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A novel cold-inducible expression system for Bacillus subtilis

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Abstract

Production of recombinant proteins at low temperatures is one strategy to prevent formation of protein aggregates and the use of an expensive inducer such as IPTG. We report on the construction of two expression vectors both containing the cold-inducible *des* promoter of *Bacillus subtilis*, where one allows intra- and the other extracellular synthesis of recombinant proteins. Production of recombinant proteins started within the first 30 min after temperature downshock to 25 °C and continued for about 5 h. © 2007 Elsevier Inc. All rights reserved.

Keywords: Cold shock; Des promoter; lacZ; htpG; a-Amylase; Penicillin-binding protein

One of the major drawbacks during high-level production of recombinant proteins in bacteria is the inability of many proteins to reach their native conformation. Under conditions of overproduction, proteins tend to accumulate within refractile aggregates designated inclusion bodies [1]. Since inclusion body formation is believed to arise from the unproductive association of folding intermediates [2], one experimental approach to prevent formation of these aggregates is to increase the intracellular concentration of molecular chaperones to favour "on-pathway" folding reactions and prevent the accumulation of kinetically trapped folding intermediates [3]. An alternative method to limit the aggregation of recombinant proteins consists in cultivating the cells at low temperatures [4]. Besides reducing formation of inclusion bodies, low-temperature expression lowers the degradation of proteolytically sensitive proteins [5,6].

To ensure high level production of recombinant proteins, two different strategies can be used: fusion of the coding region of the protein of interest to an inducible promoter, e.g., using an IPTG- or xylose-inducible promoter [7], or, alternatively, to make use of a cold-shock inducible promoter. When mid-exponential phase bacterial cells are rapidly transferred from 37 to 25 °C or even a lower temperature, the synthesis of most cellular proteins greatly decreases, while that of cold-shock proteins is transiently upregulated [8]. In Bacillus subtilis, one of these cold-shock proteins is a membrane-bound desaturase (Δ 5-Des) encoded by the des gene [9]. This enzyme catalyzes the introduction of a *cis* double bond at the $\Delta 5$ position of a wide variety of saturated fatty acids. It has been shown that the des gene is tightly regulated during cold shock. While the des mRNA is barely detectable at 37 °C, its synthesis is transiently induced upon a temperature downshift [10]. Expression of the des gene does not depend on de novo protein synthesis, but on a two-component signal transduction system which consists of the sensor kinase DesK and the response regulator DesR [11]. It is assumed that the kinase senses a temperature downshift through changes in the physical state of the cytoplasmic membrane. The C-terminal kinase domain of DesK undergoes autophosphorylation, and the phosphoryl group is then transferred to the response regulator DesR. Phosphorylated DesR binds to two adjacent DNA-binding sites leading to the recruitment of RNA polymerase to the des promoter and activation of transcription [12]. The Δ 5-desaturase directly introduces double bonds into membrane lipids leading to a return to the original fluidity of the membrane. This is sensed by DesK which changes from a kinase to phosphatase activity

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leading to a dephosphorylation of DesR with a concomitant turn off of the des gene [13].

Based on these data, we developed a cold-inducible expression system for B. subtilis making use of the des promoter. We show here that cold-induction results in a significant induction of reporter genes largely preventing formation of aggregates of an aggregation-prone protein. Cold-inducible expression systems have also been developed for Escherichia coli which are based on a different principle [14,15]. Cold-inducible expression systems provide an inexpensive alternative technology especially for industrial production of recombinant proteins complementing the widely used IPTG- and xylose-inducible systems.

Materials and methods

Materials

Bacteria, plasmids and growth conditions

Bacterial strains and plasmids used are listed in Table 1. Cells were grown in Luria Broth (LB) medium at 37 or 25 °C under aeration. Antibiotics were added where appropriate (ampicillin at 100 µg/ml, neomycin at 10 µg/ml and chloramphenicol at 10 µg/ml).

Methods

Construction of a transcriptional fusion between the des promoter and the lacZ reporter gene

The promoter region of the des gene was fused to the lacZ reporter gene using the integration vector pDG1728 [16]. The resulting transcriptional fusion is sandwiched

Table 1

Strains and plasmids	used in	this study
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between amyE-front and amyE-back allowing its integration into the B. subtilis chromosome at the amyE locus. The des promoter region was amplified using oligonucleotides (ON) ON1 (GGCCATGAATTCTCCGGCATCCC GATCATCGC; restriction site underlined) and ON2 (GGCCATAAGCTTTCTCATTGTGTGTCTCGGTTC AG). The amplicon was cleaved with EcoRI and HindIII and inserted into pDG1728 cut with the same enzymes resulting in pDG1728-des. This recombinant plasmid was transformed into strain WW02, and transformants were selected on LB plates containing chloramphenicol and screened for the loss of the neomycin resistance marker, and one positive transformant (AL03) was kept for further studies.

Construction of a des null mutant

To construct a des knockout, the gene including flanking regions was amplified using the primer pairs ON3/ON4 (GGCCATGTCGACTGAACCGAGACACACAATG; GGCCATGAGCTCATAGTTGAGCACCTTTGG), and the amplicon was cleaved with SalI and SacI and cloned into pBluescript SKII⁺ treated with the same enzymes. Next, the recombinant plasmid was treated with HindIII and BclI to remove a 61-bp internal fragment of des which was replaced by the neo marker using pBgaB as template and the primer pair ON5/ON6 (GGCCATAAGCTT AGGTCGAGATCAGGGAATGAGTT; GGCCATTGA TCAGATCAATTCTGACAGCCATG). Using the primer pair ON3/ON4, the modified gene was amplified and transformed into B. subtilis 1012. Neomycin-resistant transformants were selected and checked by Southern-blot for chromosomal replacement of the des by the neo gene (data not shown). One strain (AL02) was kept for further studies.

Strains	Genotype	
E. coli		
DH10B	F^- mcrA Δ(mrr hsdRMS mcrBC) φ80d lacZ ΔM15 deoR recA1 araD139 Δ(ara leu) ₇₆₉₇ galU galK λ^- rpsL endA1 nupG	BRL
B. subtilis		
1012	leuA8 metB5 trpC2 hsrM1	[26]
WW02	1012 amyE::neo	[27]
AL02	1012 des::neo	This work
AL03	1012 amyE::Pdes-lacZ	This work
AL04	AL02 des::neo amyE::Pdes-lacZ	This work
AL05	AL02 htpG::erm	This work
AL06	AL02 pbpE::erm	This work
Plasmids		
pBluescript SK ⁺	Cloning vector	Stratagene
pBgaB	Integration vector containing the $bgaB$ gene	[28]
pDG1728	Vector allowing integration of DNA sequences at the <i>amyE</i> locus	[16]
pHT01	Derivative of pNDH33 without a direct repeat	[22]
pKTH10	Recombinant vector containing the $amyQ$ gene	[17]
pAL10	Expression vector allowing cold-inducible intracellular production of recombinant proteins	This work
pAL12	Expression vector allowing cold-inducible secretion of recombinant proteins	This work
pNDH33-htpG	htpG fused to an IPTG-inducible promoter	[22]

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