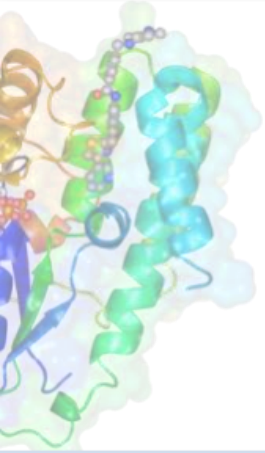


From a Drug Repositioning to a Fragment-Based Drug Design Approaches to Tackle Acute Lymphoblastic Leukemia (T-ALL)

Thomas Miller, PhD

Cancer Research Center of Marseille (CRCM) & Institut Paoli-Calmettes



From a Drug Repositioning to a Fragment-Based Drug Design Approaches to Tackle Acute Lymphoblastic Leukemia (T-ALL)

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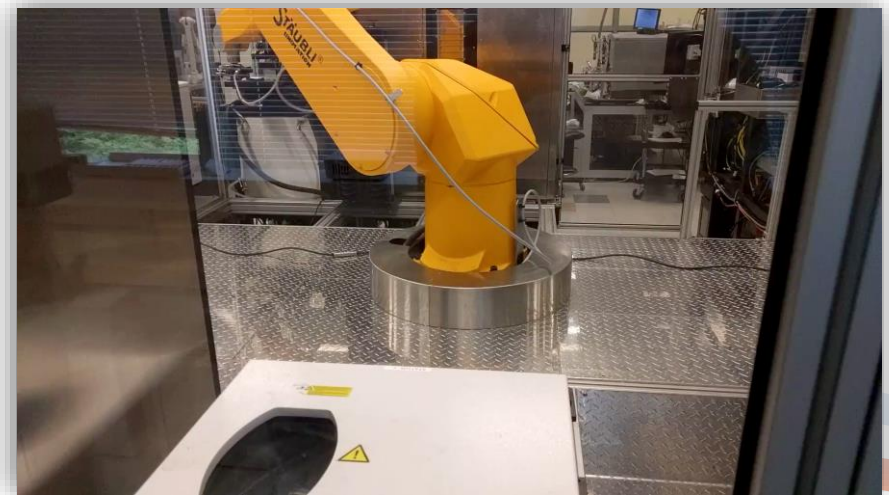
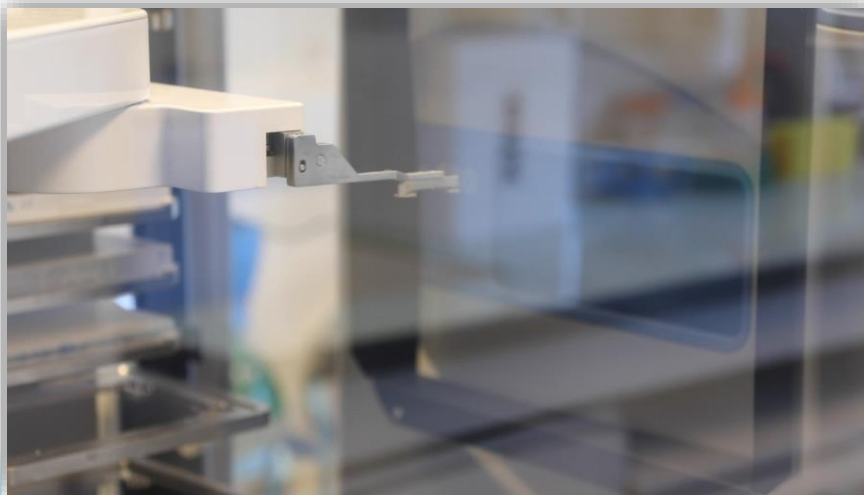
Cancer Research Center of Marseille (CRCM) & Institut Paoli-Calmettes

Introductions

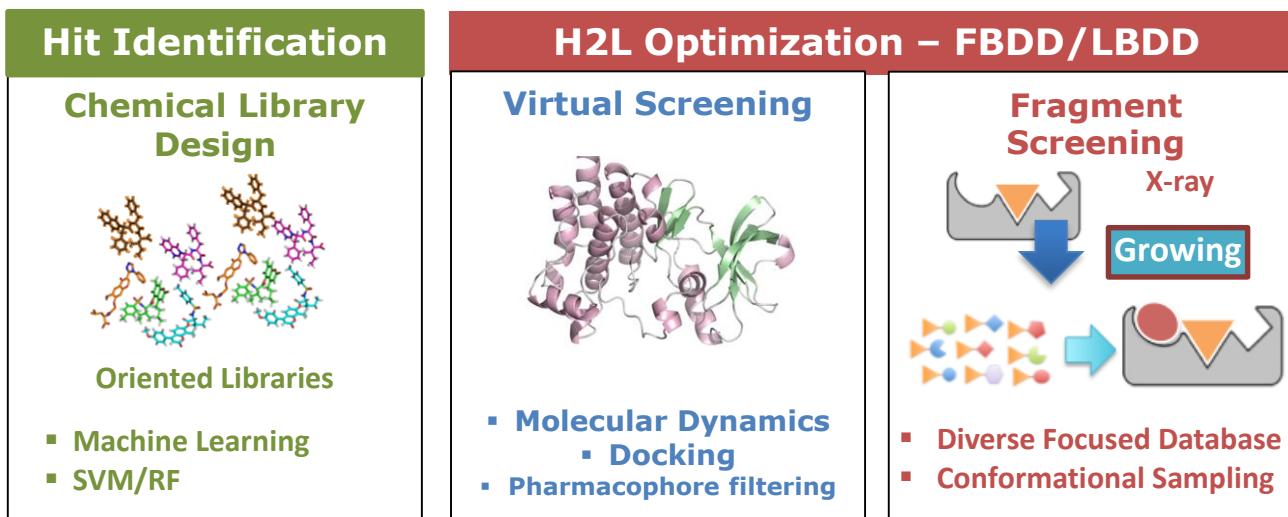
Current : iSCB team/HiTS
platform Marseille



Former: ADST team, NCATS, NIH
Bethesda MD USA



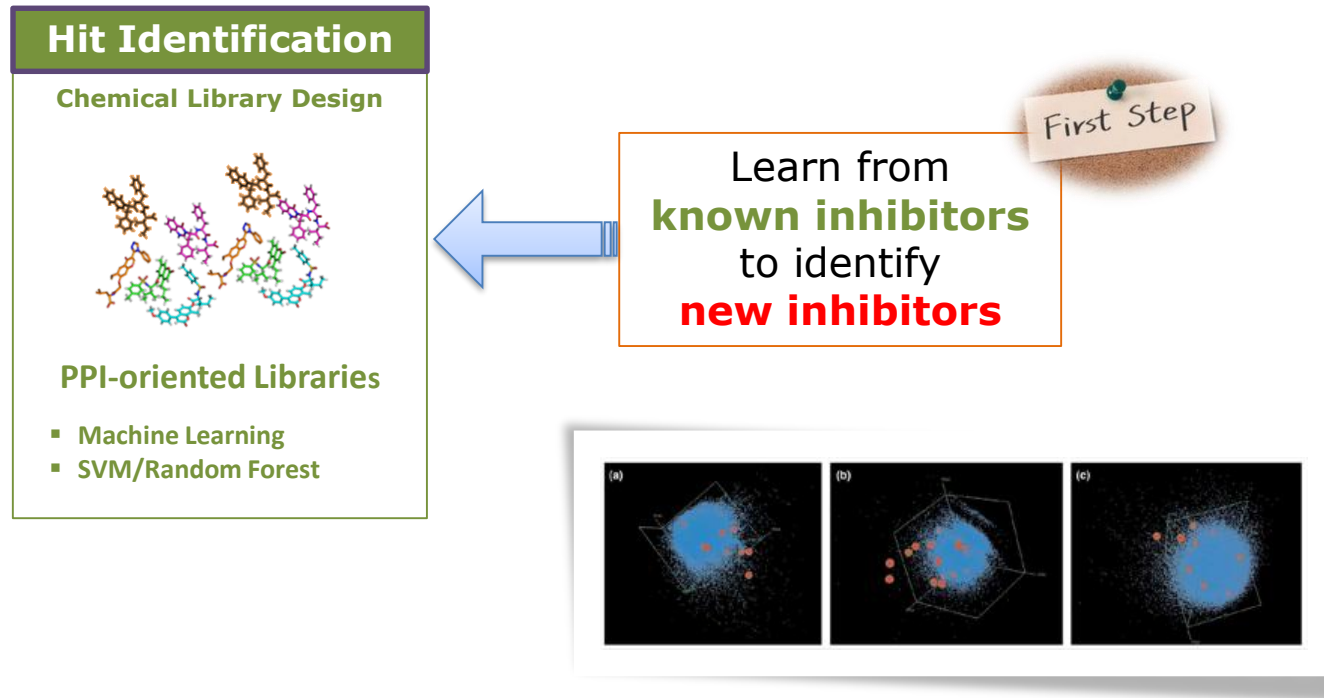
Strategies in Drug Discovery



1 "Fr-PPIChem" *: A PPI-Oriented Chemical library

2 "DOTS" *: An integrated strategy To accelerate the Hit-to-Lead Process

Strategies in Drug Discovery



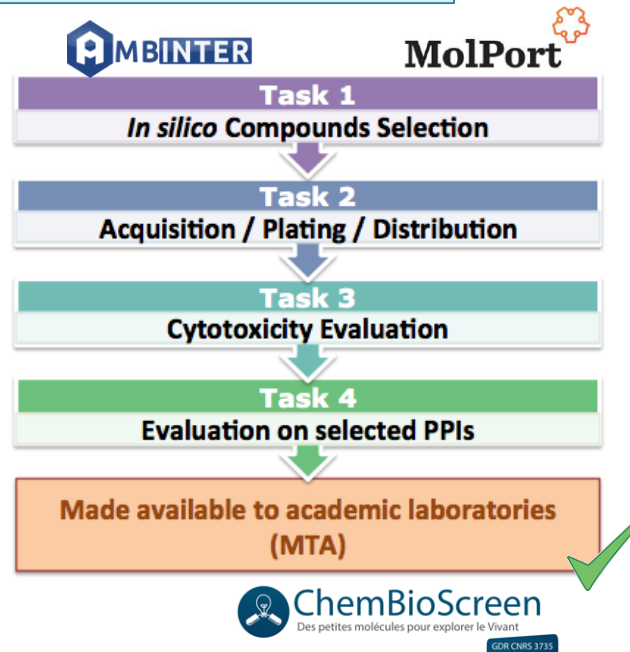
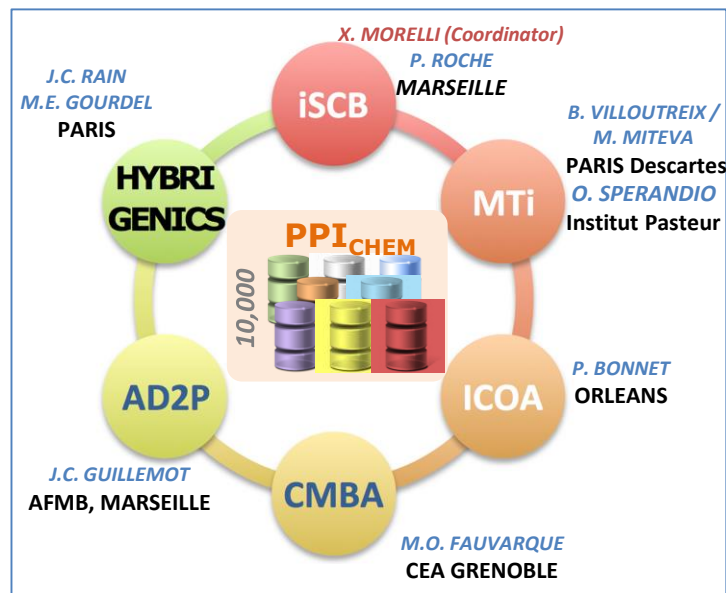
"Fr-PPIChem" : a French National Initiative



New Larger Version of 2P2I_{3D}

v2.0 (MedChem) using updated algorithms, ADMET Filters

➤ National Program (ANR) => 10.314 Compounds PPI-Library



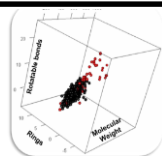
"Fr-PPIChem" : a French National Initiative

PDB PROTEIN DATA BANK

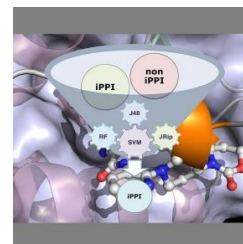
2P2I_{DB}



Machine Learning



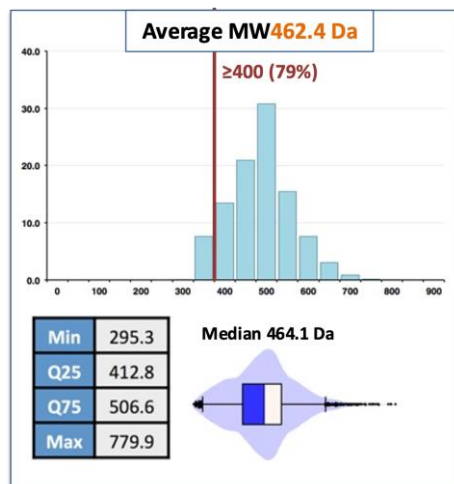
PPI-focused Chemical Libraries



Fr-PPI_{Chem}



10,314 cpds



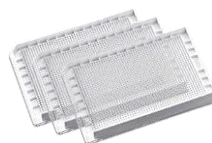
Requests

ChemBioFrance

Xavier Morelli (xavier.morelli@inserm.fr)
Philippe Roche (philippe.roche@inserm.fr)
Caroline Barette (caroline.barette@cea.fr)

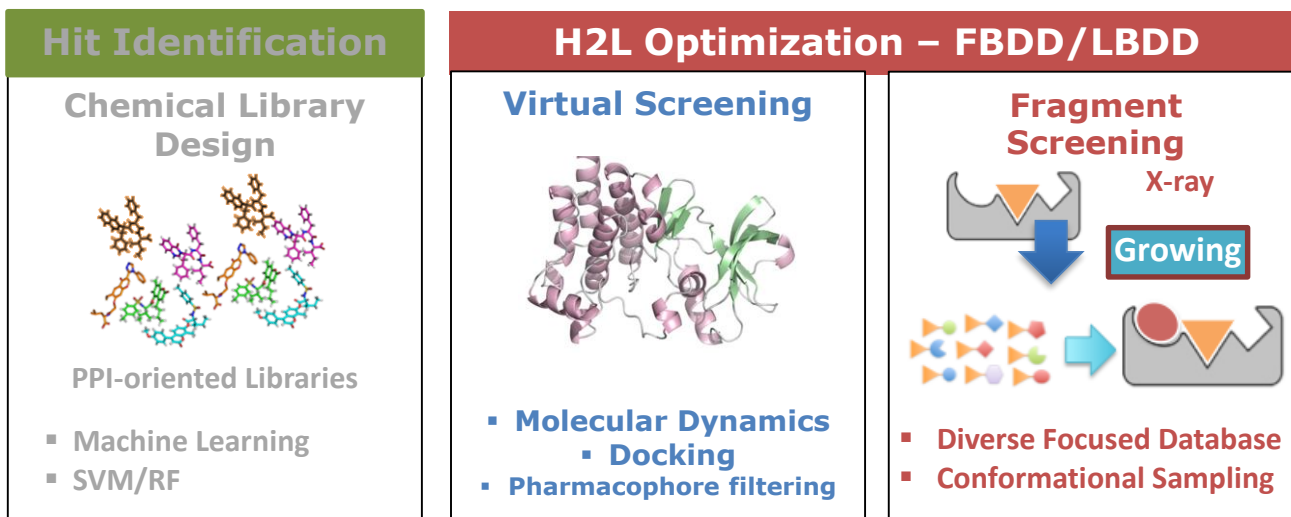


<https://chembiofrance.cn.cnr.fr/fr/composante/chimiotheque#prestation-0-5>



33 Plates (3 μ l @10mM per well)
>40 sets of plates distributed (4 USA)
>20 Targets screened @ iSCB

Strategies in Drug Discovery



1 “Fr-PPIChem”:
PPI-Oriented
Chemical library

2 “DOTS”:
An integrated strategy
To accelerate the
Hit-to-Lead Process

Finding a new drug is an almost unattainable challenge...

