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## Search For Potent And Selective Aurora A Inhibitors Based On General Ser/Thr Kinases Pharmacophore Model

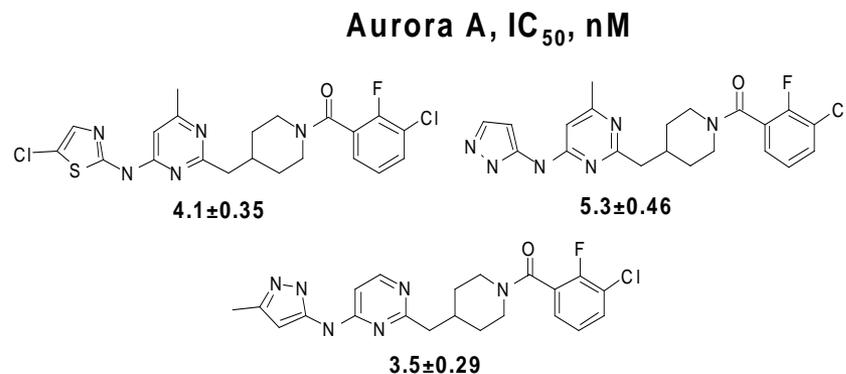
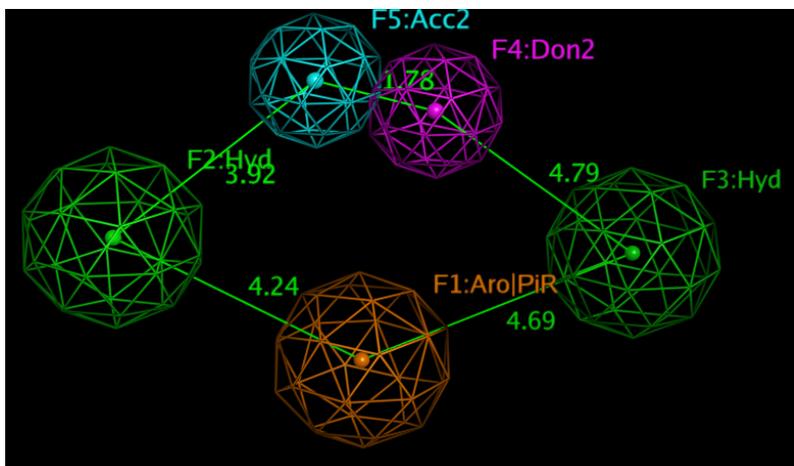
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# Search For Potent And Selective Aurora A Inhibitors Based On General Ser/Thr Kinases Pharmacophore Model



**Abstract:** Based on the data of compounds known from the literature to be active against various types of Ser/Thr kinases a general pharmacophore model for these types of kinases was developed. Search for the molecules fitting to this pharmacophore among ASINEX proprietary library revealed a number of compounds, which were tested and appeared to possess some activity against such Ser/Thr kinases as Aurora A, Aurora B and Haspin.

Our work on optimization of these molecules to Aurora A kinase allowed us to achieve several hits in 3-5 nM range of activity, with rather good selectivity and ADME properties.

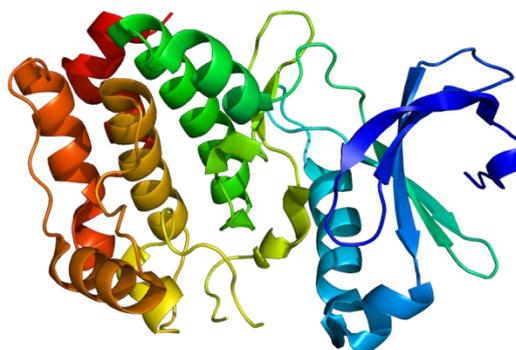
Thus we showed the possibility of performing the fine-tuning of the general Ser/Thr pharmacophore designed to desired types of kinase to get active and selective compounds.

