

NEW INHIBITORS OF THE COAGULATION FACTOR XIIA: DOCKING AND EXPERIMENTAL VERIFICATION

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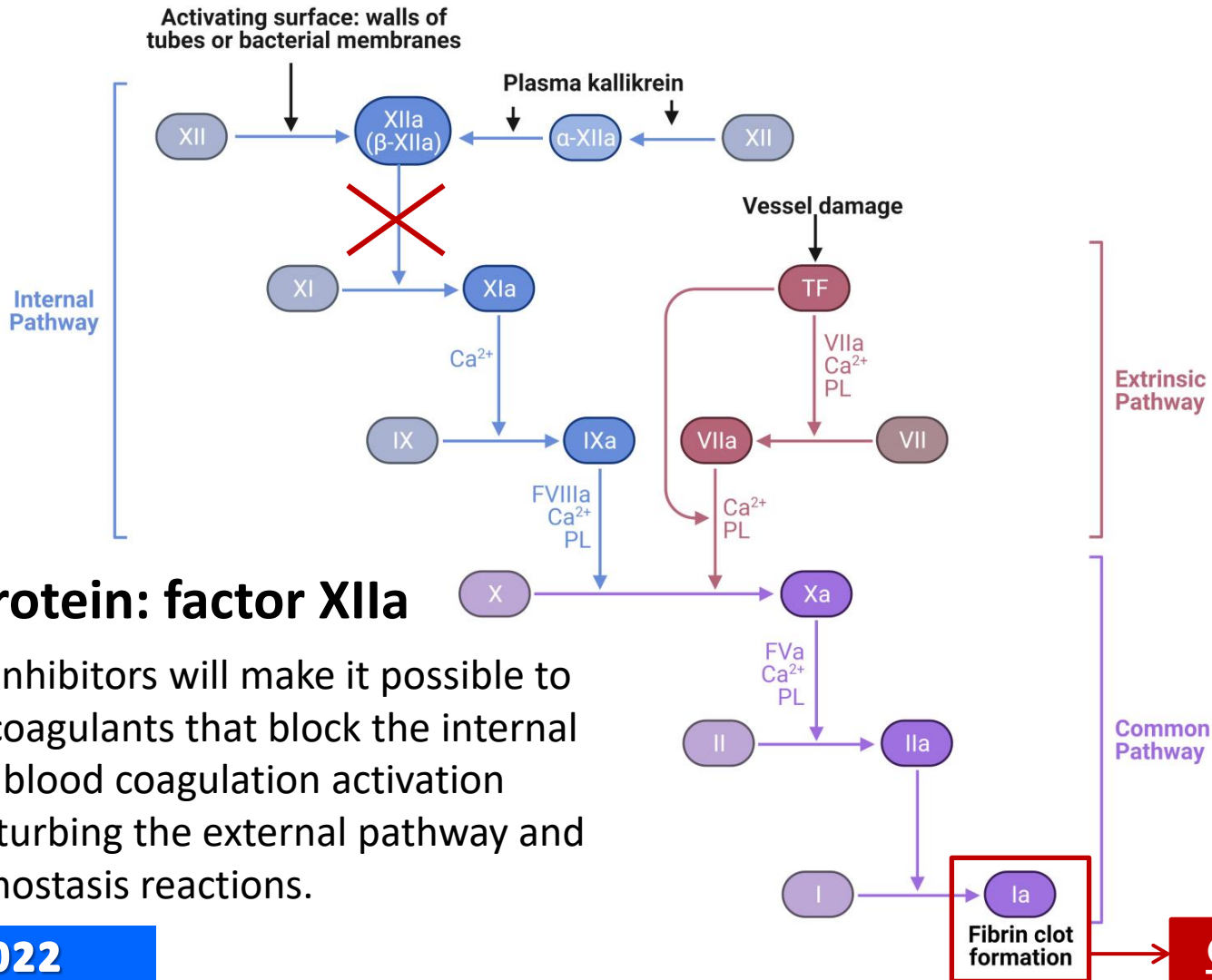
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Reactions of the plasma blood coagulation system



