

WHAT WE'RE CURRENTLY DOING



Chemical Information





Molecular Network



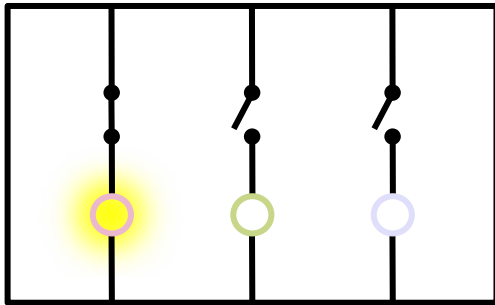
Phenotype Information

WHAT WE'RE CURRENTLY DOING


 Chemical Information


 Molecular Network

 Phenotype Information

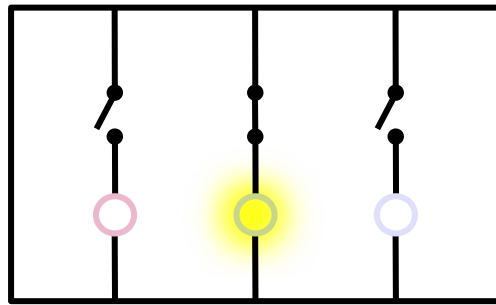
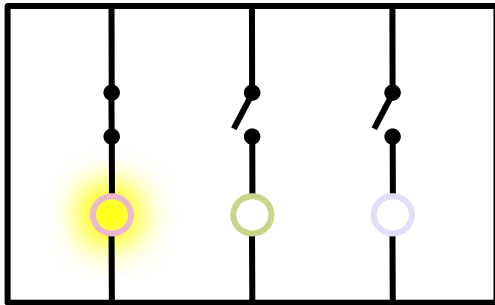


WHAT WE'RE CURRENTLY DOING


 Chemical Information


 Molecular Network

 Phenotype Information

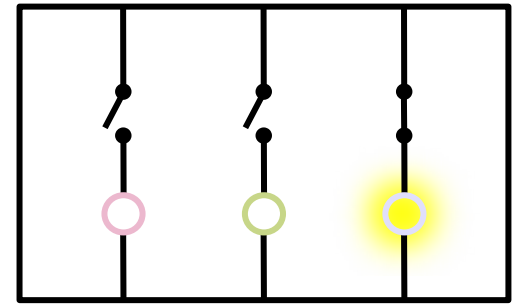
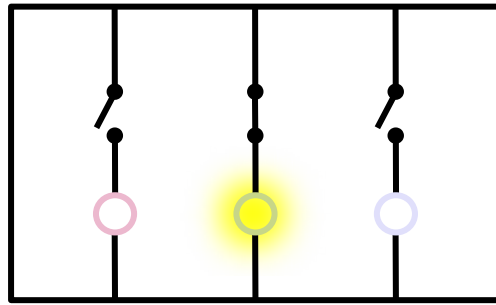
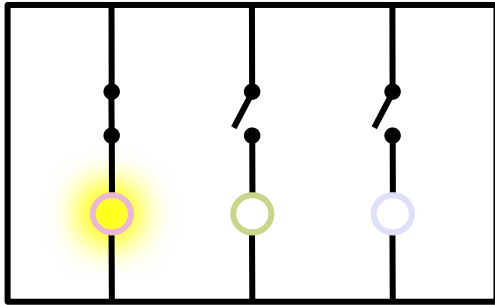