**Keywords:** general pharmacophore model, Ser/Thr kinases, Aurora A



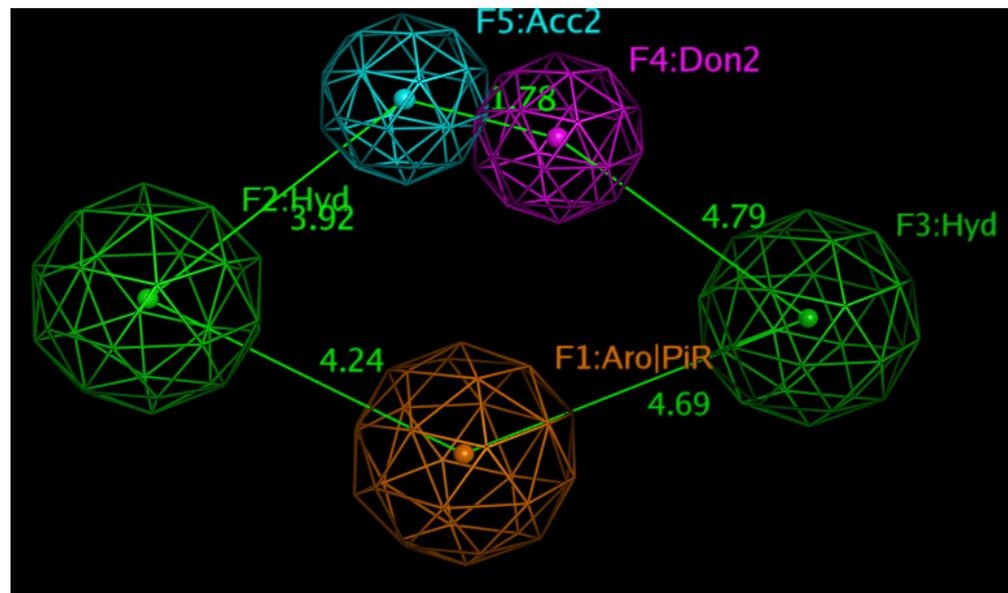
# Introduction

- **Serine/threonine protein kinase** are kinase enzymes that phosphorylate the OH group of serine or threonine. At least 125 of the 500+ human protein kinases are serine/threonine kinases (STK).
- Inhibitors of Ser/Thr kinases can possess potential therapeutic uses—from treating cancer to immune disorders. Since they were found in a number of mycobacterial organisms they can be also used for treatment of bacterial infections such as tuberculosis.
- A number of attempts have been done to construct a pharmacophore model for various Ser/Thr kinase inhibitors, such as STPK inhibitors of tuberculosis, mTor kinase inhibitors, Aurora A and B inhibitors, B-Raf inhibitors etc.
- We hypothesized that it is possible to find out some general features of all serine-threonine inhibitors and to construct a general pharmacophore model. Then this general model can be adjusted to specific kinds of serine-threonine kinase.



## Results and discussion. General pharmacophore model construction

Known inhibitors of various serine-threonine kinases including both found in ASINEX proprietary collection and in literature were used to construct the general pharmacophore model. Pharmacophore elucidating functionality of MOE version 2010.10 was used.



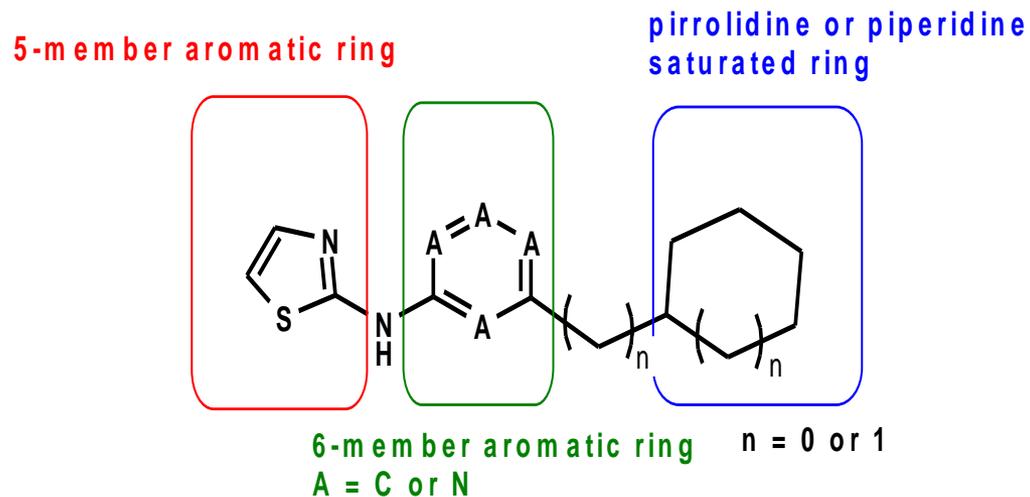
*General pharmacophore of serine-threonine kinase inhibitors (MOE 2010.10)*

The general pharmacophore found looks like a kind of rhomb with two opposite hydrophobic centers, one aromatic center and a couple of H-bond donor and acceptor projections in one corner. The length of a rhomb side is about 4 – 5 Å



# Results and discussion. Application of the general pharmacophore to ASINEX proprietary library

Application of the general pharmacophore to ASINEX proprietary library allowed to find a scaffold fitting to its requirements.



