



# Rutas bio-sintéticas de antibióticos y toxinas

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A stylized, layered mountain range graphic in shades of teal and blue, located in the bottom right corner of the slide.

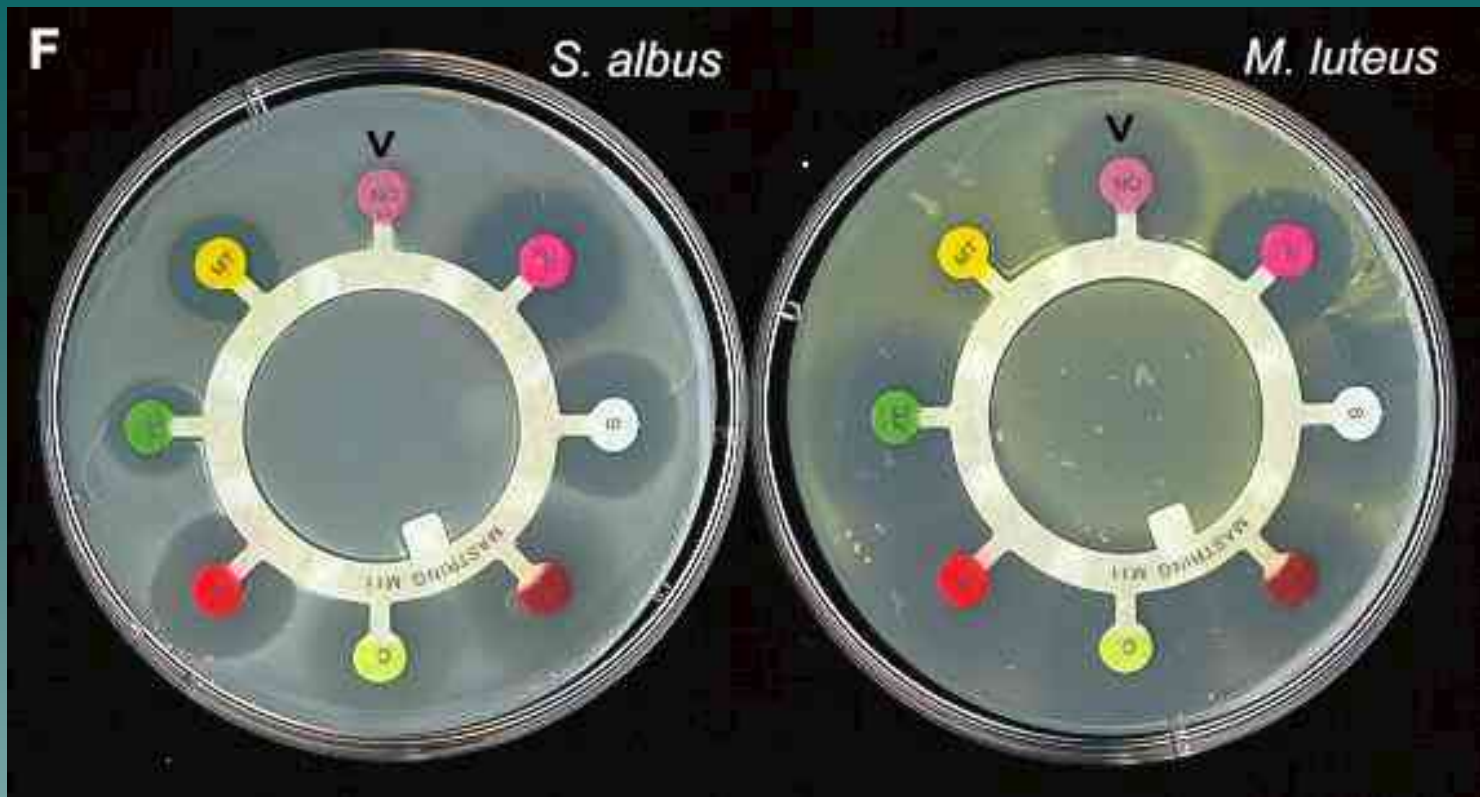
# Antibiotic Production

- ◆ There are approx. 8000 known antibiotics
  - ◆ More than 100 are produced commercially by microbial fermentation
  - ◆ 70% of these are produced by *Streptomyces* spp.
  - ◆ Strain selection and development has resulted in very high product yields (e.g., 1100 g/L Penicillin)
- 
- A stylized silhouette of a mountain range in shades of teal, located in the bottom right corner of the slide.

# Common antibiotics and their sources

- ◆ Bacitracin
  - ◆ Cephalosporin(s)
  - ◆ Chloramphenicol
  - ◆ Cycloheximide
  - ◆ Hygromycin
  - ◆ Penicillin
  - ◆ Streptomycin
  - ◆ Tetracycline(s)
  - ◆ Vancomycin
- ◆ *Bacillus subtilis*
  - ◆ *Cephalosporium* sp.
  - ◆ *S. venezuelae*
  - ◆ *S. griseus*
  - ◆ *S. hygromyces*
  - ◆ *P. chrysogenum*
  - ◆ *S. griseus*
  - ◆ *S. aureofaciens*
  - ◆ *S. orientalis*

	Some clinically important antibiotics			
	Antibiotic	Producer organism	Activity	Site or mode of action
	Penicillin	<i>Penicillium chrysogenum</i>	Gram-positive bacteria	Wall synthesis
	Cephalosporin	<i>Cephalosporium acremonium</i>	Broad spectrum	Wall synthesis
	Griseofulvin	<i>Penicillium griseofulvum</i>	Dermatophytic fungi	Microtubules
	Bacitracin	<i>Bacillus subtilis</i>	Gram-positive bacteria	Wall synthesis
	Polymyxin B	<i>Bacillus polymyxa</i>	Gram-negative bacteria	Cell membrane
	Amphotericin B	<i>Streptomyces nodosus</i>	Fungi	Cell membrane
	Erythromycin	<i>Streptomyces erythreus</i>	Gram-positive bacteria	Protein synthesis
	Neomycin	<i>Streptomyces fradiae</i>	Broad spectrum	Protein synthesis
	Streptomycin	<i>Streptomyces griseus</i>	Gram-negative bacteria	Protein synthesis
	Tetracycline	<i>Streptomyces rimosus</i>	Broad spectrum	Protein synthesis
	Vancomycin	<i>Streptomyces orientalis</i>	Gram-positive bacteria	Protein synthesis
	Gentamicin	<i>Micromonospora purpurea</i>	Broad spectrum	Protein synthesis
	Rifamycin	<i>Streptomyces mediterranei</i>	Tuberculosis	Protein synthesis



**Figure F.** Antibiotic-sensitivity testing. Petri dishes were spread-inoculated with *Staphylococcus albus* (white growth) or *Micrococcus luteus* (yellow growth) before antibiotic assay "rings" were placed on the agar surface. The coloured disks at the end of each spoke of the rungs are impregnated with different antibiotics. Clockwise from the top (arrow) these are: Novobiocin, Penicillin G, Streptomycin (white disk), Tetracycline, Chloramphenicol, Erythromycin, Fusidic acid (green disk) and Methicillin. Clear zones of suppression of bacterial growth around the individual antibiotic disks are evidence of sensitivity to these antibiotics.

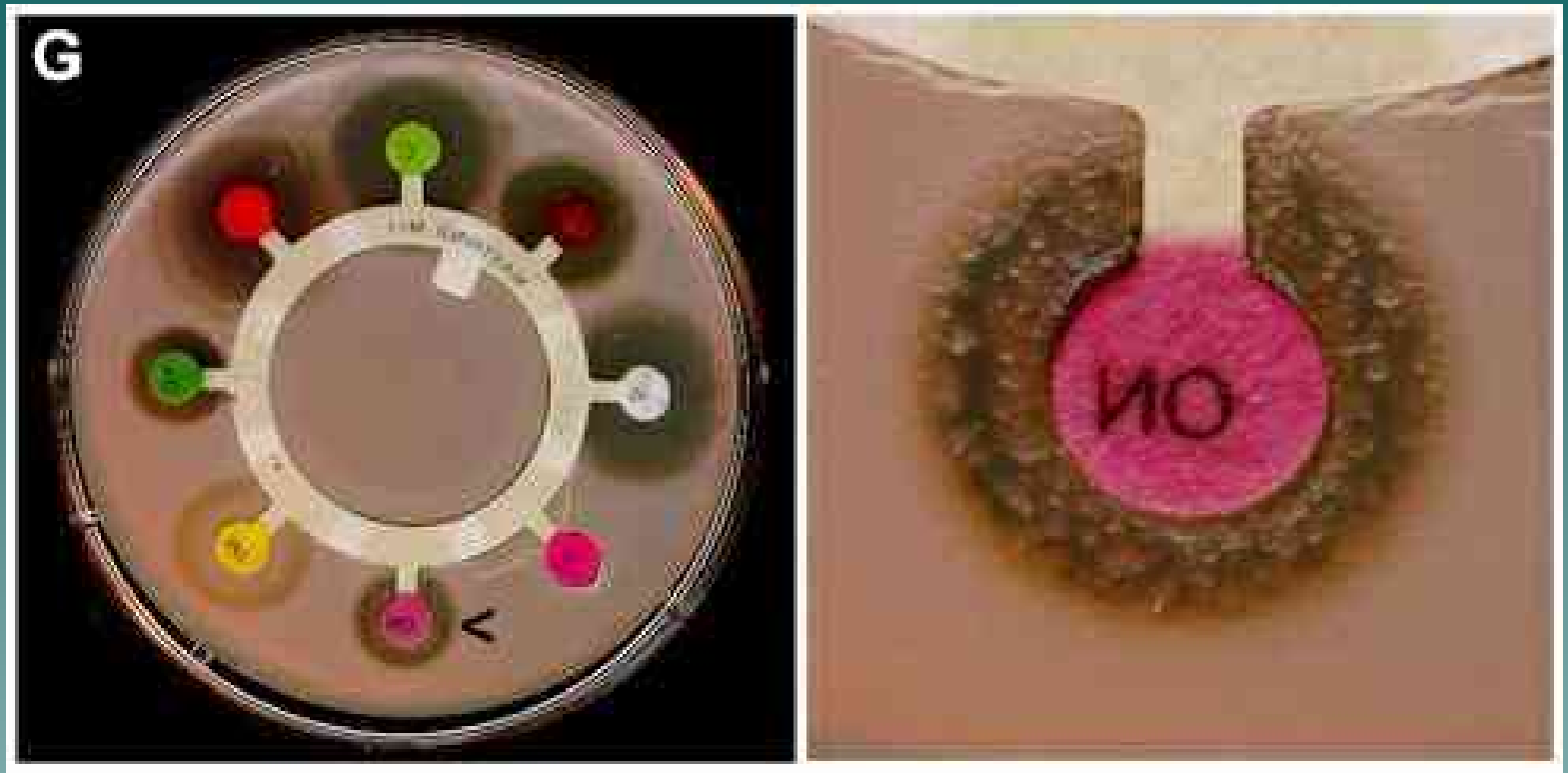



Figure G. Effects of different antibiotics on growth of a *Bacillus* strain. The right-hand image shows a close-up of the novobiocin disk (marked by an arrow on the whole plate). In this case some individual mutant cells in the bacterial population were resistant to the antibiotic and have given rise to small colonies in the zone of inhibition.

➤ One of the most important examples concerns multiple-resistant strains of ***Staphylococcus aureus*** in hospitals.

➤ Some of these strains cause serious nosocomial (hospital-acquired) infections and are resistant to virtually all the useful antibiotics, including **methicillin**, **cephalosporins** and other **beta-lactams** that target peptidoglycan synthesis, the **macrolide antibiotics** such as **erythromycin** and the **aminoglycoside antibiotics** such as **streptomycin** and **neomycin**, all of which target the bacterial ribosome.

➤ The only compound that can be used effectively against these staphylococci is an older antibiotic, **vancomycin**, which has some undesirable effects on humans. Recently, some clinical strains of *S. aureus* have developed resistance to even this compound.

A stylized, layered silhouette of a mountain range in various shades of teal, located in the bottom right corner of the slide.

# Types of antibiotics

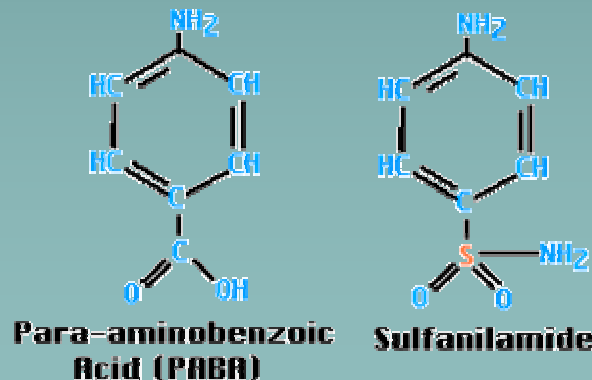
## Sulfa Drugs and Folic Acid Analogs

**Sulfanilamide** was the first antibacterial agent. Many other sulfa drugs (such as sulfamethoxazole) have since come into use.

Both bacteria and their human hosts require folic acid for nucleic acid synthesis (it is converted into purines and thymidine) as well as protein synthesis (precursor of the amino acids methionine and glycine). However, bacteria **synthesize** their folic acid starting with **para-aminobenzoic acid (PABA)**, while we must ingest our folic acid already formed; that is, for us it is a vitamin.

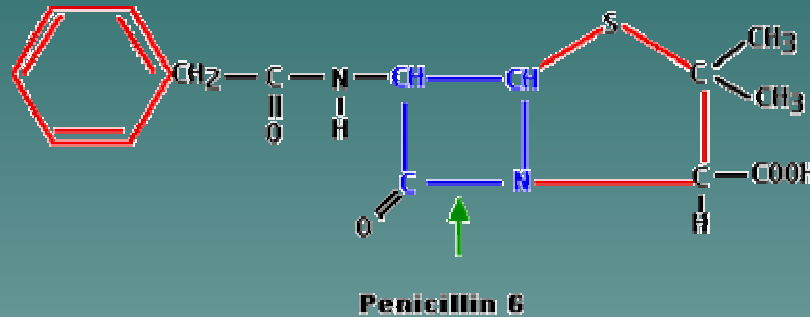
Sulfanilamide, and the other sulfa drugs, are analogs of PABA; they compete with PABA and, when chosen, block the synthesis of folic acid. Mammals ignore PABA and its analogs and thus can tolerate sulfa drugs.

### Sulfa Drugs





The **beta-lactams** get their name from the characteristic ring structure — shown here in blue — that they all share. (The green arrow shows the bond that is broken by the beta-lactamases that are synthesized by many penicillin-resistant bacteria.)



The beta-lactam antibiotics bind to and inhibit enzymes needed for the synthesis of the peptidoglycan wall. While they have little effect on resting bacteria, they are lethal to dividing bacteria as defective walls cannot protect the organism from bursting in hypotonic surroundings.

# Aminoglycosides

Examples are:


streptomycin

kanamycin

neomycin

gentamycin


The 70S bacterial ribosome differs in several ways from the 80S eukaryotic ribosome. The aminoglycosides bind to the **30S** subunit of the bacterial ribosome and interfere with the formation of the initiation complex cause misreading of the mRNA.

A stylized, dark teal silhouette of a mountain range is positioned in the bottom right corner of the slide, extending from the right edge towards the center.

# Tetracyclines

These are natural products derived from soil actinomycetes or their semi-synthetic derivatives. Examples:  
chlortetracycline (trade name = "aureomycin")  
oxytetracycline (trade name = "terramycin")  
doxycycline

Tetracyclines bind to the **30S** subunit of the bacterial ribosome. They prevent the transfer of activated amino acids to the ribosome so protein synthesis is halted.



# Macrolides, Lincosamides, Streptogramins

All these antibiotics bind to the 23S rRNA molecule in the large (**50S**) subunit of the bacterial ribosome where they block the elongation of the growing peptide chain.

Because of their similar action, the development of antibiotic resistance to one usually extends to all the others.



# Fluoroquinolones

Ciprofloxacin (Cipro®), levofloxacin and moxifloxacin are examples. Cipro is the preferred antibiotic for people who have been intentionally exposed to anthrax, although some other antibiotics appear to be equally effective.

The fluoroquinolones block the action of two bacterial topoisomerases — enzymes that relieve the coils that form in DNA when the helix is being opened in preparation for replication or transcription or repair

The topoisomerases in eukaryotes are not affected.

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# Others

## Polypeptides

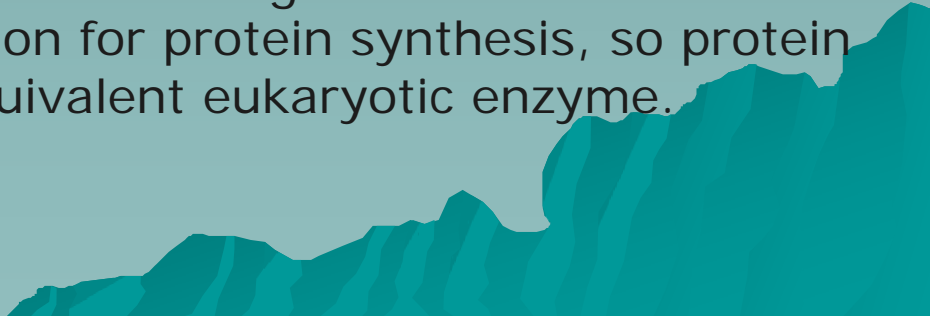
The most common of these are the **polymixins**. They behave as detergents, increasing the permeability of the membranes that encase bacteria and causing the contents of the bacterial cell to leak out.

## Rifampin

This semi-synthetic antibiotic binds to the bacterial **RNA polymerase** and prevents it from carrying out its role in transcription. Its affinity for the equivalent eukaryotic enzyme is much lower. Rifampin is also known as rifampicin.

## Mupirocin

This antibiotic blocks the action of the bacterial **isoleucine tRNA synthetase**, the enzyme responsible for attaching the amino acid isoleucine (Ile) to its tRNA in preparation for protein synthesis, so protein synthesis is inhibited. It spares the equivalent eukaryotic enzyme.

A stylized, dark teal silhouette of a mountain range is positioned in the bottom right corner of the slide, partially overlapping the text of the Mupirocin section.

## Cycloserine


Cycloserine inhibits synthesis of the bacterial cell wall but by a different mechanism than the beta-lactam antibiotics discussed above.

Cycloserine is an analog of **D-alanine** and blocks the incorporation of D-alanine into the peptide bridges in the bacterial cell wall ([look back](#)). It is derived from an actinomycete.

## Aminocyclitols

These products of another [actinomycete](#) achieve their effect by interfering with the **30S** subunit of the bacterial ribosome.

**Spectinomycin** (trade name = Trobicin) is an example. It is particularly effective against the [gonococcus](#), the bacterium that causes the sexually-transmitted disease (STD) gonorrhea.

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## Glycopeptides


Glycopeptides also interfere with the synthesis of the bacterial cell wall but by a different mechanism than the beta-lactams.

**Vancomycin** is a widely-used glycopeptide in the U.S. It binds to the **D-alanines** on the precursors of the peptidoglycan cross bridges preventing their cross-linking ([look back](#)). It has become the antibiotic of last resort as resistance to the other antibiotics has become more and more common.

## Oxazolidinones

The first of these new antibiotics, **linezolid** (trade name = **Zyvox**), was approved by the U.S. Food and Drug Administration on 19 April 2000. It is effective against many [gram-positive bacteria](#) that have developed resistance to the older antibiotics.

