

ARTIFICIAL INTELLIGENCE FOR ANTIBIOTIC DISCOVERY

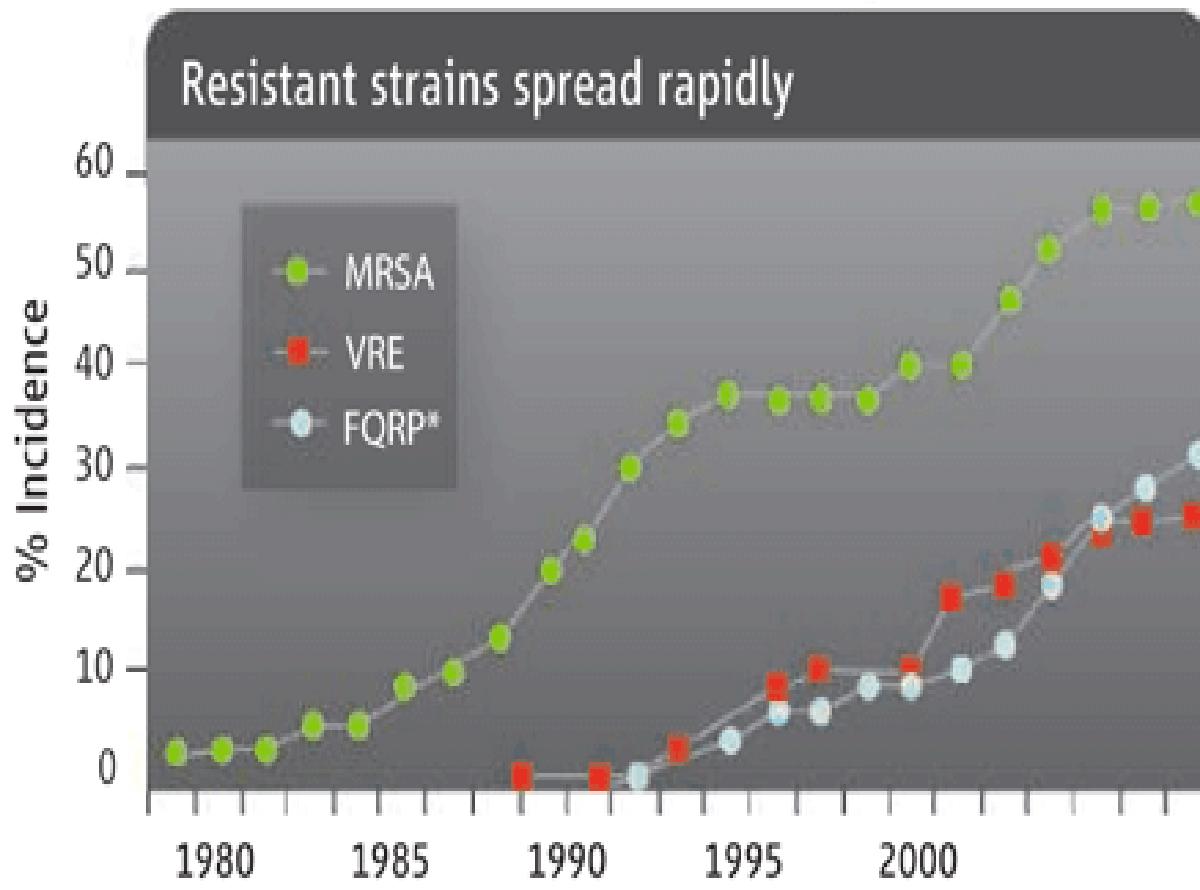
JAMES J. COLLINS

INSTITUTE FOR MEDICAL ENGINEERING & SCIENCE
DEPARTMENT OF BIOLOGICAL ENGINEERING
MASSACHUSETTS INSTITUTE OF TECHNOLOGY

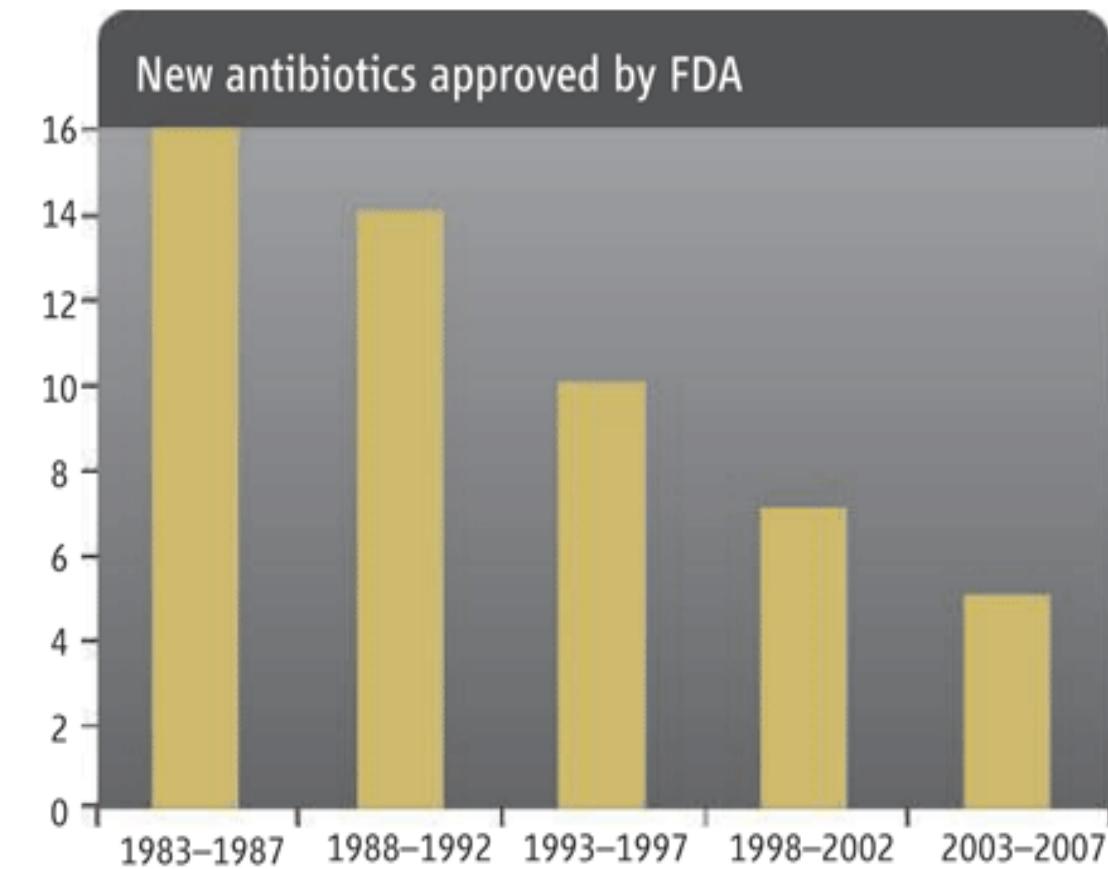
BROAD INSTITUTE OF HARVARD AND MIT

WYSS INSTITUTE FOR BIOLOGICALLY INSPIRED ENGINEERING
