

Integrating Machine Learning and Quantitative Structure-Activity Relationships (QSAR) Modeling Approaches to Develop Artificial Intelligence (AI)- assisted Interactive Physiologically Based Pharmacokinetic (iPBPK) Modeling Web Dashboard.

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Case studies

- A. Development of Artificial Intelligence (AI)-Assisted PBPK Model in Cancer Nanomedicine**

- B. Development of a Multi-Organ Toxicity Predictive Model Using Multi-Task Learning in Deep Neural Network**

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Challenge in tumor delivery of nanomedicine

- NPs are becoming an increasingly popular tool for biomedical imaging and drug delivery.

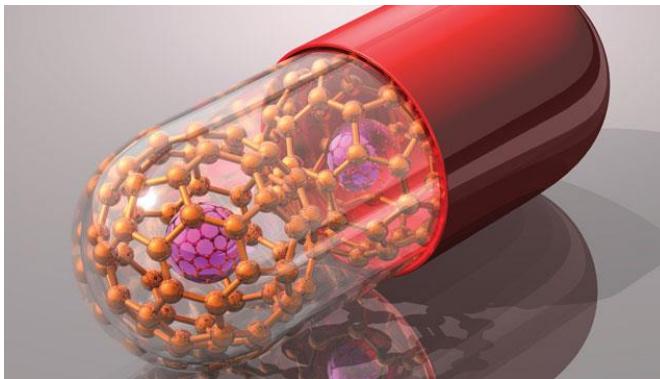
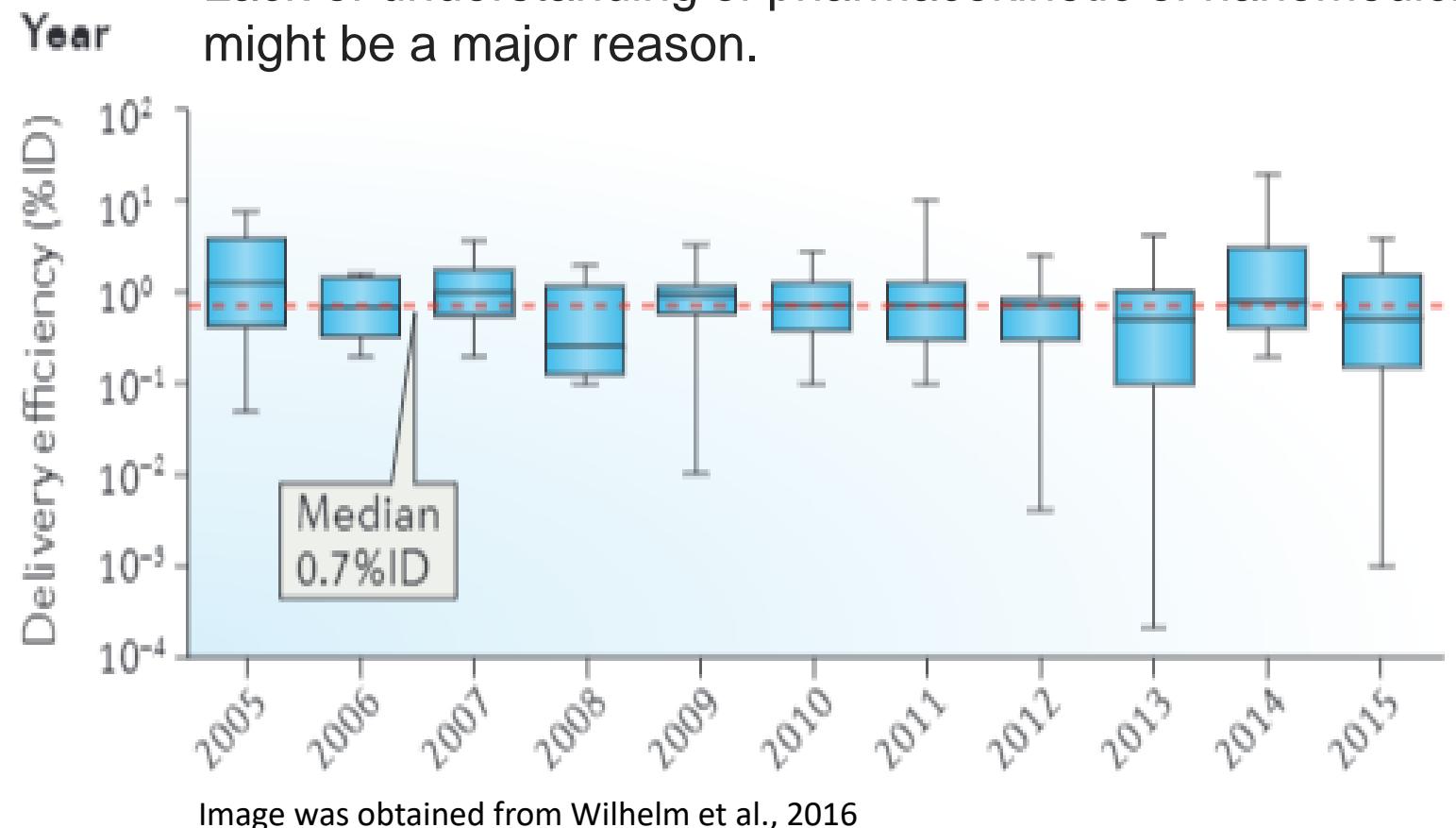


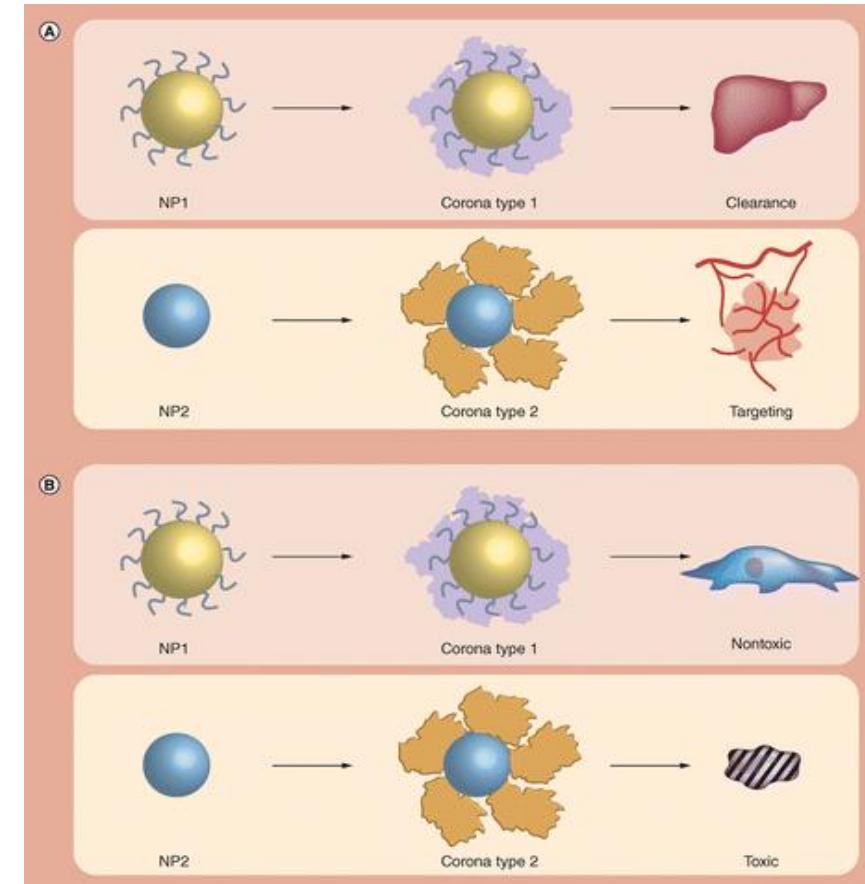
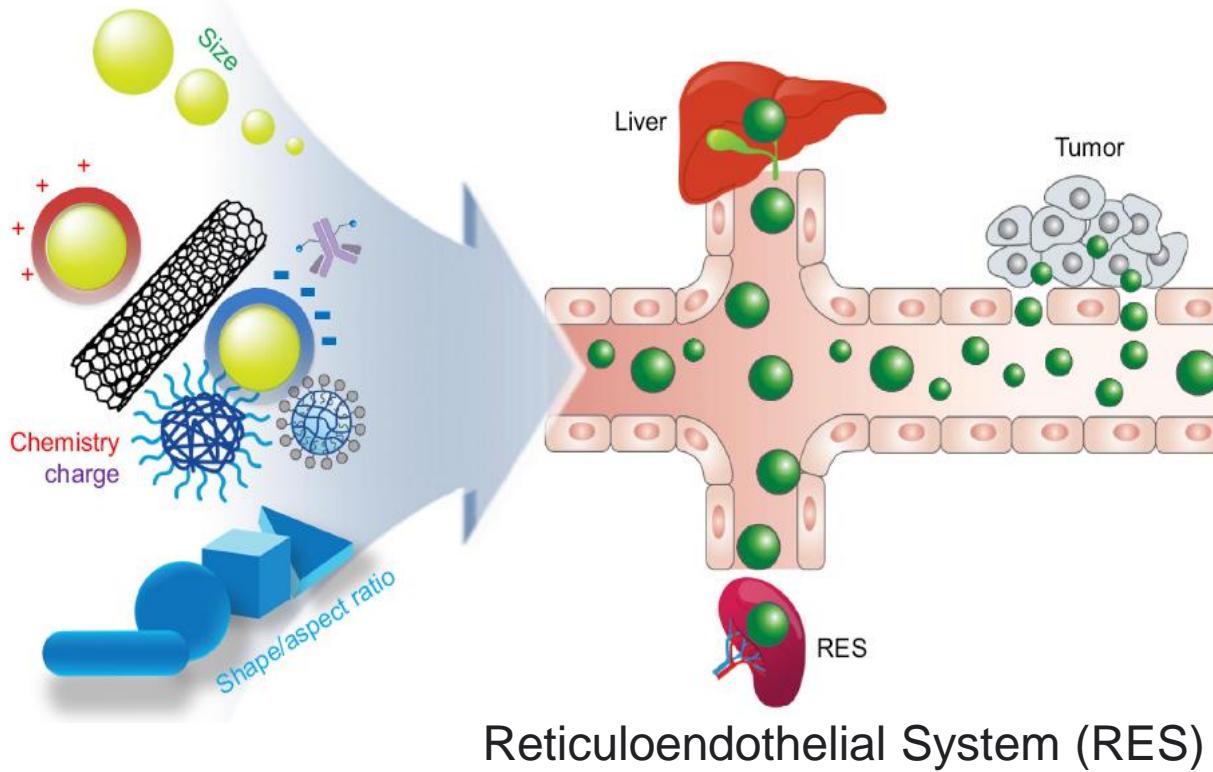
Image source: <https://www.the-scientist.com/cover-story/nanomedicine-37087>

