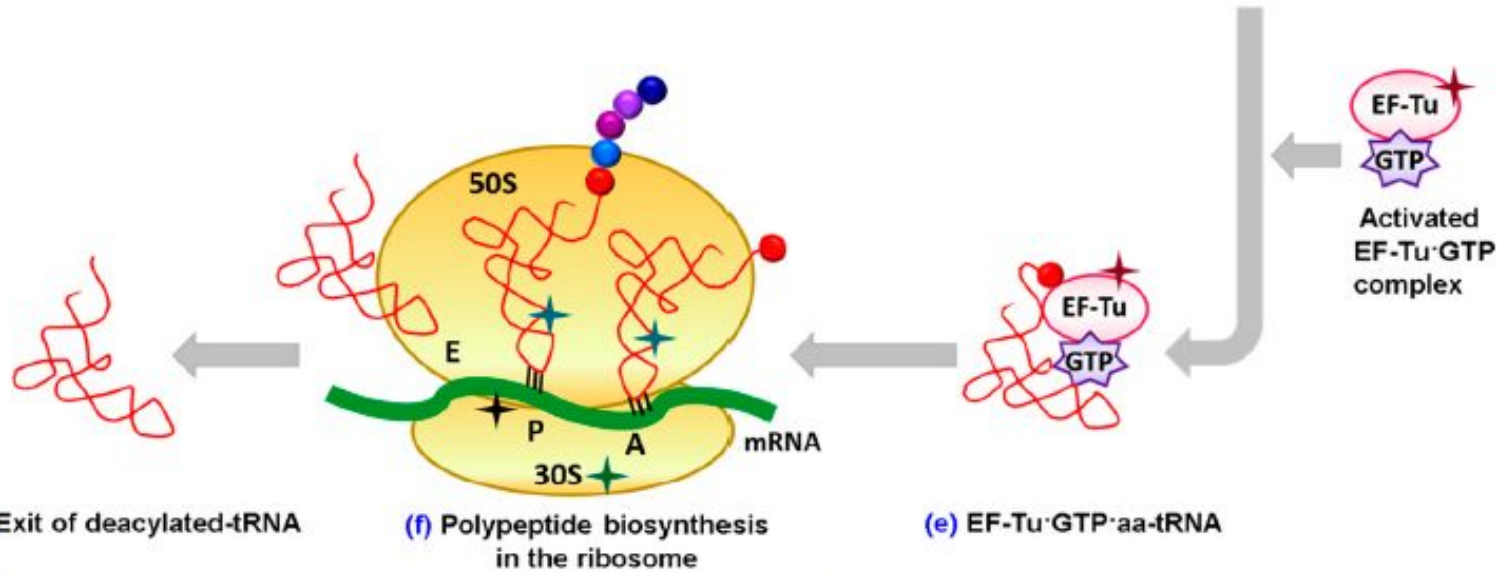
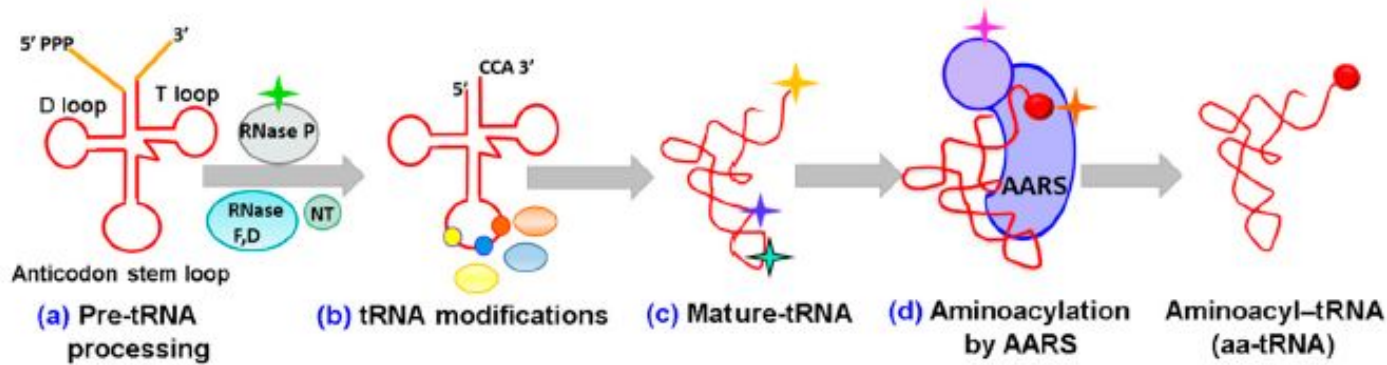


Antibiotics affecting codon  
phase-dependent binding of  
aminoacyl-tRNA to the ribosome.

Done by: Maulenova R.,  
Moldakozhayev A.,  
Naizabayeva D.

# Stages of tRNA modifications and functioning



★ Neomycin B, Bis-benzimidazole

★ Purpuromycin

★ Tobramycin

★ Colicins, Anticodon Stem Loop Inhibitors

★ AN2690

★ Agrocin 84 (TM84)

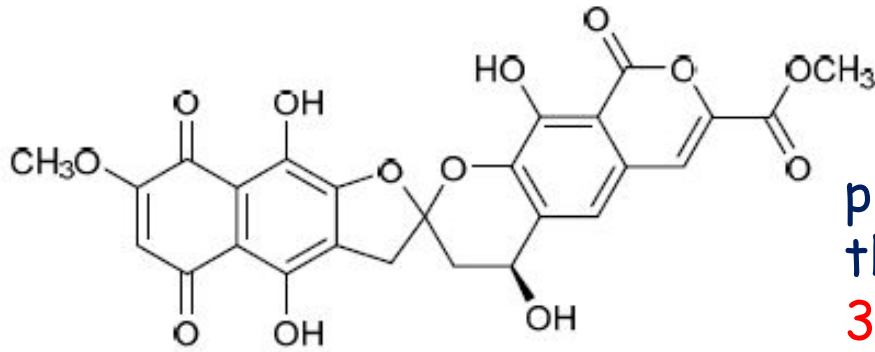
★ Aurodox, Encyloxin, GE2270A

★ Tetracycline, Aminoglycosides (Streptomycin, Neomycin B, Tobramycin)