HARVARD UNIVERSITY

MORE RESISTANCE, FEWER NEW ANTIBIOTICS

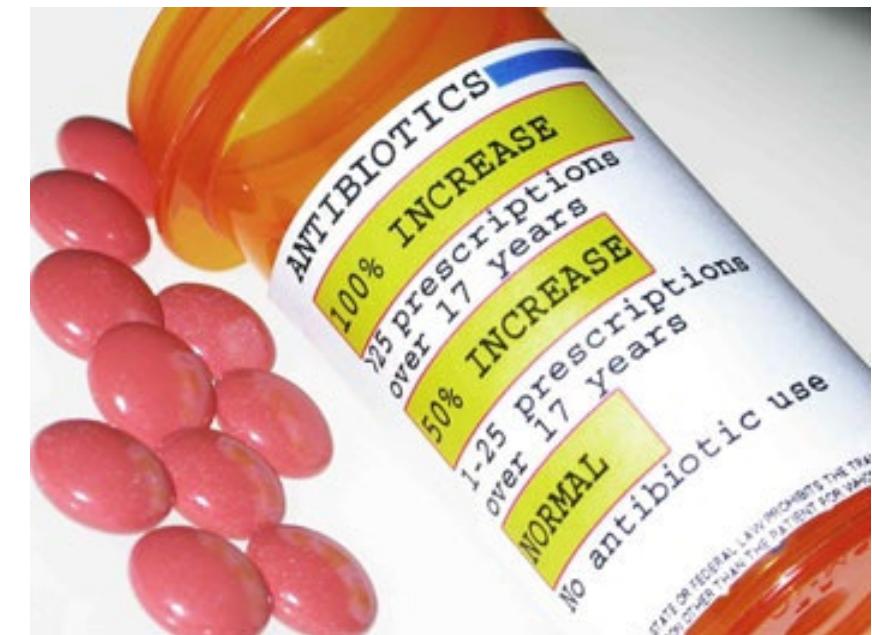
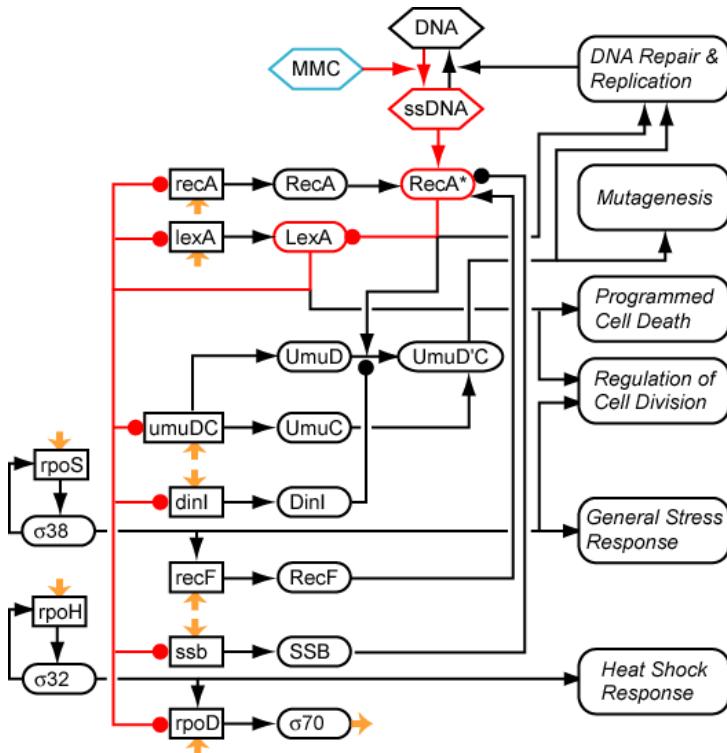
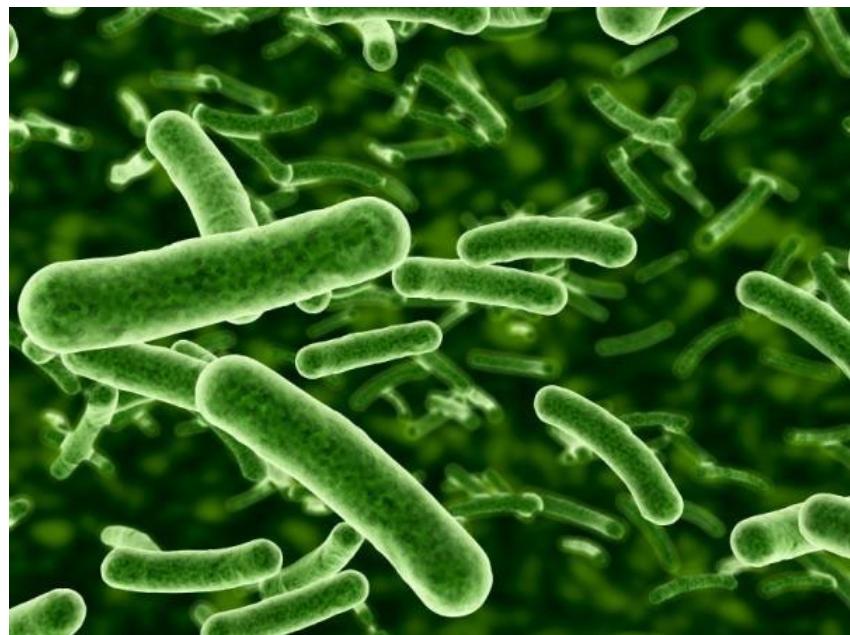


*Fluoroquinolone-resistant *Pseudomonas aeruginosa*



Taubes, Science, 2008

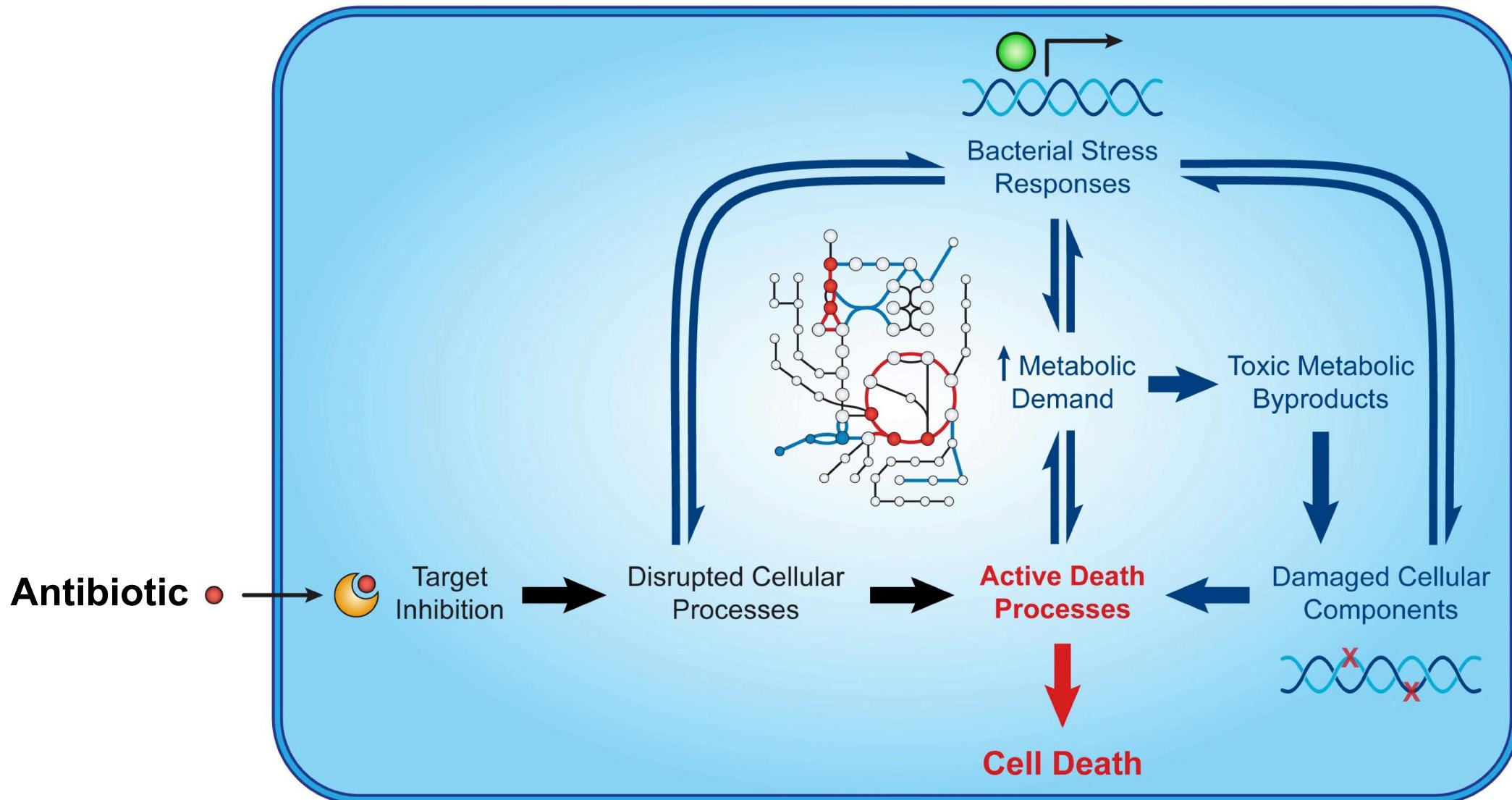
MACHINE LEARNING AND ANTIMICROBIAL RESEARCH