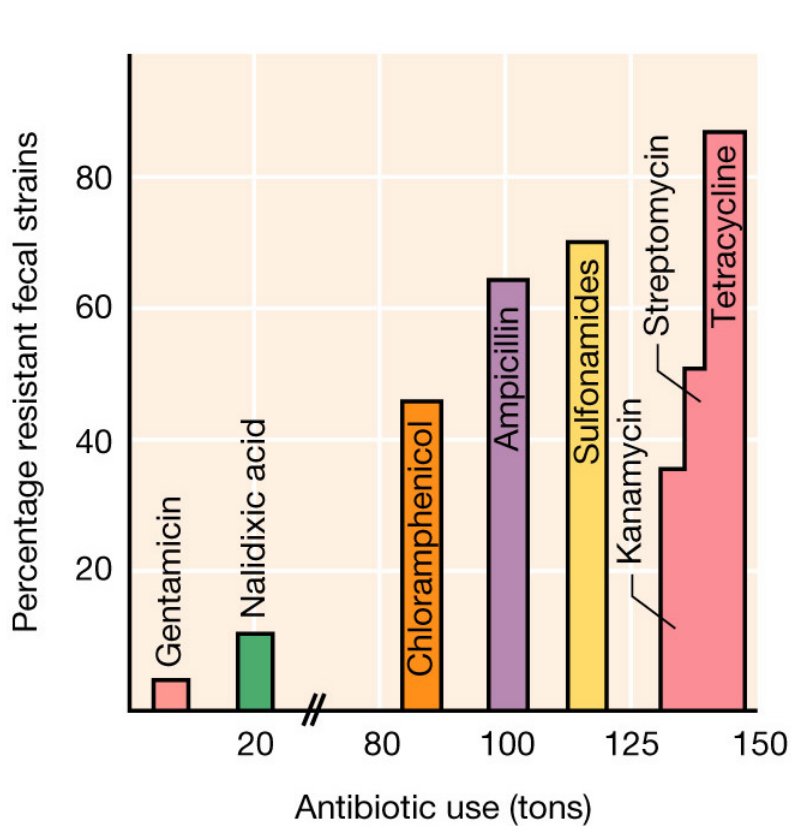
Zyvox attacks a previously-unexploited chink in the bacterium's armor: the proper assembly of the complex of **fMet-tRNA** with the messenger RNA (**mRNA**) and the two ribosomal subunits. Eukaryotes do not begin translation with [fMet](#).

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# Industrial evolution of penicillin production

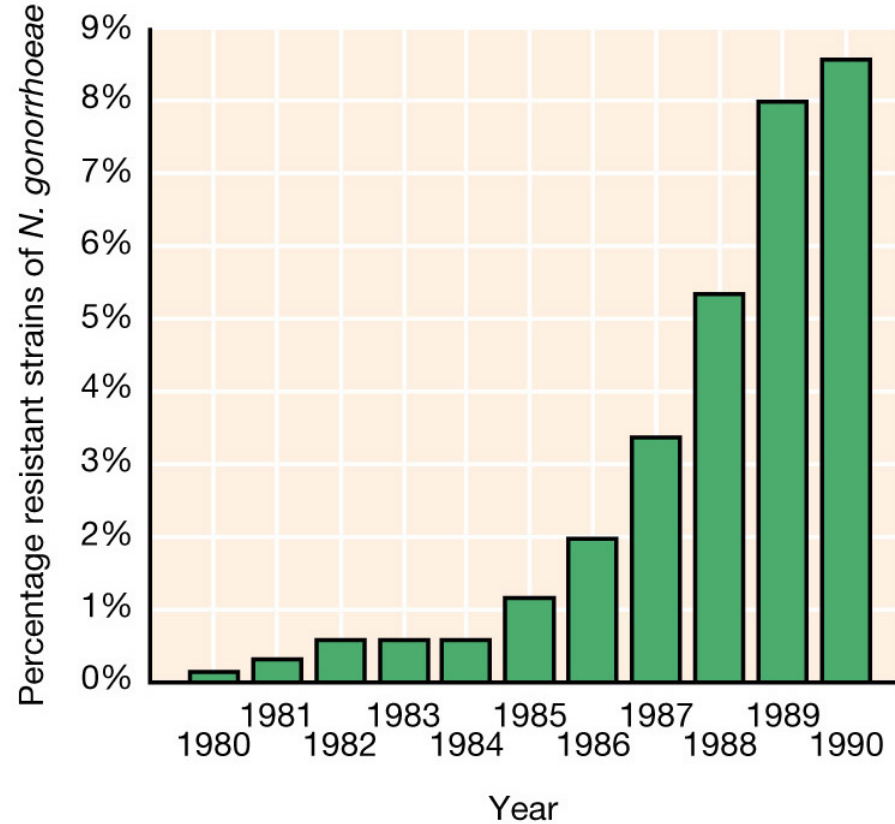
Date	Yield (units/mL)	Development
1929	2-20	Wild-type ( <i>P. notatum</i> )
1941	40-80	Better WT
1943	80-100	New WT ( <i>P. chrysogenum</i> )
1944	100-200	Colony selection
1944	300-500	X-irradiation
1945	800-1000	UV-irradiation
1949	1500-2000	Chemical mutagenesis
1951	2400	Chemical mutagenesis
1953	2700	Strain selection
1960	5000	Strain selection
1970	10000	Strain selection



(a)

The emergence of antimicrobial drug-resistant bacteria.

(a) Relationship between antibiotic use and the percentage of bacteria isolated from diarrheal patients resistant to the antibiotic. Those agents that have been used in the largest amounts, as indicated by the amount produced commercially, are those for which drug-resistant strains are most frequent.

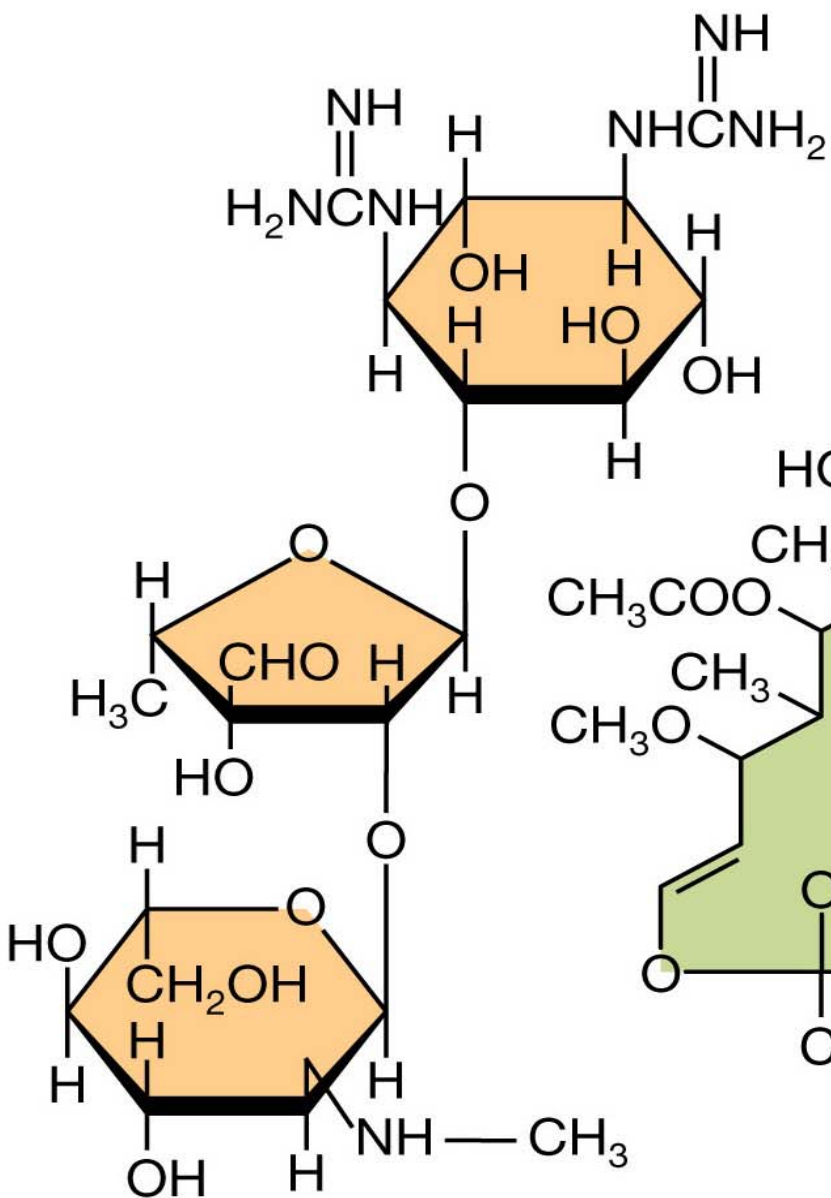


(b)

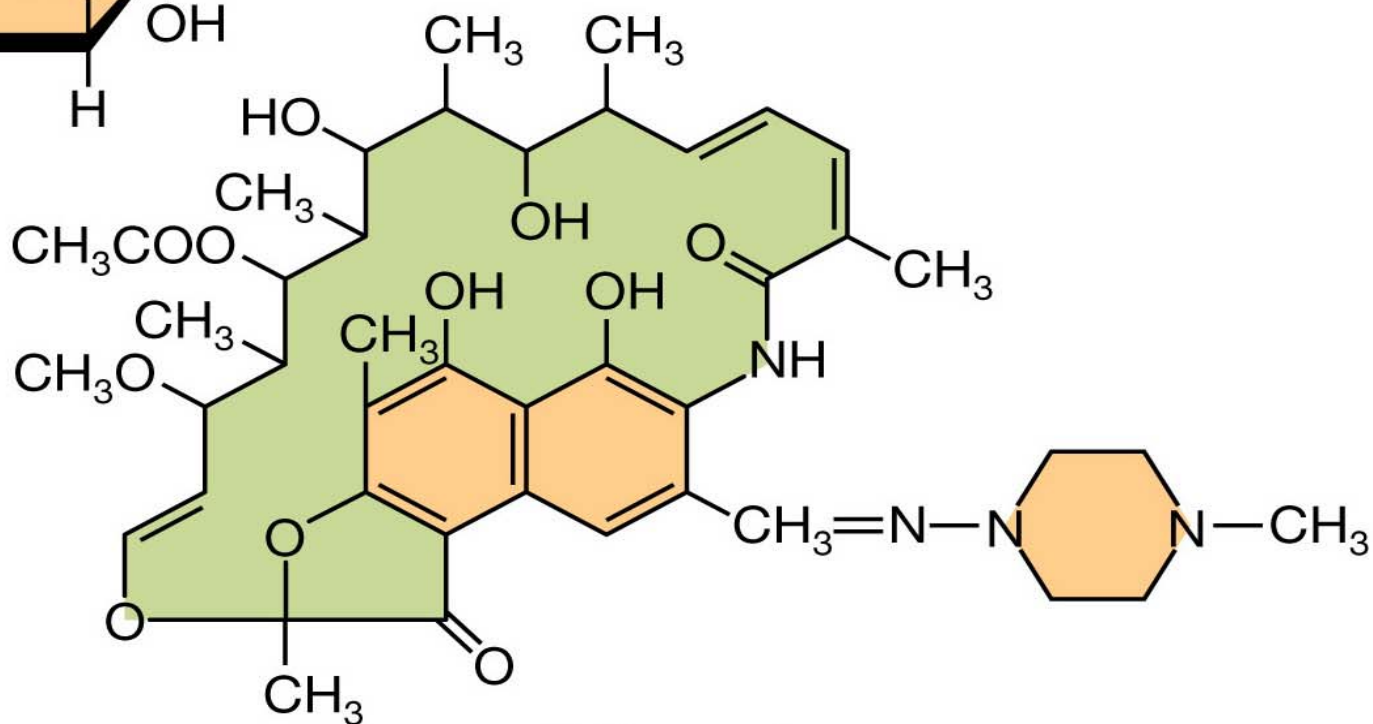
(b) Percentage of reported cases of gonorrhea caused by drug-resistant strains. The actual number of reported drug-resistant cases in 1985 was 9000. This number rose to 59,000 in 1990. Greater than 95% of the reported drug-resistant cases are due to penicillinase-producing strains of *Neisseria gonorrhoeae*. Since 1990, penicillin has not been recommended for treatment of gonorrhea because of emerging drug resistance. (Source: Centers for Disease Control, Atlanta, GA).

## Representative structure

Classification of antibacterial chemotherapeutic agents according to chemical structure. A representative example is shown for each group.

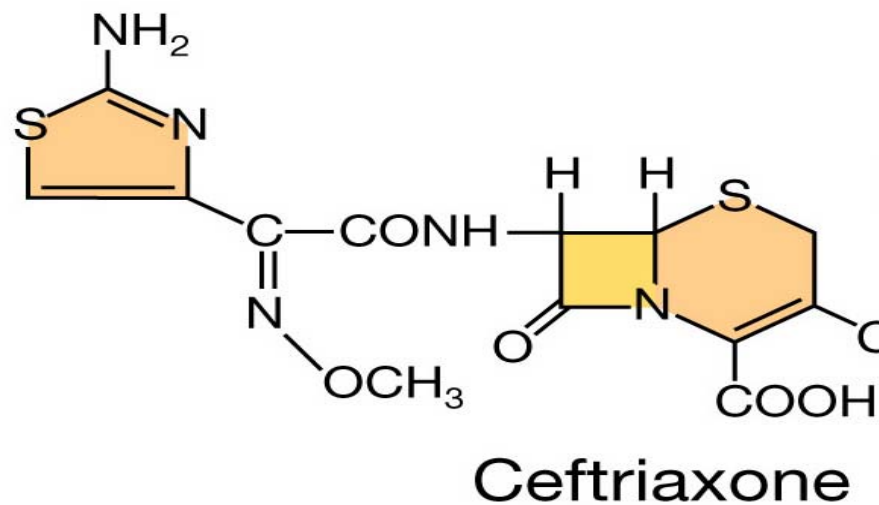


# Streptomycin

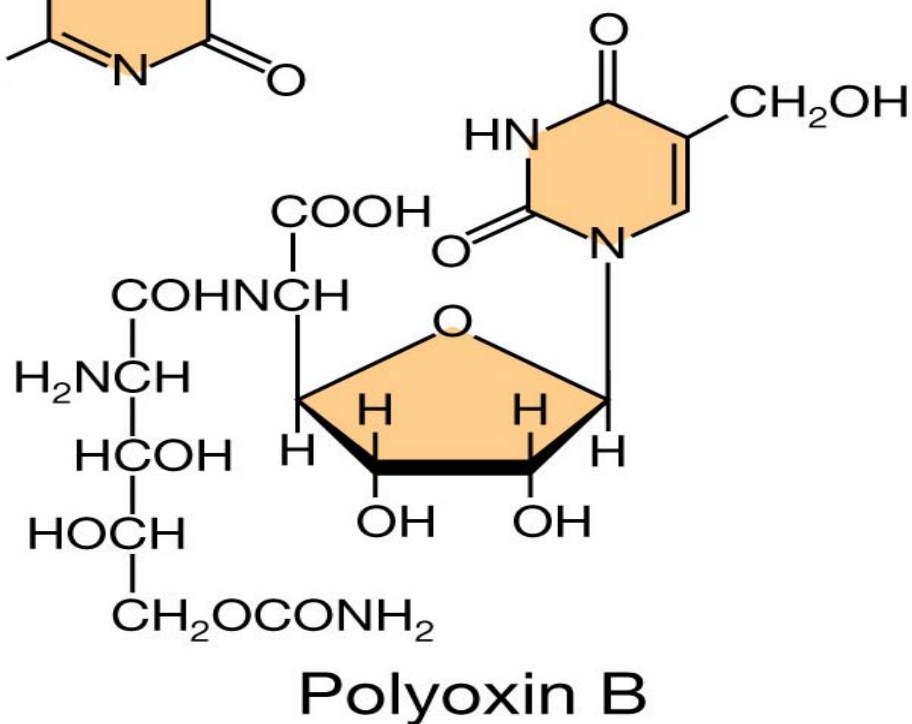
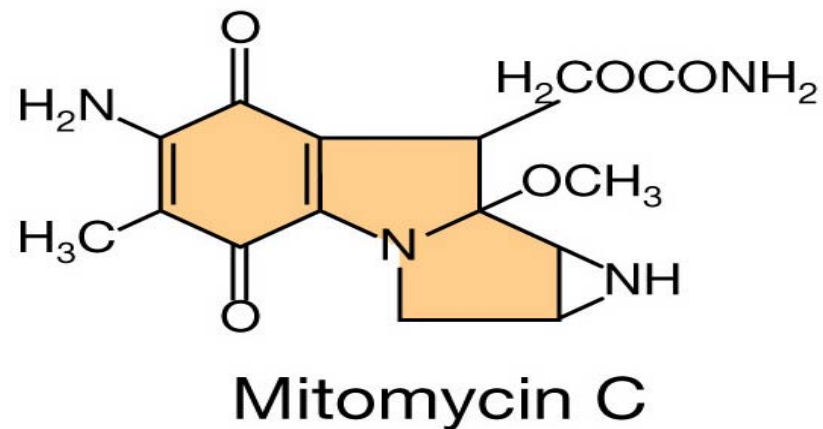


# Rifampin

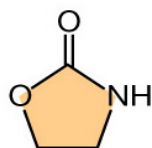
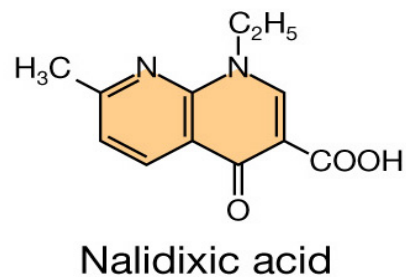
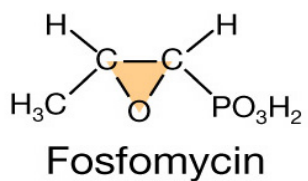
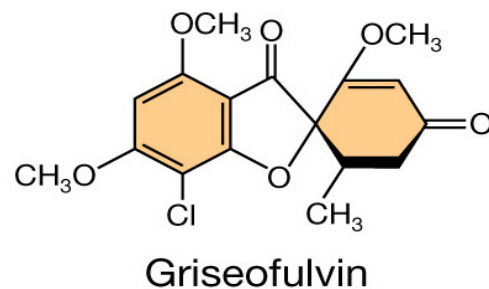
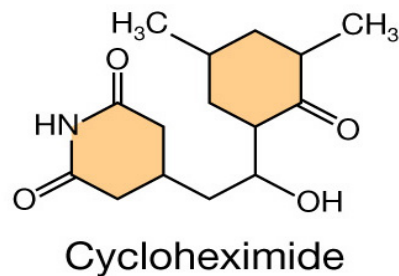
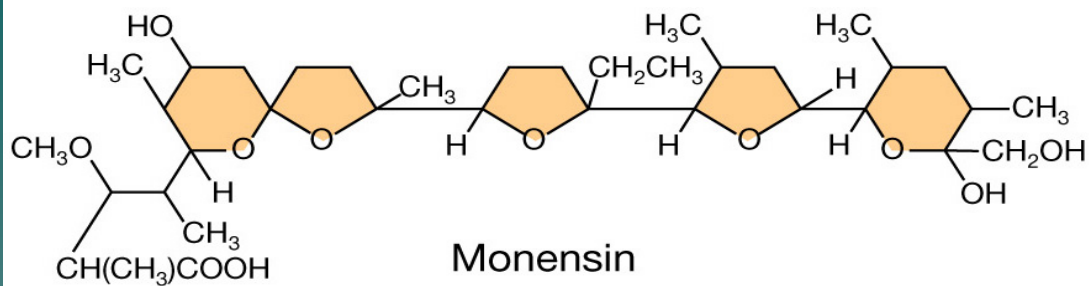
# Representative structure



Classification of antibacterial chemotherapeutic agents according to chemical structure. A representative example is shown for each group.

