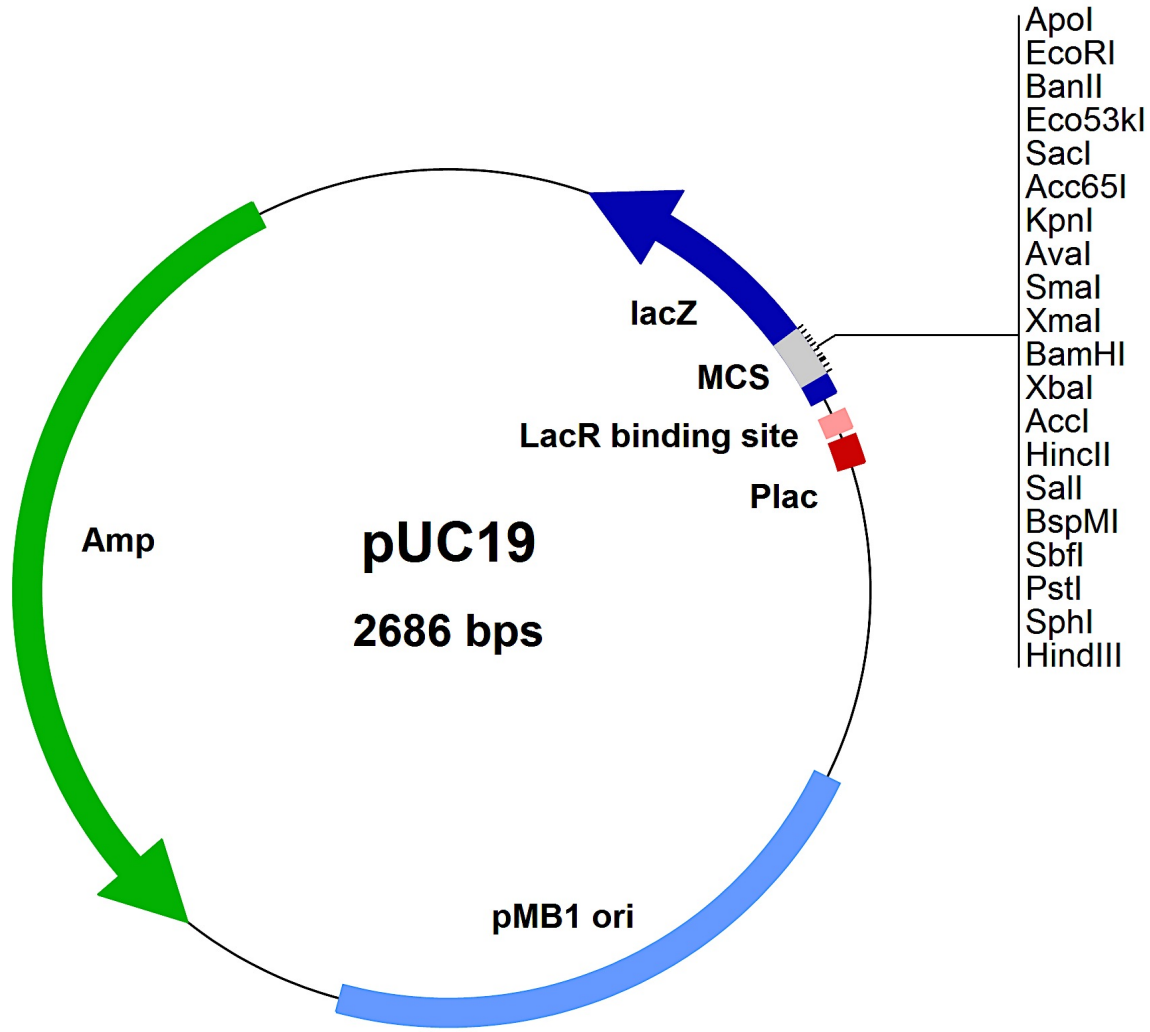
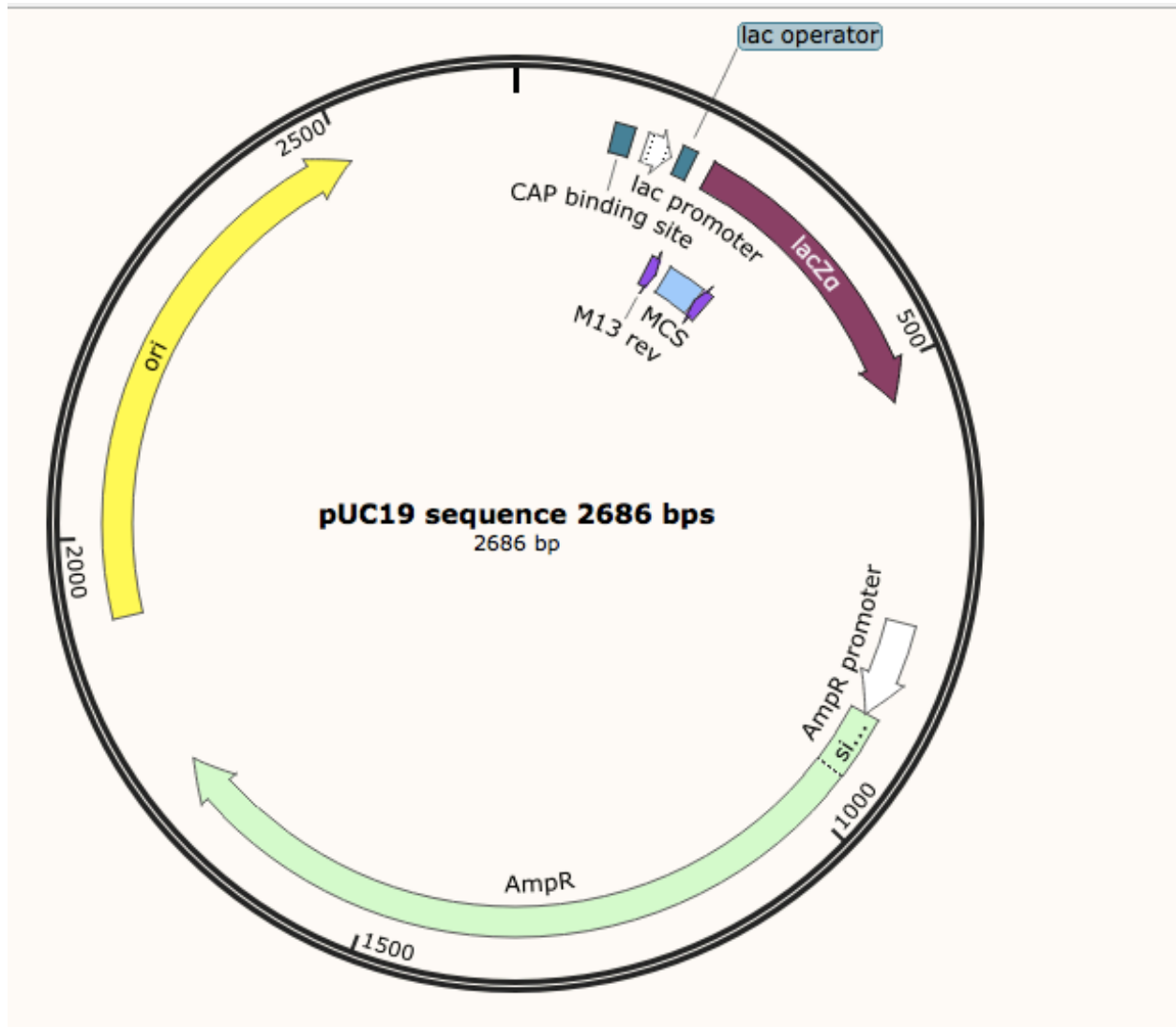
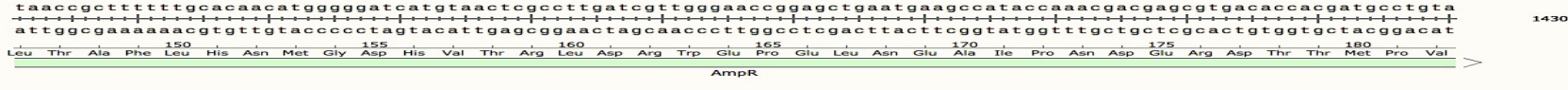
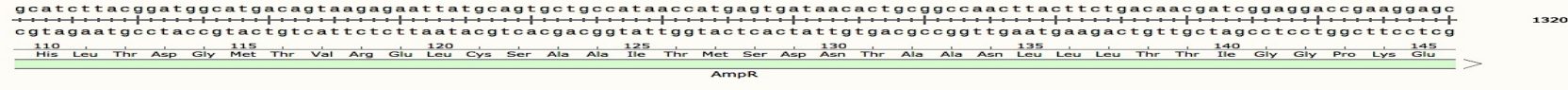
