



Chapter 6

Manipulation of Gene Expression in Prokaryotes



6.1. Gene Expression from Strong and Regulatable Promoters

Gene Expression from Strong and Regulatable Promoters

- Constitutive expression of foreign protein
 - Can be detrimental to the host because of the energy drain
 - Can cause plasmid instability
- Regulatable promoters
 - *E. coli lac, trp, tac* (-10 *lac* + -35 *trp*) promoter
 - Bacteriophage λ *pL* promoter
 - Bacteriophage T7 gene10 promoter

Lac Operon

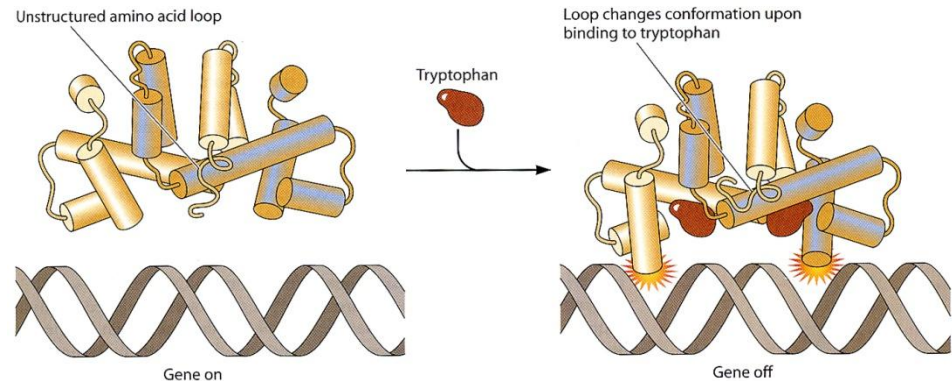
- Genes: *LacZ*, *LacY*, *LacA* for lactose utilization
- Repressor: *LacI*
 - Represses gene expression by binding to lac operator
 - DNA binding activity is inhibited by binding to lactose or isopropyl- β -D-thiogalactopyranoside (IPTG)
- Activator: CAP (catabolite activator protein)
 - Activates gene expression by binding to the promoter
 - DNA binding activity is increased by cAMP binding under low glucose conditions

Lac Promoter

Glucose concentration	Lactose concentration	cAMP concentration	<i>lac</i> promoter-operator region	Level of transcription
Low	Low	High	<p>The diagram shows the <i>lac</i> promoter-operator region with three segments: Promoter (yellow), Operator (purple), and Gene (orange). A CAP protein (red circle) bound to cAMP (grey oval) is bound to the Promoter. A lac repressor (red rounded rectangle) is bound to the Operator.</p>	Low
High	Low	Low	<p>The diagram shows the <i>lac</i> promoter-operator region with three segments: Promoter (yellow), Operator (purple), and Gene (orange). A lac repressor (red rounded rectangle) is bound to the Operator.</p>	Low
High	High	Low	<p>The diagram shows the <i>lac</i> promoter-operator region with three segments: Promoter (yellow), Operator (purple), and Gene (orange). No proteins are bound to the region.</p>	Low
Low	High	High	<p>The diagram shows the <i>lac</i> promoter-operator region with three segments: Promoter (yellow), Operator (purple), and Gene (orange). A CAP protein (red circle) bound to cAMP (grey oval) is bound to the Promoter. A lac repressor (red rounded rectangle) is bound to the Operator. RNA polymerase (red oval) is bound to the Promoter and has an arrow pointing to the right, indicating transcription.</p>	High

Trp Promoter

- Genes: genes for Trp biosynthesis
- Trp Repressor
 - Binding to the operator in the presence of Trp
 - Leaky expression



- *tac*, *trc* promoter
 - -35 trp promoter --- 16 (*tac*) or 17 (*trc*) bp --- -10 lac promoter
 - IPTG inducible
 - Stronger and more effective than the original promoters

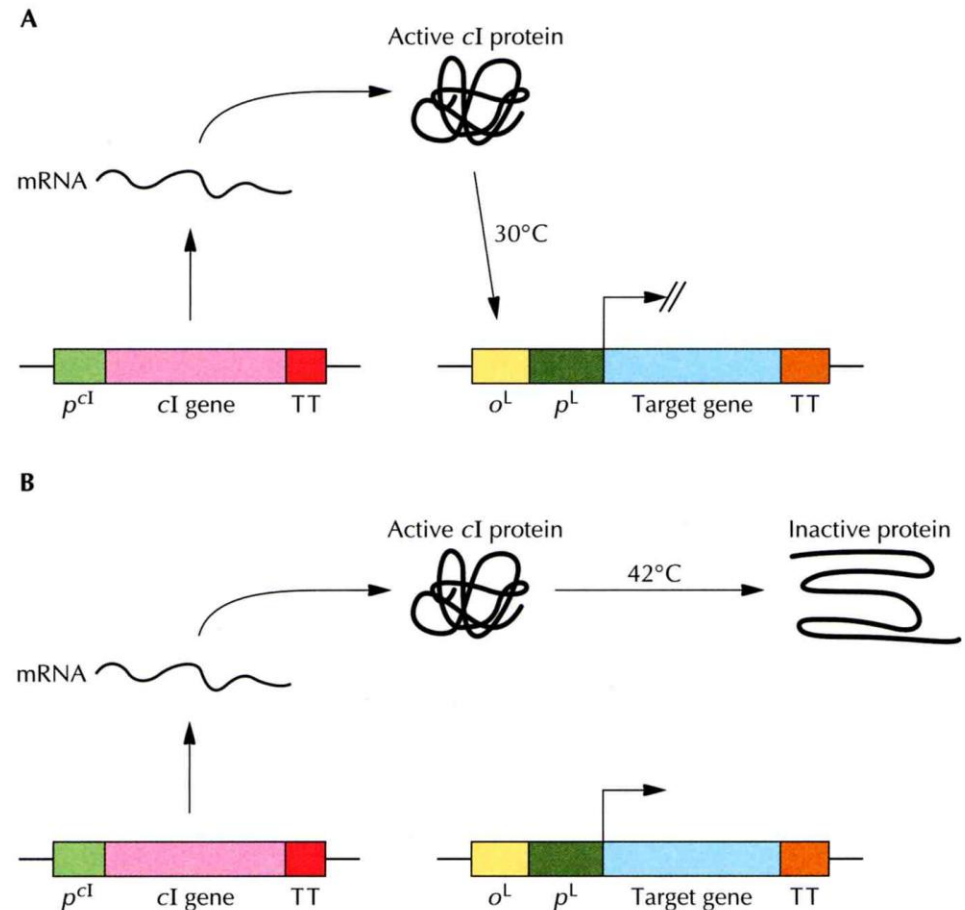
Bacteriophage λ P^L Promoter

■ Repressor : cI

□ cI857:

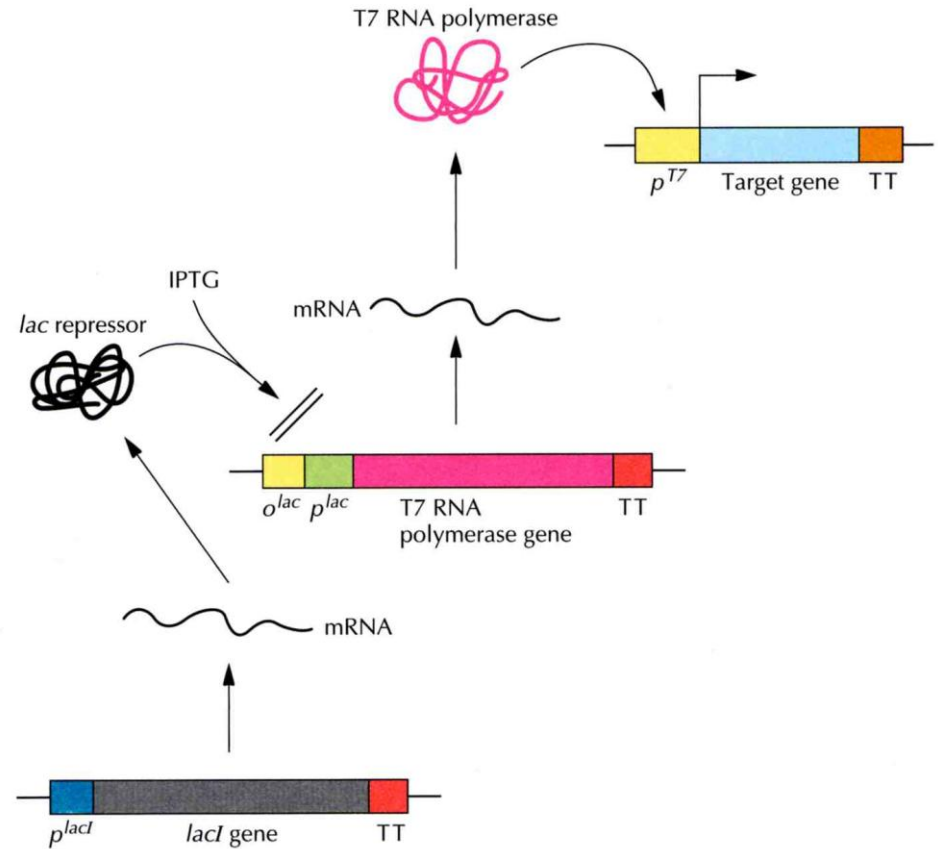
temperature-sensitive cI

- Active at low temperature (28~30°C)
- Inactive at high temperature (42°C)

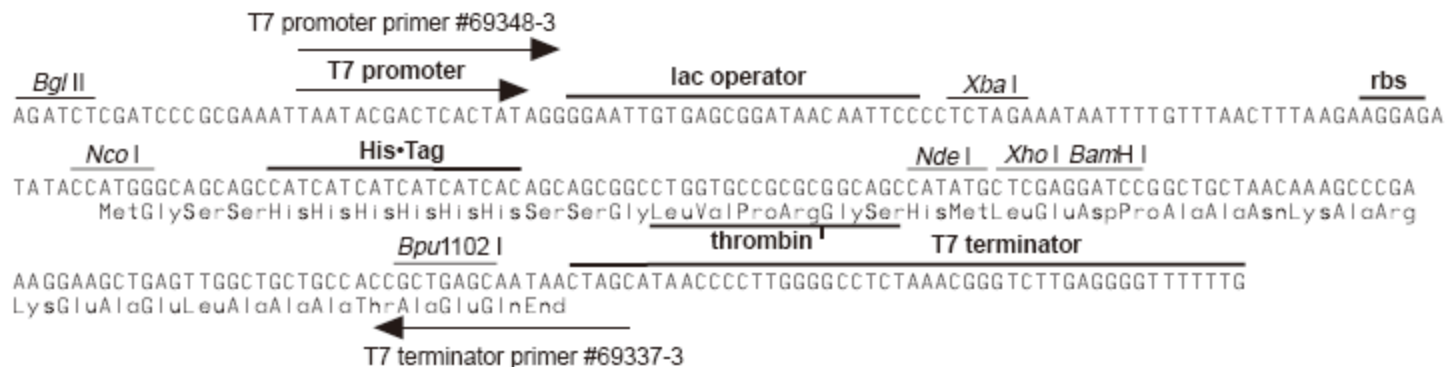
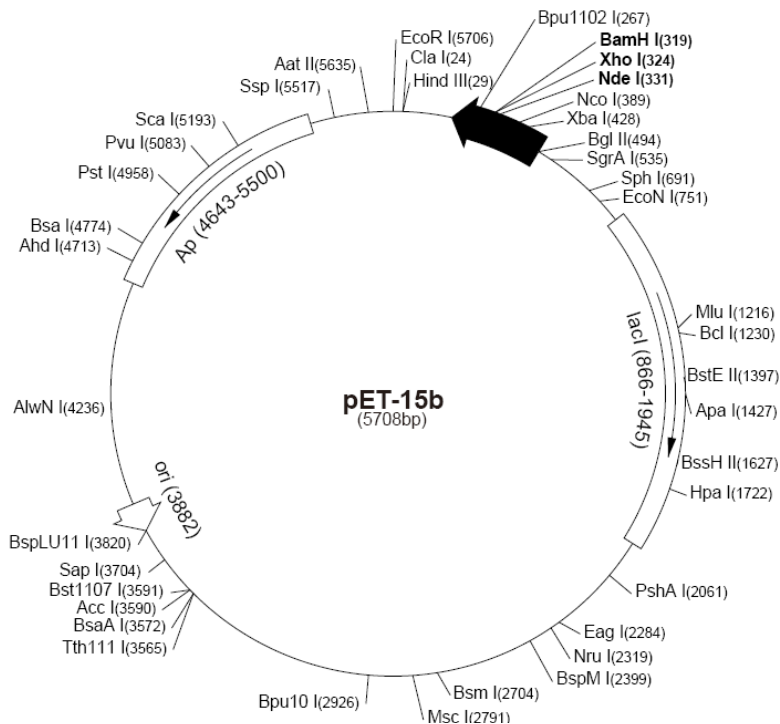