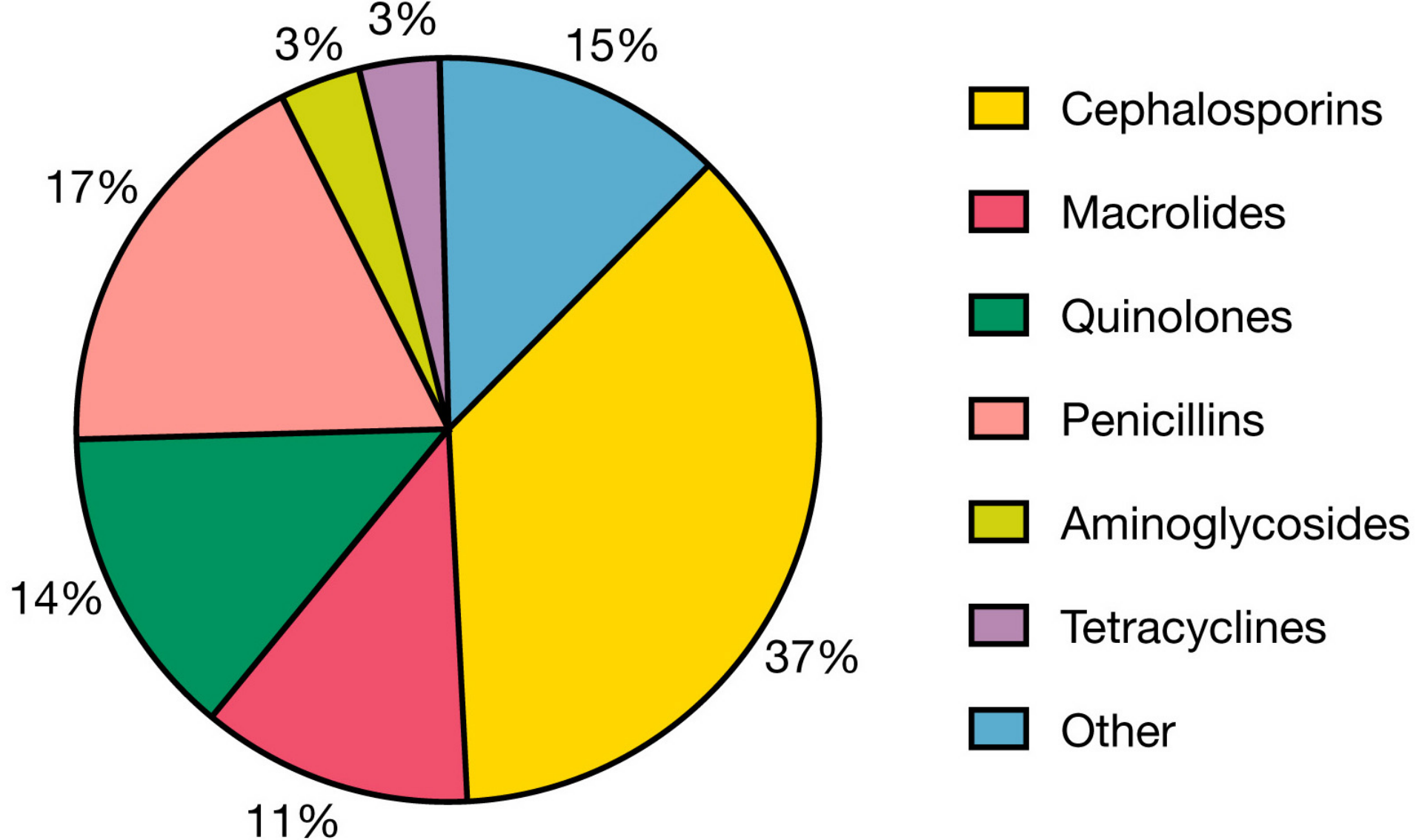


# Representative structure



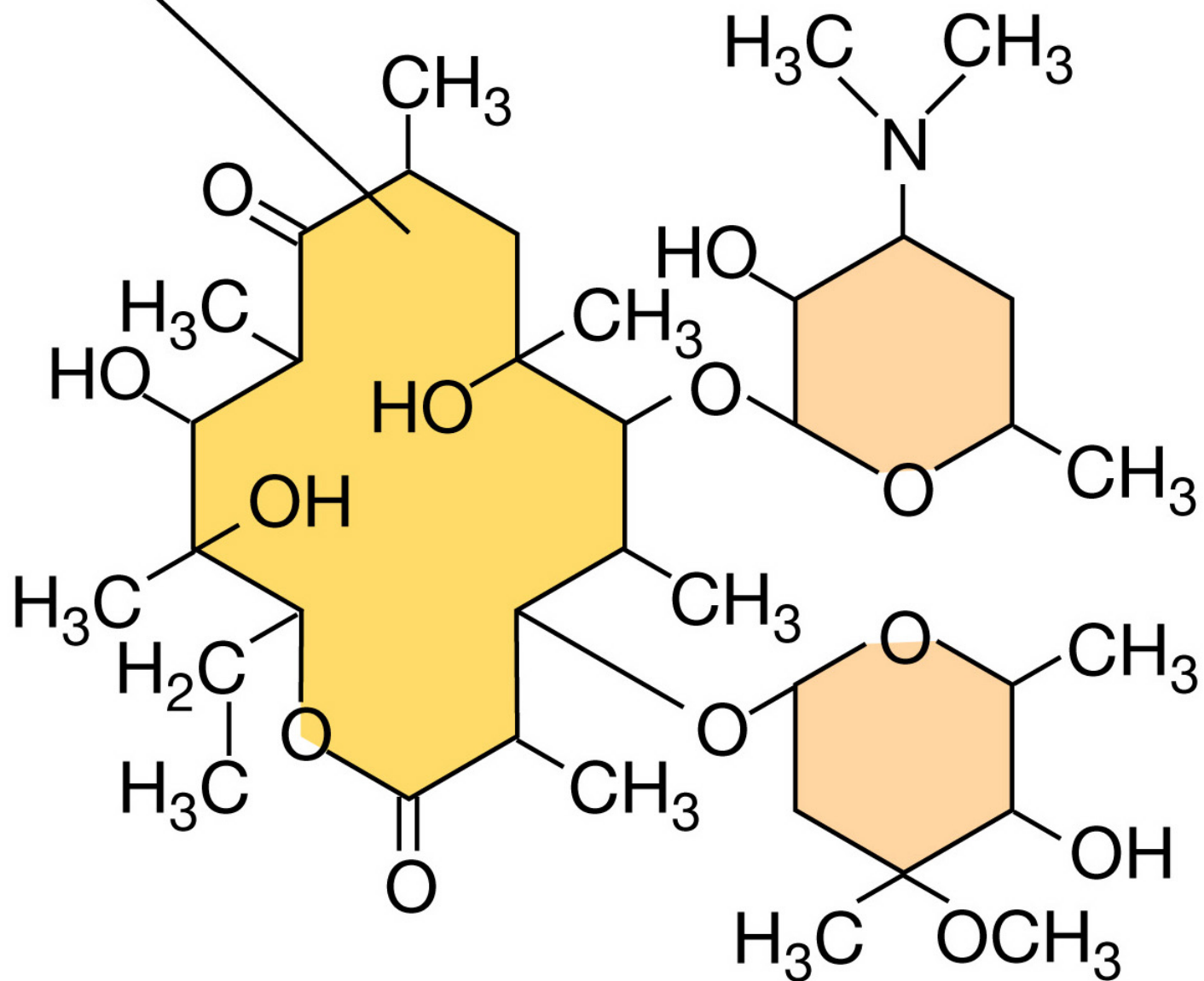
2-Oxazolidinone

Antibacterial chemotherapeutic agents according to chemical structure.



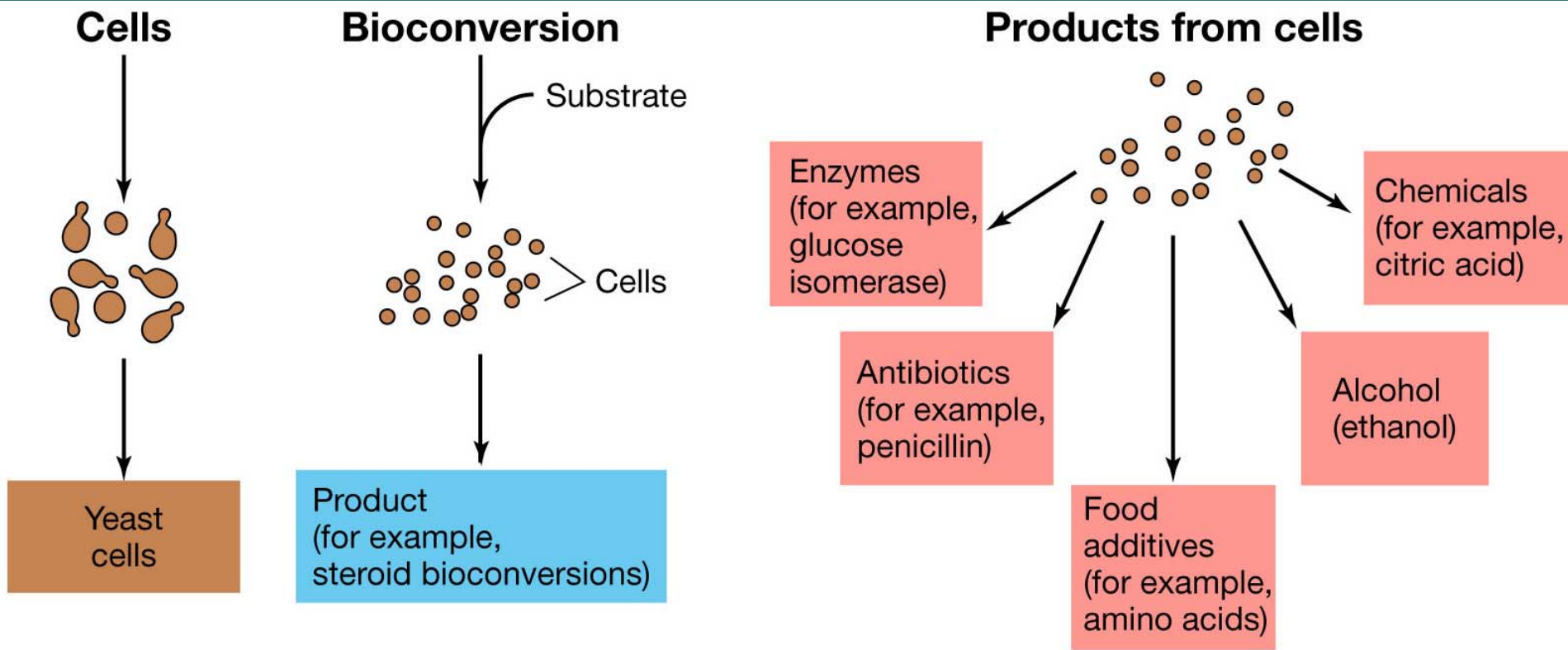
Annual worldwide production and use of antibiotics. Each year more than 500 metric tons of chemotherapeutic agents are manufactured.

Macrolide  
ring



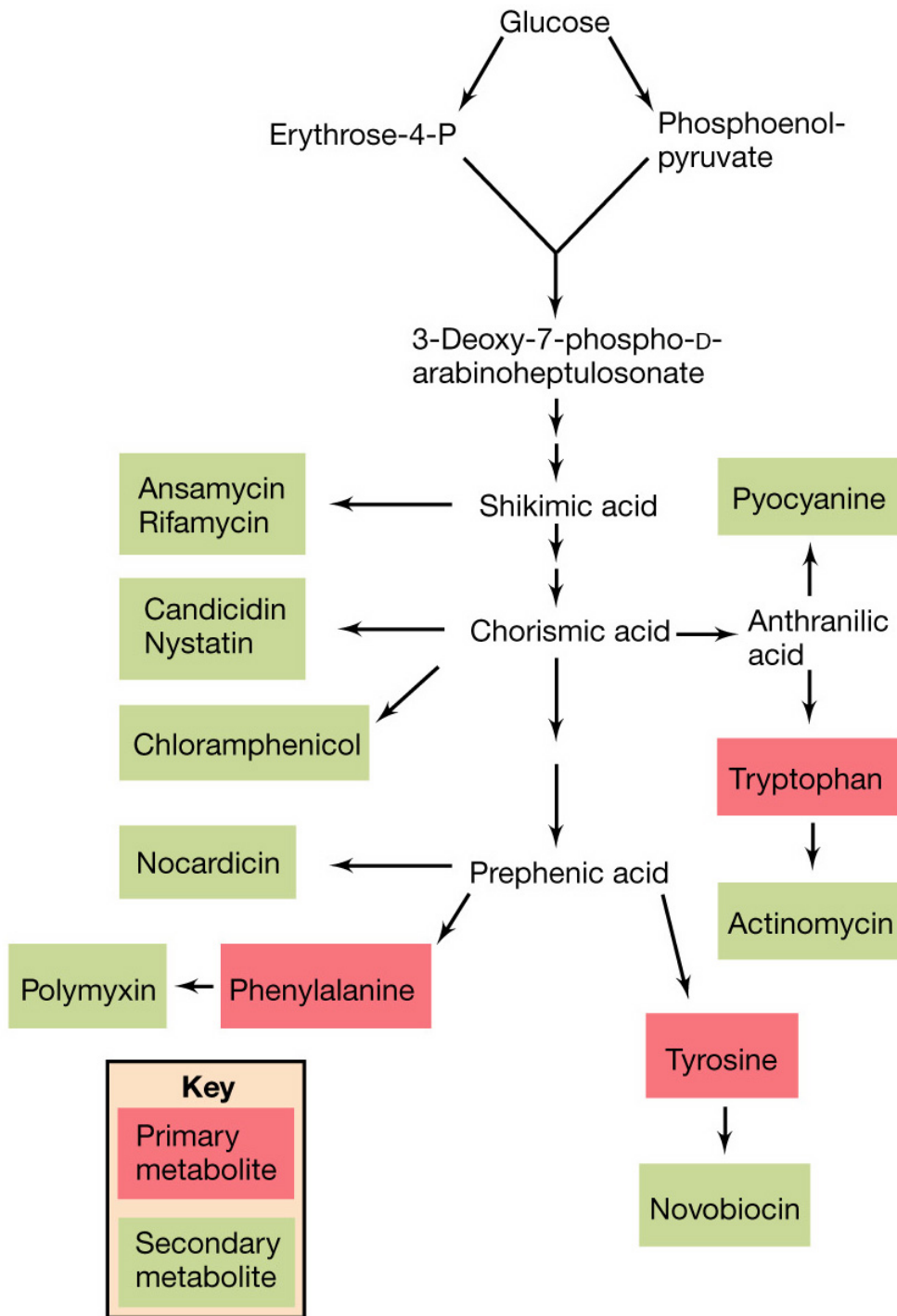
Structure of erythromycin, a macrolide antibiotic.





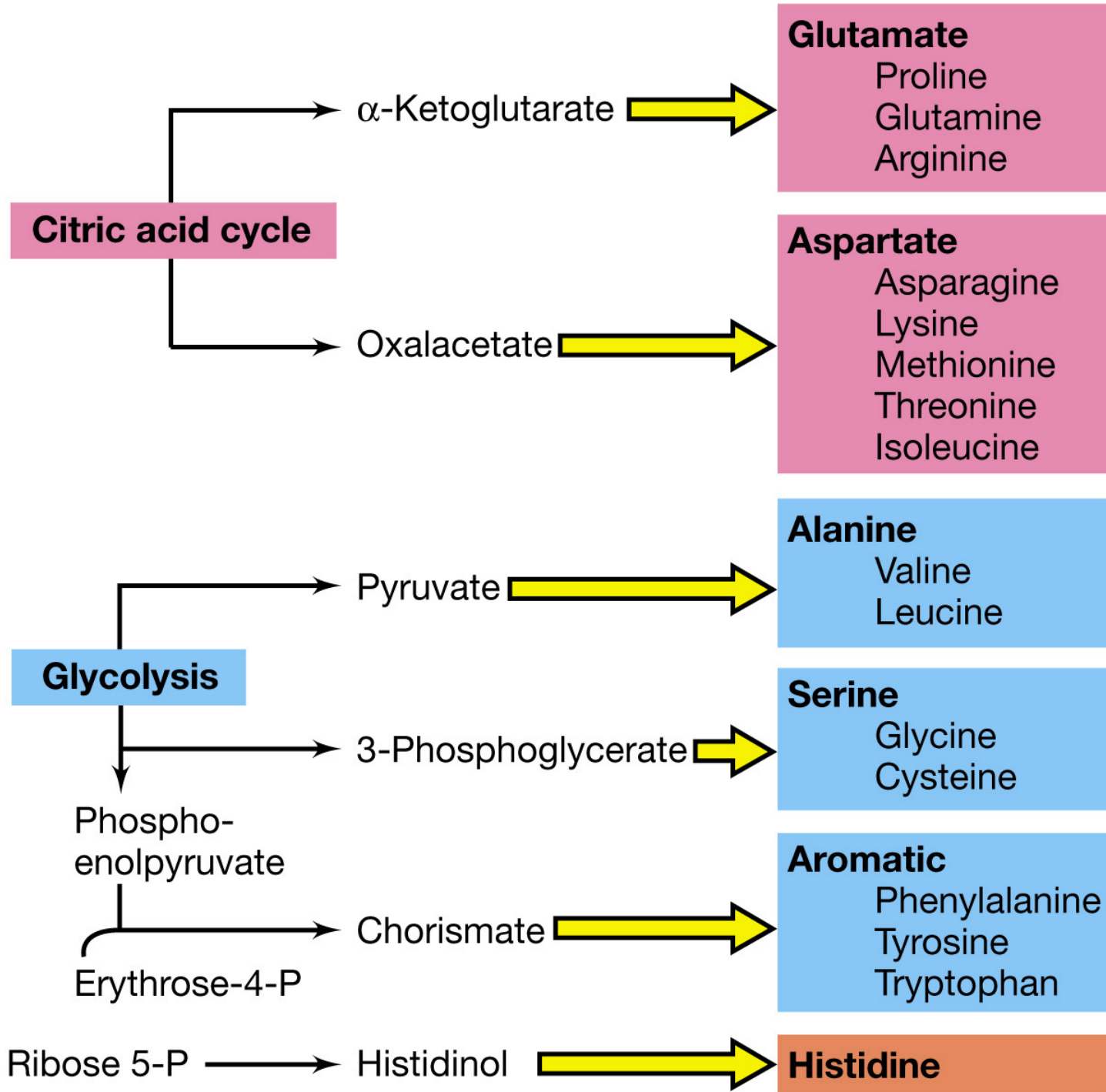
Products of industrial microbiology/biocatalysis. The products may be the cells themselves or products made from cells. In the case of bioconversion, cells are used to chemically convert a specific substance from one form to another.





Relationship of the primary metabolic pathway for the synthesis of aromatic amino acids and formation of a variety of secondary metabolite antibiotics containing aromatic rings.

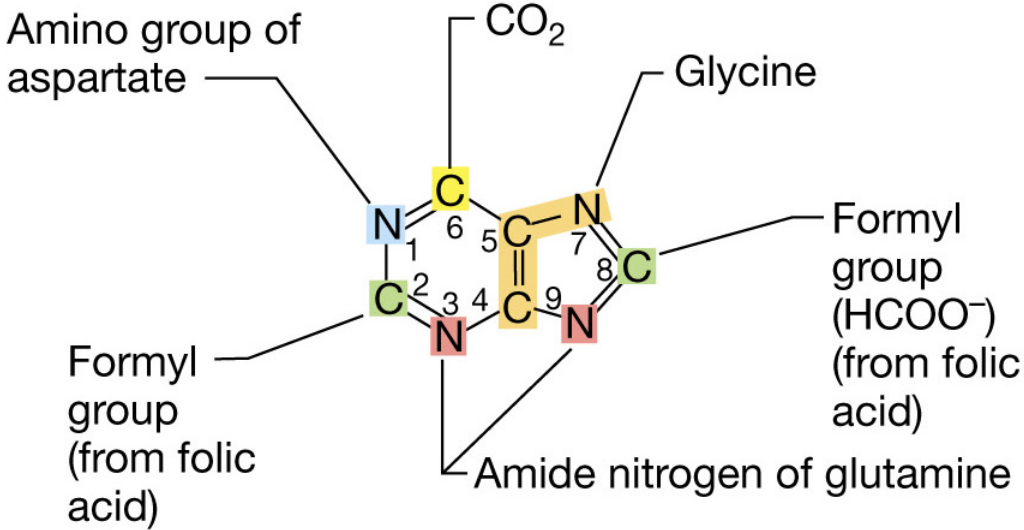
This is a composite scheme of processes occurring in a variety of microorganisms: No one organism produces all these secondary metabolites, and many individual steps exist between amino acid and antibiotic in all cases.



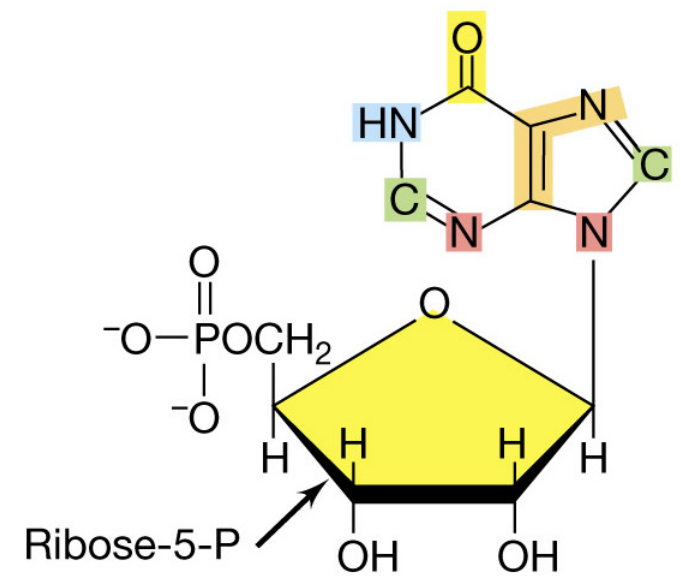
Amino acid families.