“The best way to discover a new drug is to start with an old one...”



**Sir James W. Black
(Prix Nobel 1988)**



“Off-target concept”:

The drug acts on new targets, out of the original scope, with a new therapeutic indication

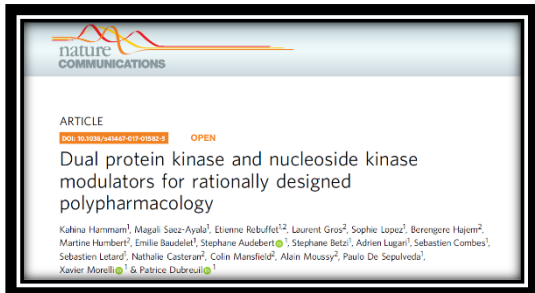
Both the targets & the indications are new.

Drug	Original Indication	New Indication
Aspirin	Inflammation, pain	Antiplatelet, Cancer
Bupropion	Depression	Smoking cessation
Disulfiram	Alcoholism	Melanoma
Doxepin	Depressive disorder	Antipruritic
Gemcitabine	Antiviral	Cancer
Methotrexate	Cancer	Psoriasis, rheumatoid arthritis
Minoxidil	Hypertension	Hair loss
Sildenafil	Angina	Erectile dysfunction (Viagra®)
Thalidomide	Morning sickness	Leprosy
Tretinoin	Acne	Leukemia

Drug Repurposing in Cancer: Turning Liabilities into Opportunities...



Masitinib (Masivet®)
 Patrice Dubreuil & AB Science



Nucleoside Kinase Activator



Oral Targeted Therapy (TKI)

Mastocytosis

Therapeutic benefit: Sensitize patients to chemotherapy

Reducing toxicity of therapeutic agents

Neurological diseases

Gastrointestinal stromal tumors (GIST)

Maintaining effectiveness at lower doses

Counteracting drug resistance initiated *via* down modulation

Hamman et al, Nat Commun, 2017

dCK vs. Cancer ...



dCK is a key enzyme in the Nucleoside Salvage Pathway (SP)...



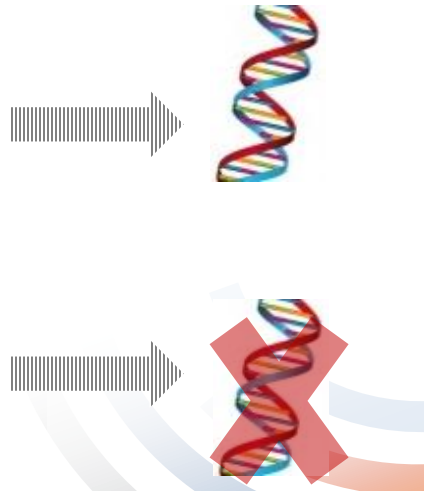
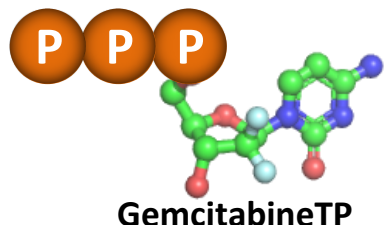
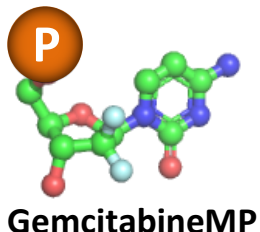
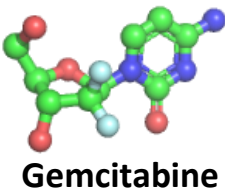
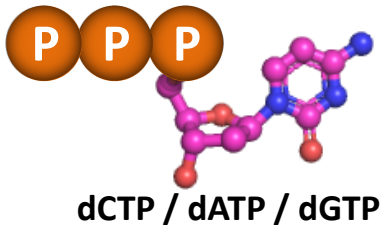
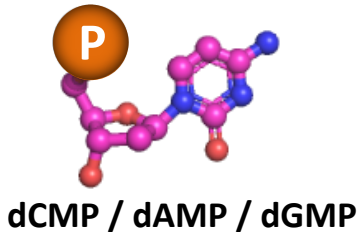
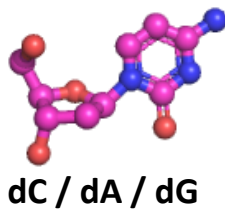
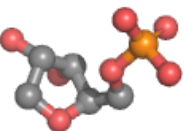
dCK vs. Cancer ...



dCK is a key enzyme in the Nucleoside Salvage Pathway (SP)...

... which recycles bases and nucleosides originating from the degradation of RNA and DNA to achieve the biosynthesis of deoxyribonucleotides, required for DNA replication and repair

... which catalyzes the phosphorylation of physiologic & nucleoside analogue drugs (but dT)

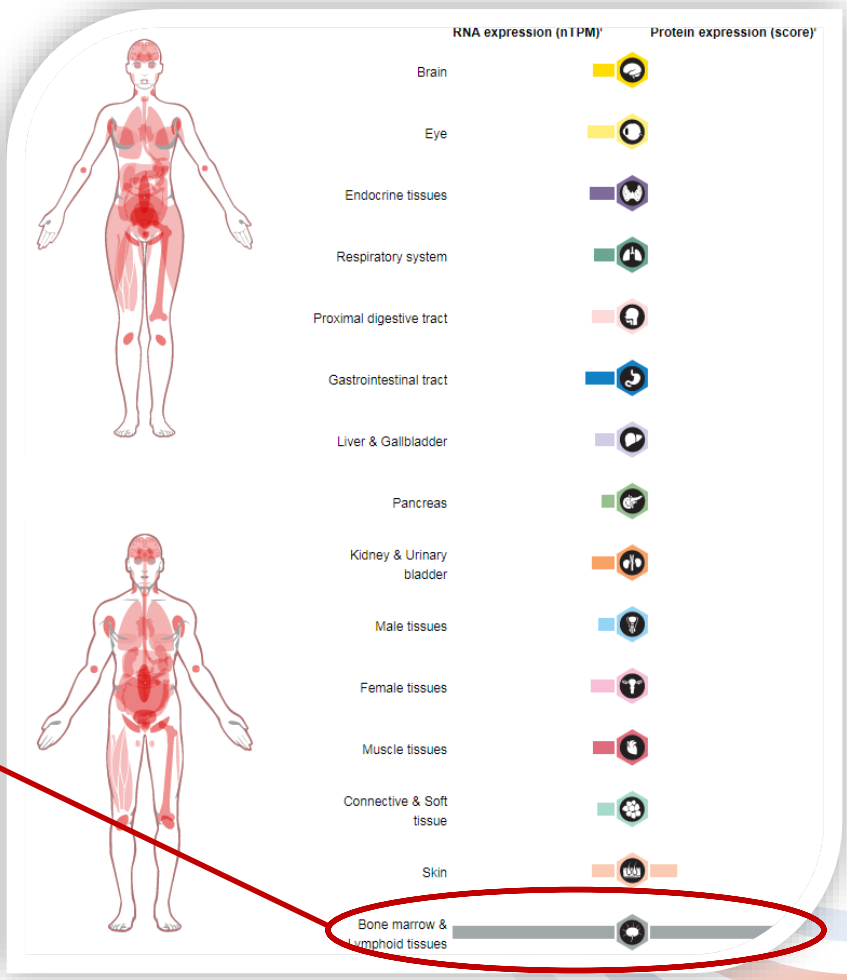
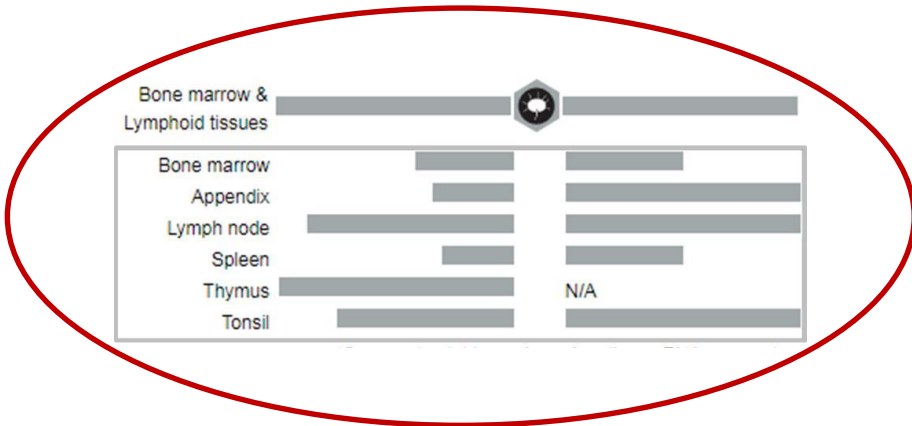


+ numerous anticancer and antiviral nucleoside analogues

dCK vs. Cancer ...



Highly expressed in bone marrow and lymphoid tissues

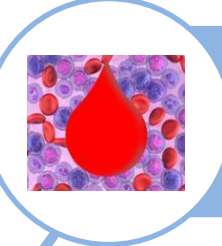


Bone marrow & lymphoid tissues

dCK vs. Cancer ...

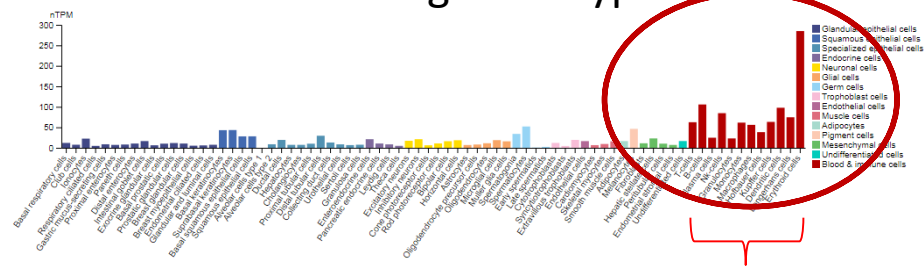


Highly expressed in bone marrow and lymphoid tissues



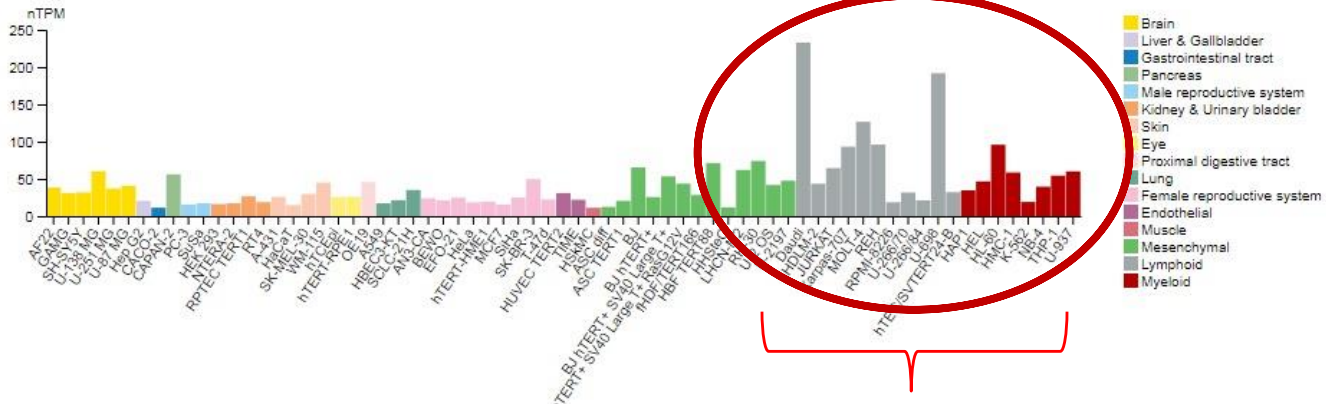
Highly expressed in lymphoid and myeloid cancer cell lines

