



Data Fusion by Matrix Completion for Exposome Target Interaction Prediction

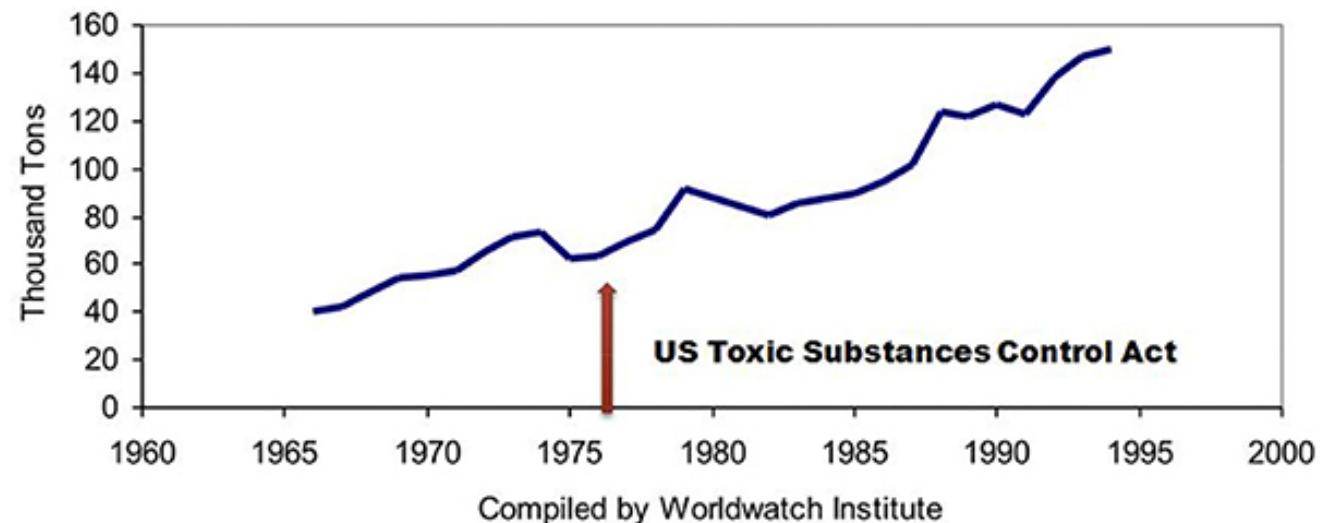
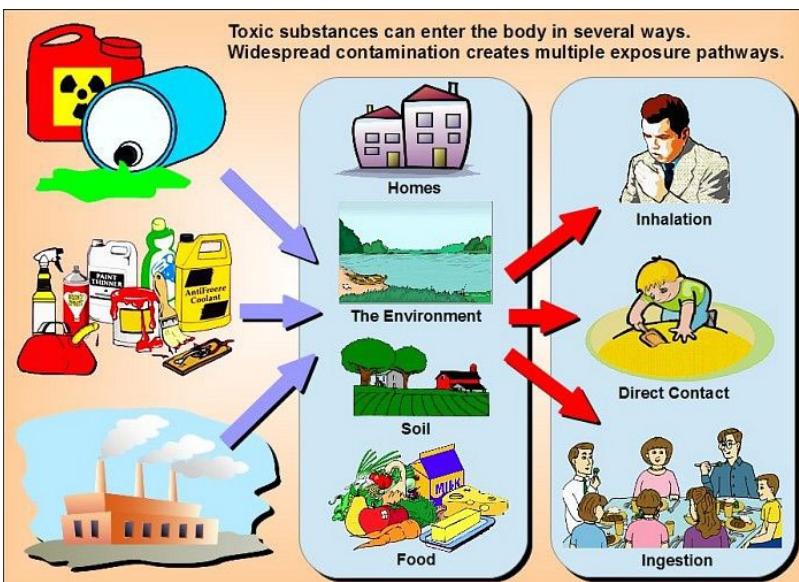
Kai Wang