Target protein: factor XIIa

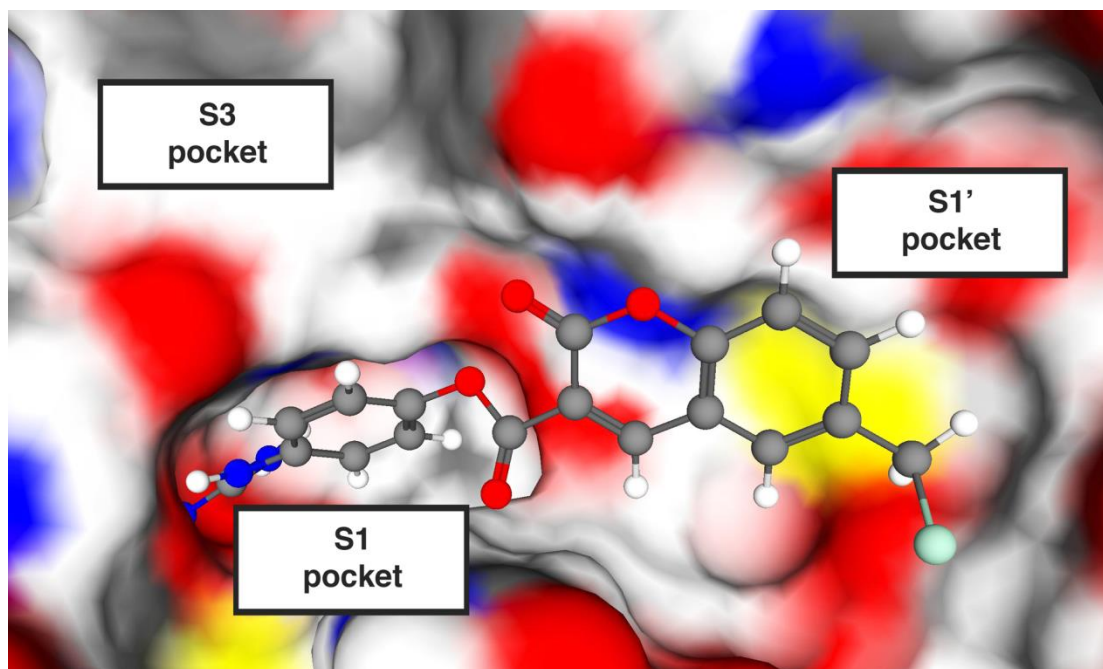
Factor XIIa inhibitors will make it possible to create anticoagulants that block the internal pathway of blood coagulation activation without disturbing the external pathway and plasma hemostasis reactions.

Anticoagulants on the basis of factor XIIa inhibitors

- ▶ Blocking the internal pathway of blood coagulation activation without disturbing the external pathway and plasma hemostasis reactions
- ▶ With factor XII deficiency, laboratory animals are either completely protected from the development of induced thrombosis or suffer significantly less from its consequences
- ▶ FXII-deficient patients demonstrate a normal hemostatic ability
- ▶ Factor XII deficiency protects against thrombosis without causing spontaneous bleeding

Factor Xlla model

Development of new inhibitors of blood coagulation factor Xlla using molecular modeling methods



Active site of the protein is represented by binding pockets S1, S1' and S3

Protein model

- ▶ Novel PDB structure **6B74** of factor Xlla crystallized with **inhibitor**.
Resolution = 2.3 Å
- ▶ adding hydrogen atoms using the APLITE program
- ▶ native ligand docking
- ▶ development of selection criteria