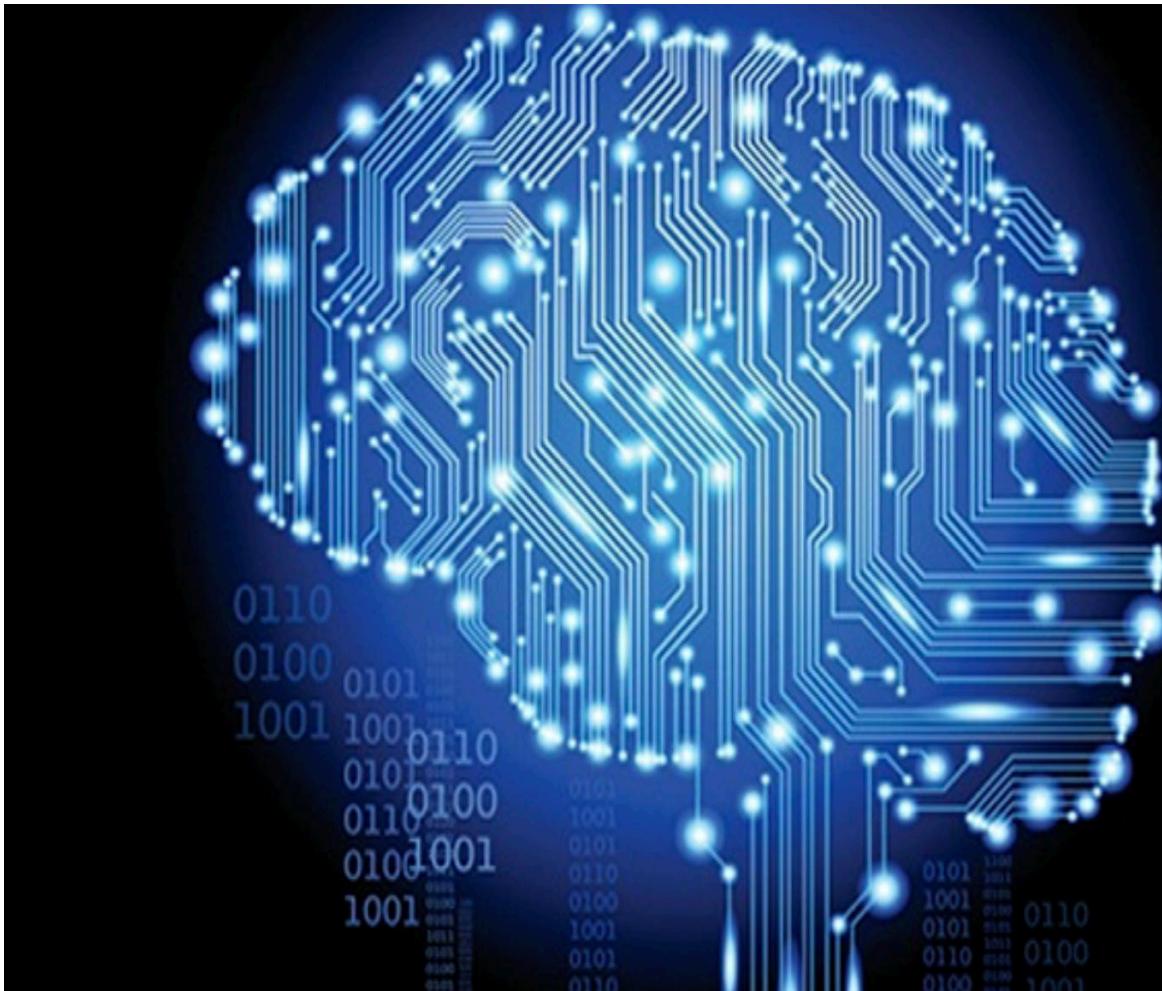
TS Gardner et al., *Science*, 2003

D di Bernardo et al. *Nature Biotechnology* 2005

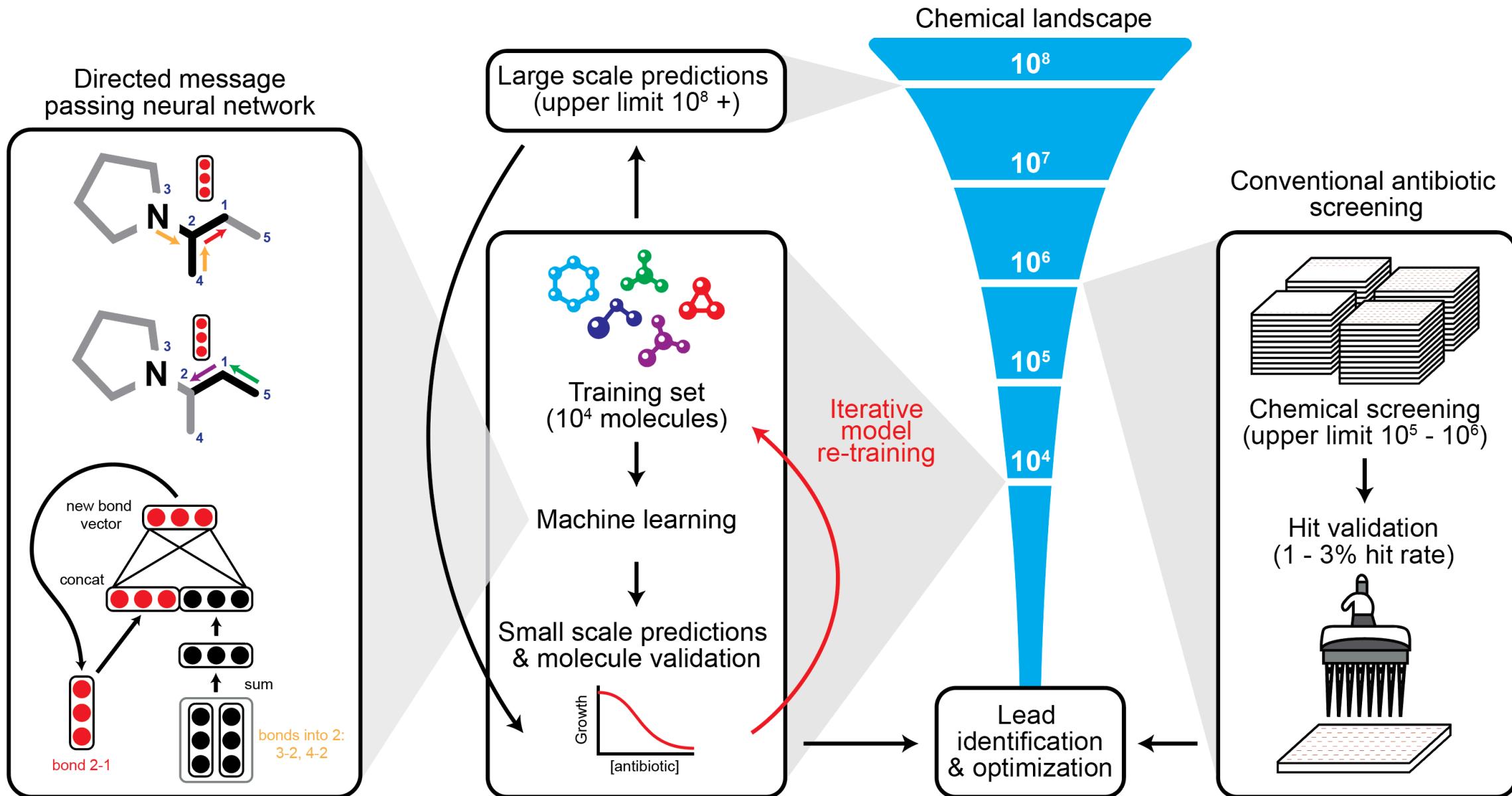
How ANTIBIOTICS Kill BACTERIA: An EXPANDED VIEW