- The poor tumor delivery efficiency of nanomedicines has been a major barrier in the translation of nanomedicine to potent drug candidates.

- Lack of understanding of pharmacokinetic of nanomedicine might be a major reason.

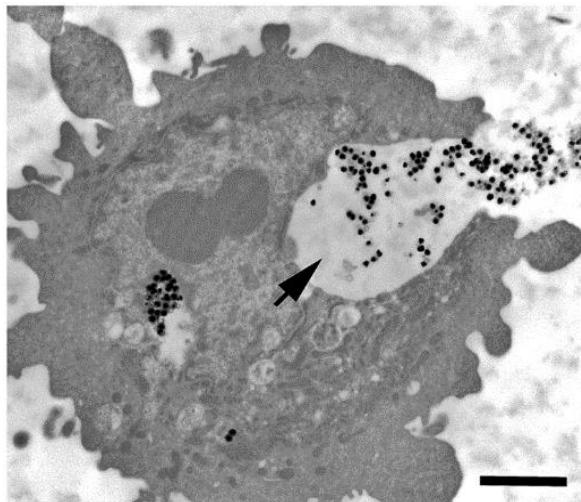


Biodistribution of Nanoparticles (NPs)



- The pharmacokinetics of nanomedicine is very different with the traditional drugs.
- One of important mechanisms to affect the NPs' biodistribution is **phagocytosis**.
- Different physicochemical properties of NPs, such as size, materials, biochemistry, and shape, may relate to the NPs' phagocytosis and biodistribution.

Theoretical parameter: Endocytosis of NPs



Monteiro-Riviere et al. 2013. Toxicology Letters

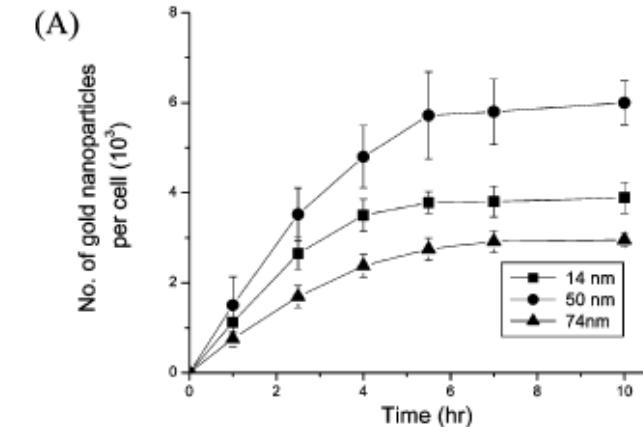
- Hill function to simulate endocytosis of gold nanoparticles

$$K_{up,i}(t) = \frac{K_{max,i} \times t^{n_i}}{t_{50,i}^{n_i} + t^{n_i}}$$

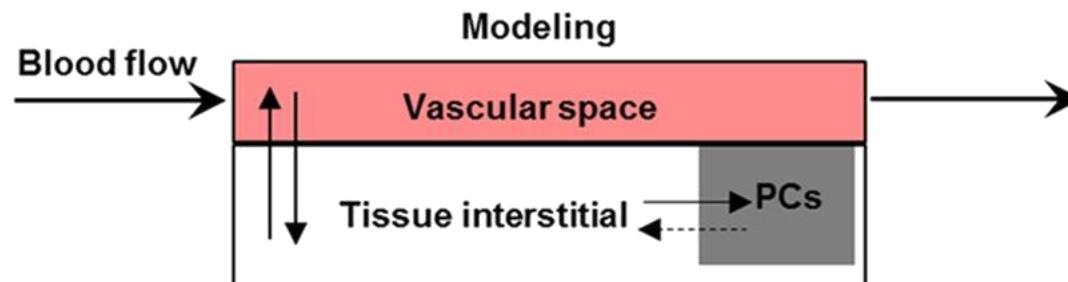
$K_{max,i}$: maximum uptake rate

$K_{50,i}$: time reaching half maximum rate

n_i : Hill coefficient



Chithrani et al. 2006. Nano Letters



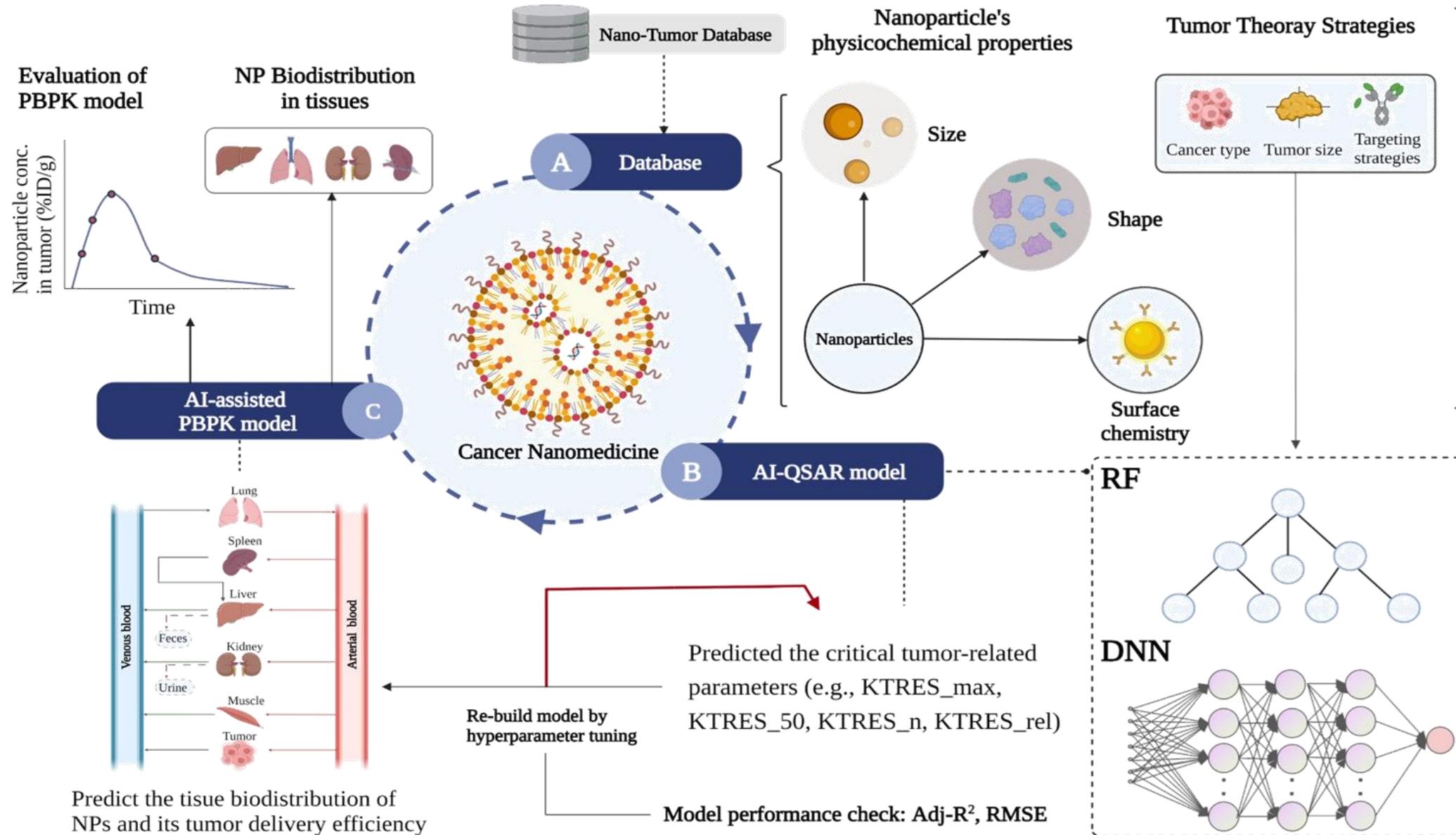
PCs represent phagocytic cells in organs or tumors;
 A_{Ti} represents amount of NPs in the tissue interstitium of the organ;
 $K_{re,i}$ is the release rate constant of NMs by PCs
Physiological based pharmacokinetic (PBPK) model