★ Erythromycin, Chloramphenicol, Linezolid, Blasticidin S

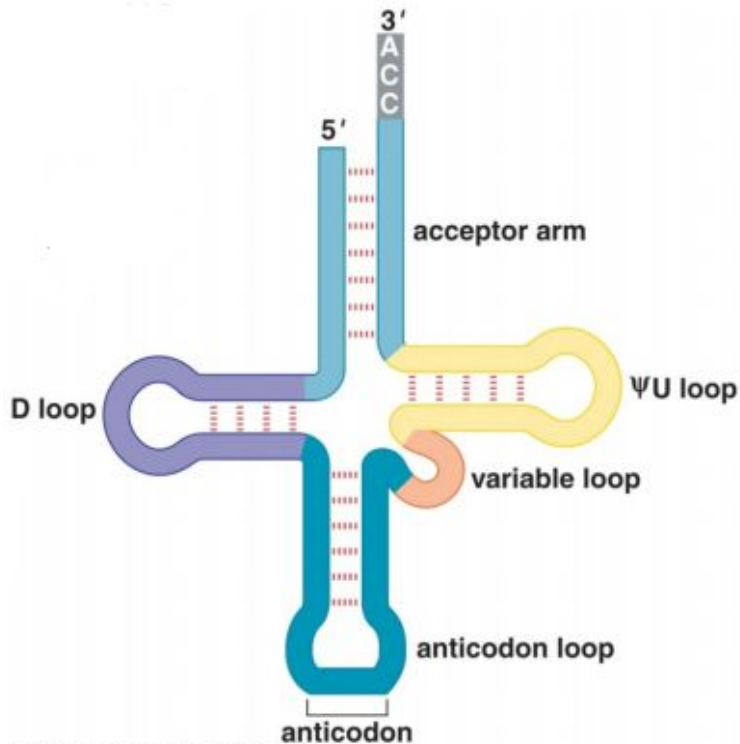
★ Viomycin, Thermorubin

# Blocking the CCA 3'-End of tRNA



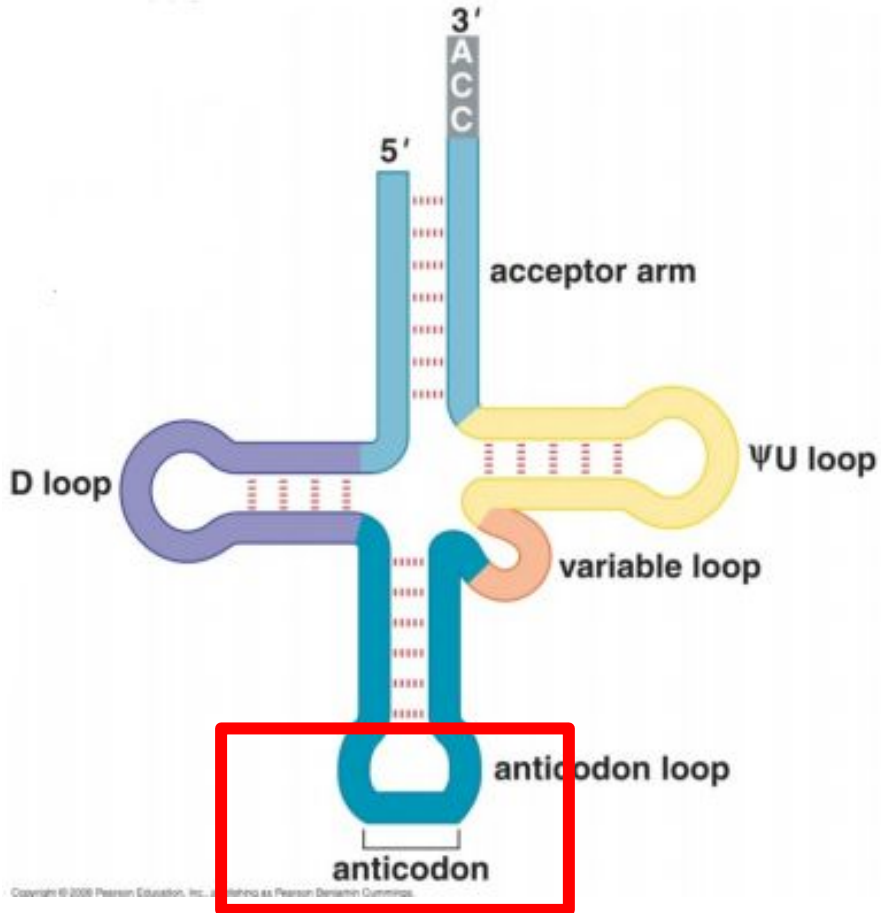
Purpuromycin

**Purpuromycin** is an antibiotic produced by *Actinoplanes ianthinogenes* that has been shown to bind to the **3'-acceptor stem** of all tRNAs with high affinity thereby **preventing the aminoacylation** of tRNA by its cognate amino acid



**Purpuromycin** is active against Gram-positive bacteria, such as *Bacillus subtilis*, *Candida albicans* and protozoa, such as *Trichomonas* sp.

# Cleavage of tRNAs



**Colicins** are antibacterial toxins secreted out into the extracellular medium by members of the **enterobacteriaceae** family, such as ***E. coli*** (about 30% of *E. coli* contain them).

Function: DNase activity, RNase activity, depolarization of the cytoplasmic membrane, and inhibition of murein synthesis.

