

# ARTIFICIAL INTELLIGENCE FOR ANTIBIOTIC DISCOVERY

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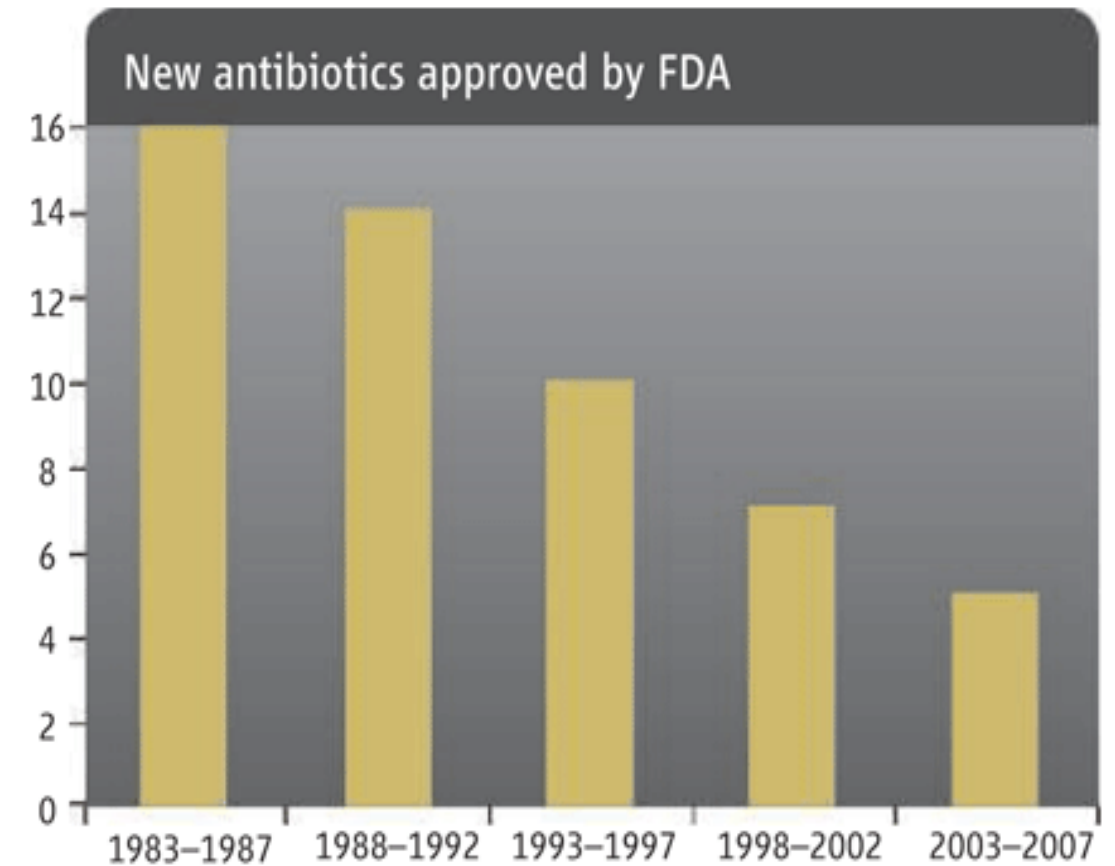
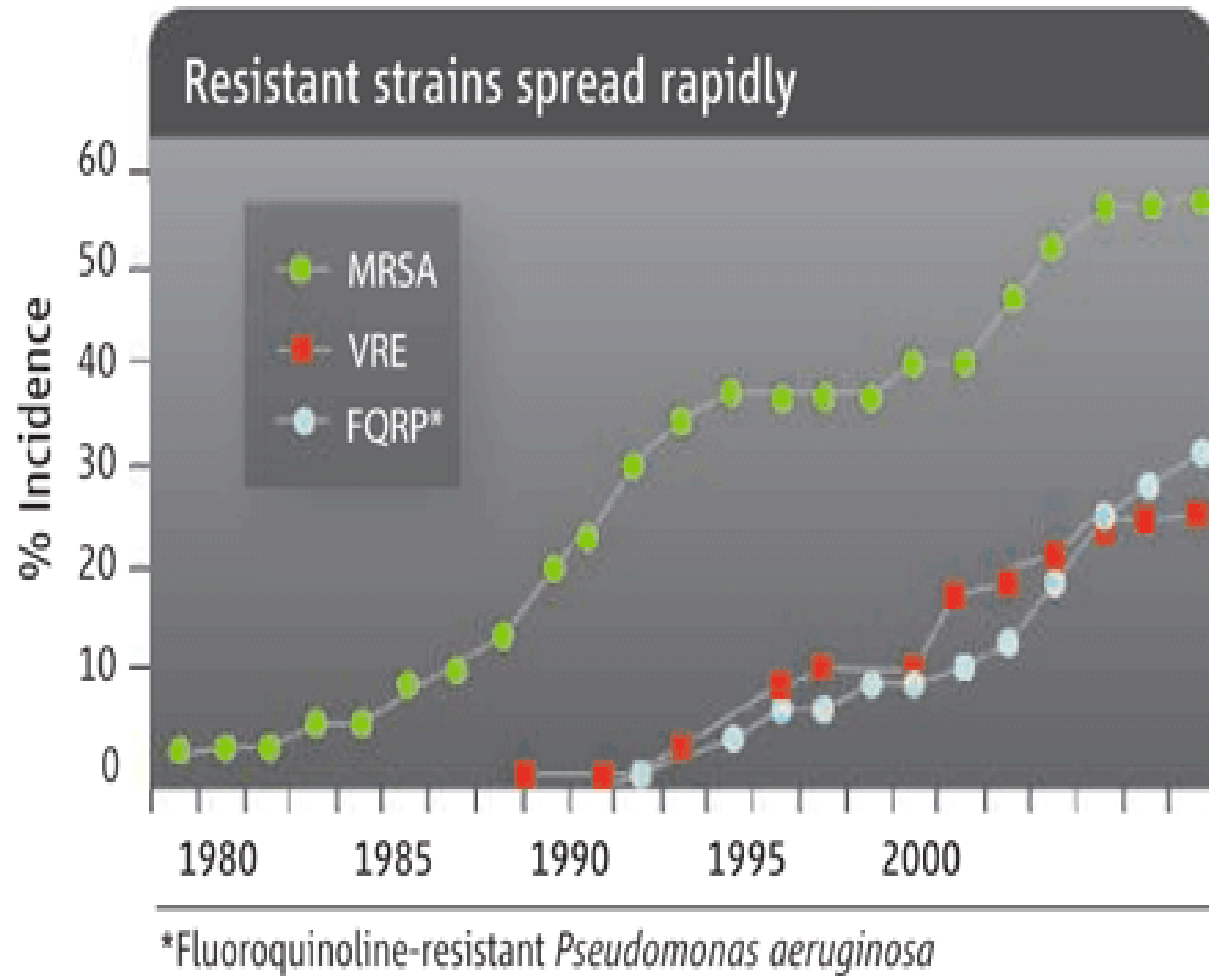
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WYSS INSTITUTE FOR BIOLOGICALLY INSPIRED ENGINEERING  
HARVARD UNIVERSITY

