

# DNA topoisomerases as targets for antibacterial chemotherapy

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# Maxwell lab

## Enzymes:

Gyrases - *E. coli*, *M. tuberculosis*,  
*S. aureus*, *Arabidopsis*, *Plasmodium*  
Topo IV - *E. coli*, *S. aureus*  
Topo VI – *Methanosarcina mazei*, *Arabidopsis*,  
*Plasmodium*  
Other topo IIs (yeast, human)

## Drugs & toxins:

Quinolones (e.g. ciprofloxacin)  
aminocoumarins (e.g. novobiocin)  
simocyclinones, naphthoquinones,  
CcdB (toxin-antitoxin)  
Microcin B17  
Chemical libraries

## Tools:

X-ray crystallography, NMR, etc.  
Enzymology, Molecular Biology, Genetics, HTS  
Chemical Genetics

## Natural products:

Novel plant toxins/  
insect gut bacteria  
Galleria

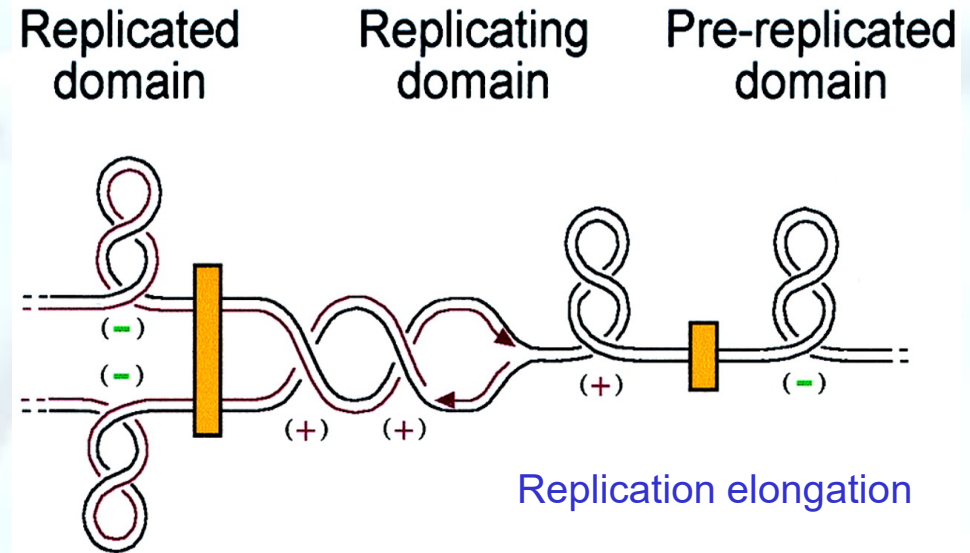
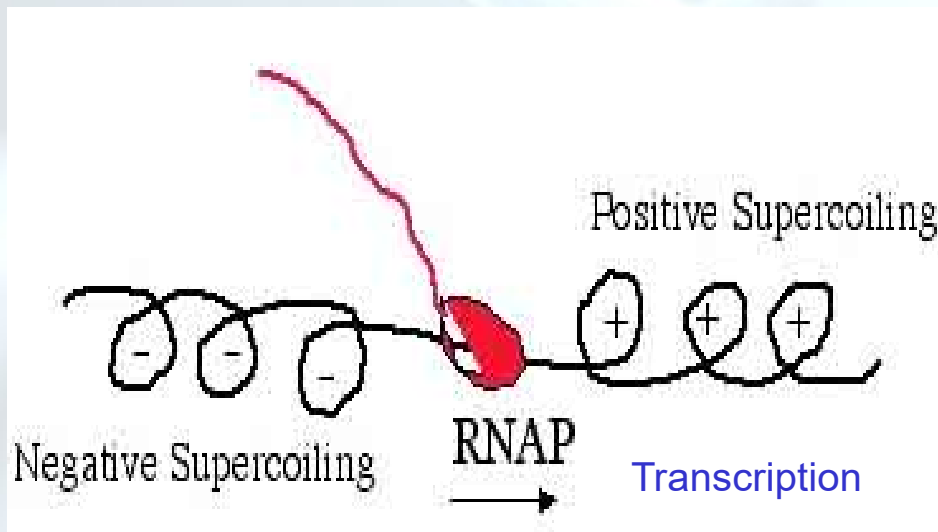
## Outputs:

Fundamental knowledge on topoisomerase structure/mechanism/physiological role  
Understanding drug action and antibiotic resistance; development of novel agents



# DNA topology & DNA topoisomerases

Various biological processes lead to DNA topological problems, e.g. introduction of unwanted supercoiled and catenanes



