

Volgograd State Medical University

Research Center of Innovative Medicines

***Laboratory for Information Technology in Pharmacology
and Computer Modeling of Drugs***



MULTI-TARGET NEURAL NETWORK
MODEL OF ANXIOLYTIC ACTIVITY
OF CHEMICAL COMPOUNDS BASED
ON CORRELATION CONVOLUTION OF
ENERGY SPECTRA OF MULTIPLE
DOCKING

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WHO – Anxiety disorders



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Anxiety disorders

27 September 2023

301 million people

Key facts

- Anxiety disorders are the world's most common mental disorders, affecting 301 million people in 2019.
- More women are affected by anxiety disorders than men.
- Symptoms of anxiety often have onset during childhood or adolescence.
- There are highly effective treatments for anxiety disorders.
- Approximately 1 in 4 people with anxiety disorders receive treatment for this condition.

Overview

Everyone can feel anxious sometimes, but people with anxiety disorders often experience fear and worry that is both intense and excessive. These feelings are typically accompanied by physical tension and other behavioural and cognitive symptoms. They are difficult to control, cause significant distress and can last a long time if untreated. Anxiety disorders interfere with daily activities and can impair a person's family, social and school or working life.

An estimated 4% of the global population currently experience an anxiety disorder (1). In 2019, 301 million people in the world had an anxiety disorder, making anxiety disorders the most common of all mental disorders (1).

Although highly effective treatments for anxiety disorders exist, only about 1 in 4 people in need (27.6%) receive any treatment (2). Barriers to care include lack of awareness that this is a treatable health condition, lack of investment in mental health services, lack of trained health care providers, and social stigma.

العربية

中文

Français

Русский

Español

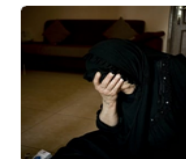
Related

[Comprehensive mental health action plan 2013–2030](#) >

[Mental Health Gap Action Programme \(mhGAP\)](#) >

[More on mental health](#) >

Fact sheets



Mental disorders

8 June 2022



Depressive disorder 3

Purpose

To construct a multi-target model of the dependence of the anxiolytic activity of chemical compounds on their multiple docking energies using correlation convolution and artificial neural networks

Tasks

- **Construction of optimized 3D models of ligands**
- **Selection and validation of relevant biotargets**
- **Formation of spaces for multiple docking**
- **Performing multiple ensemble docking to relevant biotargets**
- **Correlation convolution of multiple docking energy spectra**
- **Formation of training set**
- **Training of neural networks and selection of the best model**

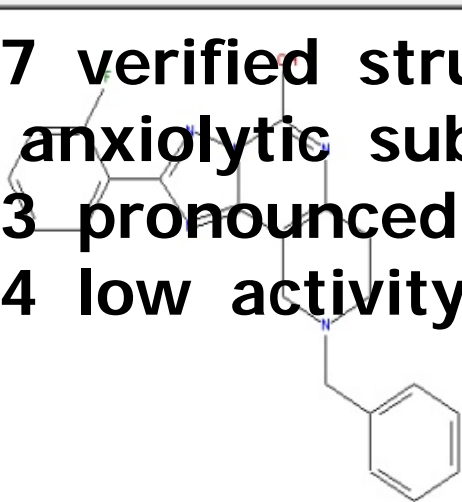
Verified database of anxiolytic substances

Anxiolytic Substances v11 - Corrected

24 January 2022

Structure

537 verified structures
of anxiolytic substances:
273 pronounced activity
264 low activity



Activity

Standard Type

ED50

pChEMBL Value

Assay

Assay ChEMBL ID

CHEMBL779993

Assay Description

Anxiolytic response measurement in rats.