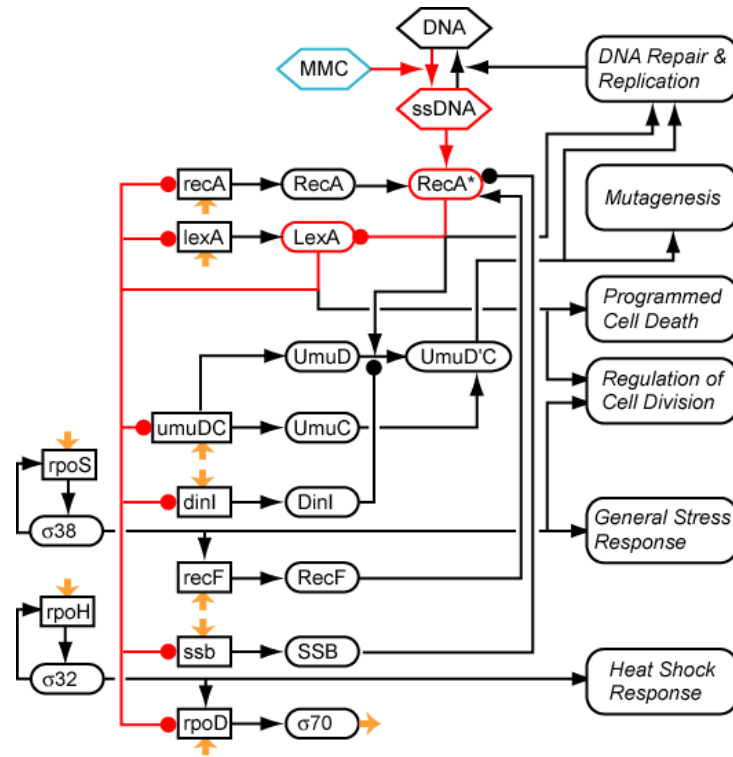
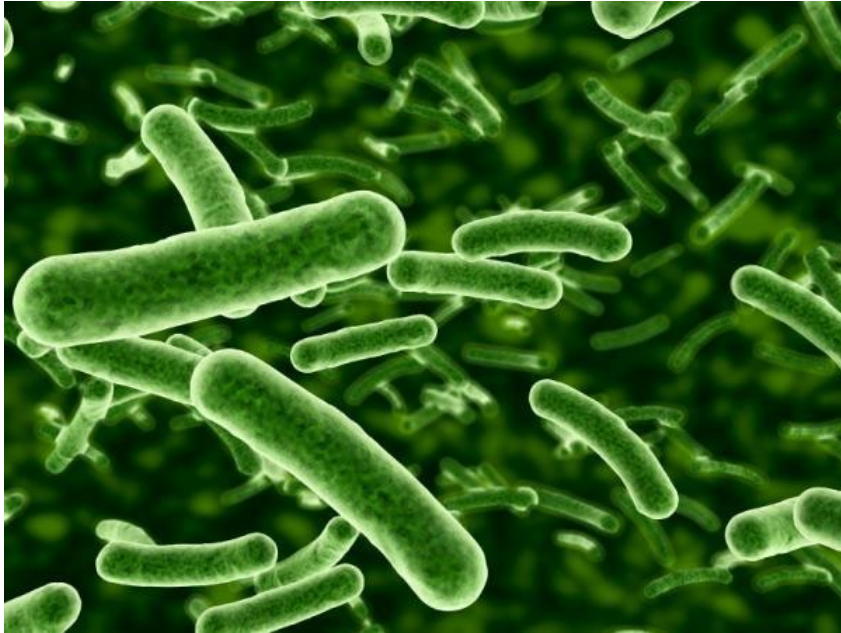


Massachusetts  
Institute of  
Technology

# MORE RESISTANCE, FEWER NEW ANTIBIOTICS



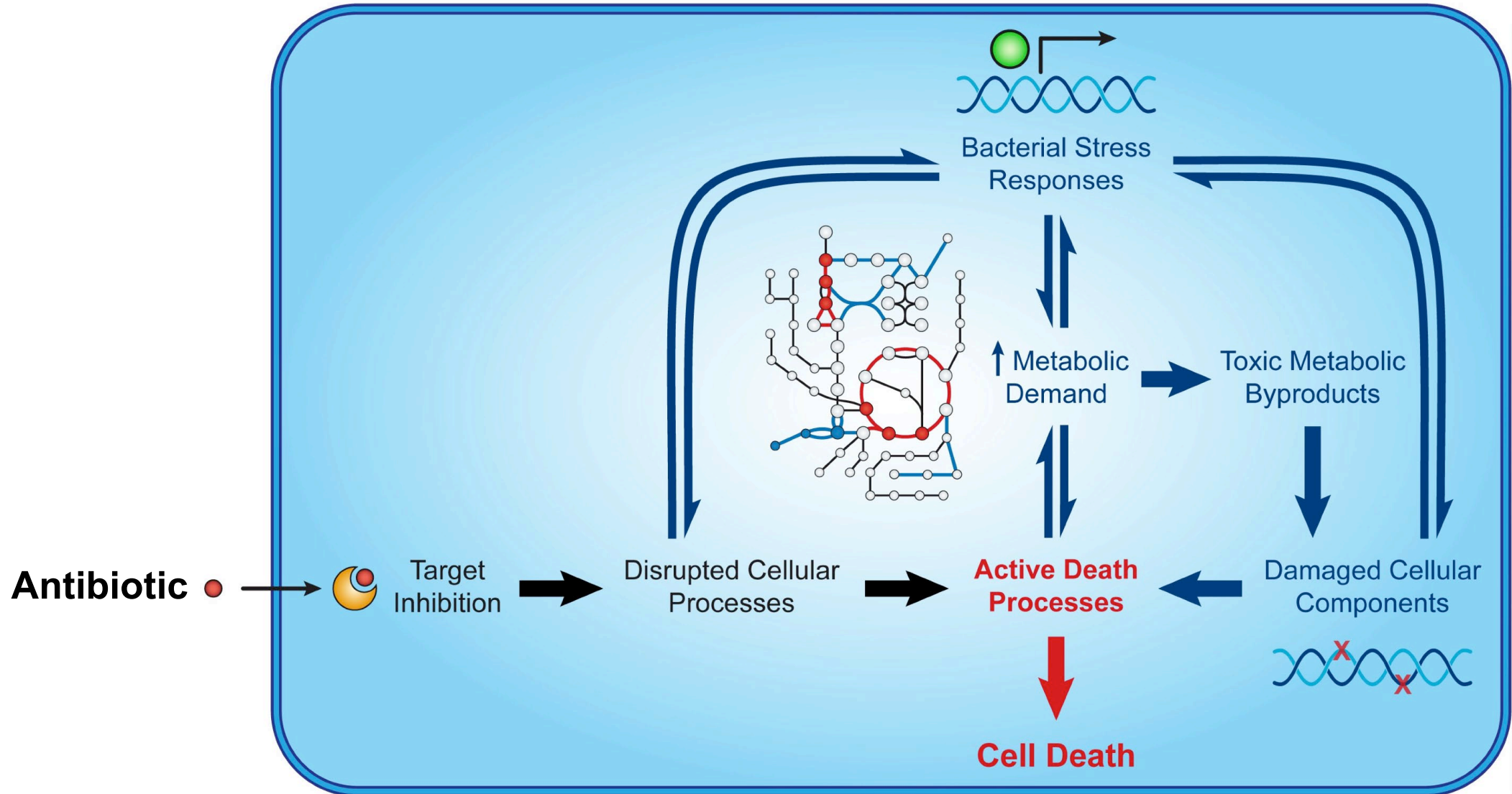
# MACHINE LEARNING AND ANTIMICROBIAL RESEARCH