- Simplified equation in PBPK model

$$\frac{dA_{Ti}}{dt} = -K_{up,i} \times A_{Ti} + K_{re,i} \times A_{PCi}$$

Lin et al., 2016. Nanotoxicology

A hybrid method (AI-assisted PBPK model)



Variables in the Nano-Tumor Database

1. Categorical variables

- Material: Inorganic/organic NPs → 1/0
- Shape: Spherical/Rod/circle → 1...3
- Cancer type: Brain/Breast/...
- Tumor model (TM)
- Targeting strategy (TS): Active/Passive → 1/0

2. Numerical variables

- Hydrodynamic diameter [nm]
- Zeta potential [mV]

3. Target variables

- Critical Kinetic parameters related to phagocytosis

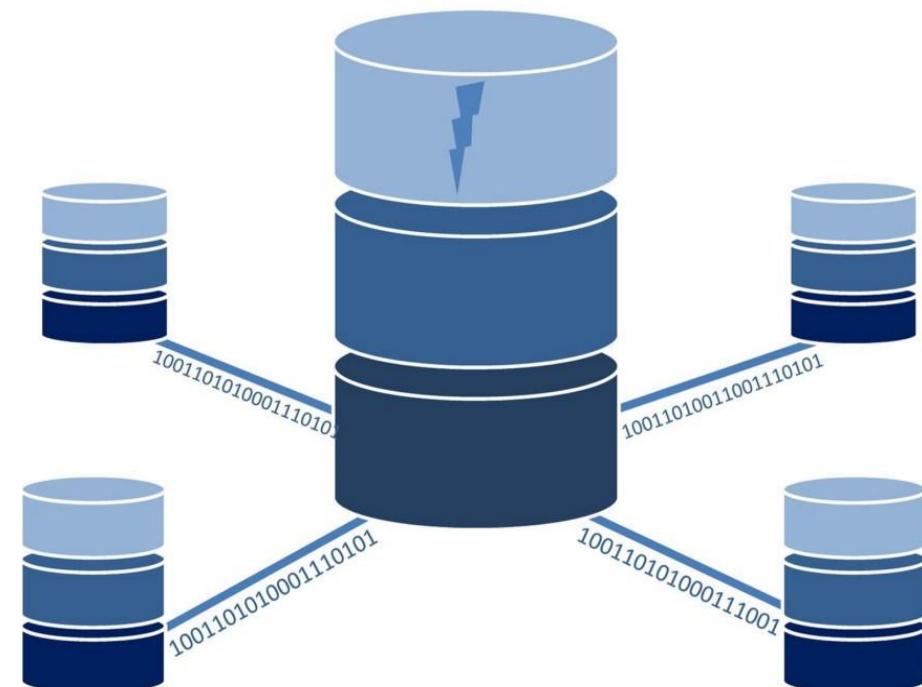
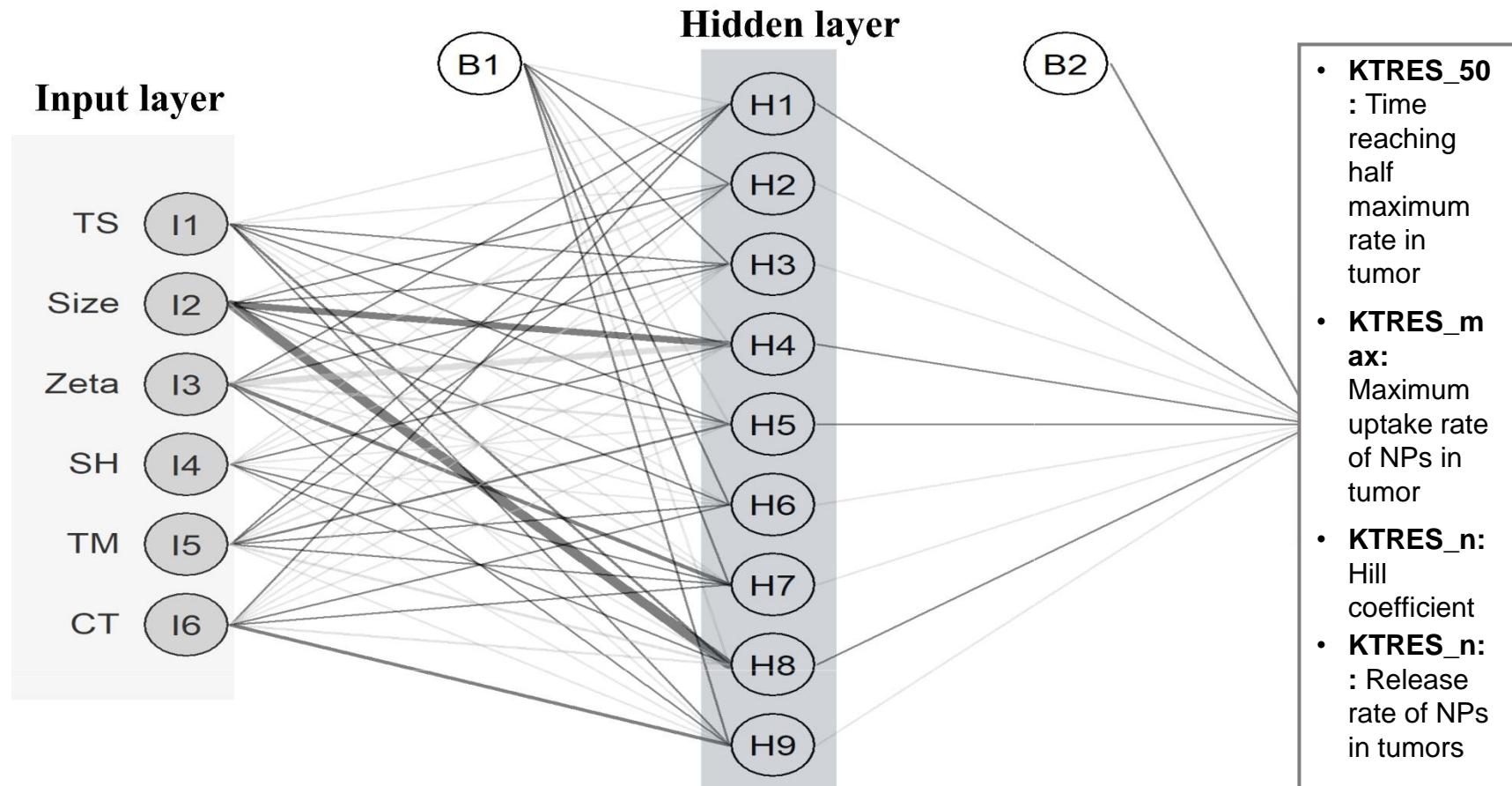


Image by Tumisu from Pixabay

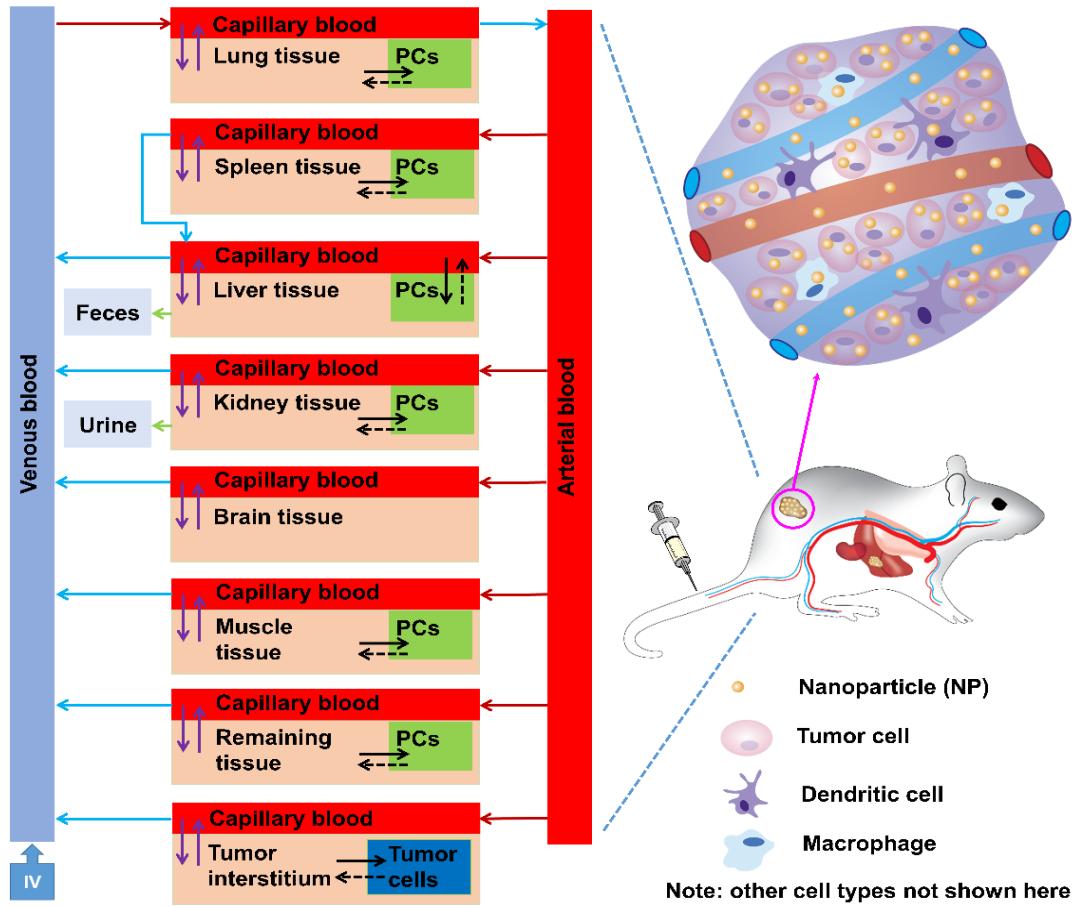
Development of AI-QSAR model



1. To avoid overfitting, we constrained the model architecture to 3 layers, which of the layers contain lower than 512 nodes
2. Shuffled 5-fold cross-validation was used to test the generalization of the model.
3. Bayesian optimization was used to tune the hyper-parameters of the models.