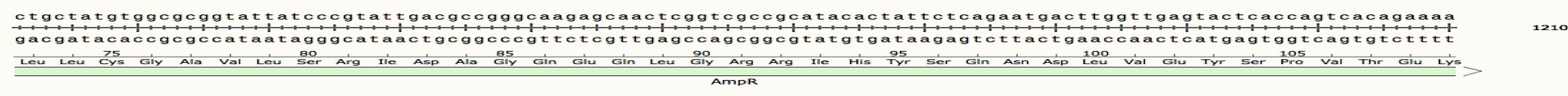
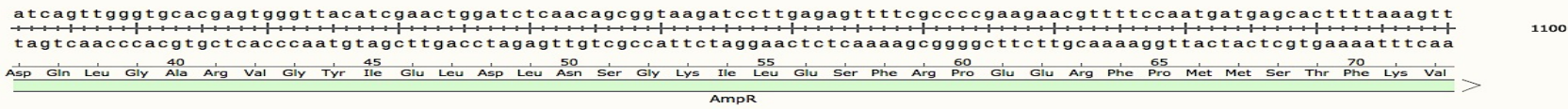
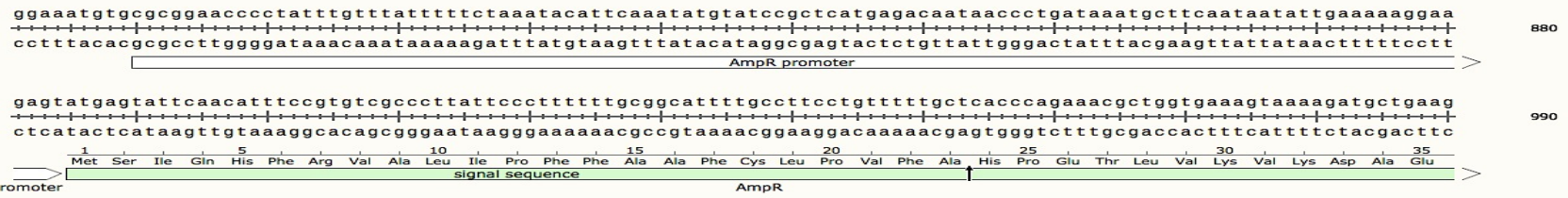
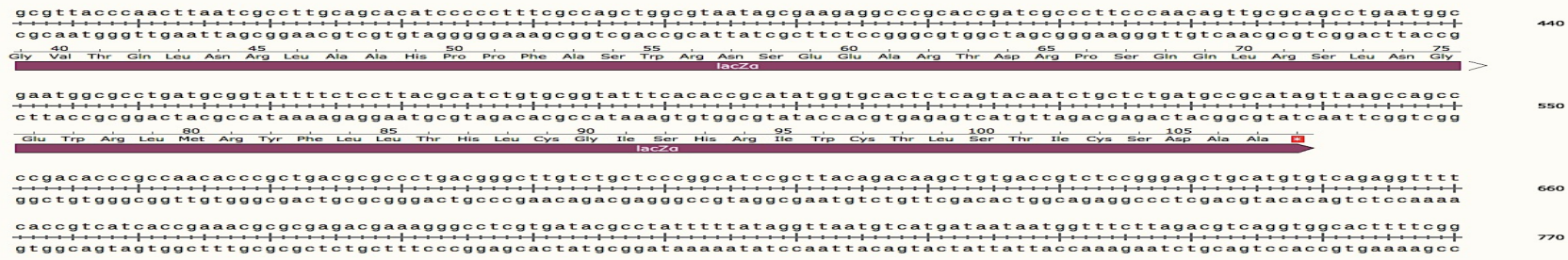