Virtual screening

Creating a protein model

Creating 3D structures of
chemical compounds

SOL program docking $\sim 10^4$

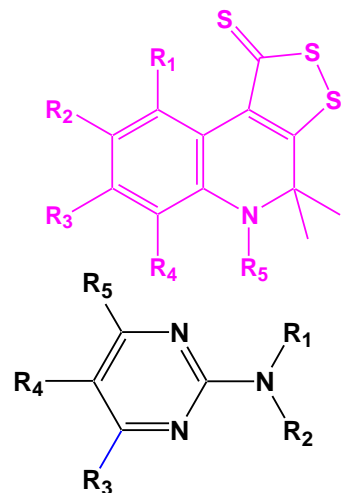
Binding enthalpy calculations by the quantum-chemical PM7
semi-empirical method with the COSMO solvent model
 $\Delta H = H_{complex} - H_{protein} - H_{ligand}$: $\sim 10^2 - 10^3$

Experimental *in vitro* verification (≈ 20 compounds)

Database of drug-like compounds of Voronezh State University

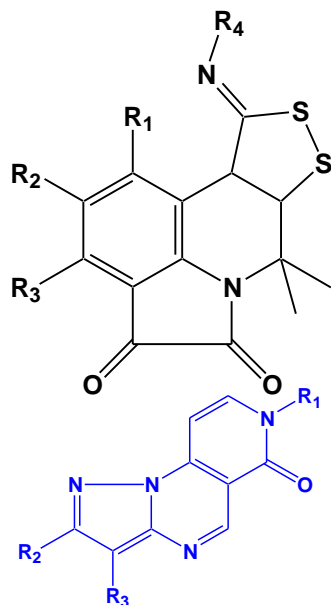
More than 200 000 compounds – virtual, but have been synthesized

Antibacterial,
antifungal activity



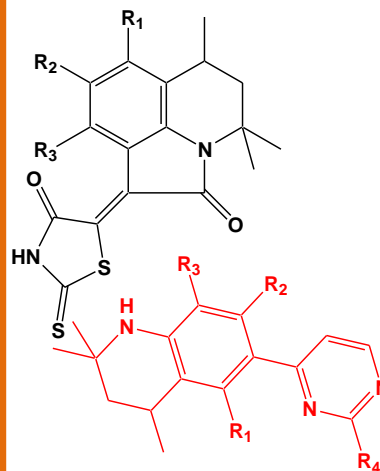
Derivatives of
hydroquinoline,
quinolinthiones,
pyrimidines,
quinazolines

Protein kinase
inhibitors



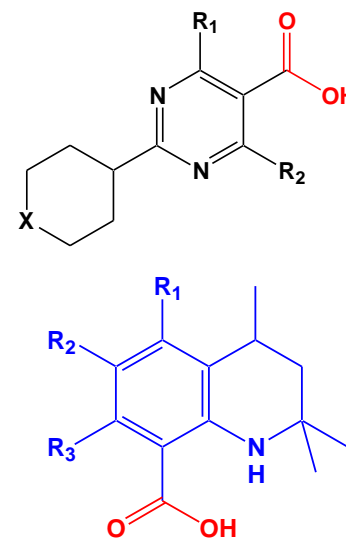
Derivatives of
aminopyrimidines,
aminoquinazolines,
pyrroloquinolines