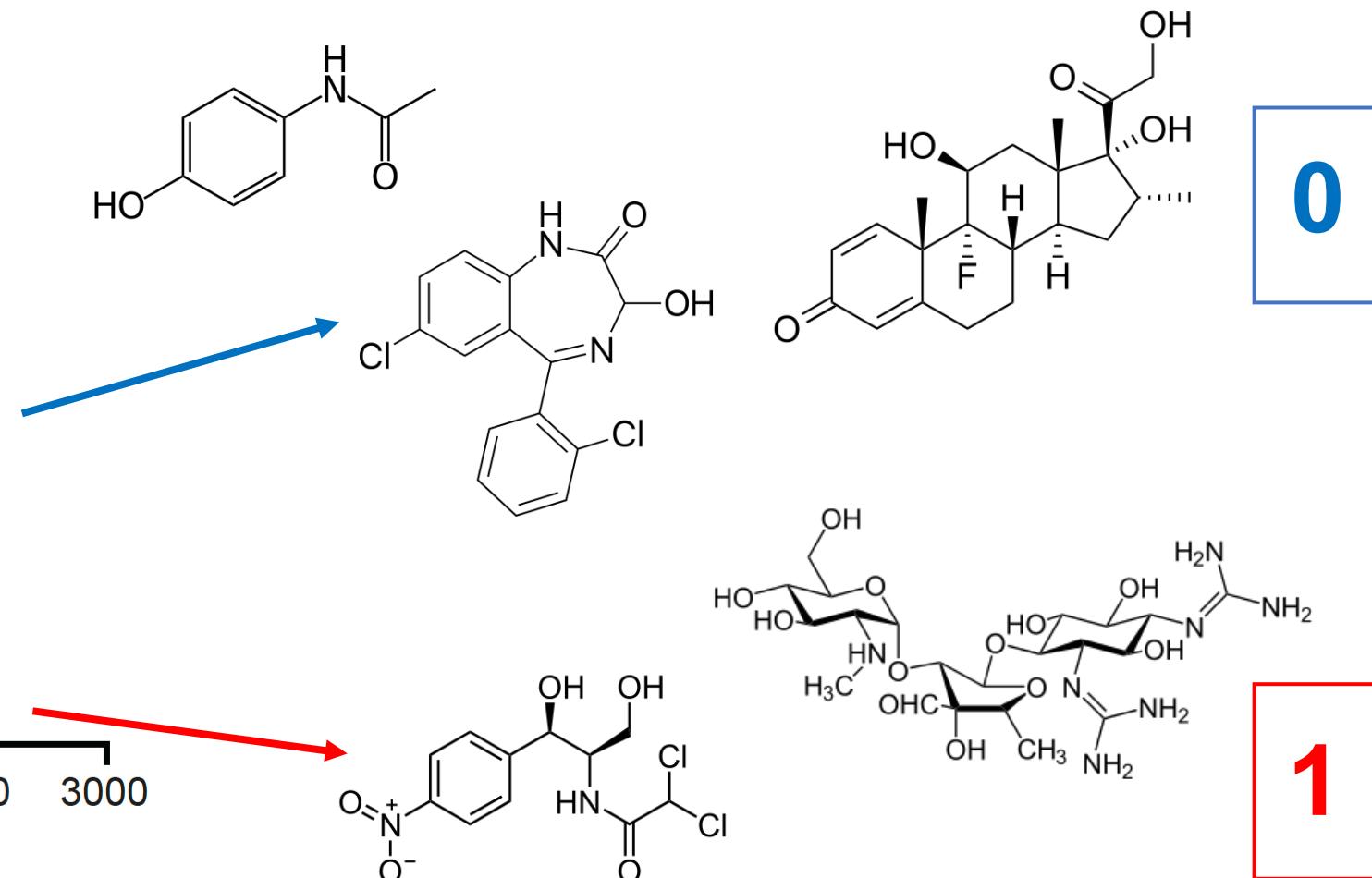
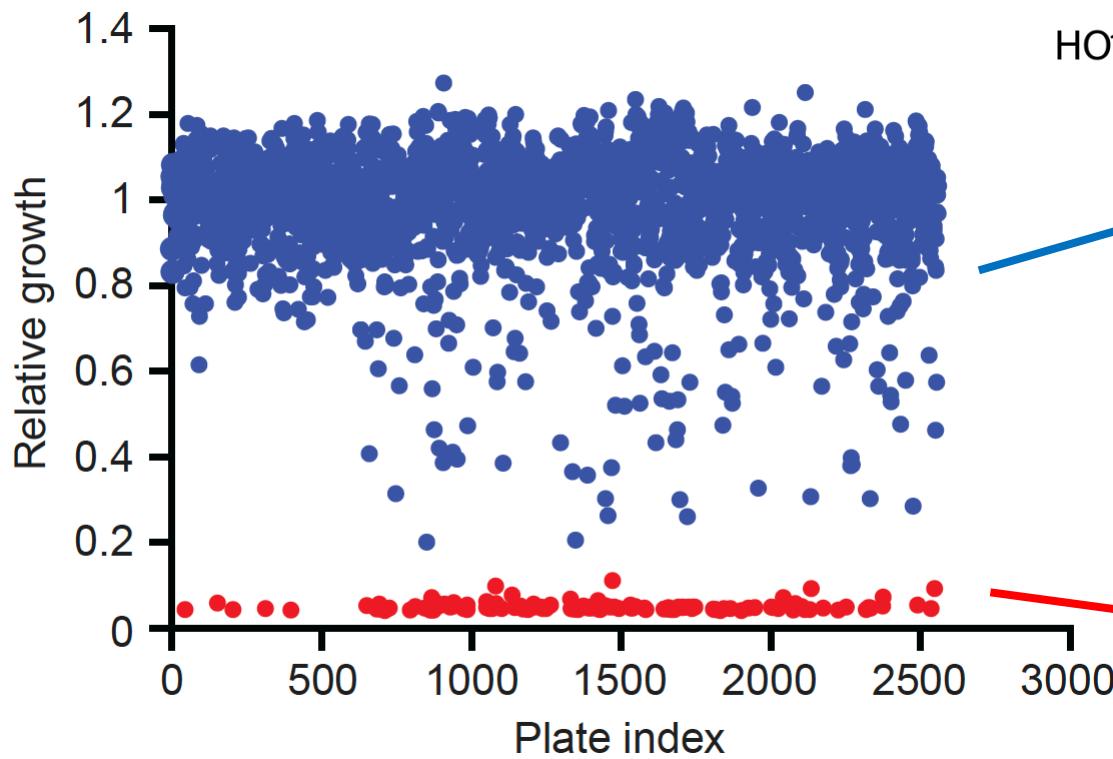
DEEP LEARNING & BIG DATA: BIOTECH & MEDICINE



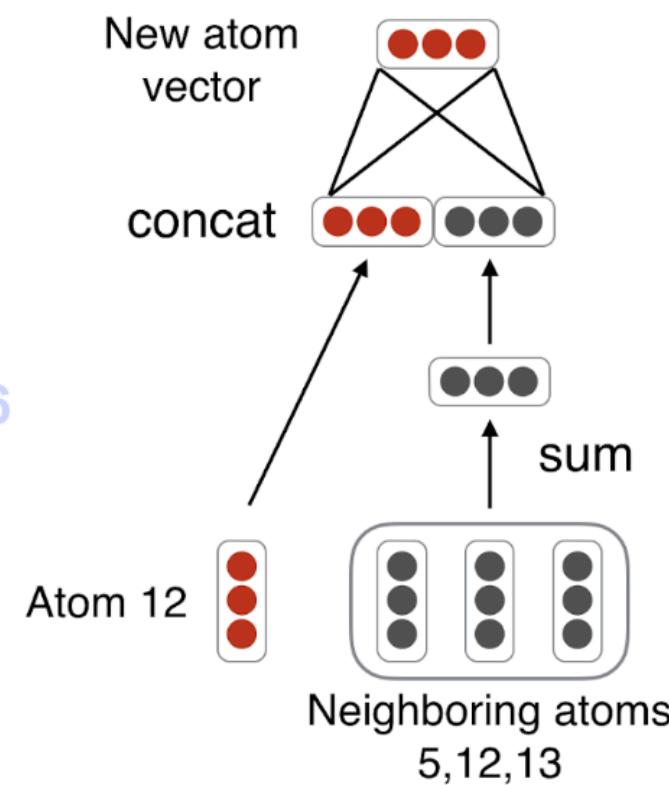
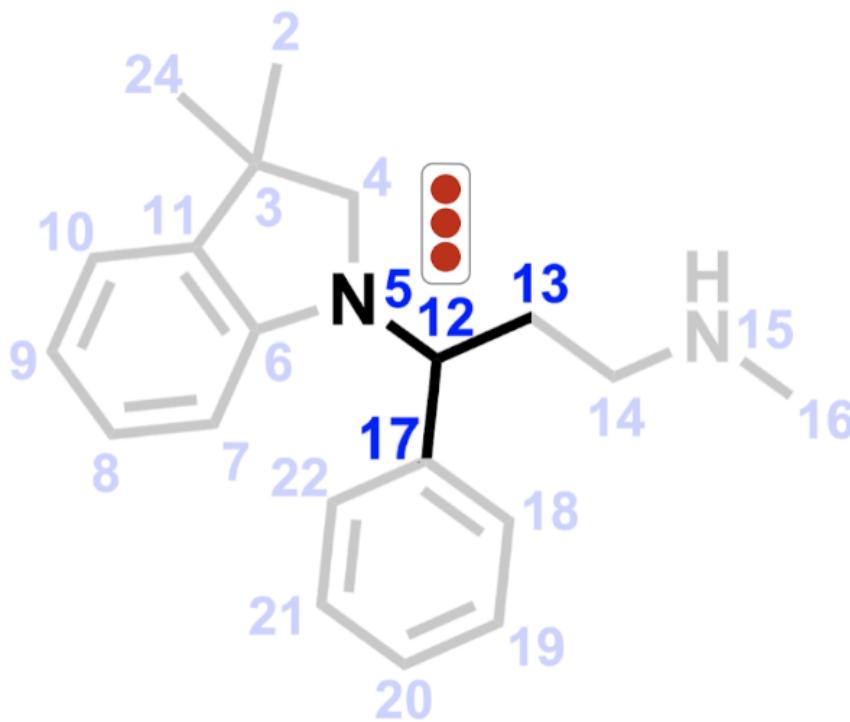
DEEP LEARNING APPROACH TO ANTIBIOTIC DISCOVERY



