

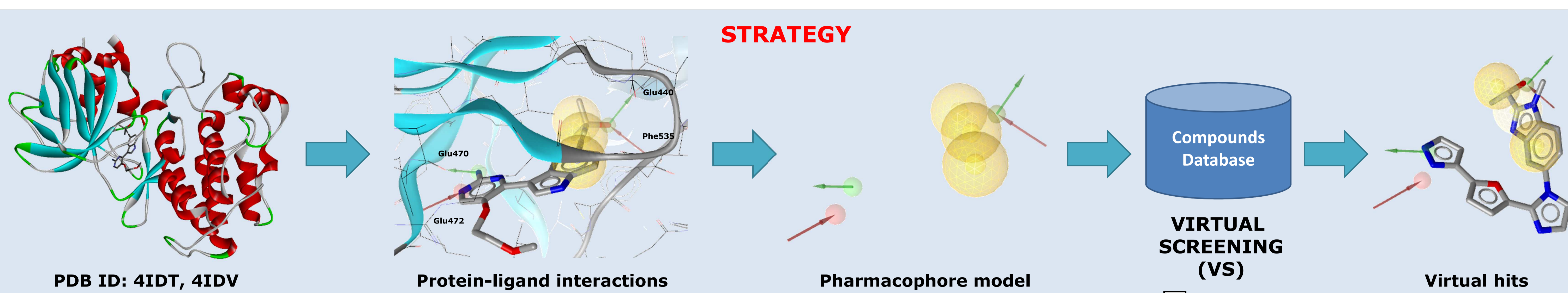
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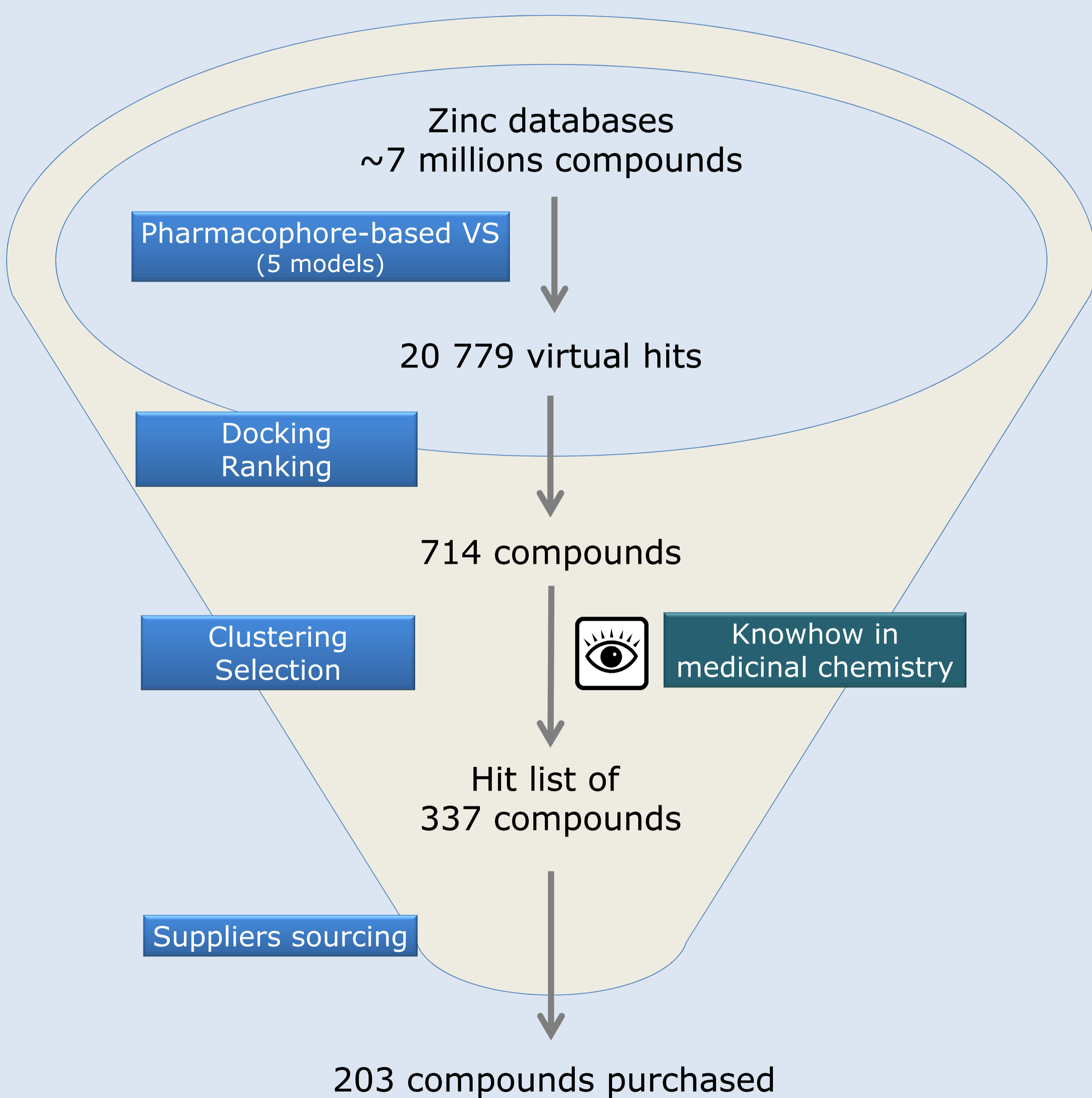


