Biotechnologie Moléculaire 3ème année

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Manipulation of Gene Expression in Prokaryotes

Gene Expression from Strong and Regulatable Promoters The *E. coli lac* promoter



Effect of the concentrations of glucose, lactose and cAMP on the level of transcription (cAMP concentration is high when the glucose level is low)

IPTG, a non-metabolisable analogue of lactose, can also be used for induction

Gene Expression from Strong and Regulatable Promoters Regulation of gene expression by the phage λp^{L} promoter



Gene Expression from Strong and Regulatable Promoters Dual plasmid systems for controlling the phage λp^{L} promoter



Gene Expression from Strong and Regulatable Promoters

Regulation of gene expression by the bacteriophage T7 promoter



Gene Expression from Strong and Regulatable Promoters Expression in microorganisms other than *E. coli*

Promoter	β-Galactosidase activity (U) in:				
	Escherichia coli	Rhizobium meliloti	Rhizobium leguminosarum	Pseudomonas putida	
None	16	110	130	150	
Nm	1,400	21,800	13,900	16,300	
lac	2,000	9,050	6,250	9,800	
tac	11,300	2,850	1,150	2,950	
S1	40	3,300	1,200	3,350	

Adapted from Labes et al., Gene 89:37-46, 1990.

 β -galactosidase expressed by various gram-negative bacteria

Gene Expression from Strong and Regulatable Promoters

"Universal" gram-negative bacterial expression vectors



Uses two overlapping promoters (p) from transposon 5 (Tn5). Allows efficient expression in *E. coli* and 6 other species

DNA constructs can be made which express a foreign protein fused to another protein (on a single polypeptide)

This can be useful to:

- Improve the expression level of the foreign protein
- Facilitate purification of the foreign protein

Fusion Proteins An example of a fusion protein cloning vector



After transcription and translation a tribrid protein is produced comprising:

 The N-terminal region of the outer membrane protein (*ompF*)
 The cloned gene (gene)
 A truncated β-galactosidase gene (*lacZ*)

To facilitate the purification of foreign proteins in *E. coli*

Fusion partner	Size	Ligand	Elution condition
77.	14 kDa	IgG	Low pH
His tail	6–10 aa	Ni ²⁺	Imidazole
Strep-tag	10 aa	Streptavidin	Iminobiotin
PinPoint	13 kDa	Streptavidin	Biotin
MBP	40 kDa	Amylose	Maltose
B-Lactamase	27 kDa	Phenyl-boronate	Borate
GST	25 kDa	Glutathione	Reducing agent
Flag	8 aa	Specific MAb	Low calcium

Adapted from Nygren et al., Trends Biotechnol. 12:184-188, 1994.

Abbreviations: aa, amino acids; kDa, kilodaltons; ZZ, a fragment of *Staphyloccus aureus* protein A; His, histidine; Strep-tag, a peptide with affinity for streptavidin; PinPoint, a protein fragment which is biotinylated in vivo in *E. coli*; MBP, maltose binding protein; GST, glutathione *S*-transferase; Flag, a peptide recognized by enterokinase; MAb, monoclonal antibody.

To facilitate the purification of foreign proteins in *E. coli*



The genetic construct used to produce a fusion protein consisting of a marker peptide and interleukin-2

To facilitate the purification of foreign proteins in *E. coli*



Increasing Expression Level

The number of plasmids per cell (the copy number) can be increased However the higher the copy number, the more of the cell's resources are diverted to the production of plasmid encoded proteins and the metabolic activities of the host strain are constrained.

An alternative is to express unidirectional tandem gene arrays cloned in a single plasmid.

Increasing Expression Level

Unidirectional tandem gene arrays



The individual gene fragments are typically generated by PCR using primers that append Type IIS restriction enzyme sites which can be cleaved to create the sticky ends (as in Golden Gate cloning)

Unfortunately tandem gene arrays are often not genetically stable due to homologous recombination causing deletions

Using a strong promoter to optimise transcription of mRNA may not be sufficient to maximise the yield of the cloned product.

It may also be necessary to optimise:

- 1. Translation of the protein
- 2. Stability of the protein

For example, the expression vector pKK233-2 contains:

- An ampicillin resistance gene as a selectable marker
- The strong *tac* promoter
- The efficient *lac*Z ribosome-binding site (rbs)
- An ATG start codon optimally located 8 nucleotides downtream from the ribosome-binding site
- The transcription terminators T1 and T2 from bacteriophage λ to increase mRNA stability



It is also important to ensure that the region comprising the region of mRNA comprising the ribosome binding site and the start codon (AUG) are not sequestered in a double-stranded region, otherwise translation will not be efficient.