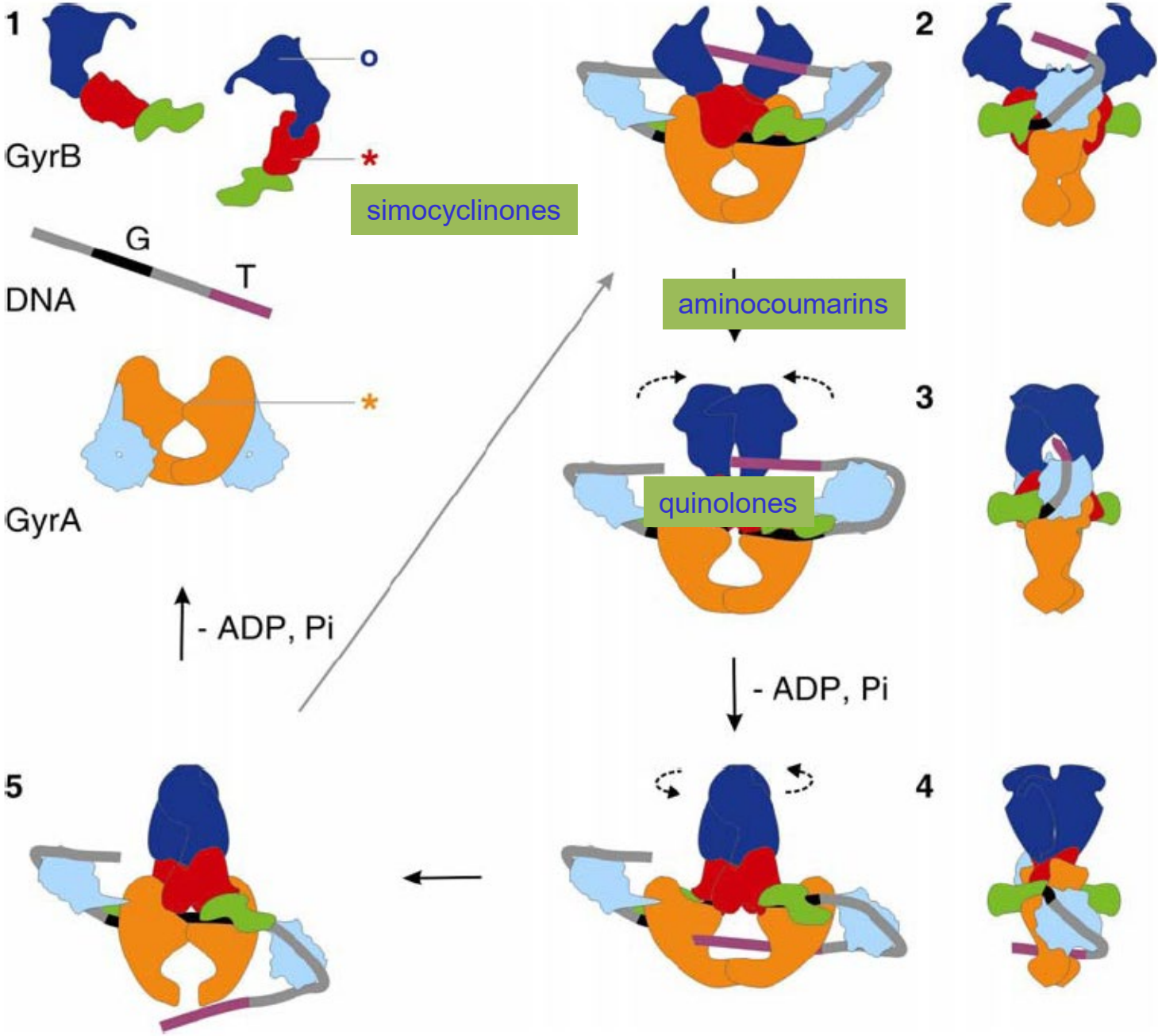
Topos, two types:

I → single-strand breaks

II → double strand breaks

- Currently no antibiotics targeted to type I topos.
- Type II topos, DNA gyrase & topo IV highly successful antibiotic targets