*TS Gardner et al., Science, 2003*

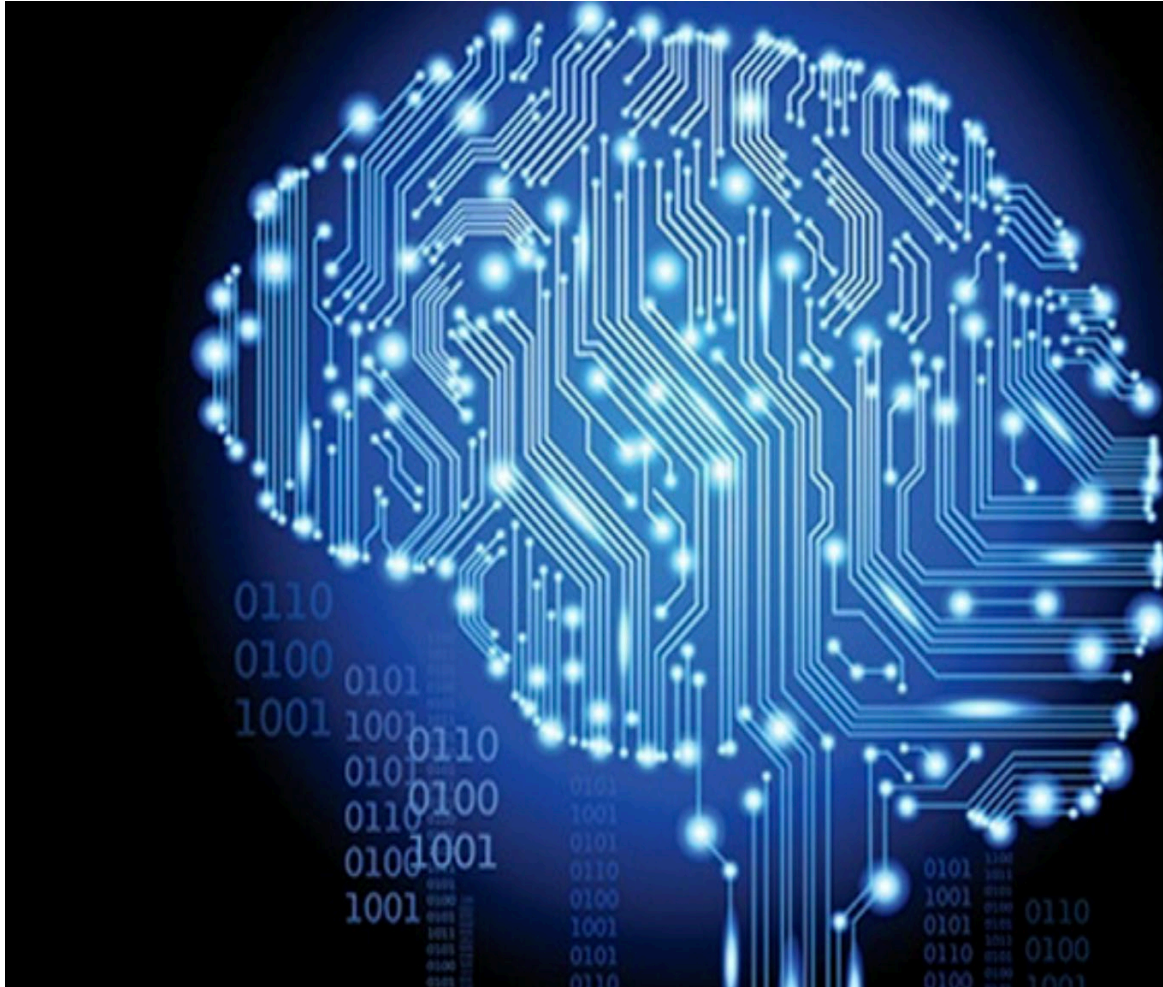
*D di Bernardo et al. Nature Biotechnology 2005*