WHAT WE'RE CURRENTLY DOING

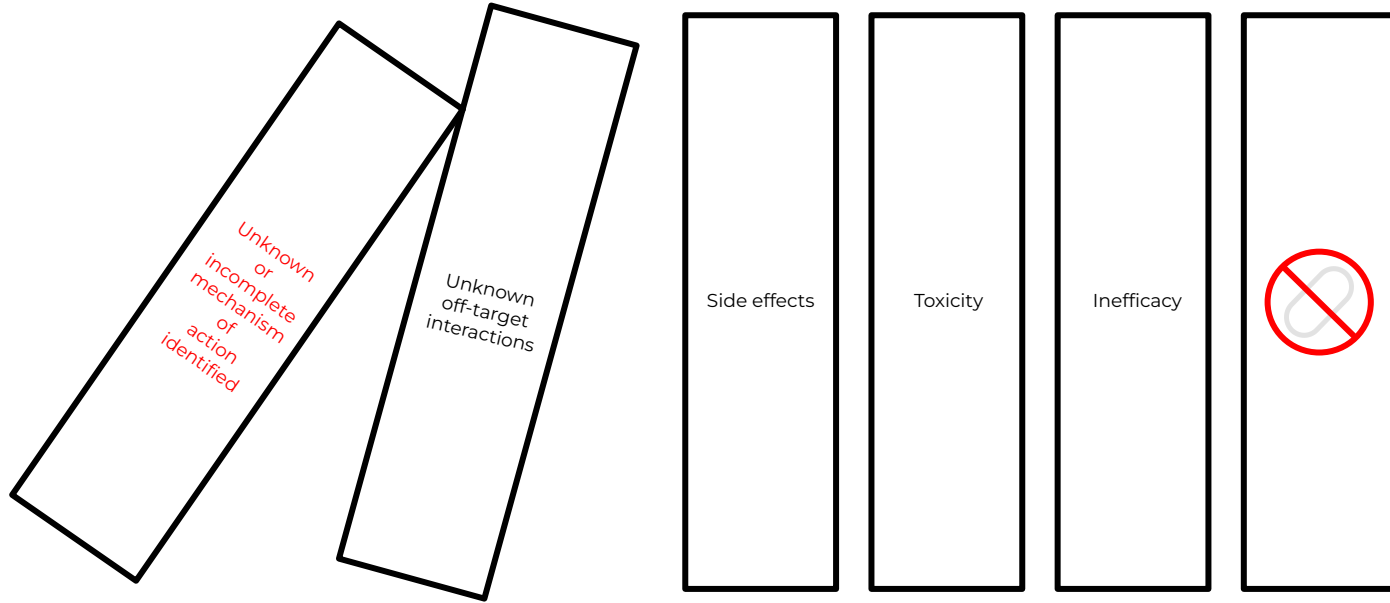
 Chemical Information

 Molecular Network

 Phenotype Information

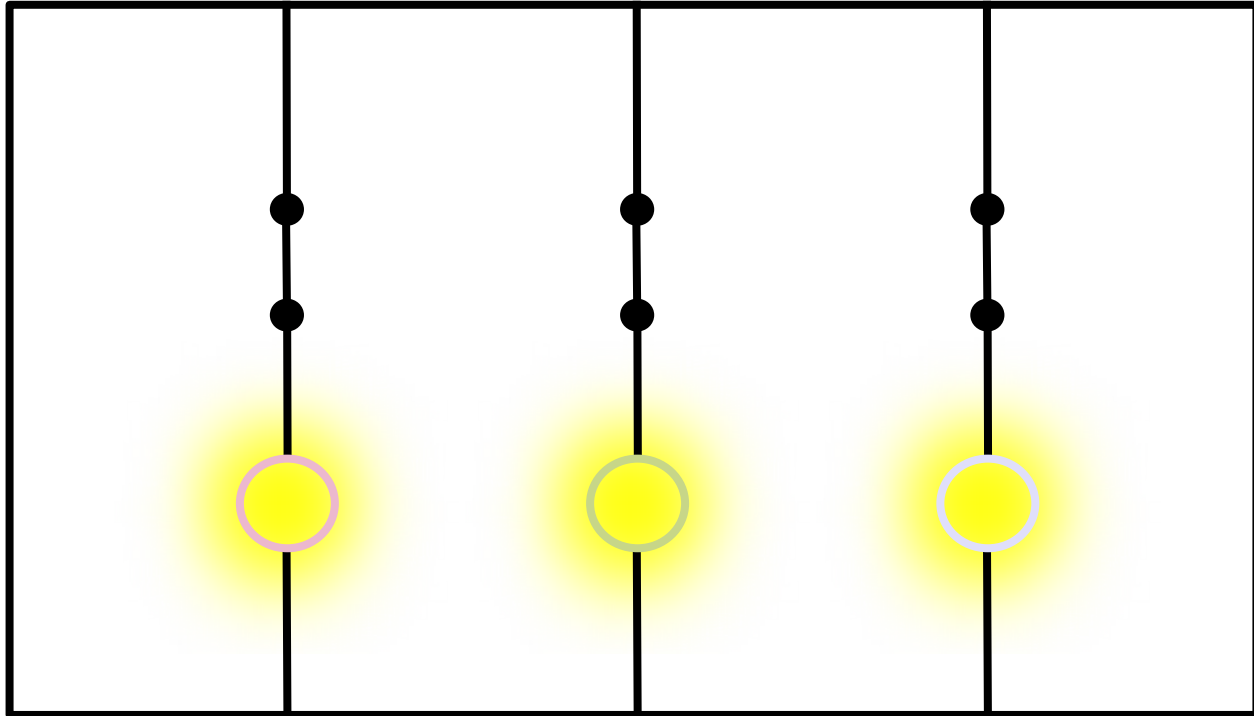


THE RESULT



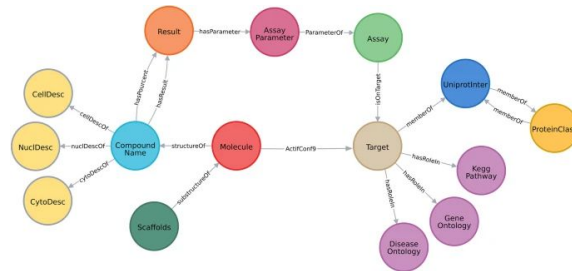
90% DRUG FAILURE RATE.

BUT WHAT IF...





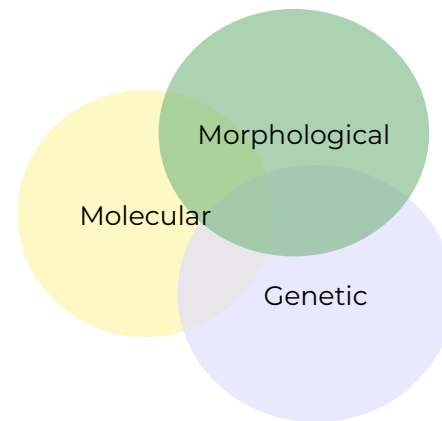
A graph database
for a
multi-dimensional
understanding of
MoAs



The screenshot shows the ChEMBL database interface. On the left, there is a 'Filters' sidebar with a tree view of compound types: N/A (19498), Antibody (994), Cell (47), Enzyme (118), Gene (77), Oligonucleotide (170), Oligosaccharide (92), Protein (22682), Small molecule (1920338), Unclassified (4), and Unknown (18016). Below this is a 'Max Phase' filter with options 0 (2147608), 1 (1479), and 2 (2346). The main area displays a grid of compound cards, each with a chemical structure, a ChEMBL ID, and key properties like Name, Max Phase, Full Mw, and Atopp. The interface includes a search bar at the top, a 'Records per page' dropdown set to 12, and a 'Showing 1-12 out of 2,157,379 records' indicator.