Note how the carbon skeletons for most amino acids are derived from either the citric acid cycle or from glycolysis.

Synthesis of the various amino acids in a family frequently requires many separate enzymatically catalyzed steps starting from the parent amino acid (shown in bold).

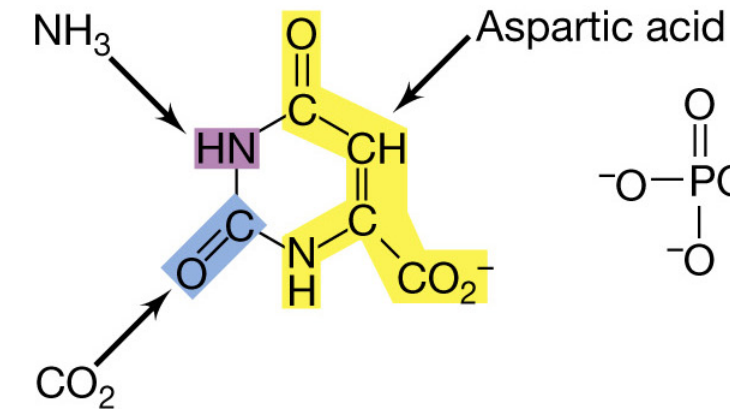


(a)

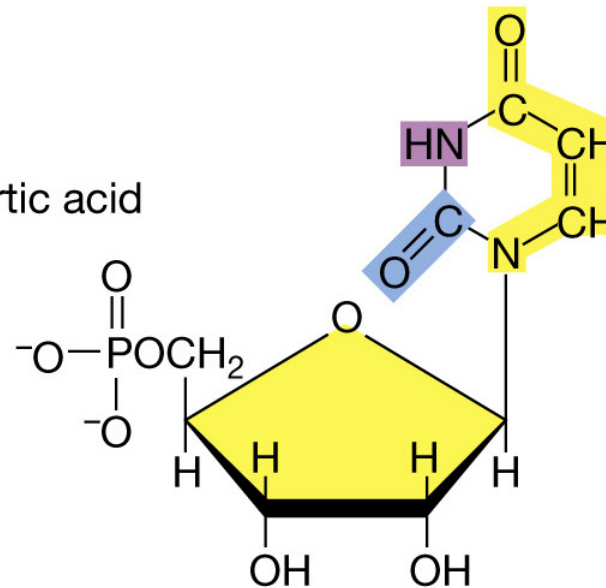


(b)

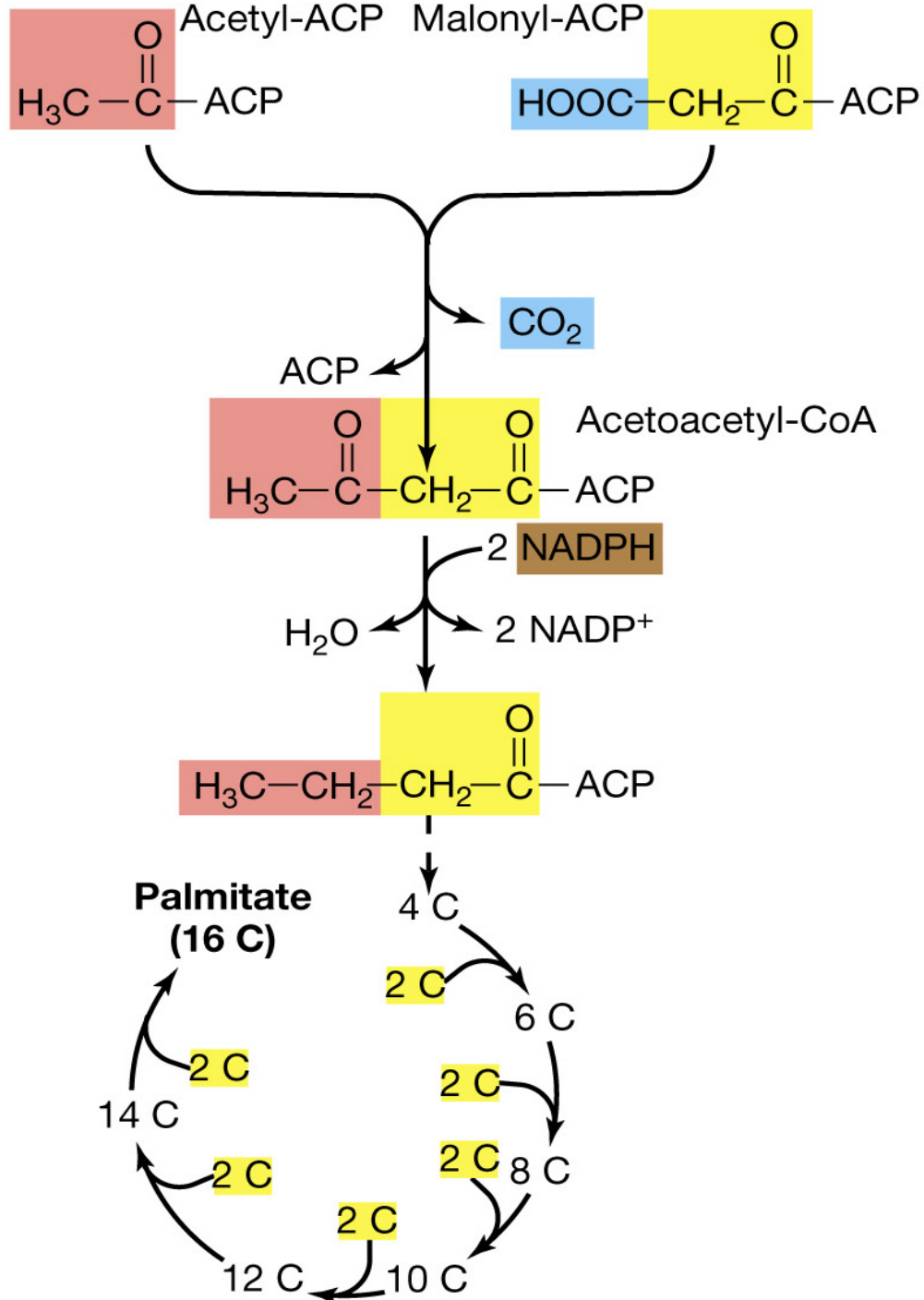
Biosynthesis of purines and pyrimidines. (a) The precursors of the purine skeleton. (b) Inosinic acid, the precursor of all purine nucleotides. (c) The precursors of the pyrimidine skeleton, orotic acid. (d) Uridylate, the precursor of all pyrimidine nucleotides. Uridylate is formed from orotate following a decarboxylation and the addition of ribose-5-phosphate.



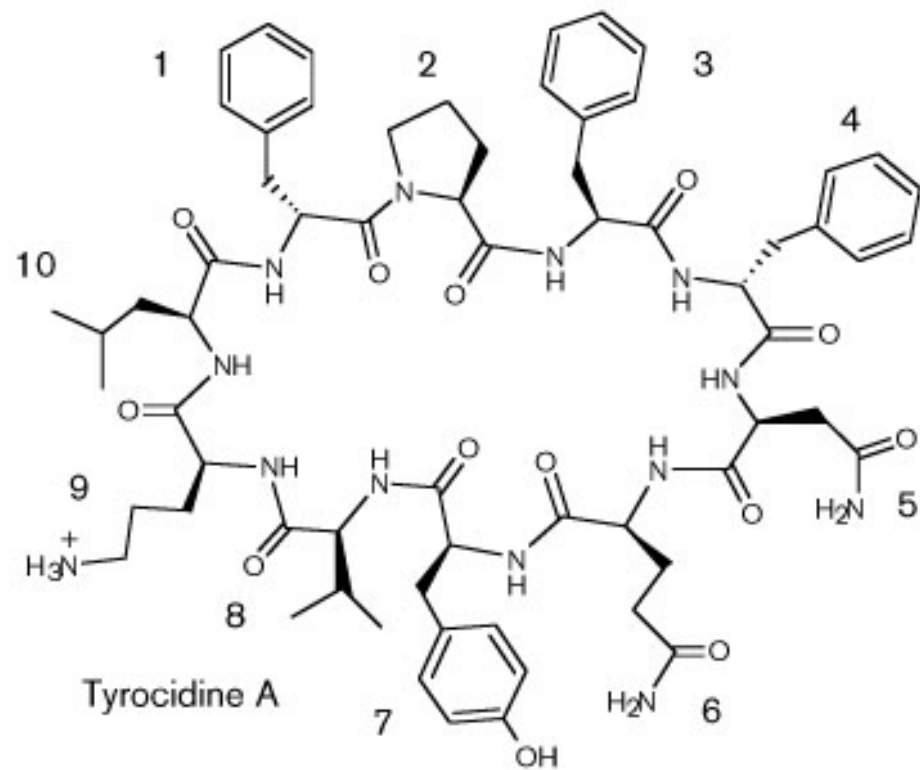
(c)



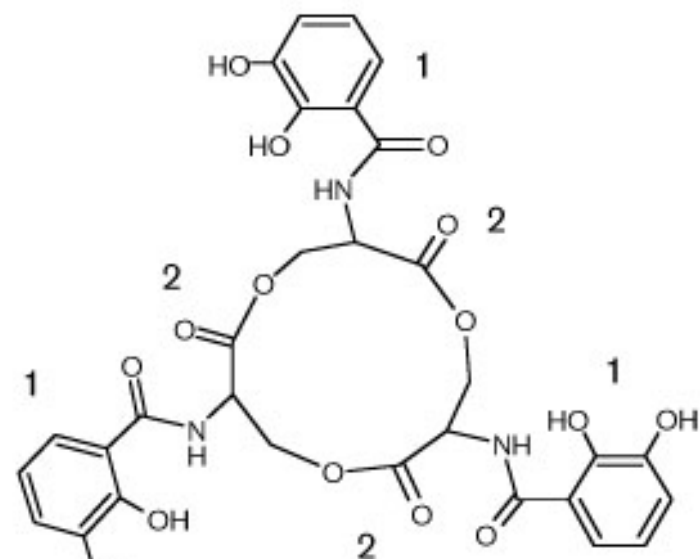
(d)



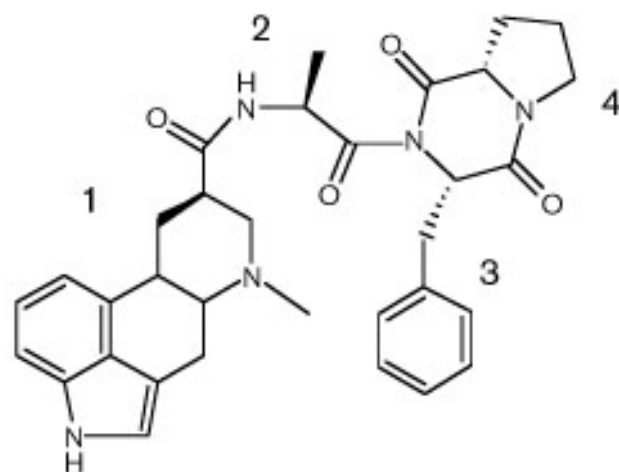
The biosynthesis of fatty acids; shown is the biosynthesis of the C16 fatty acid, palmitate. The condensation of acetyl-ACP and malonyl-ACP forms acetoacetyl-CoA. Each successive addition of an acetyl unit comes from malonyl-CoA.



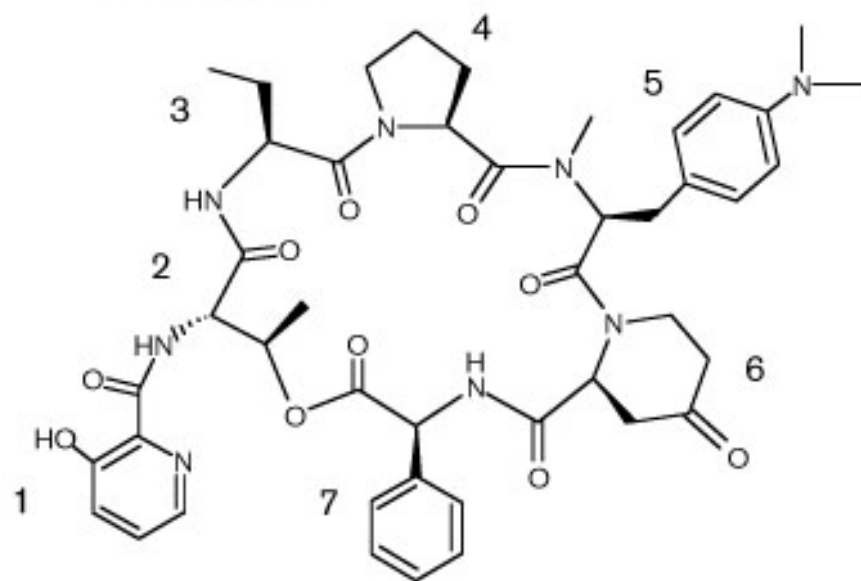
Tyrocidine A



Enterobactin



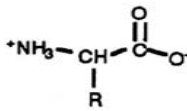
Ergotamine



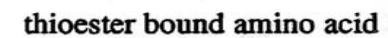
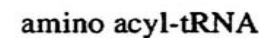
Pristinamycin I<sub>A</sub>

◆ ¿how is the biosynthesis of those antibiotics?

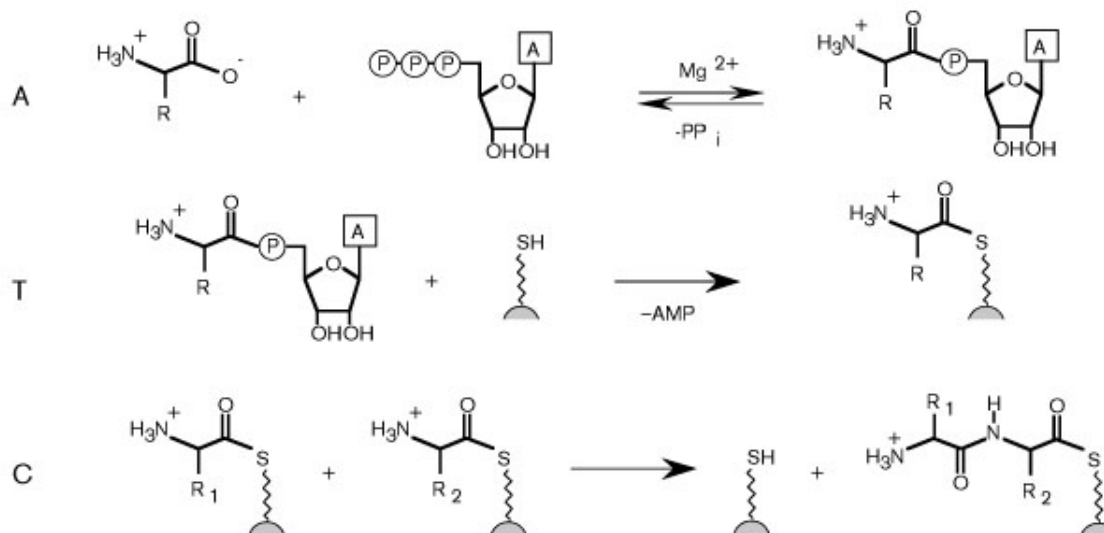
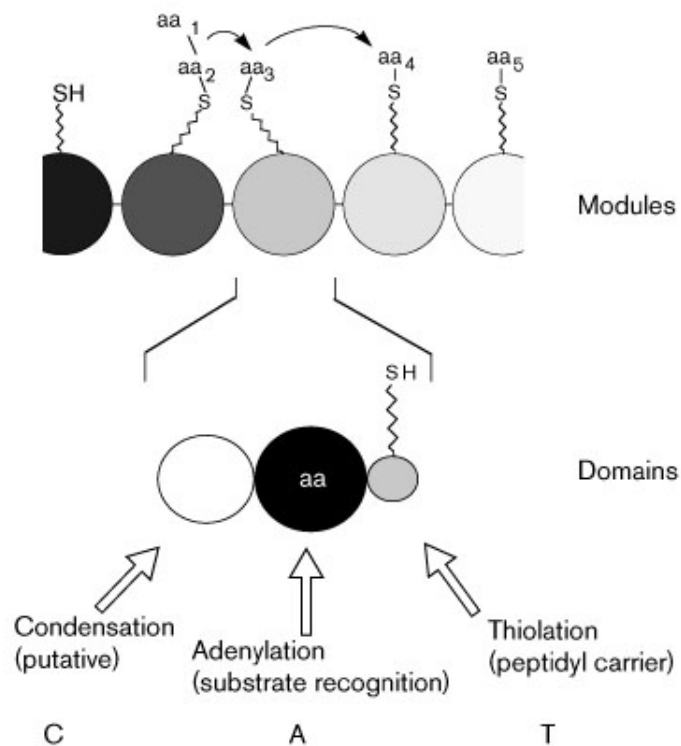
- ◆ Non ribosomal peptide synthetase
- ◆ Polyketide synthase



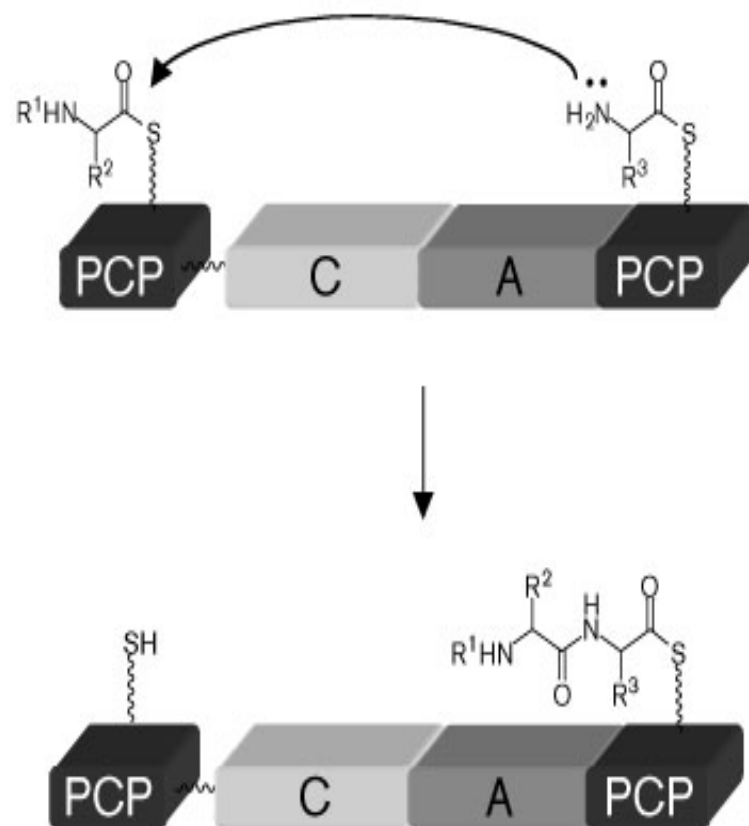
**enzyme** tRNA synthetase or  
multifunctional peptide synthetase