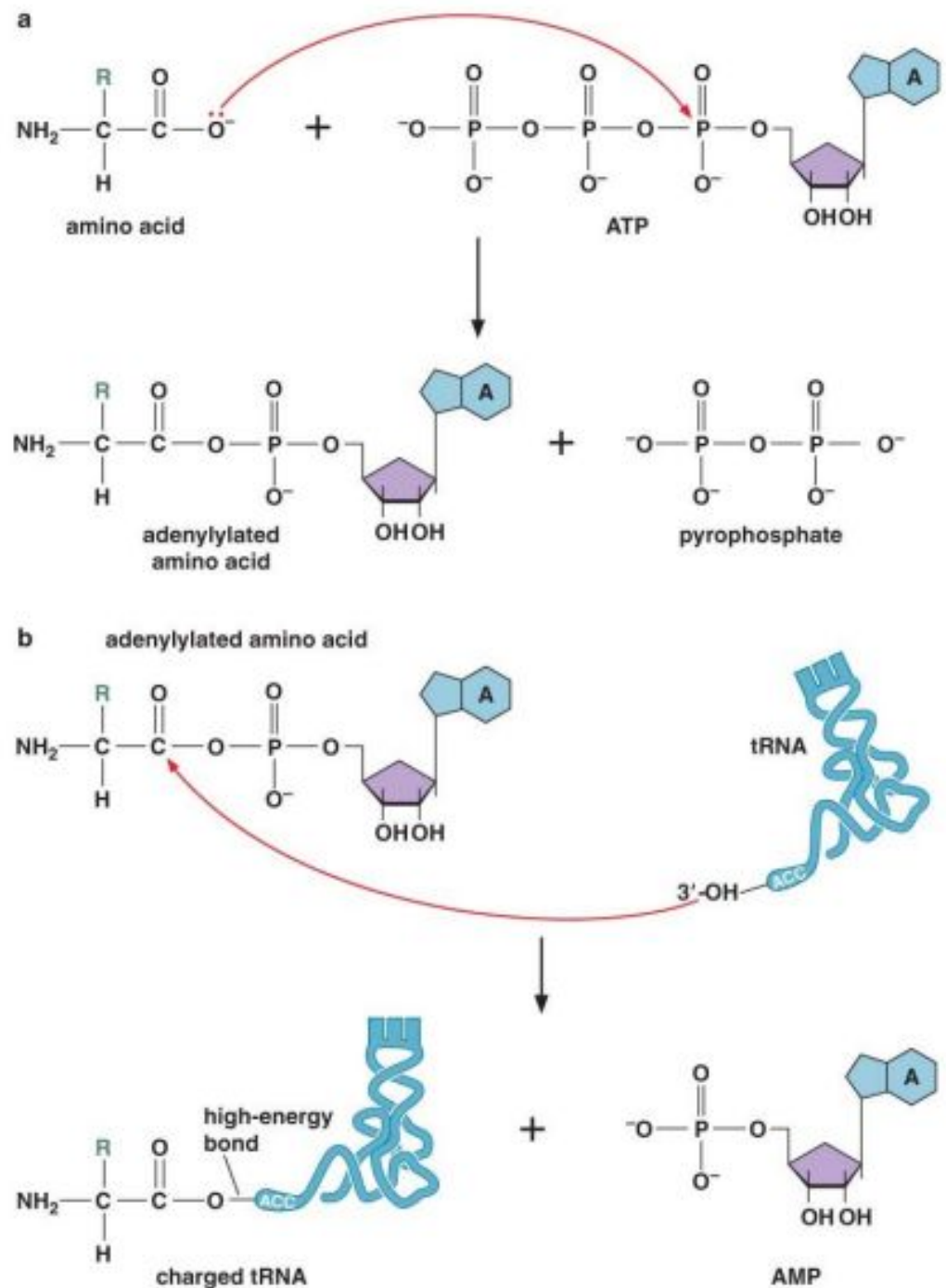
For example, **Colicin E5 RNase** targets tRNAs specific for **tyrosine**, **histidine**, **asparagine** and **aspartic acid** by **cleaving anticodon QUN** that contains the hyper-modified queuosine nucleotide (Q) at the wobble position 34.

# Aminoacyl-tRNA A formation

**Aminoacyl-tRNA synthetases** Charge tRNAs in two steps:

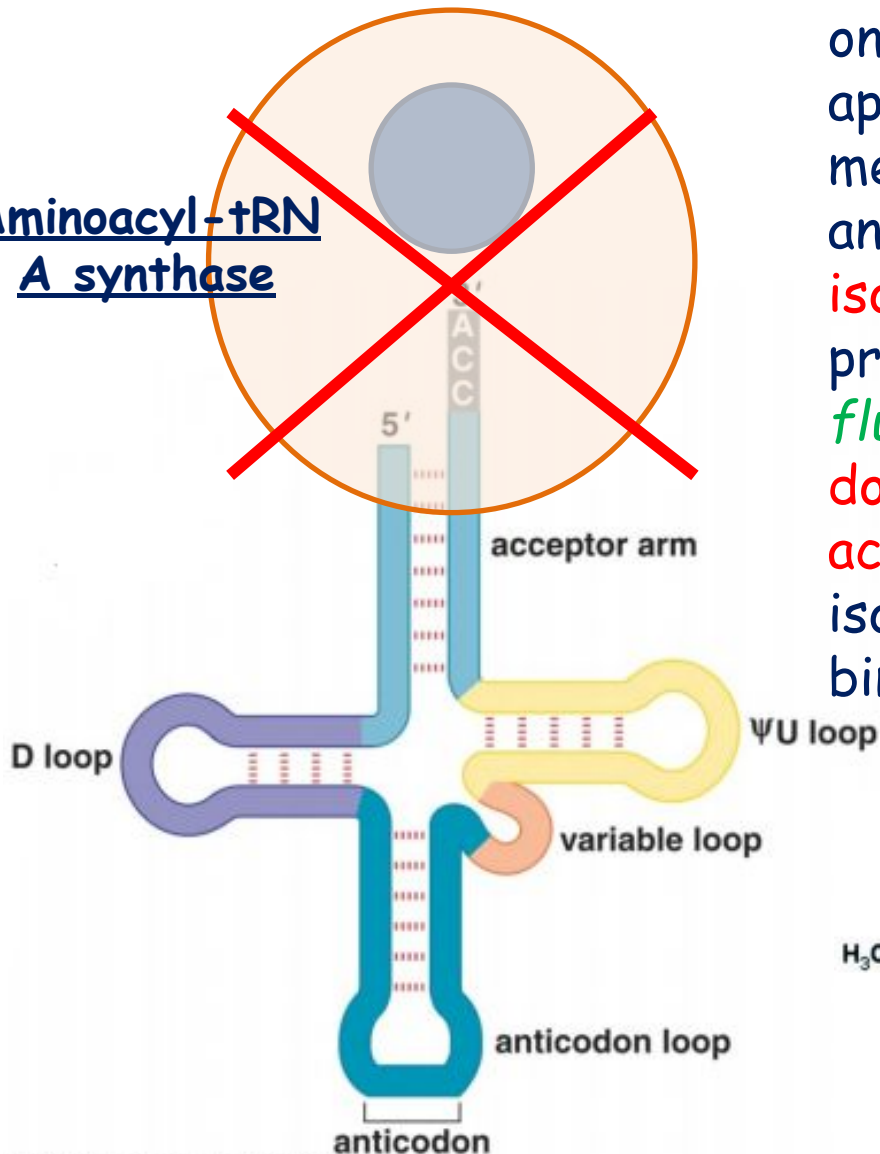
**1) adenylylation:** Amino acid react with ATP and AMP is transferred to amino acids