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

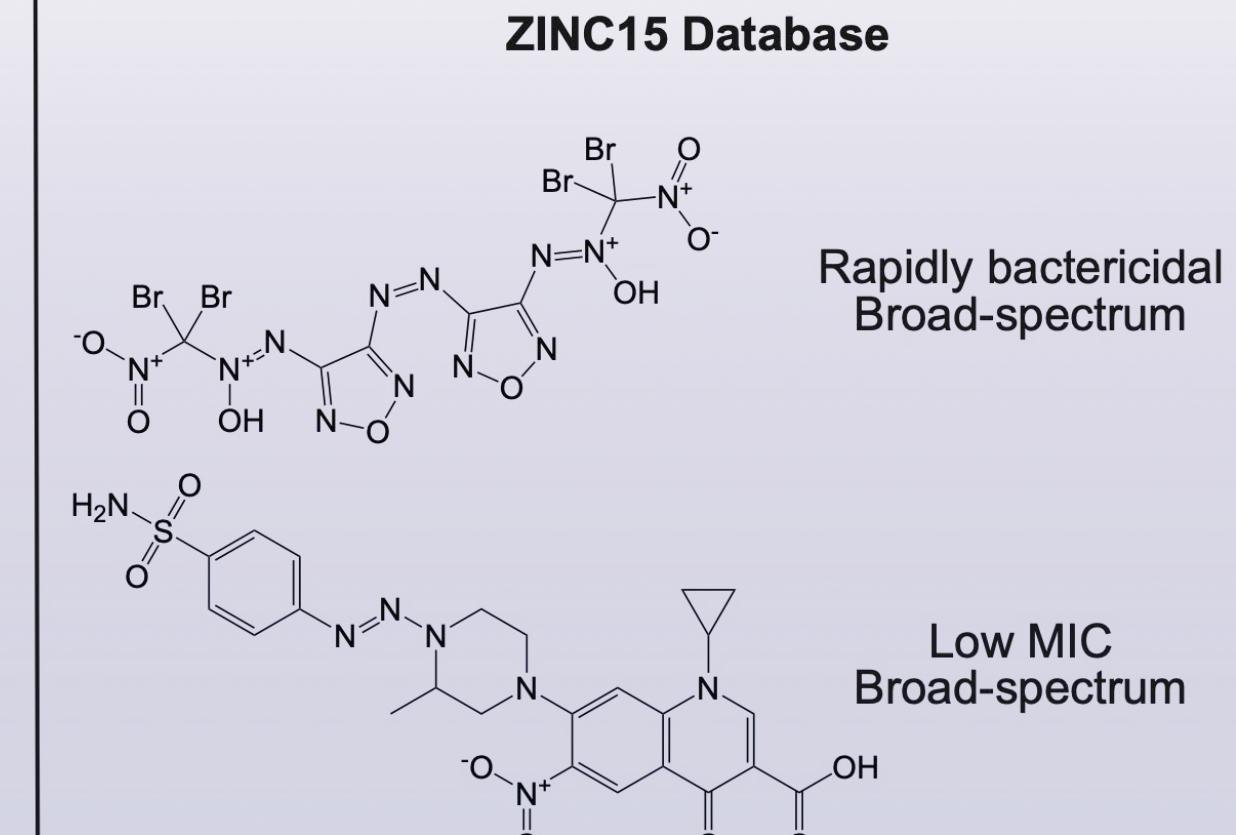
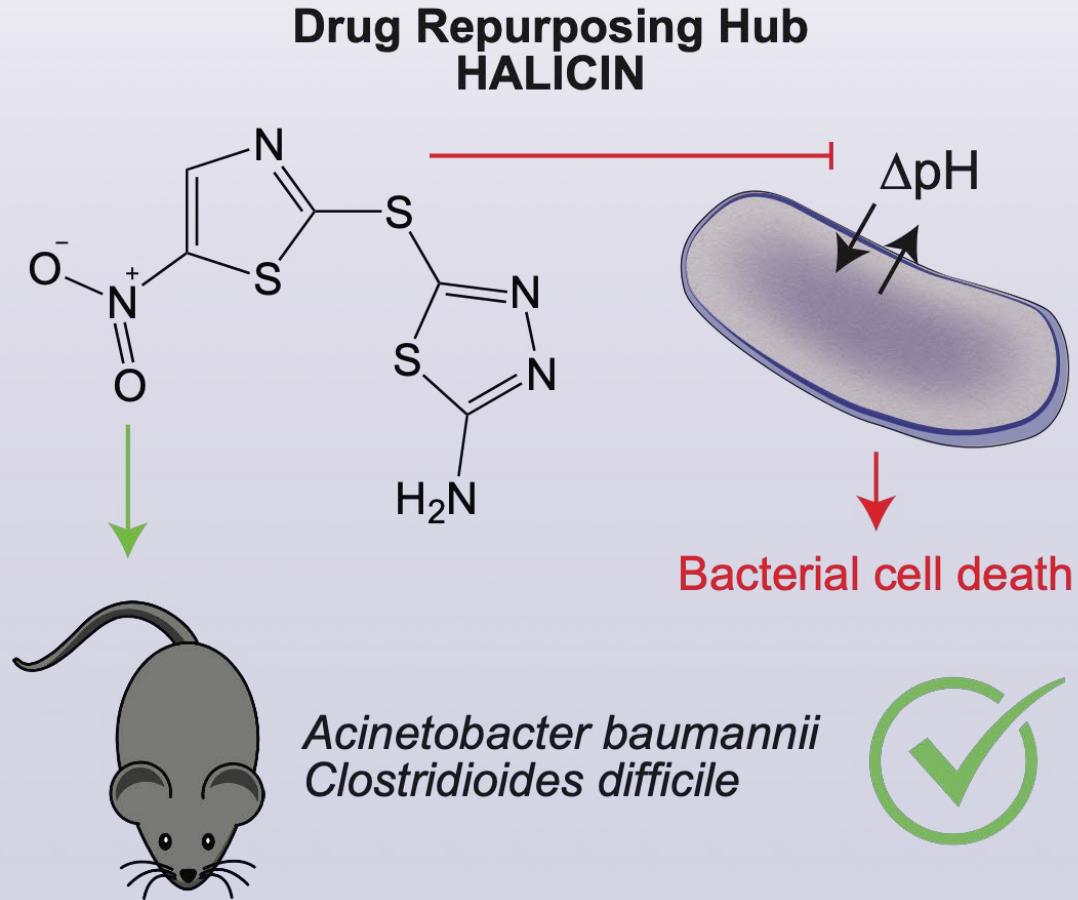


DEEP NEURAL NETWORK FOR MOLECULAR PROPERTY PREDICTION



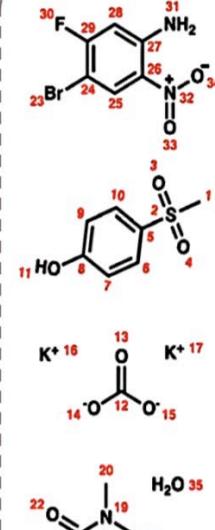
Message Passing Neural Network: train on a dataset containing molecules along with known property values for each molecule