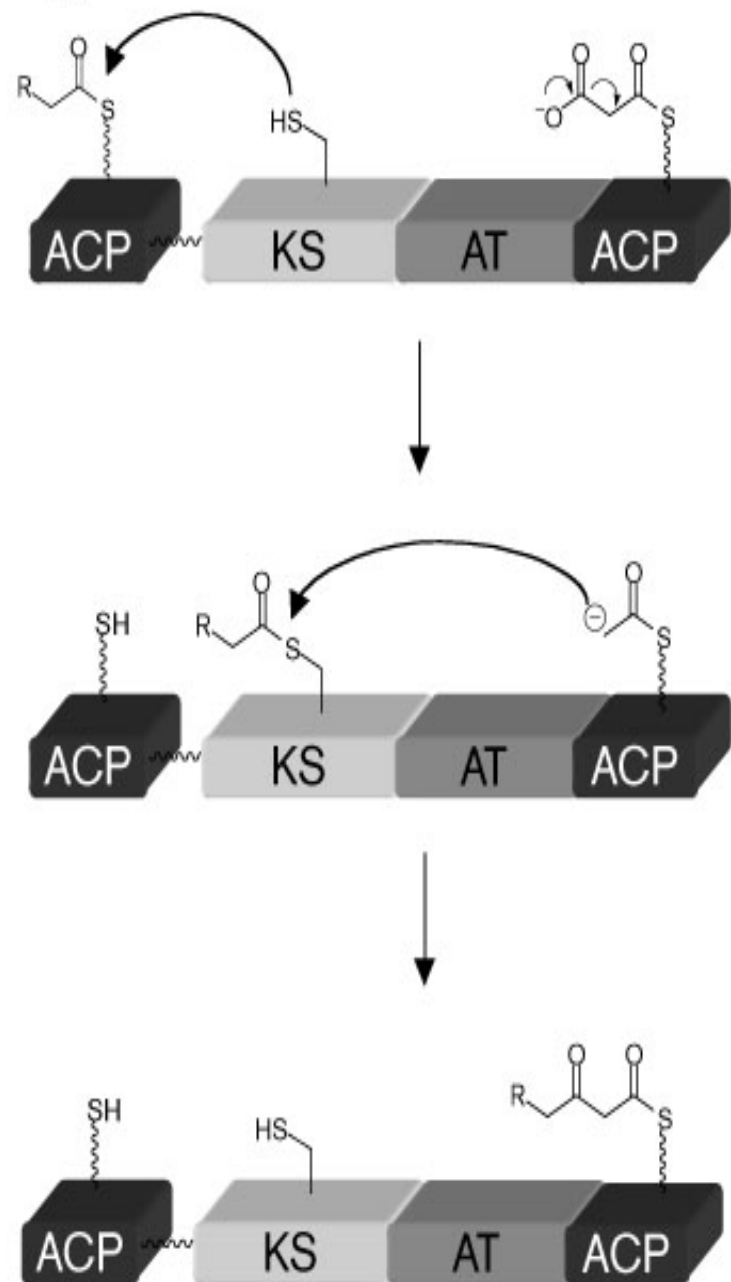




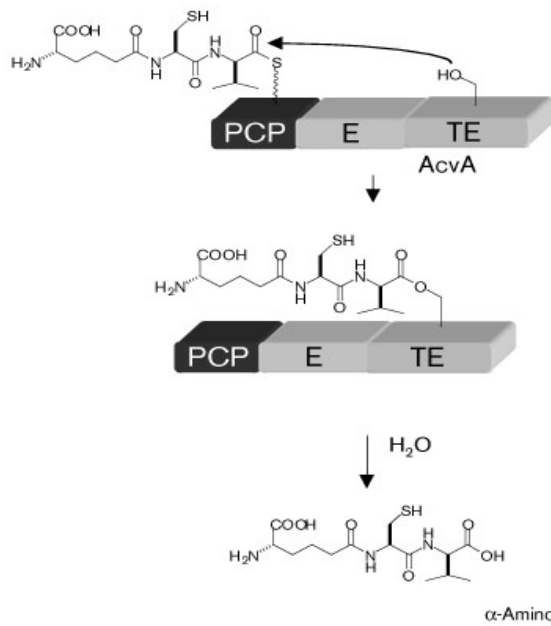
(a) Core NRPS domains



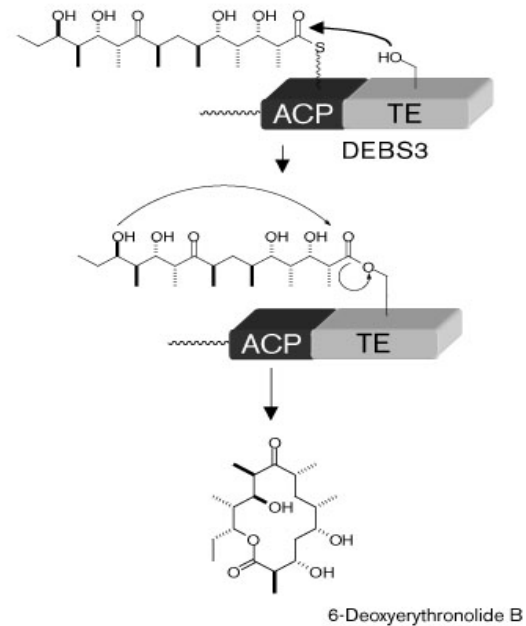
(b) Core PKS domains



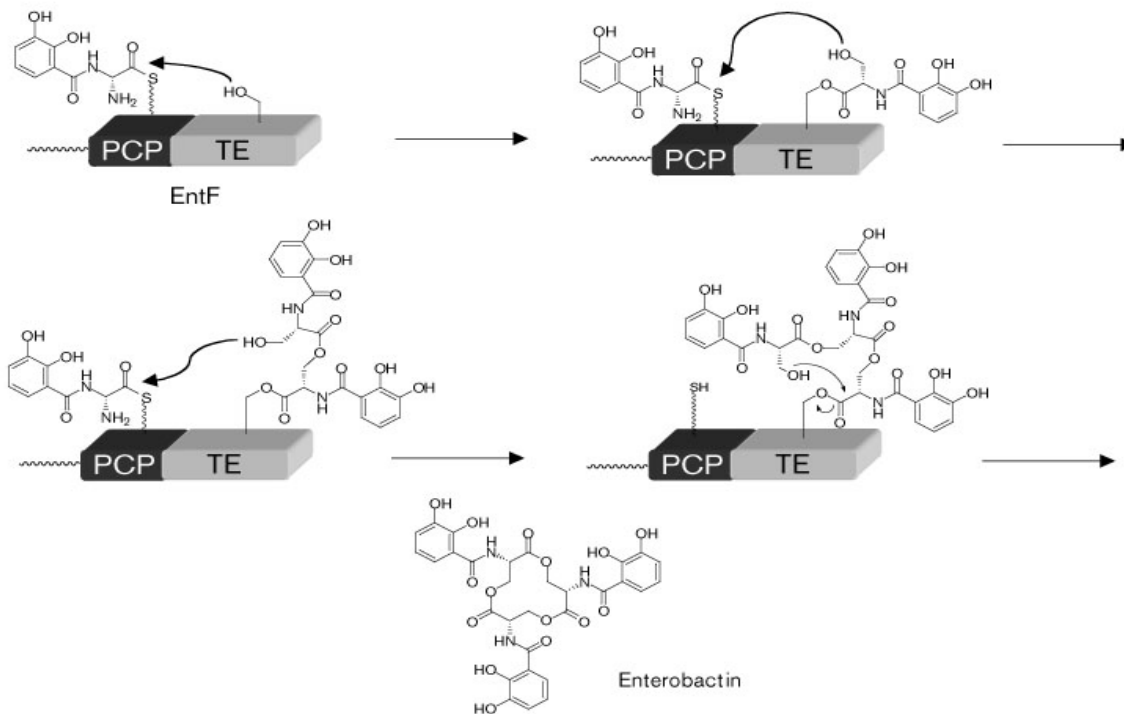
(a)

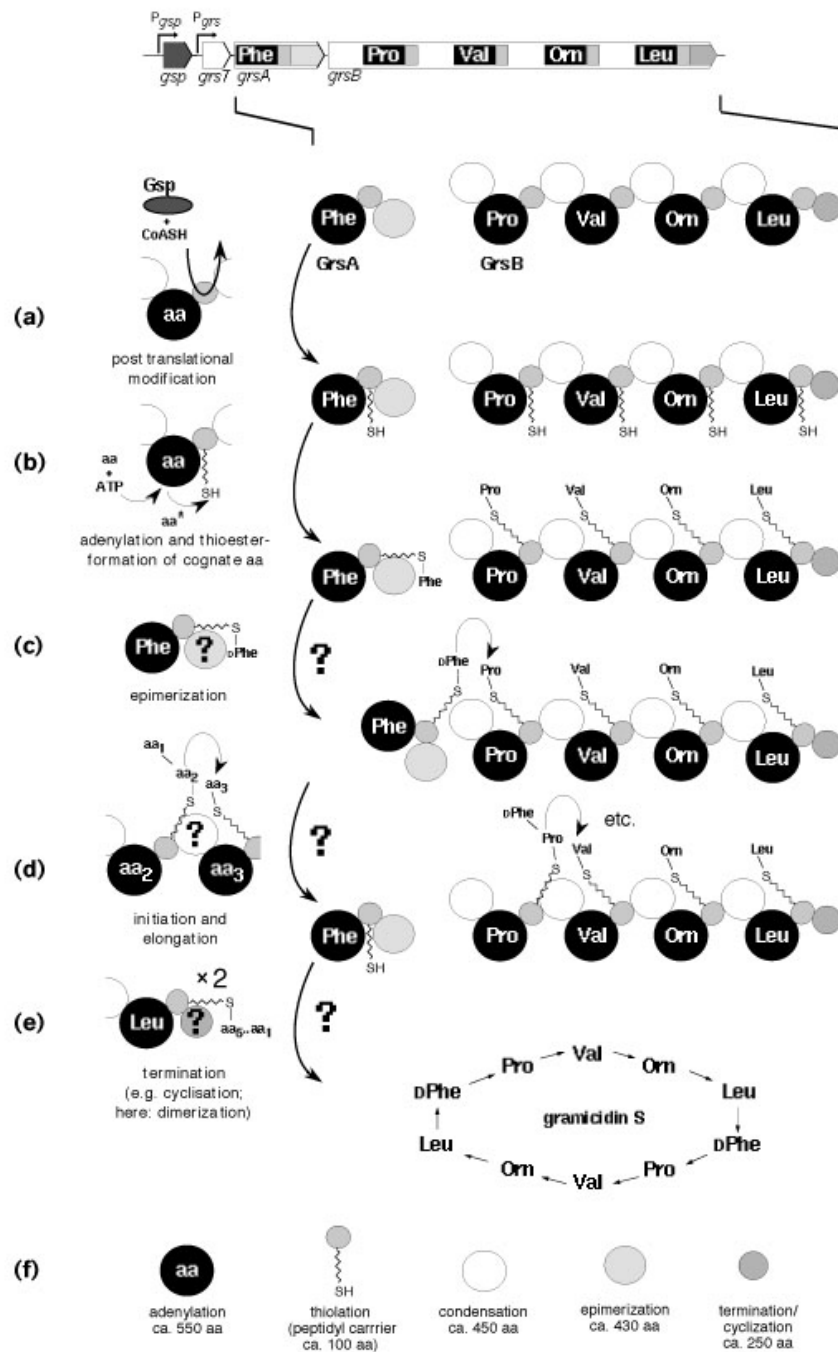


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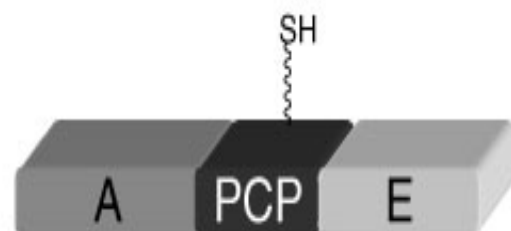


(c)

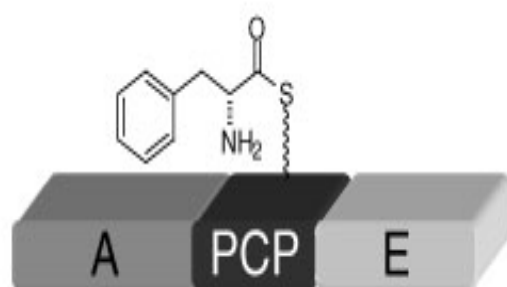
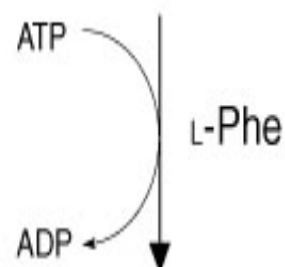




**(a) NRPS starter unit**



GrsA  
(gramicidin S)

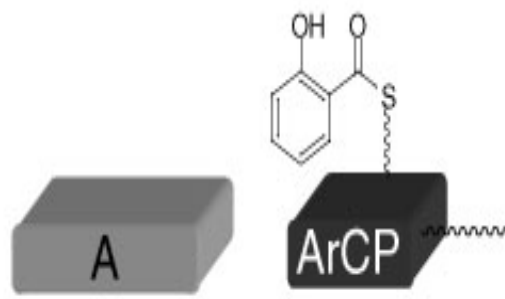
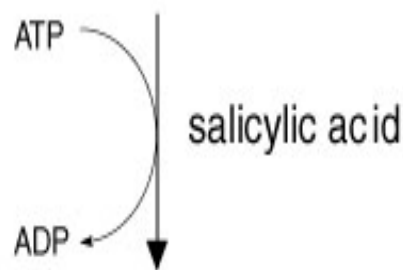


GrsA

**(b) Aryl-N-capped NRPS starter unit**



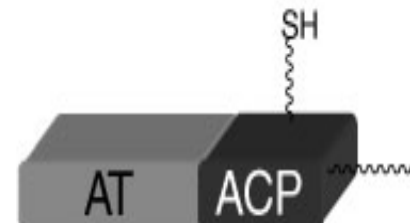
YbtE  
HMWP2  
(yersiniabactin)



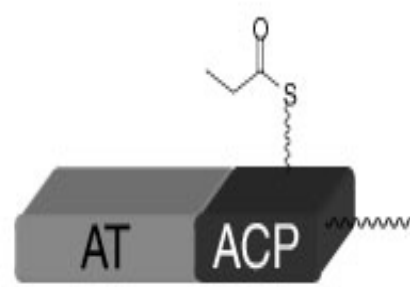
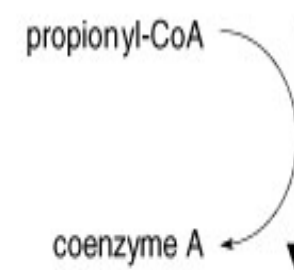
YbtE

HMWP2

**(c) PKS starter unit**

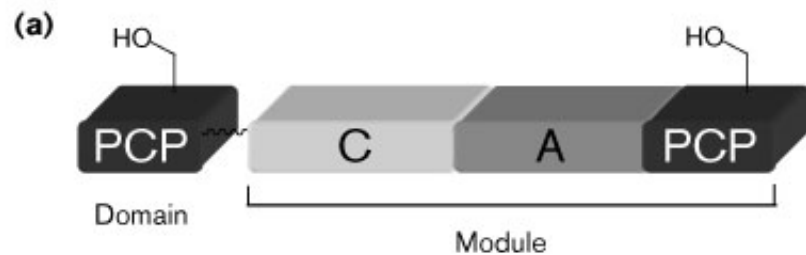


DEBS1  
(6-deoxyerythronolide B)

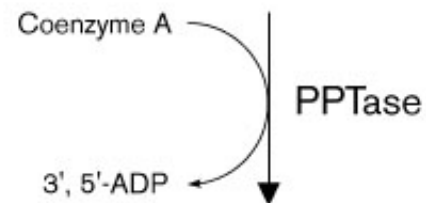
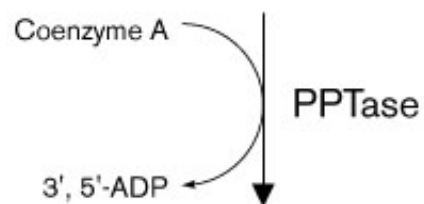


DEBS1

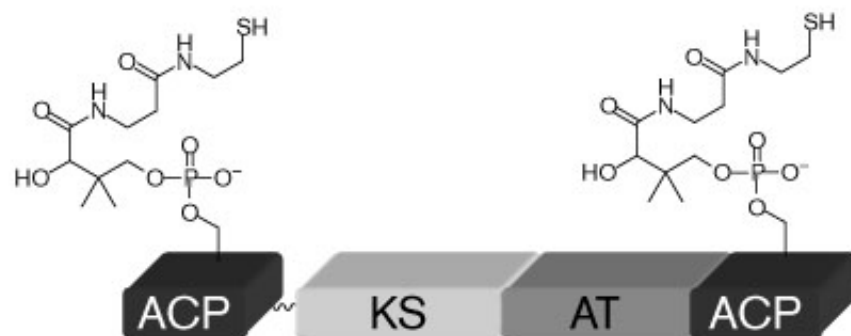
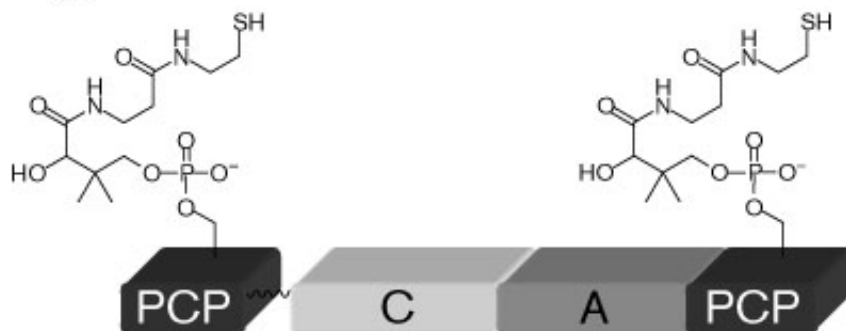
Core NRPS domains



Core PKS domains



(b)



# Polyketide synthase

**PKSs Tipo II** están compuestas de enzimas discretas y monofuncionales (KS, ACP etc.), ejemplo PKSs aromaticos.

**PKSs Tipo I** las enzimas están covalentemente unidas.

PKSs Tipo I están compuestas de una o mas polipéptidos multienzimas

**PKSs Modulares tipo I** tienen un set de enzimas (KS-AT-dominios de reducción -ACP) para cada ciclo de extensión de la cadena.

**Iterative type I PKSs** have one set of enzymes that is used repeatedly.

These systems make simpler polyketides than modular PKSs.

KS – AT - (DH) - ACP

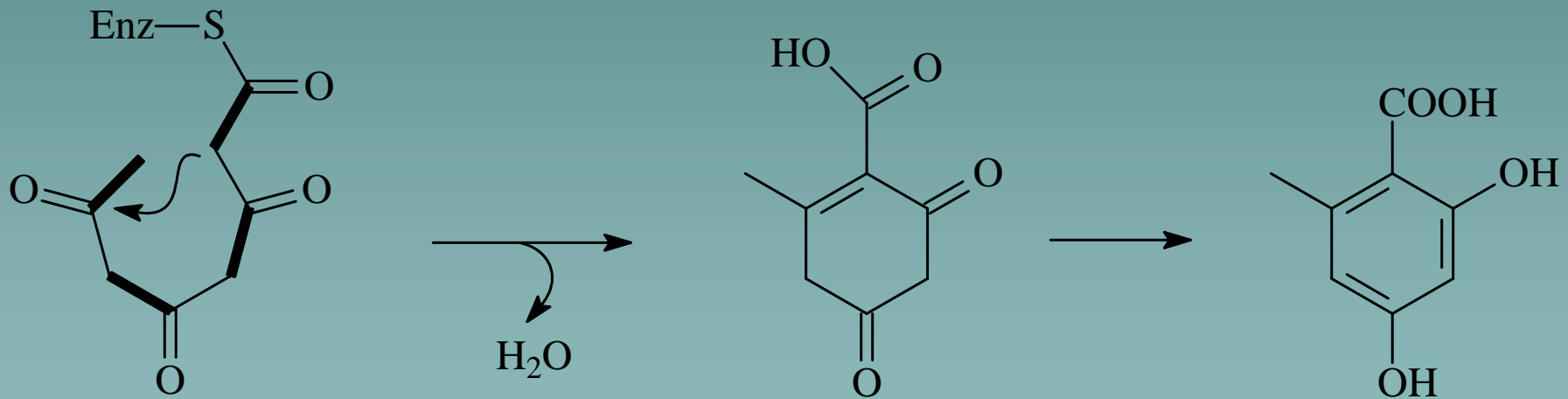




Orsellinic acid occurs in the antibiotic avilamycin, and the anticancer enediyne calicheamicin.