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09/19/2023

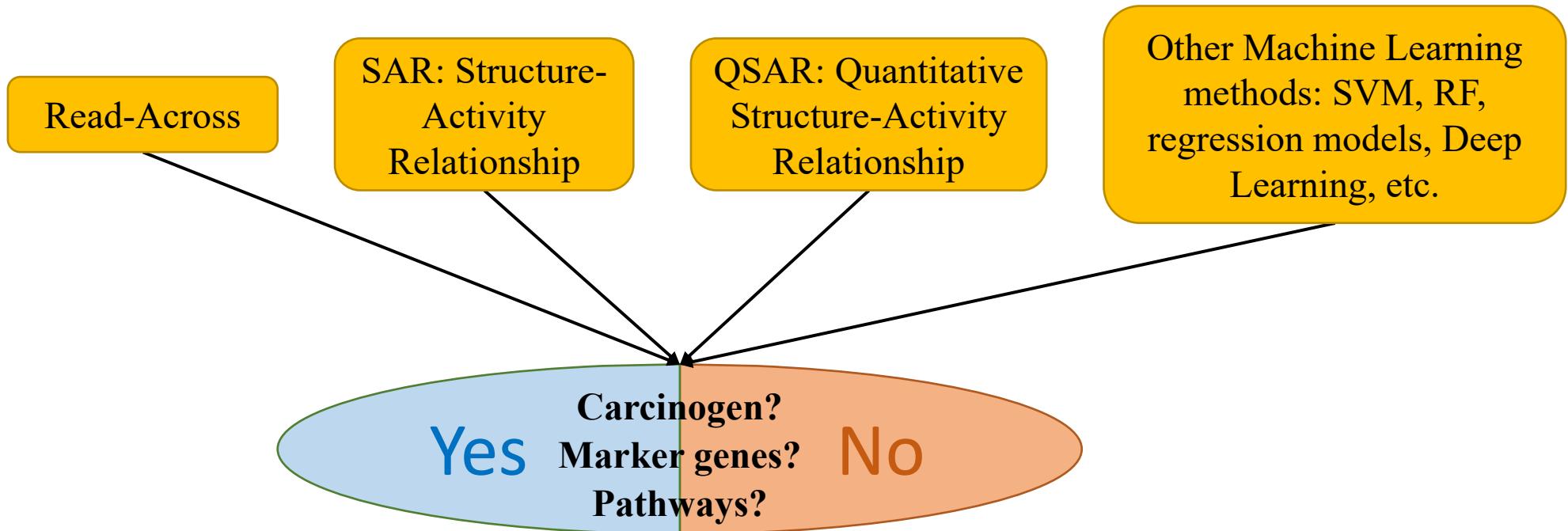
Environmental Exposures

- Environmental exposures alone or in combination with genetics can affect **human health** and **cause diseases**.
- New chemical compounds are created and released into our environment every year. About **2,000 new chemicals** are introduced into commerce annually in the U.S., at a rate of about **seven new chemicals** a day.
- Comprehensive toxicity assessment of all chemicals is an **impossible** task.



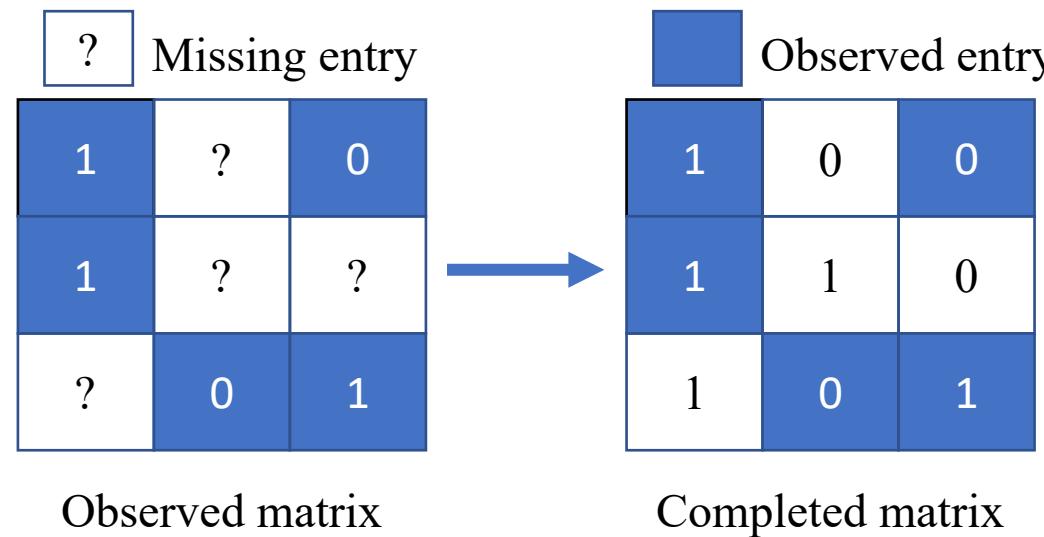
Computational Toxicology Methods

- Computational methods can be used to prioritize chemicals for *in vivo* or *in vitro* toxicity study.



Matrix Completion drug target prediction

Matrix completion is the task of filling in the missing entries of a partially observed matrix, which is equivalent to performing data imputation in statistics.



Drug target interaction: physical interaction

Environmental chemical: any types of interactions, transcription level, post translational modifications, etc.

Coupled Matrix-Matrix Completion

Input:

Main:

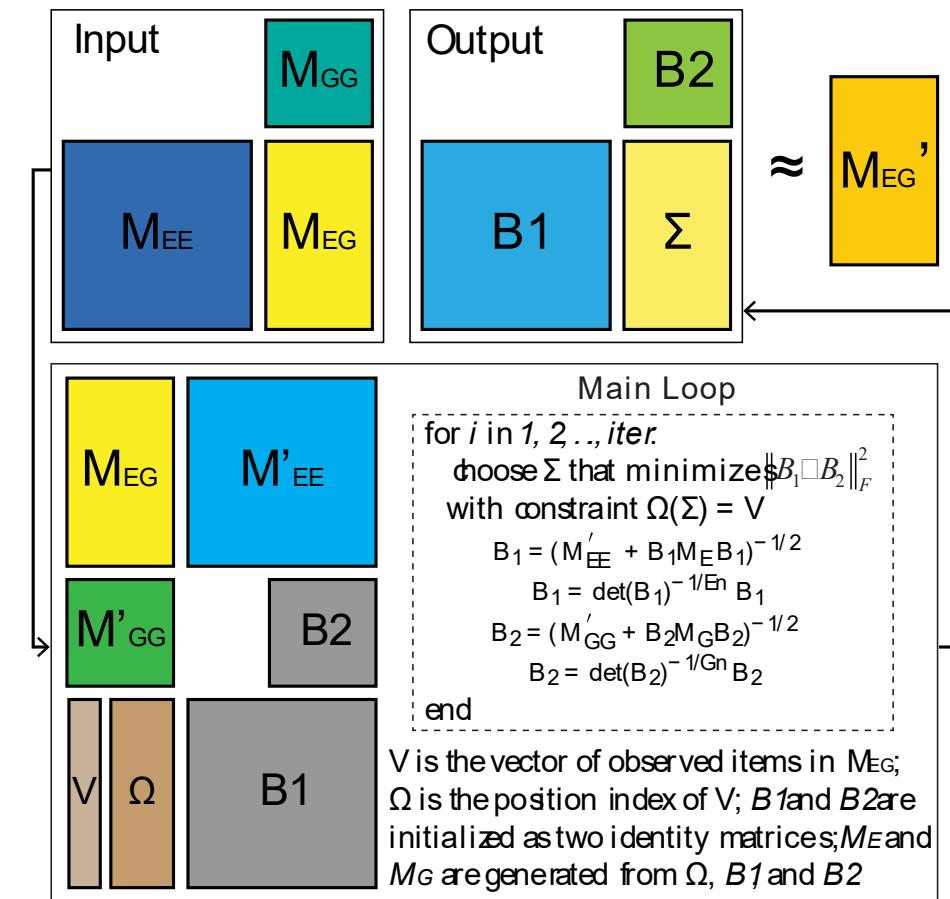
- Chemical-gene interaction matrix

Coupled:

