

# Biotechnology of drugs – Basics of genetic engineering I.

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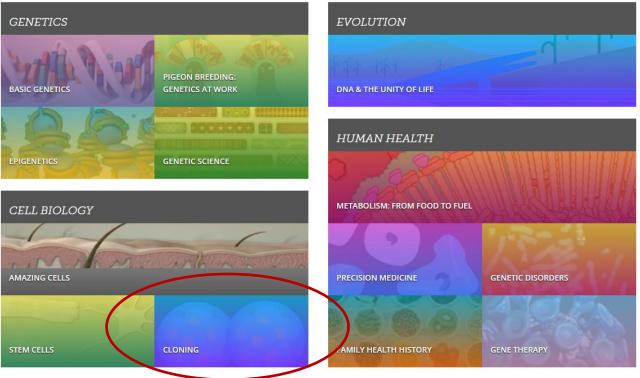
Department of Molecular Pharmacy FaF MU





Learn.Genetics

#### https://learn.genetics.utah.edu/



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#### **Biotechnological product can be created:**

#### 1. By classical biotechnological procedure

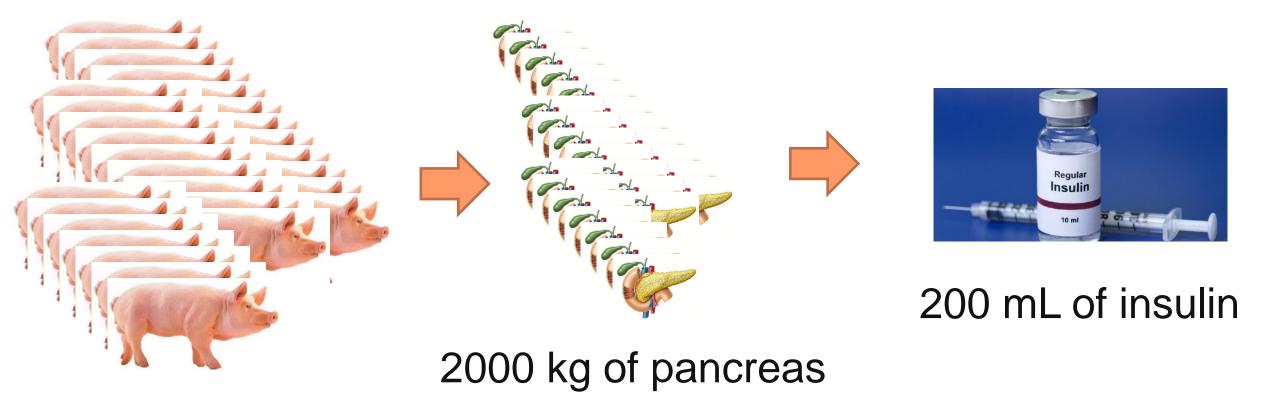
- interbreed
- mutagenesis of a population of cells
- selection of cells/organisms with suitable properties
- they cannot force the organism to produce a protein that is not its own

#### 2. By genetic engineering - recombinant DNA technology

- gene cloning → recombinant DNA technology
- genetic manipulation in vitro
- breaks down barriers between species → heterologous systems

## The goal of recombinant technologies

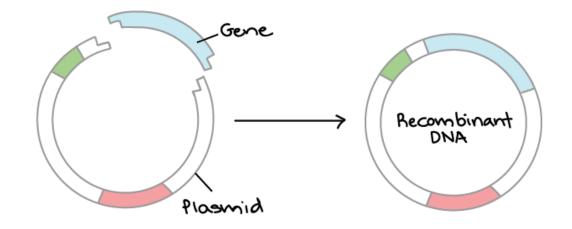
The effort to genetically encode the synthesis of a commercially advantageous product into an organism that will produce it cheaply and with high yield.



#### **Basic principles of recombinant technologies**

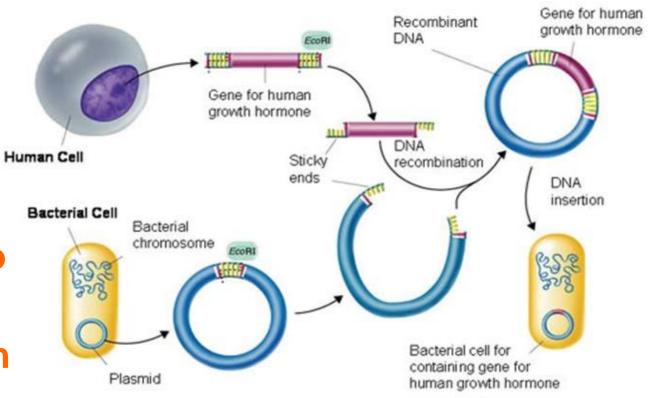
Recombinant technology - gene cloning - needs the incorporation of foreign DNA into the host cell

- introduction of DNA (transformation) into the cell
- ensuring the survival of forein DNA
- > ability to replicate
- > assurance of expression



# What is needed for gene cloning

- own gene DNA fragment
- vector plasmid, phage, cosmid
- host recipient of recombinant DNA
- > insert = gene incorporated into
  the vector
- recombinant DNA = vector with insert



https://microbenotes.com/gene-cloning-requirements-principle-steps-applications/

#### **Consequence of gene cloning = transformation**

A permanent heritable change in the genetic material of a cell caused by the acceptance and incorporation of foreign genetic information