# HOW ANTIBIOTICS KILL BACTERIA: AN EXPANDED VIEW

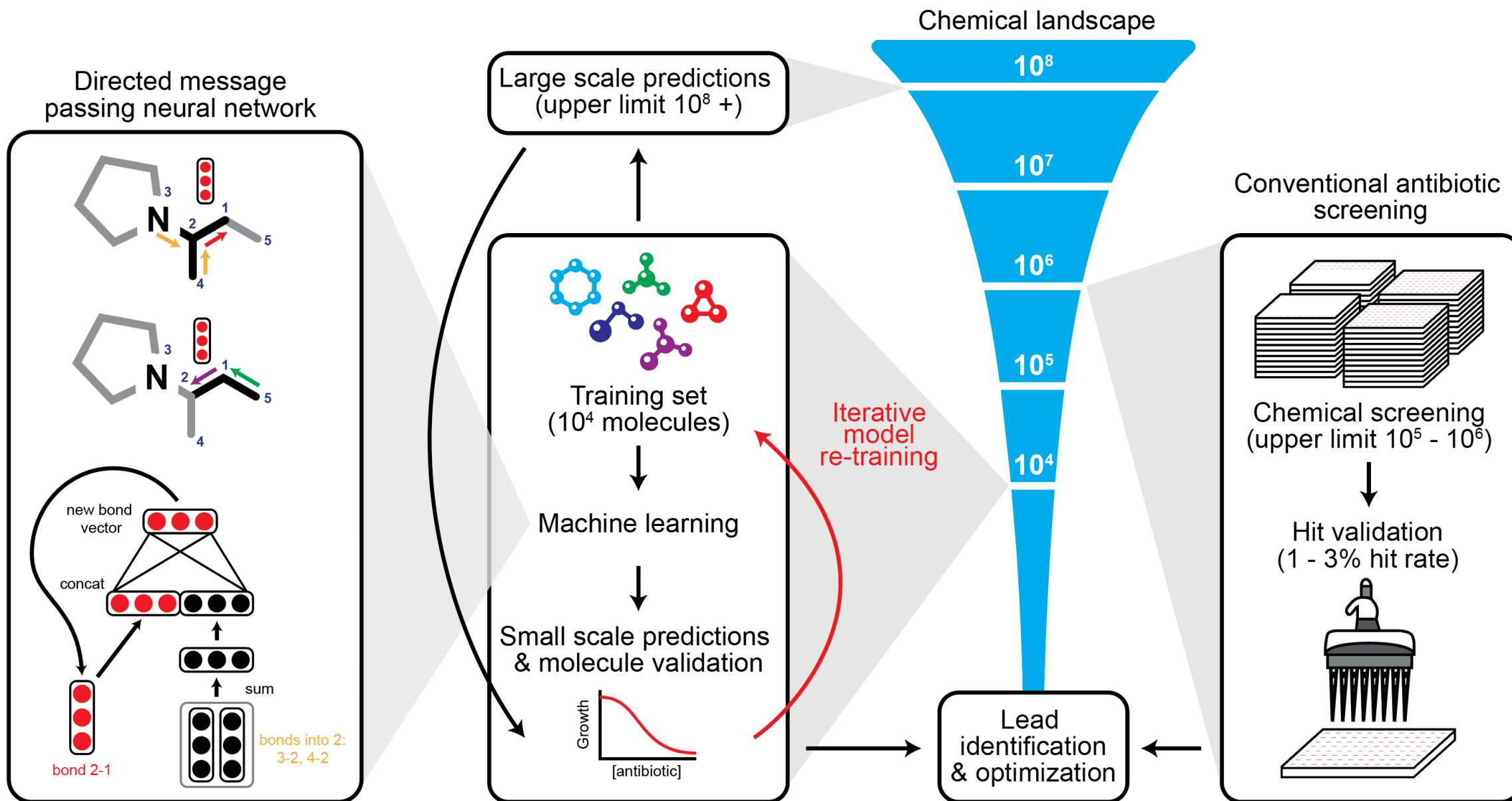




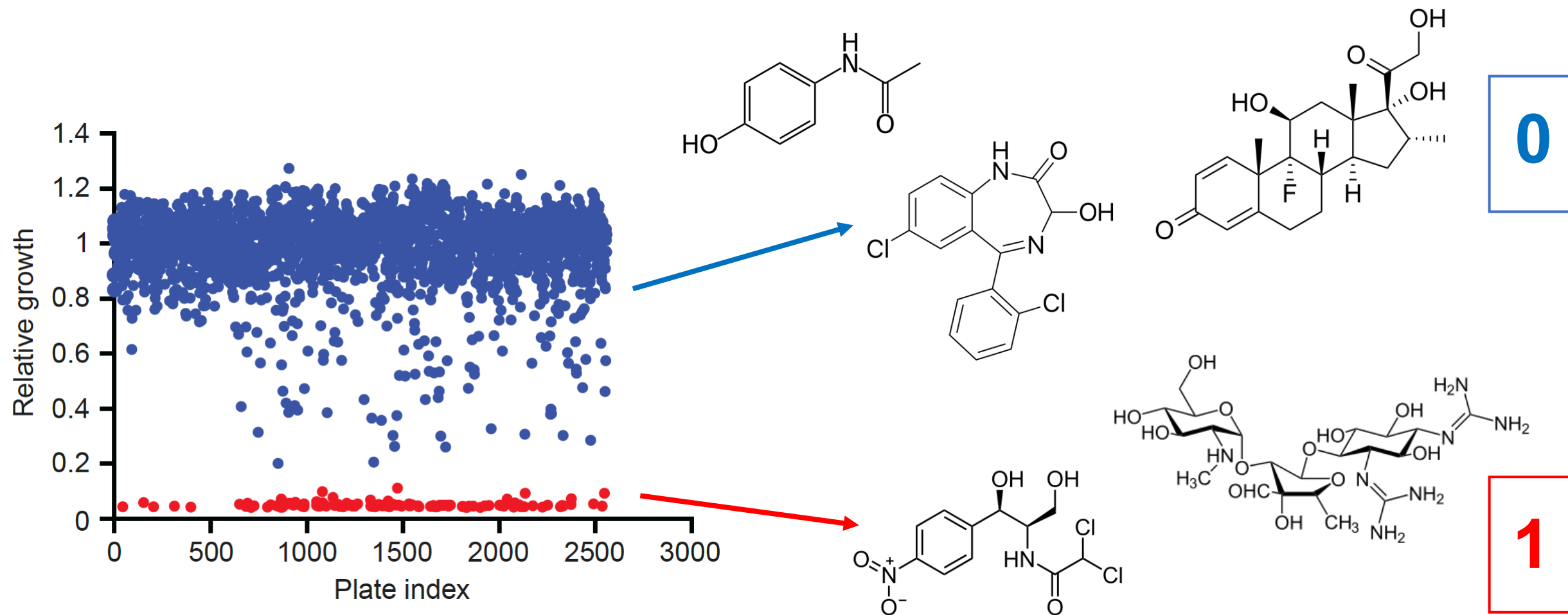
# DEEP LEARNING & BIG DATA: BIOTECH & MEDICINE



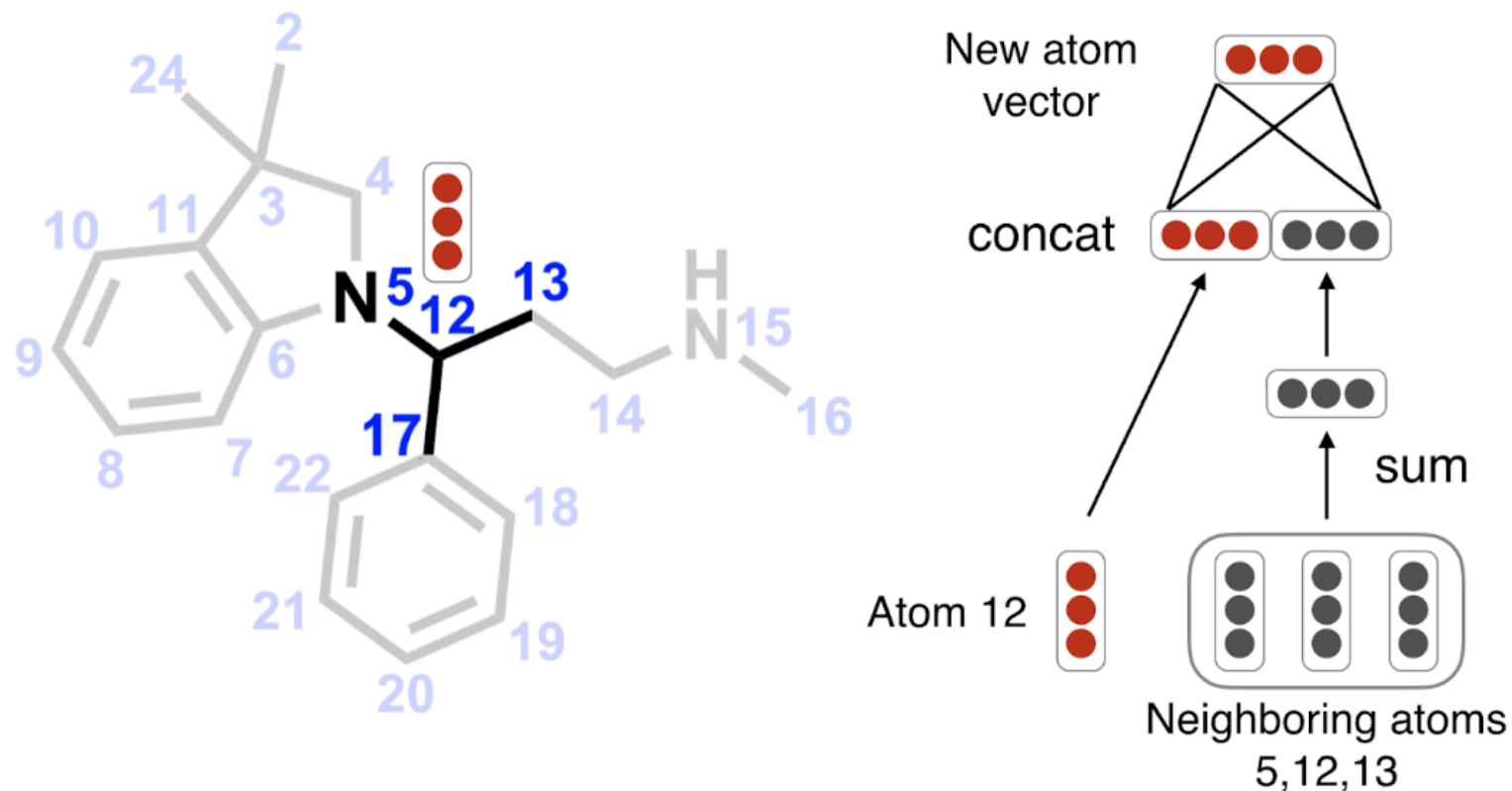
# DEEP LEARNING APPROACH TO ANTIBIOTIC DISCOVERY