INTRODUCTION

The primary objective of the TAKTIC (TrAnslational Kinase Tumor Inhibitor discovery Consortium) project is to identify and develop highly specific small molecule inhibitors targeting the kinases IKK α , IKK β and NIK.¹ NF- κ B-inducing kinase (NIK) is a serine/threonine protein kinase essential for the activation of a second major NF- κ B (NF κ B2) pathway which controls several critical biological functions such as inflammation or cell survival.²⁻³ The X-ray structure of NIK, a mitogen-activated protein kinase kinase kinase (MAP3K), has recently been published and was used in this work for a structure-based ligand design approach.⁴ Using this strategy, we identified a series of new small molecule inhibitors of NIK.

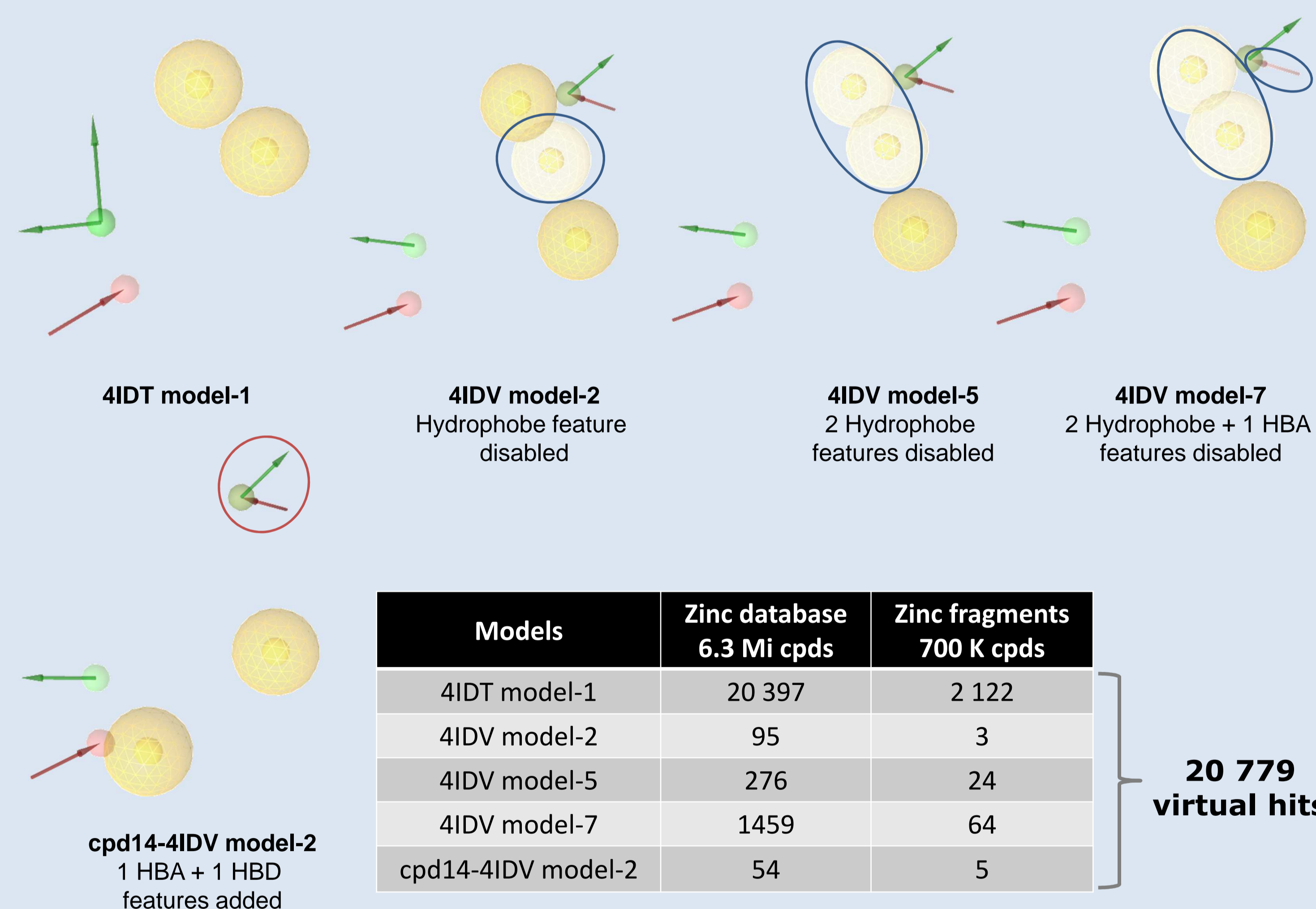


HIT DISCOVERY FLOWCHART



STRUCTURE-BASED PHARMACOPHORE MODELS AND VS

Two xray structures (PDB entries 4IDT and 4IDV) were analyzed and used to generate ten pharmacophore models by using the LigandScout software.⁵ Models were tuned and optimized in order to obtain an optimal hit rate for the virtual screening. After validation by using a set of active ligands,⁴ five models were selected for virtual screening of two 3D molecular structure databases (Zinc databases: drug-now and frag-now).⁶

