crude oil alkane pollution through the study of enzyme activity. *Biomac* (2018), doi:[10.1016/j.ijbiomac.2018.07.160](https://doi.org/10.1016/j.ijbiomac.2018.07.160)

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

Promoting the treatment of crude oil alkane pollution through the study of enzyme activity

Meng Long^{a,b}, Li Wen^{a,b}, Bao Mutai^{a,b*}, Sun Peiyan^c

^a Key Laboratory of Marine Chemistry Theory and Technology, Ministry of Education, Ocean University of China, Qingdao 266100, China

^b College of Chemistry & Chemical Engineering, Ocean University of China, Qingdao 266100, China

^c Key Laboratory of Marine Spill Oil Identification and Damage Assessment Technology, North China Sea Environmental Monitoring Center, State Oceanic Administration, Qingdao 266033, China

Abstract Microbes appear to play a key role in bioremediation of petroleum hydrocarbons pollution and little attention has been paid to the enzyme activity in the process of alkane bioremediation. Oil field bacterium identified as *Pseudomonas synxantha* LSH-7⁺ was chosen as the tested strain. Periodically collected samples were analyzed by GC-FID (Gas Chromatography- Flame Ionization Detector) and RT-qPCR (Quantitative-Real-Time-PCR). GC-FID results showed this bacterial strain has great degradation ability on crude oil *n*-alkanes and RT-qPCR data indicated the differences between the three genes expression including *AlkB*-, *Cytochromes P450*-, and *almA*- related when grown on different-chain alkanes. Meanwhile, enzyme activity like alkane hydroxylase, alcohol dehydrogenase, dehydrogenase, protease, phosphatase, catalase and lipase were measured. Extracellular alkane hydroxylase was induced in a higher degree than intracellular in the early incubation time, alcohol dehydrogenase increased/decreased along with alkane hydroxylase, and the pH of the medium obviously decreased. Other enzymes were also described including dehydrogenase activity that reached a highest point that was slower than alcohol dehydrogenase, protease activity started multiplying after a period of culture while biomass was immediately increased, catalase activity dramatically enhanced in the presence of alkanes, phosphatase activity was closely linked to pH approximately but lipase activity was found to be moderate.

Keyword crude oil alkane bioremediation; pseudomonas; gene expression and enzyme activity

Introduction

Oil pollution especially in marine environment has become a serious environmental problem, for example, Deepwater Horizon oil spill in the Gulf of Mexico (Kostka et al. 2011) and the Spanish Atlantic coast polluted by crude oil (Fernández et al. 2011).

* Corresponding author: mtbao@ouc.edu.cn; Tel/Fax: +86-532-66782509. Postal address: Songling Road 238, Ocean University of China, Qingdao, China.

Download English Version:

<https://daneshyari.com/en/article/8326832>

Download Persian Version:

<https://daneshyari.com/article/8326832>

[Daneshyari.com](https://daneshyari.com)