

SII-3-5

Bench-scale Bioremediation of Pentachlorophenol-contaminated Soil

Hee-Mock Oh*, Sung-Gie Lee and Byung-Dae Yoon

Environmental Microbiology Research Unit

Korea Research Institute of Bioscience and Biotechnology

Twelve active isolates were finally selected for the further studies on the biodegradability and for the use in the slurry-phase and land-farming treatment of pentachlorophenol (PCP)-contaminated soil. Especially, *Pseudomonas* sp. Bu34 was able to degrade a much higher concentration of PCP than any previously reported PCP-degrading bacteria or fungi. The soil brought from a PCP-contaminated site contained 50 - 100 mg/kg wet soil of PCP and was treated with the active bacterial isolates in a bench-scale slurry reactor. PCP in the contaminated soil was decreased up to 90% at 8 days after incubation of the isolates and completely disappeared within 16 days. In land-farming treatment, 1,000 mg/kg soil PCP in indigenous microorganisms (Control), indigenous and added bacterial treatment, and indigenous and added bacterial treatment with air supply were degraded to 59, 84, and 92% at 54 days after incubation, respectively. Therefore, it was demonstrated that PCP removal rate in slurry-phase treatment was higher than that in land-farming treatment. Air supply and microbial addition to the treatment system appeared to be much effective in enhancing biodegradation of PCP in soil.

SII-4-1

Crystal Structure of Hepatitis C Virus RNA Helicase; Mechanism of Unwinding Duplex RNA

Byung-Ha Oh*, Hyun-Soo Cho, Nam-Chul Ha, Lin-Woo Kang
Department of Life Sciences, Pohang University of Science and
Technology

Hepatitis C Virus (HCV) is the major etiologic agent of non-A non-B hepatitis. Its genome encodes RNA helicase domain composed of about 450 amino acids which unwinds RNA/RNA, DNA/DNA, and RNA/DNA heteroduplexes. We have determined crystal structure of HCV RNA helicase domain at 2.3 Å resolution by multiple isomorphous replacement method. One of the crystal packing interface between a pair of symmetry related molecules is a channel lining with strictly conserved arginine residues. The channel is close to the NTP binding site, and has a size adequate to accommodate only single stranded RNA. Based on the structure and the unusual crystal packing, a molecular mechanism of unwinding duplex RNA by a putative functional dimer of HCV RNA helicase will be presented. [This study made use of X-ray facility at Pohang Light Source]