# Cloning vectors







gcaatggcaacaacggttgcgcaaacatttaactggcgaactacttactctagcttcccggcaacaattaatagactggatggaggcggataaagttgcaggaccacttct  
cgttaccgttgttgcaacgcggttgataattgaccgcttgatgaatgagatcgaaggccggttgttaattatctgacctaccctccgctatttcaacgctcctggggaaga  
Ala Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr Leu Ala Ser Arg Gln Gln Leu Ile Asp Trp Met Glu Ala Asp Lys Val Ala Gly Pro Leu Leu  
185 190 195 200 205 210 215  
AmpR

gcgctcggcccttccggctggctggtttattgctgataaatctggagccgggtgagcgtgggtctcgcgggtatcattgcagcactggggccagatggtaagccctcccgta  
cgcgagccgggaaggccgaccgaccaataaacgactatttagacctcggccactcgcacccagagcgcctatagtaacgctcgtgaccccggtctaccattcgggagggcat  
Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile Ala Asp Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala Ala Leu Gly Pro Asp Gly Lys Pro Ser Arg  
220 225 230 235 240 245 250 255  
AmpR

tcgtagttatctacacgacggggagtcaggcaactatggatgaacgaaatagacagatcgcctgagataggtgcctcactgatgaagcattggtaactgtcagaccaagtt  
agcatcaatagatgtgctgcccctcagtcggtgatacctacttgccttattctgtctagcgactctatccacggagtgactaattcgtaacattgacagctctggttcaa  
Ile Val Val Ile Tyr Thr Thr Gly Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu Ile Gly Ala Ser Leu Ile Lys His Trp  
260 265 270 275 280 285  
AmpR

tactcatatatacttttagattgattttaaacttcatttttaaatttaaaggatctaggtgaagatccttttggataatctcatgacaaaaacccttaacgtaggttttc  
atgagtataatgaaatctaactaaatttgaagtaaaaaataaatttctctagatccacttctaggaaaaactattagagtagctgggttttagggaattgcactcaaaag  
gttccactgagcgtcagaccccgtagaaaagatcaaaggatcttcttgagatccttttttctgcgcgtaaatctgctgcttgcacaaaaaaaaccaccgctaccagcgg  
caagggtgactcgcagctcggggcatcttttctagtttcttagaagaactctaggaaaaaaagacgcgcatttagacgacgaacgcttgttttttgggtggcgtggtgcgc  
ori

tggtttggttgcgggatcaagagctaccaactcttttccgaaggtaactggcttcagcagagcgcagataccaaaactgcttcttctagtgtagccgtagttaggccac  
accaaaaaacaggcctagttctcgatgggtgagaaaaaggctccattgaccgaaagctgctcgcgctctatgggttatgacaagaagatcacatcggcatcaatccgggtg  
ori