Foreign cloned genes often contain codons which are rarely used by *E. coli* (such as AGG, AGA, AUA, CUA and CGA). The host cell therefore my not produce enough transfer RNA to translate these codons reducing protein yield.

This problem can be overcome in several ways:

- 1. If the cloned gene is eukaryotic it can be cloned and expressed in a eukaryotic expression system
- 2. By creating a synthetic gene which only use codons which are commonly used in *E*. *coli* (codon optimisation)
- 3. Using *E. coli* engineered to overexpress rare tRNAs

Increasing Expression Level Using *E. coli* cells engineered to overexpress rare tRNAs



Increasing Expression Level Increasing protein stability

Amino acid added	Half-life	
Met, Ser, Ala Thr, Val, Gly Ile, Glu Tyr, Gln Pro Phe, Leu, Asp, Lys Arg	>20 h >20 h >30 min ~10 min ~7 min ~3 min ~2 min	Stability of β-galactosidase with different amino acids at its N-terminus
0		

Adapted from Bachmair et al., Science 234:179-186, 1986.

Changing the amino acid at the N-terminus of a protein can often significantly change it's half-life in *E. coli*

So called PEST sequences rich in proline (P), glutamate (E), serine (S) and threonine (T) in the protein can make the protein more susceptible to proteolysis and stability of the protein can be improved by genetic manipulation of the PEST sequences.

Increasing Expression Level Protein Folding

Many proteins overexpressed in *E. coli* accumulate in insoluble, intracellular, biologically inactive inclusion bodies



Active protein can often be recovered from inclusion bodies after using protein solubilisation and refolding procedures.

However, in many cases it is preferable to produce soluble, active protein from the beginning.

There are a number of ways to achieve this.

Increasing Expression Level Producing folded protein by making a fusion protein



For example, proteins fused to the 11.7 kDa protein Thioredoxin can remain soluble even when up to 40% of the cellular protein consists of the fusion protein.

Increasing Expression Level Producing folded disulfide bonded proteins

Periplasmic Space



Disulfide bonds do not form in the reducing environment of cytoplasm.

They do, however, form in the oxidising environment of the *E*. *coli* periplasm or the eukaryotic endoplasmic reticulum.

Increasing Expression Level Producing folded disulfide bonded proteins



Tissue plasminogen activator (tPA), an important therapeutic "clot-buster" drug, contains 17 disulfide bonds.

Expression of active tPA in *E. coli* required that it be directed to the periplasm by fusion of a leader peptide and that disulfide bond formation be catalysed by overproduction of the *E. coli* enzyme DsbC.

Disulfide bonds do not form in the reducing environment of cytoplasm.

They do, however, form in the oxidising environment of the *E*. *coli* periplasm or the eukaryotic endoplasmic reticulum.

Heterologous Protein Production in Eukaryotic Cells

Eukaryotic Proteins in Prokaryotes Where is the problem?

Prokaryotic expression systems sometimes fail to produce functional versions of eukaryotic proteins

This is mostly due to:

1. Improper folding

2. Inability to modify certain amino acid side chains (post-translational modification), e.g.

• To form disulphide bonds: in eukarotes this is catalysed by the enzyme protein disulphide isomerase (PDI). Aberrant disulfide bond formation changes the configuration of the protein and can reduce stability and abolish activity.

• To add sugars (glycosylation) – about 30% of all mammalian proteins are glycosylated

Post-Translation Modification in Eukaryotes Glycosylation

A T/S — T/S Yeast — T/S ----- T/S В T/S Insects - T/S C ■− T/S -T/SMammals - T/S - T/S

S, serine; T, threonine; red circles, mannose; dark blue squares, N-acetylglucosamine; light blue squares, N-acetylgalactosamine; green squares, galactose; orange squares, sialic acid.

Examples of O-linked glycosylation (of Serine or Threonine) in different organisms

Post-Translation Modification in Eukaryotes Glycosylation



Examples of N-linked glycosylation (of Asparagine) in different organisms

Post-Translation Modification in Eukaryotes Glycosylation



Examples of N-linked glycosylation (of Asparagine) in different organisms

Eukaryotic Expression Vectors General features



A generalized eukaryotic expression vector.

