

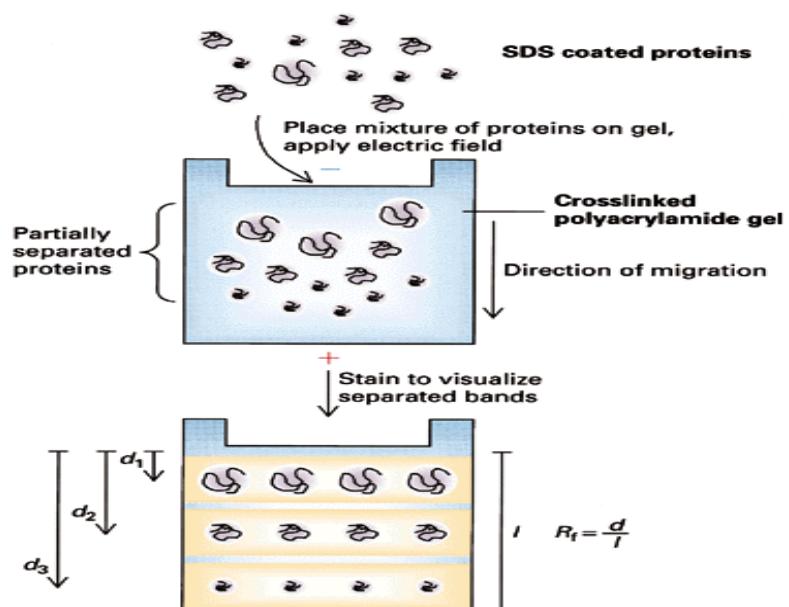
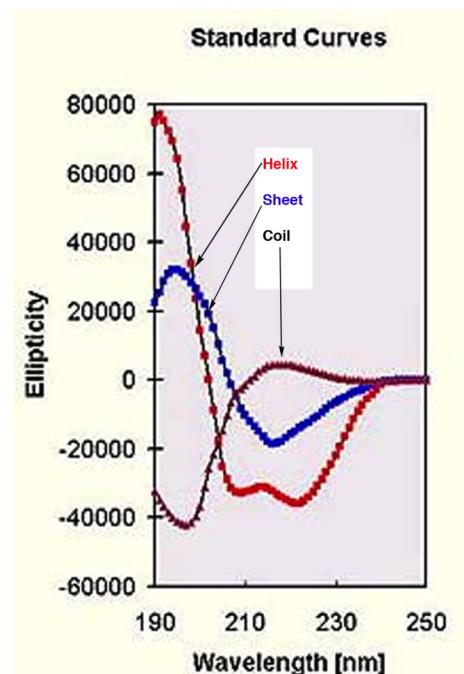
46 data points from 205 to 250 nm