cactcaagaactctgtagaccgcctacatacctcgtctctgctaatcctgttaccagtggtgctgctgccagtgccgataagtcggtcttaccgggttgactcaagacg  
gtgaagttcttgagacatcgtggcggatgtatggagcggagacgattaggacaatggtcaccgacgaggtcaccgctattcagcacagaatggcccaacctgagttctgc  
ori

atagttaccggataaaggcgcagcggctcgggctgaacggggggttctgtgcacacagcccagcttggagcgaacgacctacaccgaactgagatacctacagcgtgagctat  
tatcaatggcctattccgcgctcgcagcccagcttgccccccaagcacgtgtgtcgggtcgaacctcgcttgcctggatgtggcttgactctatggatgtcgcactcgata  
ori

atagttaccggataaaggcgcagcggctcgggctgaacggggggttctgtgcacacagcccagcttggagcgaacgacctacaccgaactgagatacctacagcgtgagctat  
tatcaatggcctattccgcgctcgcagcccagcttgccccccaagcacgtgtgtcgggtcgaacctcgcttgcctggatgtggcttgactctatggatgtcgcactcgata  
ori

gagaaagcgcacgcttcccgaaggagaaaggcggacaggtatccggtaagcggcagggtcggaaacaggagagcgcacgaggggagcttccaggggaaacgcctgggtat  
ctcttctcgggtgcaagggttccctcttccgcctgtccataggccattcgcctgccagccttgcctctcgcggtgctccctcgaagggtccccttgcggaccata  
ori

ctttatagtcctgtcgggtttcgcaccctctgacttgagcgtcgatttttgtgatgctcgtcaggggggaggcctatggaaaaacgccagcaacgcggccttttttacg  
gaaatcaggacagcccaagcgggtggagactgaaactcgcagctaaaaaacactacgagcagctccccgcctcggatacctttttcgggtcgttgcgcgggaaaaatgc  
ori

gttccctggccttttgcctggccttttgcctcacatgttcttccctcgttattccctgattctgtggataaacgctattaccgctttagtgagctgataaccgctcgcgcga  
caaggaccgaaaaacgaccggaaaaacgaggtgtacaagaaaggacgcaataggggactaagacacctattggcataatggcggaaactcactcgactatggcggagcggcgt

gccgaacgaccgagcgcagcaggtcagtgagcggaggaagcgggaaga 3'  
cggcttgcctggcctcgcgctcagtcactcgcctccttcgccttct 5' 2686

pUC19 sequence 2686 bps.dna (Circular / 2686 bp)

New Open Save Print Undo Redo Copy Paste

Try SnapGene if you wish to align DNA sequences with a reference sequence.

2686 bp

Chosen Enzymes: Unique Cutters (50) from "Nonredundant Commercial"

Enzyme	Sites	Numbers	Lines
AatII	1	753	
AccI	1	252	
Acc65I	1	272	
AflIII	1	2560	
AhdI	1	1672	
AlwNI	1	2151	
ApoI	1	284	
AvaI	1	268	
BamHI	1	263	
BanII	1	282	
BfuAI	1	238	
BmeT110I	1	269	
BpmI	1	1603	
BsaI	1	1606	
BseYI	1	2256	
BsoBI	1	268	
BspMI	1	238	
BspQI	1	2677	
BsrFI	1	1587	