Figure 7.3 Generalized eukaryotic expression vector. The major features of a eukaryotic expression vector are a eukaryotic transcription unit with a promoter (p), a multiple cloning site (MCS) for a gene of interest, and a DNA segment with termination and polyadenylation signals (t); a eukaryotic selectable marker (ESM) gene system; an origin of replication that functions in the eukaryotic cell (ori^{euk}); an origin of replication that functions in the eukaryotic cell (ori^{euk}); an origin of replication that functions in $E. \ coli \ (ori^{E})$; and an $E. \ coli \ selectable \ marker \ gene \ (Amp^r)$.

In principle, eukaryotic expression vectors do not differ from prokaryotic expression vectors

However, they contains an *E*. *coli* (*ori*^E) and a eukaryotic (*ori*^{euk}) origin of replication and *E*. *coli* and eukaryotic selectable markers

Sacharomyces cerevisiae Expression Systems Advantages

Yeast is a useful expression system because:

1. It is single celled, well characterized genetically and physiologically, and grown readily in small cultures and large-scale bioreactors.

2. Several strong promoters are known, and yeast plasmids can be constructed based on the naturally occurring yeast $2\mu m$ plasmid.

3. Yeast can carry out many post-translational modifications.

4. Yeast secretes few proteins naturally, so extracellular proteins engineered for secretion in yeast are already pretty pure, aiding purification.

5. As it is used for baking and brewing it is classed as "generally recognized as safe" by the U.S. Food and Drug Administration.

Sacharomyces cerevisiae Expression Systems Recombinant proteins produced in S. cerevisiae

VACCINES

Hepatitis B virus surface antigen Malaria circumsporozoite protein HIV-1 envelope protein

DIAGNOSTICS Hepatitis C virus protein HIV-1 antigens

 HUMAN THERAPEUTIC AGENTS
 Epidermal growth factor
 Insulin
 Insulin-like growth factor
 Platelet-derived growth factor
 Proinsulin
 Fibroblast growth factor
 Granulocyte-macrophage colonystimulating factor
 α₁ antitrypsin
 Blood coagulation factor XIIIa
 Hirudin
 Human growth factor
 Human serum albumin

Sacharomyces cerevisiae Expression Systems Classes of S. cerevisiae expression vectors

There are three main classes of *S. cerevisiae* expression vectors:

- 1. Yeast episomal plasmids (YEps).
- 2. Yeast integrating plasmids (YIps).
- 3. Yeast artificial chromosomes (YACs).

Sacharomyces cerevisiae Expression Systems Yeast episomal plasmids (YEps)



YEps work well, but are often unstable under large-scale (≥10 litres) growth conditions. This example contains:

1. A yeast gene for leucine biosynthesis (*LEU2*) – a yeast selectable marker.

2. A 2 μ m plasmid origin of replication – to allow replication of the plasmid in yeast.

3. The ampicillin resistance gene (Amp^r) – a bacterial selectable marker

4. An *E*. *coli* origin of replication (ori^E) – to allow replication of the plasmid in bacteria.

5. The cDNA for the human enzyme Cu/Zn superoxide dismutase (Cu/Zn-SOD) – the foreign protein to be expressed.

6. The promoter (GAPDp) and terminationpolyadenylation sequences (GAPDt) of the *S*. *cerevisiae* gyceraldehyde phosphate dehydrogenase gene.

Sacharomyces cerevisiae Expression Systems Promoters for S. cerevisiae expression vectors

Promoter	Expression conditions	Status
Acid phosphatase (PH05)	Phosphate-deficient medium	Inducible
Alcohol dehydrogenase I (ADHI) Alcohol dehydrogenase II (ADHII) Cytochrome c ₁ (CYC1) Gal-1-P Glc-1-P uridyltransferase Galactokinase (GAL1)	2–5% Glucose 0.1–0.2% Glucose Glucose Galactose Galactose	Constitutive Inducible Repressible Inducible Inducible
Glyceraldehyde-3-phosphate dehydrogenase (<i>GAPD</i> , <i>GAPDH</i>) Metallothionein (<i>CUP1</i>) Phosphoglycerate kinase (<i>PGK</i>) Triose phosphate isomerase (<i>TPI</i>) UDP galactose epimerase (<i>GAL10</i>)	2–5% Glucose 0.03–0.1 mM copper 2–5% Glucose 2–5% Glucose Galactose	Constitutive Inducible Constitutive Constitutive Inducible