PBPK model for tumor-bearing mice

Physiological based pharmacokinetic (PBPK) model for tumor-bearing mice



PCs represent phagocytic cells in organs or tumors;

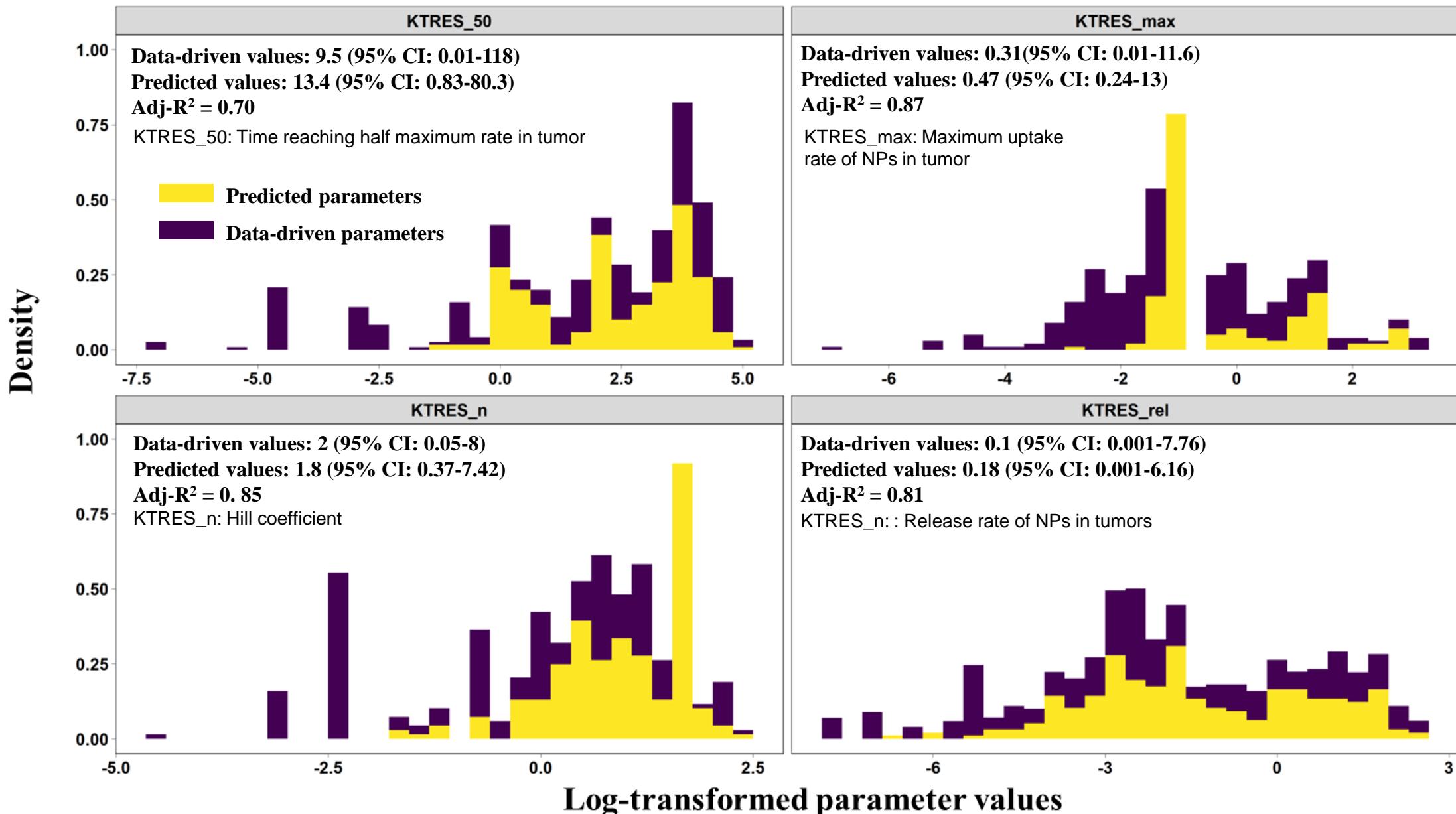
Model fitting with animal studies

Nano Tumor Database: (376 datasets from 200 studies)

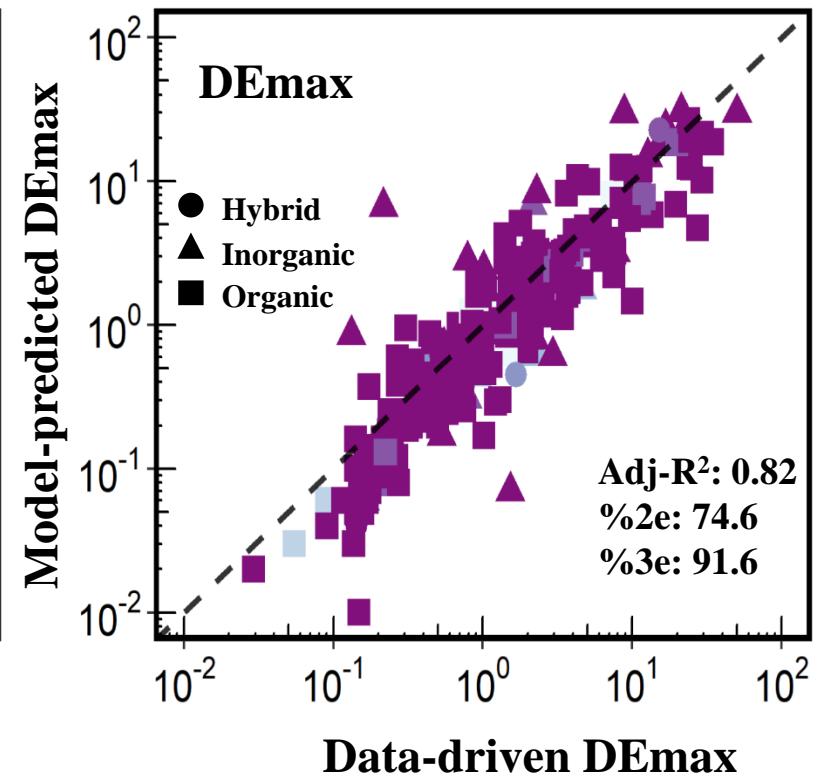
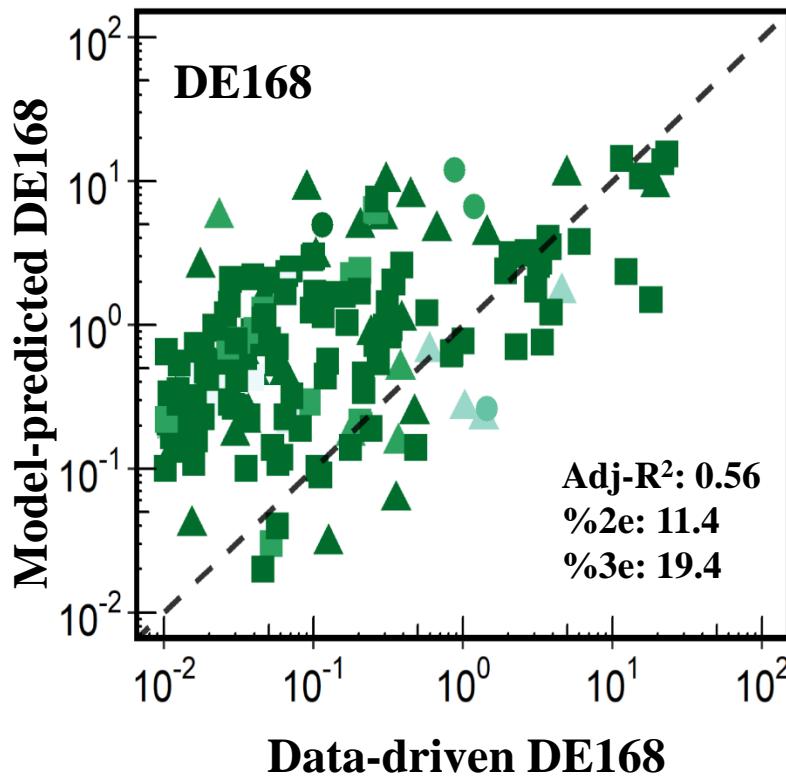
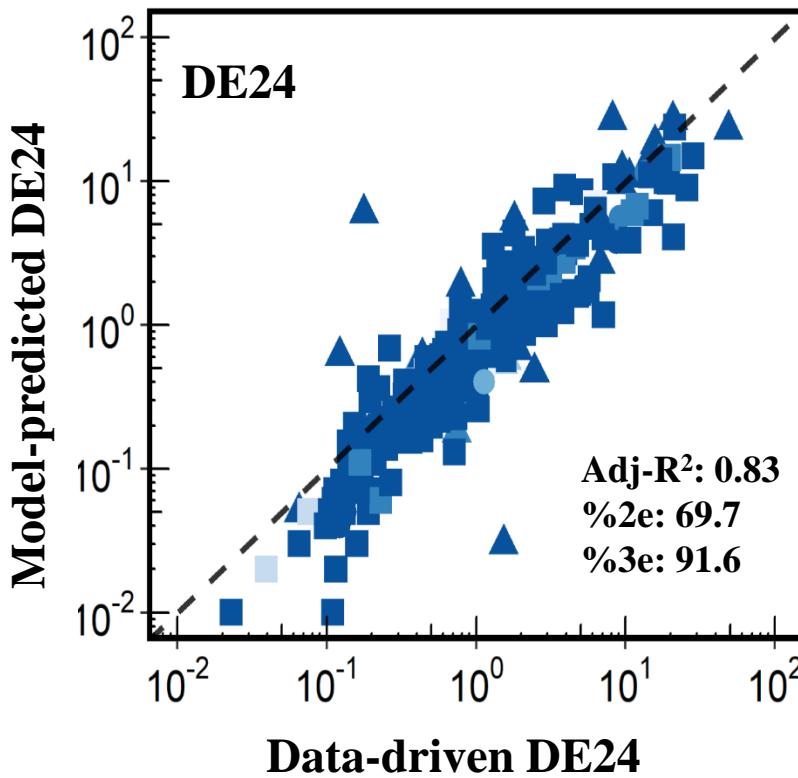
Obtain optimized model parameters