> ability to replicate

> ability to express

#### **Sources of foreign genetic information**

animal

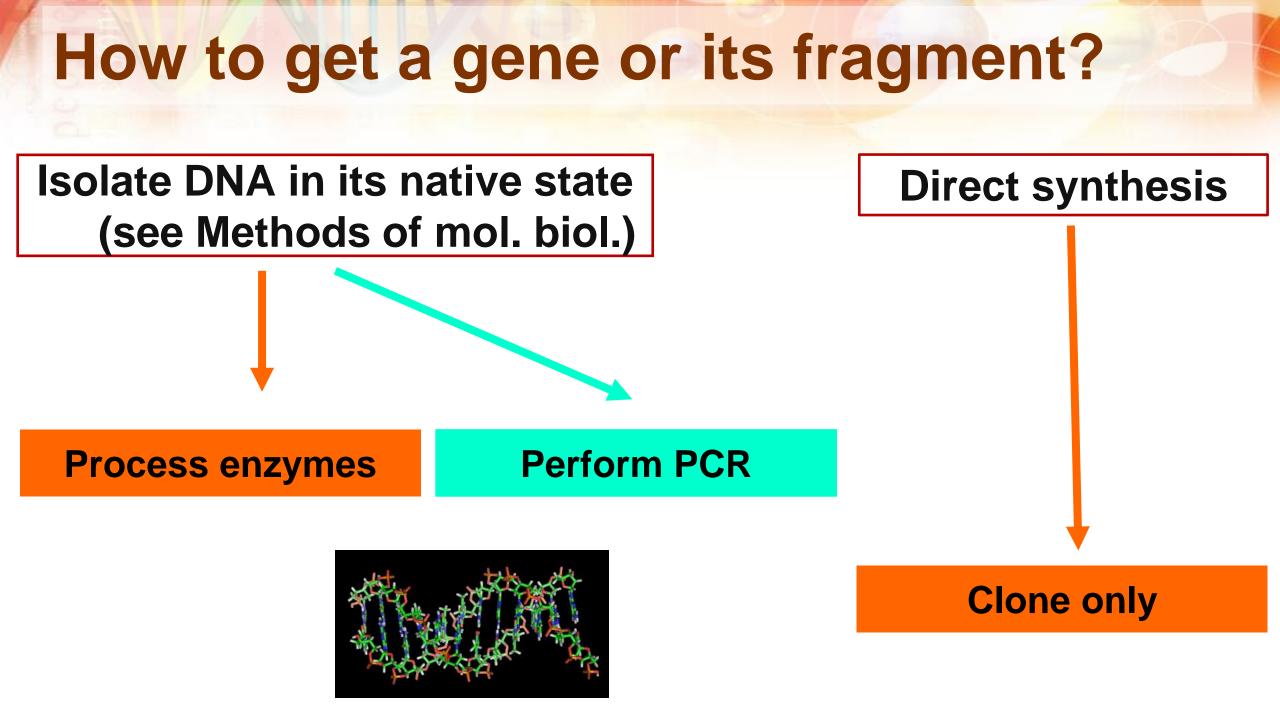
vegetal

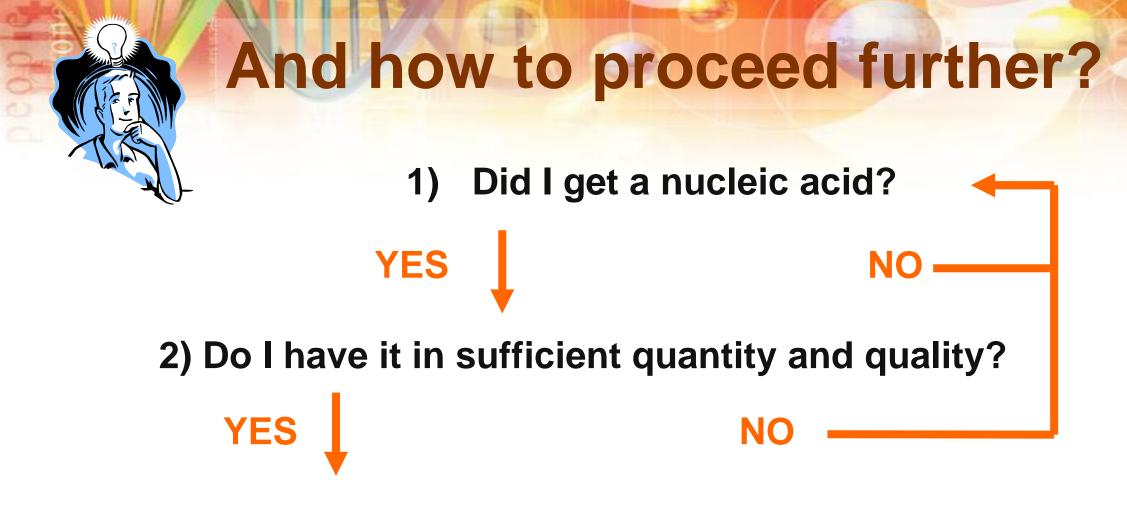
microbial

synthetic

The recombinant protein can be:

- Final product (e.g. hormone, antibody)
- Tool for further synthesis (e.g. various enzymes)





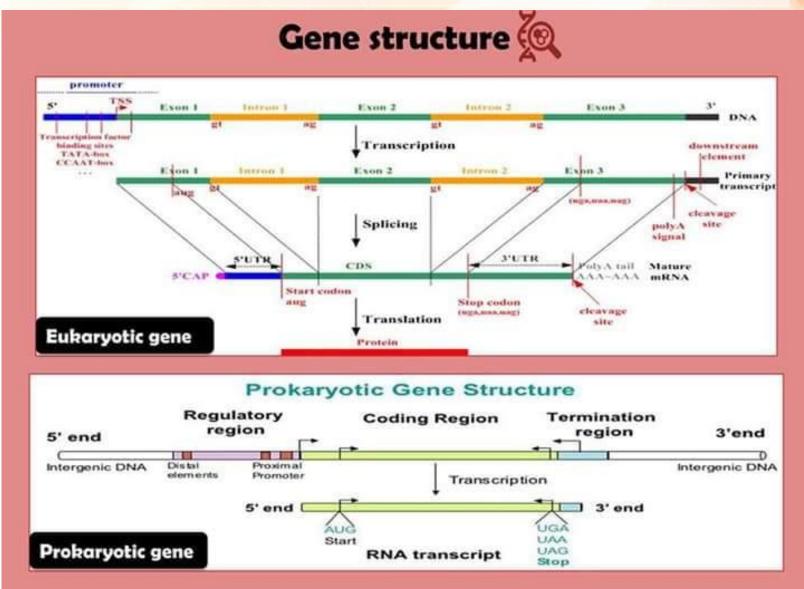
a) Process enzymes (see Methods of mol. biol.)b) Perform PCR (see Methods of mol. biol.)