Sacharomyces cerevisiae Expression Systems Yeast integrating plasmids (YIps)



This vector contains:

1. A yeast gene for leucine biosynthesis (LEU2) – a yeast selectable marker.

2. The Gene of Interest (GOI) with transcription and translation control elements

3. The ampicillin resistance gene (Amp^r) – a bacterial selectable marker.

4. An *E*. *coli* origin of replication (ori^E) – to allow replication of the plasmid in bacteria.

5. Two segments from the ends of a non-essential yeast gene (A1 and A2).

Note, there is no yeast origin of replication. The GOI and LEU2 are only replicated if they are integrated into the yeast chromosome by homologous recombination.

The system is genetically stable, but protein yield is low as there is only a single copy.

Sacharomyces cerevisiae Expression Systems Yeast artificial chromosomes (YACs)



The pYAC plasmid contains:

1. The ampicillin resistance gene (Amp^r) – a bacterial selectable marker.

2. An *E*. *coli* origin of replication (ori^E) – to allow replication of the plasmid in bacteria.

3. A yeast autonomous replicating sequence (ARS) – allows replication of the YAC in yeast.

4. A centromere (CEN) – to enable the YAC to segregate like a chromosome.

5. A gene from the uracil biosynthesis pathway (URA3) – a selectable marker in yeast.

6. A gene from the tryptophan biosynthesis pathway (TRP1) – a selectable marker in yeast.

7. Two yeast chromosome telomere sequences (T) – to allow the YAC to be maintained stably.

Large DNAs (>100 kb) can be stably maintained as a separate chromosome in double mutant *ura*3 and *trp*1 cells.

Pichia pastoris and Other Yeast Expression Systems

S. cerevisiae is not always ideal:

- Yields of protein may be poor.
- Proteins may be hyperglycosylated (>100 mannose residues in N-linked side chains.
- Proteins designed for secretion may be maintained in the periplasmic space.
- Ethanol is produced at high cell densities, which is toxic to the cells, reducing protein yield.

Pichia pastoris (a methylotropic yeast) has a number of advantages:

- It has a highly efficient, tightly regulated methanol inducible promoter (the *AOX1* promoter) in the presence of methanol the expressed gene may make up to 30% of cellular protein.
- *P. pastoris* do not make ethanol and can therefore achieve high cell densities with high protein yield.
- *P. pastoris* normally does not secrete many proteins, simplifying the purification of secreted recombinant proteins.

Other yeasts, including *Hansenula polymorpha*, *Shizosaccharomyces pombe* and *Candida utilis* have also been used to make recombinant human proteins.

Several species of the filamentous fungus *Aspergillus* have been used to produce enzymes for the food, beverage, pulp and paper industries.

Pichia Pastoris Expression Systems An integrating *P. pastoris* expression vector



The plasmid contains:

1. The ampicillin resistance gene (Amp^r) – a bacterial selectable marker.

2. An *E. coli* origin of replication (ori^E) – to allow replication of the plasmid in bacteria.

3. The Gene of Interest (GOI).

4. The promoter (AOX1p) and termination-polyadenylation sequences (AOX1t) of the *P. pastoris* alcohol oxidase 1 gene.

5. A gene from the histidine biosynthesis pathway (HIS4) – a selectable marker in *P. pastoris*.

6. The 3'-end of the AOX1 gene of *P. pastoris* – to enable recombination into the chromosomal AOX1 gene.

Pichia Pastoris Expression Systems An integrating *P. pastoris* expression vector



Integration of DNA into a specific *P. pastoris* chromosome by single or double recombination

Baculovirus-Insect Cell Expression Systems Baculoviruses infect invertebrates, including many insect species







In infected cells clusters of nucleocapsids (virions) are trapped (occluded) in a protein matrix.

The protein matrix is called **polyhedrin** and the entire package is called a **polyhedron**.

The promoter for the polyhedrin gene is exceptionally strong, and can be used to drive expression of foreign proteins in insect cells.