# TRAINING DATA SET: GROWTH INHIBITION AGAINST *E. COLI*



# DEEP NEURAL NETWORK FOR MOLECULAR PROPERTY PREDICTION

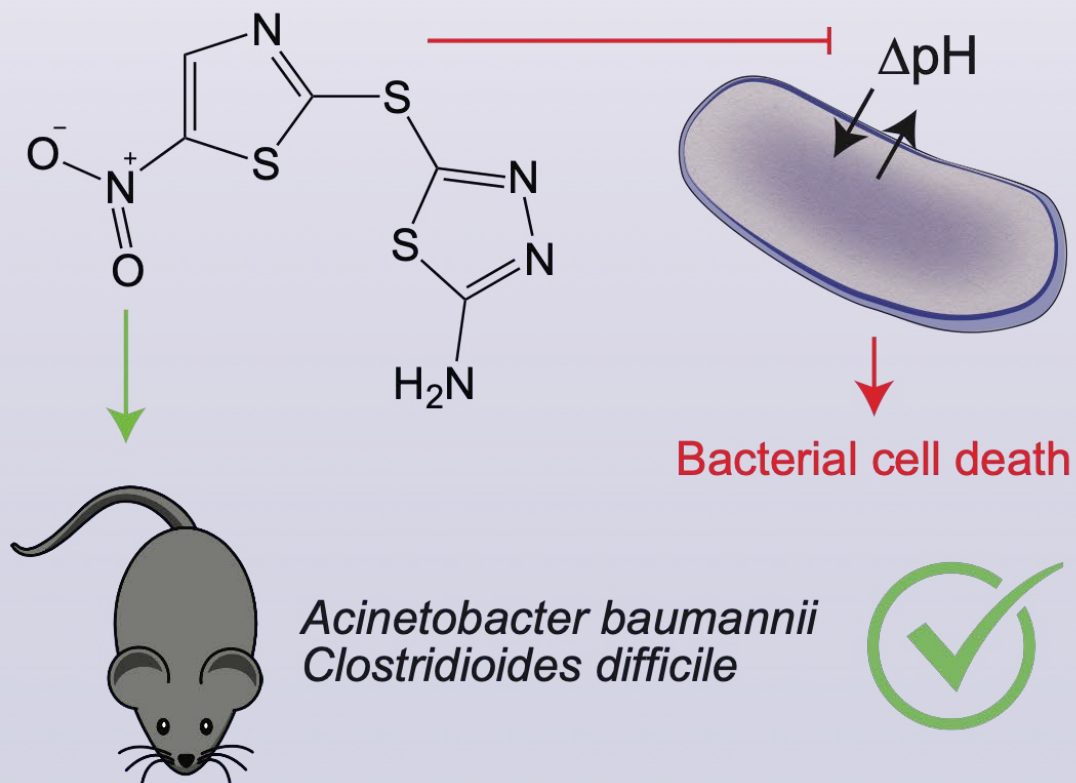


Message Passing Neural Network: train on a dataset containing molecules along with known property values for each molecule

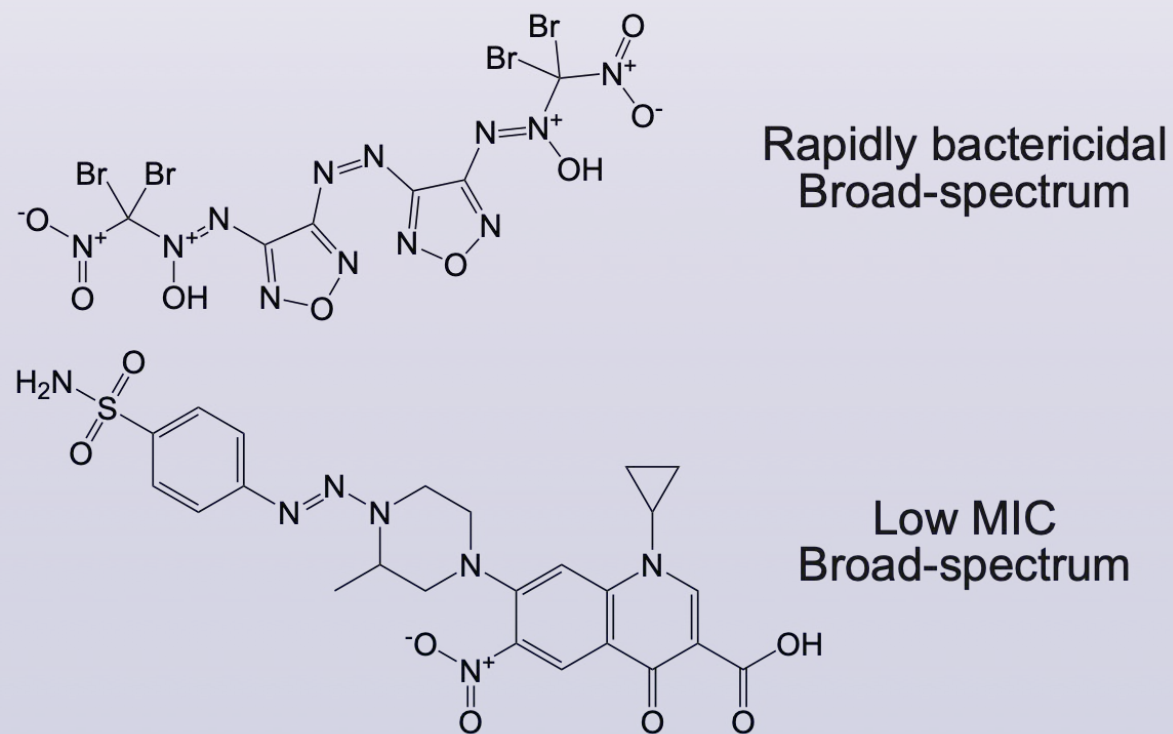


# DEEP LEARNING APPROACH TO ANTIBIOTIC DISCOVERY

## Drug Repurposing Hub HALICIN

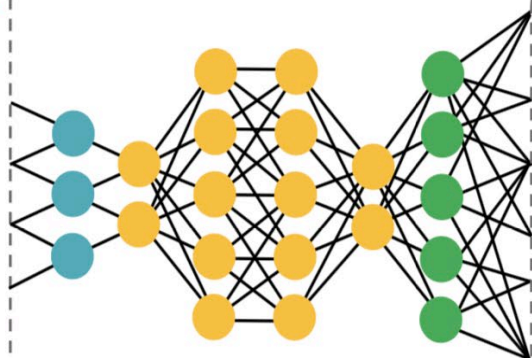
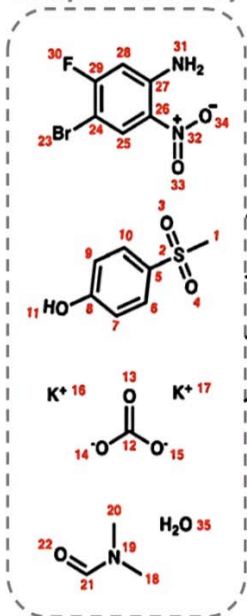


## ZINC15 Database

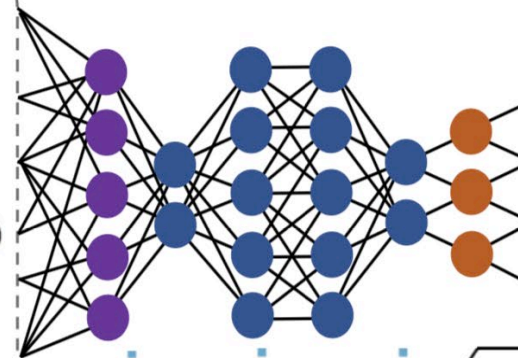
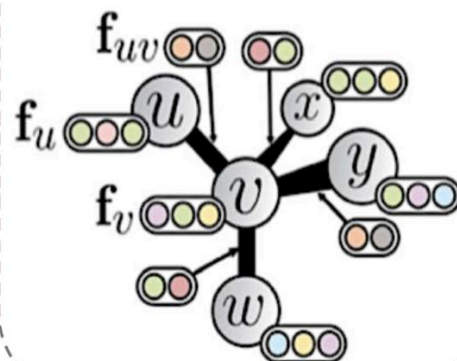