#### Factors influencing the expression of cloned genes

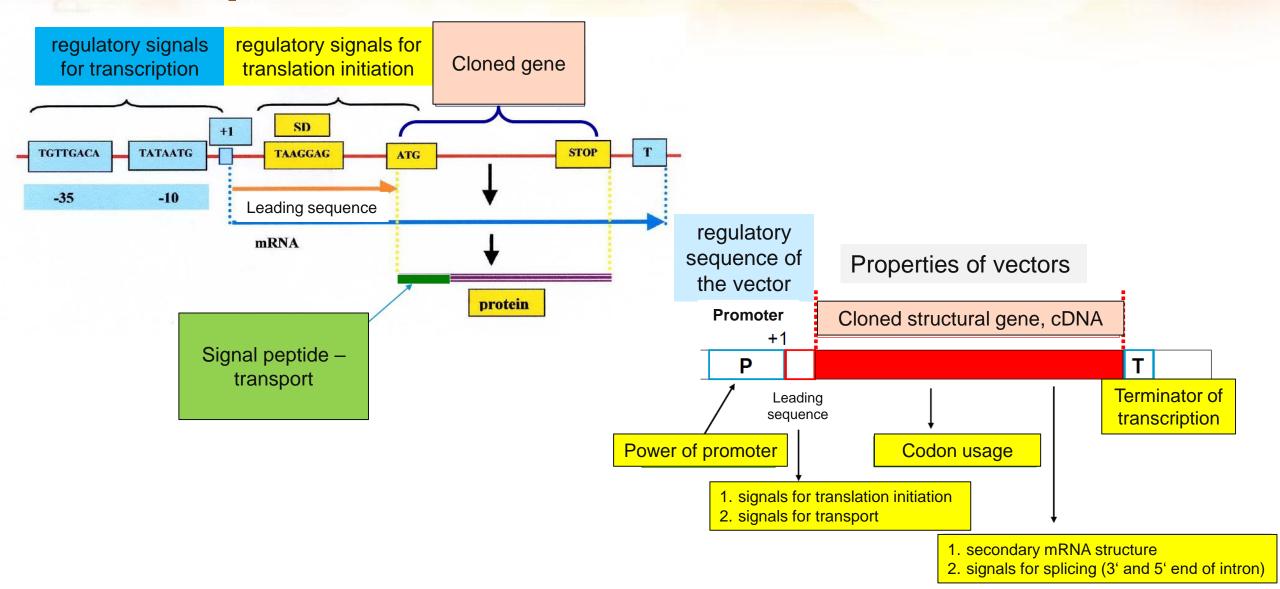
#### **Regulatory sequences for gene Properties of vectors** expression 1. Transcription 1. The number of copies of the vector in the cell The power of the promoter a. Transcription terminator b. 2. Vector stability Stability of mRNA C. Physiology of the host cell 2. Translation a. Structure of the ribosome binding site 1. Growing conditions b. Codon usage 3. Transport of proteins 2. The enzyme apparatus of the host cell a. Character of the signal sequence

### **Insert properties**

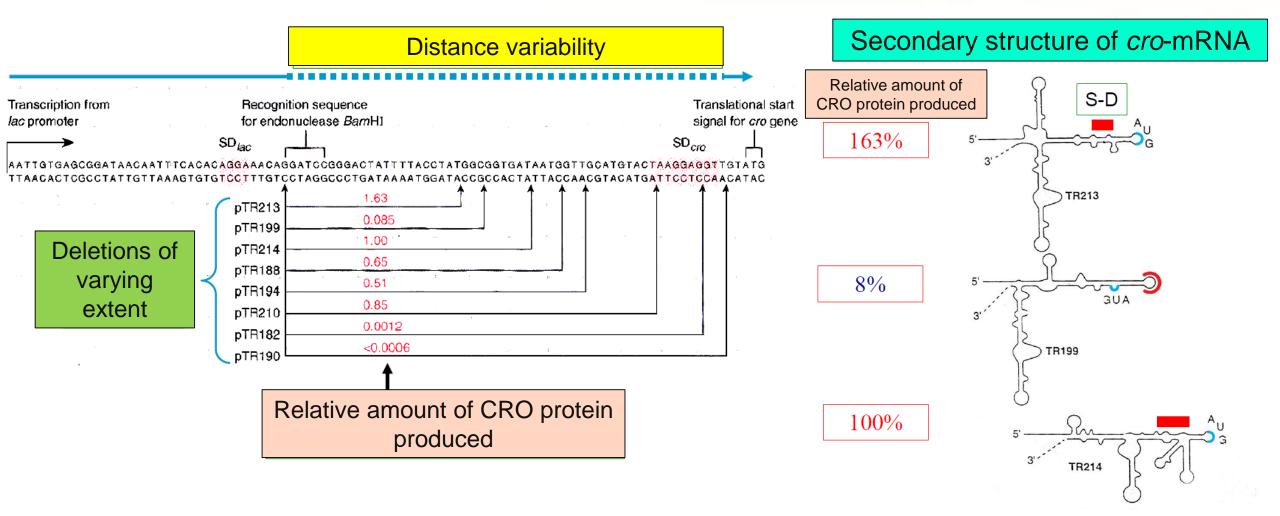
- It contains a coding sequence
- Contains the start of translation (ATG start codon)
- Contains the end of translation (stop codon)
- It carries the appropriate codons for specific amino acids



# Signals influencing prokaryotic structural gene transcription and translation

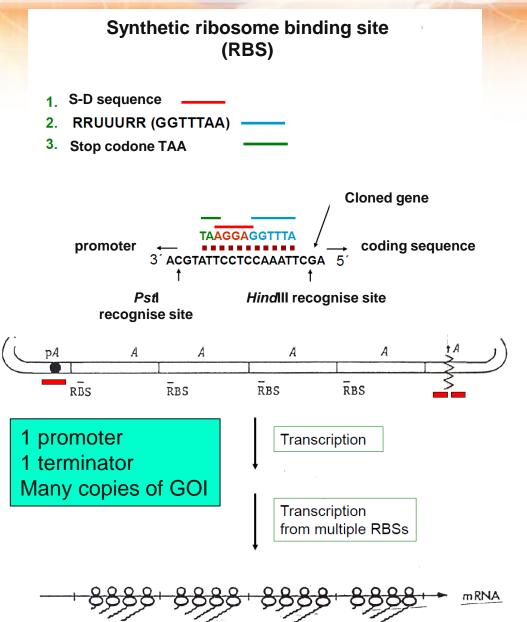


#### The effect of the distance between the promoter and the start of translation on the secondary structure of mRNA and on the amount of expressed protein