Two forms of baculovirus are formed during the infection cycle

Baculovirus-Insect Cell Expression Systems

Baculoviruses transfer vector based on

Autographa californica multiple nuclear polyhedrosis virus (AcMNPV)



The expression unit of the vector contains:

1. A multiple cloning site (MCS) in which to clone the gene to be expressed.

2. The MCS lies between the polyhedrin gene promoter (Pp) and polyhedrin gene transcription-termination sequences (Pt).

3. Flanking regions of AcMNPV DNA (5' AcMNPV and 3' AcMNPV) to allow homologous recombination into the AcMNPV genome.

Baculovirus-Insect Cell Expression Systems Replacing the AcMNPV polyhedrin gene with an expression unit



Insect cells in culture are cotransfected with AcMNPV DNA and the transfer vector containing the cloned gene.

Within some of the double infected cells a double crossover event occurs and the expression unit is integrated into the AcMNPV genome.

Recombinant virus particles are harvested and heterologous proteins is harvested 4-5 days after infection of host insect cells with high-titre recombinant virus.

Baculovirus-Insect Cell Expression Systems Increasing the yield of recombinant baculovirus



Linearisation of the AcMNPV genome before transfection into insect cells increases recombination efficiency from <1% to ~30%

To do this a baculovirus was engineered with two *Bsu*36I restriction enzyme sites on either side of the polyhedrin gene.

Baculovirus-Insect Cell Expression Systems *E. coli*-insect cell baculovirus shuttle vectors (bacmids)





Bacmids make it possible to carry out all the genetic manipulations to generate a baculovirus expression vector in *E. coli*.

Transfection of insect cells is only required for the production of the heterologous protein

Note: the recombinant bacmid has a disrupted lacZ gene and cells transfected with this construct cannot produce functional β -galactosidase facilitating their identification.

Baculovirus-Insect Cell Expression Systems Some recombinant proteins produced using baculovirus systems

- α-Interferon
 Adenosine deaminase
 Anthrax antigen
 β-Amyloid precursor protein
 β-Interferon
 Bovine rhodopsin
 Bluetongue virus neutralization antigen
 Cystic fibrosis transmembrane
 conductance regulator
 Dengue virus type 1 antigen
 Erythropoietin
- G-protein-coupled receptors HIV-1 envelope protein HSV capsid proteins Human alkaline phosphatase Human DNA polymerase α Human pancreatic lipase Influenza virus hemagglutinin Interleukin-2 Lassa virus protein
- Malaria proteins Mouse monoclonal antibodies Multidrug transporter protein Poliovirus proteins Pseudorabies virus glycoprotein 50 Rabies virus glycoprotein Respiratory syncytial virus antigen Simian rotavirus capsid antigen Tissue plasminogen activator

Problems with using baculovirus systems to produce mammalian proteins:

1. Glycosylation – insect cells do not routinely add galactose or terminal sialic acid to N-linked glycoproteins. However, a stable insect cell line has been developed with integrated mammalian α -2,6-sialyltransferase and mammalian β -1,4-galactotransferase genes.

2. Proteolytic processing – mammalian cells contain a number of enzymes (not found in insect cells) to convert larger inactive proproteins into active proteins by proteolysis (proprotein convertases).

Mammalian Cell Expression Systems Advantages

Mammalian cell expression systems are important for producing mammalian proteins with a full complement of posttranslational modifications.

Can be achieved by:

Stable gene expression

Plasmid integration in the host genome or replication of the transferred genes as an artificial chromosome. Expression is long-term (stable).

Cell lines include:

• Chinese hamster ovary (CHO) cells

Transient gene expression

The foreign DNA is not integrated into the genome or replicated and is degraded or diluted on cell division. Expression lasts for a limited period of time (transient).

Cell lines include:

- African green monkey kidney (COS) cells
- Baby hamster kidney (BHK) cells
- Human embryonic kidney (HEK-293)

Mammalian Cell Expression Systems

A generalized mammalian expression vector



The vector contains:

- 1. A multiple cloning site (MCS) in which to clone the gene to be expressed.
- 2. The ampicillin resistance gene (Amp^r) a bacterial selectable marker.
- 3. The gene for a eukaryotic selectable market (SMG)
- 4. An *E*. *coli* origin of replication (ori^E) to allow replication of the vector in bacteria.
- 5. A eukaryotic origin of replication (ori^{euk}) to allow replication of the vector in mammalian cells.
- 6. Eukaryotic promoter (p), polyadenlyation (pa) and termination of transcription sequences (TT)

Mammalian Cell Expression Systems Translational control elements



The gene to be expressed can be fitted with a variety of sequences that enhance translation and facilitate secretion and purification, including:

- 1. A Kozak sequence (K) to allow efficient initiation of translation CC(A/G)CCATGG (where ATG is the start codon).
- 2. A signal sequence (S) to assure that the protein is secreted (if required).
- 3. A protein affinity tag (T) to facilitate purification of the protein.
- 4. A proteolytic cleavage site (P) to allow the tag (T) to be removed from the purified protein.
- 5. A stop codon (SC) to terminate translation.