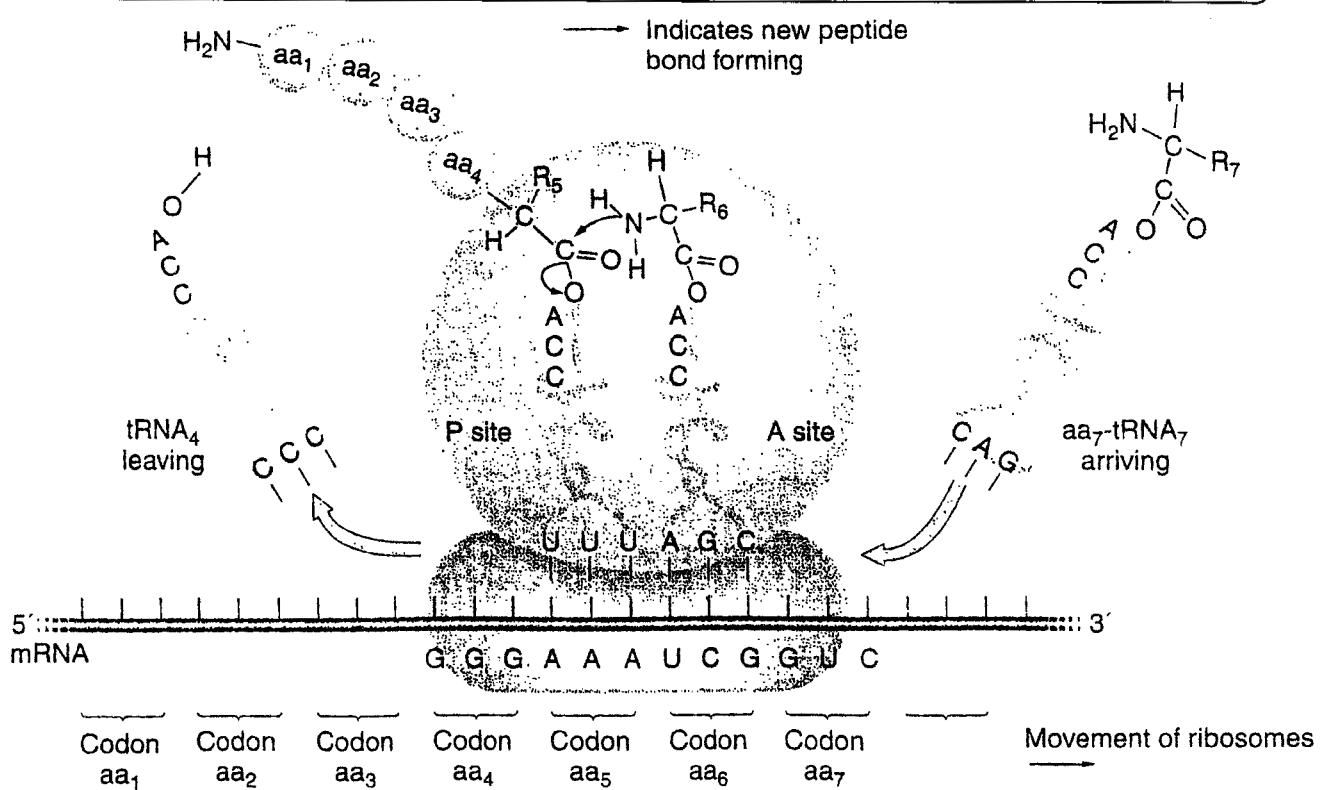
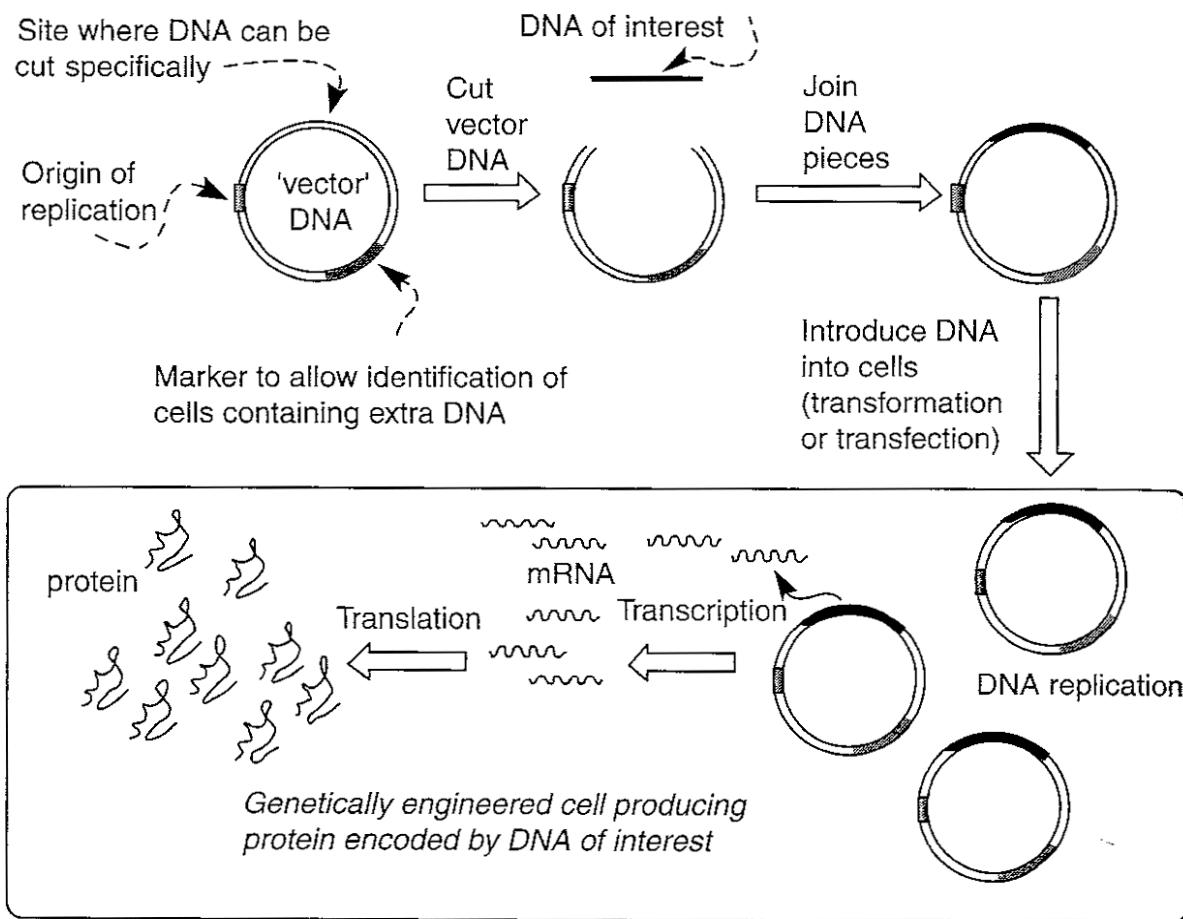
Molecular weight	g/mol	:	26698.00
Sample concentration	mg/cm ³	:	.500000
Sample concentration	g/cm ³	:	.00050000
Sample concentration	mol/cm ³	:	.00000002
Cell path	mm	:	.500000
Cell path	cm	:	.050000
Number of residues		:	269.000000
Conversion factor [cm ² /dmole]		:	106792000.00