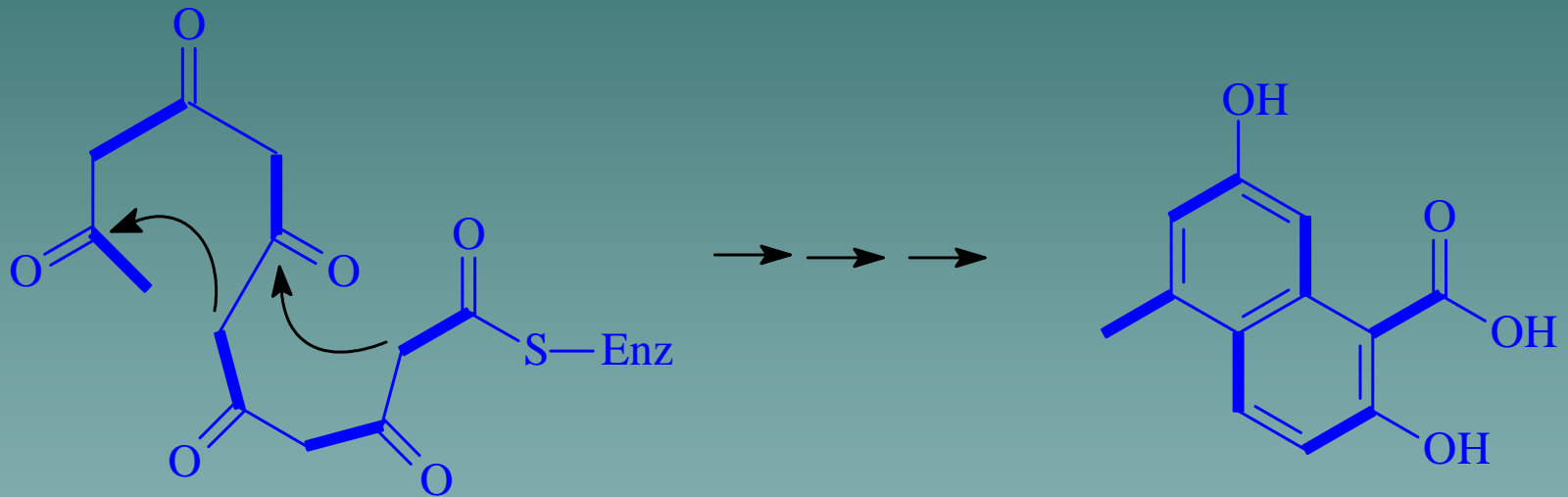
It is synthesised by an iterative (non-modular) type I PKS.

KS – AT - (DH) - ACP



Orsellinic acid

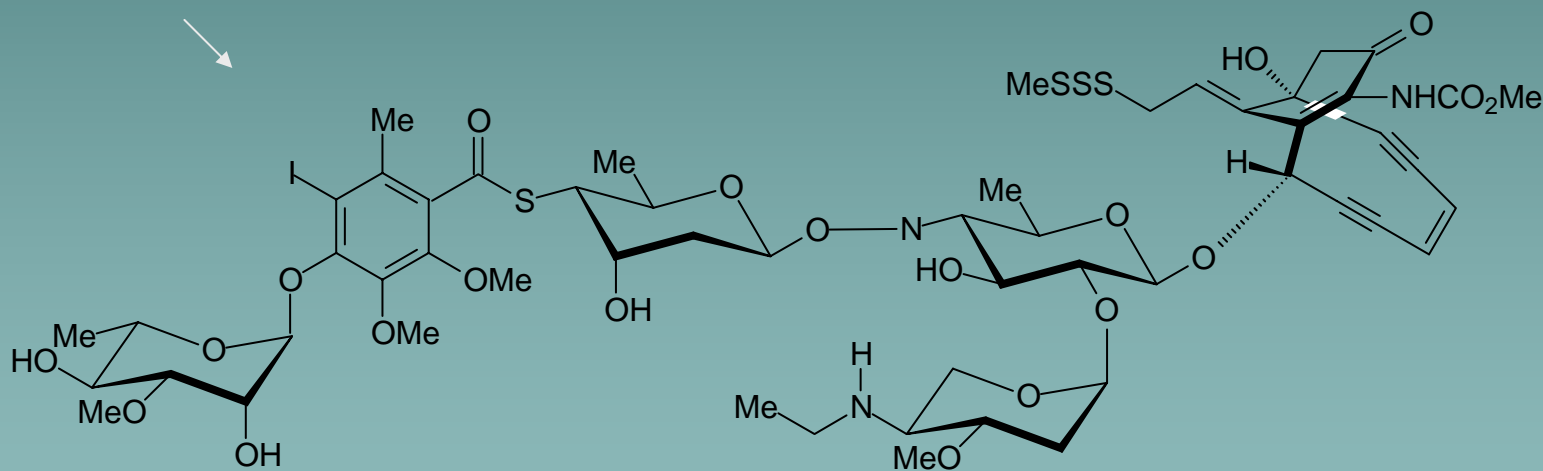
Napthalinic acid is also synthesised by an iterative type I PKS.



Eneidyne antibiotics contain two polyketide-derived moieties.  
Both are synthesized by iterative type I PKSs.

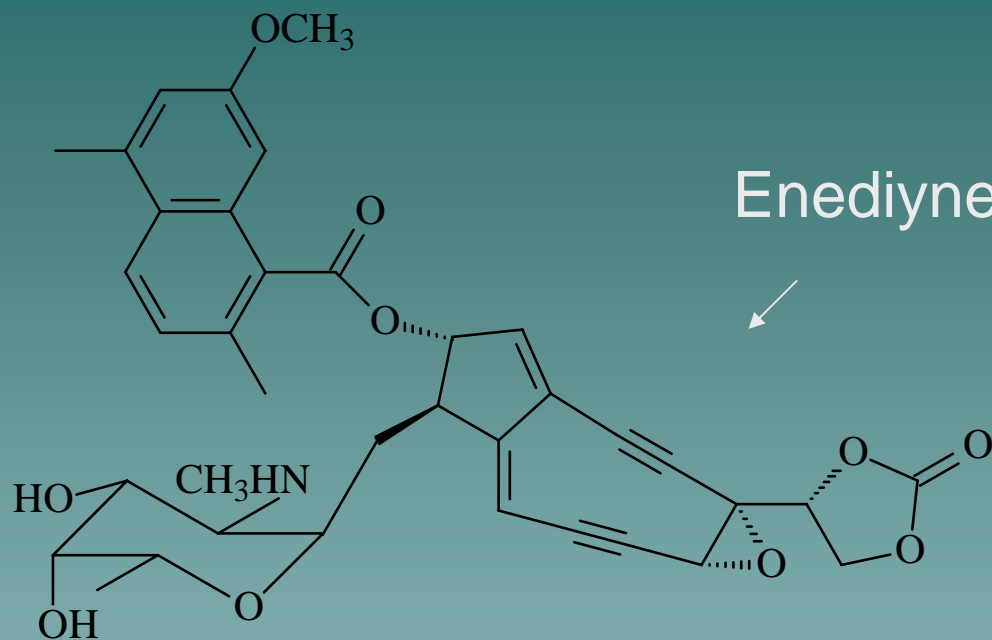
Orsellinic acid

Eneidyne core



Calicheamicin

Napthalinic acid

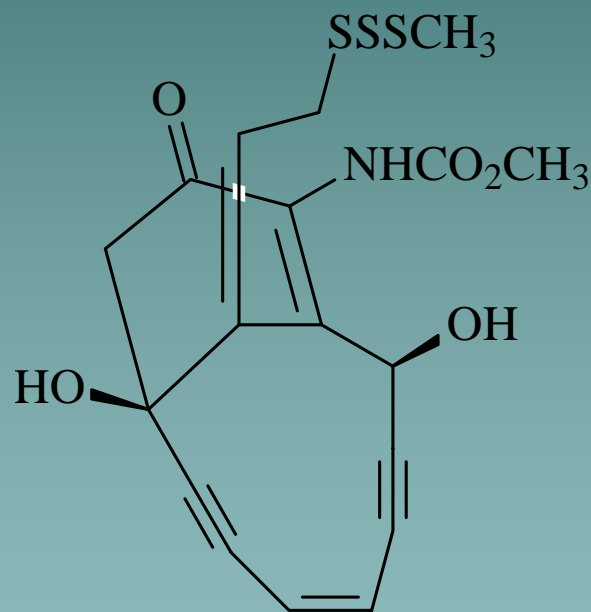


Enediyne core

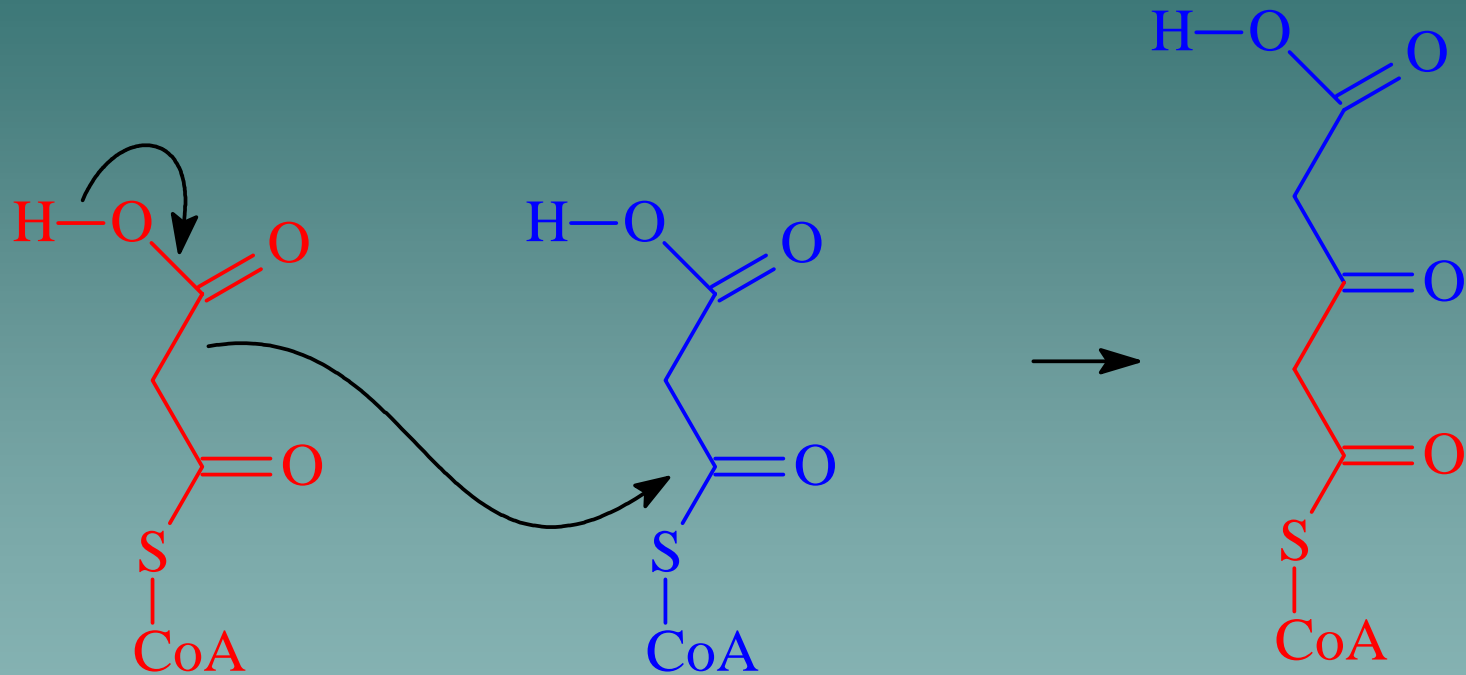


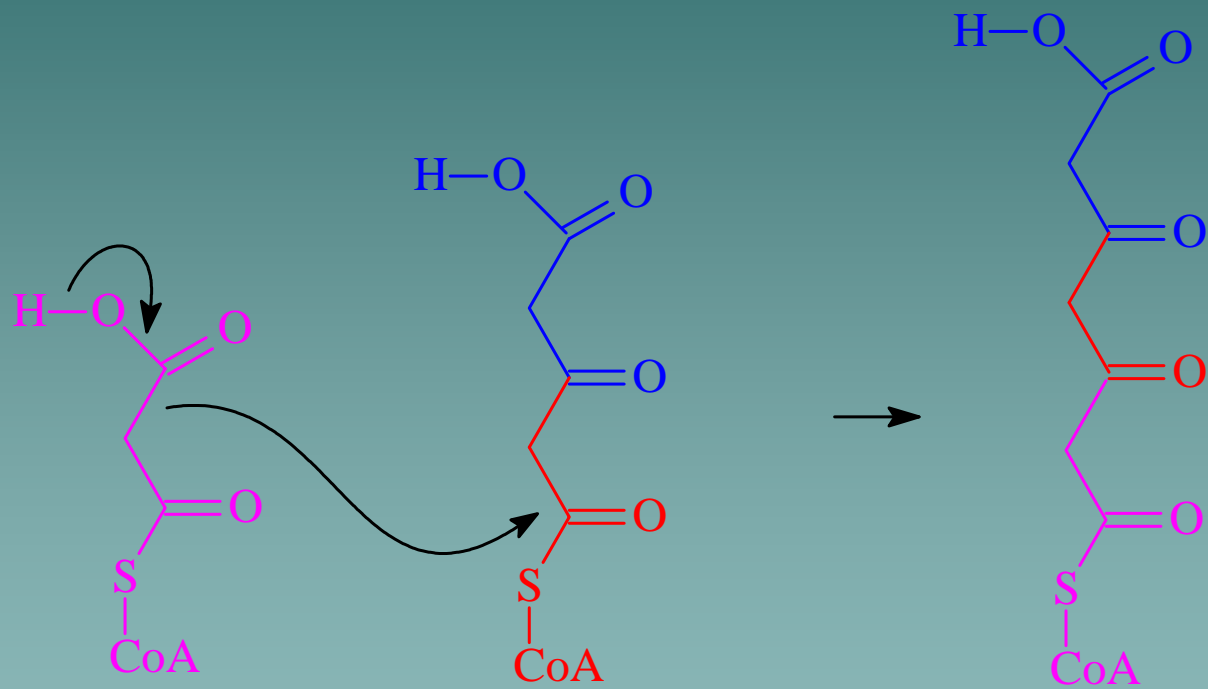
Enediyne C1029

Enediyne cores are derived from polyunsaturated acyl chains that are synthesised by iterative type I PKSs.

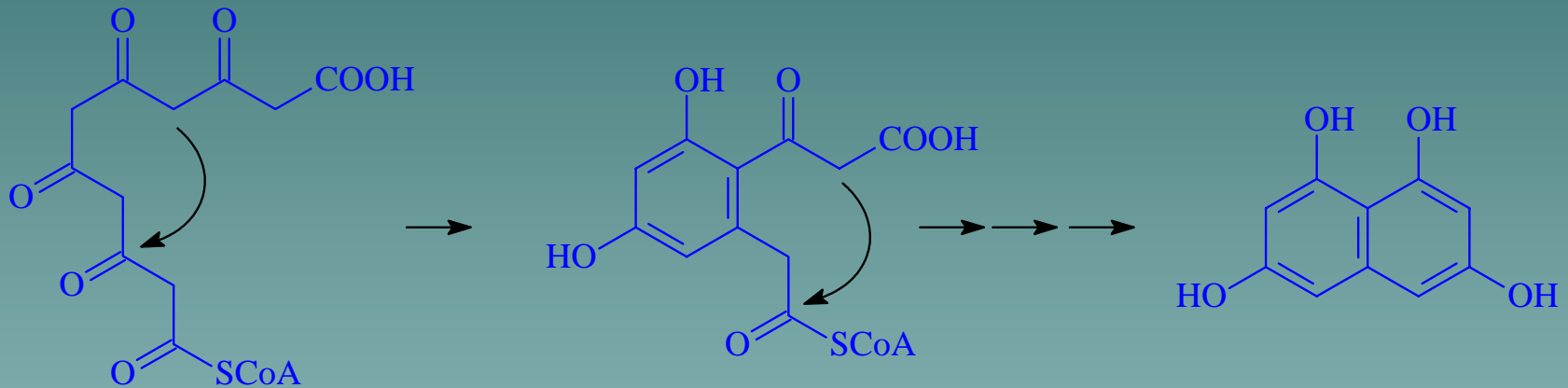


**Type III PKSs** are composed of KS domains only. They assemble the polyketide chain on CoA pantetheine thiol (not ACP).





Type III PKSs synthesise pigments (e. g. flower colours),  
chalcones, flavonoids, flaviolins.