Mol_ID Brutto formula Mol weight Salt Component

1

C₂₁H₁₈FN₅O

375.3

Compound Codes & Names

MolFileName

CHEMBL99259

CODE - Molecule ChEMBL ID

CHEMBL99259

Molecule Name

Compound Key

8g

Activity Processing

Gen_Assay_IDs

Rat-07a

Gen_Assay_CHEMBLID

CHEMBL3256343 CHEMBL784521 CHEMBL8

References

Document_ChEMBL_ID

CHEMBL1125833

Document Journal

J. Med. Chem.

Document_Year

1991

15.07.2022

РОССИЙСКАЯ ФЕДЕРАЦИЯ



СВИДЕТЕЛЬСТВО

о государственной регистрации базы данных

№ 2022621744

Соединения с анксиолитической активностью

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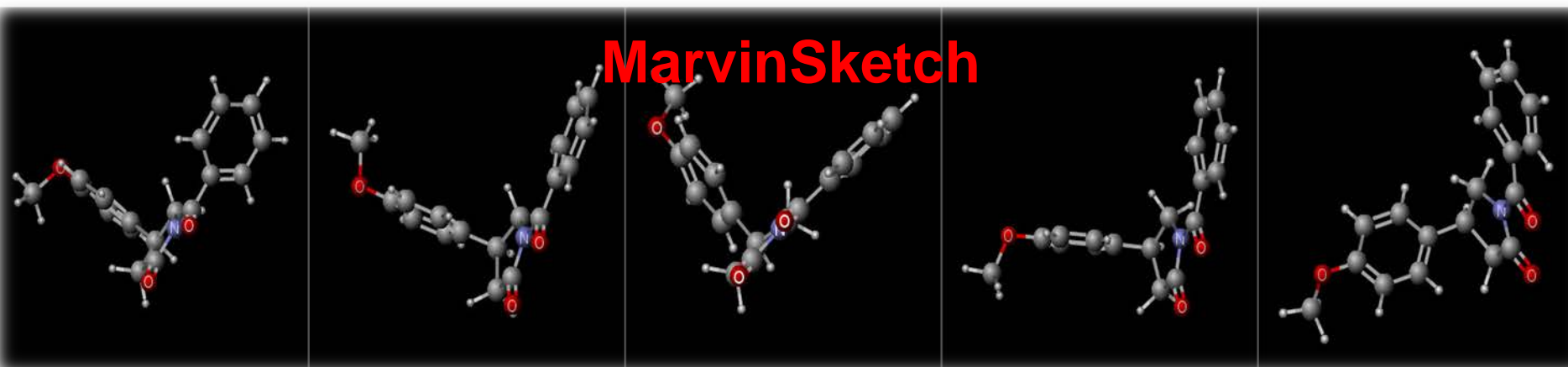
Руководитель Федеральной службы по интеллектуальной собственности

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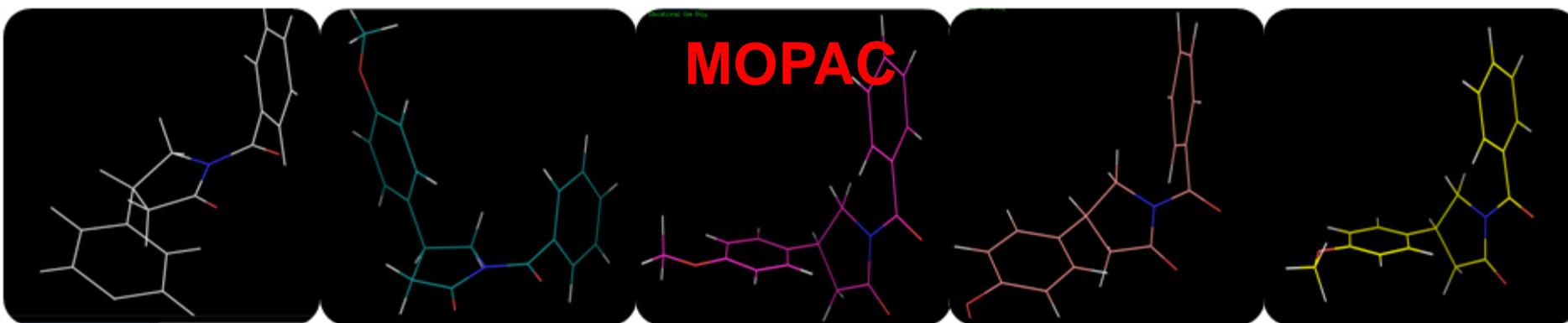