Zero shift applied : 2.

Least squares percentage(s) and scale factor
57.92 26.22 15.85 .77

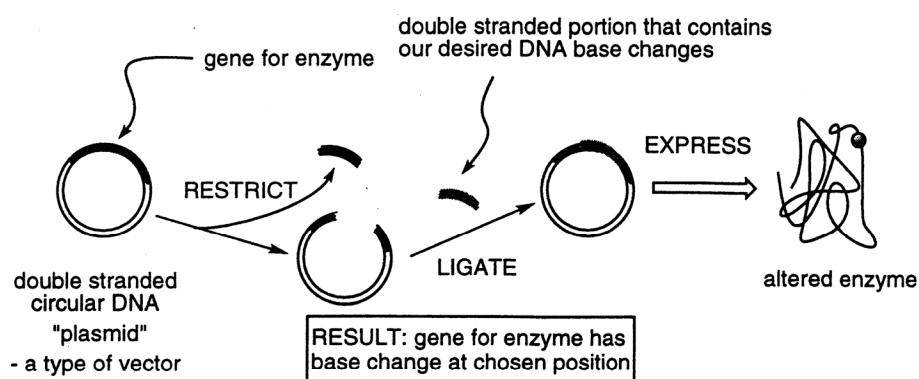
rmsd	helix	sheet	coil	scale
68.08	44.58	20.18	12.20	.77
Rfac %	helix	sheet	coil	
3.15	57.92	26.22	15.85	