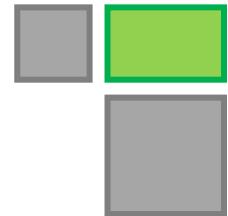
- Chemical-chemical similarities
- Gene-gene similarities

Output:

- Chemical-gene interaction matrix with imputed values



The Comparative Toxicogenomics Database

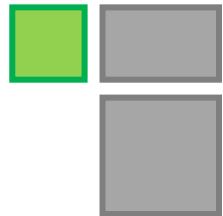


- A curated database that promotes understanding about the effects of **environmental chemicals** on **human health**.
- **4,864** chemicals exposures
- **22,606** genes
- **201,375** known chemical-gene interaction

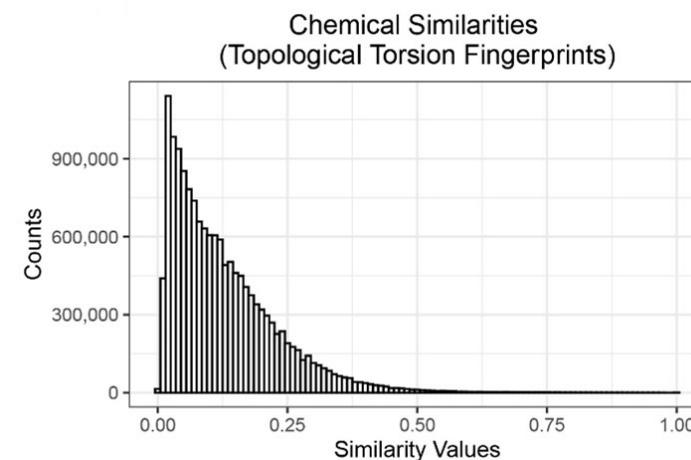
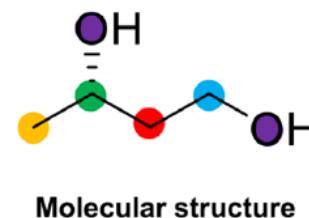
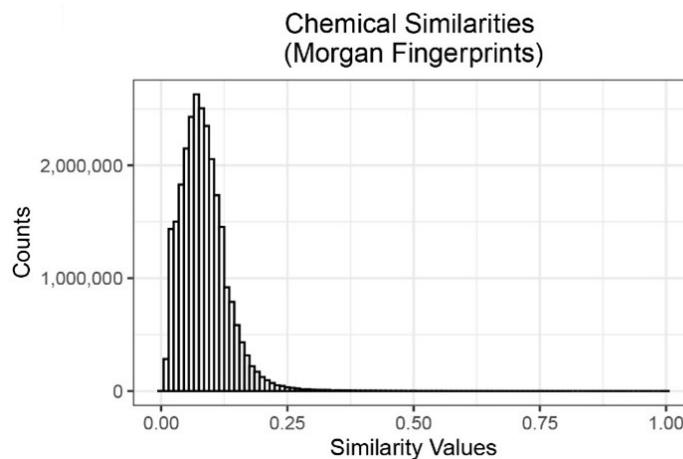
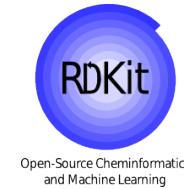
$201,375 / 109,955,584 \approx 0.183\%$