T7 Promoter

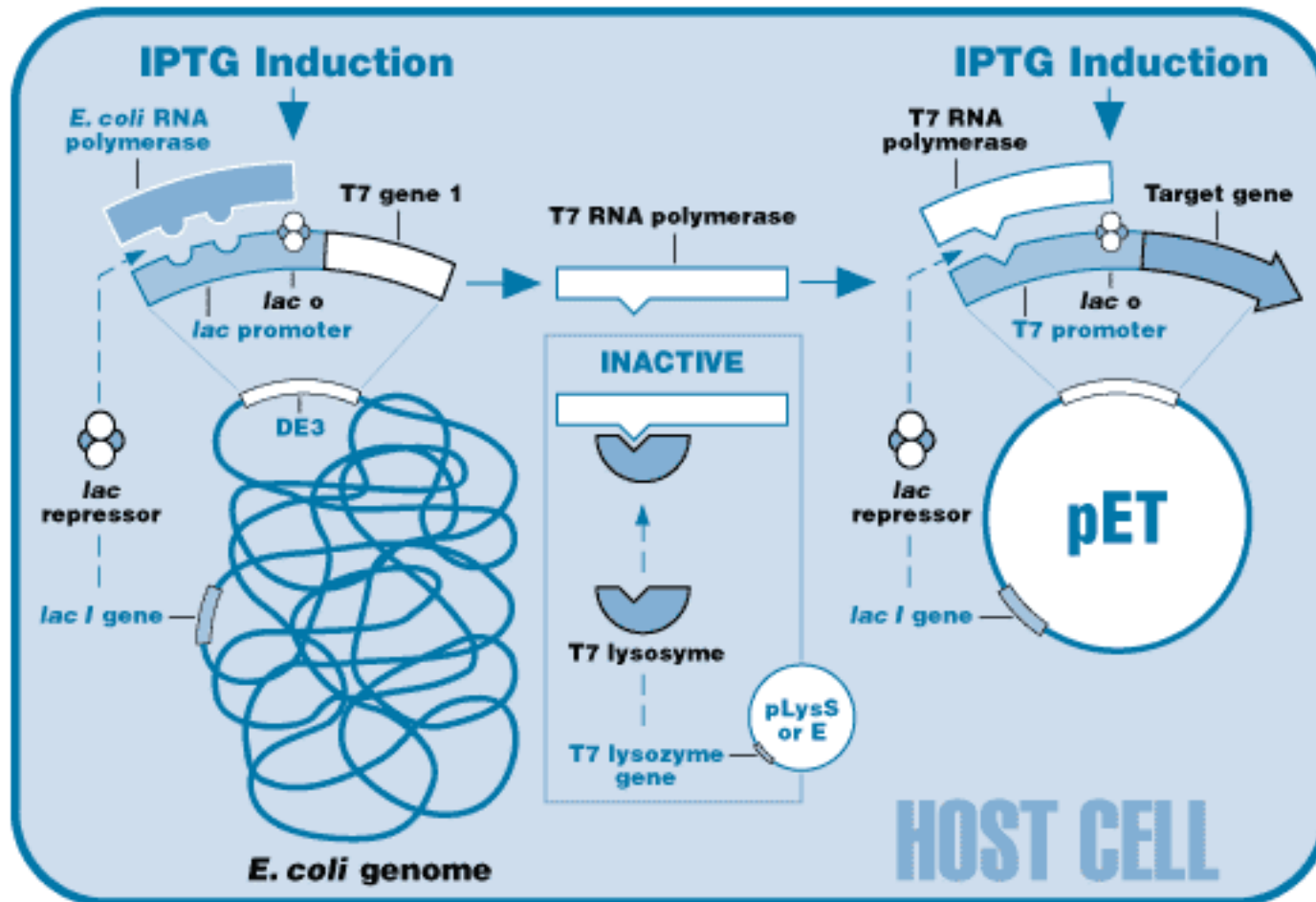
- T7 promoter
 - Transcription by T7 polymerase
- T7 polymerase under the control of the *lac* promoter
 - Induction of T7 promoter-regulated genes by IPTG



pET Expression Vector



PET Expression System



E. coli DE3 (pLysS)

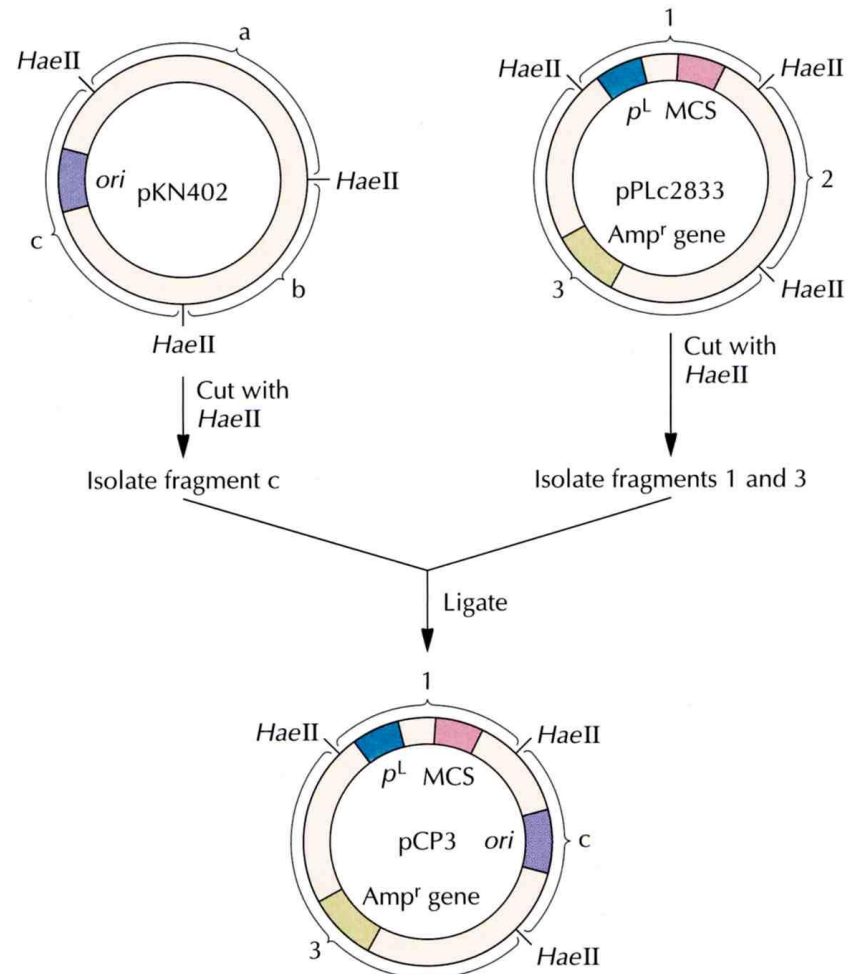
Temperature-Dependent Regulation of Plasmid Copy number