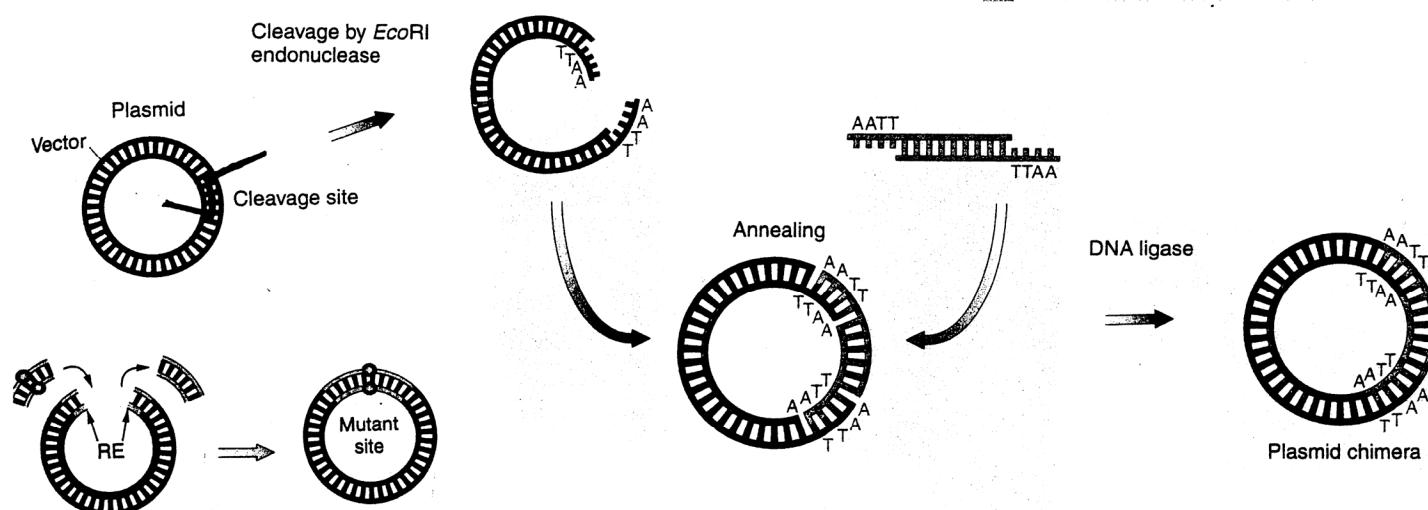


.....|UUU|UUA|GGGIAAIUCGIGUCI →|UUU|UUA|GGGIAAIU**G**CIGUCI
 Phe Leu Gly Lys Ser Val → Phe Leu Gly Lys Cys Val

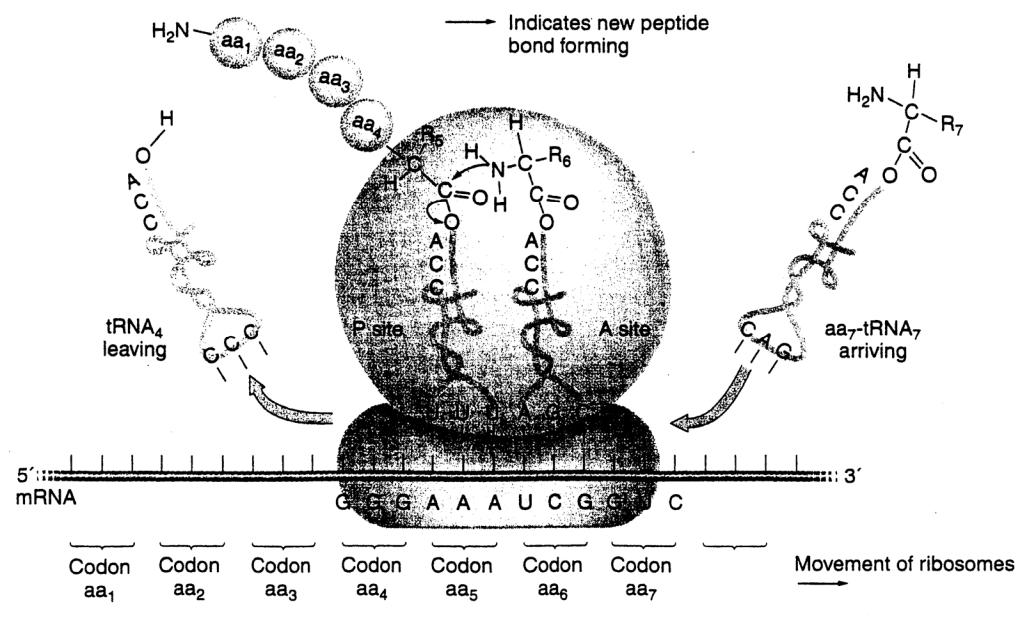
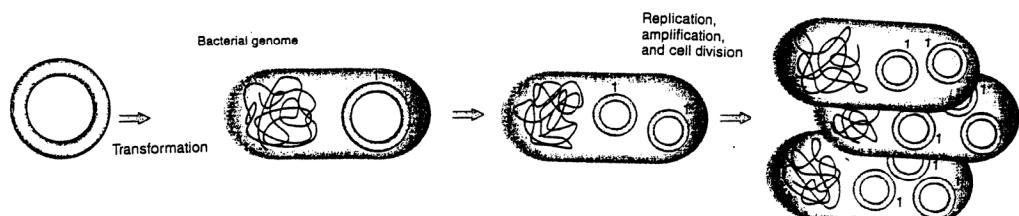
Site-Directed Mutagenesis