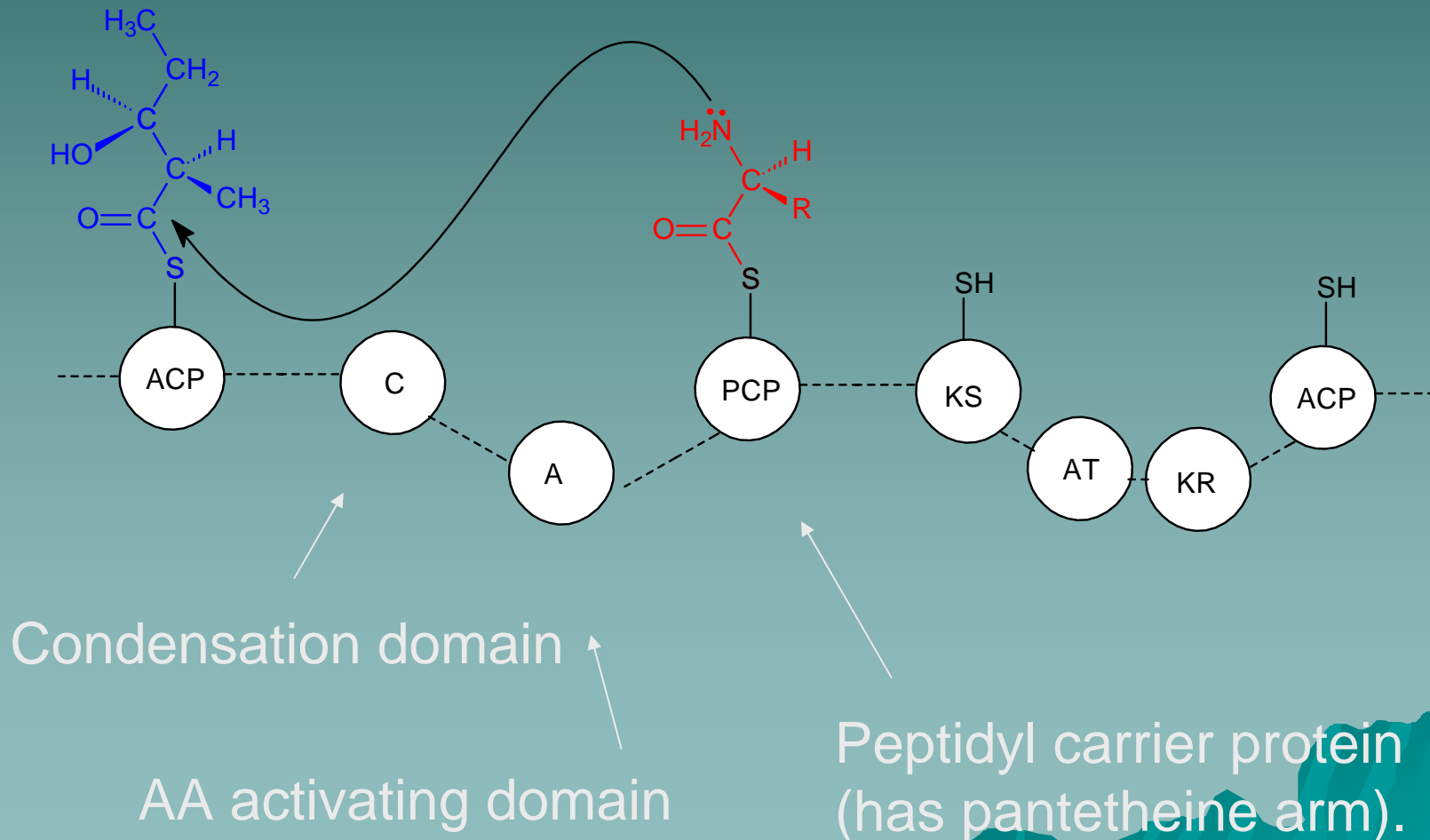


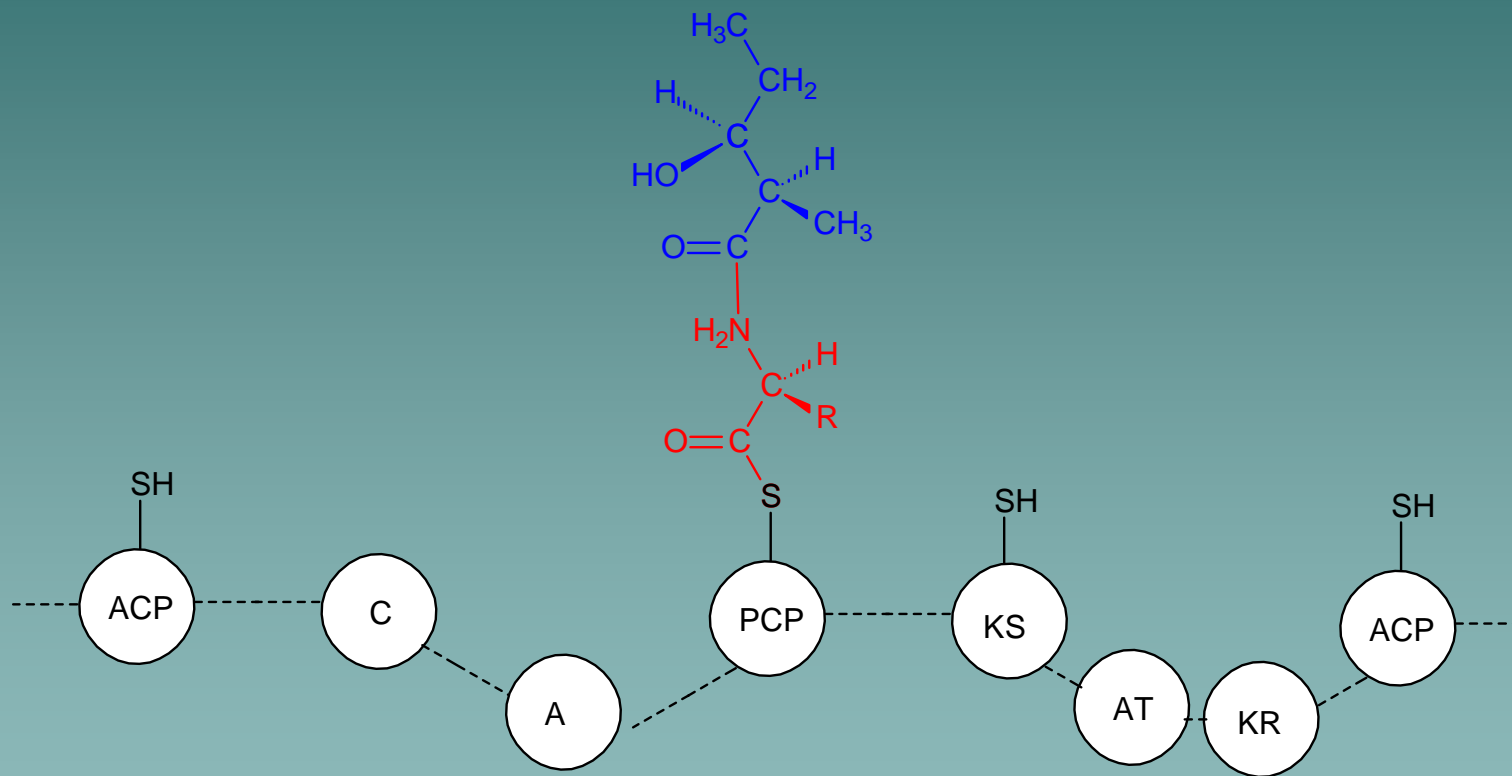
Flaviolin

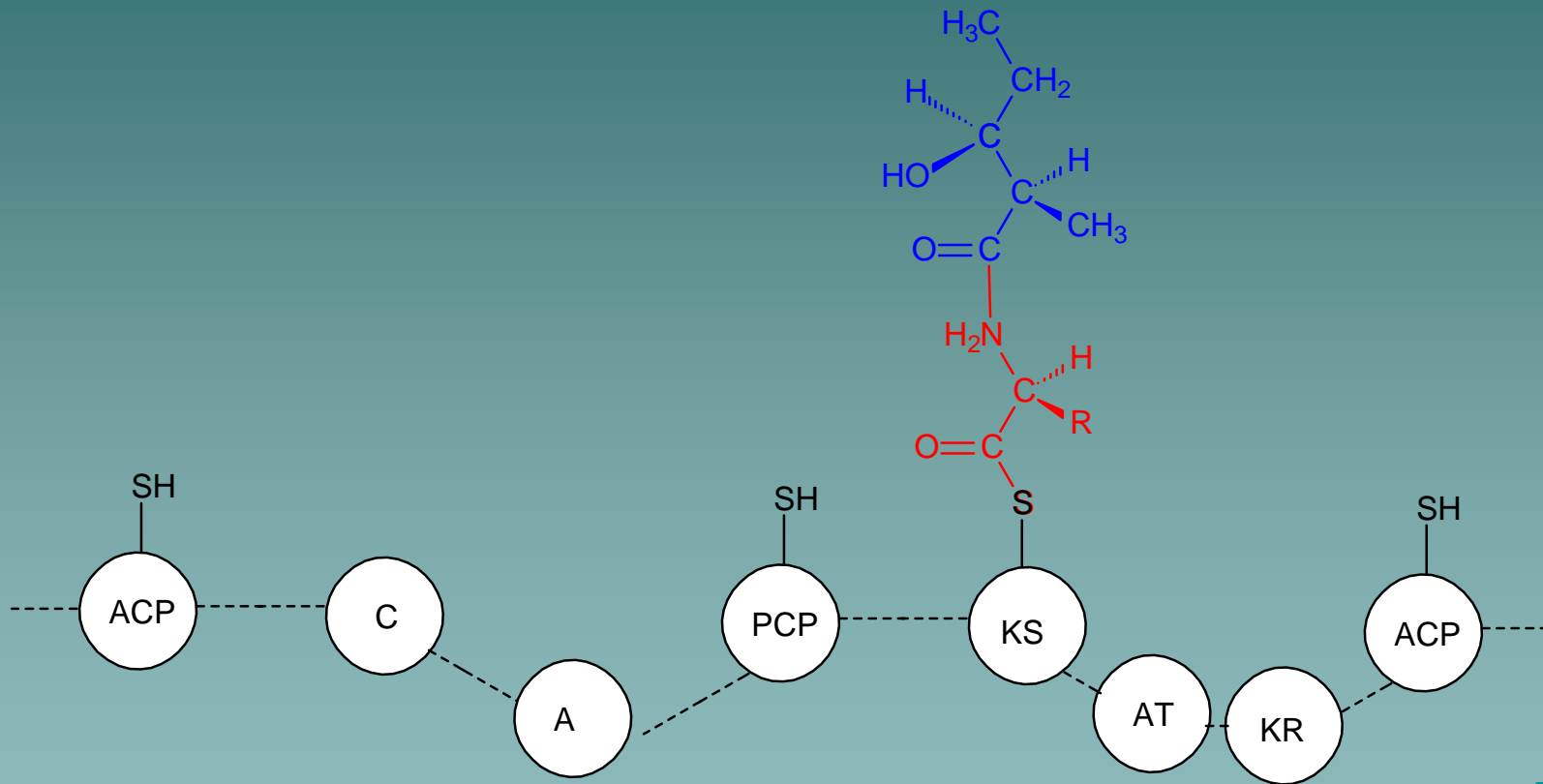


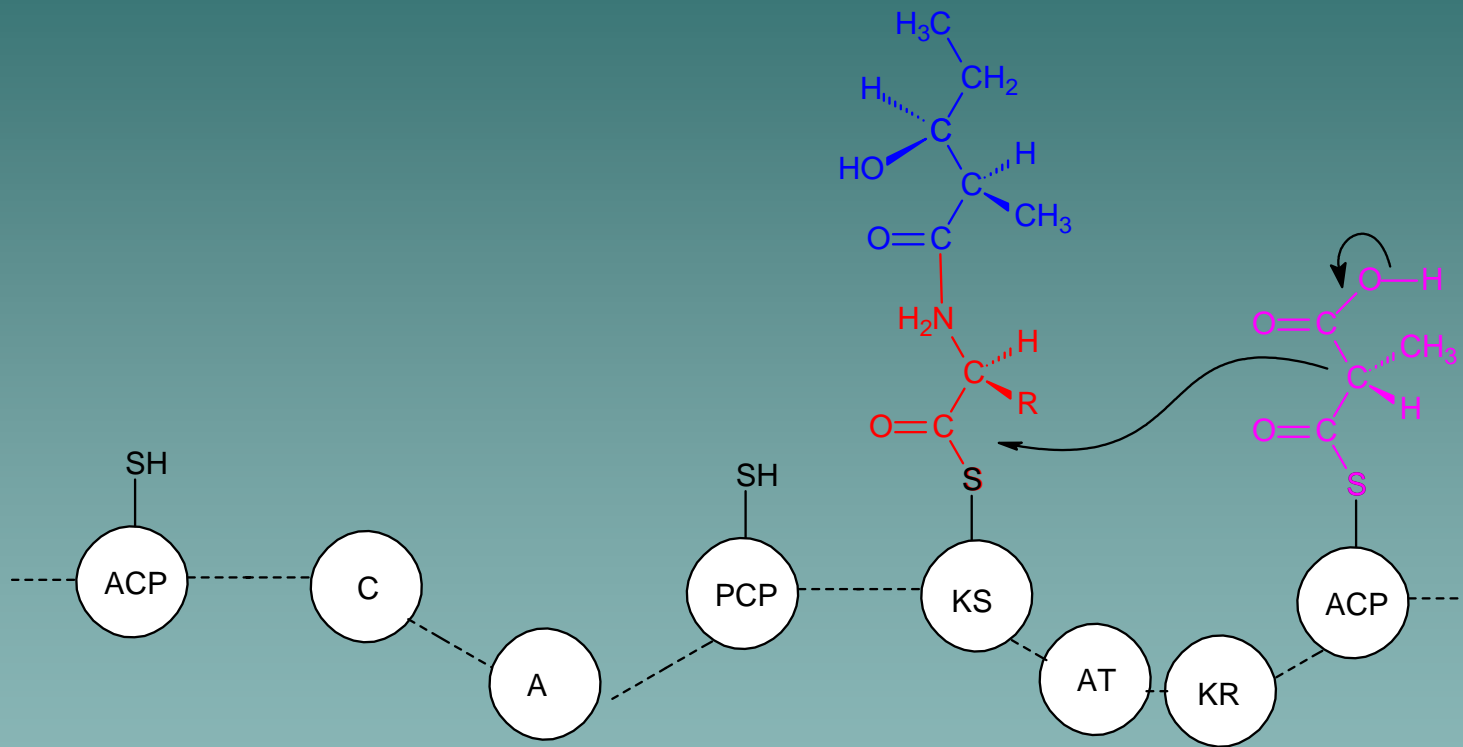
# Mixed PKS-NRPS (non-ribosomal peptide synthetase) systems.

Modular PKSs that can use amino acid extenders as well as dicarboxylic acid extenders.

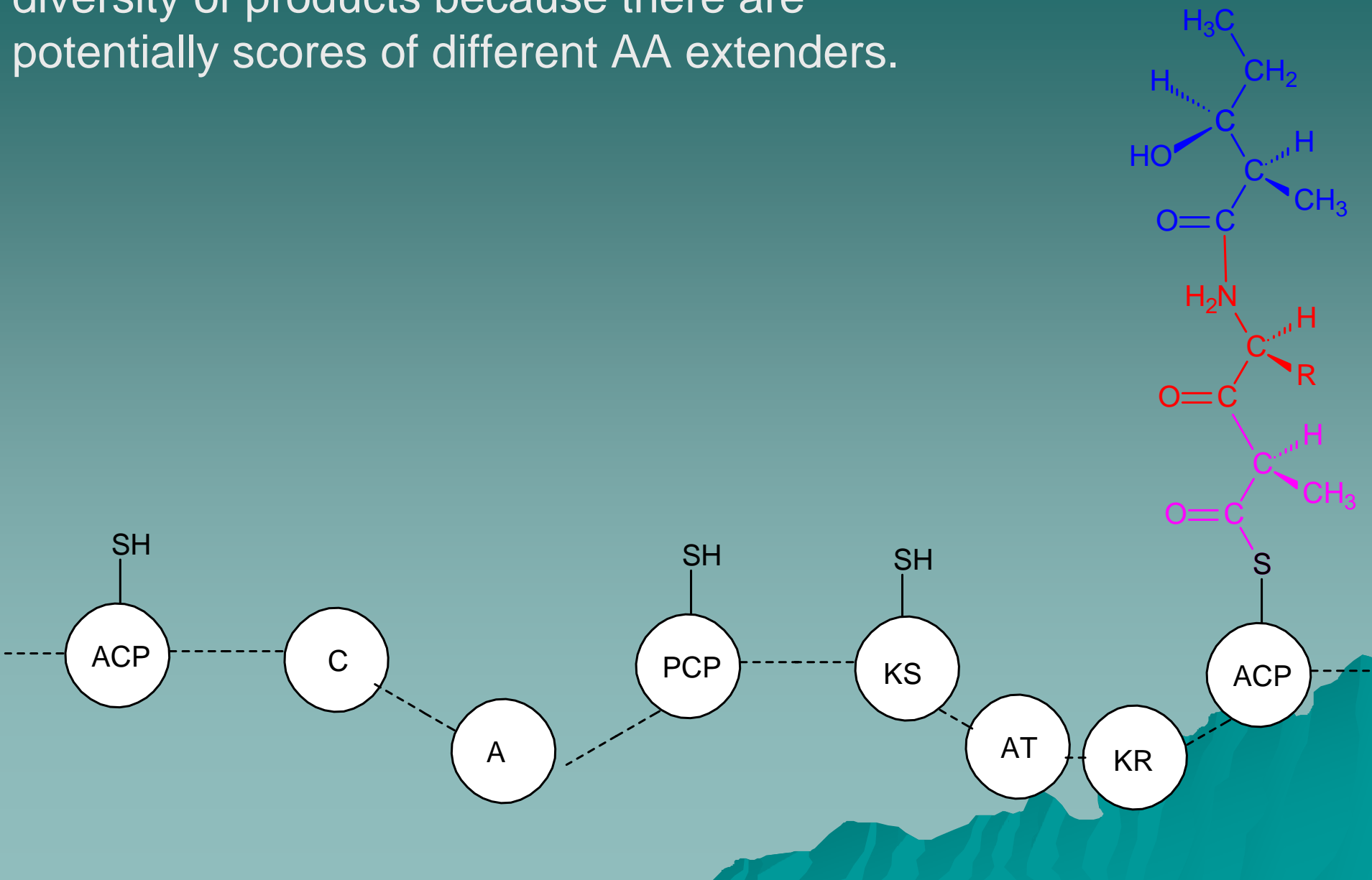








In principle, mixed PKS-NRPSs could make a greater diversity of products because there are potentially scores of different AA extenders.



# Types of PKS


**Type III** (KS domains only, build chains on CoA)

**Type II** (composed of small discrete enzymes, synthesise aromatic polyketides).

**Type I** (covalently linked enzymes[e.g. KS-AT-X- Y-ACP])

**Type I iterative** – one set of enzymes is used repeatedly.  
Relatively simple chains are made.

**Type I modular** – a set of enzymes for every cycle includes mixed PKS-NRPSs.  
Complex polyketides are made.  
Can be reprogrammed.

A stylized, dark teal mountain range graphic is located in the bottom right corner of the slide, partially overlapping the text of the 'Type I modular' section. The mountains are depicted with sharp, jagged peaks and are rendered in a monochromatic teal color that matches the slide's background.

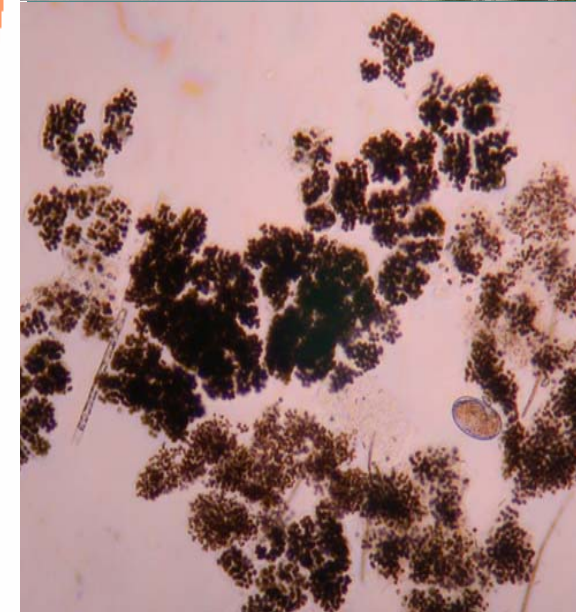
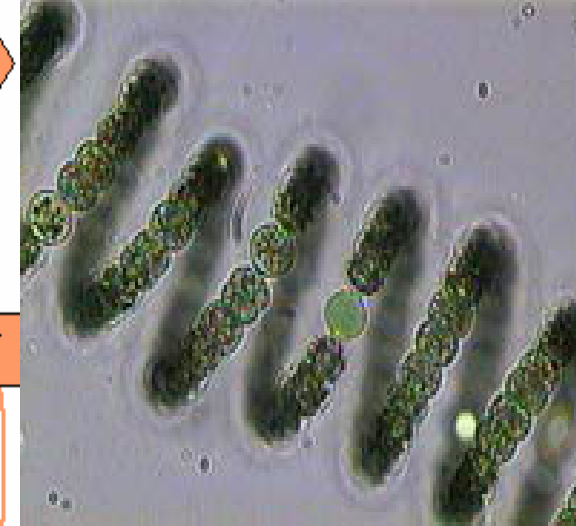
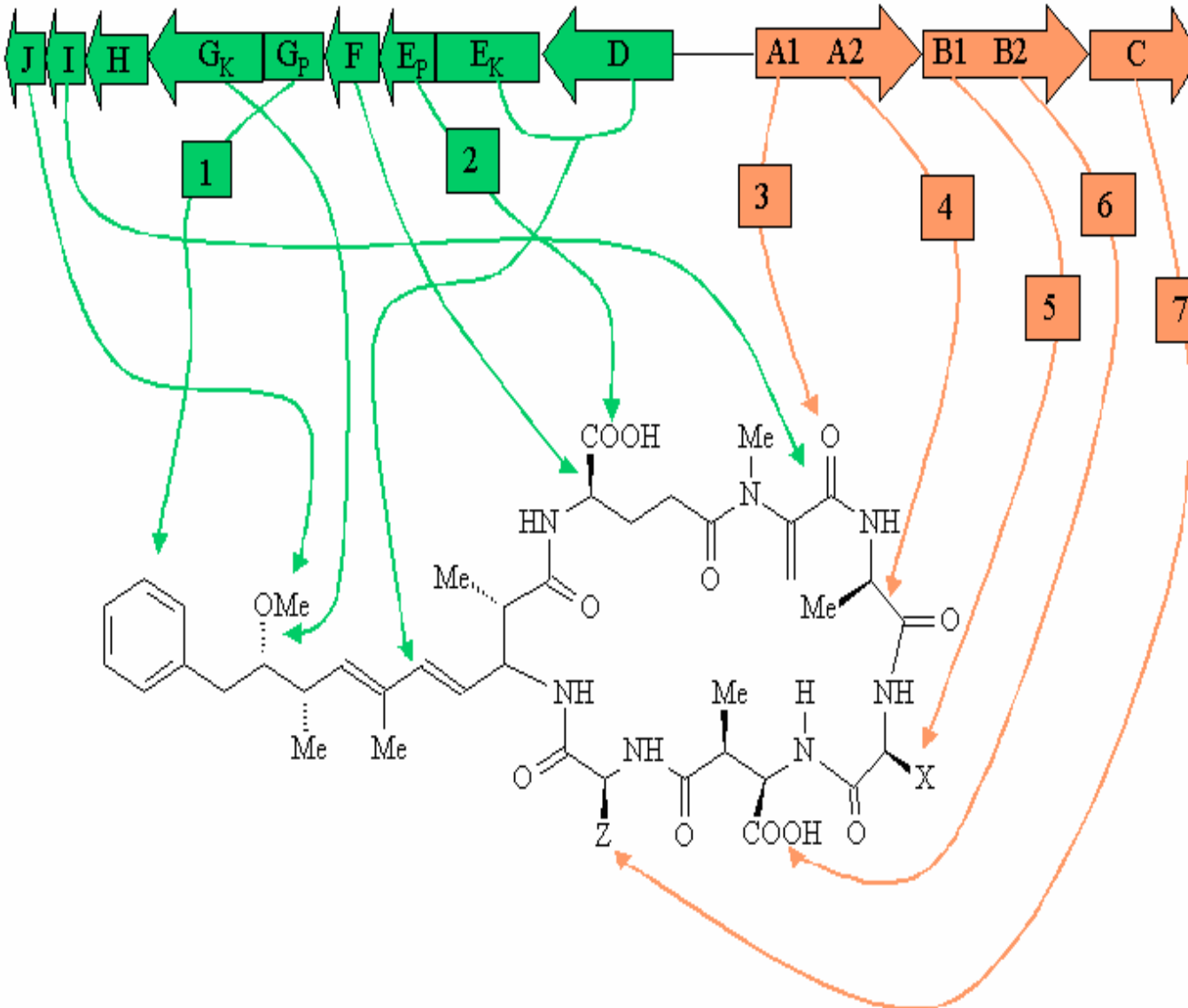
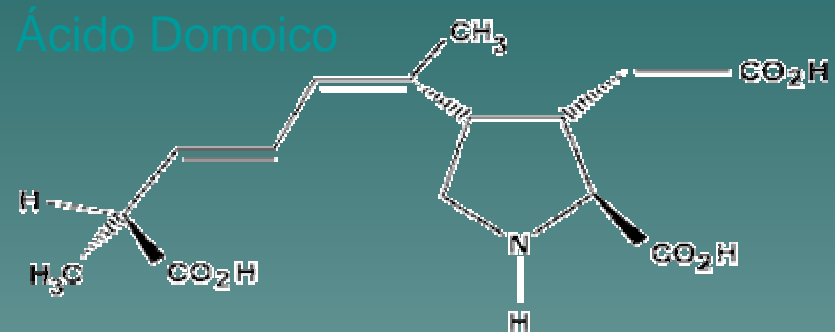
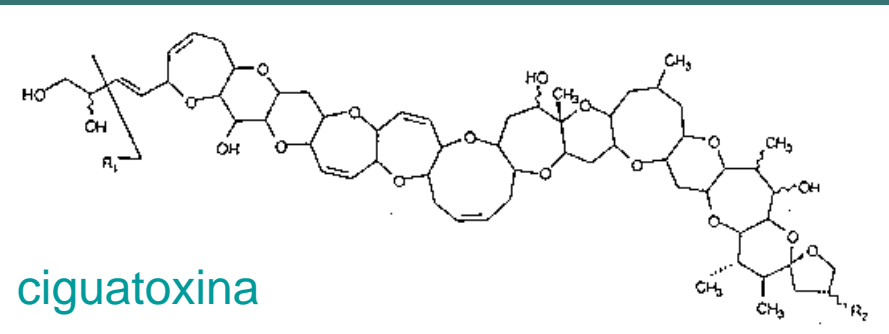
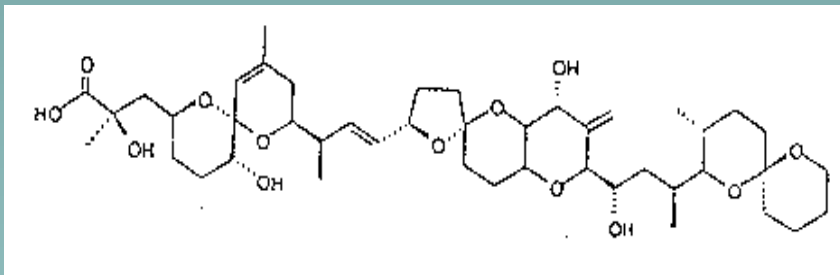