**2) tRNA charging:** transfer of aAmino acid to the 3' end of tRNA via 2'- or 3'-OH and release of AMP

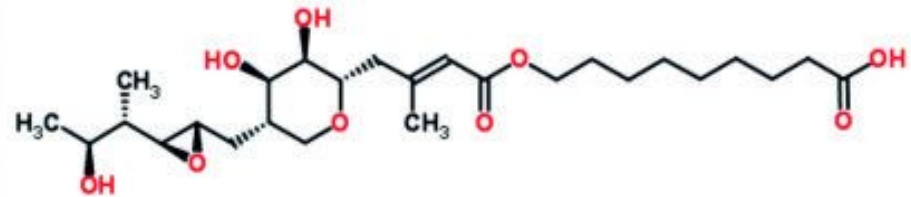


# Antibiotics Inhibiting Aminoacylation of tRNA

Aminoacyl-tRNA  
A synthase



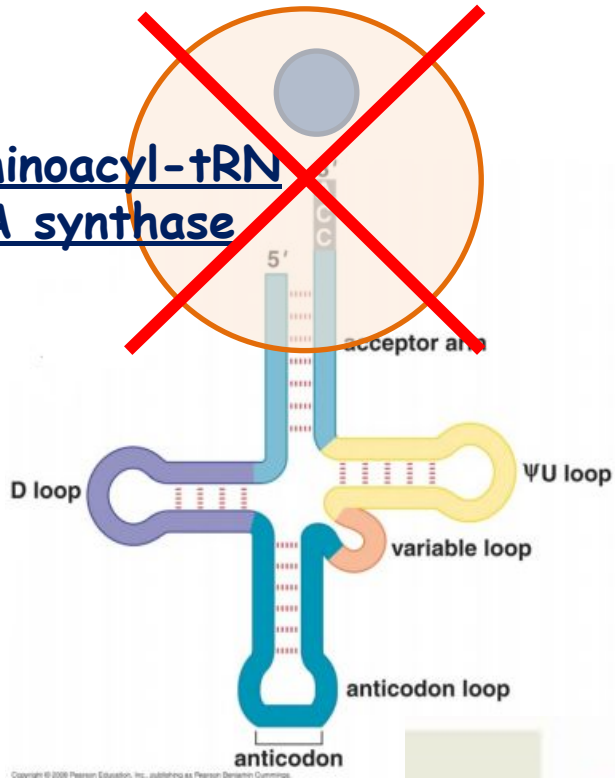
**Pseudomonic acid** or **mupirocin** is one of the most effective topically applied antibiotics used to combat methicillin resistant *S. aureus*. This antibiotic is a naturally occurring **isoleucyl-tRNA synthetase inhibitor** produced by *Pseudomonas fluorescens* strains and works by **docking onto the enzyme catalytic active site** and competing with the isoleucine and ATP substrates for binding.



**Mupirocin**

# Inhibition by Trapping tRNA in a LeuRS Editing Domain

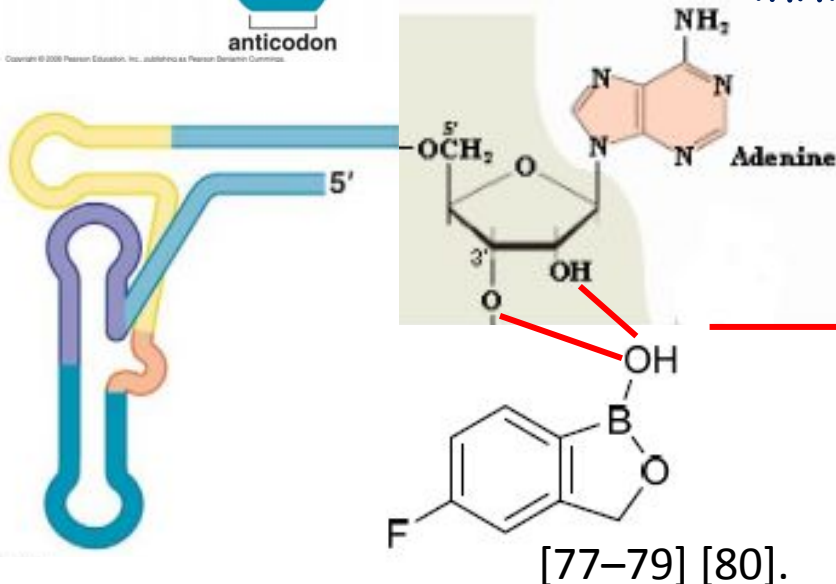
Aminoacyl-tRNA synthase



*LeuRS*, *valyl-tRNA synthetase* (*ValRS*) and *IleRS*, possess an additional proofreading domain called the *CP1 domain*.