Construction of optimized 3D models of ligands

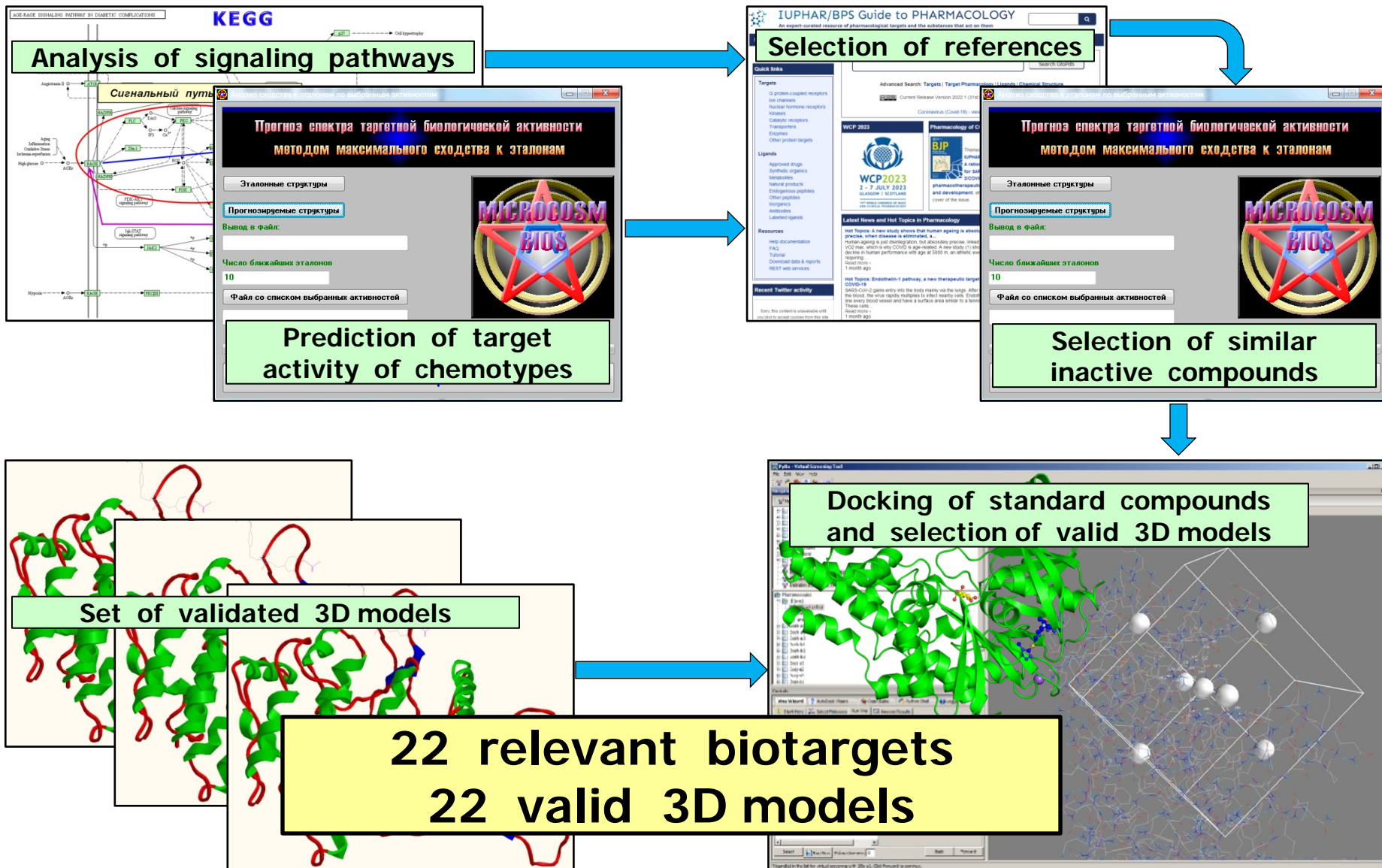
Molecular mechanics 10 conformers



Quantum chemistry PM7

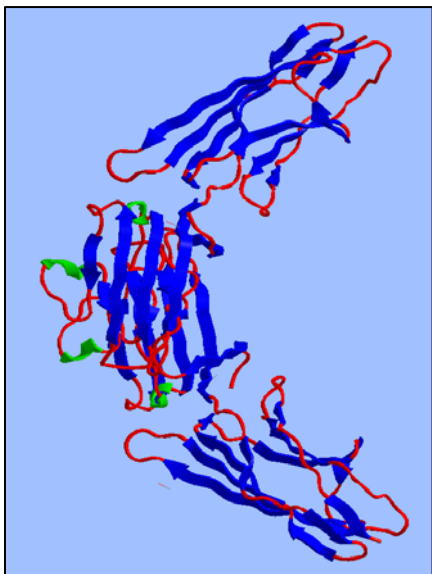


Selection and validation of relevant biotargets



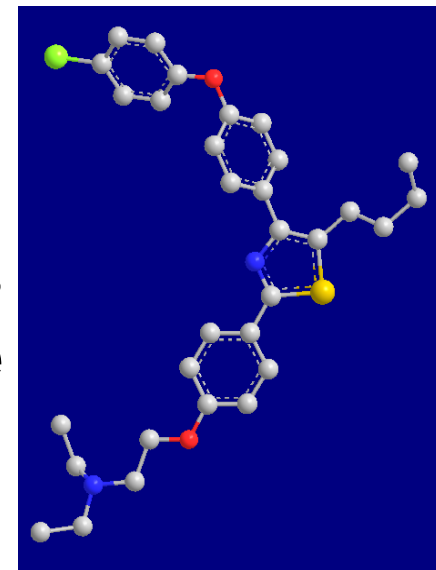