Finalized PBPK model

Similarity between predicted and data-driven parameters

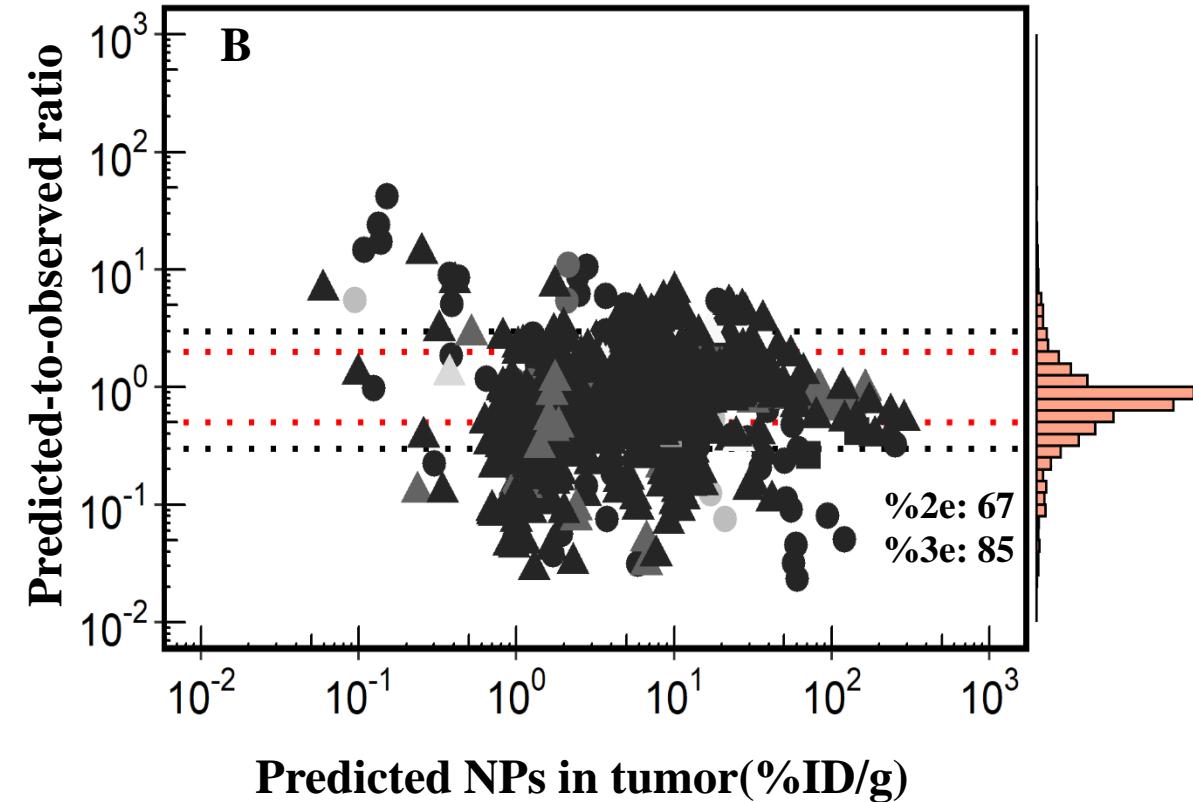
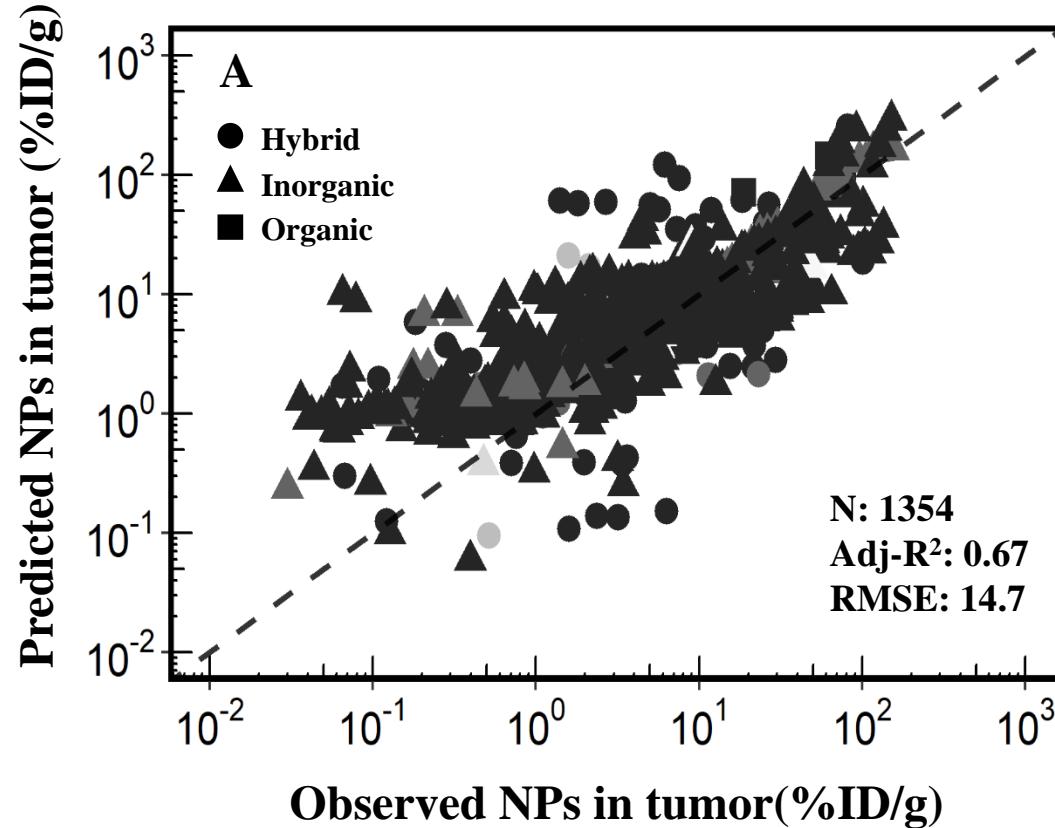


