

豹纹鮰棘鲈尾部溃烂症病原菌的鉴定与药敏试验

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Identification and drug sensitive test of bacterial Pathogens from Plectropomus leopardus with tail fester diseaseGU Liangbin^{1,2}, XU Liwen¹, FENG Juan¹, SU YouLu¹, LIU Guangfeng¹, GUO Zhixun¹

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2011年海南会文烟堆某豹纹鮰棘鲈(*Plectropomus leopardus*)养殖场突发大量死亡, 病鱼体色发白, 尾鳍溃烂。从病鱼病灶处分离出1株菌X11YD05, 回归感染证实为患病豹纹鮰棘鲈病原菌, 对平均体质量为30.8 g的斜带石斑鱼(*Epinephelus cooides*)的半数致死量(LD50)为 $1.67 \times 10^5 \text{ cfu} \cdot \text{g}^{-1}$ 。为进一步确定病原菌的分类学特征, 扩增其16S rDNA、toxR、rpoD、mreB、topA和rctB等6种管家基因, 构建系统发育树。结果发现, X11YD05与哈维弧菌(*Vibrio harveyi*)有较高同源率。结合生理生化试验, 确定X11YD05为哈维弧菌。16种药敏试验表明该菌仅对氟氯沙星、诺氟沙星、环丙沙星、恩诺沙星、复方新诺明等5种抗生素敏感。

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关键词: 豹纹鮰棘鲈, 哈维弧菌, 烂尾症, 16S rDNA**作者相关文章****Abstract:**

In 2011, there was an outbreak in a *Plectropomus leopardus* farm in Wenhui Town, Yandui Village, Hainan Province. The color of sick fish turned white and the tail fin festered. We isolated a Strain X11YD05 from sick *P. leopardus* which was proved to be pathogen by artificial infection and the LD50 was $1.67 \times 10^5 \text{ cfu} \cdot \text{g}^{-1}$ in *Epinephelus cooides* using Karber method. To determine the molecular characteristics of X11YD05, we amplified the 16S rDNA, transmembrane regulatory protein (toxR), RNA polymerase $\delta 70$ factor (rpoD), actin-like cytoskeleton protein (mreB), topoisomerase (topA) and replication origin-binding protein (rctB) by PCR. Sequence analysis indicates that X11YD05 belonged to *V. harveyi*. Drug sensitivity test shows that X11YD05 was sensitive to norfloxacin, ofloxacin, ciprofloxacin, enrofloxacin and cotrimoxazole.

Key words: *Plectropomus leopardus* *Vibrio harveyi* tail fester disease 16S rDNA**收稿日期:** 2014-10-10 **修回日期:** 2014-12-22 **出版日期:** 2015-08-05**PACS:** S 941.4**基金资助:**

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