■ pKN402

- Increase in copy number at high temperature
- 28°C: 82 → 42°C: 521

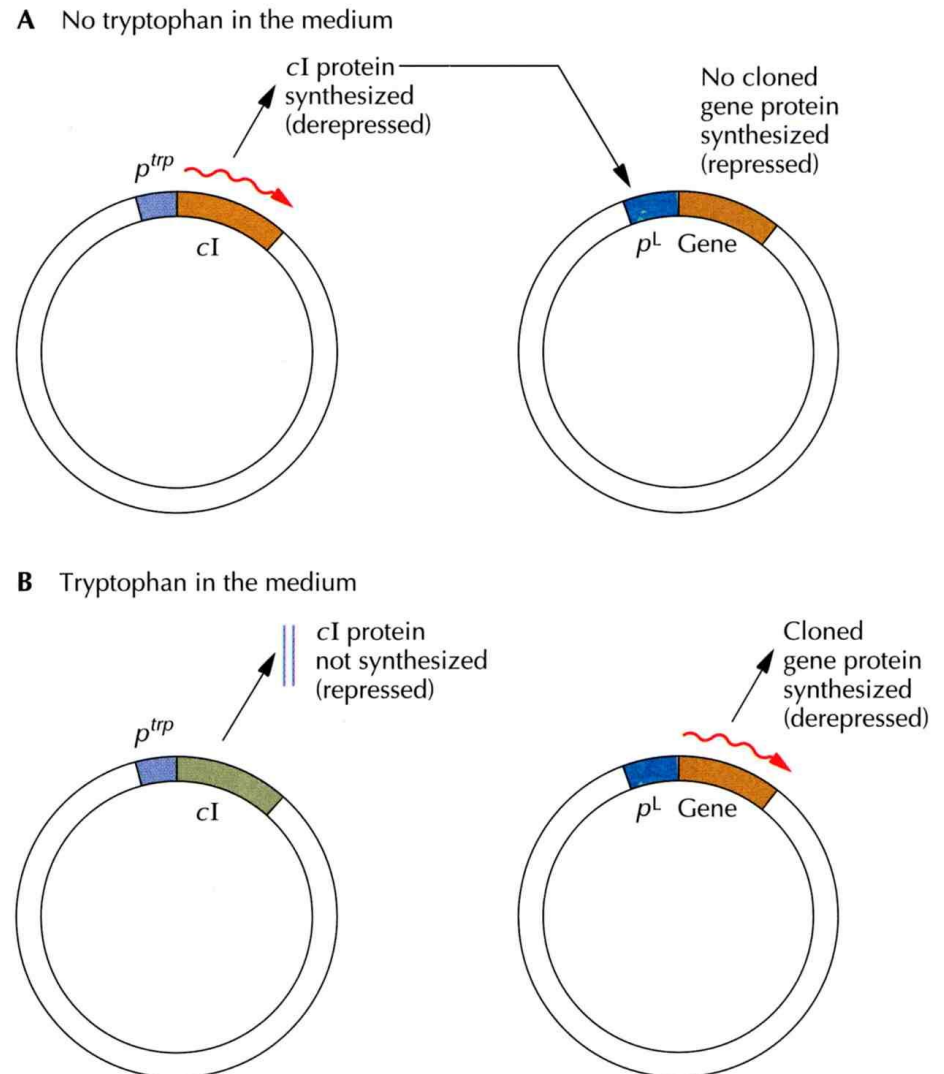
■ pCP3

- Contains pKN402 replication origin
- High level protein expression at 42°C



Dual Plasmid System

- Two plasmids
 - *cI* under the control of *trp* promoter
 - Protein expression under the control of p^L promoter
- Control gene expression
 - Without Trp
 - Repression of protein expression
 - With Trp
 - Induction of protein expression
- Inexpensive system for large scale protein production



Expression in Other Microorganisms

- Universal gram (-) bacterial expression vector
 - Tn5 promoter in a broad-host-range plasmid pRK290
 - Efficient gene expression in different bacterial hosts
- Modification of promoter strength
 - *Lactococcus lactis* constitutive promoter
 - Screening strong promoters from spacer region (between -35 and -10) library

TTGACNNNNNNNNNNNNNNNTGRTTATAAT



6.2. Fusion Proteins



Fusion Proteins

■ Fusion protein

- In frame fusion of a target protein with a stable host protein
- Resistance to proteolysis

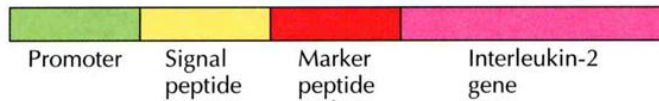
■ Cleavage of fusion proteins

- Linkage of two proteins with nonbacterial protease recognition sequence
- Ile-Glu-Gly-Arg : C-terminal end cleavage by factor X_a

Uses of Fusion Proteins

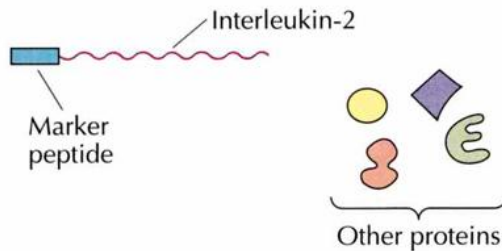
- Raising antibody
- Simplifying purification
 - His tail (6-10 aa): Ni²⁺ binding, elution by imidazole
 - Strep-tag (10 aa): Streptavidin binding, elution by iminobiotin
 - MBP (40 kDa): Amylose binding, elution by maltose
 - GST (25 kDa): Glutathione binding, elution by reducing agent
 - Flag (8 aa): Flag Ab binding, elution by low calcium
 - ZZ (14 kDa): IgG binding, elution by low pH

