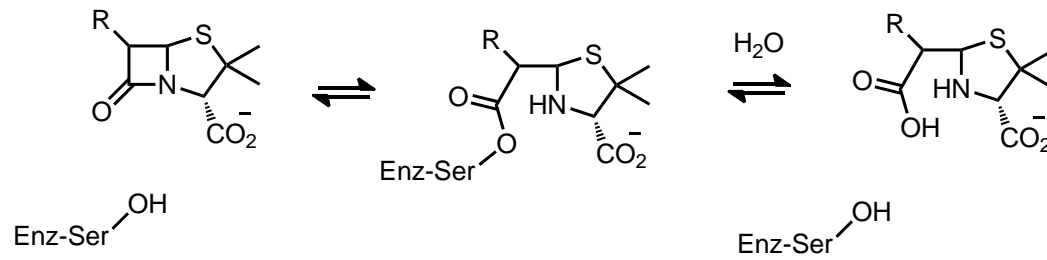


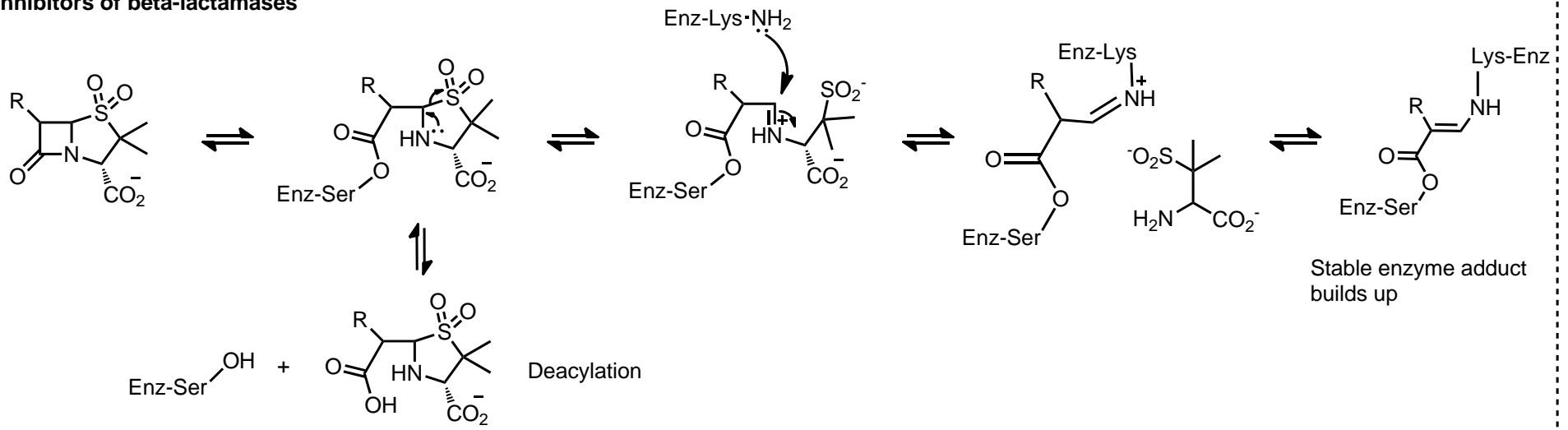
11.10.15 - Lecture 15- Antibiotics and RNA catalysts

Penicillin antibiotics-

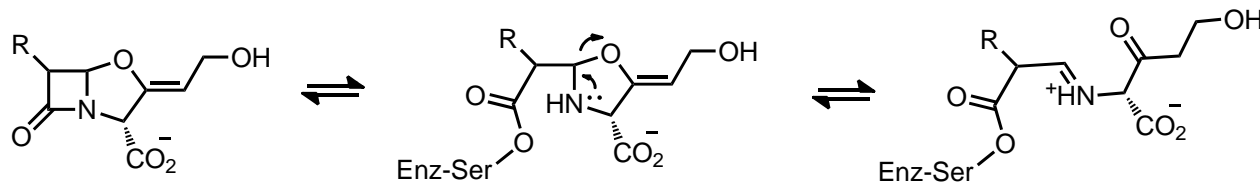
-inactivated by beta-lactamase's ability to perform a fast deacylation of the inhibitor-enzyme adduct