# Results and discussion. Activity of compounds fitting to general pharmacophore model against selected Ser/Thr kinases

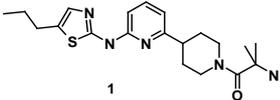
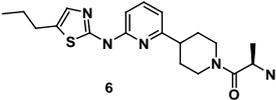
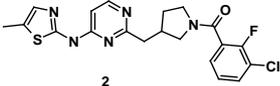
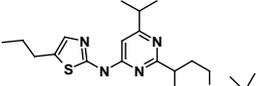
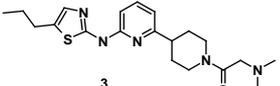
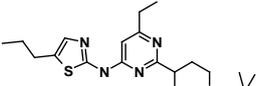
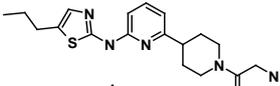
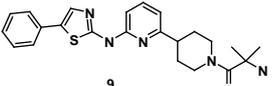
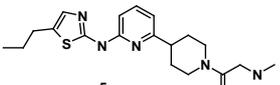
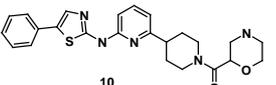
To confirm the activity against Ser/Thr kinases a set of compounds from ASINEX library belonging to the found scaffold was tested against Aurora A, Aurora B and Haspin kinases

*The best  
compound  
against Aurora A  
kinase is  
2 (100 nM)*



# Results and discussion. Activity of compounds fitting to general pharmacophore model against selected Ser/Thr kinases

To confirm the activity against Ser/Thr kinases a set of compounds from ASINEX library belonging to the found scaffold was tested against Aurora A, Aurora B and Haspin kinases

Structure	Aurora A	Aurora B	Haspin	Structure	Aurora A	Aurora B	Haspin
IC <sub>50</sub> , uM			IC <sub>50</sub> , uM				
 1	2.06	1.45	1.49	 6	1.67	1.75	1.12
 2	0.10	3.46	105.2	 7	2.79	2.25	3.67
 3	1.95	1.76	7.06	 8	3.58	2.05	2.90
 4	2.08	1.72	0.08	 9	3.17	8.02	0.27
 5	1.44	0.91	0.43	 10	8.38	0.84	7.80

**The best compound against Aurora A kinase is 2 (100 nM)**



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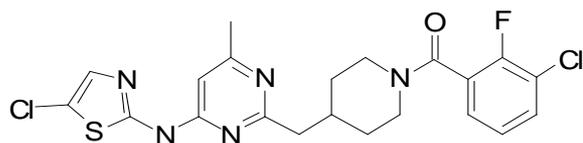
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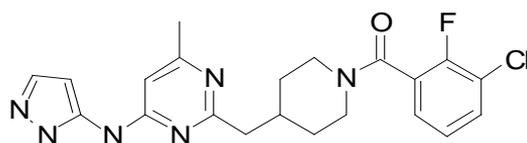
pharmaceuticals

# Results and discussion. Optimization of compound **2** for Aurora A activity

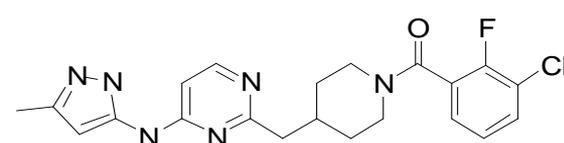
- To confirm the possibility of fine-tuning of the general pharmacophore for selected type of kinases we chose Aurora A kinase and tried to adjust the scaffold found to its specific requirements.
- With this goal a library of analogues of compound **2** was synthesized
- As a result of optimization compounds **29**, **30** and **32** were achieved



**29**, 4.1 nM



**30**, 5.3 nM



**32**, 3.5 nM



# Results and discussion. Selectivity and ADME properties of compounds 29, 30 and 32

	29	30	32
	IC <sub>50</sub> , nM	IC <sub>50</sub> , nM	IC <sub>50</sub> , nM
Aurora A	4.1	5.3	3.5
Aurora B	232.2	18300.0	4182.5
c-Kit	17000.00	21000	
EGFR			
CSK	9500.0		
EPHA2	9000.0		
GSK-3		109.0	292.8
Haspin			
JNK3	85.18	1634.0	572.5
LYN	3270.0		
PIM1			
PLK1			
RON	970.0		
CYP1A2			
CYP2C19		9597.0	965.7
CYP 2C9		3900.0	326.6
CYP3A4		15070.0	596.2
HERG			
Solubility (nephelometry)	< 10 ug/ml	>100 ug/ml	>100 ug/ml

- Compound 30 has**
- **Good selectivity,**
  - **Low cytochrome inhibition,**
  - **Low HERG channels inhibition and**
  - **Good solubility**



# Conclusions

- A general pharmacophore model applicable to various serine-threonine kinases was designed.
- A possibility of fine tuning of this pharmacophore to particular types of serine-threonine kinases was exemplified by design of potent and selective Aurora A inhibitors
- Several compounds with 3-5 nM potency, good selectivity and satisfactory ADME parameters were obtained



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