#### **Possibilities of ensuring high proteosynthesis**

- The rate of proteosynthesis depends on the amount of mRNA in the cell
- Modifications in the 3'-UTR or 5'-UTR can increase the mRNA half-life
- The **PuPuUUUPuPu** sequence near the Shine-Dalgarn sequence is essential for translation of eukaryotic genes in *E. coli*
- Construction of homopolycistronic expression cassettes
- Proteins in the periplasm have a longer half-life than in the cytoplasm
- Targeted inhibition of specific proteases
- Formation of fusion proteins protein from the host organism (stabilizing partner) + protein of interest



# Effect of codon usage

CODON USAGE IN E. COLI GENES<sup>1</sup>

- Different species use codons with different frequencies  $\rightarrow$  related to the amount of tRNA
- Solution:
  - Cloning rare tRNAs together with a gene of interest
  - Substitution of rare codons for common ones by site-directed mutagenesis

	Codon	Amino	9 <b>%</b> 3	Ratio <sup>4</sup>	Codon	Amino	%	Ratio	Codon	Amino	%	Ratio	Codon	Amino	198	Ratio	
		ac id <sup>2</sup>				ac id				ac id				ac id			
U	UUU	Phe (F)	1.9	0.51	UCU	Ser (8)	1.1	0.19	UAU	Туз (Ү)	1.6	0.53	UGU	Cys (C)	0.4	0.43	U
	UUC	Phe (F)	1.8	0.49	UCC	Ser (8)	1.0	0.17	UAC	Туз (Ү)	1.4	0.47	UGC	Cys(C)	0.6	0.57	С
	UUA	Leu (L)	1.0	0.11	UCA	Ser (8)	0.7	0.12	UAA	STOP	0.2	0.62	UGA	STOP	0.1	0.30	Α
	UUG	Leu (L)	1.1	0.11	UCG	Ser (8)	0.8	0.13	UAG	STOP	0.03	0.09	UGG	T1p (V)	1.4	1.00	G
С	CUU	Leu (L)	1.0	0.10	CCU	Pro (P)	0.7	0.16	CAU	His (H)	1.2	0.52	CGU	A1g (R)	2.4	0.42	U
	CUC	Leu (L)	0.9	0.10	CCC	P10(P)	0.4	0.10	CAC	His (H)	1.1	0.48	CGC	Aig (R)	2.2	0.37	С
	CUA	Leu (L)	0.3	0.03	CCA	Pro (P)	0.8	0.20	CAA	Gln (Q)	1.3	0.31	CGA	Aig (R)	0.3	0.05	Α
	CUG	Leu (L)	5.2	0.55	CCG	P10(P)	2.4	0.55	CAG	Gln (Q)	2.9	0.69	CGG	A1g (R)	0.5	0.08	G
Α	AUU	Ile (I)	2.7	0.47	ACU	Thu (T)	1.2	0.21	AAU	Asn (N)	1.6	0.39	AGU	Ser (8)	0.7	0.13	U
	AUC	Ile (I)	2.7	0.46	ACC	Thu (T)	2.4	0.43	AAC	Asn (N)	2.6	0.61	AGC	Ser (8)	1.5	0.27	С
	AUA	Ile (I)	0.4	0.07	ACA	Thu (T)	0.1	0.30	AAA	Lys (K)	3.8	0.76	AGA	A1g (R)	0.2	0.04	Α
	AUG	Met (M)	2.6	1.00	ACG	Thu (T)	1.3	0.23	AAG	Lys (K)	1.2	0.24	AGG	A1g (R)	0.2	0.03	G
G	GUU	Val(∀)	2.0	0.29	GCU	Ala (A)	1.8	0.19	GAU	Asp (D)	3.3	0.59	GGU	Gly (G)	2.8	0.38	U
	GUC	Val (V)	1.4	0.20	GCC	Ala (A)	2.3	0.25	GAC	Asp (D)	2.3	0.41	GGC	Gly (G)	3.0	0.40	С
	GUA	Val (V)	1.2	0.17	GCA	Ala (A)	2.1	0.22	GAA	Glu(E)	4.4	0.70	GGA	Gly (G)	0.7	0.09	Α
	GUG	Val (V)	2.4	0.34	GCG	Ala (A)	3.2	0.34	GAG	Glu(E)	1.9	0.30	GGG	Gly (G)	0.9	0.13	G
	Ū				C			Α			G						

<sup>1</sup> The data shown in this table is from the Arabidopsis Research Companion on the World Wide Web (//weeds/mgh.harvard.edu). Codon

frequencies for many other bacteria can be found at http://morgan.angis.su.oz.au/Angis/Tables.html.

<sup>2</sup> The letter in parenthesis represents the one-letter code for the amino acid.

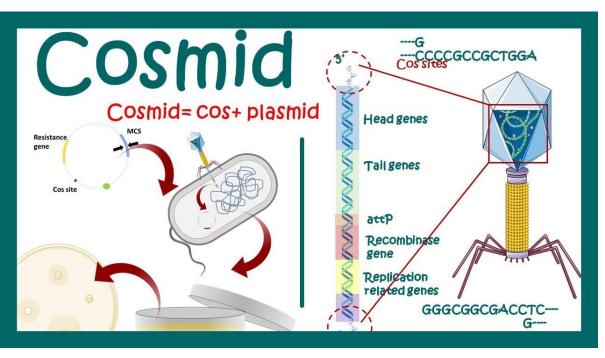
<sup>3</sup> % represents the average frequency this codon is used per 100 codons.

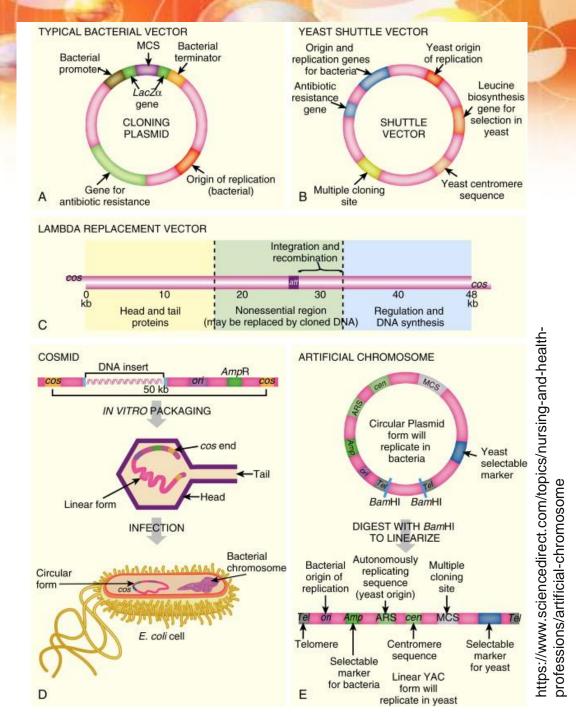
<sup>4</sup> Ratio represents the abundance of that codon relative to all of the codons for that particular amino acid.