Function: recognizing and hydrolyzing misacylated amino acids on the 3'-end of the tRNA.

Novel synthetic compound [Anchor company], **AN2690** (5-fluoro-1,3 dihydro-1-hydroxy-2,1-benzoxazole) inhibited a fungal *LeuRS* from the yeast *Saccharomyces cerevisiae*.



Contacts made with the 2' and 3'-oxygen atoms of the ribose of the 3'-terminal adenosine of tRNA leads to the formation of a stable tRNA<sub>Leu</sub>-AN2690 adduct

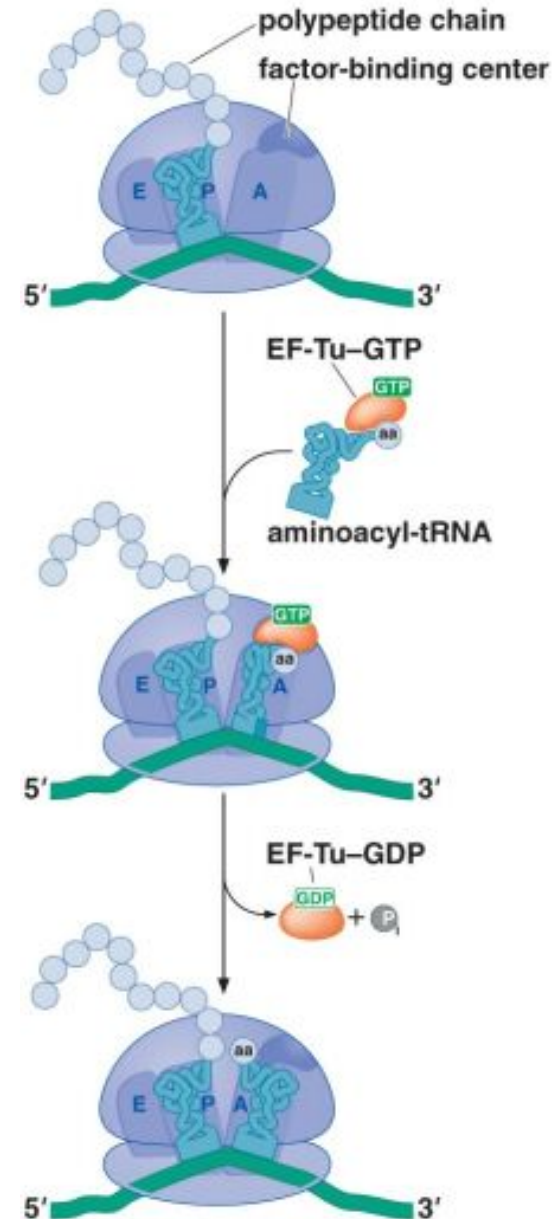
# Antibiotics Affecting Elongation Factor EF-Tu

-aminoacyl-tRNA is escorted to the ribosome by elongation factor EF-Tu

-EF-Tu binds to tRNA's 3' end, masking the coupled amino acid -> \*prevent the bound aminoacyl-tRNA from participating in peptide bond formation

\*affinity of EF-Tu is regulated by GTP status

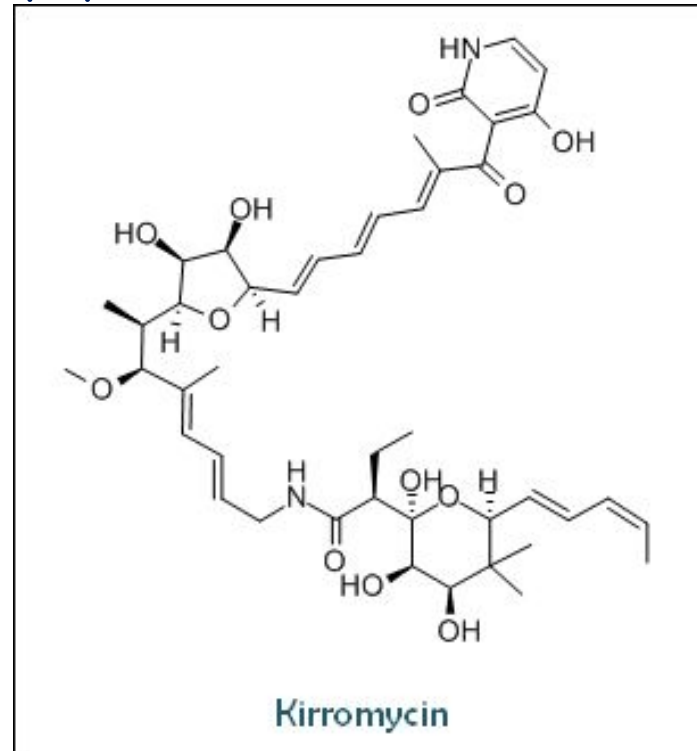
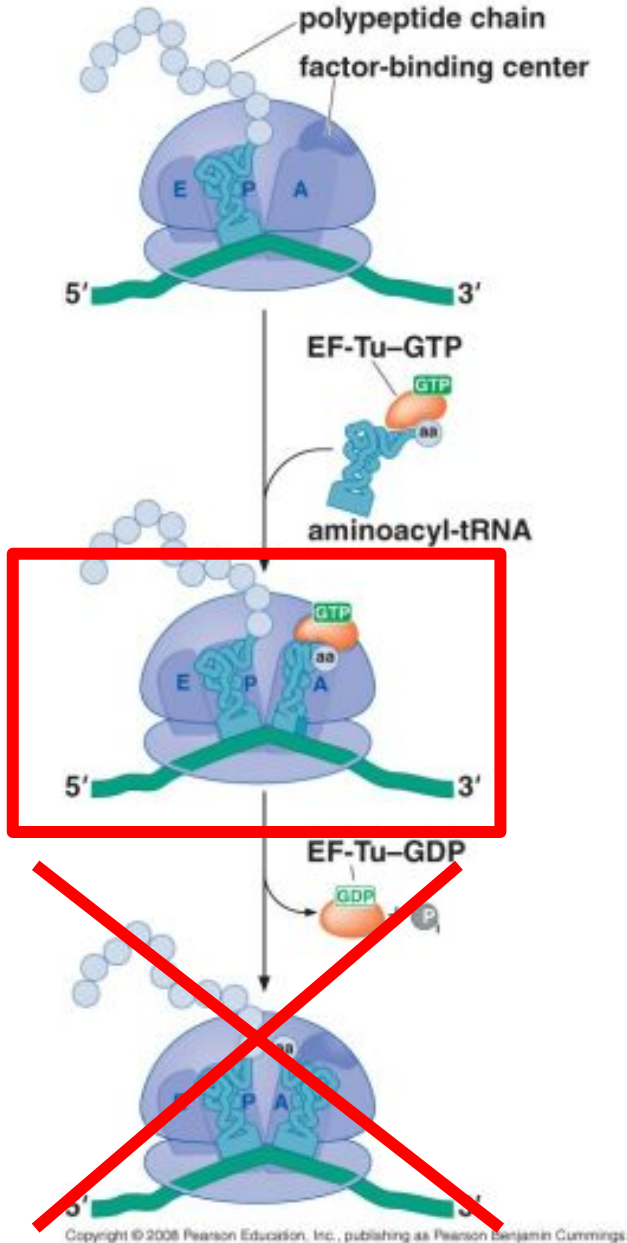
\*control of GTP hydrolysis by EF-Tu is critical to the specificity of translation