6. 5' and 3' untranslated regions (5'UTR and 3'UTR) to increase the efficiency of translation and improve mRNA stability.

Mammalian Cell Expression Systems Expressing proteins made up of two different polypeptides



Some very important proteins (for example antibodies) contain two different polypeptides (the heavy and light chain in antibodies).

The genes for the two polypeptides (Gene α and Gene β) are cloned in expression vectors with different selectable marker genes (SMG1 and SMG2).

After cotransfection and selection with the two markers, the two polypeptides are co-expressed and the protein subunits can assemble to form the functional protein.

However, loss of one of the vectors from doubly transfected cells is common and the two vectors do not always have the same copy number, causing an imbalance in the expression of the two subunits.

Mammalian Cell Expression Systems Expressing proteins made up of two different polypeptides



To overcome the problem of two vector systems two-gene expression vectors have been developed.

The gene for each subunit each has its own promoter (p1 and p2) and polyadenylation (pa1 and pa2) and transcription termination (TT) regions.

Mammalian Cell Expression Systems Expressing proteins made up of two different polypeptides



Alternatively bicistronic expression vectors can be used.

There is a single promoter (p), polyadenylation (pa) and transcription termination region (TT) for both genes. Hence, there is only a single mRNA.

The mRNA contains an internal ribosome entry site (IRES) from a mammalian virus which allows intiation of translation of the downstream gene.

Mammalian Cell Expression Systems Selectable marker systems for mammalian expression vectors

Selective agent	Action of selective agent	Marker gene	gene protein
Xyl-A Blasticidin S Bleomycin G-418 (Geneticin) Histidincl	Damages DNA Inhibits protein synthesis Breaks DNA strands Inhibits protein synthesis Produces cytotoxic effects	Adenine deaminase (<i>ada</i>) Blasticidin S deaminases (<i>Bsr</i> , <i>BSD</i>) Bleomycin-binding protein (<i>Ble</i>) Neomycin phosphotransferase (<i>Neo</i>) Histidinol dehydrogenase (<i>hisD</i>)	Deaminates Xyl-A Deaminates blasticidin S Binds to bleomycin Phosphorylates G-418 Oxidizes histidinol to histidine
Hygromycin B	Inhibits protein synthesis	Hygromycin B phosphotrans- ferase (<i>Hvh</i>)	Phosphorylates hygro- mycin B
MSX	Inhibits glutamine synthesis	Glutamine synthetase (GS)	Cells that produce excess glutamine synthetase survive
MTX	Inhibits DNA synthesis	Dihydrofolate reductase (dhfr)	Cells that produce excess dihydrofolate reductase survive
PALA	Inhibits purine synthesis	Cytosine deaminase (codA)	Lowers cytosine levels in the medium by con- verting cytosine to uracil
Puromycin	Inhibits protein synthesis	Puromycin N-acetyltransferase (Pac)	Acetylates puromycin

Table 7.2 Selective marker gene systems for mammalian cells

MSX, methionine sulfoximine; MTX, methotrexate; PALA, N-(phosphoacetyl)-L-aspartate; Xyl-A, 9-β-D-xylofuranosyl adenine.

Mammalian Cell Expression Systems

Selectable markers that allow amplification of the copy number of the vector



The dihydrofolate reductase-methotrexate (DHFR-MTX) system allows selection for increased vector copy number and enhanced protein yield.

DHFR catalyses the reduction of dihydrofolate to tetrahydrofolate, which is required for the production of purines.

MTX is a competitive inhibitor of DHFR.

Sensitivity of the cells to MTX can be overcome by making more DHFR, which can be achieved by increasing the copy number of vectors carrying the DHFR gene.

The concentration of MTX is gradually increased over time and eventually cells with very high vector copy number and protein expression level are selected.