# AI-BASED DISCOVERY AND DESIGN OF NOVEL ANTIBIOTICS

Large chemical  
compound library



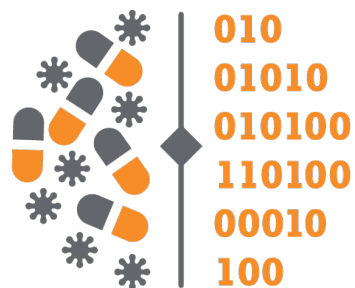
Learned molecular  
representations



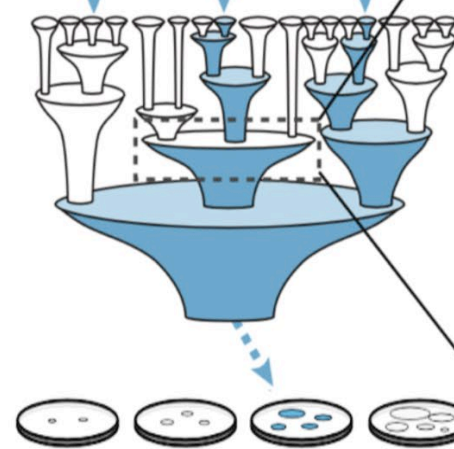
Novel antibiotic  
prediction

Feature-based  
molecular property  
optimization

“White box”  
machine learning to  
elucidate antibiotic  
mechanism of action

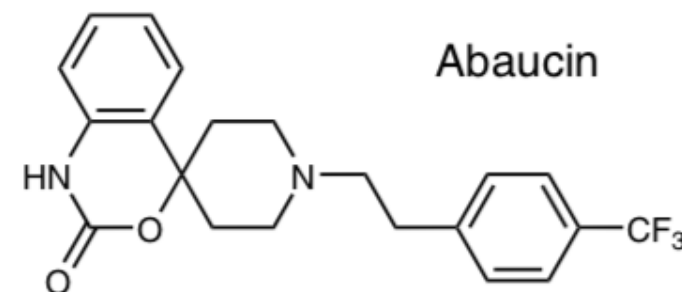
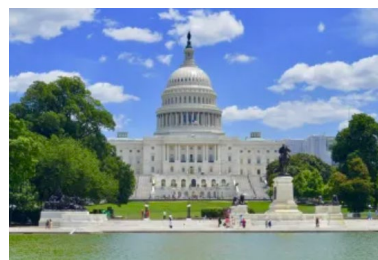


**Antibiotics-AI Project**



Predicted antibiotic efficacy

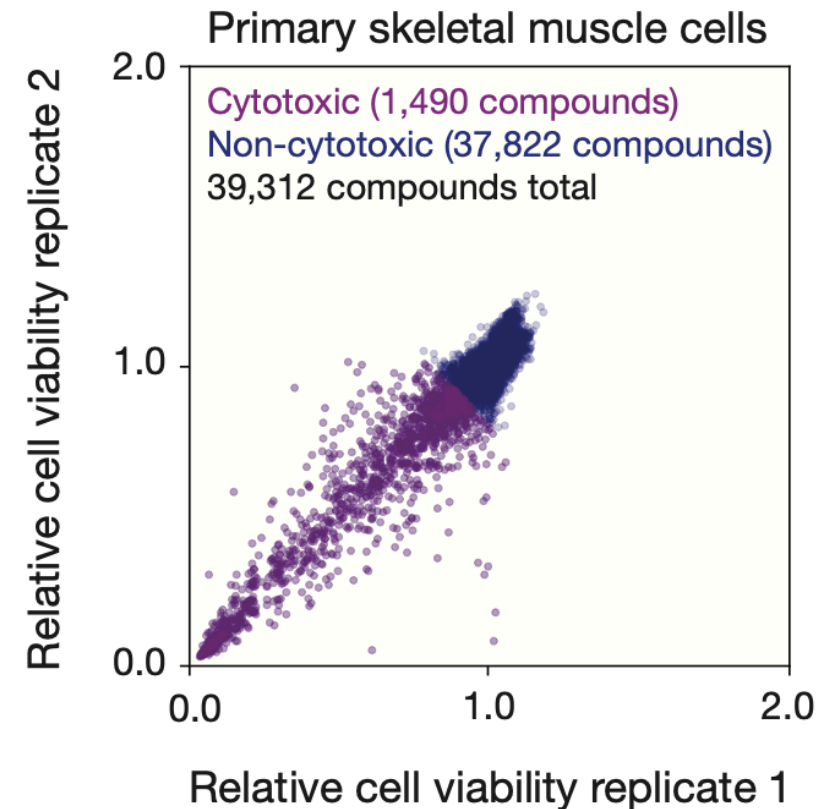
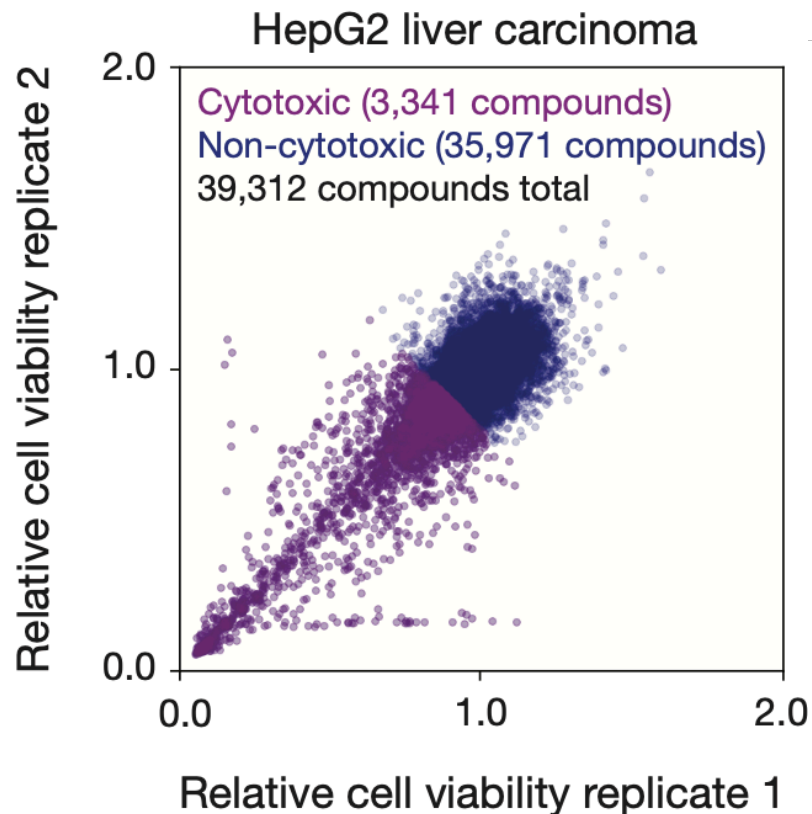
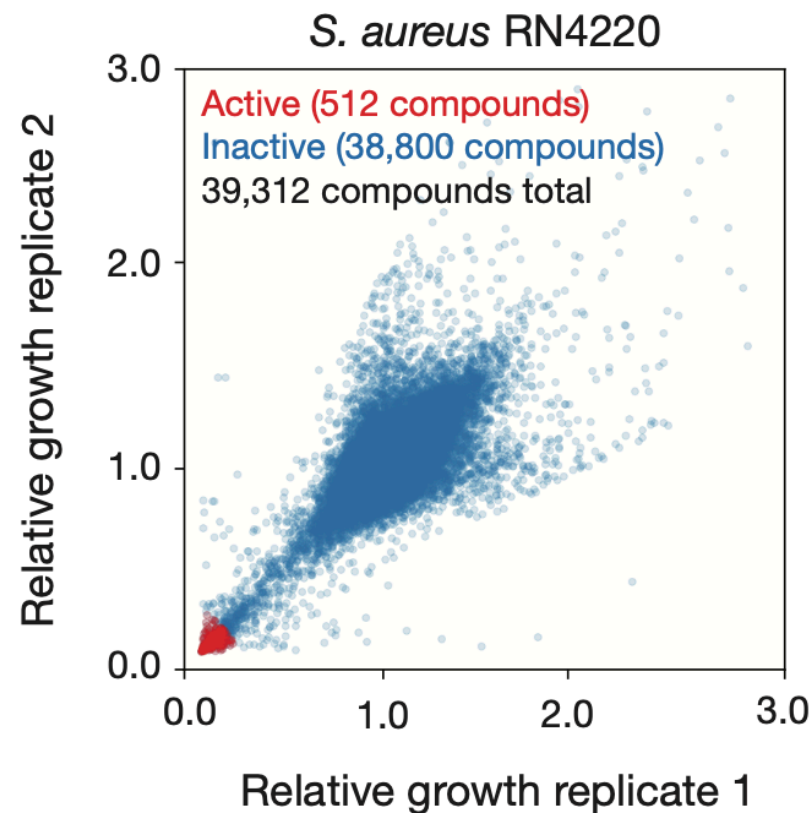
# AI-BASED DISCOVERY OF ABAUCIN