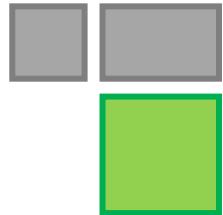
Chemical Similarities



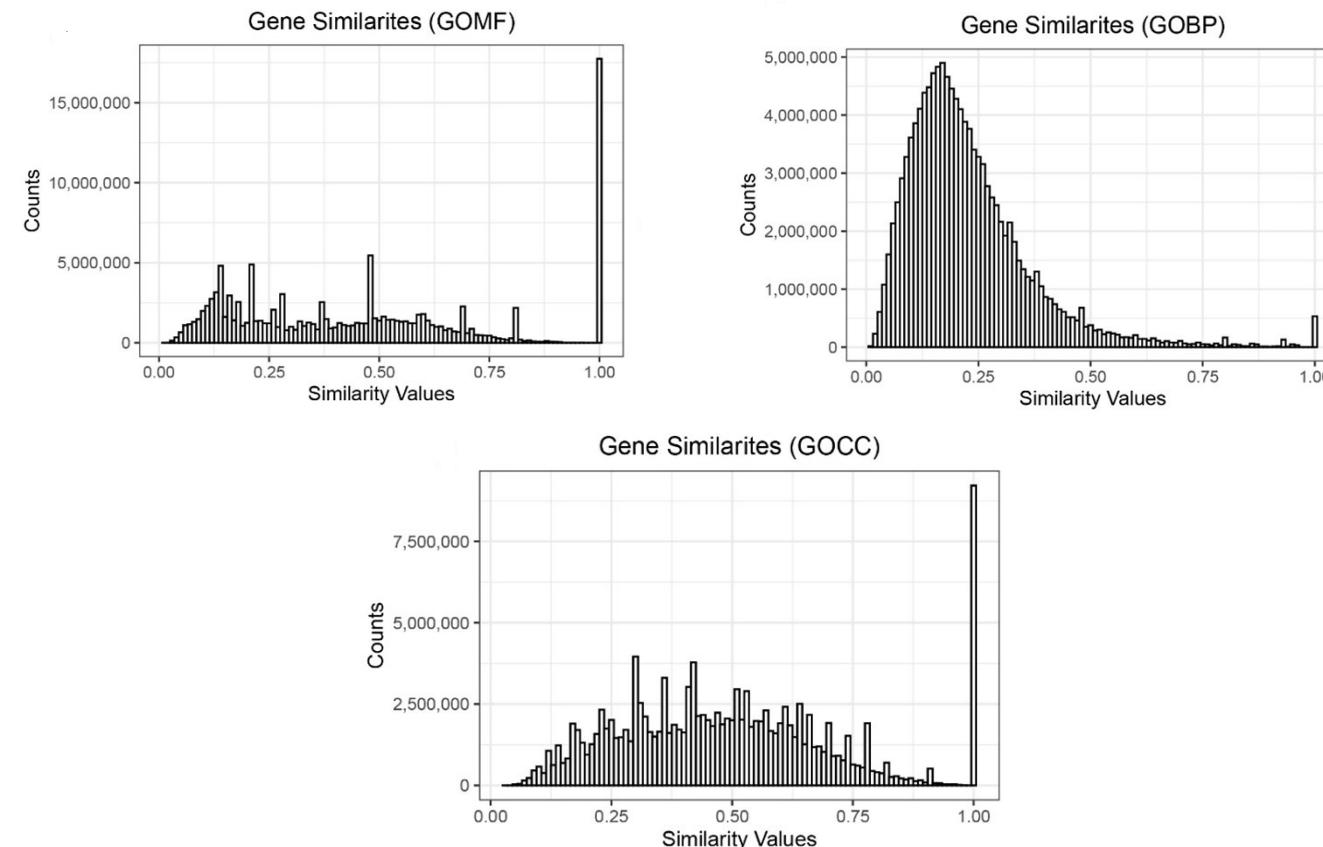
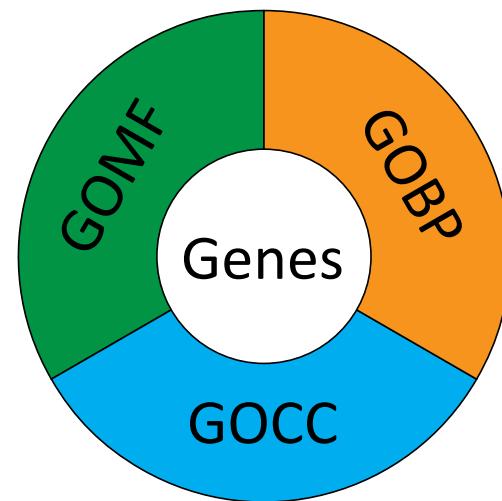
- RDKit: Open-Source Cheminformatics Python package
- Two types of fingerprints were used to calculate chemical similarities.



Gene Similarities

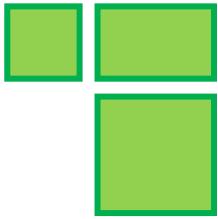
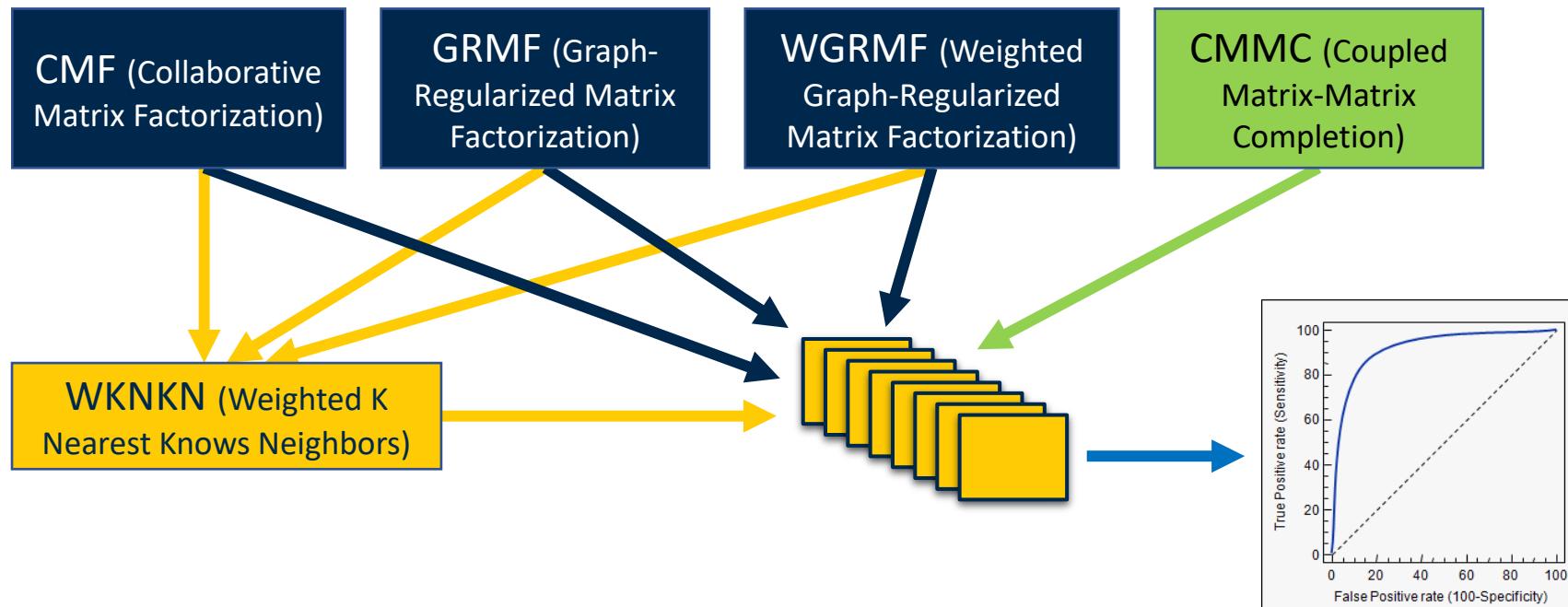


- GOSemSim: an R package for measuring semantic similarity among GO terms and gene products.

