

Complete genome sequencing of *Bacillus* sp. TK-2, analysis of its cold evolution adaptability

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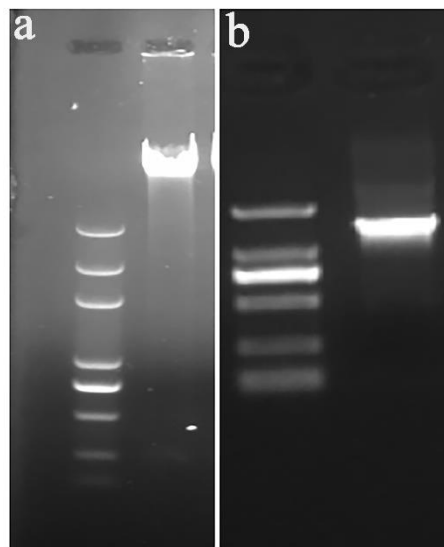
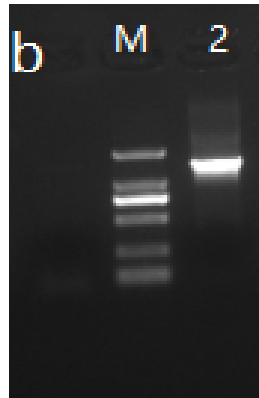
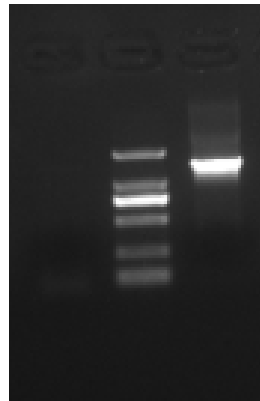
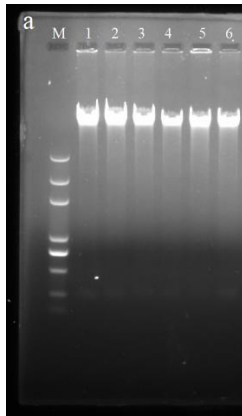
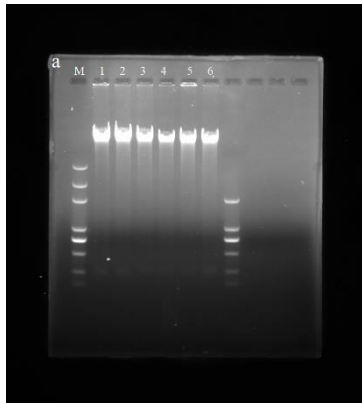
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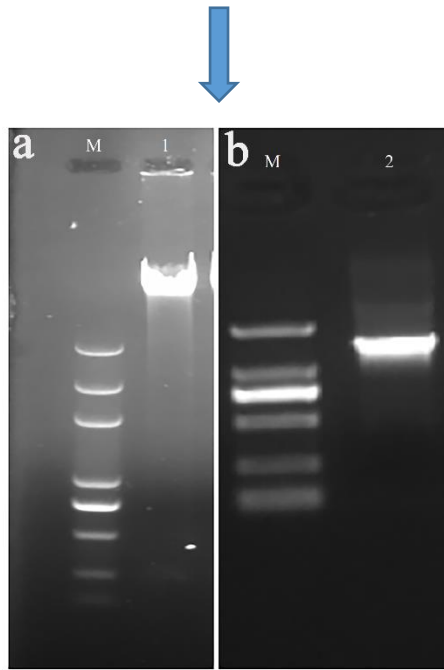


Figure S1. Agarose gel electrophoresis. (a) DNA extracted from *Bacillus* sp. TK-2. **M(Marker):** (DL15000: 15000、8000、5000、3000、2000、1500、1000、750bp), 1: TK-2. (b) PCR product of 16S rDNA of *Bacillus* sp. TK-2. **M(Marker):** (DL2000: 2000、1000、750、500、250、100bp), 2: TK-2.

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1  tgcagtcgag cgaatggatt aagagcttgc tctatgaag ttagcggcgg acgggtgagt
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1381 accggaagtc ggtggggtaa cctttt

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Figure S2. Nucleotide sequence of 16S rDNA from *Bacillus* sp. TK-2.