Inhibitors of beta-lactamases

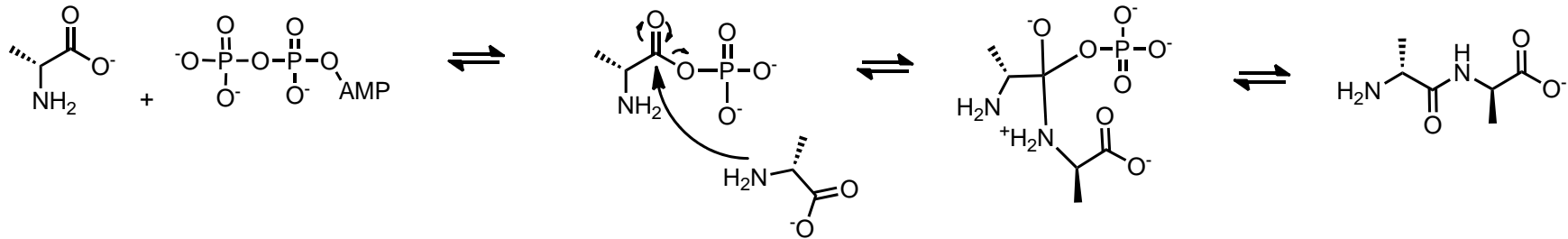


Using an alternative leaving group-



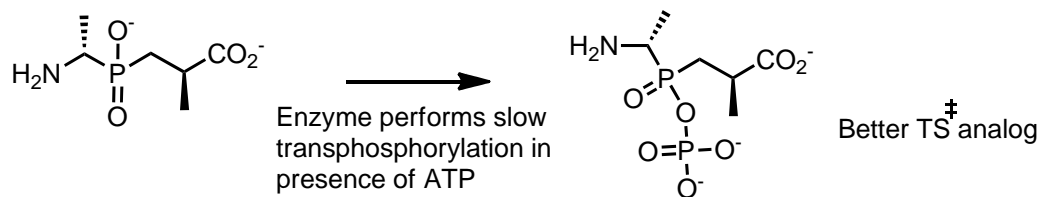
In general, beta-lactam antibiotics are not affected by drug pumps as their target resides in the periplasmic space

Targeting D-Ala D-Ala ligase



Design of an inhibitor of alanine ligase

- inhibition is dependent on presence of ATP
- off rate $t_{1/2} \sim 17$ days
- $k_{off}^{INH} \ll k_{off}^{PDT}$, 10^8 - 10^9 slower off rate
- example of slow binding/slow off rate inhibitor



Previously discussed antibiotic mechanisms targeting D-Ala D-Ala

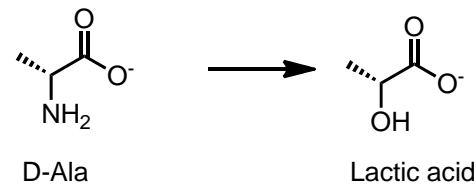
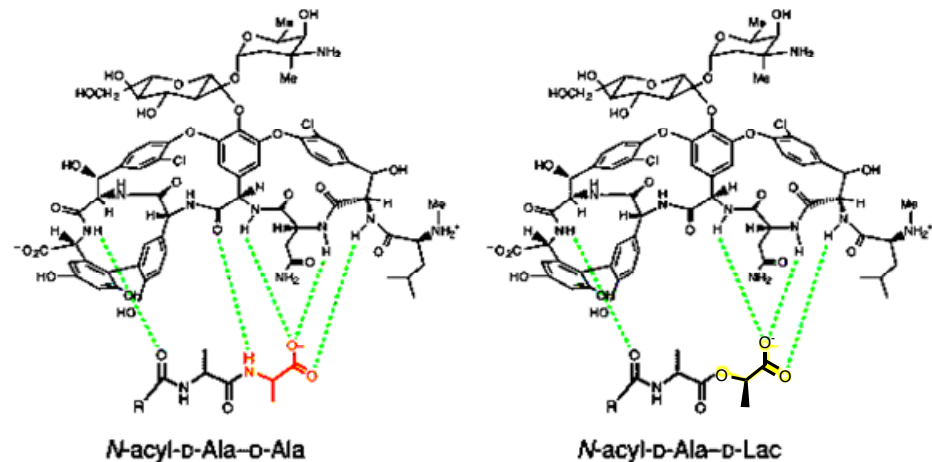
- transpeptidase inhibitor
- inhibitor of alanine racemase
- inhibition of D-Ala ligase

Vancomycin-

-a natural product 'receptor' which specifically binds to D-Ala-D-Ala via a network of H-bonding and blocks transpeptidase

Vancomycin Resistance

- bacteria use lactic acid instead of D-Ala
- ester linkage is much less favorable, repulsive interactions between oxygen lone pairs, loss of H-bond