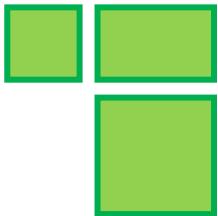
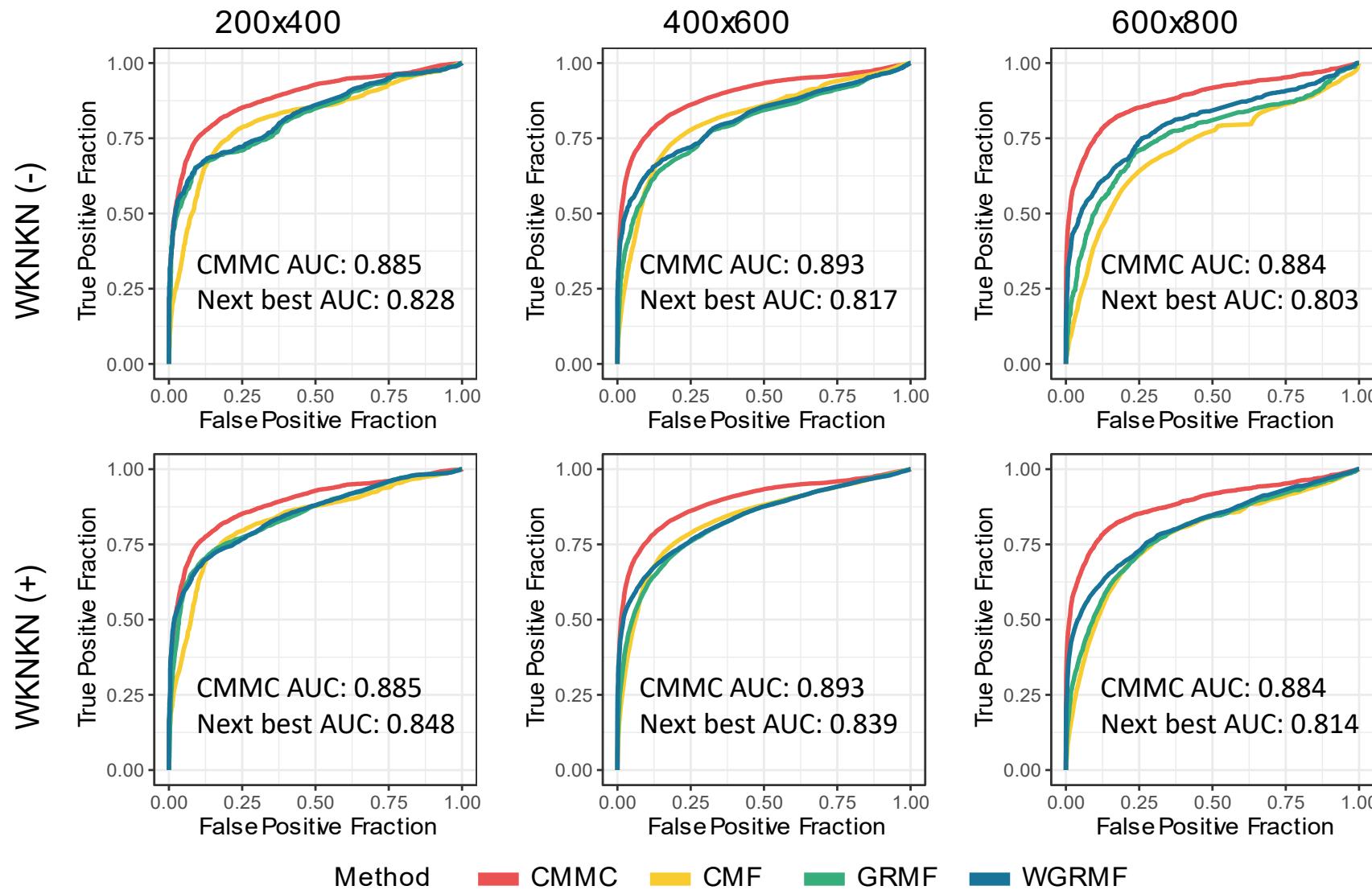


Performance Comparison

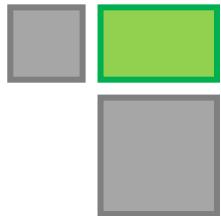
- MATLAB - CMF, GRMF, WGRMF
- CMMC - C++



