# Studying allosteric enzyme inhibition using simulated molecular dynamics

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### Outline

- 1 Background Antibiotic resistance Allosteric regulation The shikimate pathway
- 2 The Story So Far MD simulations NeSI's role
- Wrap-Up
- 4 Acknowledgements



## Houston, we have a problem.

### Resistance to antibiotics is growing...

- Mid 1940s: Mass production of penicillin
- Late 1940s: Penicillin-resistant Staphylococcus aureus
- MRSA, gonorrhoea, etc.
- 70+ years of whack-a-mole...only worse.

### The search for new antibiotics

- Try to block a bacterial enzyme
- We look for pathways not found in animals



# **Enzyme Inhibition**

### Competitive inhibition

- Binding in the active site
- Competes with substrate for room

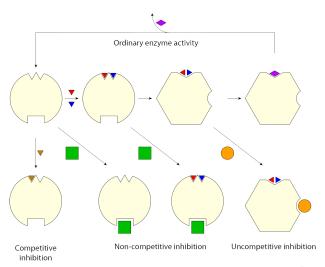
### Uncompetitive, non-competitive, or mixed inhibition

- Does not compete with substrate for binding, but:
- Reduces affinity of enzyme for substrate, catalytic activity, or both
- Often involves binding at another site ("allosteric inhibition")

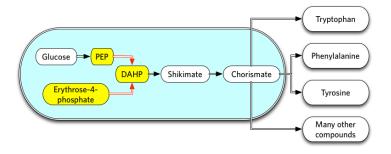


Background

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# Aromatic Amino Acid Biosynthesis

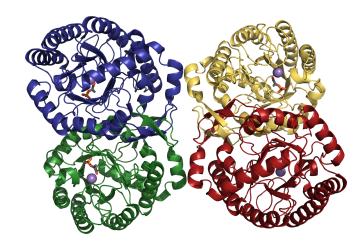


### The shikimate pathway

- Bacteria, fungi, plants, etc., but not humans or animals
- First enzyme: DAHP synthase (step in yellow and red)



# DAHP Synthase





# What can we study?

### Molecular dynamics simulations of:

- Enzymes from different bacterial species
- Wild-type vs various mutants
- Open vs closed
- Presence vs absence of inhibitor



### Simulations so far:

### Open vs closed (T. maritima)





### Presence vs absence of tyrosine

- Conformation and dynamics
- Free-energy changes



# How NeSI has helped

### Computers

- 1 Computing power on BlueGene/P and POWER 7
- 2 Software: NAMD (simulations), Amber (post-processing)

### Expertise

- Installing and configuring Amber
- 2 Expert MD advice
- 3 Avoiding the "waste-of-life" file



### To Summarise...

Background

- Hundreds of ns of time simulated
- Amber has been deployed on POWER 7
- Closed form is stable when simulated
- Open form is extremely dynamic
- Does tyrosine cause closure; if so, how?
- Next step: Free energy calculations

# Acknowledgements

- Eric Lang
- **Emily Parker**
- François Bissey

# Questions & Answers