RNA single cell type



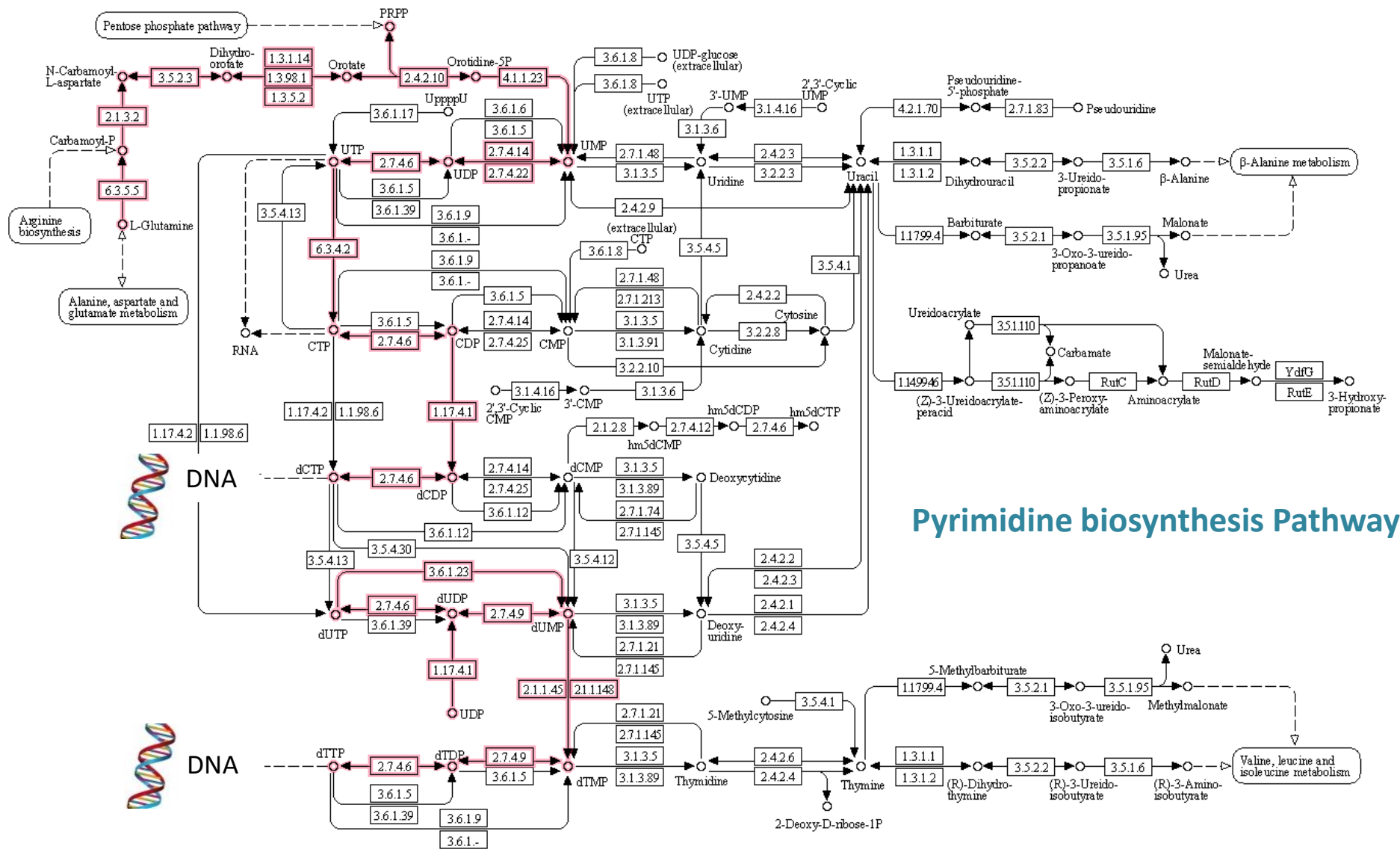
Blood & Immune cells

Cancer Cell lines

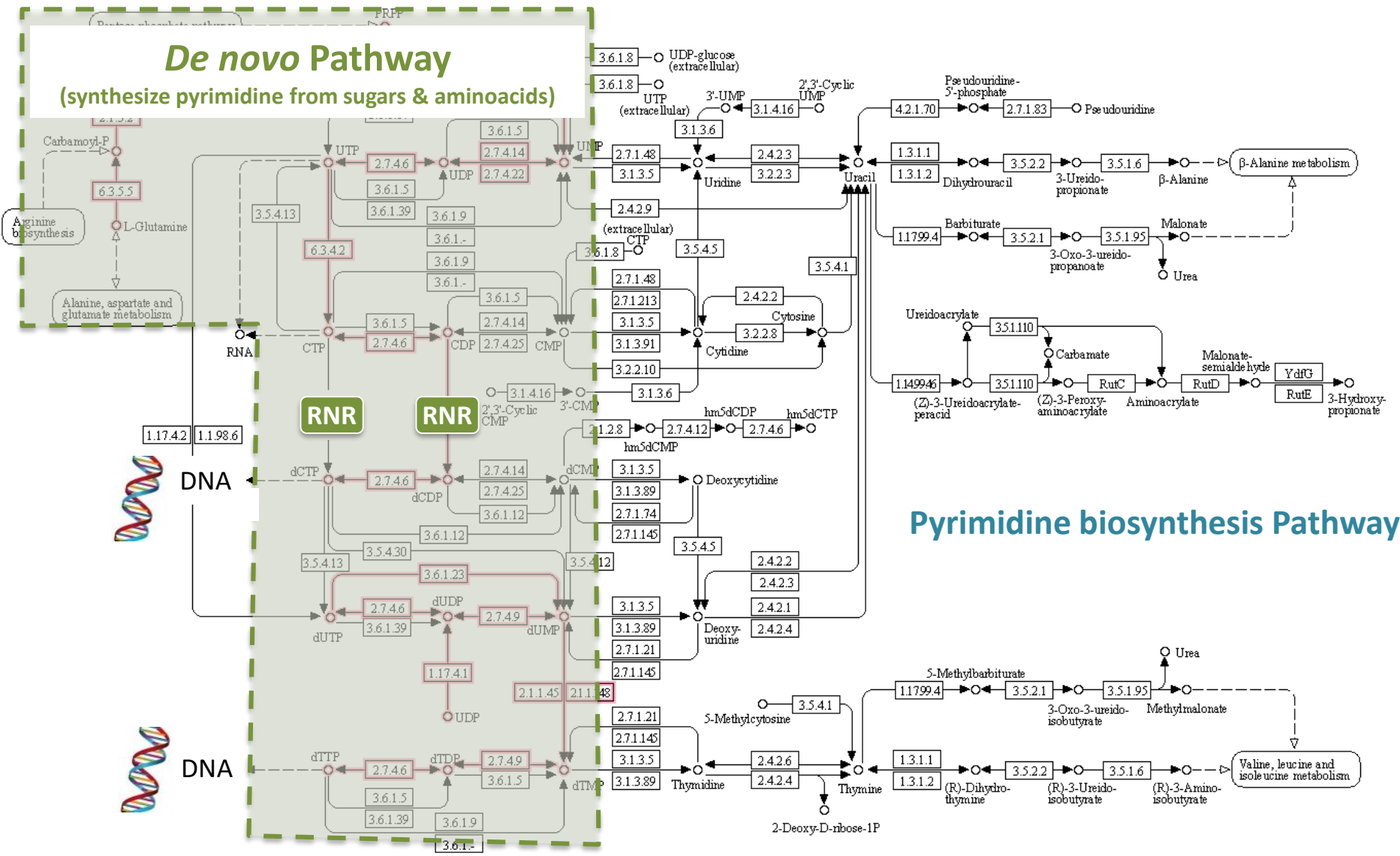


Lymphoid & Myeloid cell lines

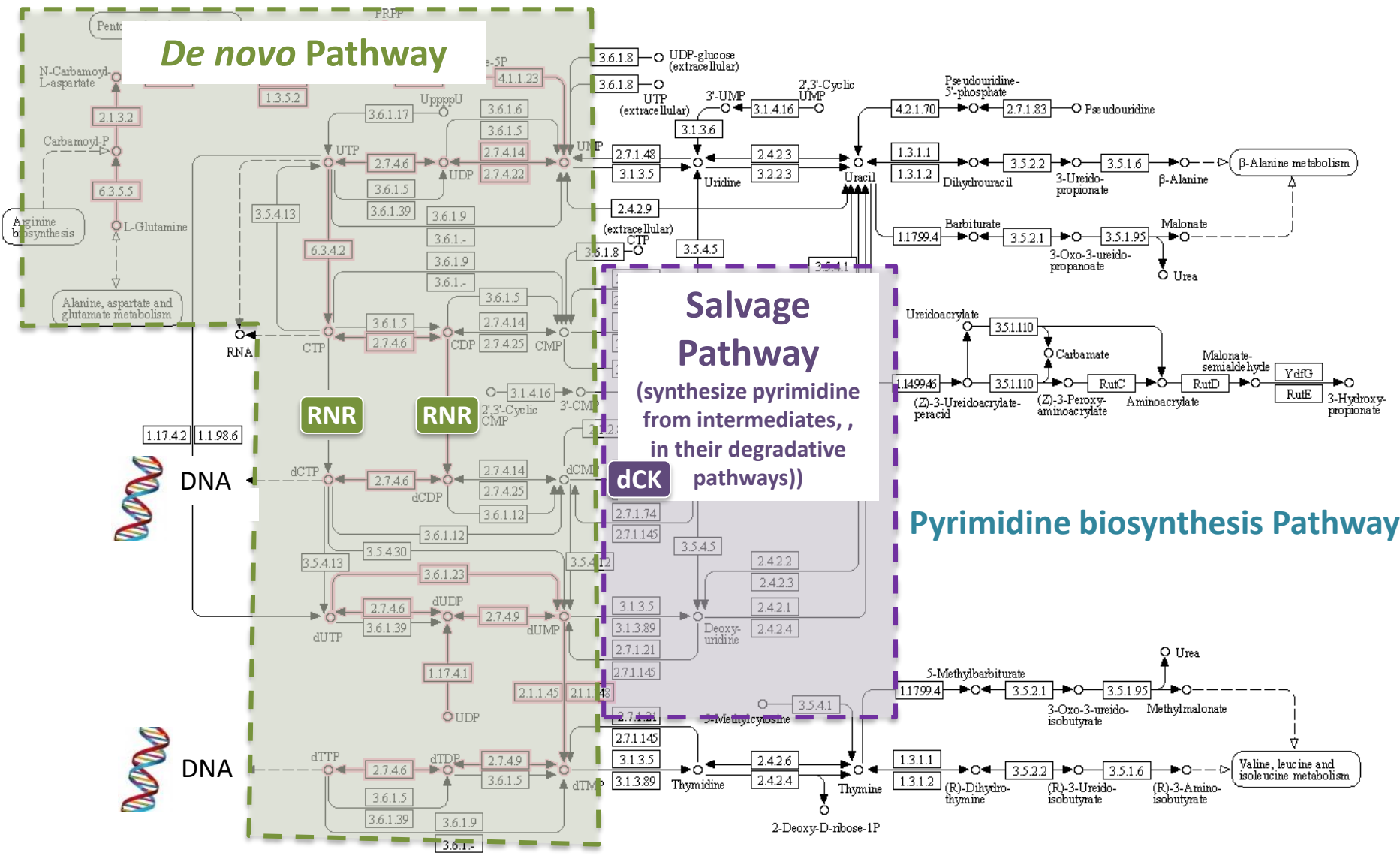
dCK vs. Deoxyribonucleotides biosynthesis Pathways...



dCK vs. De novo & Salvage Pathways...



dCK vs. De novo & Salvage Pathways...



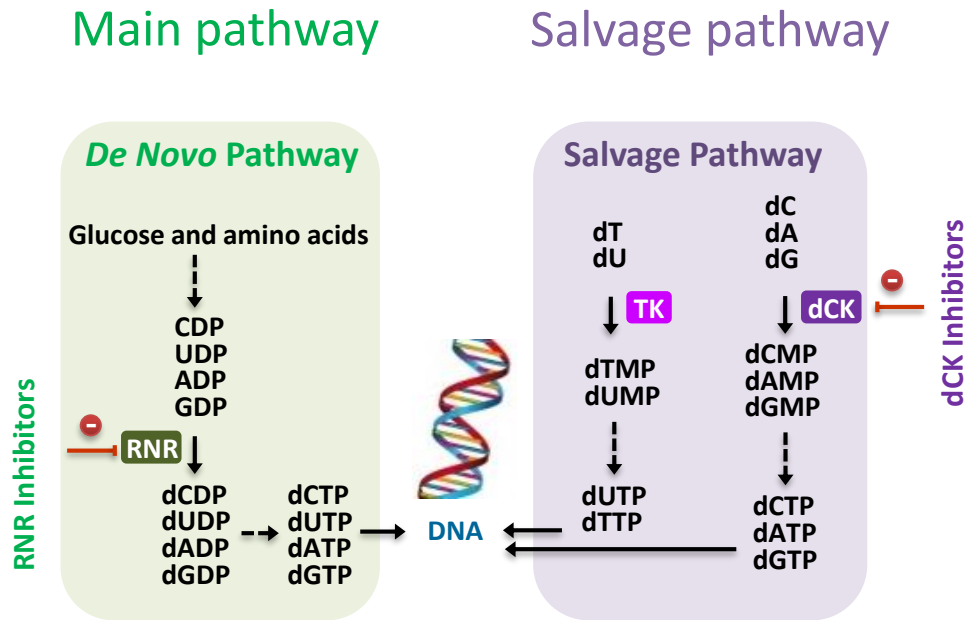
De novo Pathway

Salvage Pathway
(synthesize pyrimidine from intermediates, in their degradative pathways))

Pyrimidine biosynthesis Pathway

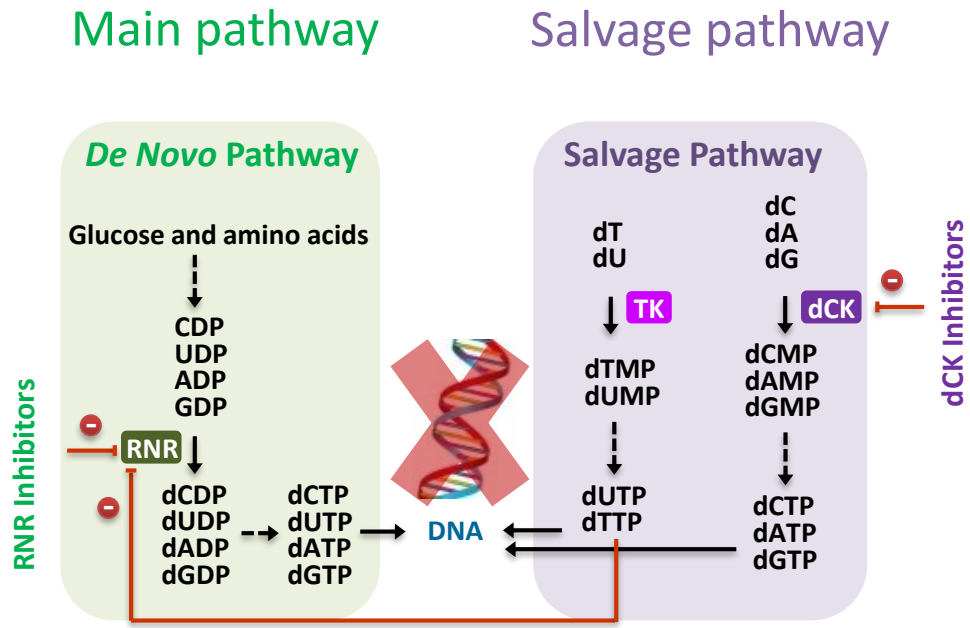
dCK vs. *De novo* & Salvage Pathways...