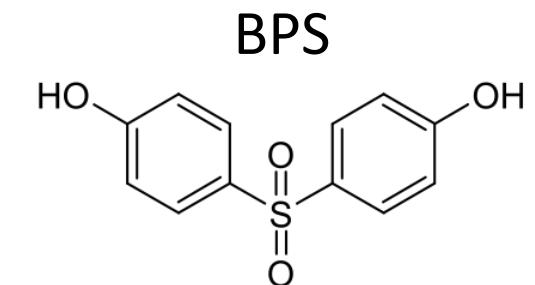
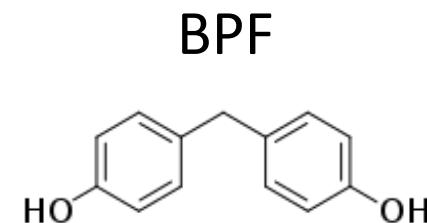
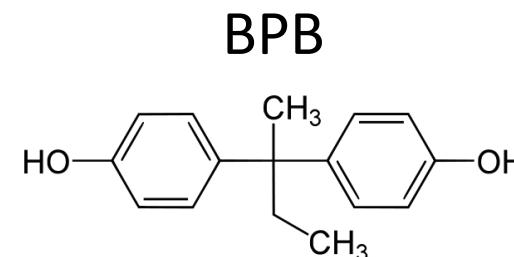
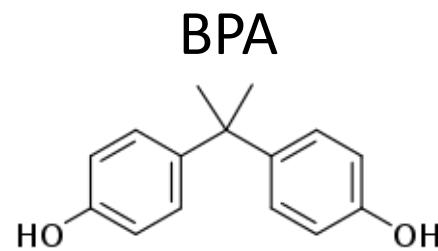
Performance Comparison



Case study - BPA with its analogs



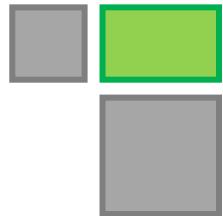
Problem: how to get the target gene of novel chemicals without any biological activity data?