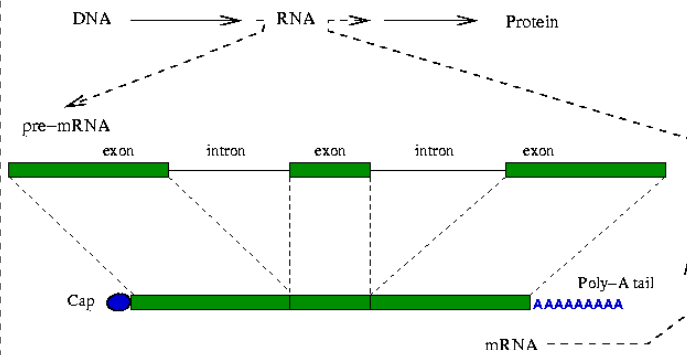


Loss of H-bond and introduction of unfavorable interaction

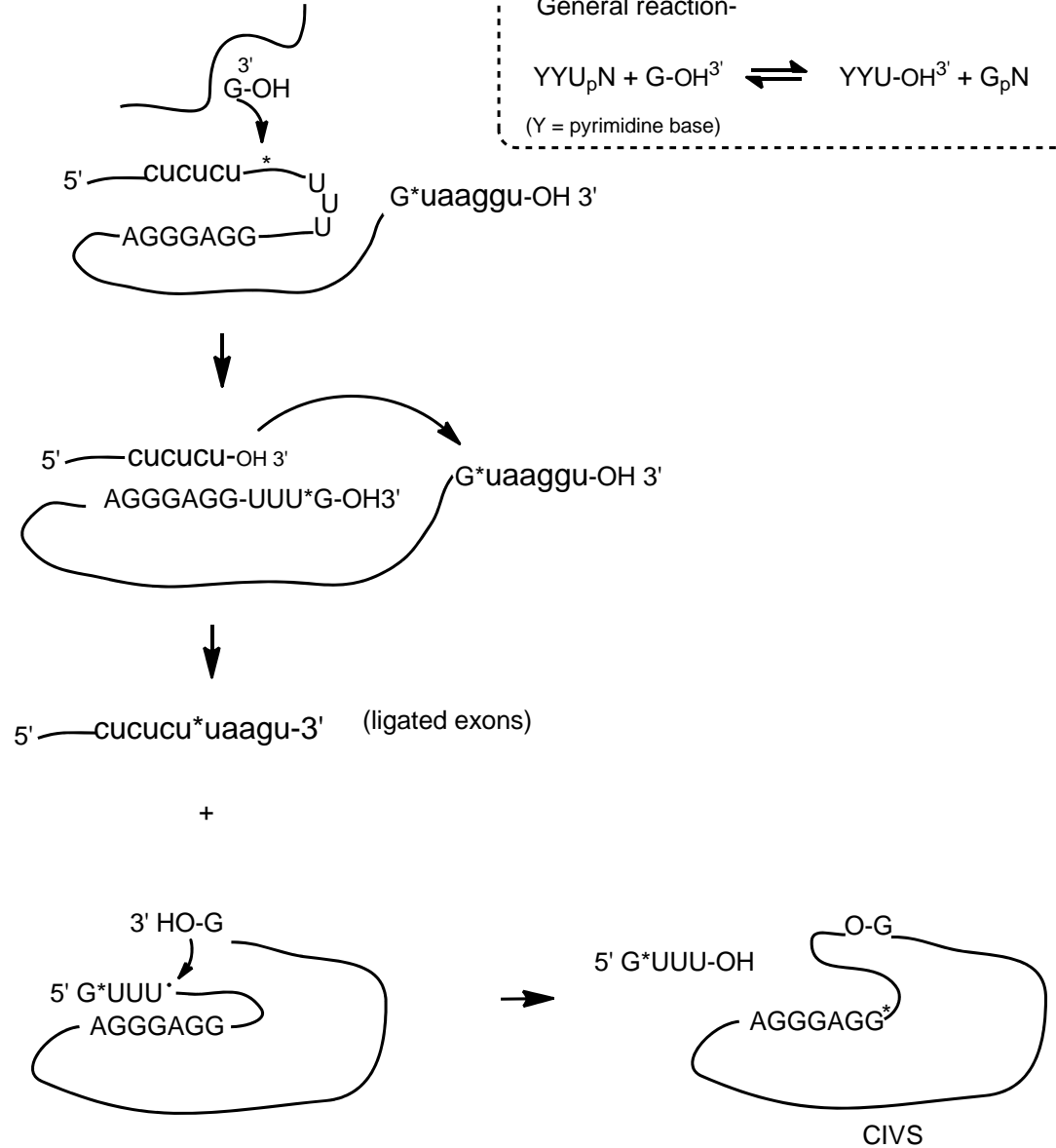
RNA catalysts

- RNA can carry out catalysis
- most often dependent on presence of Magnesium

Intron splicing

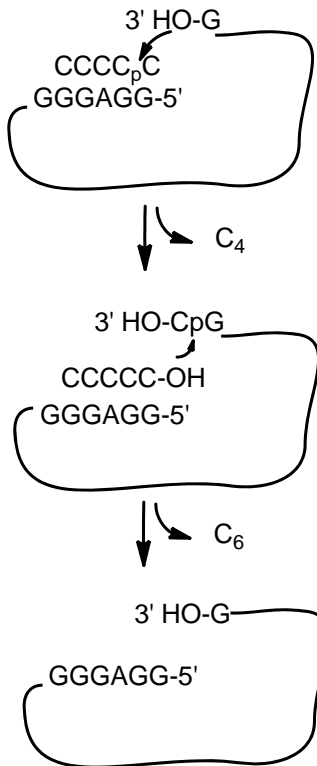


Self splicing RNA in tetrahymena



Catalytic RNA

-in order to be catalytic, the catalyst molecule must remain unchanged at the end of the reaction

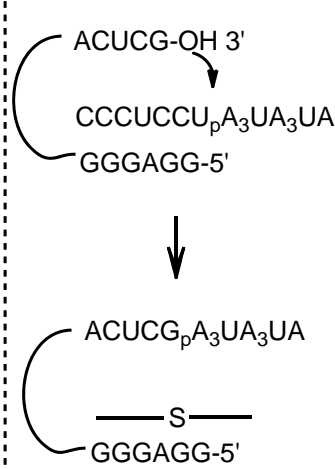


+ C₄ + C₆

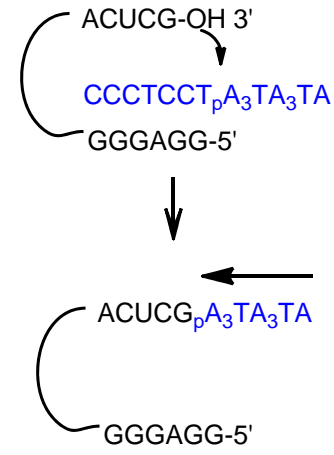
(catalytic RNA unchanged)

(catalyzes disproportionation reaction)

RNA ligase-



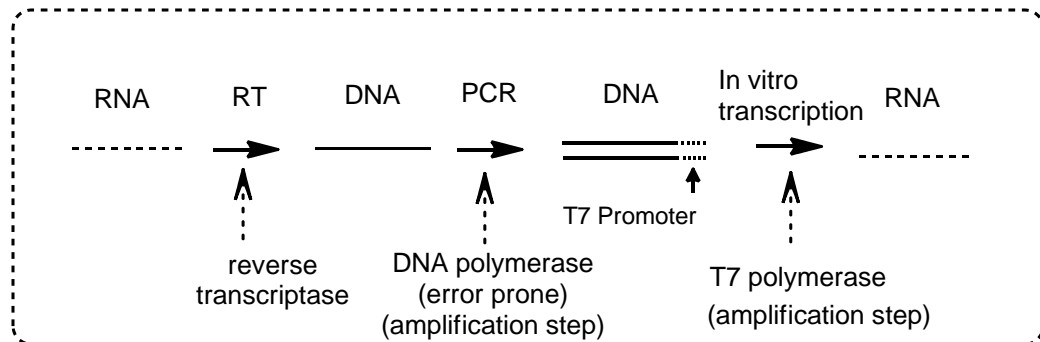
RNA which ligates to DNA substrate-



Amplification of starting RNA with RT rxn, PCR, and in vitro transcription with T7 polymerase. Sequence diversity generated by error prone PCR conditions or using an error prone DNA polymerase

RNAs with increased catalytic activities

Starting RNA	$k_{cat} - 2 \times 10^{-4} \text{ min}^{-1}$	$K_m \sim 7 \mu\text{M}$
After selection	$k_{cat} - 0.007 \text{ min}^{-1}$	$K_m \sim 2 \mu\text{M}$



An RNA ligase selected from randomized sequence

- randomize 220 bases
- library will be incomplete (4220 potential members possible); must assume many potential solutions

Using this strategy-
an RNA enzyme was discovered with
a $t_{1/2}$ of 5 minutes vs. 33 years compared
to the uncatalyzed reaction
(10^7 rate acceleration)