Inhibition of one pathway alone do not stop DNA synthesis



dCK vs. *De novo* & Salvage Pathways...

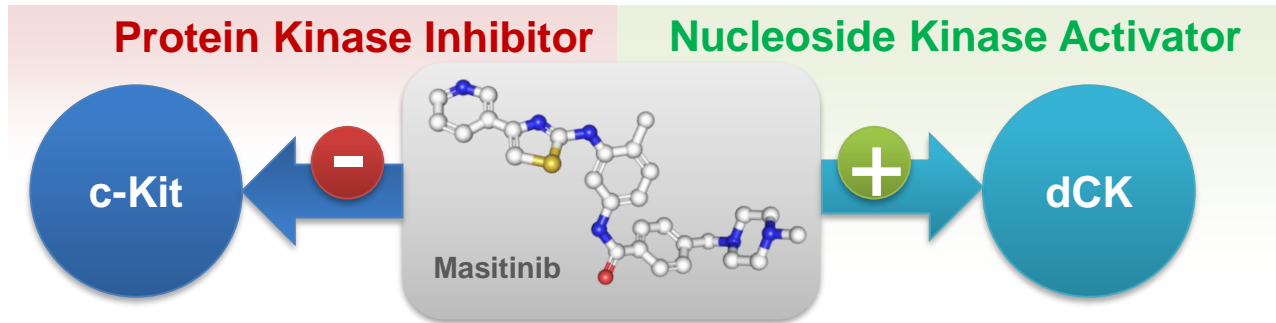
Inhibition of both pathways stops DNA synthesis



Concept of Synthetic lethality (DNP + SP)

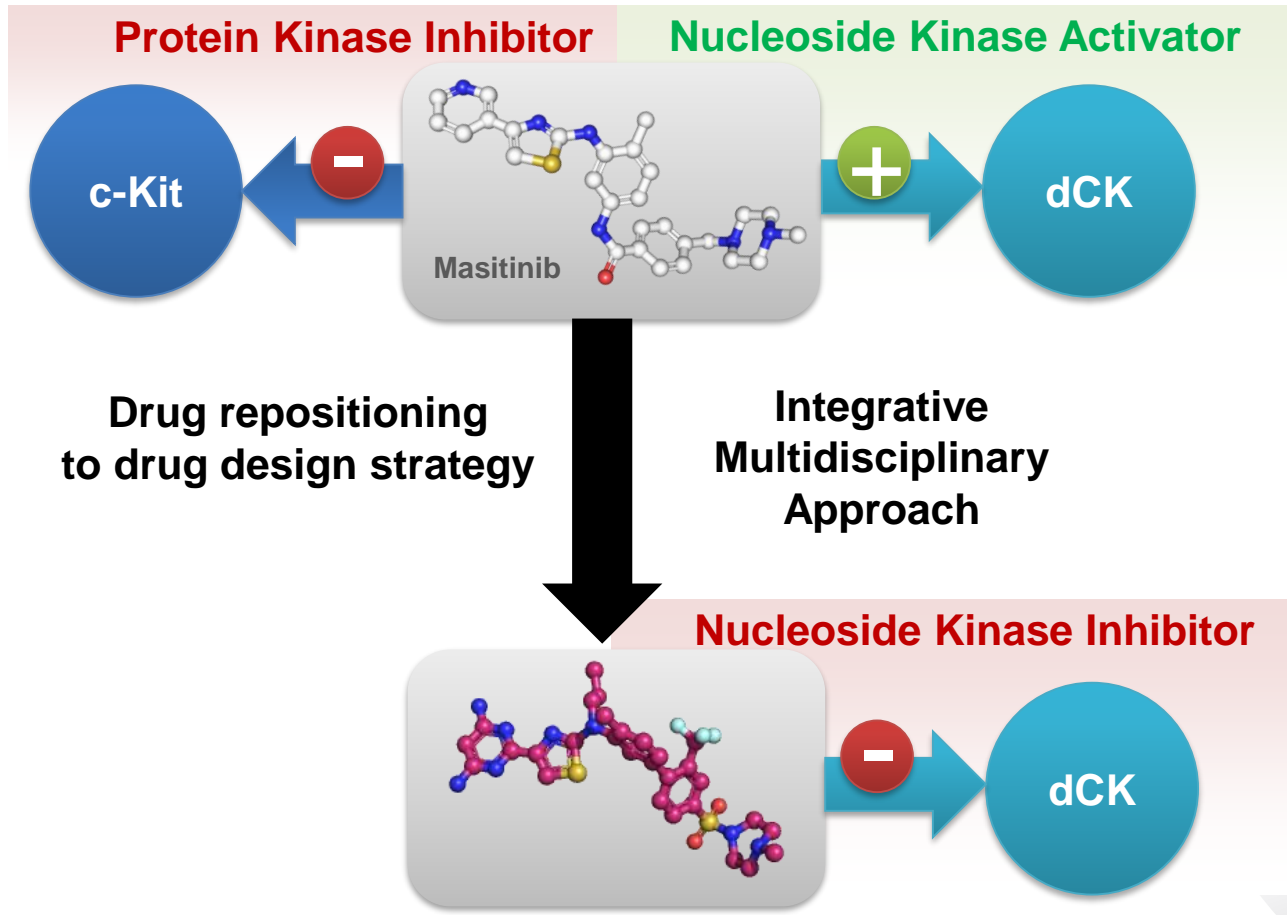
dCK vs. *De novo* & Salvage Pathways...

Hypothesis: Repositioning of masitinib to develop an inhibitor of dCK ?
Increase affinity to become an inhibitor ?



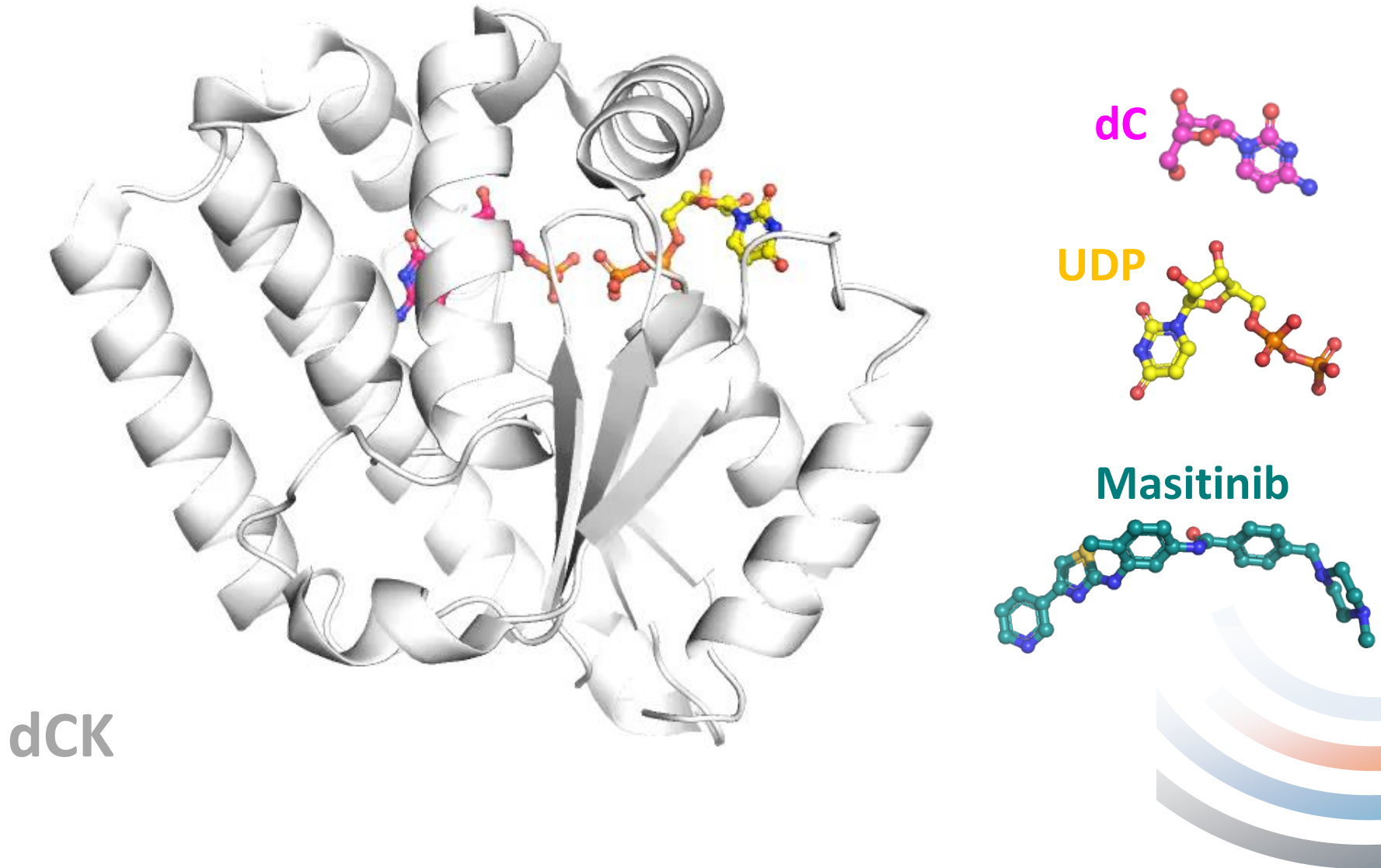
dCK vs. *De novo* & Salvage Pathways...

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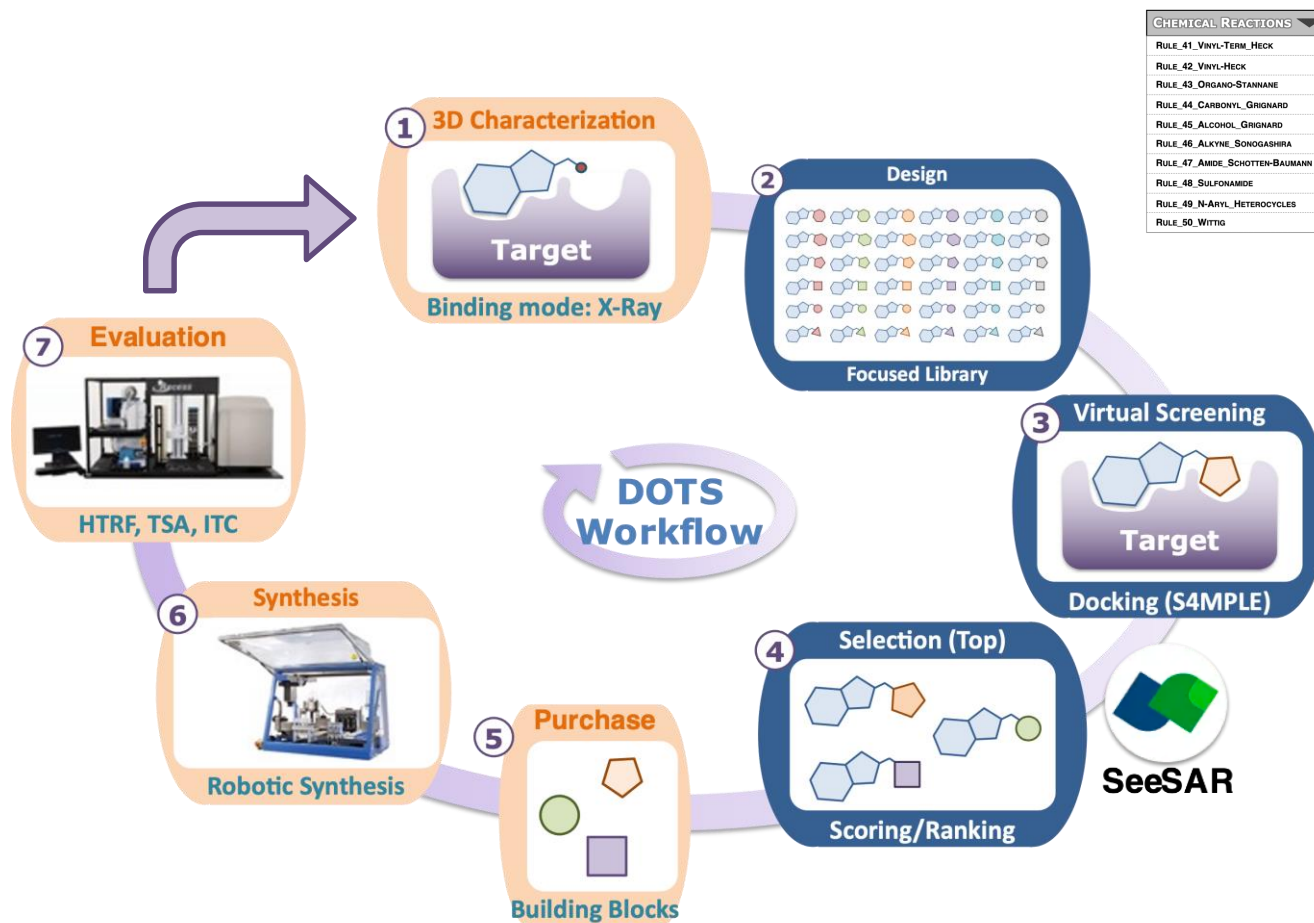