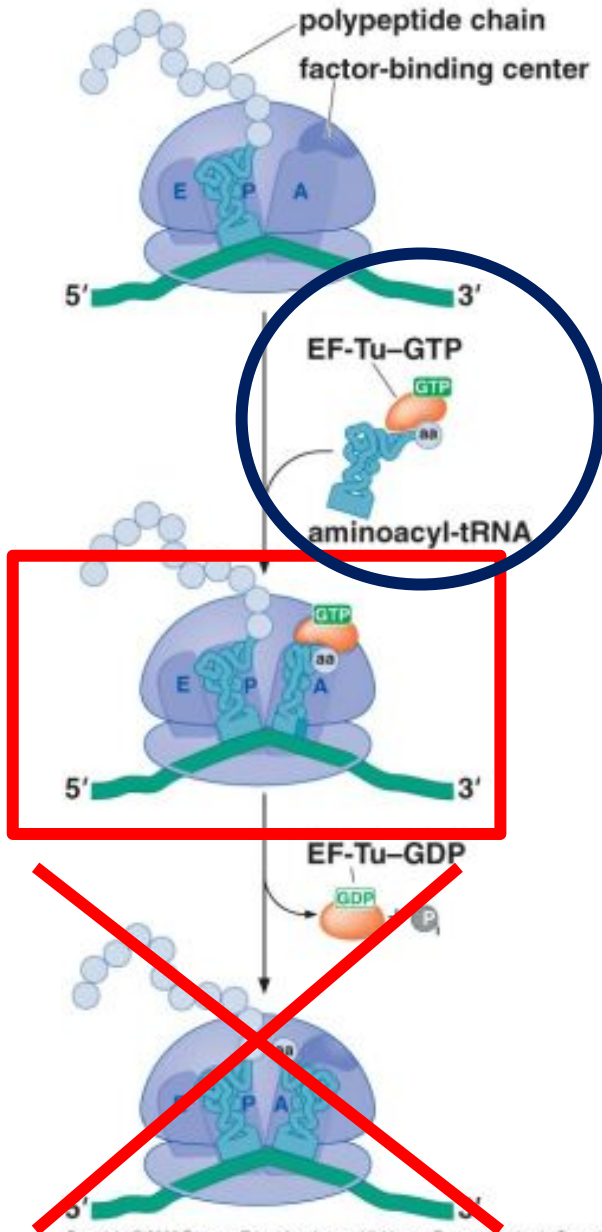
# Antibiotics Affecting Elongation Factor EF-Tu

Kirromycin is an antibiotic that binds to the ribosome\*aa-tRNA\*EF-Tu\*GDP complex. This results in inhibition of the release of the EF-Tu·GDP complex from the ribosome. **Failure of the release of EF-Tu from the ribosome** does not affect the binding of aminoacyl-tRNA to the A-site of the ribosome, but blocks the subsequent peptide bond formation step.