DEEP LEARNING APPROACH TO ANTIBIOTIC DISCOVERY

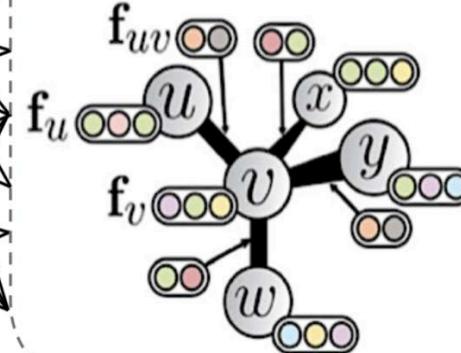


AI-BASED DISCOVERY AND DESIGN OF NOVEL ANTIBIOTICS

Large chemical compound library



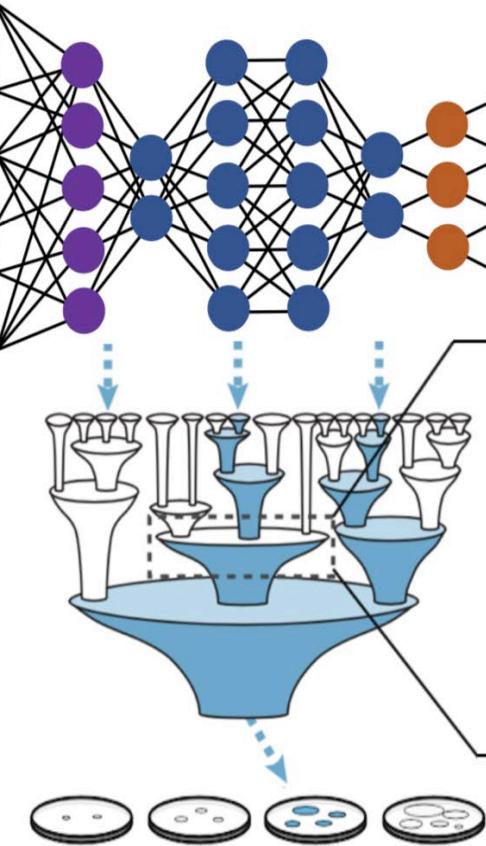
Learned molecular representations



Novel antibiotic prediction

Feature-based molecular property optimization

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

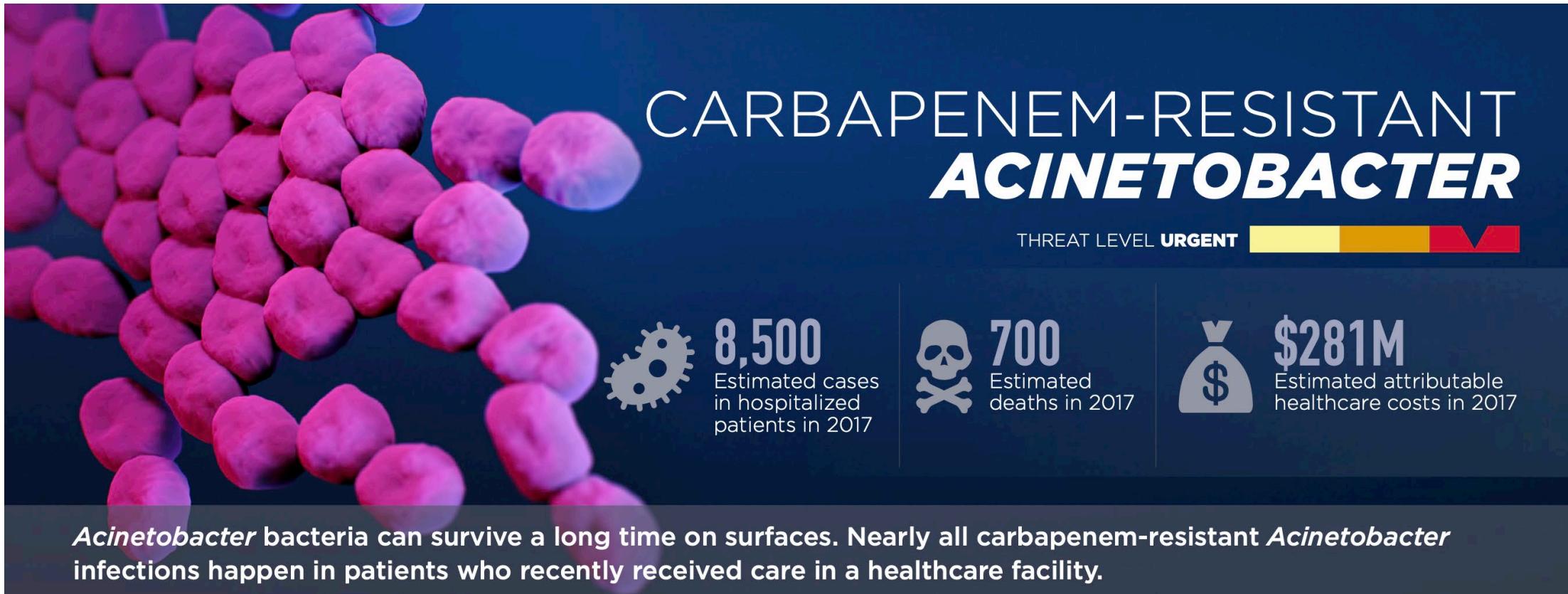


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Antibiotics-AI Project

Predicted antibiotic efficacy

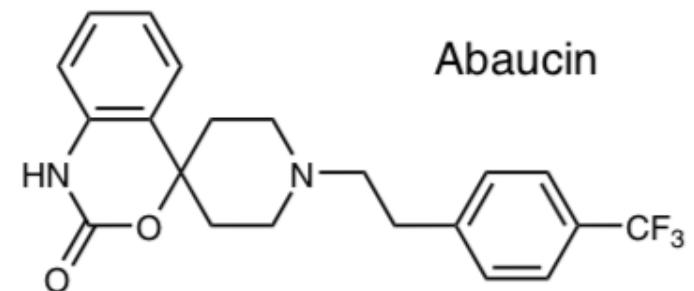
AI-BASED DISCOVERY OF ABAUCIN



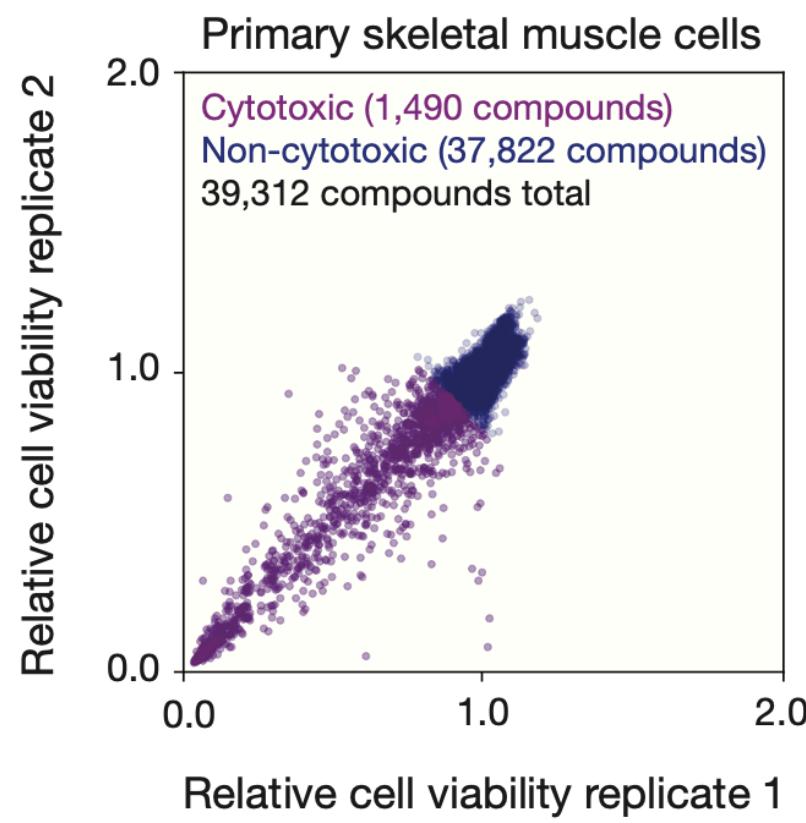
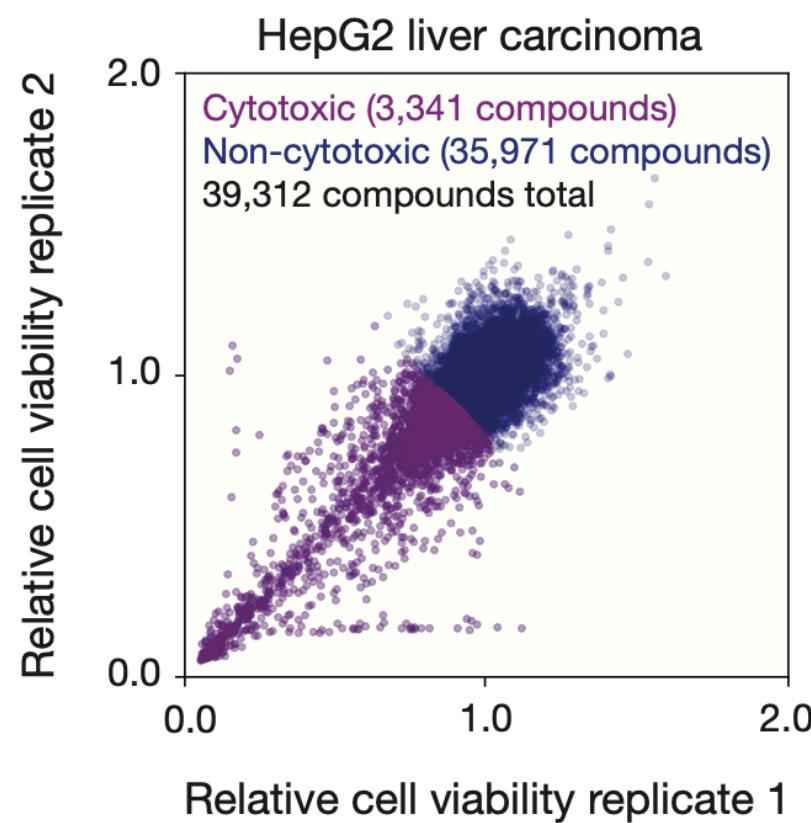
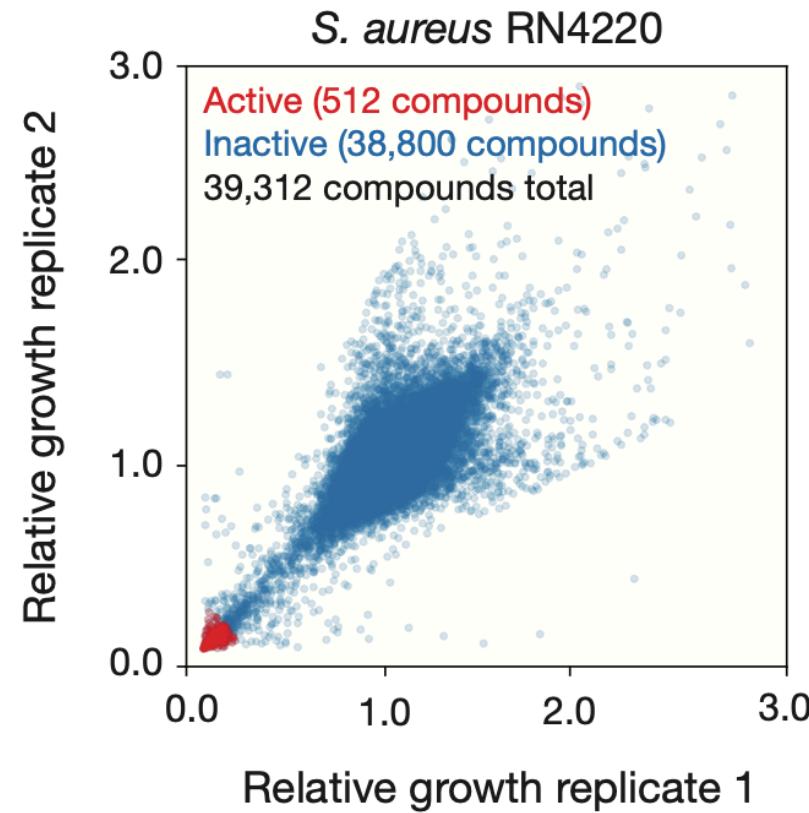
10 Downing Street