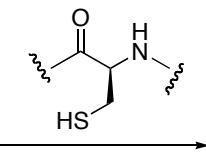
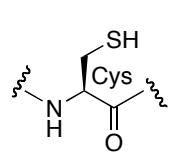
	U	C	A	G	
	UUU	UCU	UAU	UGU	Cys
U	UUC	UCC	UAC	UGC	Stop
U	UUA	UCA	UAA	UGA	Trp
U	UUG	UCG	UAG	UGG	
C	CUU	CCU	CAU	CGU	
C	CUC	CCC	CAC	CGC	
C	CUA	CCA	CAA	CGA	Arg
C	CUG	CCG	CAG	CGG	
A	AUU	ACU	AAU	AGU	
A	AUC	ACC	AAC	AGC	Ser
A	AUA	ACA	AAA	AGA	
A	AUG	ACG	AAG	AGG	Arg
G	GUU	GCU	GAU	GGU	
G	GLC	GCC	GAC	GGC	
G	GUU	GCA	GAA	GGA	Gly
G	GUG	GCG	GAG	GGG	



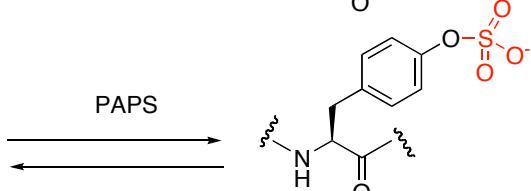
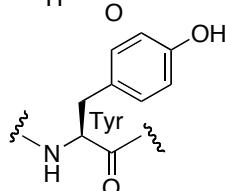
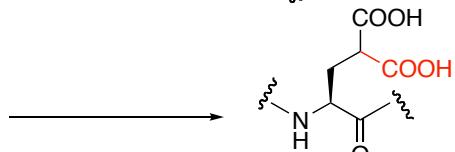
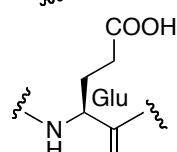
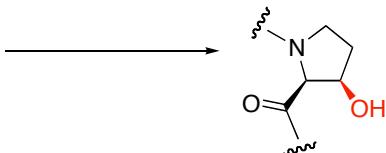
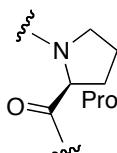
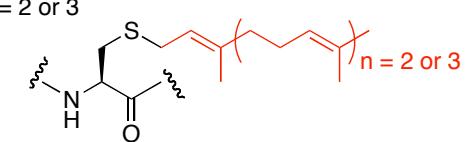
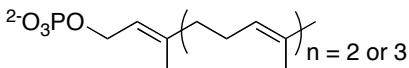
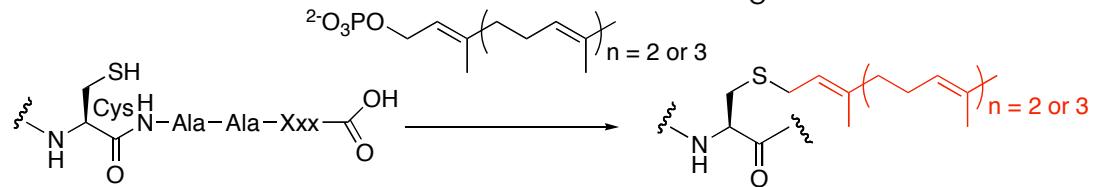
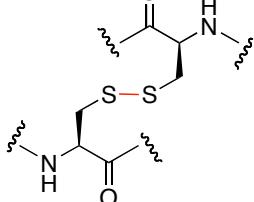
Enzyme	Source organism	Restriction site in double-stranded DNA
EcoRI	<i>Escherichia coli</i>	5' ↓ m -G-A-A-T-T-C- -C-T-T-A-A-G- 5' m ↑
EcoRII	<i>E. coli</i>	5' ↓ m -G-C-C-T-G-G-C- -C-G-G-A-C-C-G- 5' m ↑
HindIII	<i>Haemophilus influenzae</i>	5' ↓ m -G-T-Py-Pu-A-C- -C-A-Pu-Py-T-G- 5' m ↑
HindIII	<i>H. influenzae</i>	5' ↓ m -A-A-G-C-T-T- -T-T-C-G-A-A- 5' ↑m
HaeIII	<i>H. aegyptius</i>	5' ↓ -G-G-C-C- -C-C-G-G- 5'
HpaII	<i>H. parainfluenzae</i>	5' ↓ -C-C-G-G- -G-G-C-C- 5'
PstI	<i>Providencia stuartii</i>	5' ↓ -C-T-G-C-A-G- -G-A-C-G-T-C- 5'
SmaI	<i>Serratia marcescens</i>	5' ↓ -C-C-C-G-G- -G-G-C-C-C- 5'
BamI	<i>Bacillus amyloliquefaciens</i>	5' ↓ -G-G-A-T-C-C- -C-C-T-A-G-G- 5'