Anticoagulants



Hydroquinoline
derivatives linearly
bonded and fused to
other heterocycles

Plant growth
regulators

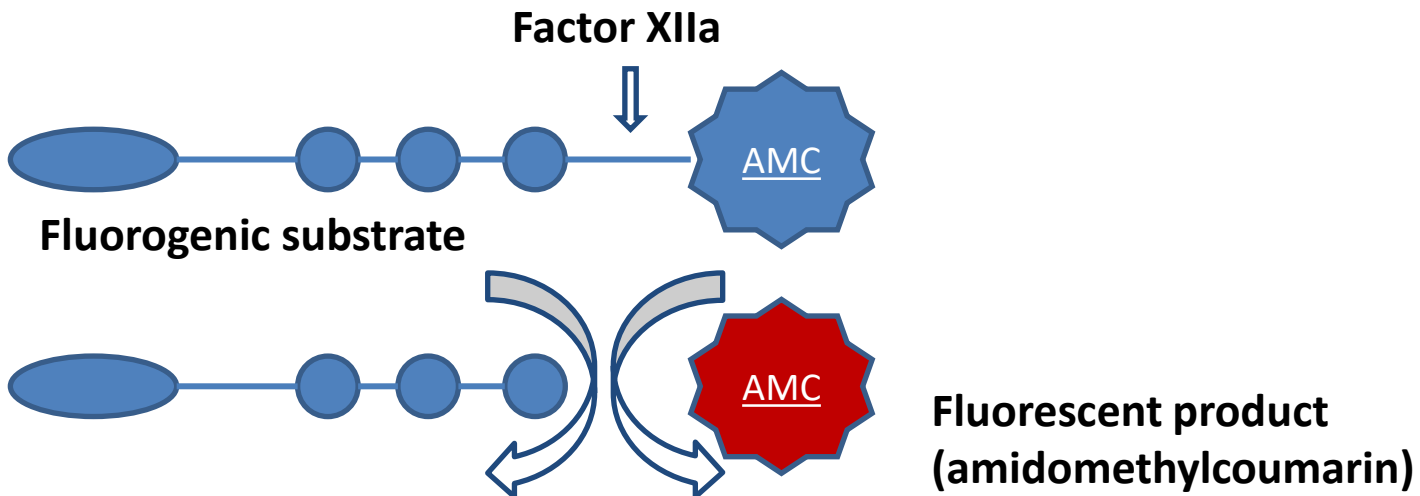


Derivatives of
hetarylcarboxylic
acids

In vitro verification

Experiments carried out in the laboratory of biophysics of the NMIC
DGOI named after Dmitry Rogachev

Substrate hydrolysis reaction with coagulation factor XIIa



Factor XIIa: final concentration 5 nM

Substrate S2302 (Chromogenics, Italy): final concentration 200 μ M

Test compounds: concentration 30 μ M

A library of chemical compounds

Focused library of chemical compounds - 59 molecules – 326 3D-structures

Selection of compounds for experimental study

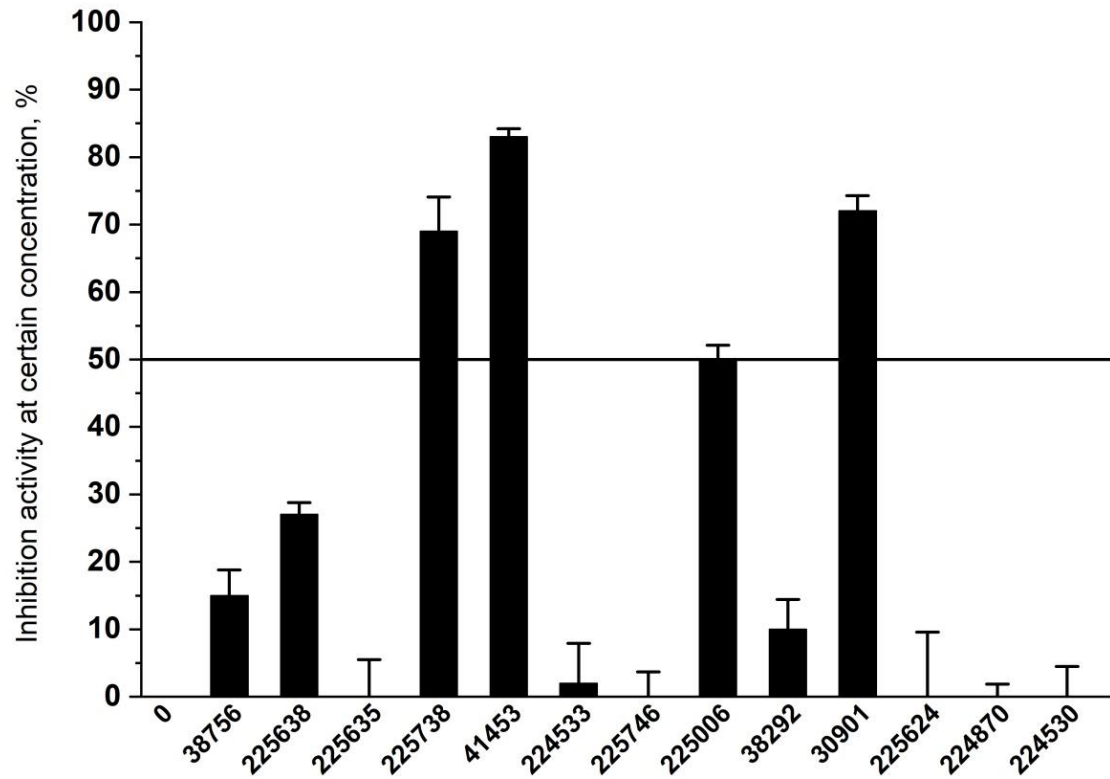
- ▶ Docking 326 conformers, 39 ligands were selected for ΔH calculations, 13 compounds were selected for experiments: 4 compounds confirmed inhibition Xlla in experiments
- ▶ Selection criteria: native ligand docking: SOL score = -5.19 kcal/mol