\* Methylation-sensitive sites

Noncutters

Remove Add

Map Sequence Enzymes Features Primers History

Description Panel





# - The useful properties of pBR322

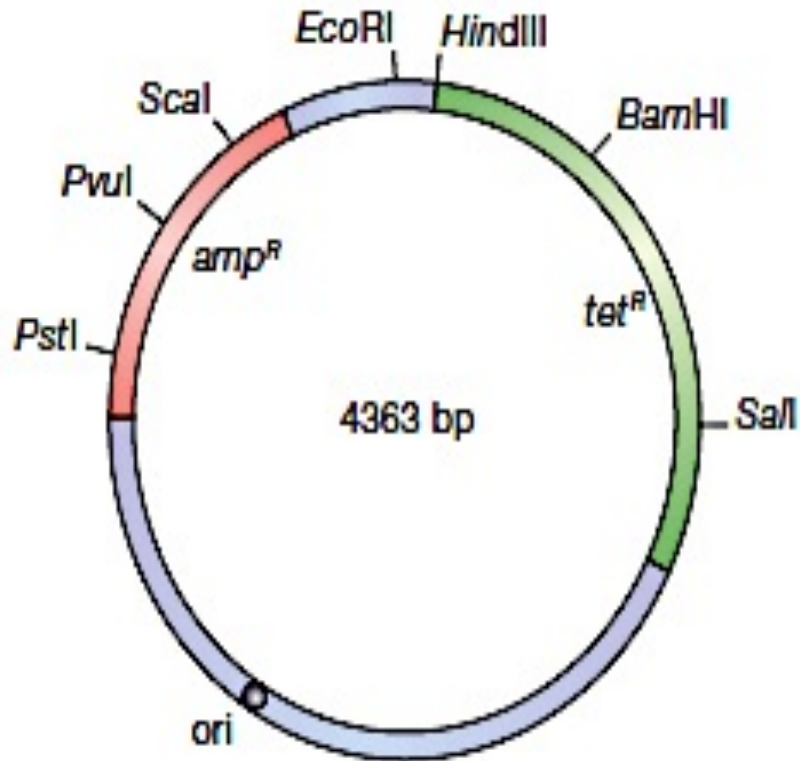
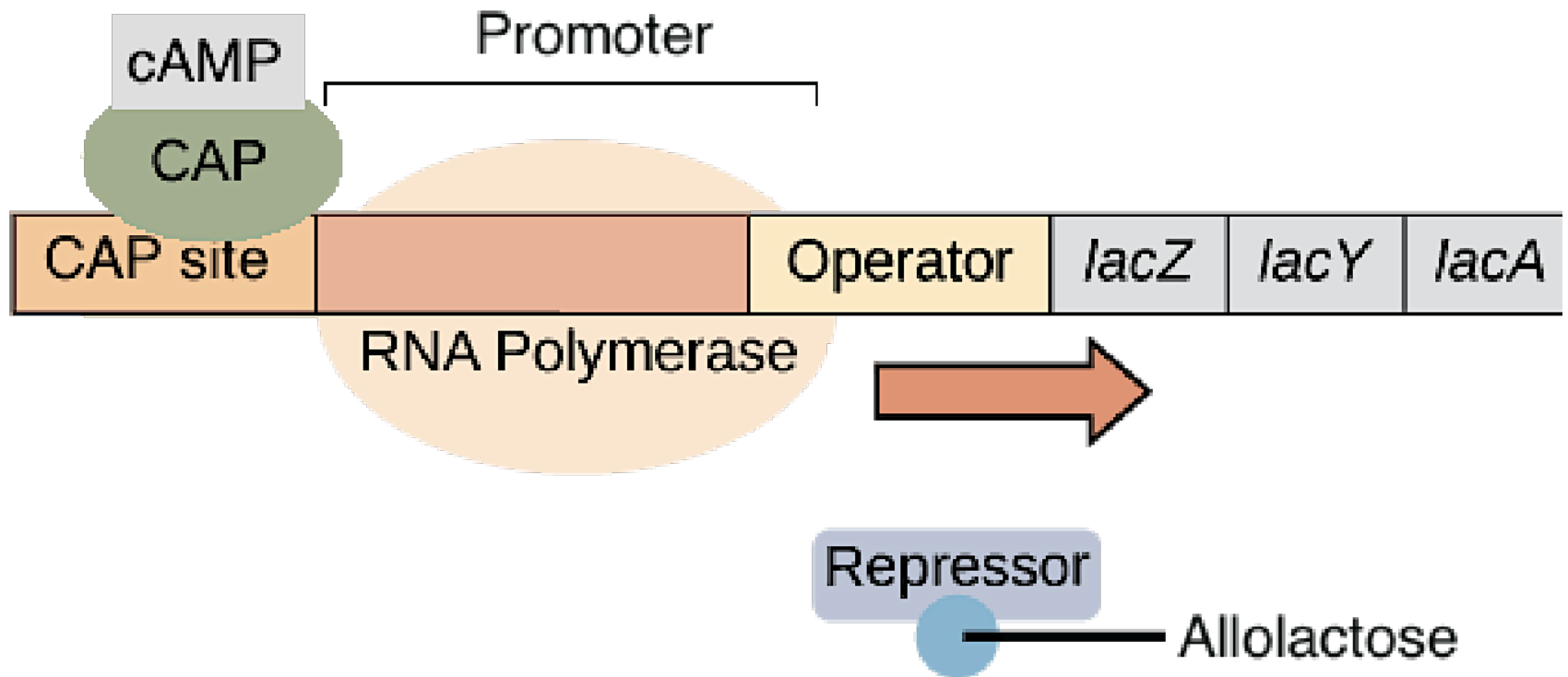


Figure 6.1

A map of pBR322 showing the positions of the ampicillin resistance (*amp<sup>R</sup>*) and tetracycline resistance (*tet<sup>R</sup>*) genes, the origin of replication (*ori*) and some of the most important restriction sites.

Glucose absent, lactose present:



# - pUC8- aLac selection plasmid

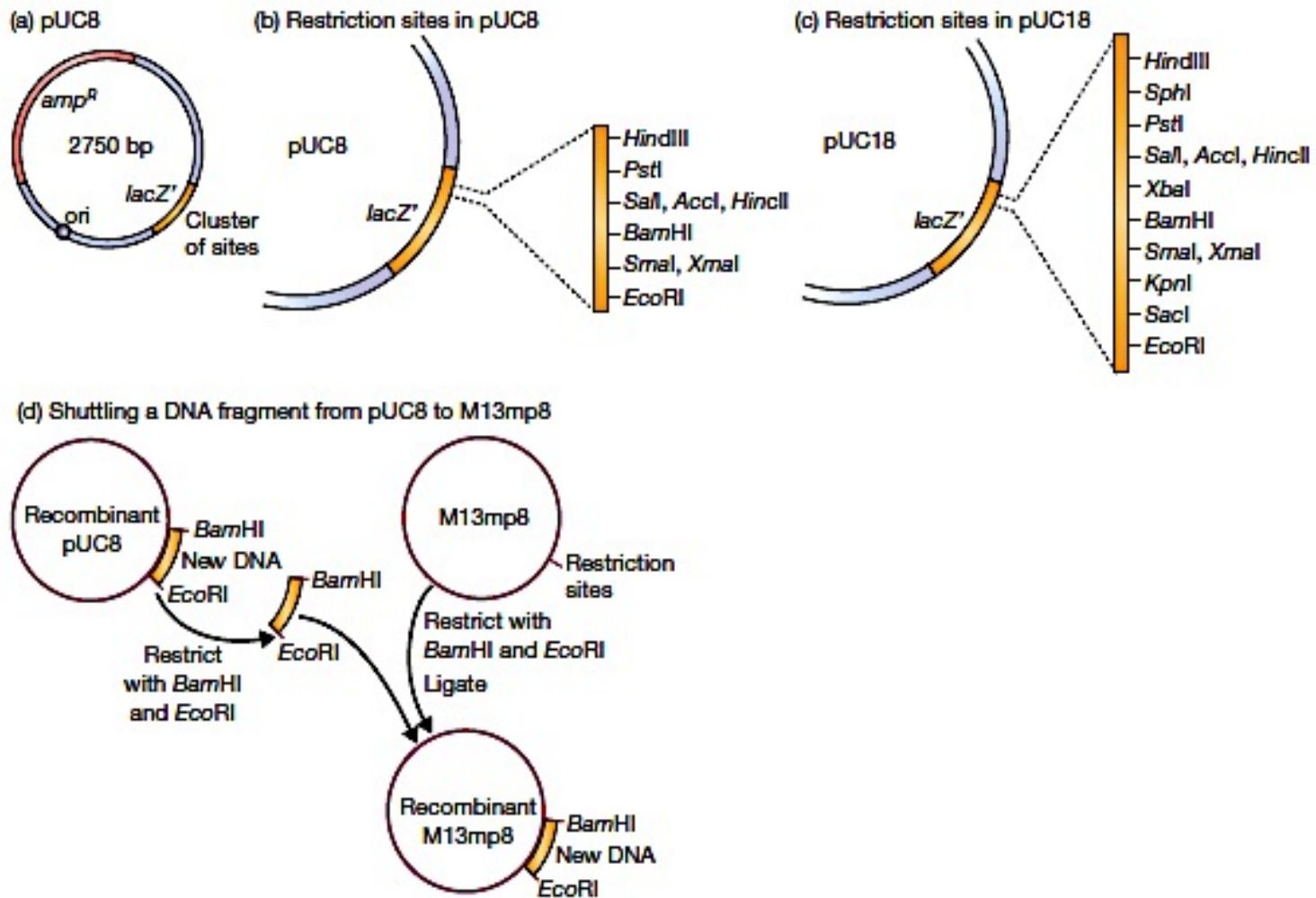


Figure 6.3

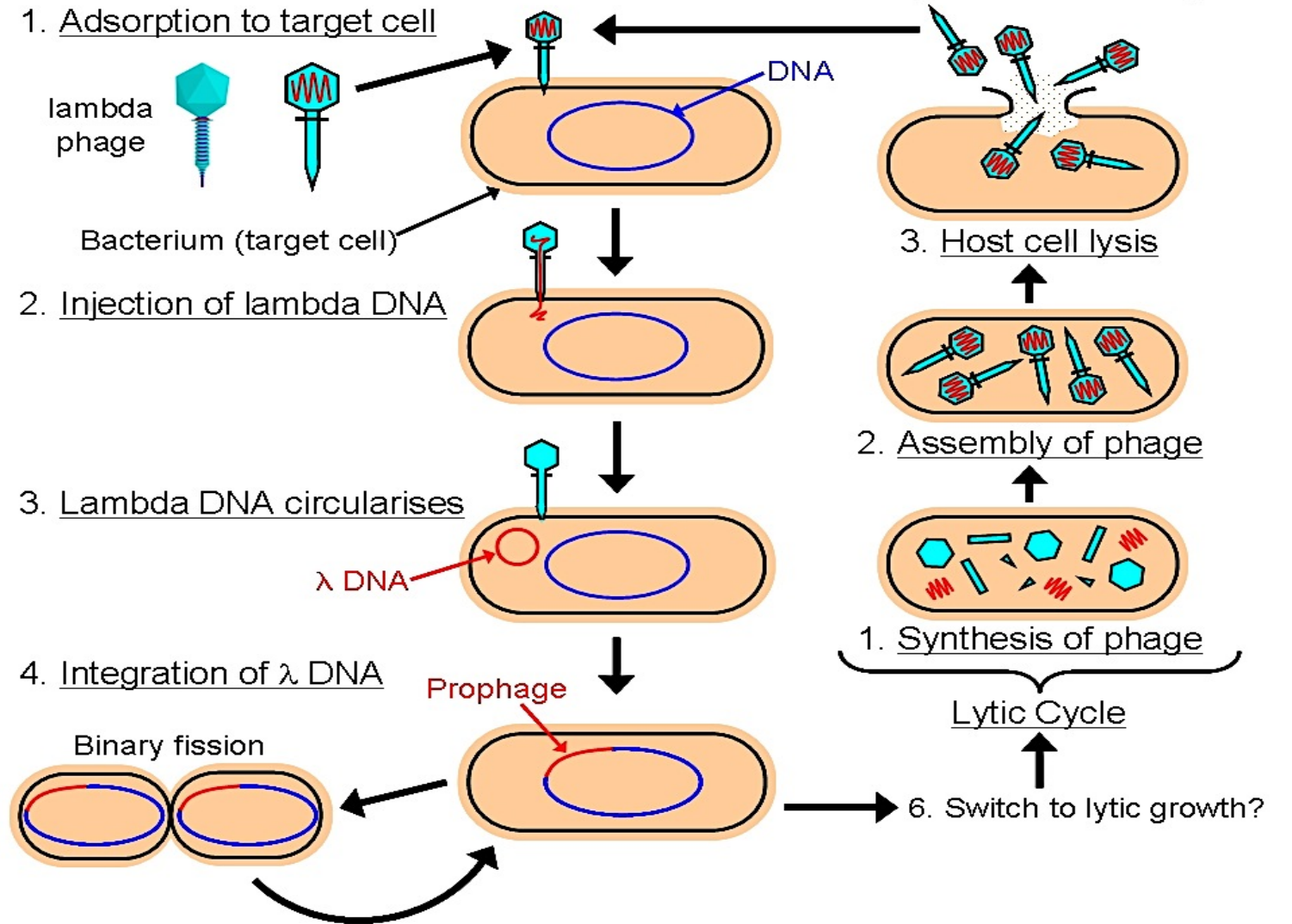
The pUC plasmids. (a) The structure of pUC8. (b) The restriction site cluster in the *lacZ'* gene of pUC8. (c) The restriction site cluster in pUC18. (d) Shuttling a DNA fragment from pUC8 to M13mp8.