*G Liu et al., Nature Chemical Biology, 2023*

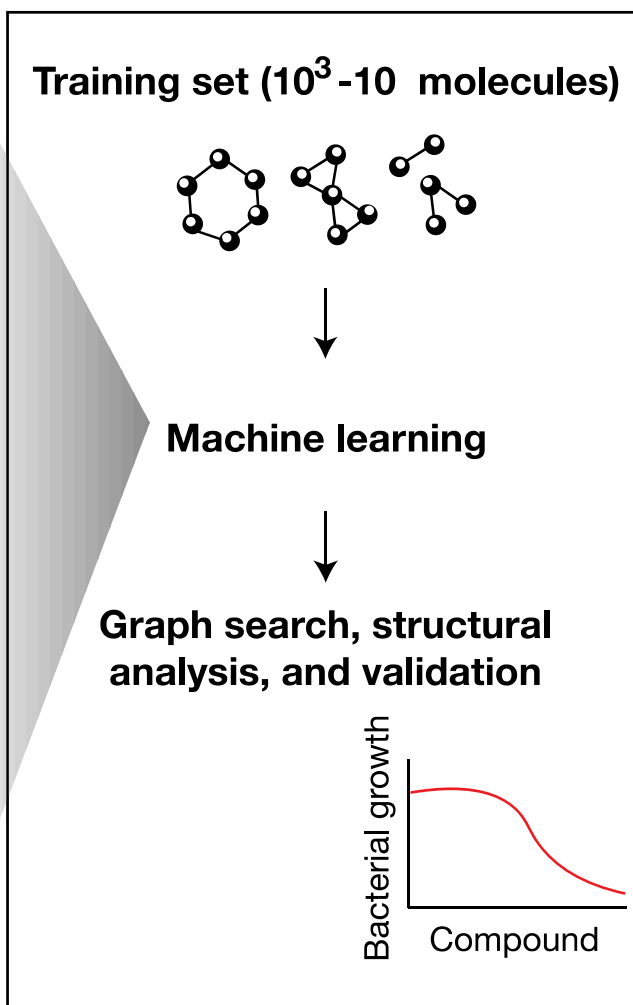
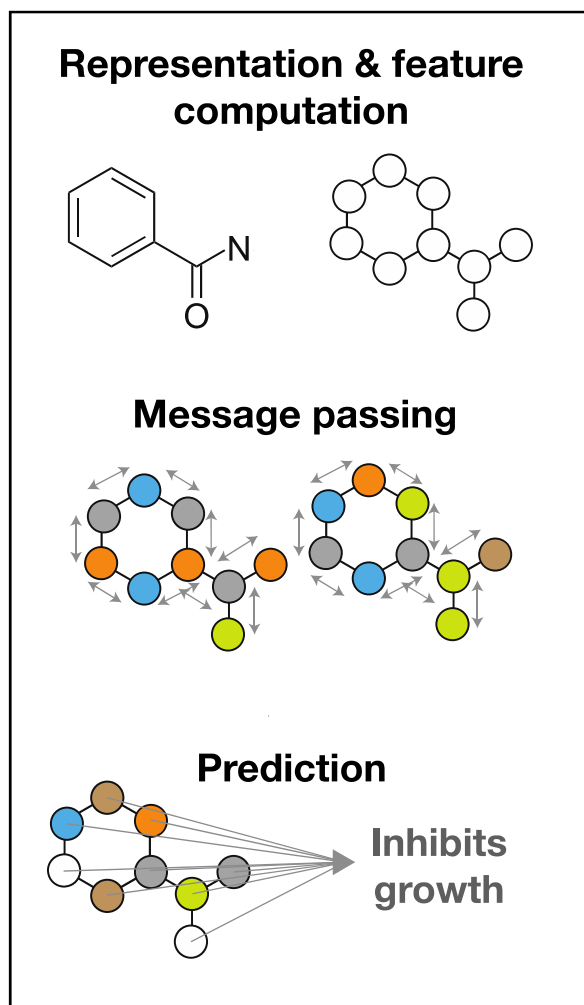


