

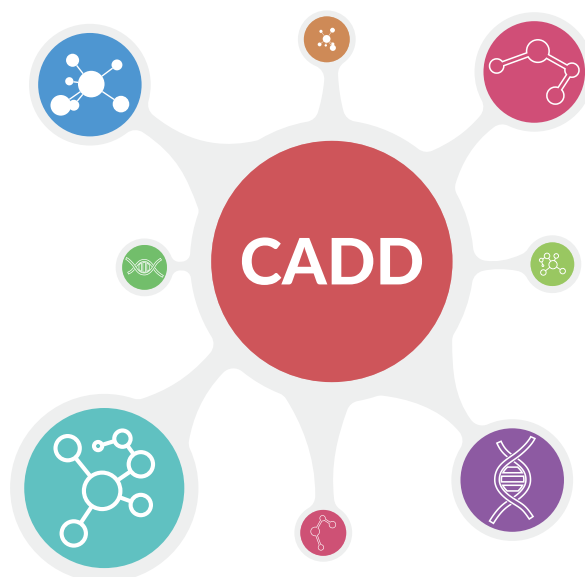


## Computer-Aided Drug Design

### Technical Description

Computer-aided drug design is based on computational chemistry and uses computer modeling techniques to simulate, calculate and predict the relationships between drugs and biological molecules in order to design and enhance lead compounds.

The most fundamental goal of CADD is to simulate the binding between targets and candidate drugs to reduce the number of compounds to be screened in vivo, thereby improving the efficiency of lead compound discovery. (In random screening, the bioactivity rate was only 0.01%-0.1%; With CADD, the bioactivity rate can reach 5%-20%)



#### Target Structure Simulation & Prediction

We offer homology modeling services to predict the 3D structure of proteins from amino acid sequences; We also provide target fishing services to find possible targets for bioactive compounds.

#### Molecular Docking/Virtual Screening

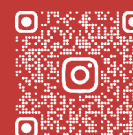
We provide molecular docking services to explore the interaction mode between protein targets and small molecules/peptides/proteins; Meanwhile, we offer virtual screening services based on target/ligand structure, thereby reducing the number of compounds to be screened by focused screening and improving the efficiency of lead compound discovery.

#### ADMET Prediction

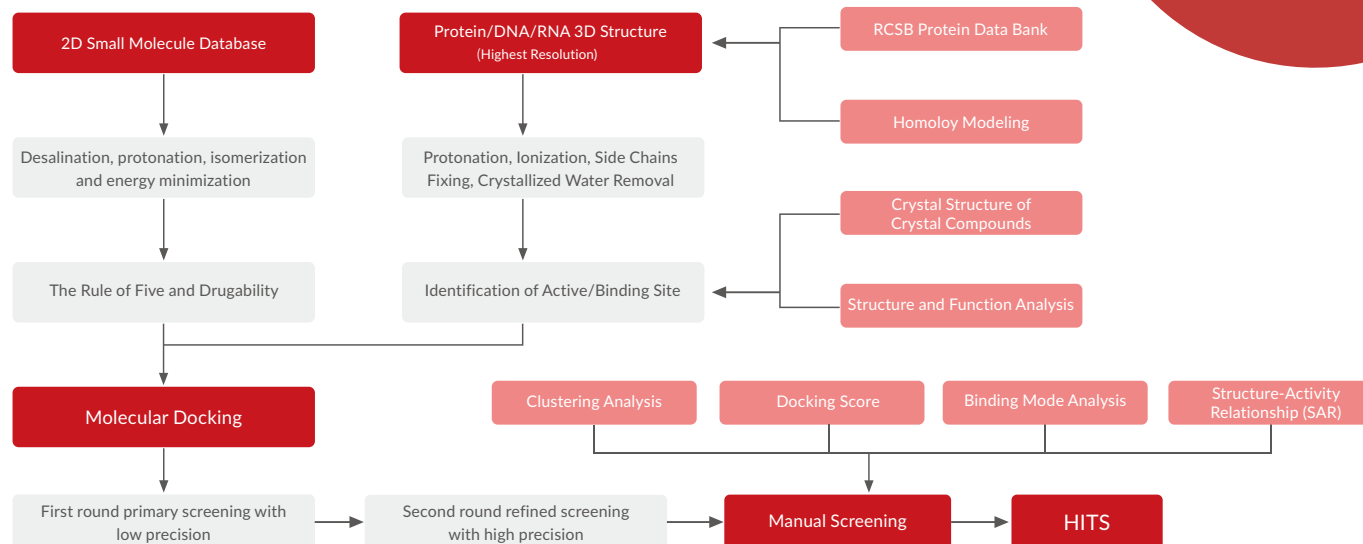
Using ADMET platform to predict the absorption, distribution, metabolism, excretion and toxicity properties of small molecules in the human body to reduce the elimination rate of candidate molecules and increase the success rate of drug development.

#### Molecular Design & Optimization/Scaffold Hopping

Based on the experiment requirements, design and optimize high-quality small molecule derivatives; Find drug molecules with novel structures through the method of scaffold hopping, and obtain hit compounds without intellectual property protection.



## II Molecular Docking-based Virtual Screening



### o Highlighted Feature

Manual screening: Based on the docking score, we selected high-affinity target compounds by taking into account the 3D structural characteristics (hydrophilicity, hydrophobicity, hydrogen bonding) of the drug binding sites. At the same time, referring to the structure-activity relationship (SAR) of existing target drugs, we reserved the most important and conservative interaction mode of compounds, explored more new target compounds by the extension of the core scaffold, and finally by cluster analysis, obtained multiple target compounds for subsequent experimental activity verification.

## III Advantages of TargetMol®



### Professional Team

Experienced technical team with more than 10 years experience in CADD



### Compound Library

A stable supply of 20,000+ compounds to meet the needs of different clients



### Screening Process

Additional manual screening significantly improves the reliability of simulated results



### Screening Plan

Informative evaluation and closure reports to help clients fully understand the screening process and results



### Featured Packages

Specialized natural compound and drug repurposing libraries, with well-defined targets and low prices



### High Cost Effectiveness

Virtual screening at a cost much lower than the market average



### Results Assurance

Free HD display of binding to meet publication requirements



### Licensed Software

Multiple licensed professional CADD software, clients are safe from software intellectual property rights issues

Targetmol Chemicals Inc.

— Drug Screening Expert (Inhibitors, Natural Products, Compound Libraries)

\* All products are for Research Use Only. Not for Human or Veterinary or Therapeutic Use.