Bioresource Technology 117 (2012) 155-163

Contents lists available at SciVerse ScienceDirect

Bioresource Technology

journal homepage: www.elsevier.com/locate/biortech

Evolution of the microbial community in a full-scale printing and dyeing wastewater treatment system

Qingxiang Yang^{*}, Jia Wang, Hongtao Wang, Xuanyu Chen, Siwei Ren, Xueling Li, Ying Xu, Hao Zhang, Xuemei Li

College of Life Sciences, Henan Normal University, Xinxiang 453007, China Key Laboratory for Yellow River and Huai River Water Environment and Pollution Control, Ministry of Education, Xinxiang 453007, China

HIGHLIGHTS

- ► We traced the treatment efficiencies of the biological system in a full scale PDW treatment plant.
- The PDW treatment system harbored diverse bacteria, achaea and fungi.
- The microbial populations underwent dramatic changes during system start-up and running period.
- ► In contrast to fungi, the diversity of bacteria and archaea increased with the system running.

ARTICLE INFO

Article history: Received 18 February 2012 Received in revised form 14 April 2012 Accepted 19 April 2012 Available online 27 April 2012

Keywords: Archaea Bacteria Denaturing-gradient gel electrophoresis (DGGE) Fungi microbial community Printing and dyeing wastewater

1. Introduction

In China, various dye-containing wastewaters constitute nearly 30% of industrial effluents, in which printing and dyeing wastewa-

G R A P H I C A L A B S T R A C T

DGGE profiles of the microbial populations in different stages of the system running.



ABSTRACT

In this study, the dynamics of bacterial, fungal and archaeal populations in two-stage biological processes of a full-scale printing and dyeing wastewater treatment system were traced using cultivation and molecular biological techniques. The enumeration results indicated that bacteria were the dominant population in the system, in which the ratio of fungi to bacteria decreased in all the treatment units, while the ratio of archaea to bacteria increased significantly, especially in samples from the second-stage biological treatment process. PCR-denaturing gradient gel electrophoresis (DGGE) analysis showed that the microbial diversity increased with system running and 64.6% of bacterial, 57.6% of fungal and 38.2% of archaeal populations remained in the system from the seed sludge during system start-up. In spite of variation in the microbial community and composition of the influents, some bacterial species such as *Thauera* sp. and Xanthomonadaceae were present simultaneously in all the collected samples.

© 2012 Elsevier Ltd. All rights reserved.

ter (PDW) is one of the most important effluents. PDW normally has a low ratio of BOD_5/COD (5 days biochemical oxygen demand/ chemical oxygen demand, around 20%), high pH value (10–13), and contains toxic, frequently changing and bio-recalcitrant components such as dyes and dyeing additives (polyvinyl alcohol, PVA) (Wu et al., 2007). Although the current treatment for this wastewater in China is a combination of physical-chemical and biological processes, various biological methods play core roles and are capable of removing 40–50% COD and 50–60% of colority



^{*} Corresponding author. Tel.: +86 373 3325528.

E-mail addresses: yangqx66@163.com (Q. Yang), tao666tao666@163.com (H. Wang), cxy549983744@126.com (X. Chen), renwei000000@163.com (S. Ren), lixueling870710@126.com (X. Li), jingjing86321@126.com (Y. Xu), kele1564@126. com (H. Zhang), xm633@163.com (X. Li).

^{0960-8524/\$ -} see front matter @ 2012 Elsevier Ltd. All rights reserved. http://dx.doi.org/10.1016/j.biortech.2012.04.059