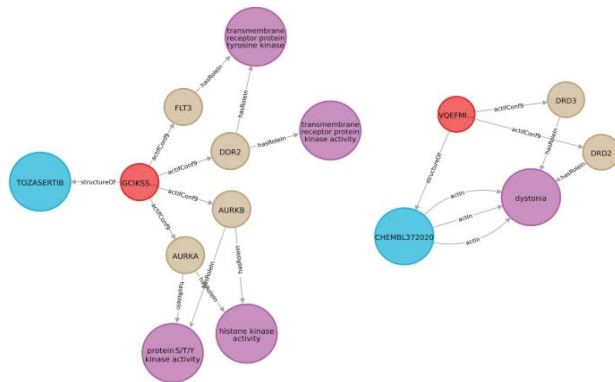
1

Mine data from public databases + client data to add to BenchSci's database



2

Standardize molecular, genetic, and morphological data

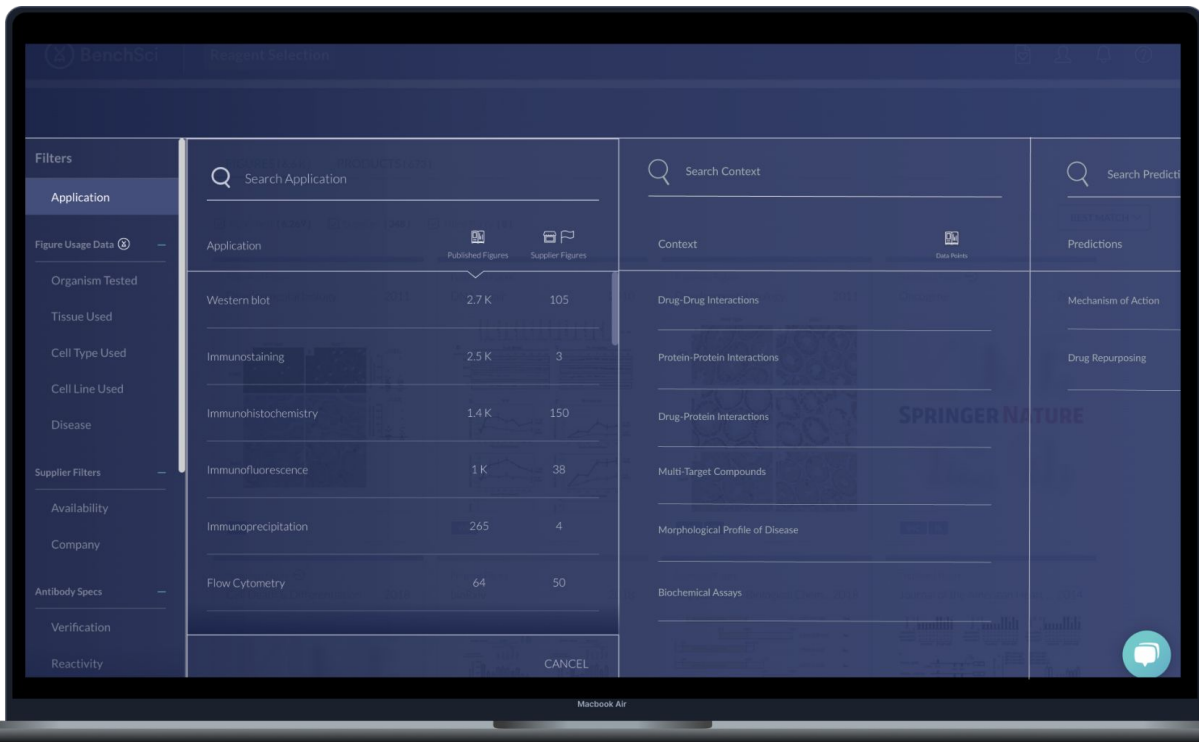


3

**Multi-dimensional
graph network
generation**

4

**Mechanism of action,
multi-target compound,
chemical-morphological
profiling and prediction**



DRUG REPURPOSING: ENGINEERING SERENDIPITY

5X

Up to 500% reduction in **time**

15 yrs → 3 yrs

37X

Up to 3700% reduction in **costs**

\$2-3 billion → \$40-80 million

SYSTEMS MEDICINE: HOLISTIC UNDERSTANDING

Mechanism of
action prediction

Morphological
profiling

Toxicity & side
effect inference

Drug repurposing
avenues