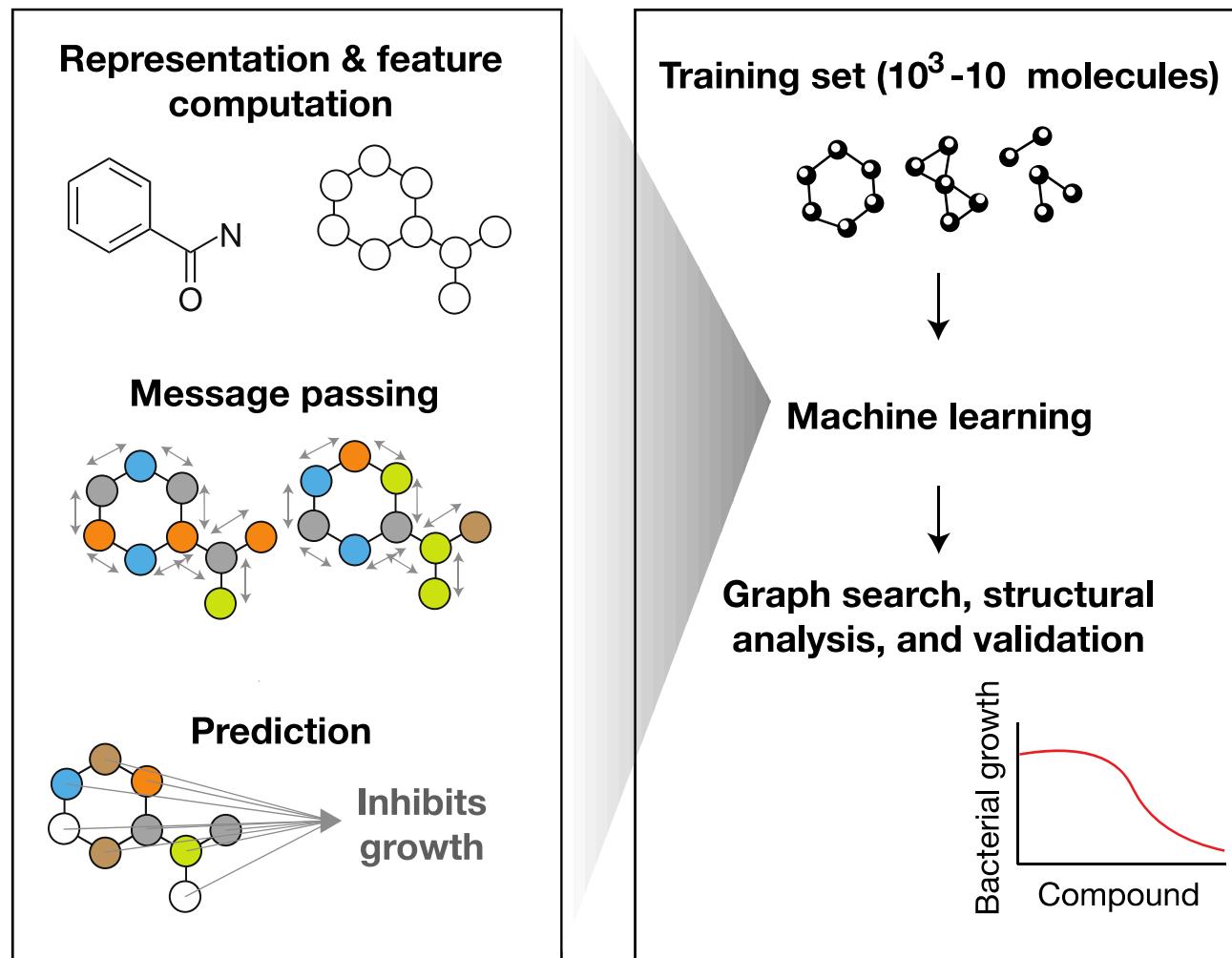
U.S. Capitol



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

$\sim 10^3$

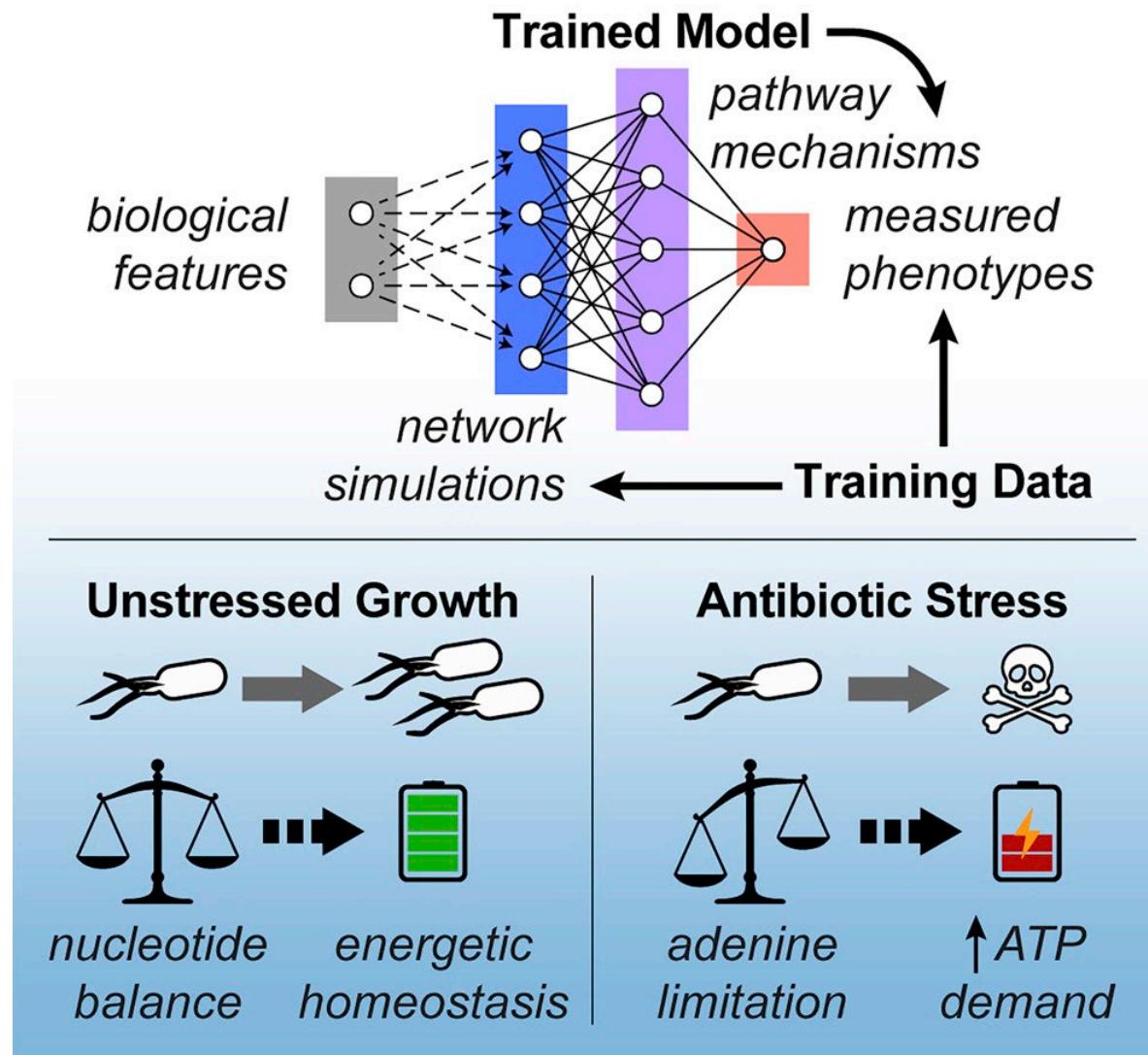
Number of strong predicted hits

$\sim 10^2$

Number of test molecules screened

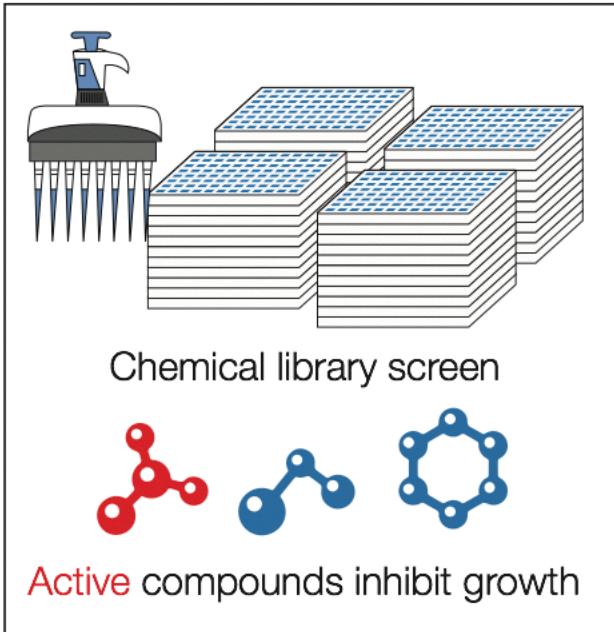
WHITE-BOX MACHINE LEARNING: IDENTIFYING ANTIBIOTIC MoA