Turning an Activator into Inhibitor

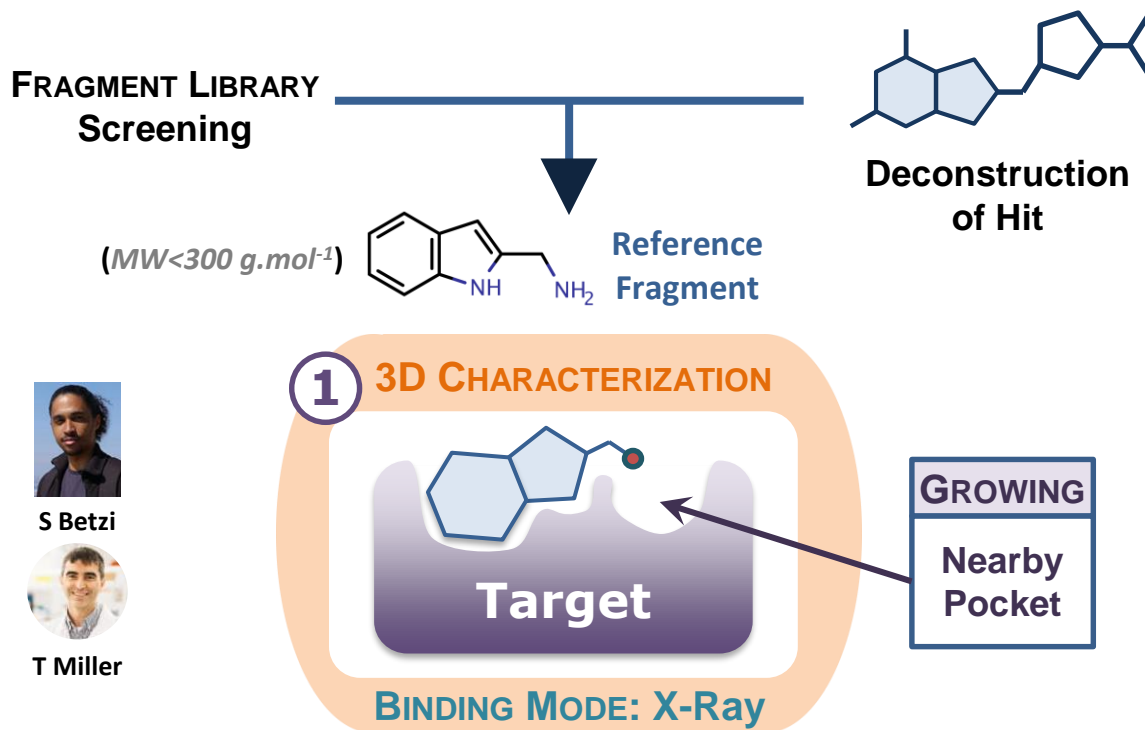
Masitinib interacts with dCK and links to a pocket that overlaps with the active site



DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS

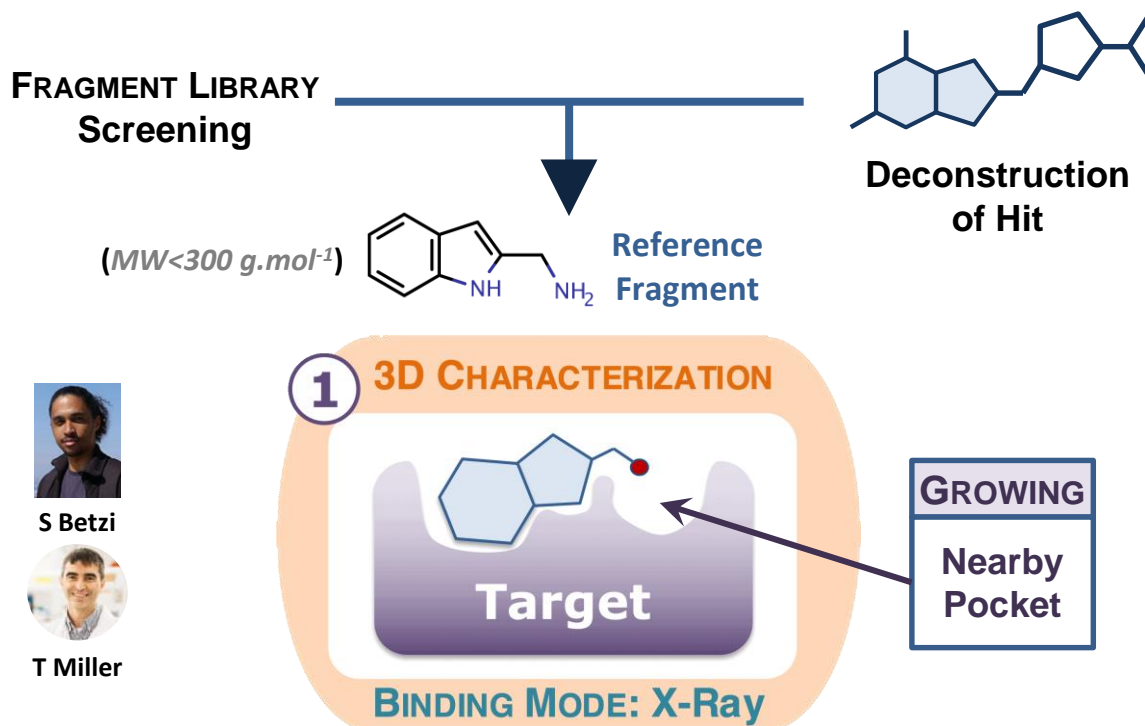


S Betzi



T Miller

DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



S Betzi



T Miller



DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



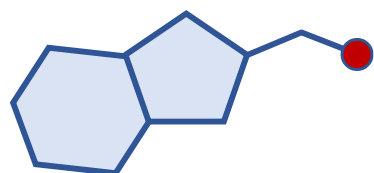
L Hoffer

VIRTUAL CHEMICAL LIBRARY DESIGN

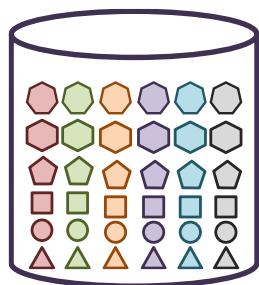
1 3D Characterization

Target

Binding mode: X-Ray



REFERENCE
FRAGMENT



BUILDING BLOCK
LIBRARY
(>6M)

CHEMICAL REACTIONS ▲

x80

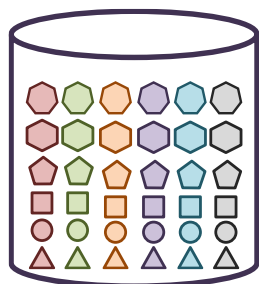
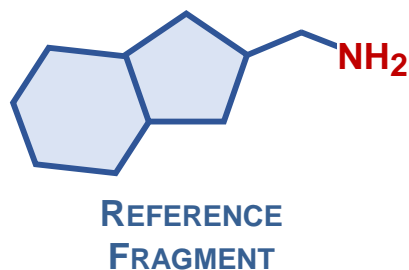
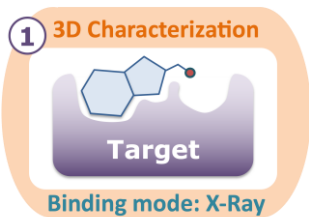
MEDCHEM
CHEMICAL REACTIONS

DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



L Hoffer

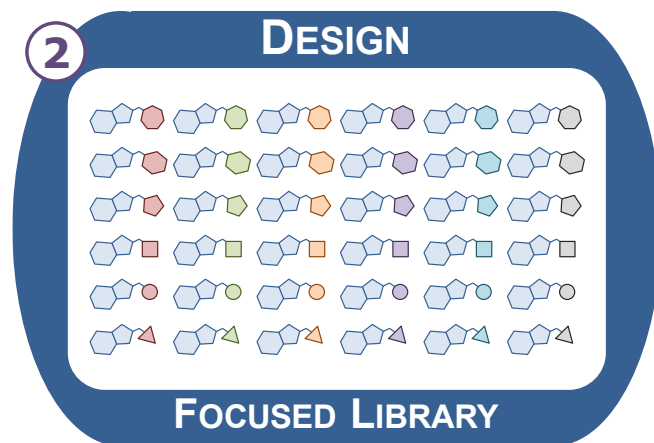
VIRTUAL CHEMICAL LIBRARY DESIGN



BUILDING BLOCK
LIBRARY
(>6M)

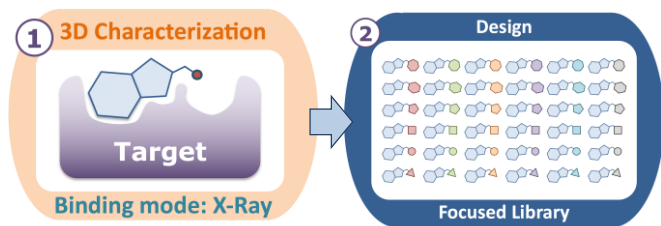
CHEMICAL REACTIONS
RULE_41_VINYL-TERM_HECK
RULE_42_VINYL-HECK
RULE_43_ORGANO-STANNANE
RULE_44_CARBONYL_GRIGNARD
RULE_45_ALCOHOL_GRIGNARD
RULE_46_ALKYNE_SONOGASHIRA
RULE_47_AMIDE_SCHOTTEN-BAUMANN
RULE_48_SULFONAMIDE
RULE_49_N-ARYL_HETEROCYCLES
RULE_50_WITTIG

MEDCHEM
CHEMICAL REACTIONS



ALL COMPOUNDS IN THE VIRTUAL LIBRARY
ARE AMENABLE TO ORGANIC SYNTHESIS

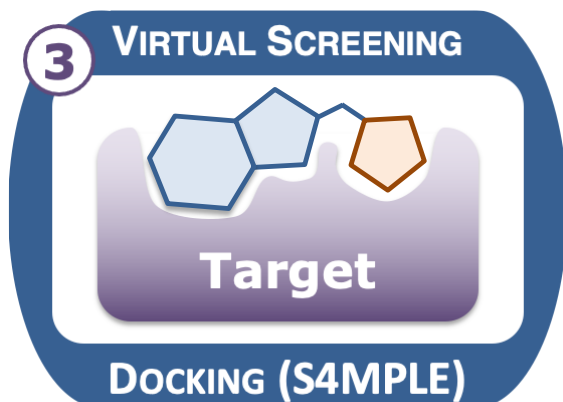
DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