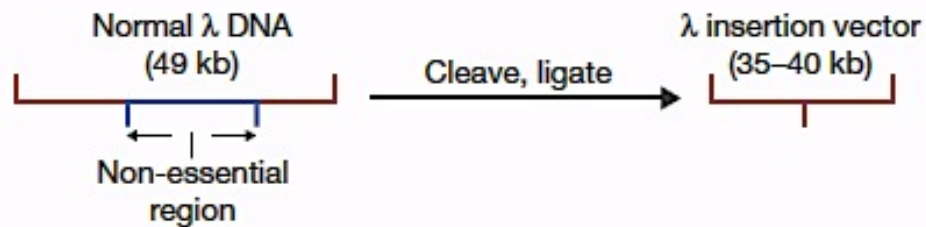
# - Blue-white screen



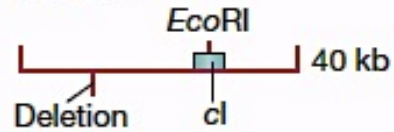
# Lysogenic Cycle of Lambda ( $\lambda$ ) Phage



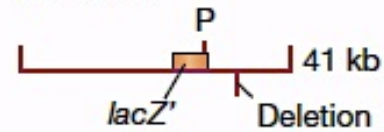
(a) Construction of a  $\lambda$  insertion vector