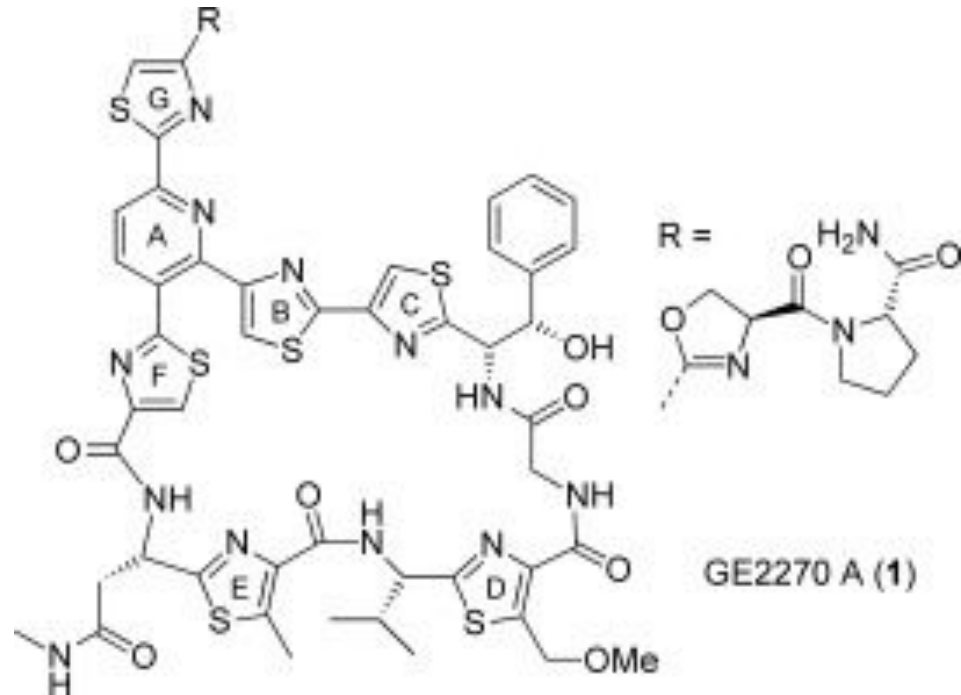




# Antibiotics Affecting Elongation Factor EF-Tu



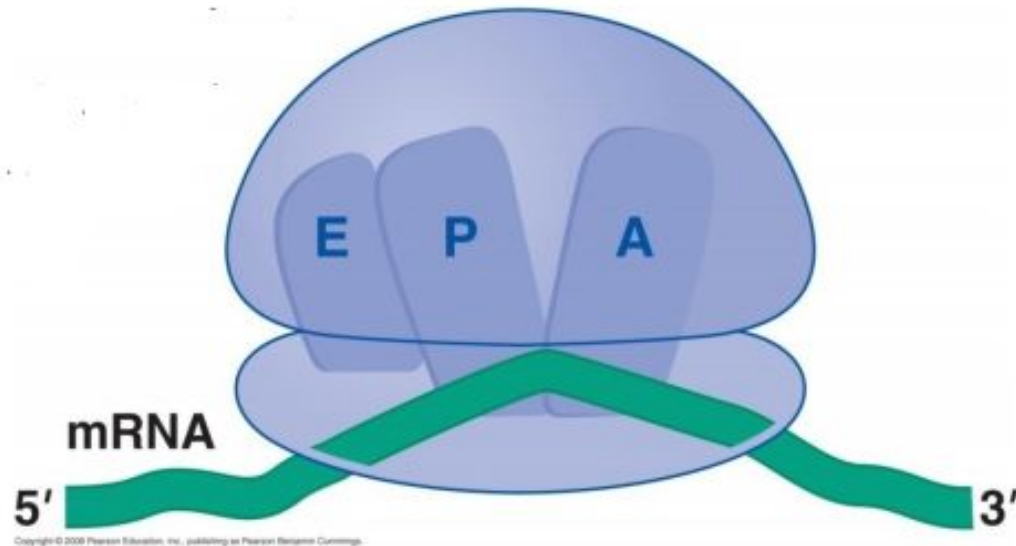
**GE2270A** is a **thiazolyl peptide** antibiotic that is active against Gram-positive bacteria. Crystal structure of *E. coli* **EF-Tu\*GDP\*GE2270** complex has confirmed that this compound directly **competes with aminoacyl-tRNA for the same binding site** on EF-Tu. It also blocks the GTP to GDP conformational change in EF-Tu.



# Targeting tRNAs in the Ribosome

The ribosome has three binding sites for tRNA

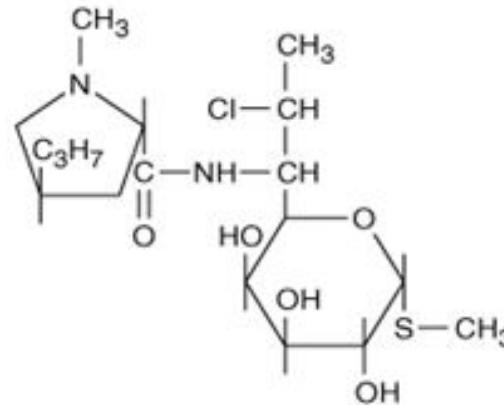
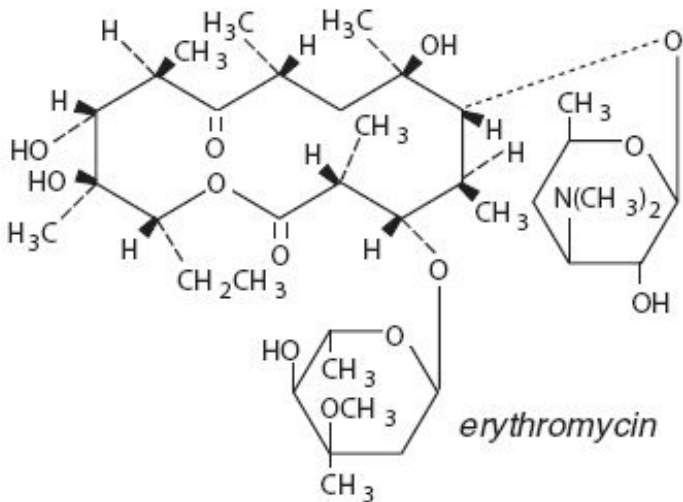
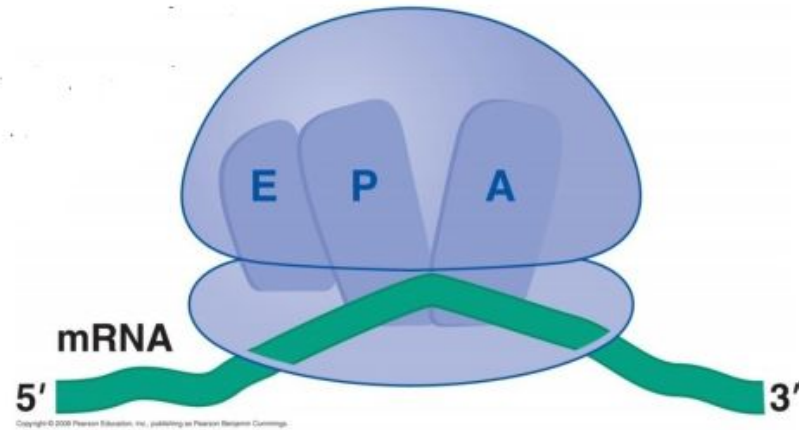
- 1) **A site**: binding site for aminoacyl-tRNA
- 2) **P site**: binding site for peptidyl-tRNA
- 3) **E (denote exit) site**: binding site for tRNA released after growing polypeptide chain has been transferred to the aminoacyl-tRNA