Purification of Fusion Proteins by Immunoaffinity Chromatographic Purification

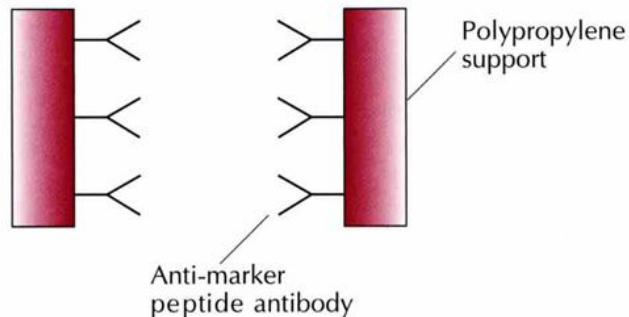


N-Asp-Tyr-Lys-Asp-Asp-Asp-Lys-C

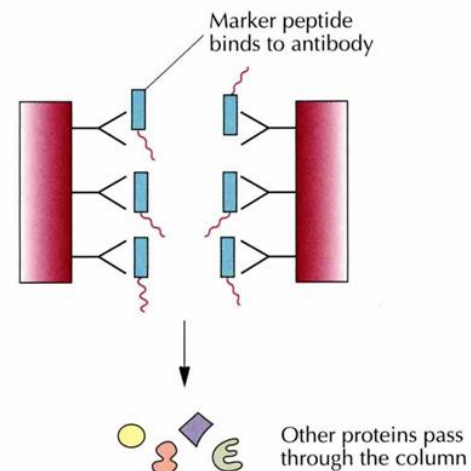
1 Concentrate secreted protein mixture



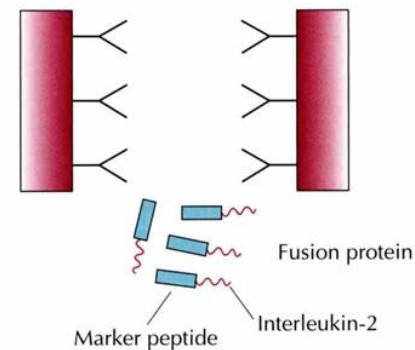
2 Prepare immunoaffinity column



3 Add secreted protein mixture to the column

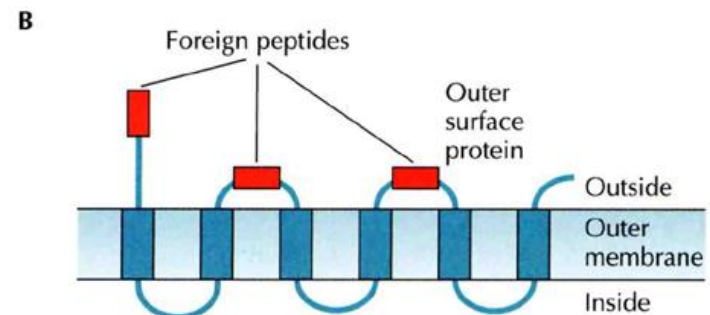
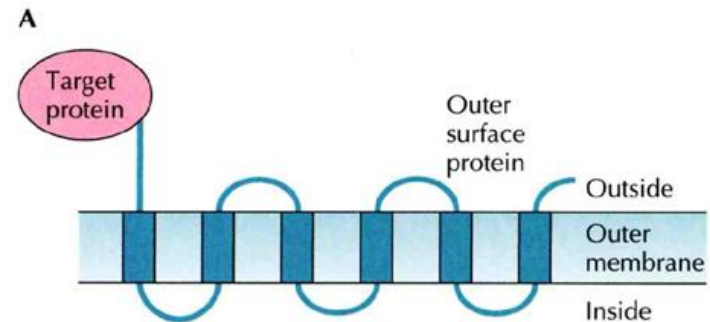
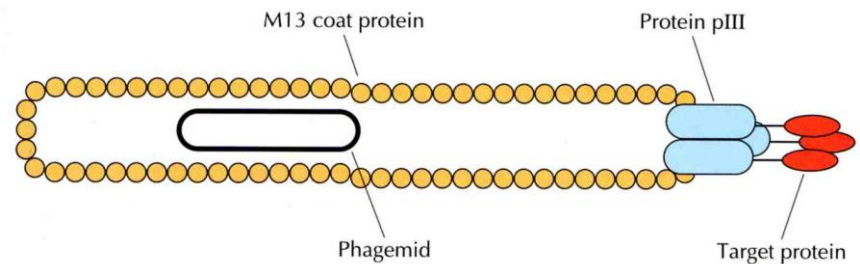


4 Elute fusion protein



Surface Display

- Display proteins as fusion proteins with a surface proteins (filament, pilus protein)
 - Phage display
 - Protein pIII of M13 phage
 - Bacterial display
 - *E. coli* OmpA, OmpF : outer membrane protein
 - *E. coli* PAL: peptidoglycan-associated lipoprotein
 - Target protein in N or C terminus, or in the middle



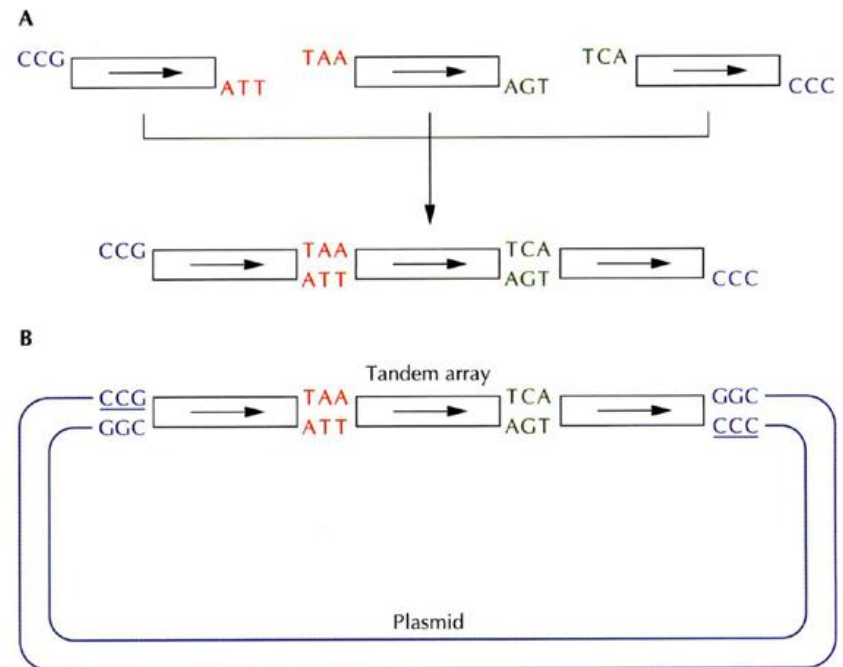
Usage of Surface Display

- Screening of cDNA libraries
- Overexpression of peptides or proteins
 - Expression of antigenic determinant of the parasite *Plasmodium falciparum* (causing malaria) by inserting into surface-exposed loops of the major outer membrane protein from *P. aeruginosa* (OprF)
 - Possible usage as vaccines