http://2014.igem.org/Team:Penn State/CodonOptimization

## Vectors

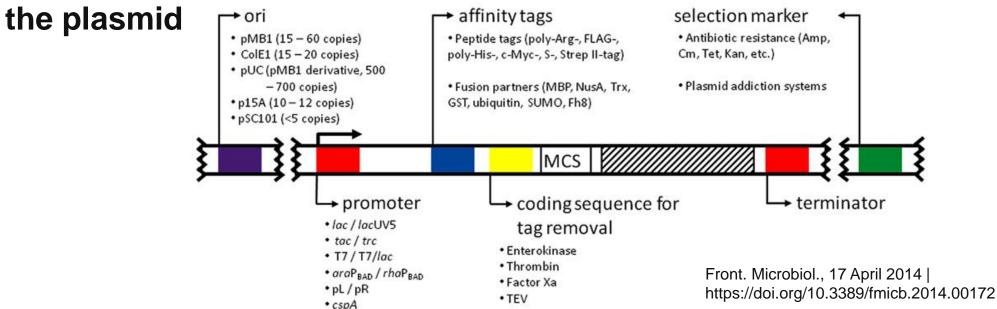
- Plasmids
- Bacteriophages
- Cosmids
- Artificial chromosomes
  - BAC (bacterial artificial chromosomes)
  - YAC (yeast artificial chromosomes)



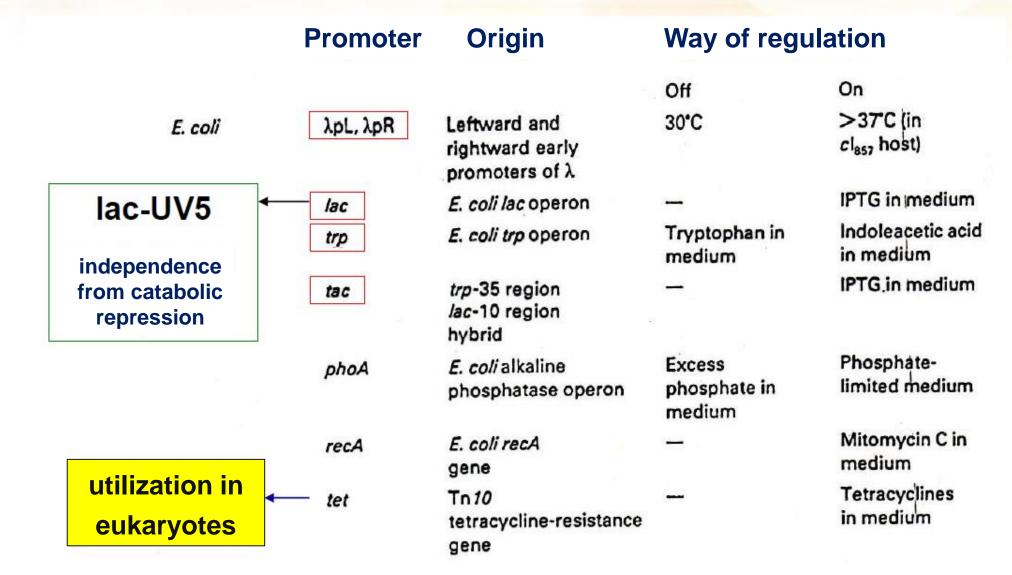


#### Anatomy of an expression vector

- origin of replication (ori), which is a condition for the production of new copies
- 2) inducible promoter, which will allow to regulate the expression of the desired protein
- 3) selection marker ensuring the preferential growth of transformed bacteria
- 4) multi-cloning site (MSC), allowing a foreign DNA fragment to be inserted into

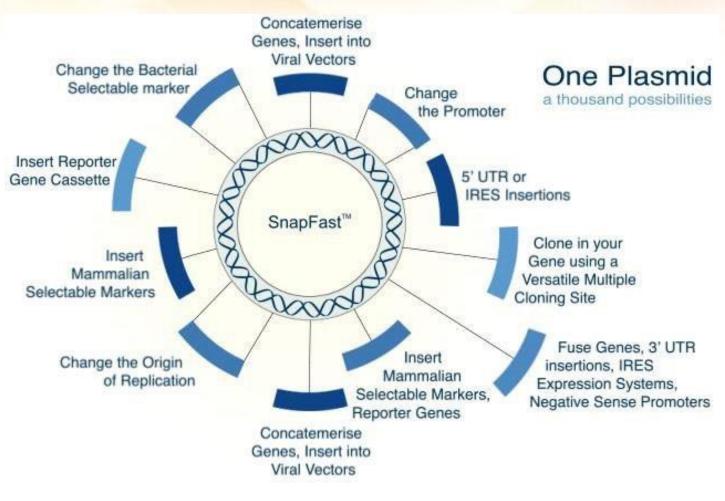


# Examples of regulatable promoters in expression vectors



## **Plasmids as vectors**

- extrachromosomal circular dsDNA
- occurrence in many bacterial species
- > 1,000 to 200,000 bp in size
- carry only genes encoding secondary traits (resistance to antibiotics)
- autonomous replication
- > insert size = up to 25 kbp

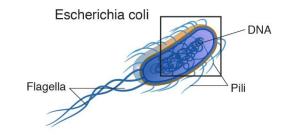


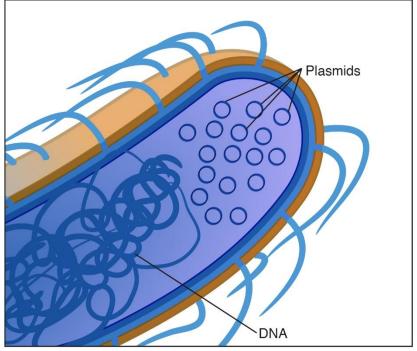
# Recent trend → plasmid synthes on demand

(§) VectorBuilder	Login or Register		My Account 🐱		Search	
Design Vector	Products & Services	GMP Manufacturing	Tools	Resources	Support	
Home > Choose Vector System > Mammalian Gene Exp	ression Vector					
	Vector Desig	n Studio How to	use this page?			
Mammalian Gene Expression Vector	Guide 🗹				Design Another Vector	
Select number of ORFs O	Click on steps on the vector matching	Step 1	Add Promoter ♥ Kozak Step 2   Add SV40 late pA 3   Add Marker ♥	d ORF ♥	Finish Design ►	