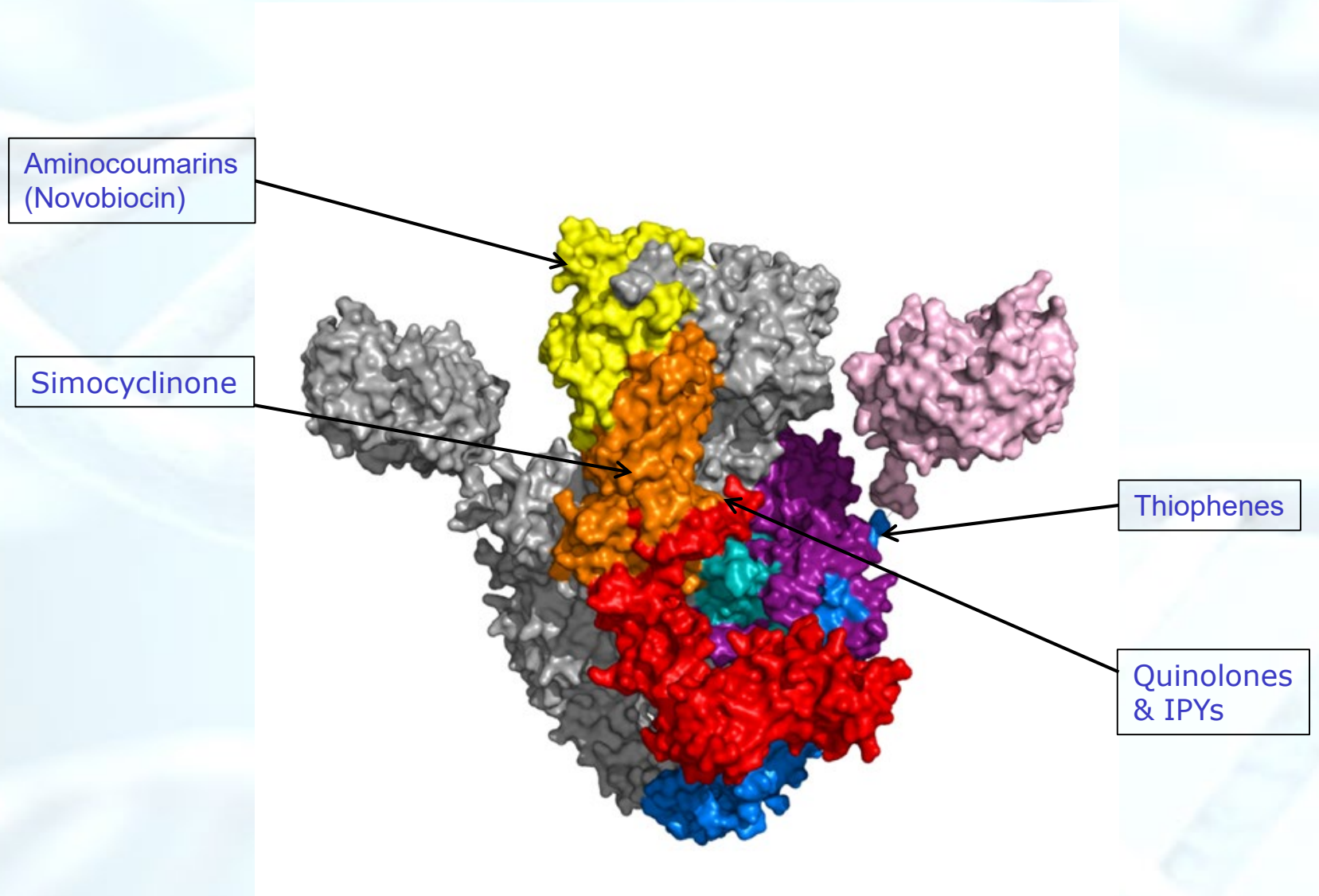
# Mechanism of DNA supercoiling by gyrase



Costenaro, *et al.*  
(2007).  
*Structure* **15**,  
329-39.

# Sites of drug action on DNA gyrase

- Bax *et al.* *J Mol Biol.* 2019; In press.
- Germe *et al.* *Nucleic Acids Res.* 2018; **46**:4114-28.
- Bush *et al.* *The Biochemist.* 2018; **40**:26-31.
- Nagaraja *et al.* *Drug Discov Today.* 2017; **22**:510-8.
- Chan *et al.* *Proc Natl Acad Sci U S A.* 2017; **114**:E4492-E500.



# Funding & Consortium/Collaboration Activities

- ENABLE – EU: IMI (ND4BB) consortium
- Mm4tb – EU consortium; now finished; 2 TB drugs in clinical trials
- AMR on the NRP
- Screening services and provision of reagents (via Inspiralis)
- Scientific Advisory Board of Antibiotic Action

# Bottlenecks

What we do is a very small slice of a very big cake and our contribution to the AMR effort has to be collaborative. We need:

- Novel compounds
- Funding (at the appropriate scale)
- Ready access to appropriate collaborators

# Proposed screening cascade for discovery of new antibacterials