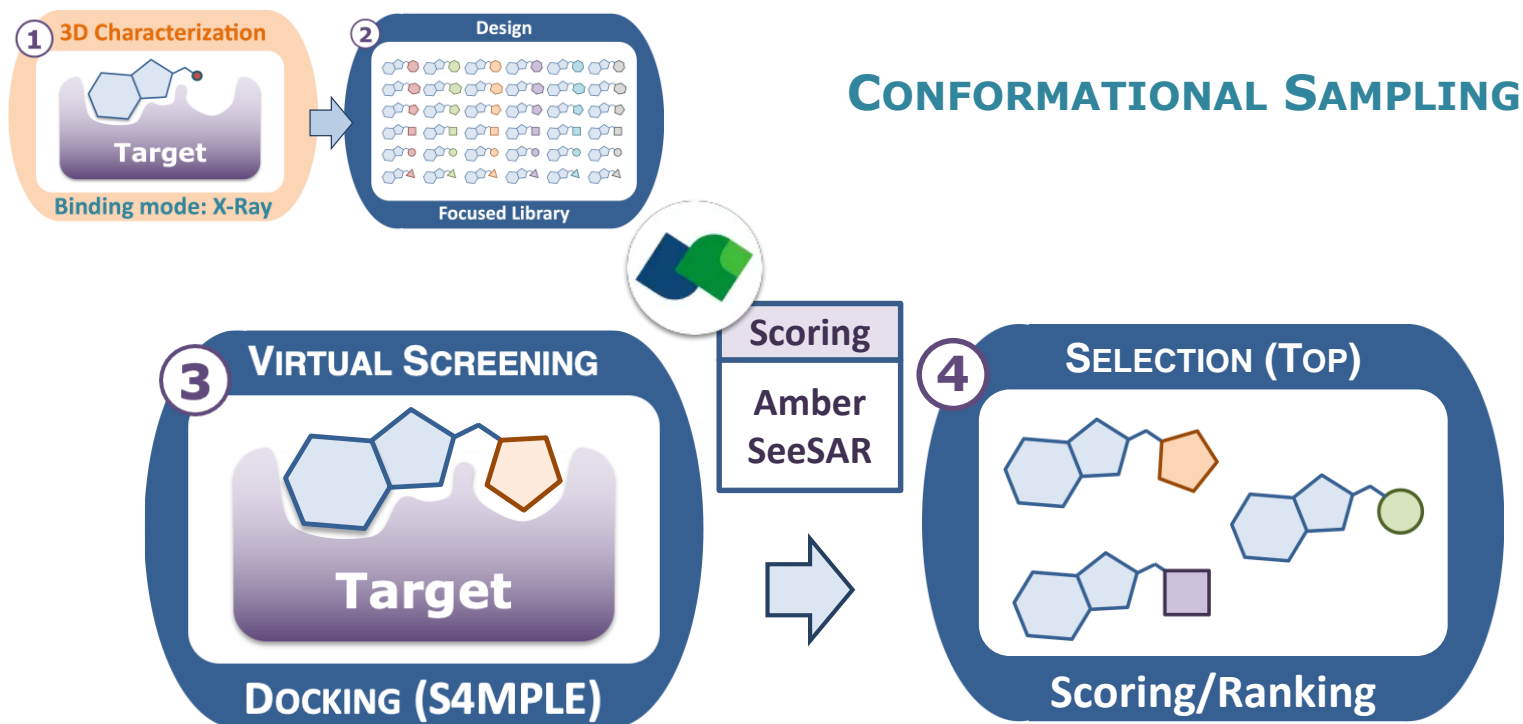
CONFORMATIONAL SAMPLING



L Hoffer

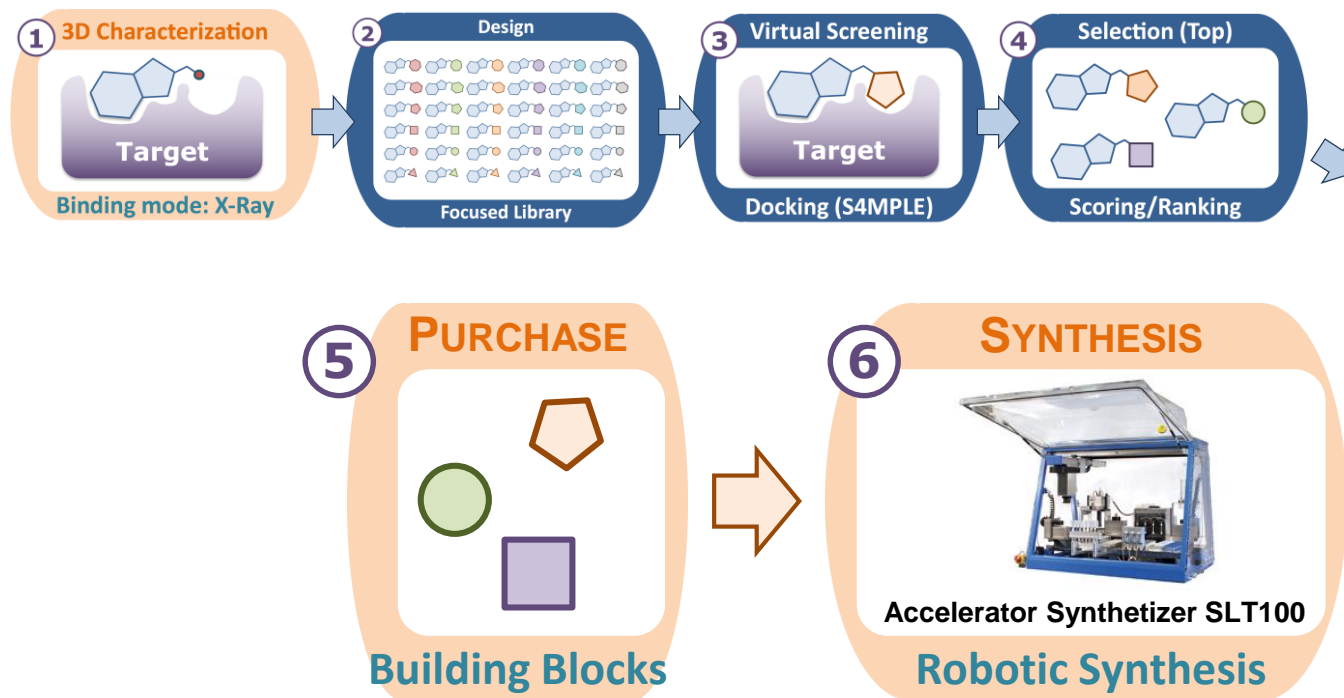


DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



L Hoffer

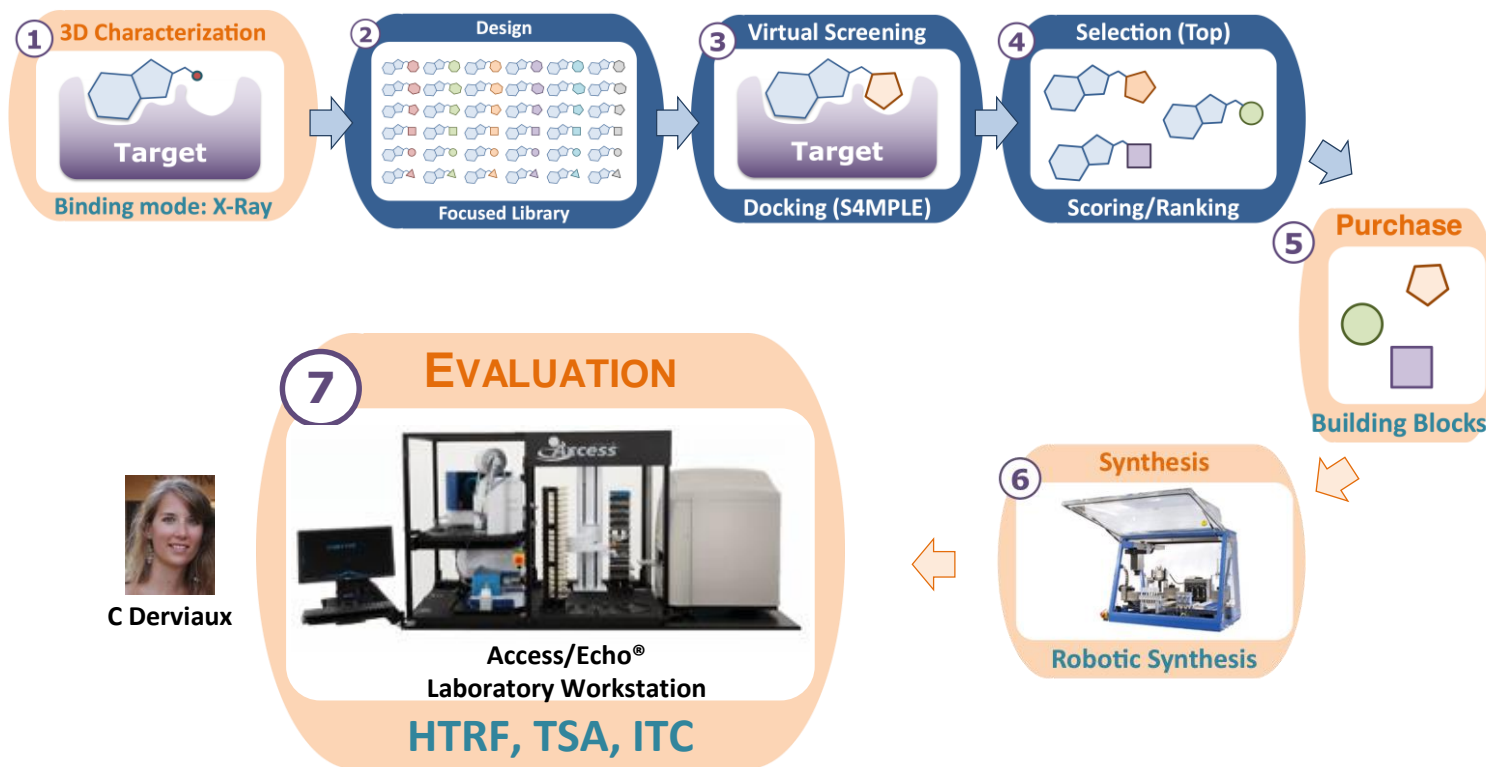
DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



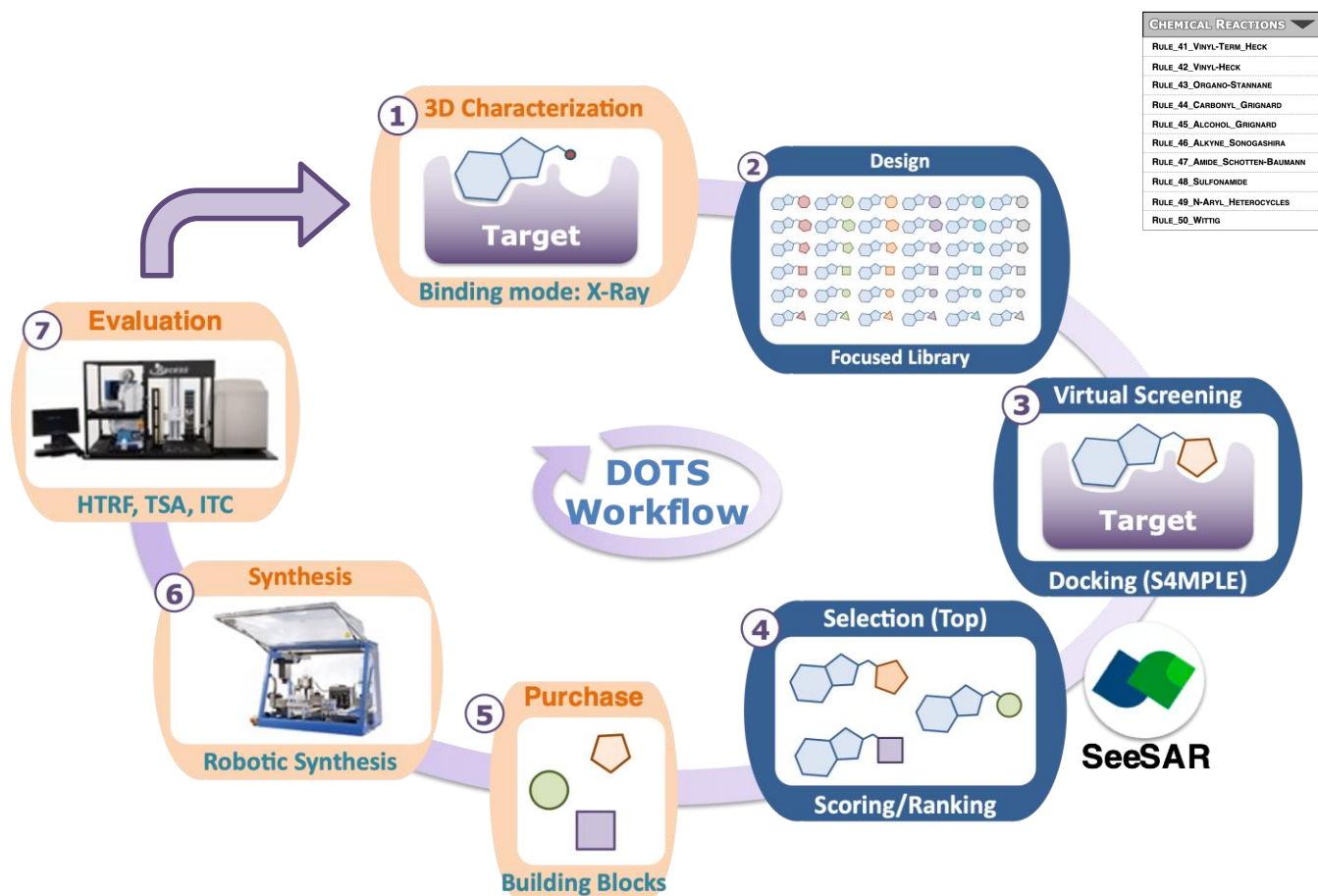
P Bremond S Combes

CHEMICAL SYNTHESIS

DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



DIVERSITY-ORIENTED TARGETED SYNTHESIS: DOTS



The ChemoDOTS (web server generating the virtual libraries) will be available Q3 2023 / Q1 2024

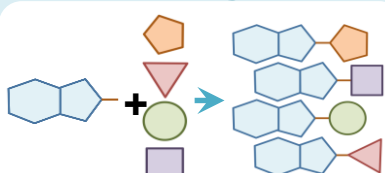
TURNING AN ACTIVATOR INTO INHIBITOR

Characterization



Binding mode: X-Ray

Design



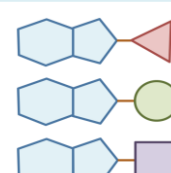
Focused Library

Virtual Screening



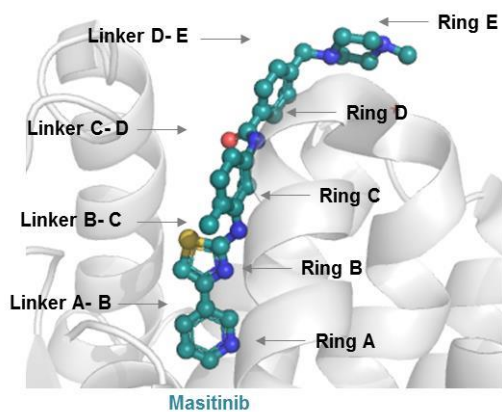
Docking

Selection



Ranking (Scoring)

Structure modifications



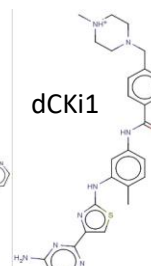
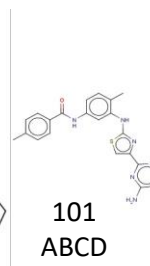
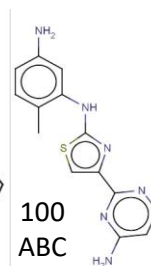
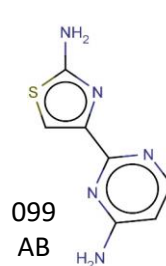
>500

Designed
compounds



Scoring

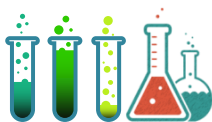
Amber
SeeSAR



Name	Src	Estimated affinity	LLE	LE	Tor.	Intra	Inter	MW	LogP
099								193.23	0.76
100								298.37	2.82
101								416.51	4.80
0_dcki_1								515.66	2.82

TURNING AN ACTIVATOR INTO INHIBITOR

Synthesis



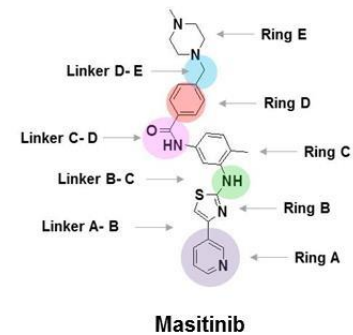
Synthesis

Evaluation

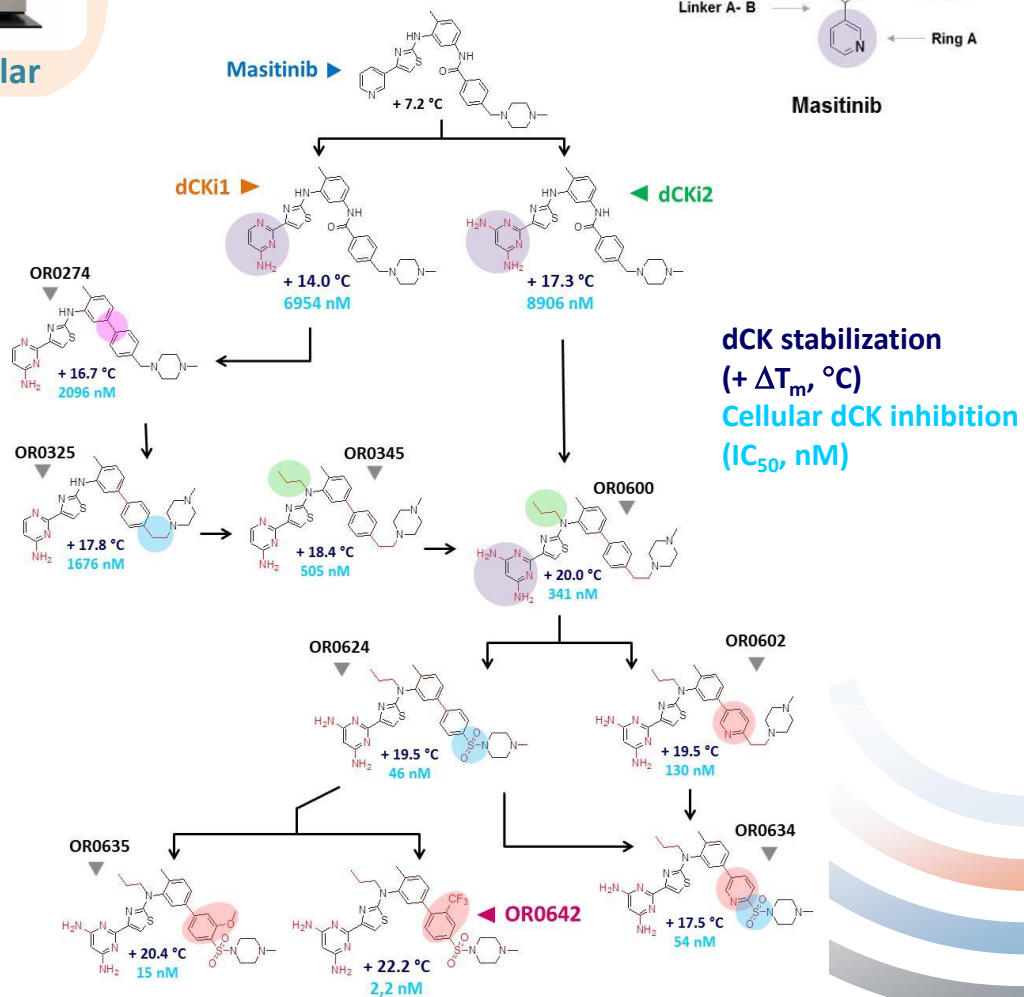
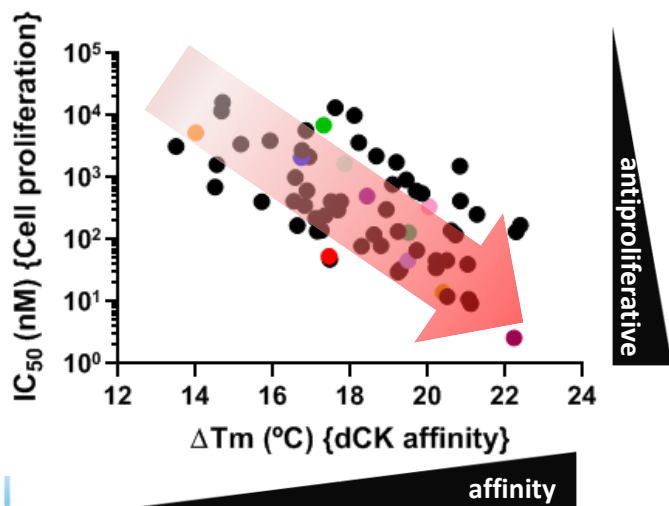