Figure 1: The structure of microcystin and its biosynthetic pathways (left). The gene organization is shown on the top. The amino acids in the X and Z positions represent variable amino acids. *Anabaena* (top right) and *Microcystis* (bottom) as seen under a microscope.

# Toxinas

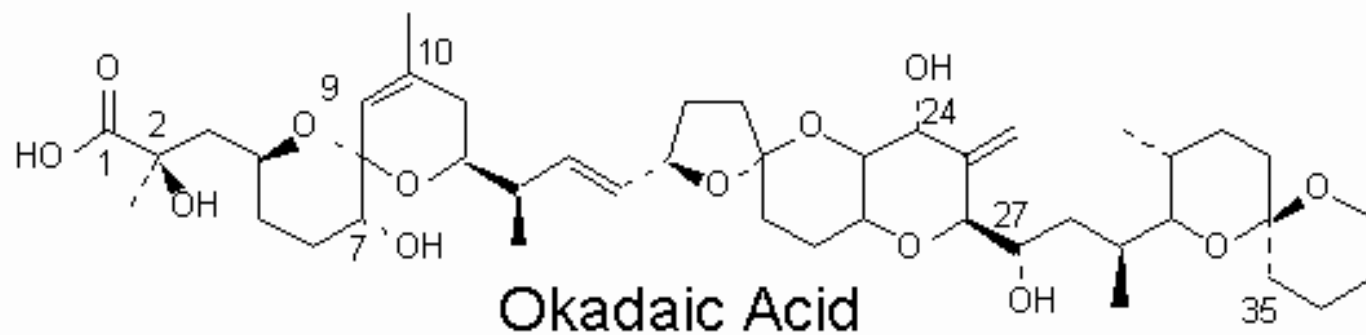


Ácido okadaico

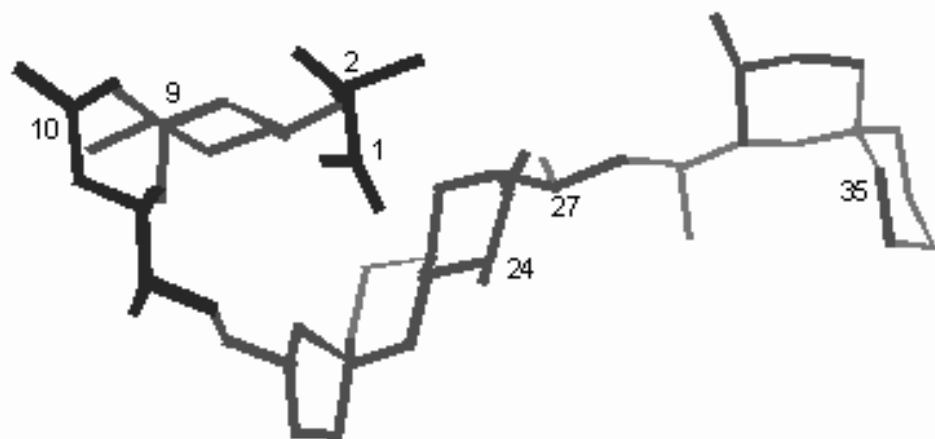




A

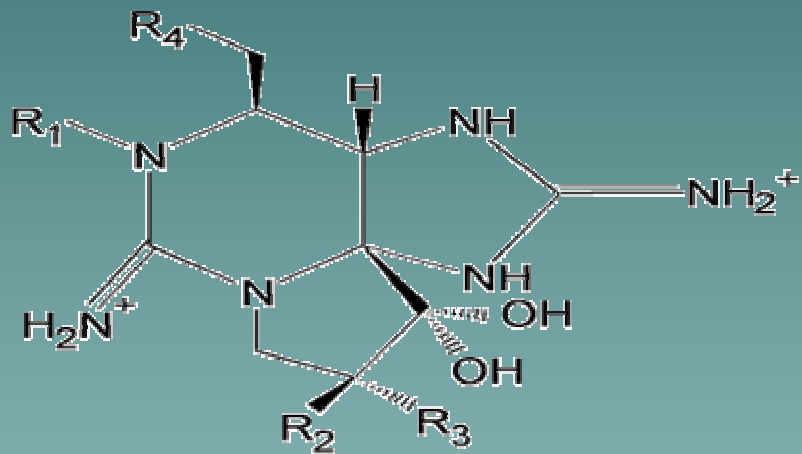


B



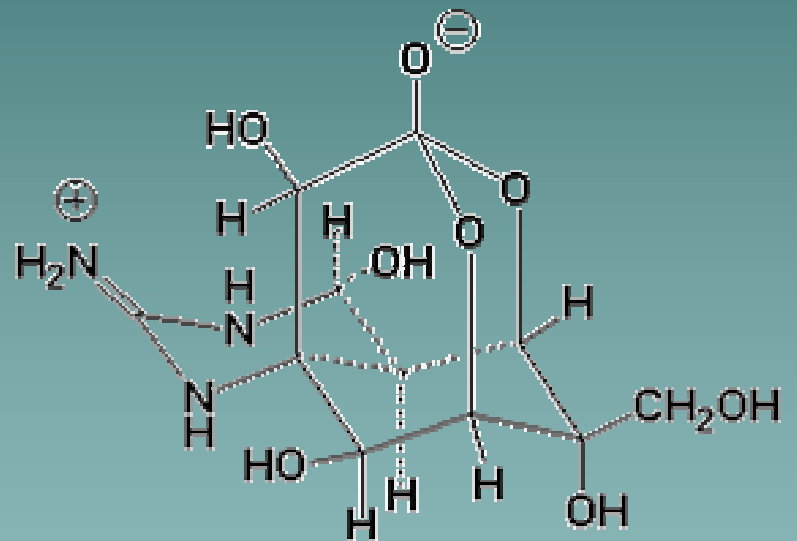
	IC <sub>50</sub>
PP-1	20 nM
PP-2A	0.2 nM
PP-2B	5.0 μM

?



PSP toxins

?



Tetrodotoxin

## Useful references.

Combinatorial biosynthesis of erythromycin and complex polyketides. Staunton, J. 1998.

Current Opinion in Chemical Biology 2: 339- 345.

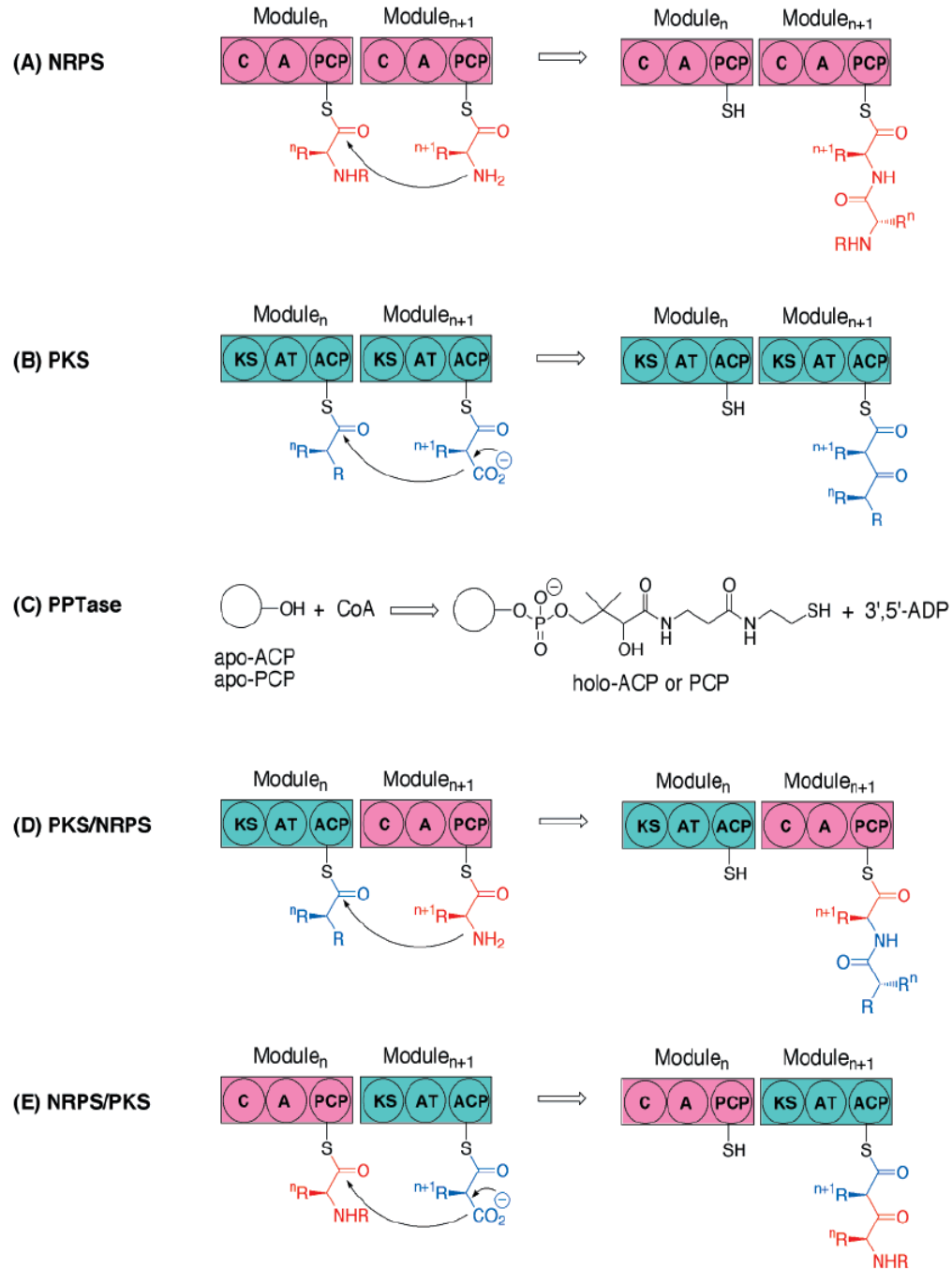
Polyketide biosynthesis beyond the type I, II and III PKS paradigms. Shen, B. 2003. Current Opinion in Chemical Biology 7: 285- 295.

Building block selectivity of polyketide synthases.

Liou, G. F. and Khosla, C. 2003. Current Opinion in Chemical Biology 7: 279-284.

Antibiotics: action, origin and resistance by C. Walsh 2003

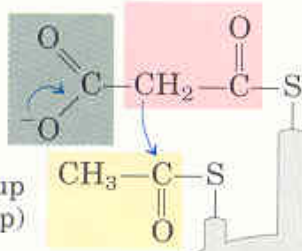




**Figure 2.** Modular organization of NRPS (A), PKS (B), hybrids PKS/NRPS (D) and NRPS/PKS (E), and post-translational modification of apo-ACP or apo-PCP into holo-ACP and holo-PCP by a PPTase (C). Hypothetical NRPS (pink) and PKS (blue) modules are shown with core domains. A, adenylation; ACP, acyl carrier protein; AT, acyltransferase; C, condensation; KS, ketoacyl synthase; PCP, peptidyl carrier protein.

# FAS

Malonyl group



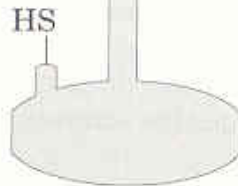
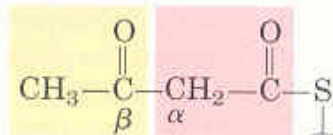
Acetyl group  
(first acyl group)



Fatty acid synthase

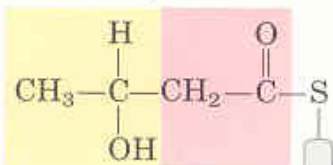
condensation

①



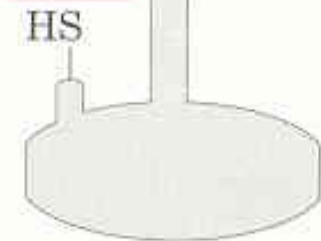
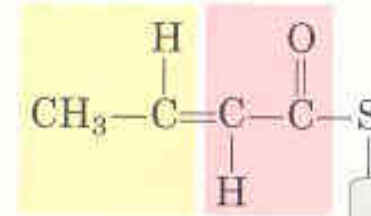
reduction

②



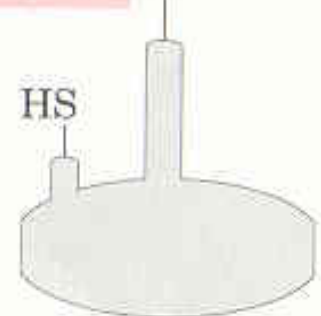
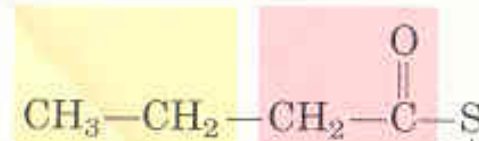
dehydration

③



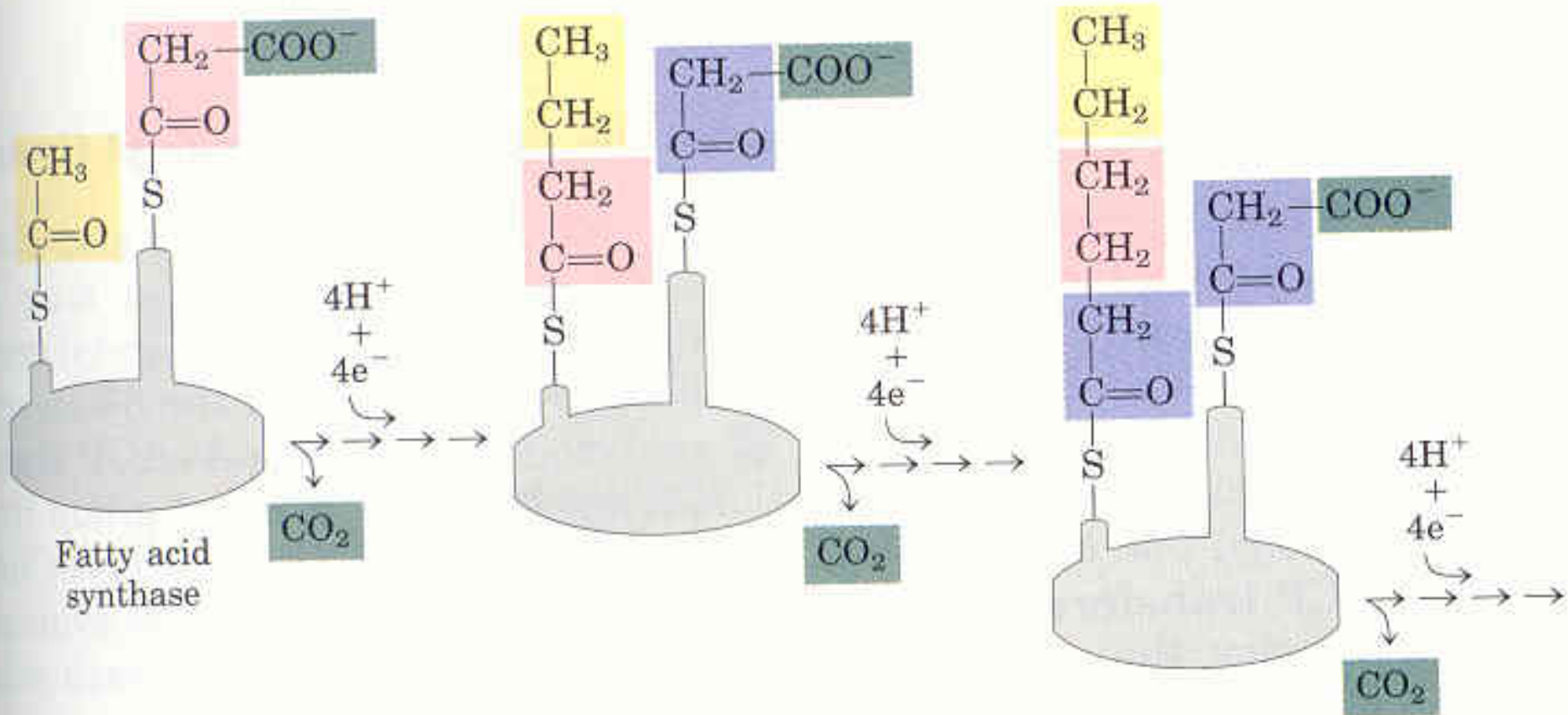
reduction

④



Saturated acyl group,  
lengthened by two carbons

# FAS: elongation



# FAS: elongation