White-Box Machine Learning

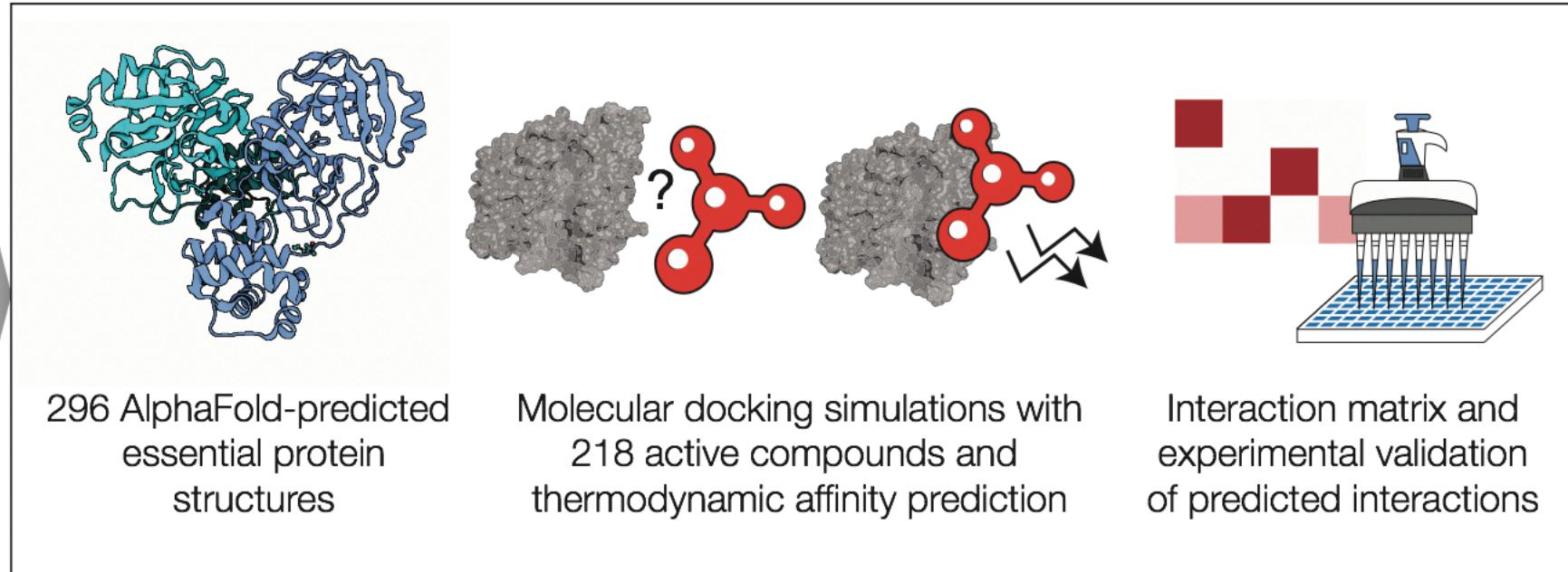


ANTIBIOTICS-AI PROJECT: LEVERAGING ALPHAFOLD

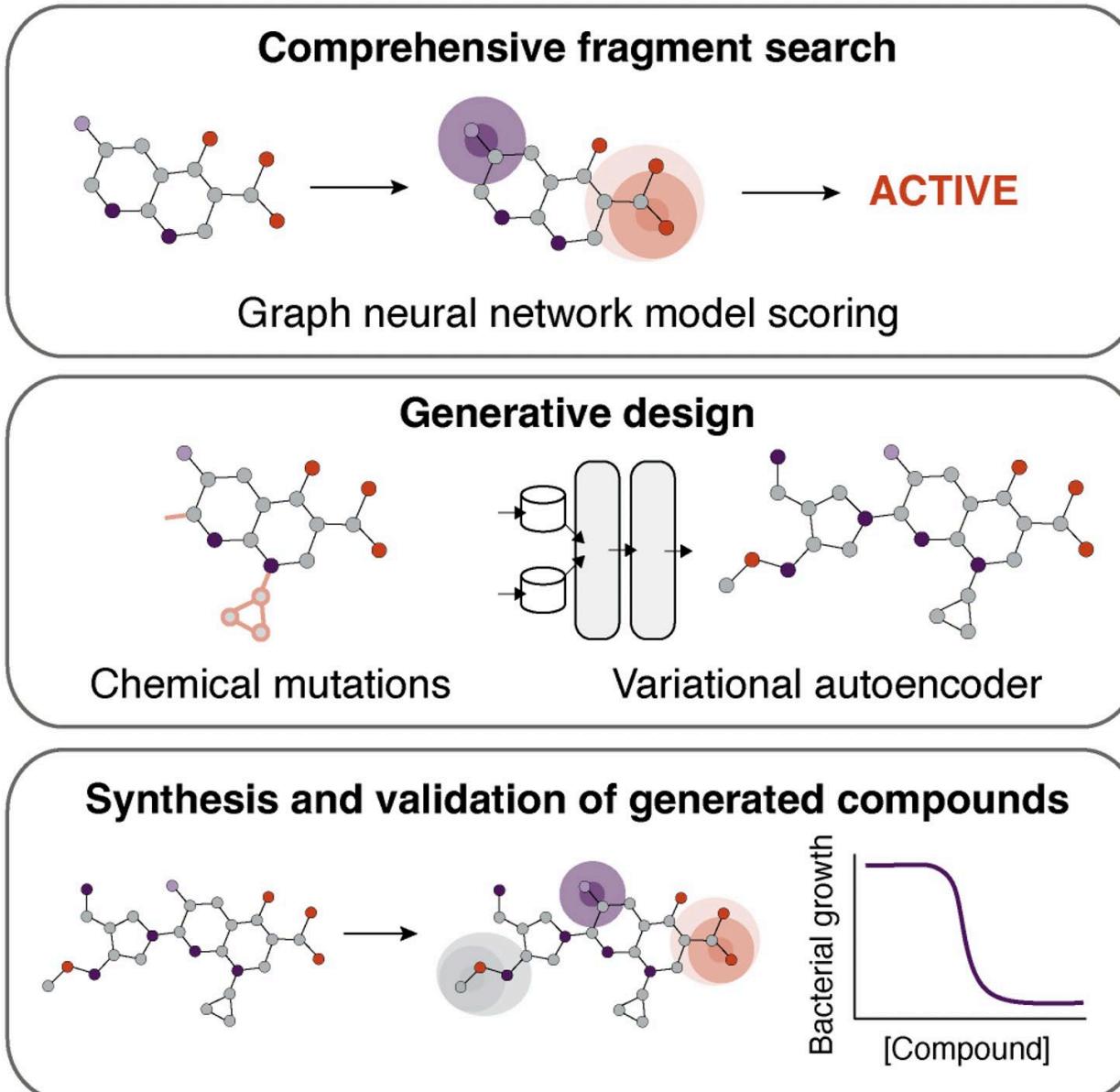
Growth inhibition screening



AlphaFold-enabled identification of protein-ligand interactions



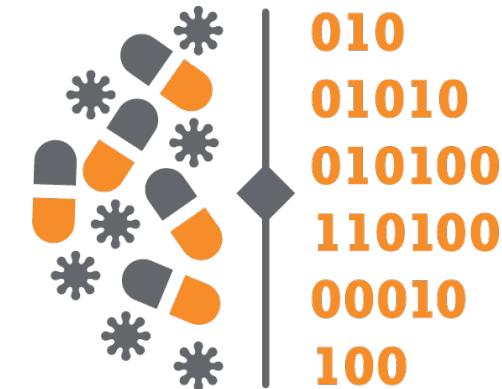
GENERATIVE AI FOR *De Novo* ANTIBIOTIC DESIGN



ANTIBIOTICS-AI PROJECT AND PHARE BIO



TED



Antibiotics-AI Project

PHAREBIO