Evaluation: TSA, cellular

Milestone modifications

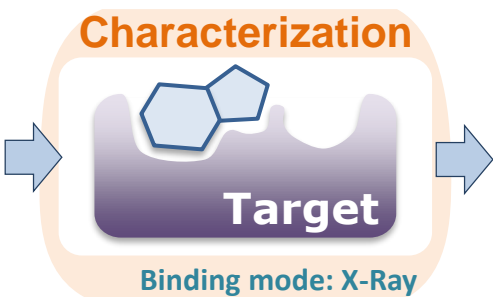


76
Synthesized
compounds



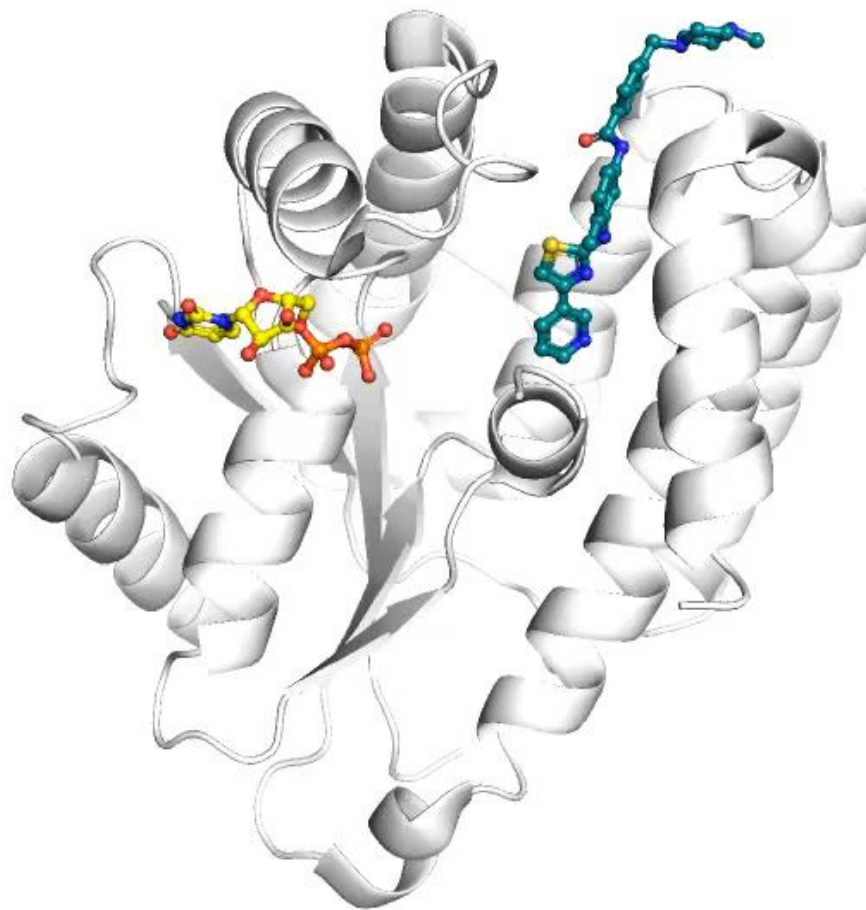
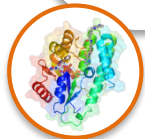
TURNING AN ACTIVATOR INTO INHIBITOR

Characterization



30

Crystal
structures



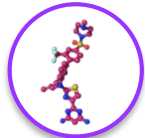
Modifications:

- Cycle A → Di-amino
- Linker B-C → *N*-propyl
- Linker C-D → Direct
- Linker D-E → Sulfonamide
- Cycle D → CF₃

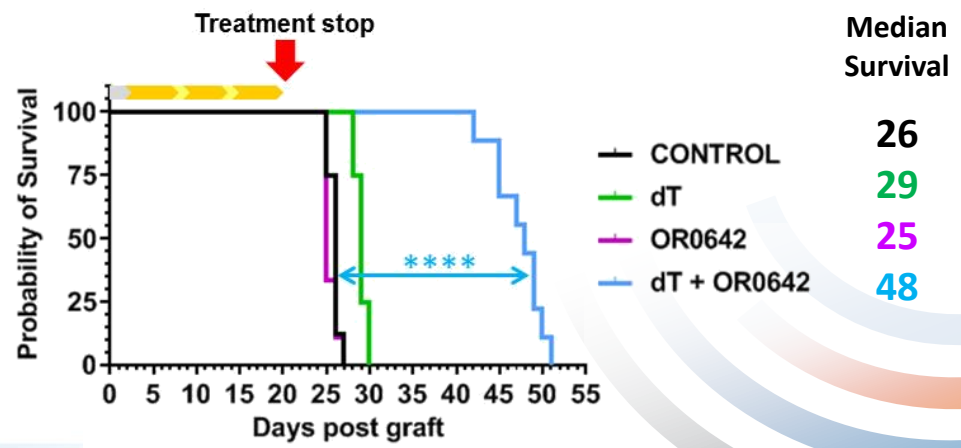
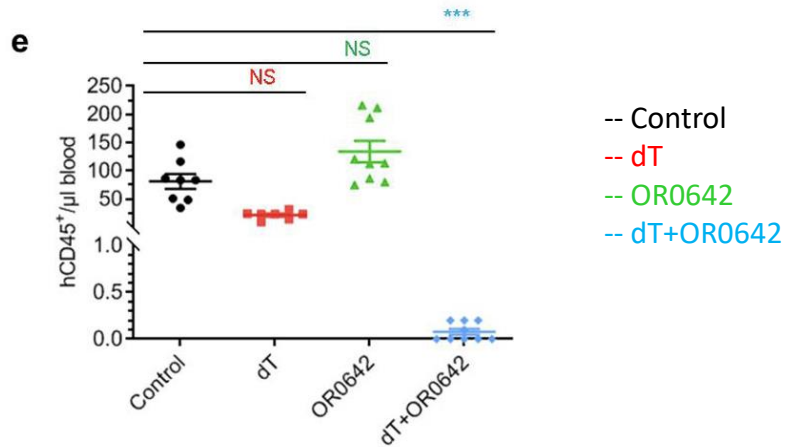
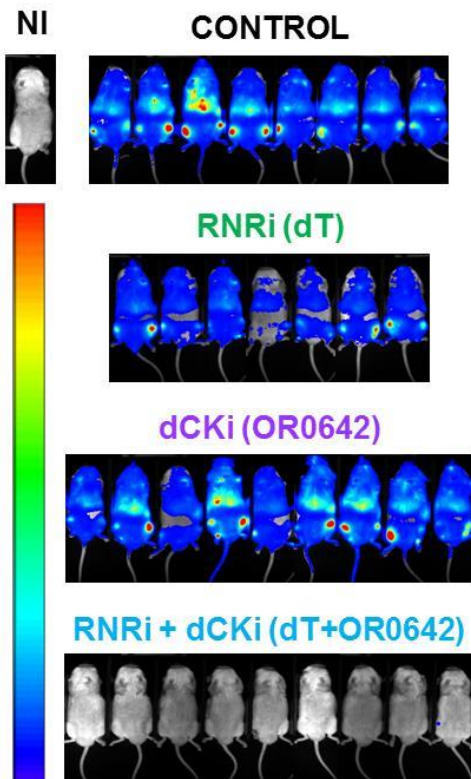
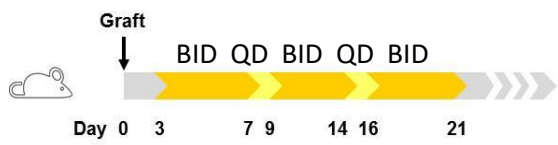
'(pre)Lead' Compound & *in vivo* validation (CCRF-CEM/T-ALL model)

CDX

In vivo cell-line-derived xenograft



OR0642
CCRF-CEM
T-ALL
EC₅₀ = 2.6nM

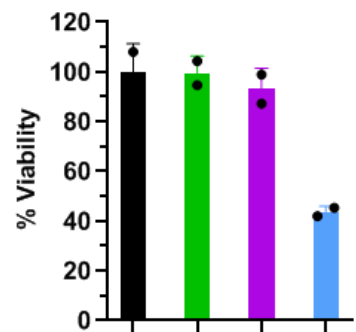
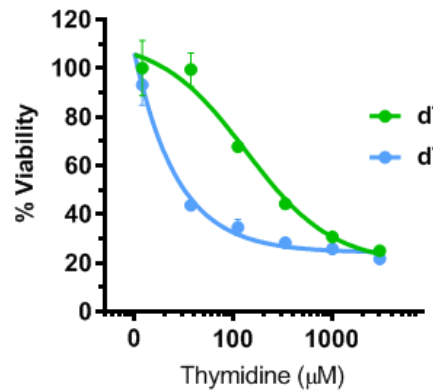


'(pre)Lead' Compound & *in vivo* validation (PDX / T-ALL)

PDX

Ex vivo patient-derived xenograft

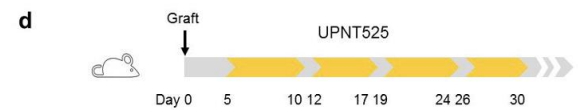
UPNT525
T-ALL



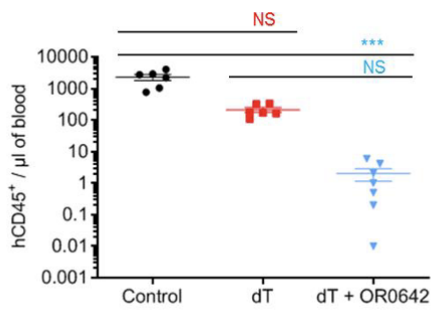
CONTROL
dT 37 μM
OR0642 250 nM
dT + OR0642

PDX

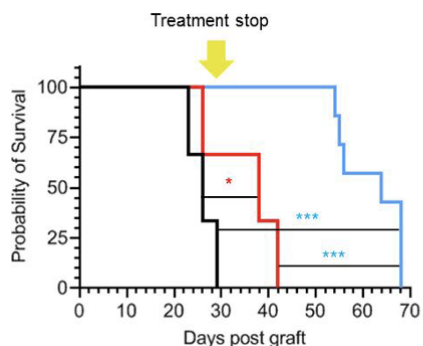
In vivo patient-derived xenograft



e



f



Yellow arrow: Vehicle, dT, dT+OR0642, BID
Grey arrow: No administration

Control
dT
dT + OR0642

Median Survival


26
38
66

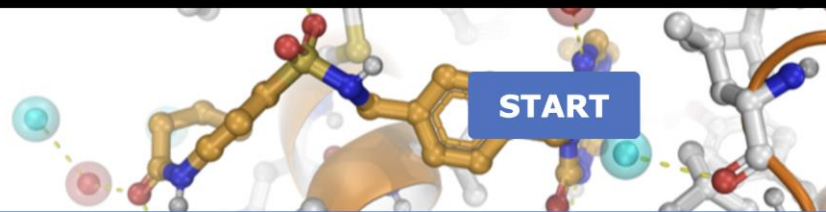
Conclusions & Perspectives

CONCLUSIONS

- From a Drug Repositioning to a Structure-Based Drug Design program
- 'chemoDOTS' transformed a 'dCK' activator, Masitinib, into an inhibitor:
from μM to sub-nM IC₅₀'s
- **OR0642** validation *in vitro*, *ex vivo* and *in vivo* (CDX & PDX)

PERSPECTIVES

- Developing orally-compatible version (**on going synthesis**)
 - PDX studies (on low & highly proliferative patient samples)
 - Biomarkers correlation (w/ single seq RNA from patient samples)
 - Preclinical regulatory studies in 2025
 - Phase I in 2025-2026
- 



Main Menu

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ChemoDOTS

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Citation

Useful Links

iSCB
HITS Platform
CRCM
IPC
CNRS
Inserm

Our Tools

2P2Idb
2P2I-inspector
Fr-PPICChem



[Step-by-step Tutorial](#) [Start a new Project](#)

The web server facility is **freely** accessible to **all users**, including commercial users.

[Structure](#) [Reaction rules](#) [Summary](#) [Raw library](#) [Post processing](#) [Download 2D](#) [Download 3D](#)

Draw Fragment

[Example 1](#) [Example 2](#) [Reset](#)



SMILES code

? How to use the server ?

