[Title of collaboration research]: Genomic Sequencing of thermo-stable microorganisms

[The members of collaboration research]: Toshifumi Tsukahara

[Reference home-page address]: http://www.jaist.ac.jp/nmcenter/labs/tsukahara-www/

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**[Contents]** Genomic sequencing of thermo-stable microorganisms is an efficient way for screening or generating thermo-stable useful enzymes. It is possible that there are various new thermo-stable microorganisms in Vietnam. We will cloning of fragments of genomic DNAs and sequencing them. All the fragments generated from a thermo-stable microorganism are inserted into a plasmid, then vectors allows for propagation and replication in E. coli. Each isolated plasmid should be sequencing by using ABI 3130xl sequencer, then sequences of fragments will be finally re-aligned based on overlaps in their sequences. By the complete genomic sequencing, we can deduce primary sequences of whole proteins in the thermo-stable microorganism. Thermo-stable proteins could be useful tools as nano-biomaterials.

The research is concerned with the basic science underlying bioremediation efforts, structural science and genome science, as well as other uses of biotechnology. Identification of new thermo-stable proteins will enhance numerous possibilities to create super-biomolecules which will be added thermo-stability to specific functional proteins. The

results of the investigations may be contributed to generate stable bioreactors for degradation of hazardous waste, production of useful metabolic compounds and biomass.