# Targeting tRNAs in the Ribosome

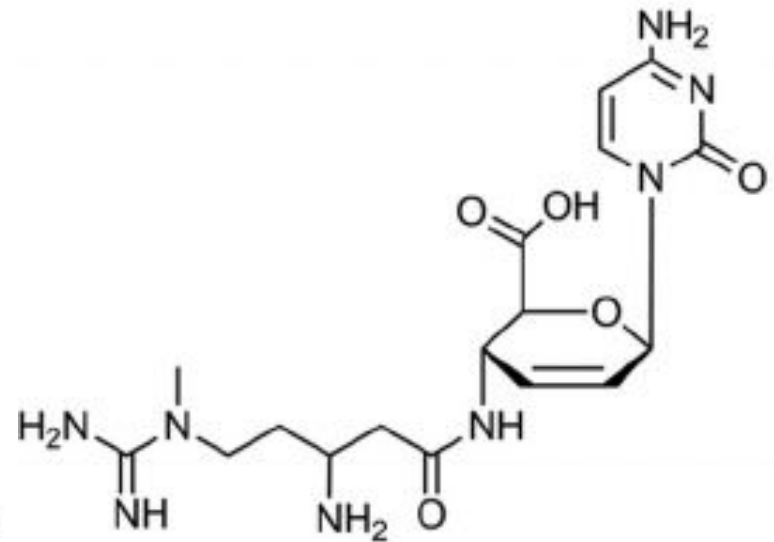
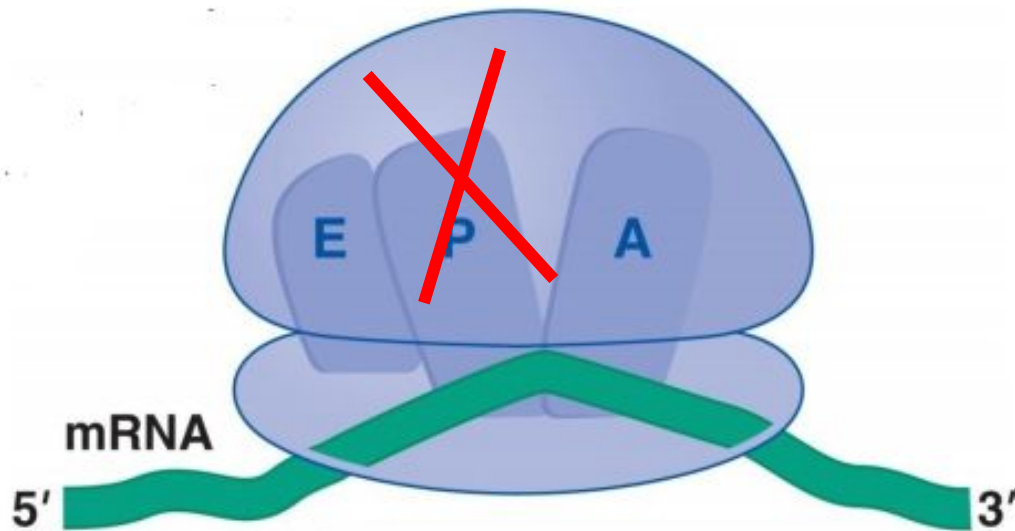
Inhibitors that prevent the binding of the initiator tRNA at the **P-site** - **oxazolidinones** (**linezolid**)

Antibiotics that prevent peptide bond formation and/or the **translocation** of tRNA from the A-site to the P-site on the ribosome - **macrolide** (**erythromycin**), **lincosamide** (**clindamycin**) and streptogramin (**dalfopristin**) class of antibiotics).



# Targeting tRNAs in the Ribosome

**Blasticidin S** is an antibiotic produced by *Streptomyces griseochromogenes*. **BlaS** has been found to be a potent inhibitor of both prokaryotic and eukaryotic cells. **BlaS** binds to the **50S subunit** of the ribosome at the **P-site** and not at the A-site like other Bla antibiotics. Upon binding the P-site, **BlaS bends the CCA 3'-end** of the tRNA bound at the P-site **to the A-site** resulting **shift** in the ribose phosphate backbone of the base C75 of the tRNA. This results in a **decrease in the flexible movement of the CCA 3'-end** of the tRNA, an important feature required by **translation**.



Blasticidin S

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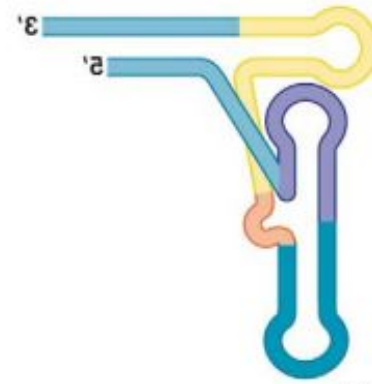
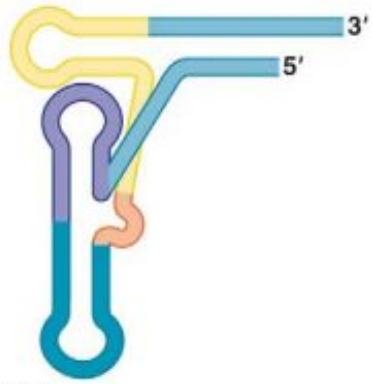
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Thanks  
for  
attention!