(b)  $\lambda$ gt10



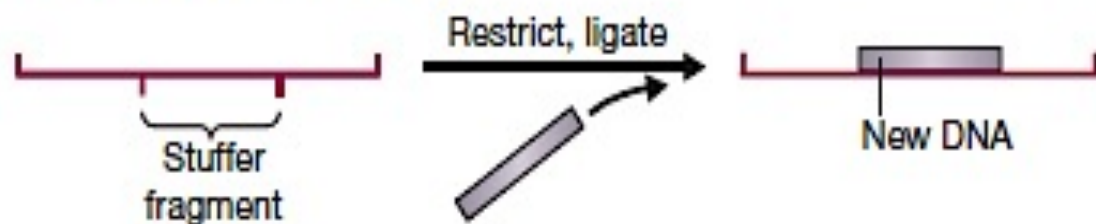
(c)  $\lambda$ ZAPII



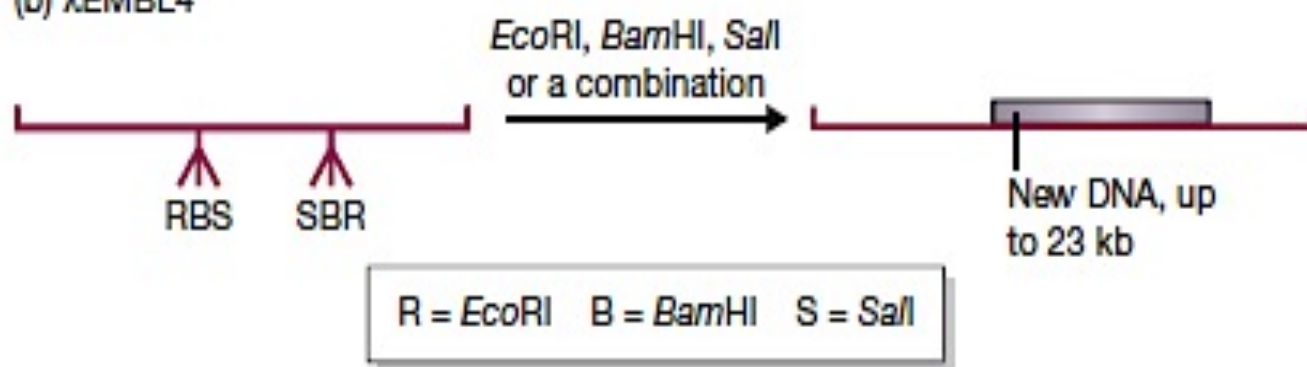
**Figure 6.12**

$\lambda$  insertion vectors. P = polylinker in the *lacZ'* gene of  $\lambda$ ZAPII, containing unique restriction sites for *SacI*, *NotI*, *XbaI*, *SpeI*, *EcoRI*, and *XhoI*.

(a) Cloning with a  $\lambda$  replacement vector



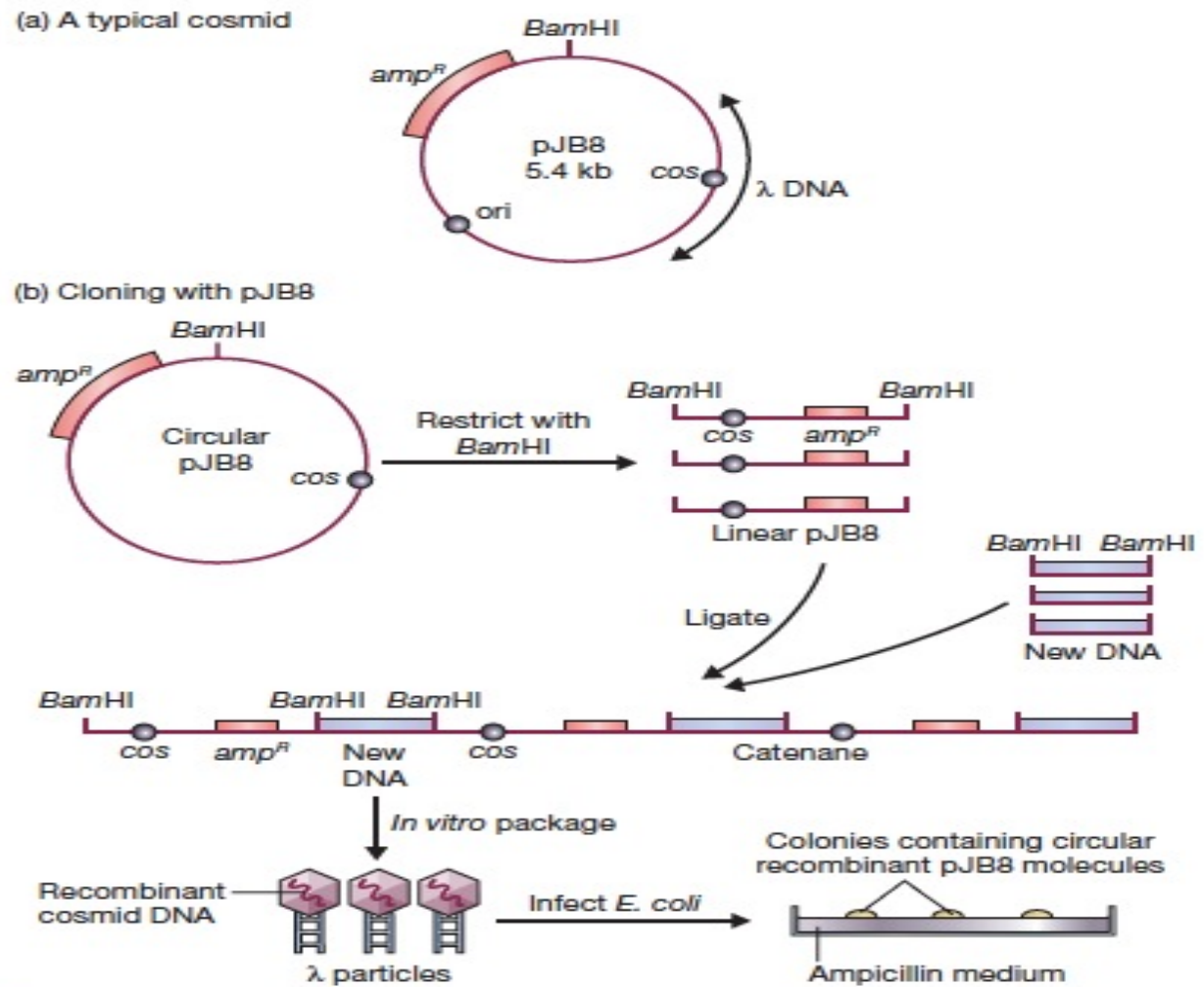
(b)  $\lambda$ EMBL4



**Figure 6.13**

$\lambda$  replacement vectors. (a) Cloning with a  $\lambda$  replacement vector. (b) Cloning with  $\lambda$ EMBL4.

# - Cosmid



**Figure 6.15**

A typical cosmid and the way it is used to clone long fragments of DNA.