Multiple docking

Ligands interact with the protein throughout its entire volume

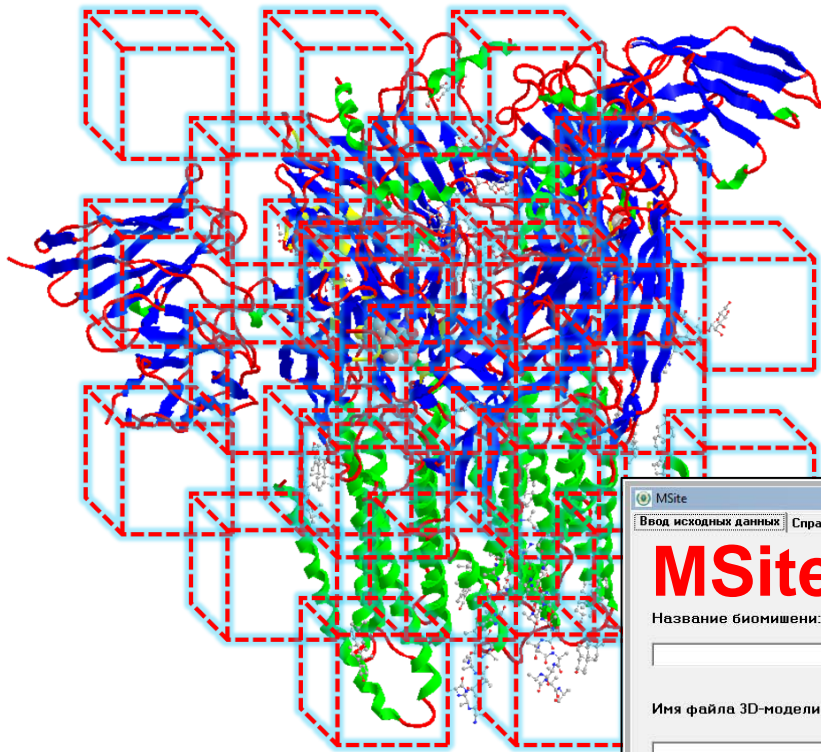


- Active concentration 10^{-10} M.
Avogadro's number $\sim 6 \cdot 10^{23}$.
 **$\sim 6 \cdot 10^{13}$ ligand molecules
interact with the target protein!**