# **Plasmid suitability criteria**

- 1) small size = ability to transform
- 2) plasmid stability
- 3) high number of copies in the cell = yield
- 4) easy handling
- 5) "shuttle" vectors = work in multiple host species (e.g. *E. coli* + mammalian cells)

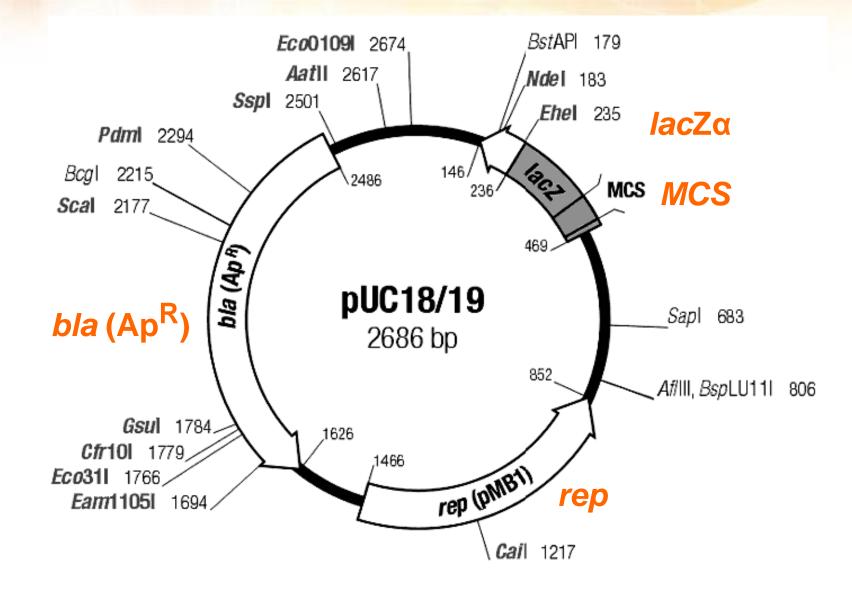




Escherichia coli

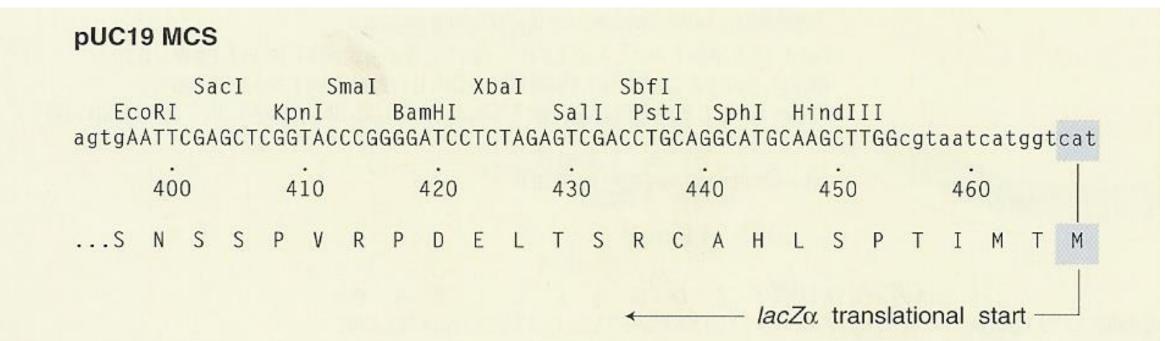
https://www.genome.gov/genetics-glossary/Plasmid