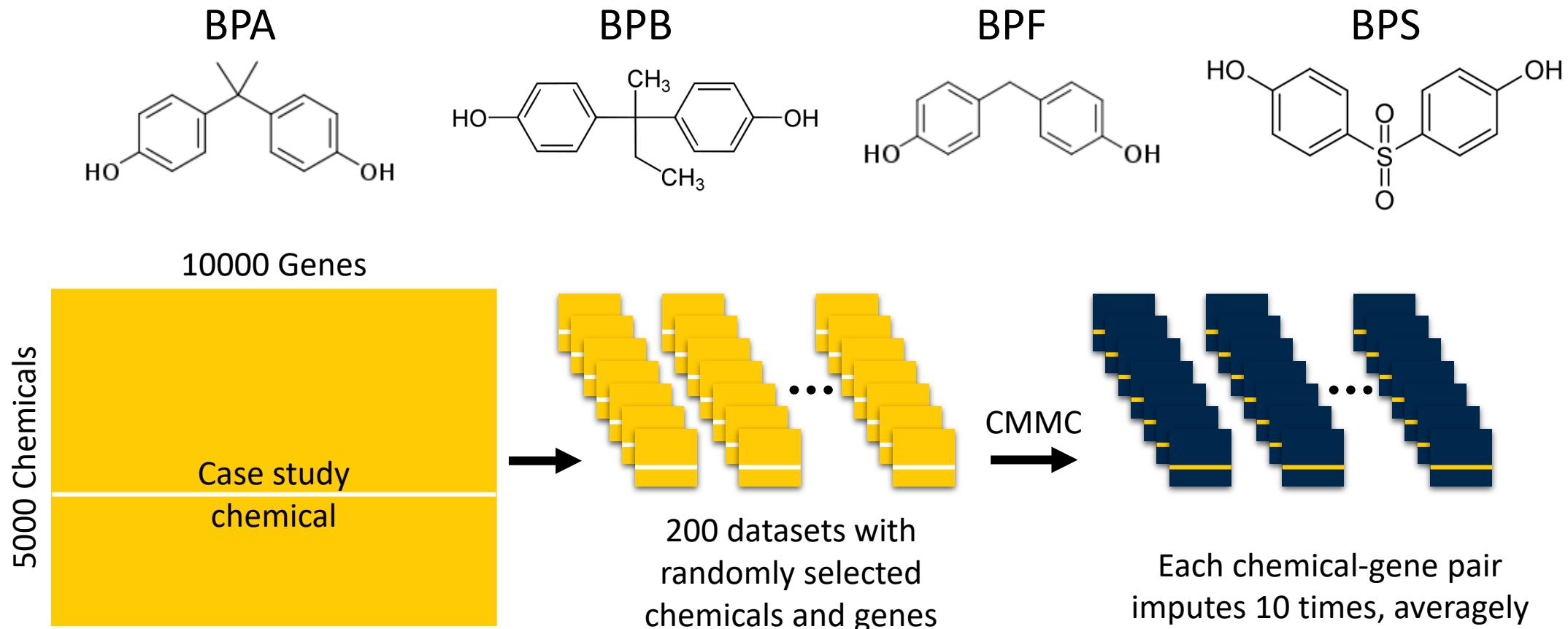
CMMC



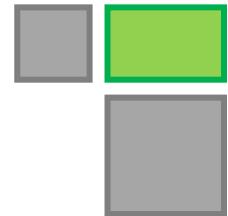
Case study - BPA with its analogs



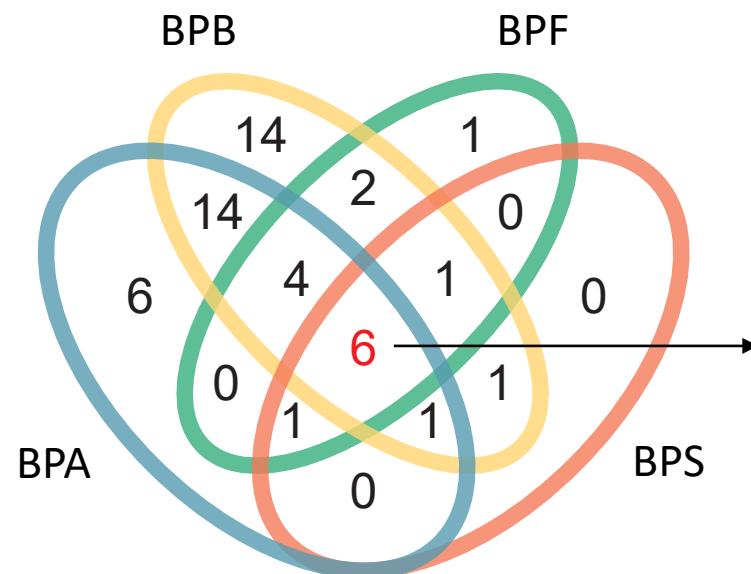
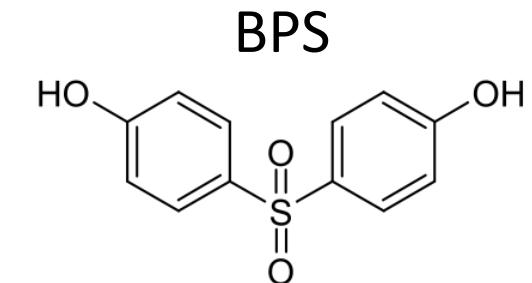
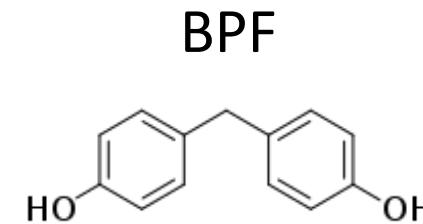
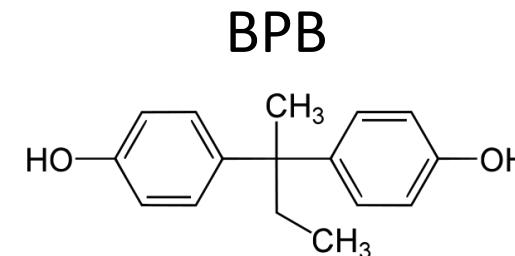
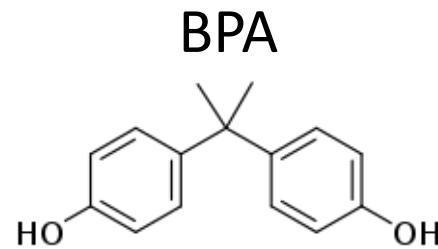
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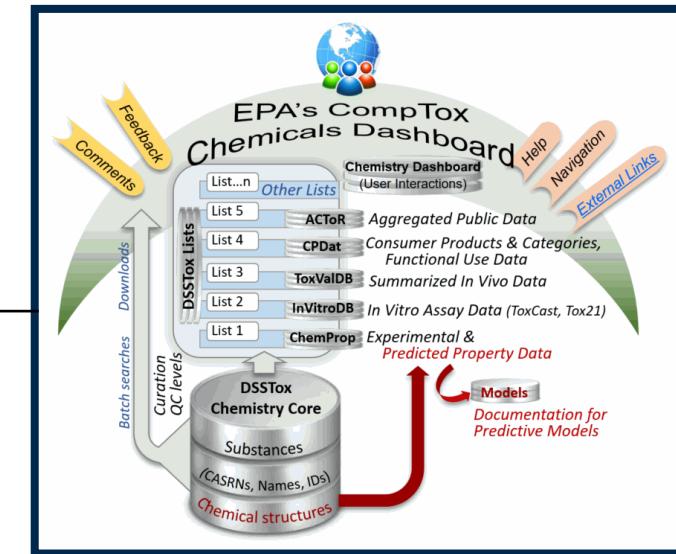
Case study - BPA with its analogs



Problem: how to get the target gene of novel chemicals without any biological activity data?



ESR1
ESR2
AR
NR1/2
PGR
PPARG
PPARD



GPER1: target gene of BPB

- G Protein-Coupled Estrogen Receptor 1: This gene encodes a **multi-pass membrane protein** that localizes to the endoplasmic reticulum and a member of the G-protein coupled receptor 1 family.
- CMMC predicted this gene as a **top interactive gene** only for **BPB**.



Article

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Bisphenol AF and Bisphenol B Exert Higher Estrogenic Effects than Bisphenol A via G Protein-Coupled Estrogen Receptor Pathway

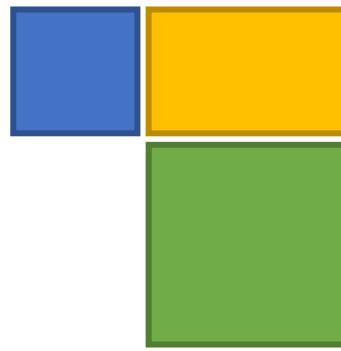
Lin-Ying Cao,^{†,‡} Xiao-Min Ren,^{*,†} Chuan-Hai Li,^{†,‡} Jing Zhang,[†] Wei-Ping Qin,^{†,‡} Yu Yang,[†] Bin Wan,[†] and Liang-Hong Guo^{*,†,‡,ID}

[†]State Key Laboratory of Environmental Chemistry and Eco-toxicology, Research Center for Eco-environmental Sciences, Chinese Academy of Sciences, 18 Shuangqing Road, Beijing 100085, P. R. China

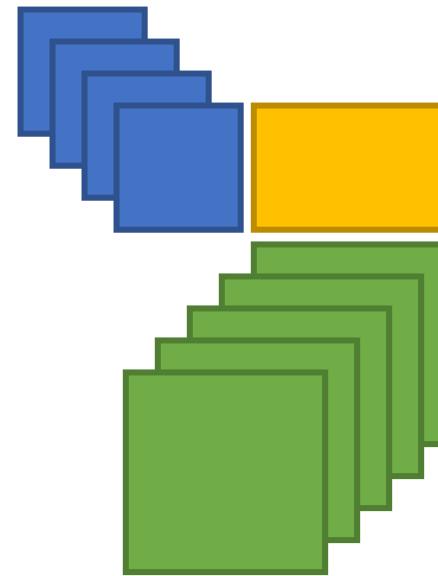
^{*}University of Chinese Academy of Sciences, Beijing 100039, P. R. China

...six BPA analogues bound to **GPER** directly, with bisphenol AF (**BPAF**) and bisphenol B (**BPB**) displaying **much higher (~9-fold)** binding affinity than **BPA**...