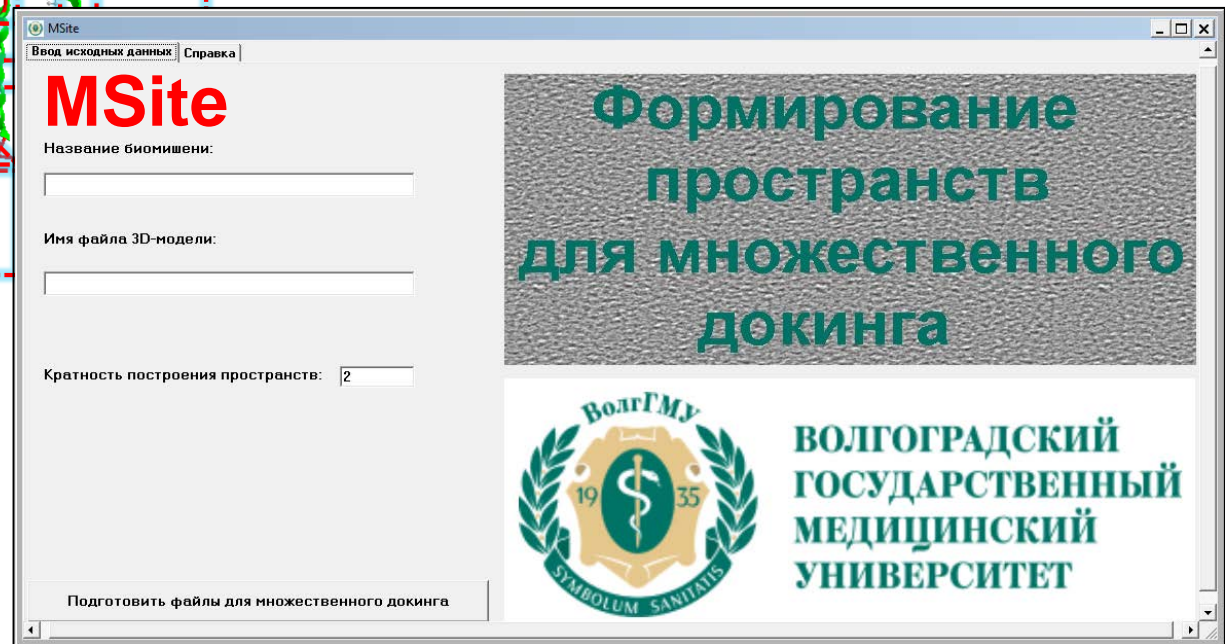
- RAGE volume $34\,902 \text{ \AA}^3$.
Ligand volume $1\,578 \text{ \AA}^3$.
**The ligand fits
into the receptor space
at least 22 times!**