Unidirectional Tandem Gene Array

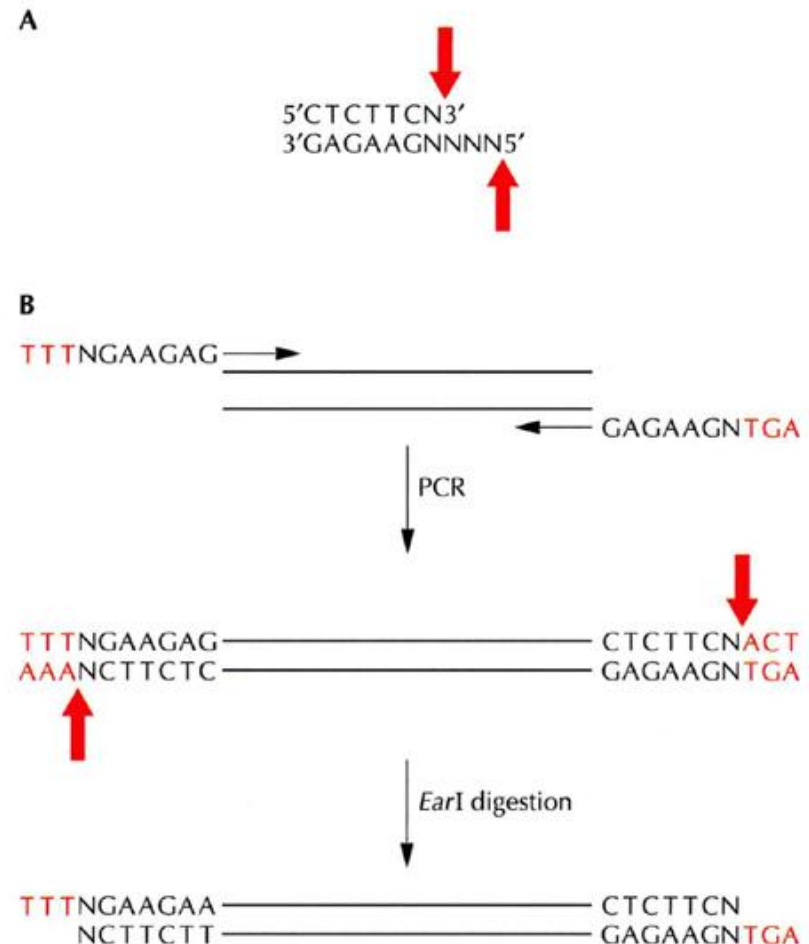
■ Protein expression levels

- Multiple copies of genes in a low-copy-number plasmids > Single copy of gene in a high-copy-number plasmid
- Generation of tandem gene array
 - Using complementary 3-nucleotide extensions



Unidirectional Tandem Gene Array

- TypellS restriction enzyme
 - Cut random sequence outside of the recognition site
- Use pfu polymerase for PCR





6.3. Translation Expression Vectors



Translation Expression Vectors

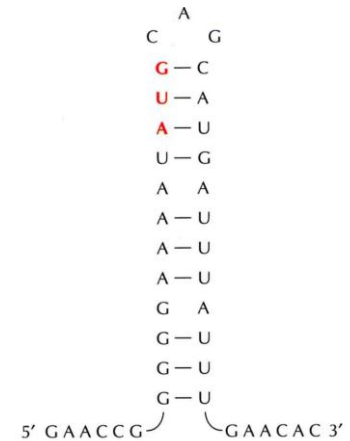
■ Translation efficiency

□ Ribosome binding site

- UAAGGAGG
- Precise distance from the starting codon
- No secondary structure formation preventing ribosome binding

□ Codon usage

- Rarely used codons in *E. coli* (AGG, AGA, AUA, CUC, CGA)
- Solutions
 - Expression in eukaryotes
 - Change codons
 - Provide tRNAs for rare codons
 - Strain is commercially available



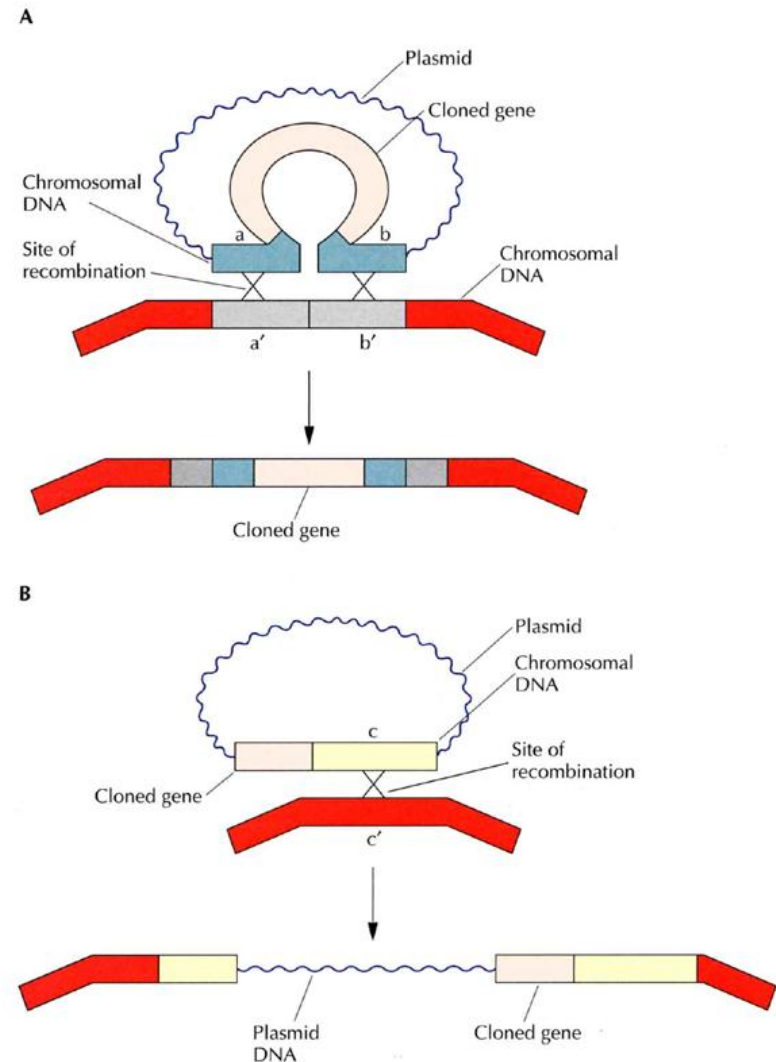


6.4. DNA Integration into the Host Chromosome



DNA Integration into the Host Chromosome

- DNA integration into chromosome
 - Stable expression
 - Integration into non-essential gene
 - Homologous recombination through chromosomal sequence
 - Double cross over or
 - Single cross over
 - Use nonreplicating plasmid for integration



Multiple Integration

- Expression of α -amylase in *B. subtilis*
 - Integration of plasmid with α -amylase gene and chloramphenicol resistance marker on *B. subtilis* chromosome
 - Isolation of clones with multiple integration by selection under high chloramphenicol

Copies/genome

α -amylase activity (U/ml)

