

عنوان مقاله:

Bacterial diversity in the intestine of sea cucumber Stichopus japonicus

محل انتشار:

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نویسندگان: M.L. Gao

H.M. Hou G.L. Zhang Y. Liu

L.M. Sun

خلاصه مقاله:

The intestinal bacterial diversity of Stichopus japonicus was investigated using 15S ribosomal RNA gene (rDNA) clone library and Polymerase Chain Reaction/Denaturing Gradient Gel Electrophoresis (PCR-DGGE). The clone library yielded a total of IAA clones, and these were sequenced and classified into 109 operational taxonomic units (OTUs) with sequence similarity ranging from AA to 100%. The coverage of the library was YY.F%, with approximately AA.Y% of the sequences affiliated to Proteobacteria. Gammaproteobacteria and Vibrio sp. were the predominant groups in the intestine of S. japonicus. Some bacteria such as Legionella sp., Brachybacterium sp., Streptomyces sp., Propionigenium sp. and Psychrobacter sp were first identified in the intestine of sea cucumber

کلمات کلیدی: Intestinal bacterial diversity, ۱۶S rDNA, PCR-DGGE, Sequencing, Stichopus japonicus

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