Evaluation results of AI-PBPK model-predicted tumor delivery efficiency

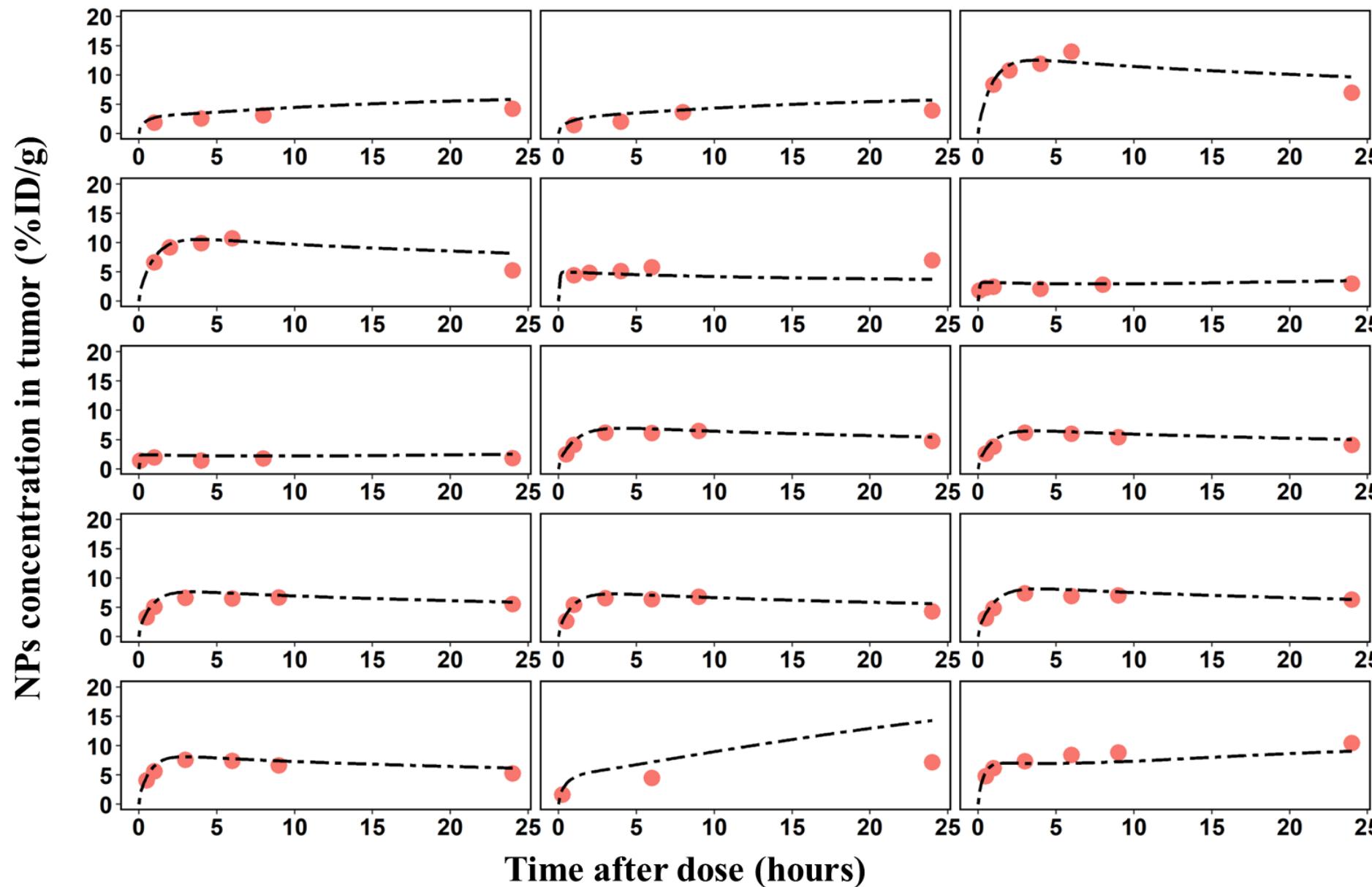


Abbreviation: DE, delivery efficiency; DE24, delivery efficiency at 24 hours;
DE168, delivery efficiency at 168 hours; DEmax, maximum of DE;
%2e, percentage of 2-fold error range
%3e, percentage of 3-fold error range

Evaluation results of AI-PBPK model-predicted time-dependent distribution of nanoparticles (NPs) to tumors



Representative evaluation results of AI-PBPK model



Summary

- This study demonstrated the feasibility of an integration of machine learning/AI technologies with a mechanistic PBPK model to predict the tumor delivery efficiency of NPs.
- Our AI-assisted PBPK model not only provides an early screening tool for estimating tumor delivery efficiency of NPs, but also can reduce the number of animals use at the early-stage preclinical trials to identify NPs with desired delivery efficiency to tumor.

Case studies

- A. Development of Artificial Intelligence (AI)-Assisted PBPK Model in Cancer Nanomedicine**

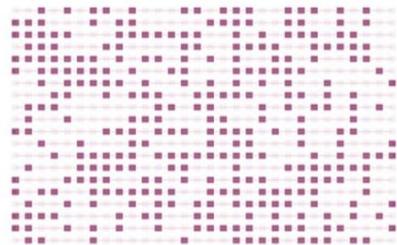
- B. Development of a Multi-Organ Toxicity Predictive Model Using Multi-Task Learning in Deep Neural Network**

Hypothesis

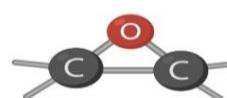
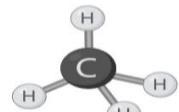
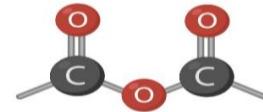
- Drug-induced organ toxicity presents a significant hurdle in drug discovery, potentially impacting multiple organs.
- Existing quantitative structure-activity relationship (QSAR) models predict single toxicity types and overlook systemic toxicity, affecting multiple organs simultaneously.
- Our hypothesis suggests that the multitask-learning QSAR model can overcome the constraints of traditional QSAR models by predicting toxicity across multiple organs.

A schematic of the multi-task QSAR model for the prediction of multi-organ toxicity.

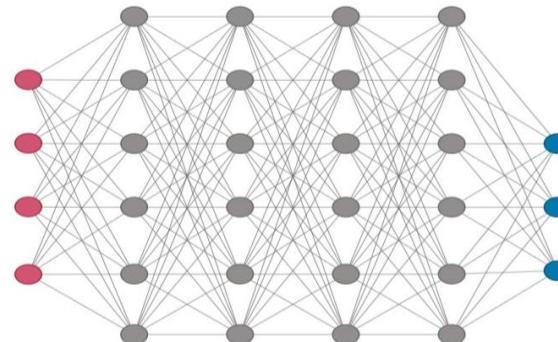
High-throughput screening (HTS) bioactivity *in vitro* data