Docking, SOL program: MMFF94 force field, Genetic algorithm

Population size 30 000; number of generations 1 000. For each ligand, 50 independent runs of the genetic global optimization algorithm were performed. The 50 solutions, best ligand poses, were grouped into clusters using a criterion RMSD < 1 Å; a reliable solution must have population of the 1st cluster > 10, 1st cluster contains ligand poses with most negative SOL score.

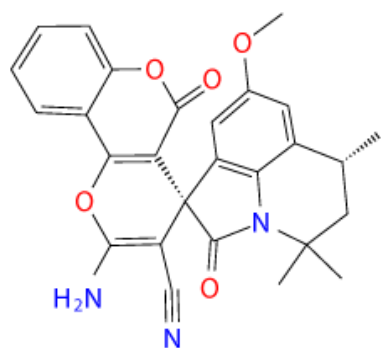
PM7+COSMO binding enthalpy ΔH must be maximum negative

Experimental results for 13 selected compounds

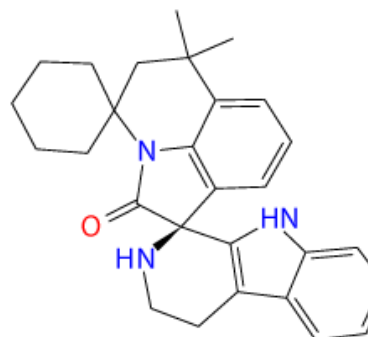


$$\text{Inhibitory activity} = \left(1 - \frac{\text{activity of XIIa with an inhibitor at a concentration } 30 \mu\text{M}}{\text{activity of XIIa without inhibitor}} \right) \times 100\%$$