Formation of spaces for multiple docking

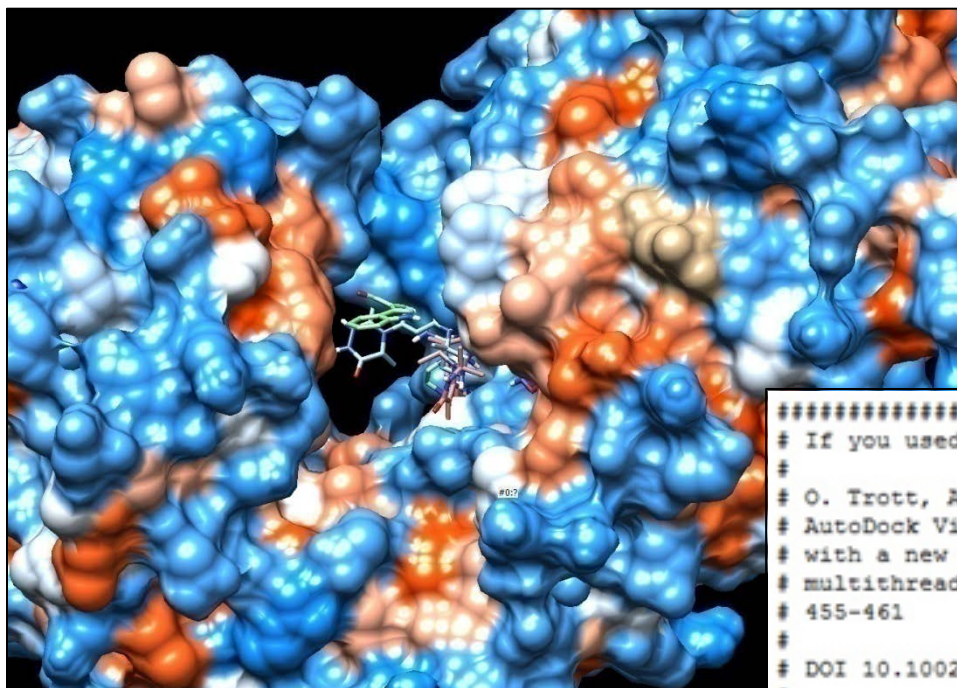


$3 \cdot 3 \cdot 3 = 27$ spaces



The screenshot shows the MSite software interface. The title bar reads "MSite". Below the title bar, there are two tabs: "Ввод исходных данных" and "Справка". The main content area displays the "MSite" logo in red. Below the logo, there are three input fields: "Название биомшени:" (empty), "Имя файла 3D-модели:" (empty), and "Кратность построения пространств:" with a value of "2". At the bottom, there is a button labeled "Подготовить файлы для множественного докинга". On the right side of the interface, there is a large text area with the title "Формирование пространств для множественного докинга" in green. Below this text area is the logo of Volgograd State Medical University, which includes the text "ВолГМУ" and "19 35" inside a shield, and "СYMBOLUM SANITATIS" at the bottom. To the right of the logo is the text "ВОЛГОГРАДСКИЙ ГОСУДАРСТВЕННЫЙ МЕДИЦИНСКИЙ УНИВЕРСИТЕТ".

Multiple ensemble docking to relevant biotargets



Each of 537 ligands
to each of 22 3D models
to each of 27 spaces
5 times in 10 conformations

$$\Delta E_{ijkl} = \min(\Delta E_{ijklm})$$

Total processed
 $537 \cdot 22 \cdot 27 \cdot 5 \cdot 10$
~ 16 000 000
docking energies

AutoDock Vina

```
#####  
# If you used AutoDock Vina, please cite:  
#  
# O. Trott, A. J. Olson,  
# AutoDock Vina: improving the speed and accuracy of docking  
# with a new scoring function, efficient optimization and  
# multithreading, Journal of Computational Chemistry 31 (2010)  
# 455-461  
#  
# DOI 10.1002/jcc.21334  
#  
# Please see http://vina.scripps.edu for more information.  
#####  
  
Output will be .....\Ligands\Flavopiridol-PM7_out.pdbqt  
Detected 12 CPUs  
WARNING: at low exhaustiveness, it may be impossible to utilize all CPUs  
Reading input ... done.  
Setting up the scoring function ... done.  
Analyzing the binding site ... done.  
Using random seed: 58910256  
Performing search ...  
0% 10 20 30 40 50 60 70 80 90 100%  
|----|----|----|----|----|----|----|----|----|  
#####
```

Correlation convolution

$$R_{ijkl} = \frac{Cov_{ijkl}}{S_{ikl} \cdot S_{jkl}}, \quad i, j = 1 \dots M, \quad k = 1 \dots K, \quad l = 1 \dots L, \quad i \neq j$$

$$W_{kn} = \frac{1}{2} \sum_{\substack{l=1 \\ l \neq i, j}}^L \sum_{\substack{i=1 \\ i \neq j}}^M R_{ijkl} \cdot X_{ilkn} \cdot X_{jlkcn}, \quad k = 1 \dots K, \quad n = 1 \dots N$$