2

500

5

2,300

7

3,100

9

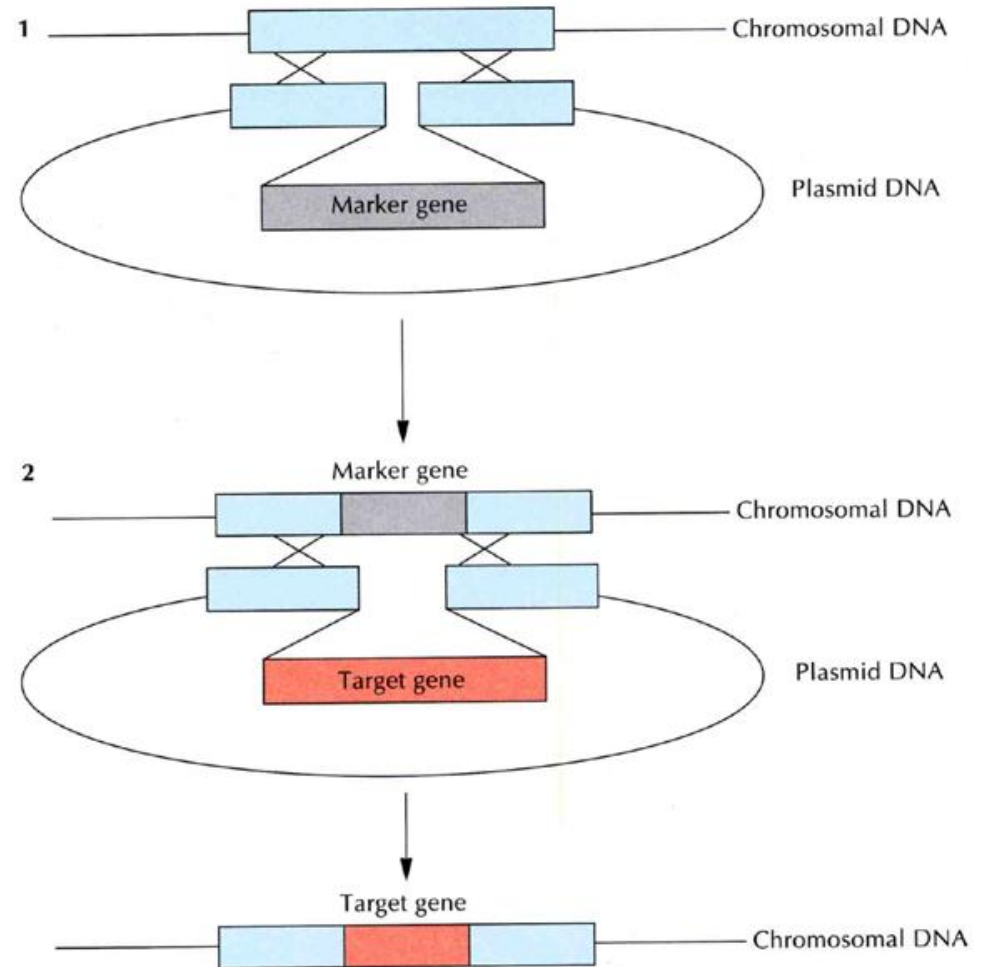
4,400

Multicopy plasmid

700

Multiple integration at specific sites

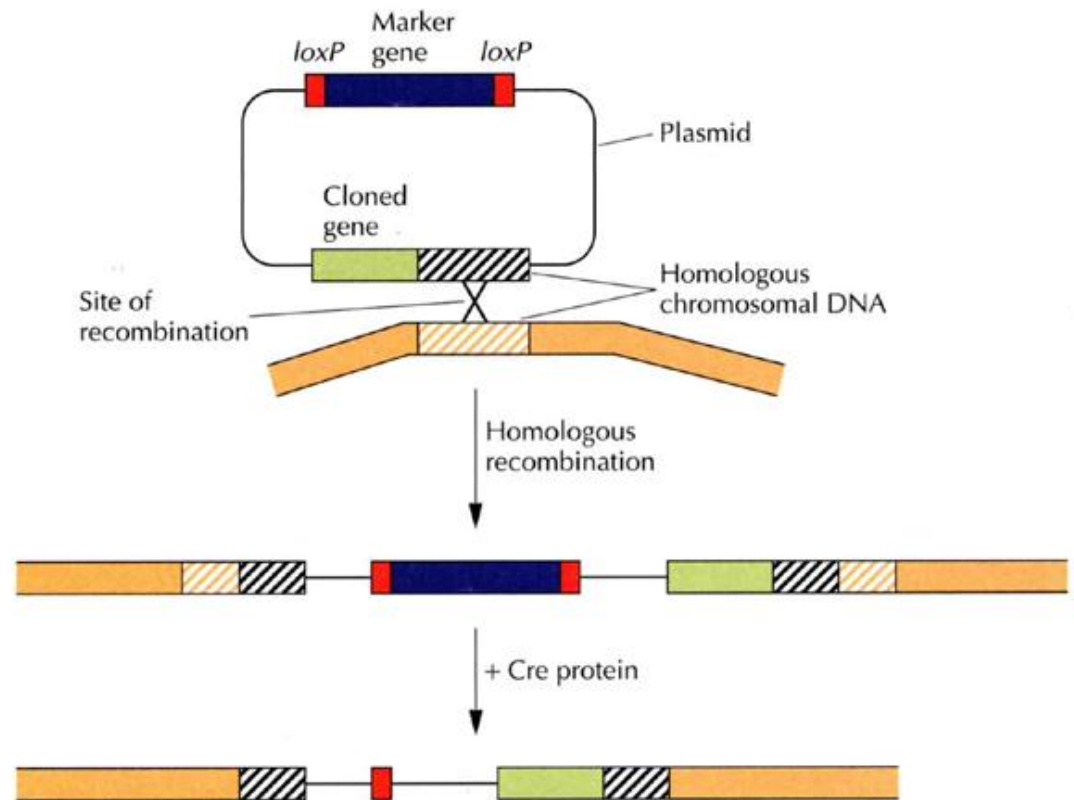
- Integration of a marker
- Replacement of the marker with a target gene



Removing Selectable Marker Genes

■ Cre-LoxP system

- Cre recombinase
- LoxP site: 34-bp recombination sites
- Removal of a marker flanked by loxP site by expression of Cre enzyme (under the control of lac promoter)





