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环境因子对脱氯功能念珠藻 *Nostoc* PD-2 降解多氯联苯过程中基因表达的影响

**Effects of environmental factors on expression of genes in *Nostoc* PD-2 during polychlorinated biphenyl-degradation process**

关键词: [多氯联苯](#) [Nostoc](#) [双加氧酶](#) [细胞色素b<sub>6</sub>f复合体铁硫蛋白](#)

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摘要: 以分离筛选自多氯联苯污染水稻田中的脱氯功能蓝藻 *Nostoc* PD-2 为材料, 两种典型的三氯代多氯联苯 PCB28 和 PCB30 为目标污染物, 在不同的氮源、碳源和培养温度条件下, 研究了脱氯功能藻种 *Nostoc* PD-2 中双加氧酶基因和细胞色素 b<sub>6</sub>f 复合体铁硫蛋白基因的表达情况及两种基因对脱氯作用的影响。结果显示, 以硝酸钠作为氮源时, 与氮气氮源组相比, PCB28 和 PCB30 降解组中的两种基因均显著上调表达, 双加氧酶基因的上调倍数分别为 1.9 和 5.7 倍, 铁硫蛋白基因的上调倍数分别为 1.1 和 1.7 倍; 添加碳酸钠时, 与对照相比, 双加氧酶基因最高上调了 2.2 倍, 铁硫蛋白基因最高上调了 3.4 倍; 提高温度对双加氧酶基因和铁硫蛋白基因的表达均有促进作用。相较于铁硫蛋白, 双加氧酶基因相对表达量与 PCB28 和 PCB30 的脱氯百分比之间的相关系数分别为 0.872 和 0.832, 表明双加氧酶基因活性与功能藻种 PD-2 脱氯降解 PCBs 的相关性更高。本研究结果表明, 不同环境因子可引起脱氯功能藻种 *Nostoc* PD-2 降解 PCBs 过程中双加氧酶基因和铁硫蛋白基因的差异表达, 添加碳源对降解效果的促进作用最明显, 脱氯功能藻种在工程应用中采用优化的环境条件有利于提高降解效率。

**Abstract:** Cyanobacterium *Nostoc* PD-2 isolated from polychlorinated biphenyl (PCB)-contaminated paddy soils was used to biodegrade two kinds of typical trichlorodiphenyls (2,4,4'-trichlorobiphenyl and 2,4,6-trichlorobiphenyl). The expressions of dioxygenase genes and cytochrome b<sub>6</sub>f complex Fe-S protein genes in *Nostoc* PD-2 during the PCB-degradation were determined under the conditions of different nitrogen sources, carbon sources, and cultural temperatures. In addition, the relationships between both genes expressions and PCB-dechlorination efficiencies were also analyzed. Results showed that, the expressions of both genes in the sodium nitrate group were significantly up-regulated comparing to that in the nitrogen group. The up-expression folds of dioxygenase gene under the exposure of PCB28 group and PCB30 group were 1.9 and 5.7, respectively. While the up-expression folds of Fe-S protein gene were 1.1 and 1.7, respectively. Compared to the control group, the maximum up-regulated folds of dioxygenase gene and Fe-S protein gene in the sodium carbonate group were 2.2 and 3.4, respectively. Elevated temperature could facilitate the expressions of dioxygenase gene and cytochrome b<sub>6</sub>f complex Fe-S protein gene. The expression of dioxygenase gene more significantly correlated with the dechlorination percentages of both PCB congeners than that of Fe-S protein gene. The correlation coefficients between dioxygenase gene expressions and the dechlorination percentages of both PCBs congeners were 0.872 and 0.832, respectively. Such findings can demonstrate a new insight to clarify the molecular mechanism of PCB-biodegradation by *Nostoc* PD-2. In addition, using sodium carbonate as carbon source can promote the efficiencies of PCB-dechlorination by *Nostoc* PD-2. Our study can provide the new application of cyanobacteria in bioremediation engineering.

**Key words:** [polychlorinated biphenyls](#) [Nostoc](#) [dioxygenase](#) [cytochrome b<sub>6</sub>f complex Fe-S protein](#)

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