$K = 22$, number of biotargets

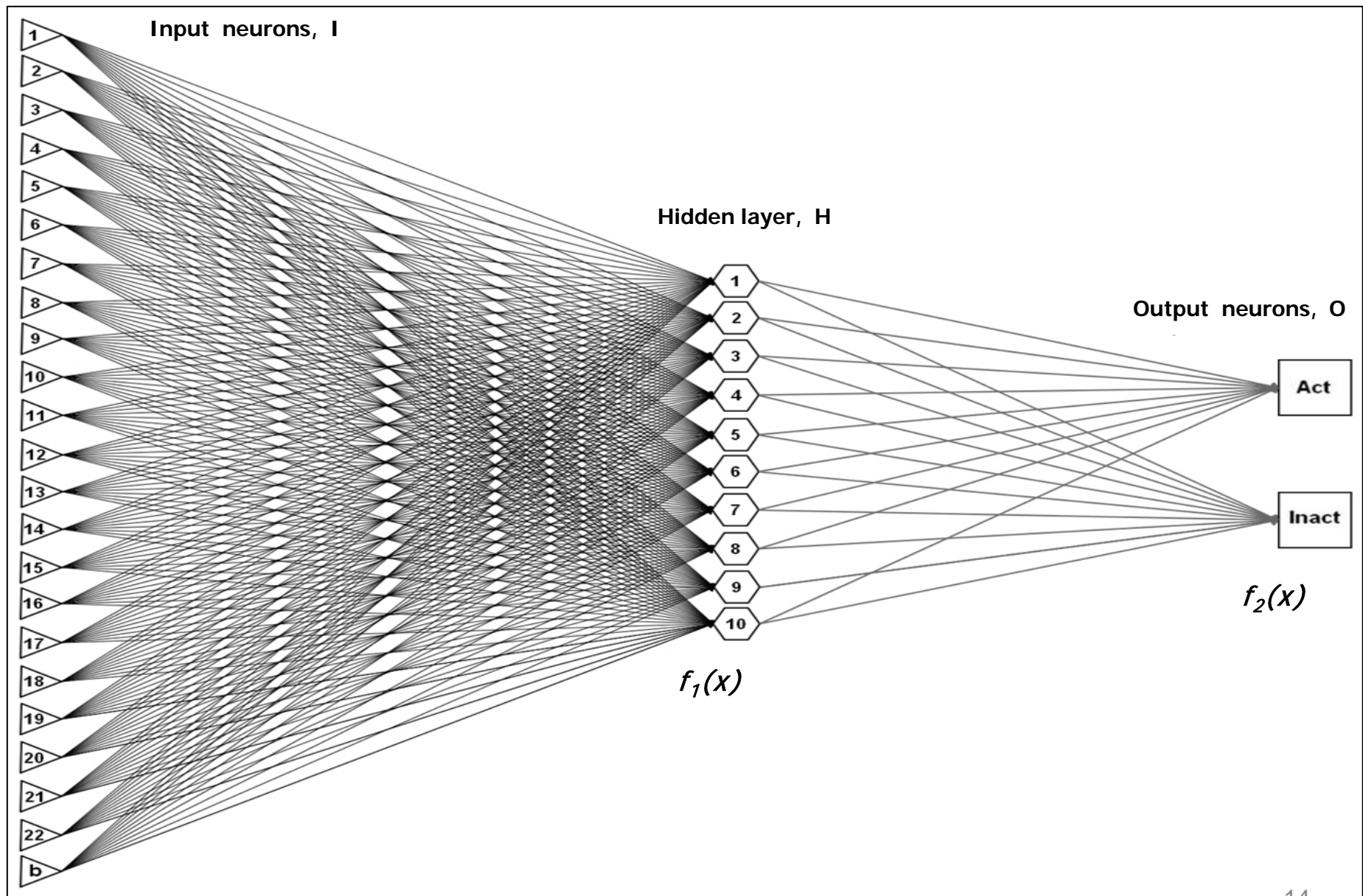
$M = 27$, number of docking spaces

$L = 5$, number of docking energies for space i

$N = 537$, number of substances

2970 docking energies \longrightarrow 22 convolution variables

Neural network with bottle-neck



Training of neural networks

7 sampling options

Net. ...	Net. name	Training ...	Test p...	Algorit...	Error fu...	Hidden ...	Output ...
1	MLP 22-1...	78.911565	80.555...	BFGS 14	SOS	Tanh	Logistic
2	MLP 22-1...	78.911565	80.555...	BFGS 14	SOS	Exponen...	Logistic
3	MLP 22-1...	79.591837	80.555...	BFGS 14	SOS	Exponen...	Tanh
4	MLP 22-1...	78.911565	80.555...	BFGS 14	SOS	Logistic	Logistic
5	MLP 22-9-L	78.231233	58.335...	BFGS 11	SOS	Identity	Tanh
6	MLP 22-1...	78.911565	80.555...	BFGS 9	SOS	Tanh	Logistic

4000 trained neural networks for each sampling option

50 automatically selected neural networks for each sampling option

Neural network training in progress...

Building network 145 (MLP 22-16-2, exp, logistic)
Cycle=40: **1 best neural network**
Classification rate: Train=85.034, Test=75

~30 000 networks were trained

Options

- Train
- Test
- Validation
- Missing

Laboratory for Information Technology in Pharmacology and Computer Modeling of Drugs



~40 Tflops

Creative team

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Thank You for Your
attention!