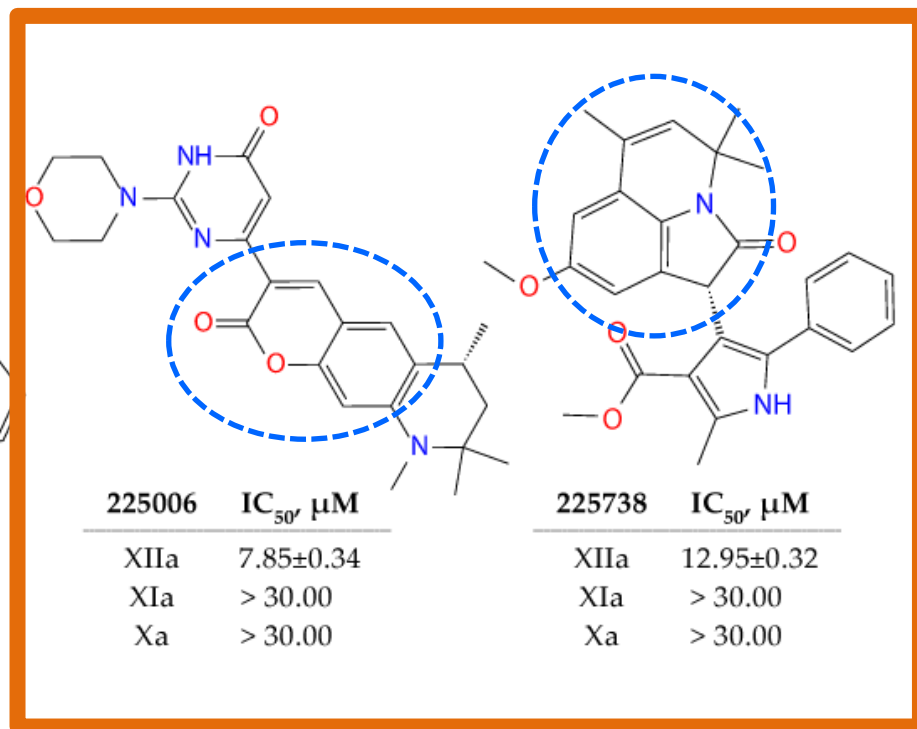
4 confirmed hits



| 30901 | IC _{50'} μM |
|-------|----------------------|
| XIIa | 19.67±1.31 |
| XIa | > 30.00 |
| Xa | 12.40±0.38 |



| 41453 | IC _{50'} μM |
|-------|----------------------|
| XIIa | 25.88±1.77 |
| XIa | > 30.00 |
| Xa | > 30.00 |



| 225006 | IC _{50'} μM |
|--------|----------------------|
| XIIa | 7.85±0.34 |
| XIa | > 30.00 |
| Xa | > 30.00 |

| 225738 | IC _{50'} μM |
|--------|----------------------|
| XIIa | 12.95±0.32 |
| XIa | > 30.00 |
| Xa | > 30.00 |

Anna Tashchilova, et al. *Molecules* (MDPI), 2022, **27**, 1234.
<https://doi.org/10.3390/molecules27041234>

Chemical classes:
225006 – coumarin derivative
225738 – pyrroloquinoline

Database of chemical compounds

Large VSU database - more than 19 000 compounds (30 000 conformers)

Selection of compounds for experimental study

Docking > **30 000** conformers, SOL score < -6.00 kcal/mol; **423** ligands were selected for ΔH calculations, **41** compounds were selected as best for experiments $\Delta H < -53$ kcal/mol; additional filtering: clustering due to ligand similarity using DataWarrior software (similarity 0.85) \Rightarrow **25** clusters \Rightarrow **28** ligands, analysis of interactions, H-bonds, **16** compounds were selected for experiments: **7** compounds are weak inhibitors in experiments, **1** compounds confirmed noticeable inhibition Xlla in experiments:

41% Xlla inhibition at 30 μ M

2 H-bonds

3 Halogen bonds

Chemical class: spiro-derivative of succinimide fused with tetrahydrofuran

Analogs of published Xlla inhibitors

100 compounds – analogs of recently published Xlla inhibitors: Marvin Korff, et al. Journal of Medicinal Chemistry 2020, 63 (21), 13159-13186

Selection of compounds for experimental study

- ▶ SOL score < -5.1 kcal/mol, 21 selected for experiments
- ▶ 3 compounds confirmed inhibitors Xlla in experiments.
- ▶ They belong to two chemical classes:

triazolo-pyrimidine

triazolo-pyridine

Conclusions

- ▶ **The initial search for inhibitors of the coagulation factor XIIa in the available database was carried out using the SOL docking program**
- ▶ **Selected compounds were tested in vitro**
- ▶ **Several experimentally confirmed hits were discovered**
- ▶ **Factor XIIa inhibitors found belong to 5 chemical classes**
 - Coumarin derivative
 - Pyrroloquinoline
 - Spiro-derivative of succinimide fused with tetrahydrofuran
 - Triazolo-pyrimidine
 - Triazolo-pyridine
- ▶ **Success of virtual screening using docking is determined by the content of the screened database of compounds**

Thank you for attention!



... Surely every medicine is an innovation;
and he that will not apply new remedies,
must expect new evils ...

Francis Bacon
(1561-1626)
OF INNOVATIONS

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