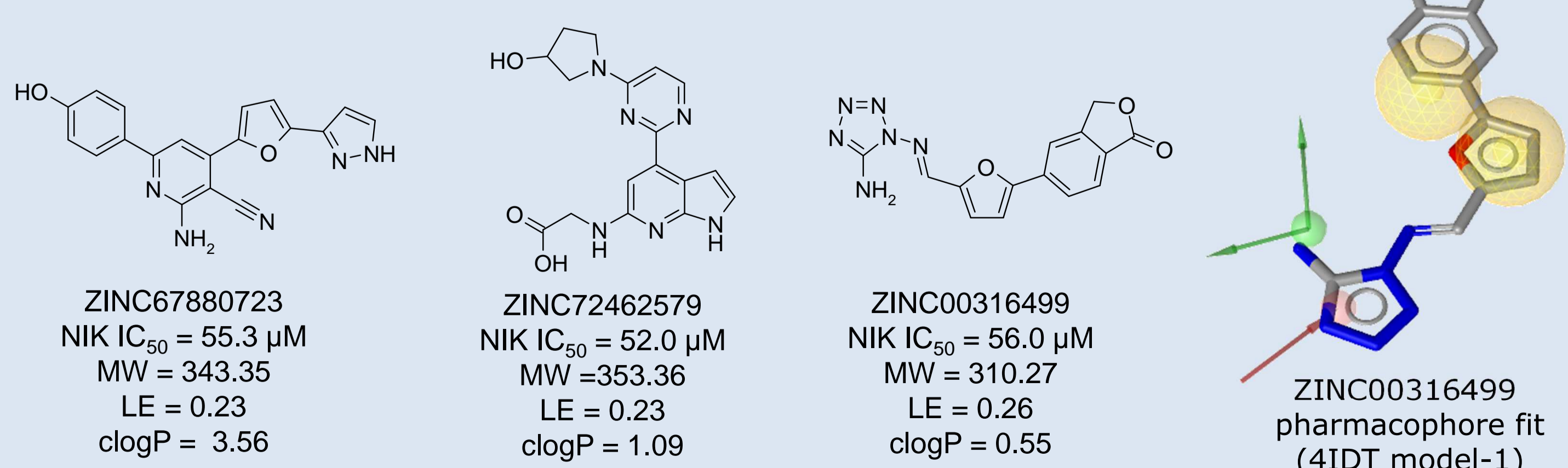


BIOCHEMICAL ASSAYS

Human full length NIK kinase was expressed in Sf9 insect cells and purified as recombinant GST fusion protein. NIK kinase activity was measured using the radiometric 33PanQinase[®] activity assay (ProQinase GmbH). In a primary screen, 203 compounds were tested for NIK inhibition (% residual activity) at 100 μ M. Verification of 27 primary hit compounds showing at least 40% NIK inhibition at 100 μ M was done by determination of IC₅₀ values (concentration range: 3nM - 100 μ M). 17 compounds showed a IC₅₀ below 100 μ M and several highly active inhibitors were identified (IC₅₀ < 10 μ M).

HITS IDENTIFIED

Among the 17 compounds having a IC₅₀ < 100 μ M, 8 main chemical series were identified. Two chemical series exhibited inhibition in the low micromolar range (IC₅₀ < 10 μ M - data not shown). Only three representative compounds are disclosed here.



REFERENCES

1. TAKTIC is a FP7-SME-2012 project (Grant number: 315746).
2. Hayden MS, Ghosh S; Cell 2008, 132, 344-362.
3. Vallabhapurapu S, Karin M; Annu. Rev. Immunol. 2009, 27:693-733.
4. Li K1, McGee LR, Fisher B, Sudom A, Liu J, Rubenstein SM, Anwer MK, Cushing TD, Shin Y, Ayres M, Lee F, Eksterowicz J, Faulder P, Waszkowicz B, Plotnikova O, Farrelly E, Xiao SH, Chen G, Wang Z.; Bioorg. Med. Chem. Lett. 2013, 23(5):1238-44.
5. <http://www.inteligand.com/>.
6. <http://zinc.docking.org/>

CONCLUSION

Pharmacophore-based virtual screening experiments were performed with a large chemical database allowing, after selection and experimental evaluation of 203 compounds, to identify micromolar hits against NIK kinase. Two chemical series (IC₅₀ < 10 μ M) are currently under investigation in a hit-to-lead validation process, kinase selectivity (profiling) and early ADMET evaluation. The objective is to enter into a lead optimization phase in order to develop new and selective NIK small molecule inhibitors.