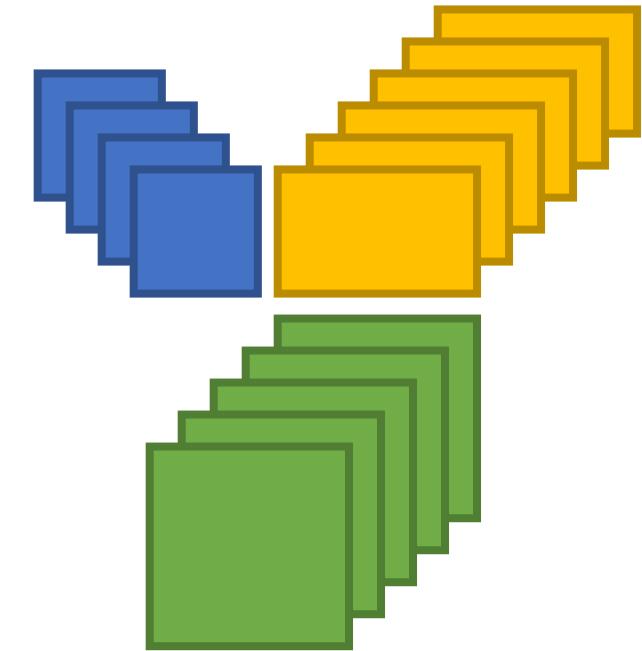
Future plan - CTMC and CTTC



CMMC



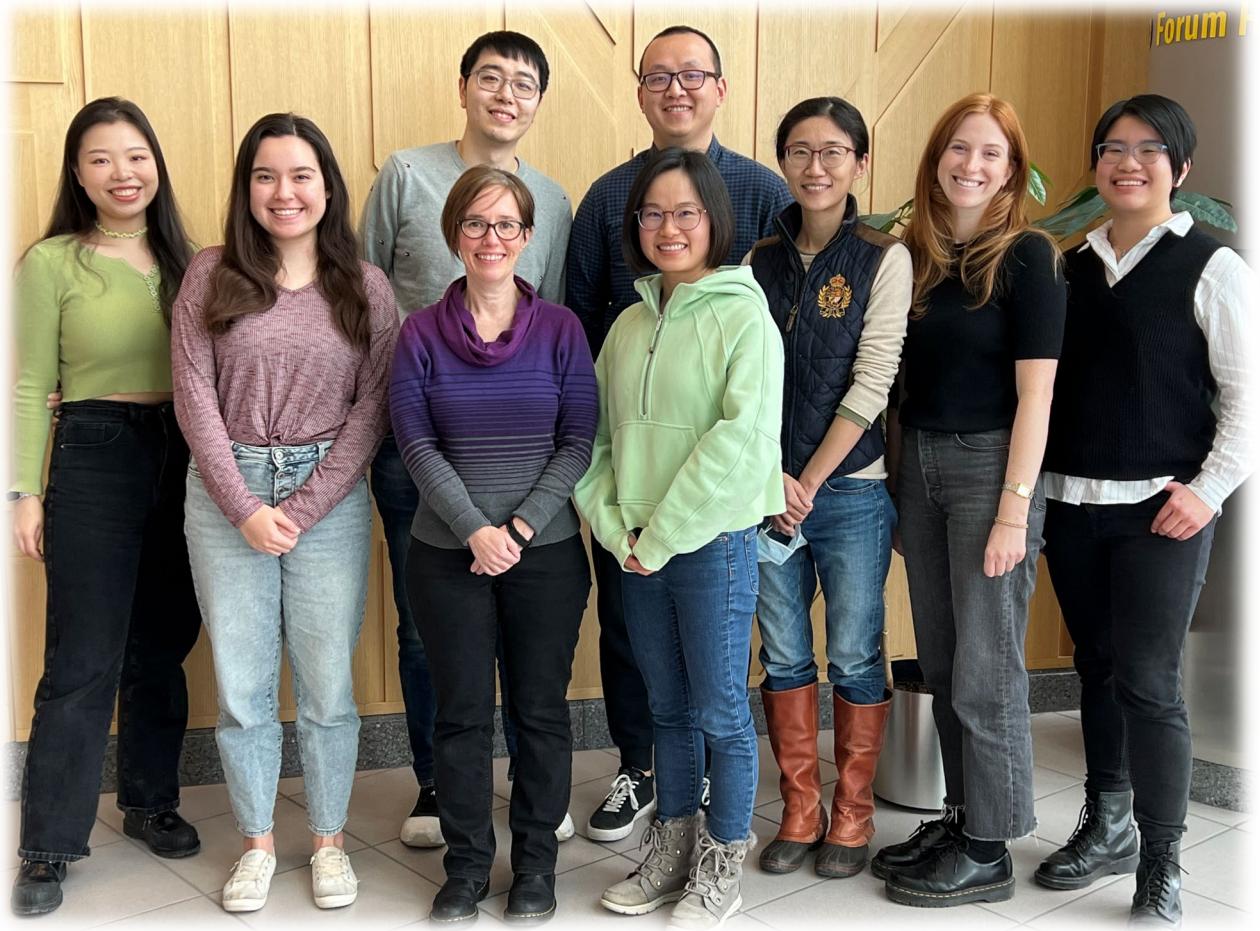
CTMC



CTTC

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