Chemical Structure



Deep neural network



- Attribute selection
- Training
- Cross-validation
- Model assessment

Multi-organ toxicity prediction



Materials and Methods

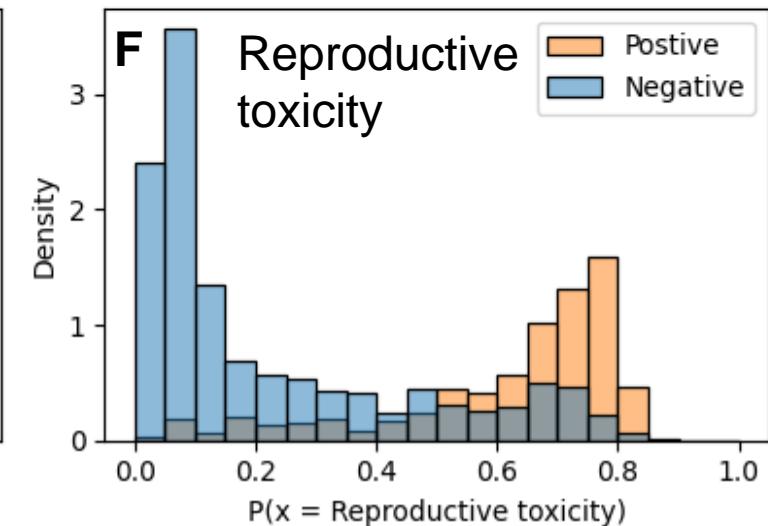
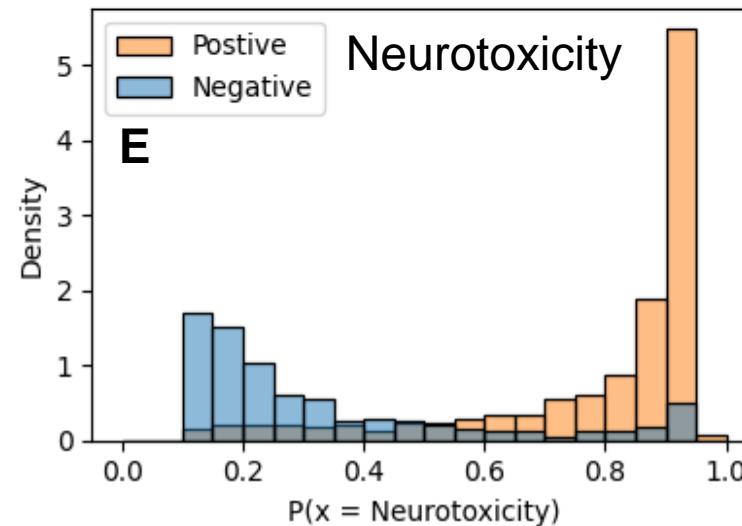
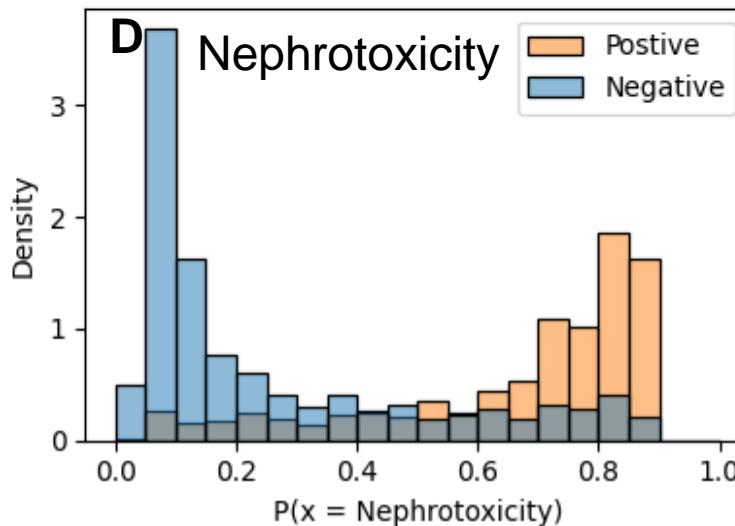
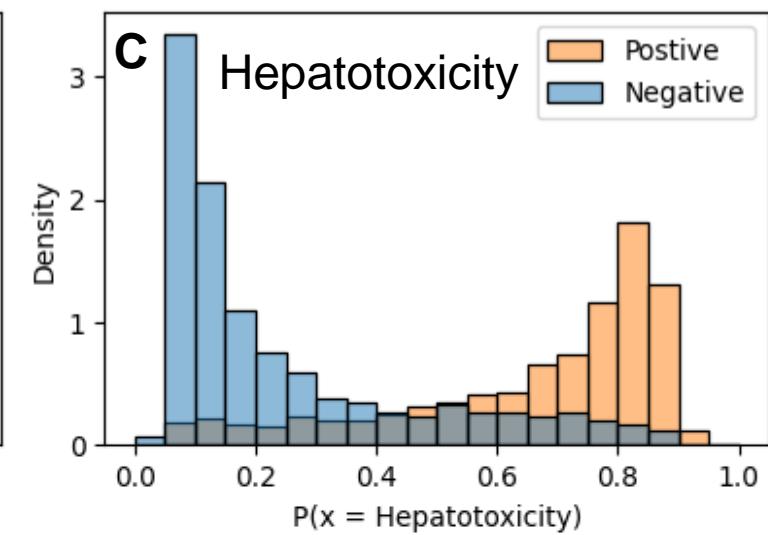
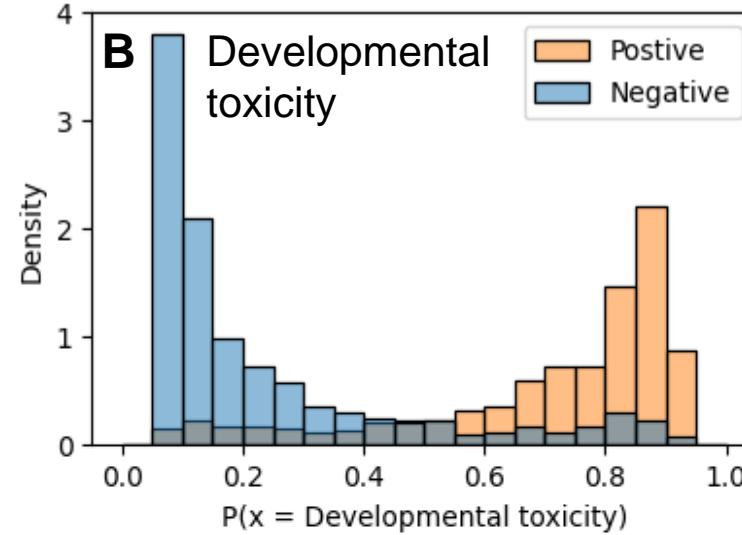
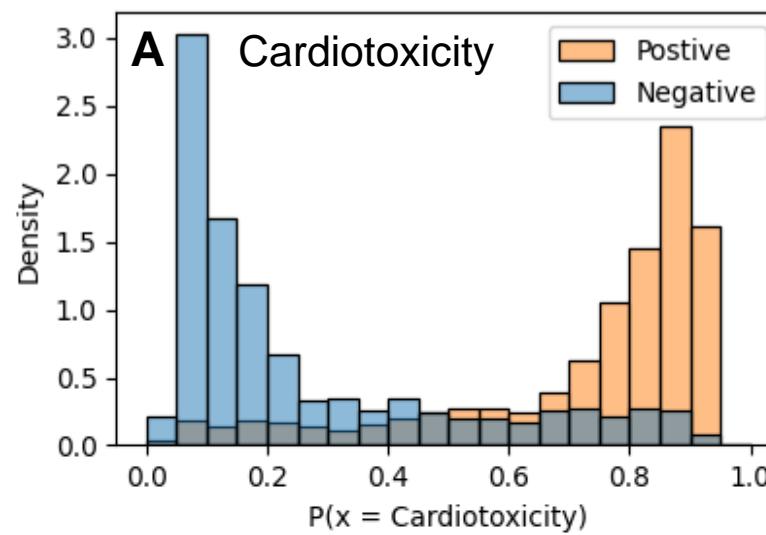
A. In vitro assays and structure data

- ❖ The *in vitro* dataset was collected from Tox21 quantitative high-throughput screening (qHTS) data via the National Center for Advancing Translational Sciences (NCTS) website: <https://tripod.nih.gov/tox21/assays/>
- ❖ In this study, we used 72 assays by filtering out the “*Chinese hamster ovary cell lines*” and other “*non-human*” originated cell lines.
- ❖ The 1024-bit ECFP4 fingerprints were generated using the Python package “RDKit”.

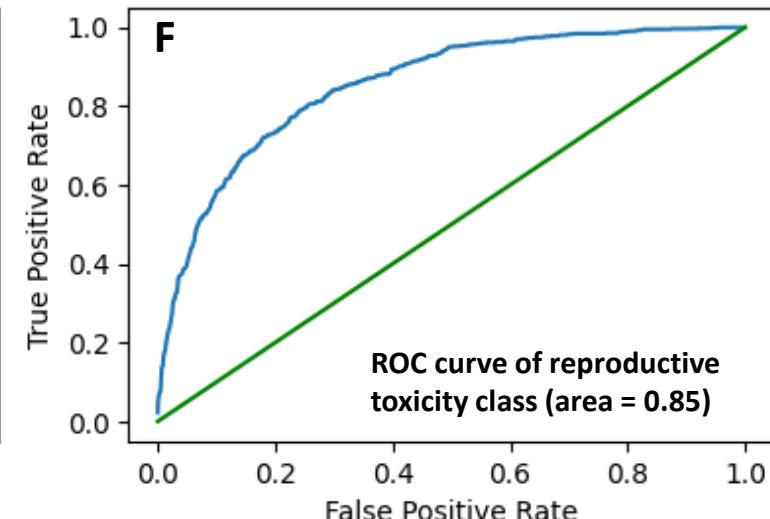
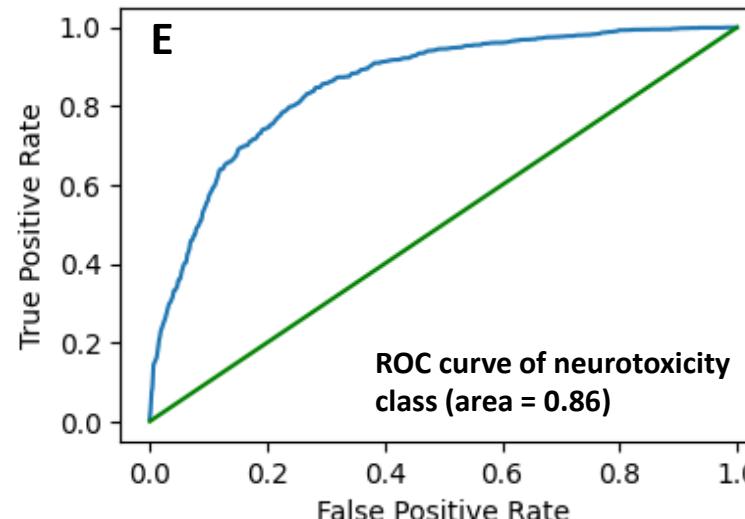
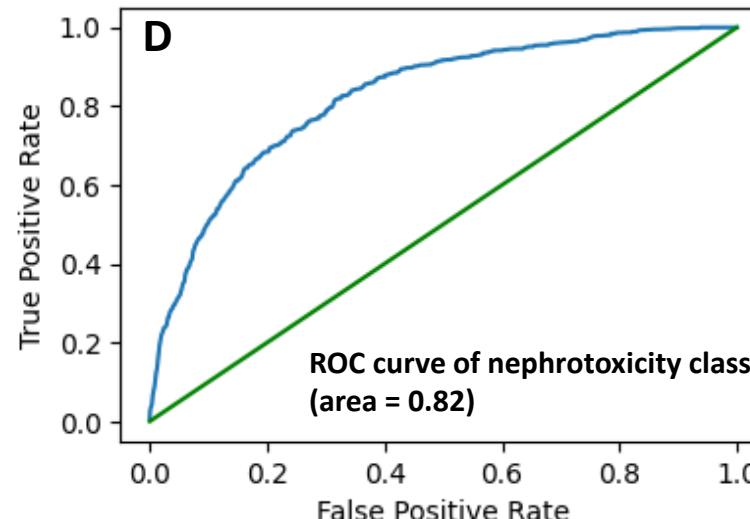
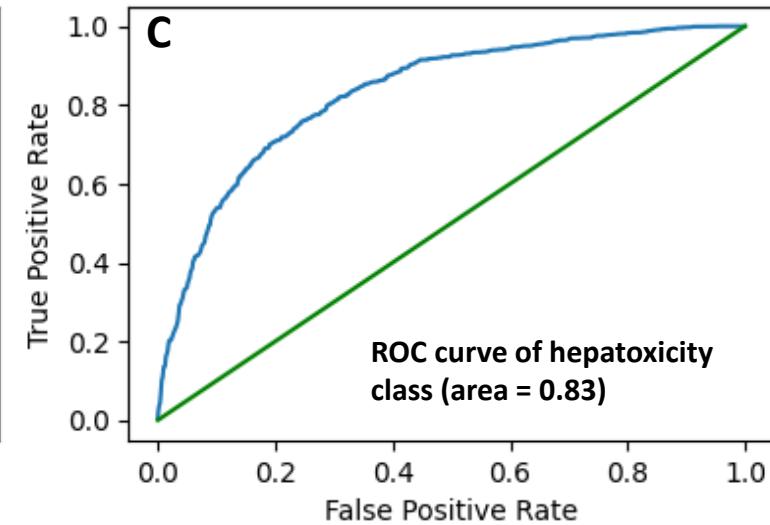
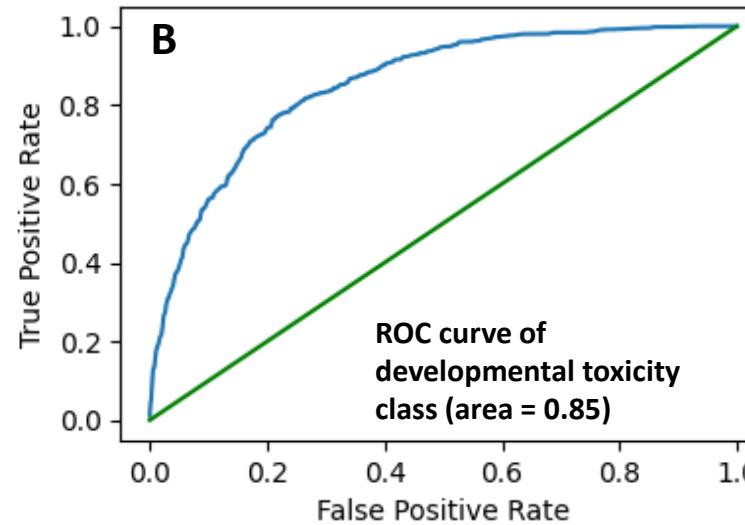
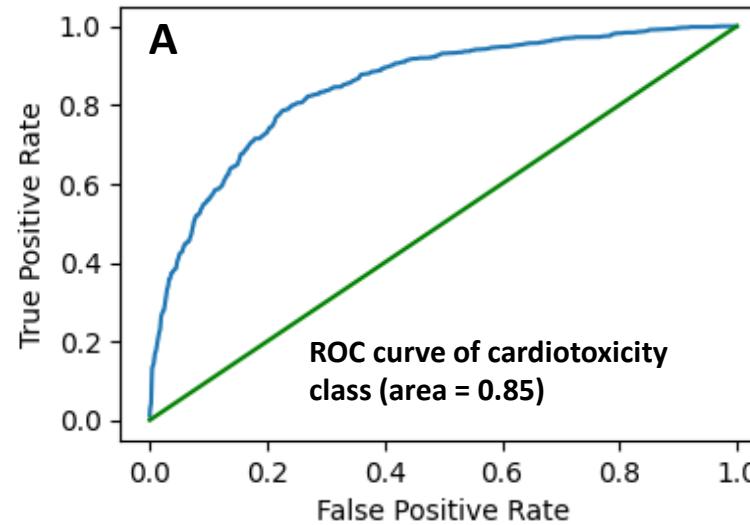
B. Human organ level toxicity data