#### Plasmids pUC18 and pUC19

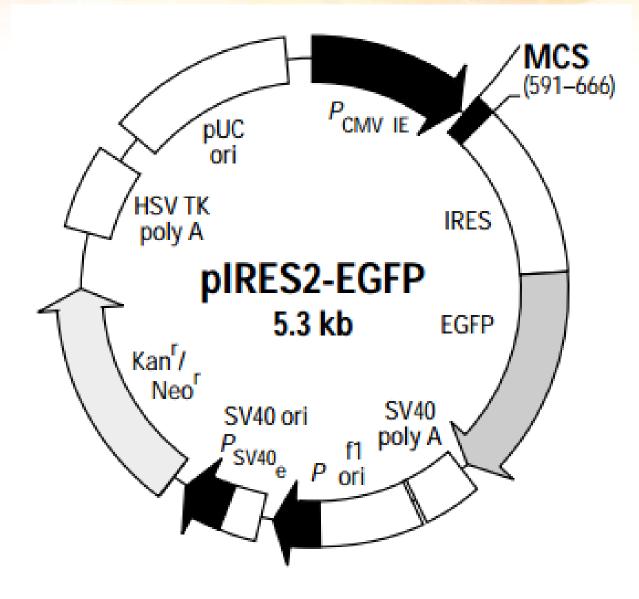


#### Polylinker pUC18 and pUC19

#### MCS = multi-cloning site

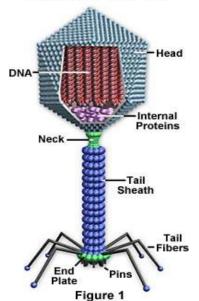


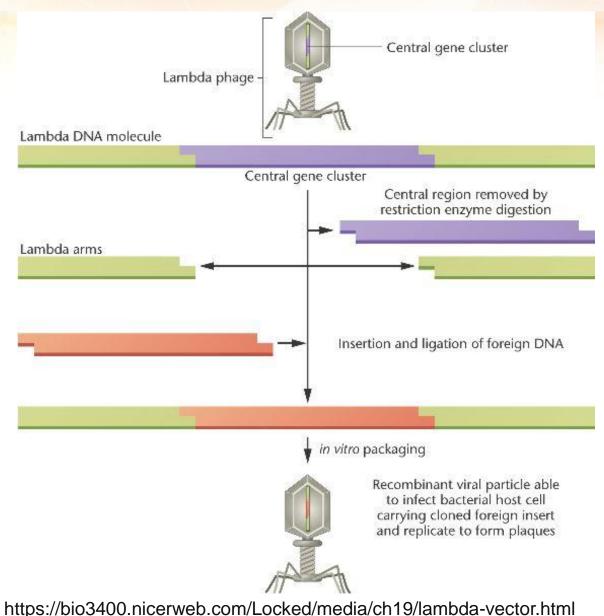
#### **Expression plasmid pIRES2-eGFP**



#### **Bacteriophage λ-based vectors**

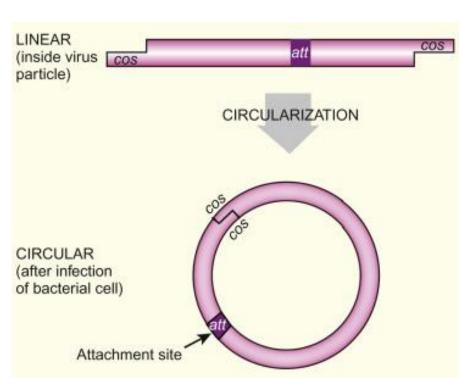
- hey replace plasmids when longer DNA fragments need to be cloned
- Insertion vectors 8 10 kbp
- Replacement vectors 8 24 kbp Bacteriophage Structure

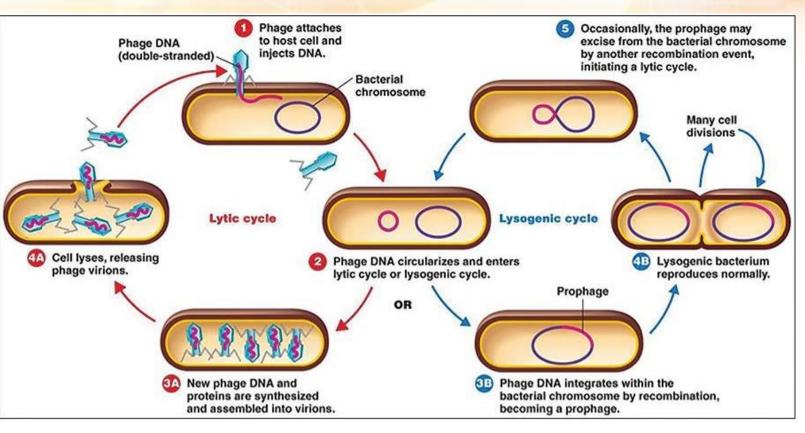




# **Bakteriophage** λ

- > 50 kbp dsDNA
- linear and circular form
- cos sites

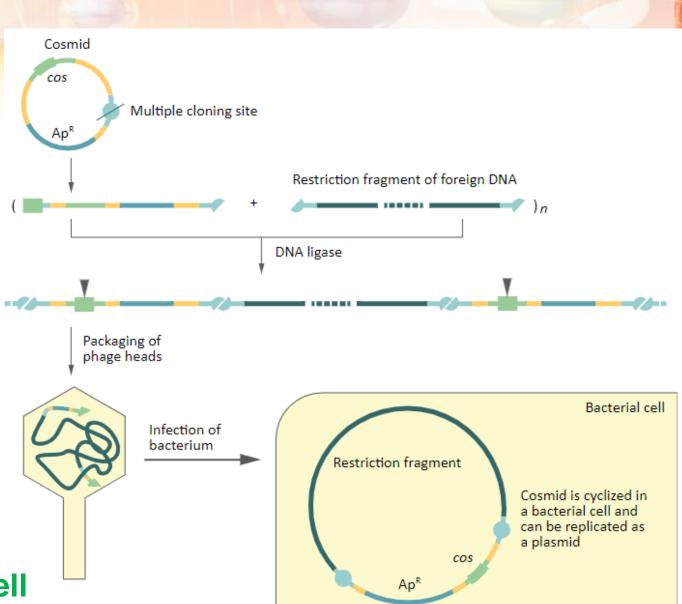




http://hotcore.info/babki/bacteriophage-lysogenic-cycle.html

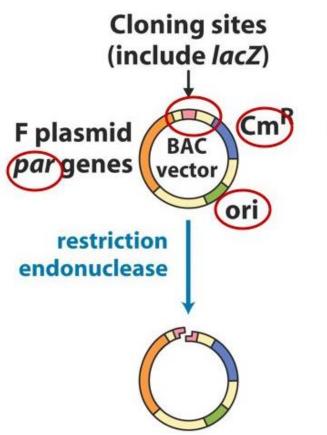
# Cosmids

- combination of plasmid and phage
- prokaryotic origin of replication oriV
- selection marker
- cloning site
- capacity 37 52 kbp
- cos sites of bacteriophage λ
- for packaging, packaging proteins must be added
- enters as a phage
- it behaves like a plasmid in the cell



# **Bacterial Artificial Chromosome**

- BAC = bacterial artificial chromosome
- Derived from plasmid F'
- Designed for cloning into bacterial cells
- They occur in the number of 1-2 copies per cell
- Cloned DNA is highly stable
- Cloning capacity up to 300 kbp (maybe more)
  - Used in the HUGO project
  - Today, they are replaced by the methods of whole-genome sequencing, next generation sequencing and third generation sequencing



Artificial Chromosomes allow for cloning of large pieces of DNA

Bacterial Artificial Chromosome

1. *ori* allows for replication in bacteria,

2. *par* helps segregate BAC evenly between daughter cells,

3. *lac Z* allows for detection of insert,

4. Cm<sup>R</sup> allows for selection of transformed cells.

https://library.uams.edu/assets/COM/BioChem/MolecularTools/ MolecularToolsSDL12.html

## **Other variants of vectors**

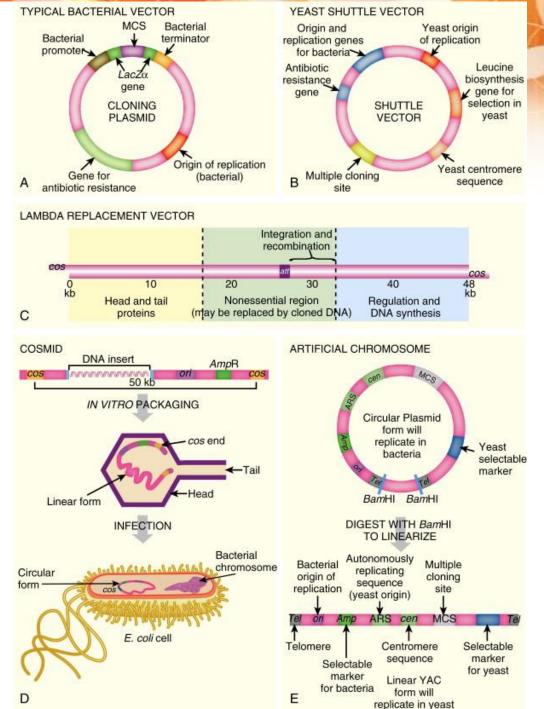
#### **Phosmid**

- Similar to a cosmid, but based on a bacterial F-plasmid
- > The host (*E. coli*) may contain only a single molecule
- Cloning capacity up to 40 kbp
- Suitable for constructing stable libraries from complex genomes
- > Highly stable; capable of maintaining human DNA for over 100 generations

#### Phagemid

- Plasmid containing origin of replication of phage M13
- > It is used to prepare ssDNA.
- > The best known examples are the pBluescript series of cloning vectors.