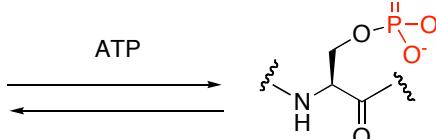
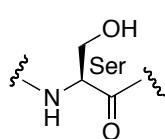
PROTEIN



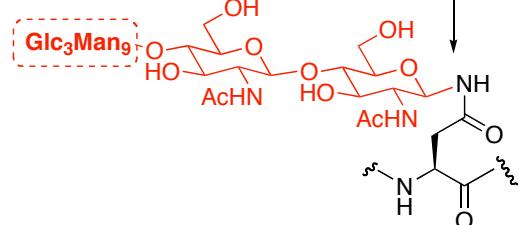
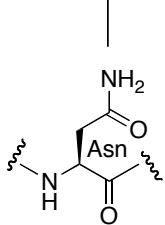
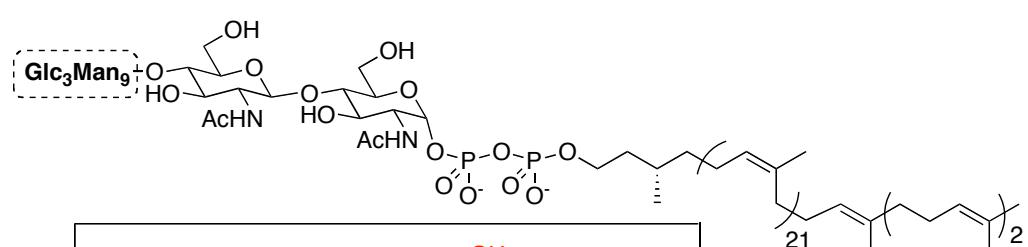
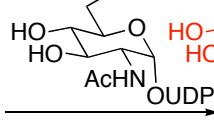
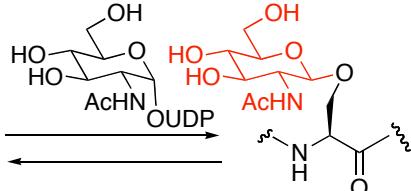
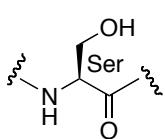
PROTEIN with PTM



PAPS



ATP



Additional Reading