6.5. Protein Stability, Folding, and Secretion



Increasing Protein Stability

- Protein half life

- A few minutes to hours

- Factors affecting protein stability

- N terminal amino acid

- M, S, A, T, V, G at N terminus of β -galactosidase: > 20 h

- R: ~ 2 min

- Internal PEST sequence

- Facilitate degradation

Protein Folding

■ Inclusion body

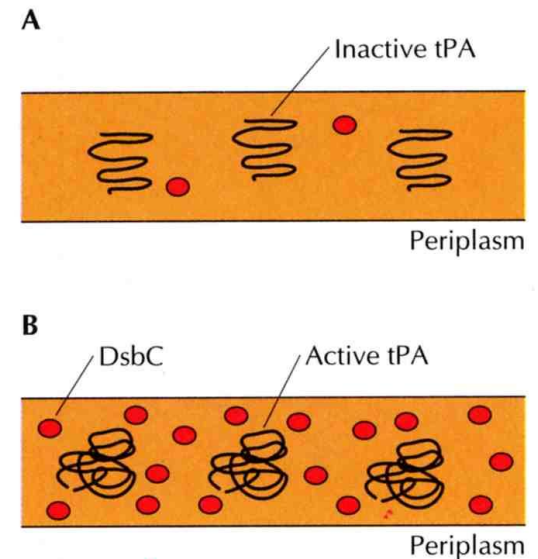
- Insoluble aggregates of expressed protein

■ Strategies to prevent inclusion body formation

- Tagging with other proteins
 - Thioredoxin, GST

- Proteins with multiple disulfide bonds

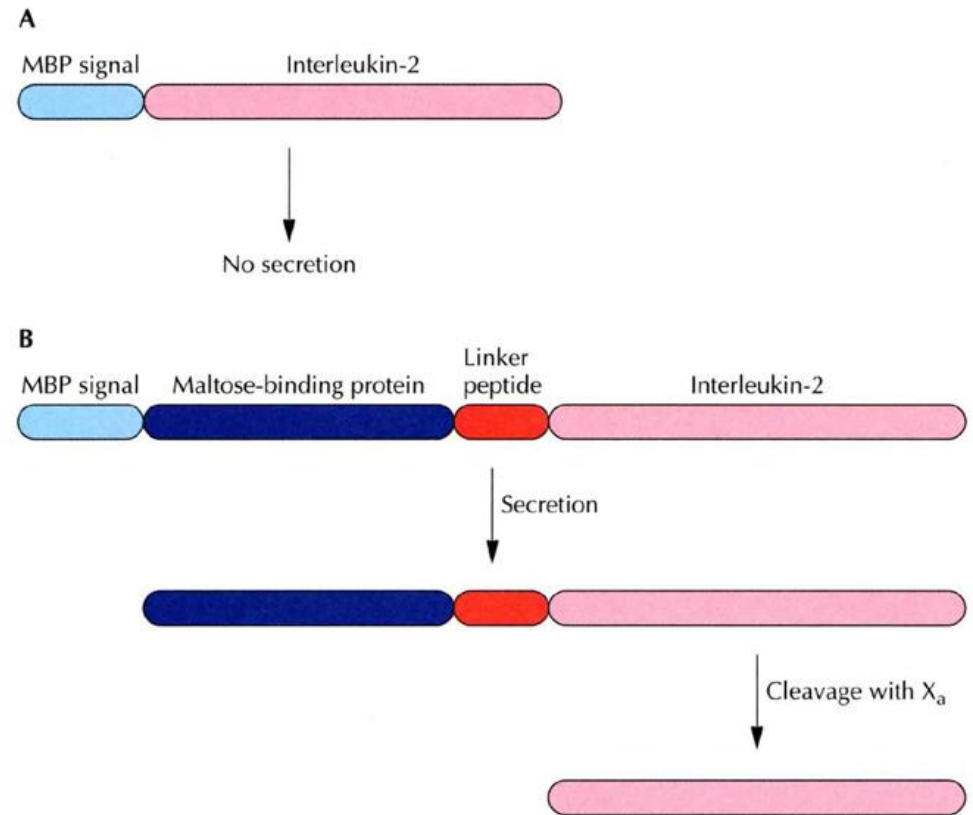
- Expression with signal peptide for secretion into periplasmic space
- Coexpression of high levels of periplasmic DsbC enzyme involved in disulfide bond formation



tPA: tissue plasminogen : 17 S-S bonds

Increasing Secretion

- Advantages of secretion of a target protein
 - Protein stability
 - Proinsulin: more stable in periplasm
 - Easy to purify
- Protein secretion
 - Addition of signal peptide
 - Fusion with a secretory protein



Permeabilization of *E. coli*

- Inducible expression of bacteriocin release protein
 - Activation of phospholipaseA
 - Permeabilization of inner and outer membrane
- Translational overload
 - Inhibition of protein secretion
 - Expression of limiting components protein secretion pathway

Bacteria with Increased Permeability

■ L-form bacteria

- Bacterial lacking cell wall
- Generation of L-form bacteria
 - Spontaneous mutations
 - Treatment with penicillin
 - Inhibition of the final step in cell wall biosynthesis
 - Treatment with lysozyme
 - Hydrolysis of saccharide linkage
- High yield of protein secretion

■ Spheroplasts, protoplasts

- Complete loss of cell wall



6.6. Production of Recombinant Proteins



Overcoming Oxygen Limitation

- Oxygen limitation
 - Slow growth
 - Enter stationary phase
 - Protease production
- Use of protease-deficient host strains
 - Proteases are also required for cell growth
 - rpoH (heat shock sigma factor) and degP (protease for high temperature growth) mutants
 - Decrease in protein degradation of secretory proteins
- Bacterial hemoglobin
 - Hemoglobin-like molecule in *Vitreoscilla* bacterium
 - Expression in *E. coli* to increase protein synthesis

Metabolic Load

- Impairment of normal cellular function of host cells by expression of foreign DNA
 - Replication and maintenance of high copy number plasmid
 - Limitation of dissolved oxygen
 - Depletion of certain aminoacyl-tRNAs and/or drain energy
 - Prevent proper localization of host proteins by foreign secretory proteins
 - Interference of host cell function by foreign proteins

Effects of a Metabolic Load

- Decrease in cell growth rate
 - Loss of plasmid or a portion of plasmid DNA
- Decrease in energy-intensive metabolic processes
 - Nitrogen fixation
 - Protein synthesis
- Changes in cell size and shape
- Increase in extracellular polysaccharide production
- Increase in translational errors



Prevention of Metabolic Load

- Prevention of metabolic load
 - Low copy number plasmid for expression
 - Integration of DNA into host chromosome
 - Inducible promoter
- Optimization for the maximum yield
 - Protein expression levels
 - Cell density

Increasing Cell Density

- Elimination of growth inhibitory waste products
 - e.g. prevention of acetate formation
 - Use glucose analogue α -glucoside to reduce glucose uptake
 - Use ptsG (enzyme II in the glucose phosphotransferase system) mutant
 - Expression of acetolactase synthase