- ❖ Human *in vivo* toxicity data in this study were collected from the studies Xu et al. (2021) and Hu et al., 2022. A total of **2,389** chemicals were collected from the database.
- ❖ Six endpoints were chosen which represent different organ-level adverse outcomes, including *cardiotoxicity*, *developmental toxicity*, *hepatotoxicity*, *nephrotoxicity*, *neurotoxicity*, and *reproductive toxicity*.

Density plot of the multi-task QSAR model



Receiver operating characteristic curve (ROC) plot of the multi-task QSAR model



Predictability of multi-task QSAR model for each toxicity endpoint

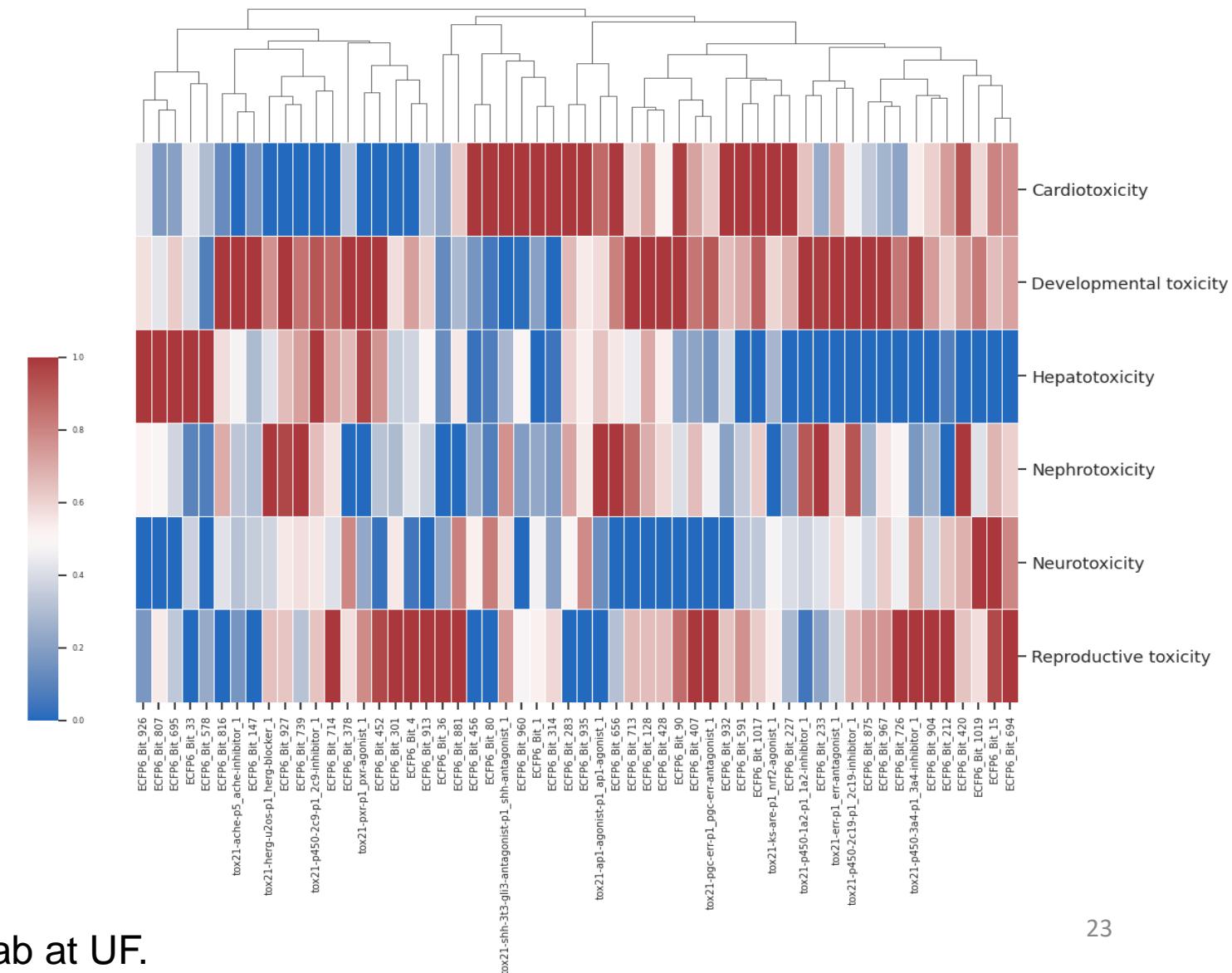
	ROC-AUC	BA	Precision	Recall	F1-Score
Cardiotoxicity	0.89	0.82	0.81	0.84	0.82
Developmental toxicity	0.90	0.84	0.83	0.82	0.83
Hepatotoxicity	0.88	0.80	0.78	0.78	0.78
Nephrotoxicity	0.87	0.80	0.77	0.79	0.78
Neurotoxicity	0.88	0.82	0.86	0.87	0.87
Reproductive toxicity	0.89	0.82	0.74	0.80	0.76
<i>Micro_Average</i>	0.88	0.82	0.80	0.82	0.81

Note: Micro-averaging values computed a global average ROC-AUC, Balanced accuracy (BA), Precision Recall and F1 score by counting the sums of the True Positives (TP), False Negatives (FN), and False Positives (FP) for each of endpoints

Summary of feature importance by using Extended-Connectivity Fingerprints (ECFPs) chemical descriptors and Tox21 assays

Tox21 assays related to organ toxicity:

- tox21-ache-p5_ache-inhibitor_1
- tox21-ap1-agonist-p1_ap1-agonist_1
- tox21-err-p1_err-antagonist_1
- tox21-herg-u2os-p1_herg-blocker_1
- tox21-ks-are-p1_nrf2-agonist_1
- tox21-p450-2c19-p1_2c19-inhibitor_1
- tox21-p450-2c9-p1_2c9-inhibitor_1
- tox21-p450-3a4-p1_3a4-inhibitor_1
- tox21-pgc-err-p1_pgc-err-antagonist_1
- tox21-pxr-p1_pxr-agonist_1
- tox21-shh-3t3-gli3-antagonist-p1_shh-antagonist_1



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National FARAD 2022

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