# DEEP LEARNING FOR ANTIMICROBIAL ACTIVITY AND CYTOTOXICITY





# EXPLAINABLE DL FOR STRUCTURAL CLASSES OF ANTIMICROBIALS



## Chemical landscape

$>10^9$

Upper limit of deep neural network predictions

$\sim 10^8$

Number of molecules in large compound database

$\sim 10^3$

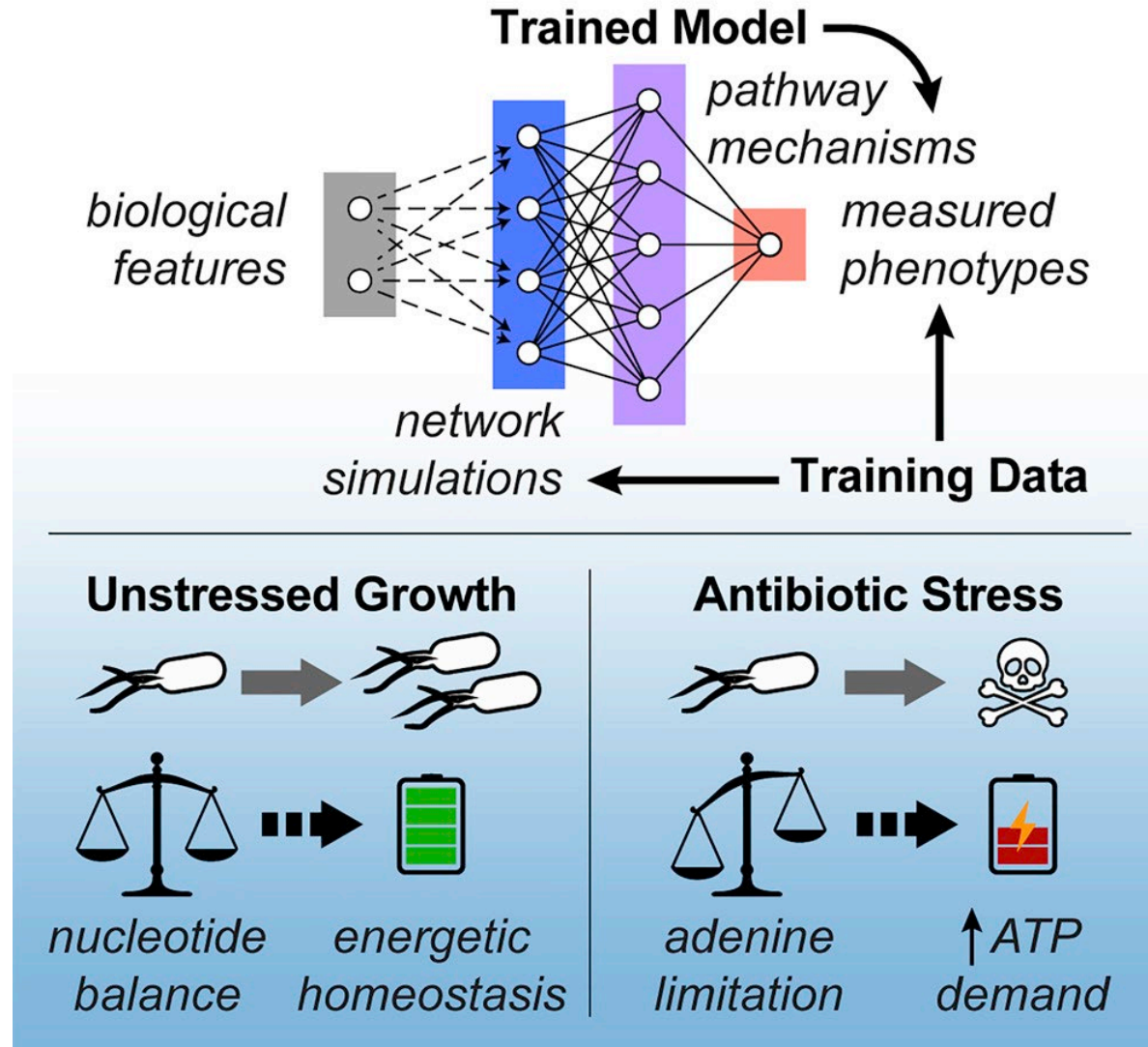
Number of strong predicted hits

$\sim 10^2$

Number of test molecules screened

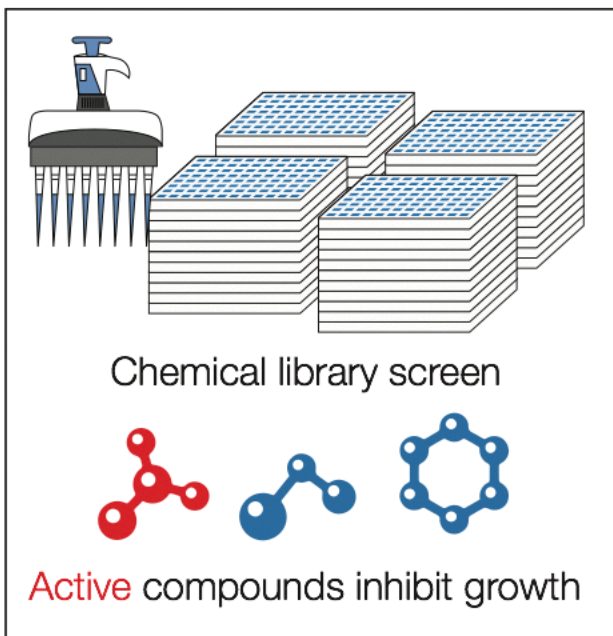
# WHITE-BOX MACHINE LEARNING: IDENTIFYING ANTIBIOTIC MoA

## White-Box Machine Learning

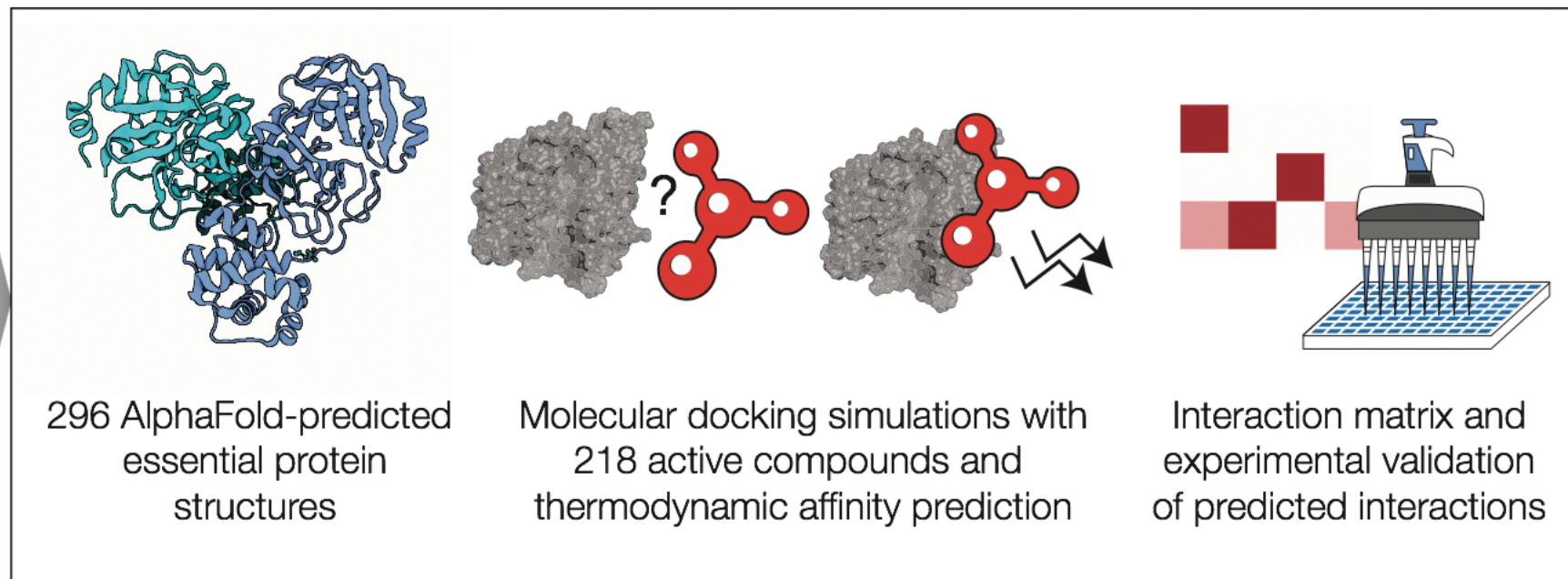


# ANTIBIOTICS-AI PROJECT: LEVERAGING ALPHAFOLD

## Growth inhibition screening



## AlphaFold-enabled identification of protein-ligand interactions



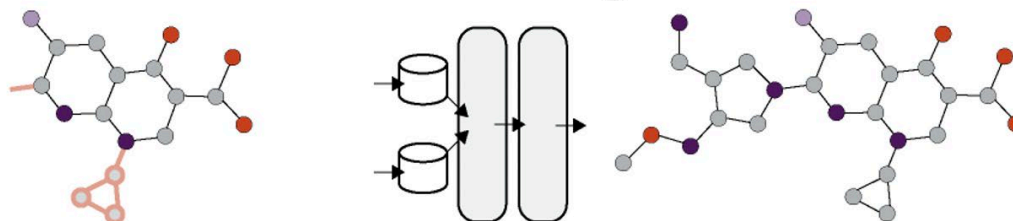
# GENERATIVE AI FOR *DE NOVO* ANTIBIOTIC DESIGN

## Comprehensive fragment search



Graph neural network model scoring

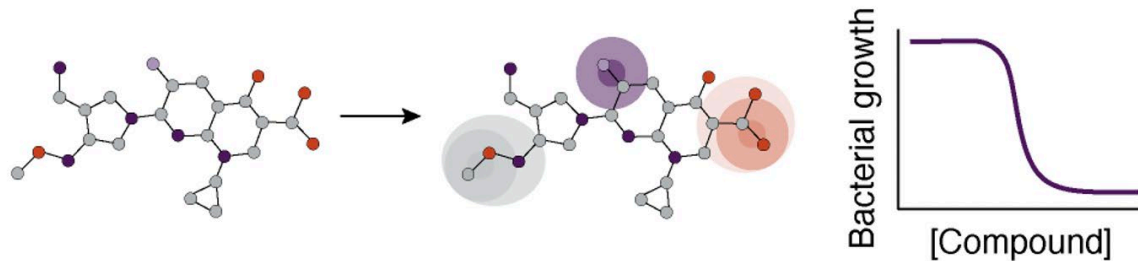
## Generative design



Chemical mutations

Variational autoencoder

## Synthesis and validation of generated compounds





# ANTIBIOTICS-AI PROJECT AND PHARE BIO



**Antibiotics-AI Project**

**TED**

**PHAREBIO**