(A) Typical bacterial cloning vector. This vector has bacterial sequences to initiate replication and transcription. In addition, it has a multiple cloning site embedded within the lacZ  $\alpha$  gene so that the insert can be identified by alpha-complementation. The antibiotic resistance gene allows the researcher to identify any E. coli cells that have the plasmid. (B) Yeast shuttle vector. This vector can survive in either bacteria or yeast because it has both yeast and bacterial origin of replication, a yeast centromere, and selectable markers for yeast and bacteria. As with most cloning vectors, there is a polylinker. (C) Lambda replacement vectors. Because lambda phage is easy to grow and manipulate, its genome has been modified to accept foreign DNA inserts. The region of the genome shown in green is nonessential for lambda growth and packaging. This region can be replaced with large inserts of foreign DNA (up to about 23 kb). (D) Cosmids. Cosmids are small multicopy plasmids that carry cos sites. They are linearized and cut so that each half has a cos site (not shown). Next, foreign DNA is inserted to relink the two halves of the cosmid DNA. This construct is packaged into lambda virus heads and used to infect E. coli. (E) Artificial chromosomes. Yeast artificial chromosomes have two forms: a circular form for growing in bacteria and a linear form for growing in yeast. The circular form is maintained like any other plasmid in bacteria, but the linear form must have telomere sequences to be maintained in yeast. The linear form can hold up to 2000 kb of cloned DNA and is very useful for genomics research.



ίΩ -0-642 https://doi.org/10.1016/C2009

#### Hosts = recipients of recombinant DNA

**Regardless of source type** 

- bacterial cells
- > yeasts and molds
- plant and animal cells
- whole plant or animal

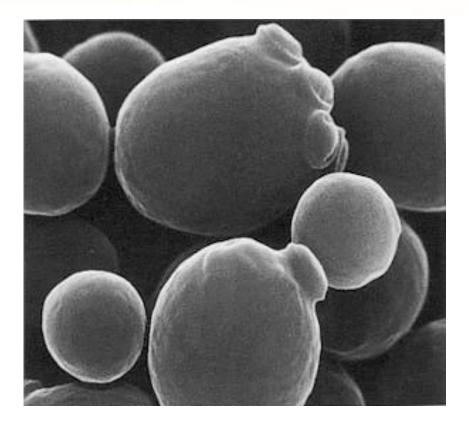
## Escherichia coli

- ➢ G- bacteria, circular chromosome 3×10<sup>6</sup> bp
- > amount of usable plasmids
- > generation time 20 min.  $\rightarrow$  rapid biomass formation
- > undemanding and cheap cultivation
- ➤ stationary phase 2×10<sup>9</sup> cells/mL
- a number of mutants (DH5α, HB101, BL21,...)
- disadvantages significantly different posttranslational modifications compared to eukaryotes



#### Saccharomyces cerevisiae

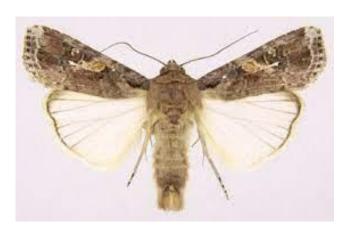
- linear chromosomes
- ➢ approximately 13×10<sup>6</sup> bp
- ➤ about 6,275 genes
- > the simplest eukaryotic organism
- identical transcription and translation apparatus with other eukaryotes
- b differences in post-translational processes, e.g. mannose hyperglycosylation
  - Schizosaccharomyces pombe
     Pichia pastoris



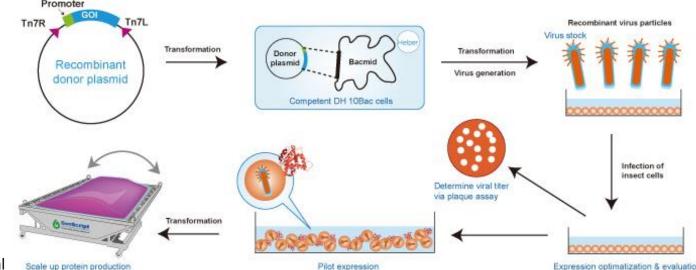
### Insect cells and baculoviruses

- similar selection principles as in lower eukaryotes or bacteria
- > use of binary "shuttle" vectors (bacteria + insects)
- replacement of the polyhedrin sequence of the virus with a recombinant gene
- the most commonly used Sf9 cell culture derived from Spodoptera frugiperda

#### baculoviruses only attack insect cells



https://www.genscript.com/insect-customized-expression-package.html scale up protection



#### **Plant cells**

vectors are typically bacterial plasmids that contain plant expression cassettes

> direct transformation
 > transformation using Agrobacterium
 > transformation by viral vector

host cells – Nicotiana tabacum, Arabidopsis thaliana, ...

**Relatively safe technology** 



#### Mammalian cells and their viruses

- systems closest to man
- It the most common producer is mammalian CHO cells (Chinesse hamster ovaries)
- > differential interspecies glycosylation
- > as vectors serve adenoviruses, retroviruses, herpesviruses



#### **Technology security issues !**

#### **Creation of recombinant proteins in different organisms**

Parameter	Bacteria	Yeast	Mammalian cell culture	Transgenic plants
Glycosylation	None	Incorrect	Correct	Generally correct
Assembles multimeric proteins	Limited	Limited	Limited	Yes
Production costs	Medium	Medium	High	Low
Protein-folding accuracy	Low	Medium	High	High
Protein yield	High	High	Medium	Medium
Scale-up costs	High	High	High	Low
Time required	Low	Low	High	Medium
Skill level required for growth	Medium	Medium	High	Low