- 1 Draw or import molecules in sketcher
 - 2 Chemical functions are automatically detected
 - 3 Select function used as attachment point for growing
 - 4 Select reaction(s) for growing
 - 5 Generate raw chemical library
 - 6 Filter library using physico-chemical properties
 - 7 Generate mol2 files with atom types and charges
- Optional steps

Progress through each step by using [Next->](#)

To cite the ChemoDOTS web server, please refer to the following publication:

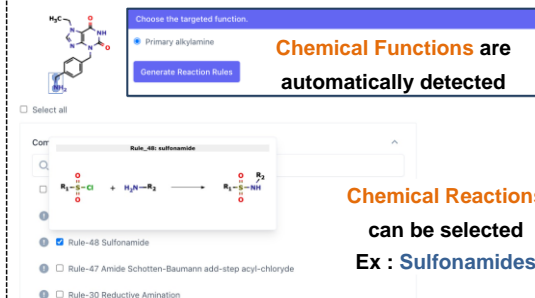
ChemoDOTS webserver functionality

Step 1 : Draw Fragment



The **activated Fragment** can be drawn or imported into the sketcher

Step 2 : Chemical Reactions



Step 3 : commercial libraries

Reaction rules

Hartenfeller 51: Buchwald-Hartwig

Building Block databases

Molport (default)

Molport (default)

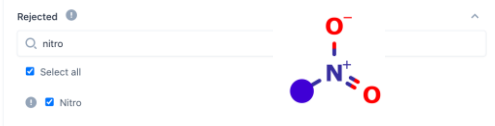
Enamine

Enamine + Molport

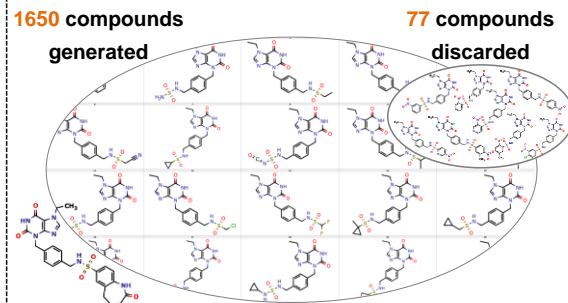
Different commercial building block libraries can be selected with **diverse** compositions and **size**

Step 4 : Undesired Structures

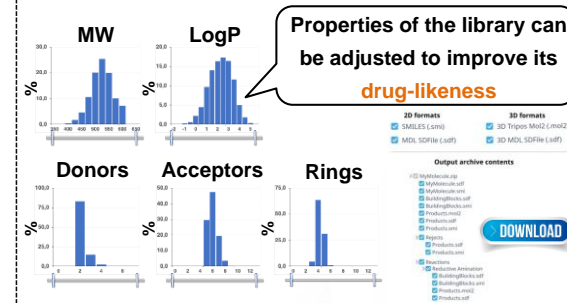
It is possible to select **undesired** sub-structures (ex **Nitro** groups). These sub-structures will be **discarded** from the final chemical library



Step 5 : Library Design



Step 6 : Post-Processing



Rapid: generates ~1000 molecules per second

Practical: all building blocks commercially available

Integrated: compounds produced in "ready to dock" format



Practical Fragments @PracticalFrag · May 28

Welcome ChemoDOTS web server!

practicalfragments.blogspot.com/2024/05/free-c...

Free resource finds functionalities on fragment hits, picks suitable reactions, predicts resulting molecules & lets you refine by cLogP, hydrogen bond donors, and much more

@Xavier_Morelli @CNRS @Inseam



3



6

360



ChemoDOTS utility: retrospective hit to lead for approved drugs

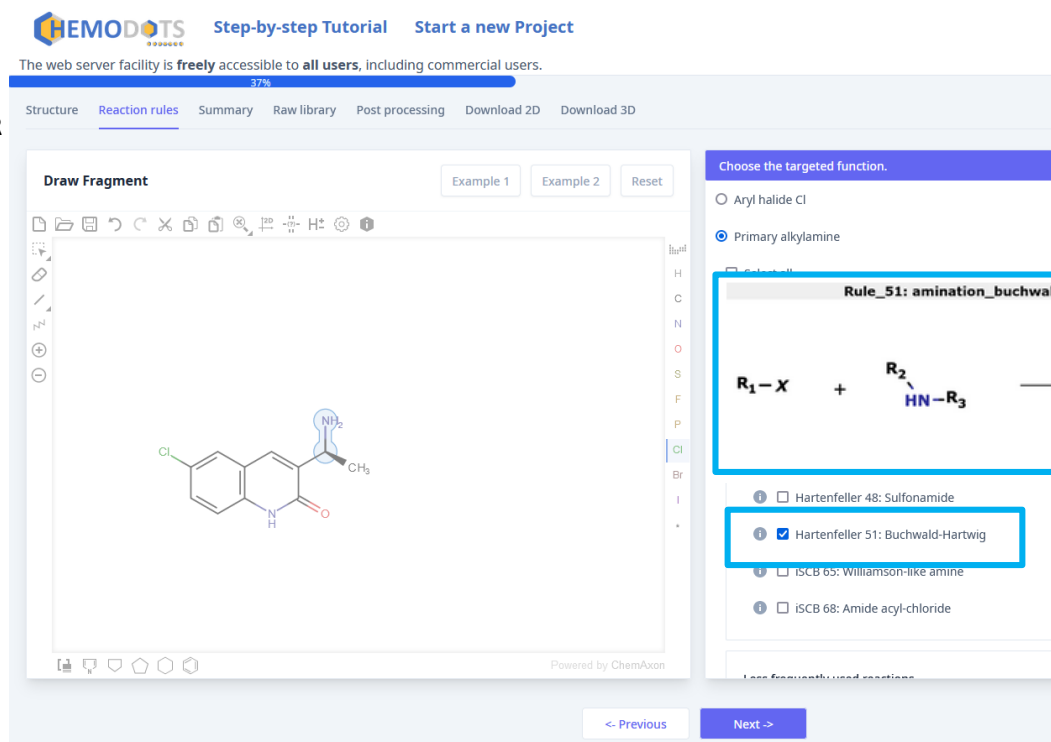
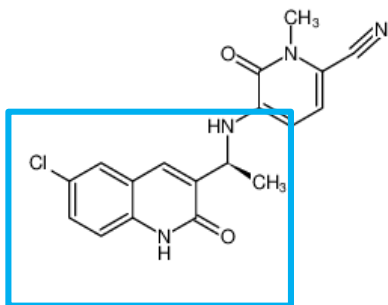
OLUTASIDENIB

(approved in 2022 - RIGEL PHARMS INC)

ANTINEOPLASTIC

ISOCITRATE DEHYDROGENASE-1 (IDH1) INHIBITOR

r/rAML IDH1mut



The screenshot displays the ChemoDOTS web interface. The main window is titled "Draw Fragment" and shows the chemical structure of Olutasidenib with a retrosynthetic arrow pointing to a fragment containing a methyl group and a nitrogen atom. The interface includes a navigation menu with options like "Structure", "Reaction rules", "Summary", "Raw library", "Post processing", "Download 2D", and "Download 3D". A sidebar on the right allows users to "Choose the targeted function", with options for "Aryl halide Cl" and "Primary alkylamine". The "Primary alkylamine" option is selected, and a specific reaction rule, "Rule_51: amination_buchwald-hartwig", is highlighted. This rule shows the reaction of an aryl halide (R₁-X) with a primary amine (R₂-NH-R₃) to form a secondary amine (R₁-N(R₂)-R₃). Below the rule, a list of reaction rules is shown, with "Hartenfeller 51: Buchwald-Hartwig" selected. The interface is powered by ChemAxon and includes navigation buttons for "< Previous" and "Next ->".

>80k synthetically accessible analogs generated in ~2 min



Step-by-step Tutorial Start a new Project

The web server facility is **freely** accessible to **all users**, including commercial users.

Structure Reaction rules Summary **Raw library** Post processing Download 2D Download 3D

Raw library download

Here you can find some statistics about the raw library generation. The downloads are available [below](#).
Duration of library generation: **00:02:05**

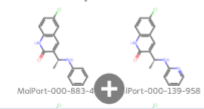
You can **bookmark** the following link to reload and **access the saved data**, which will be stored for **two weeks**:
<https://chemodots.marseille.inserm.fr/growing?tab=4&experiment=a7c39758-27d1-4a05-8109-07cd2174b6ba>

Overall

Total building blocks	501542
Reacted building blocks	16.2% (81340/501542)
Generated products	81340
Duplicate products	19.0% (15484/81340)
Final products	81.0% (65856/81340)
Overall contribution to chemical space	100.0% (65856/65856)

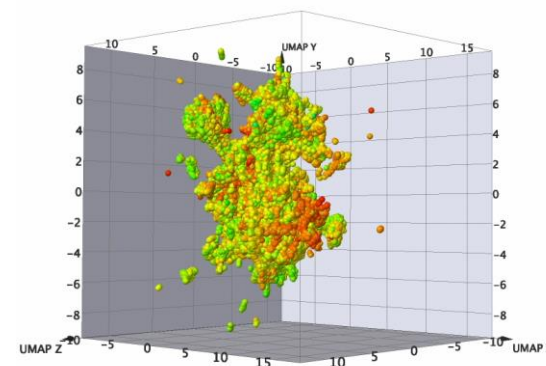
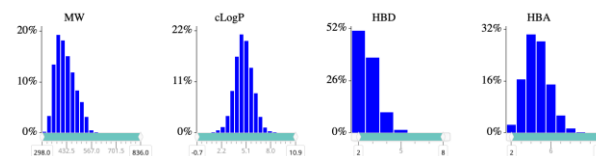
Hartenfeller 51: Buchwald-Hartwig

Generated products overview



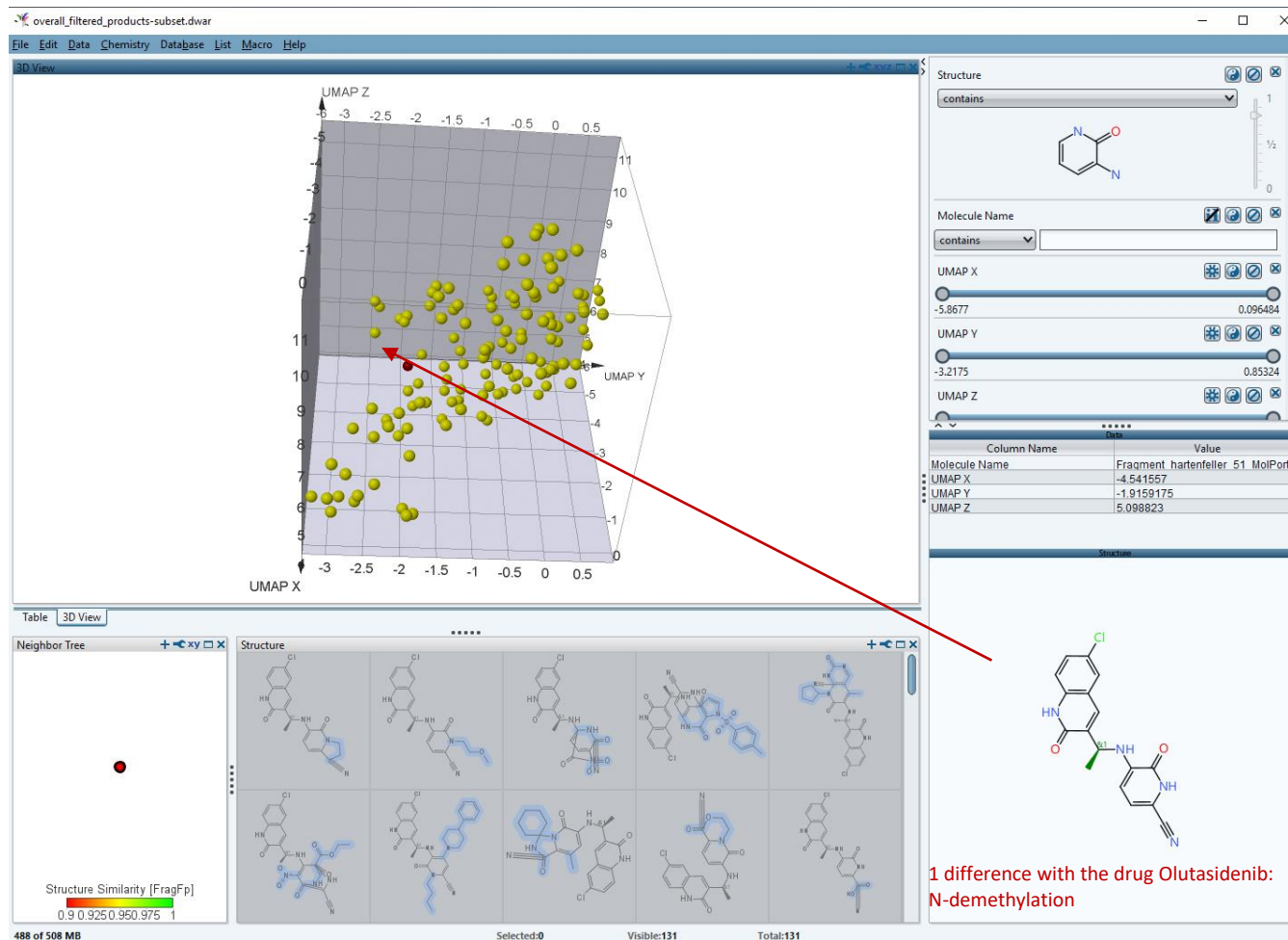
<- Previous

Next ->

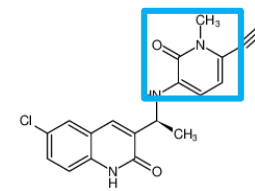


Molweight 300 400 500 600 700 800 900

131 analogs have a pyridinone group as in the drug Olutasidenib



Olutasidenib



ChemoDOTS integration with virtual screening to select synthesis priorities

HEMODOOTS

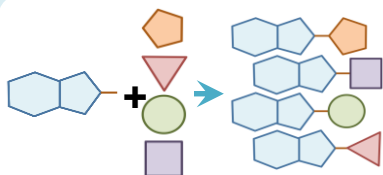
Characterization



IDH1:6U4J

Binding mode: X-Ray

Design



Ready to dock library

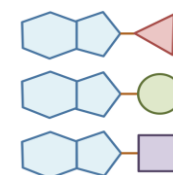
Virtual Screening



IDH1:6U4J

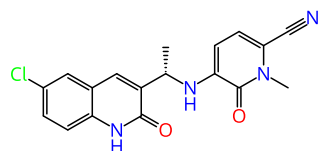
Docking

Selection

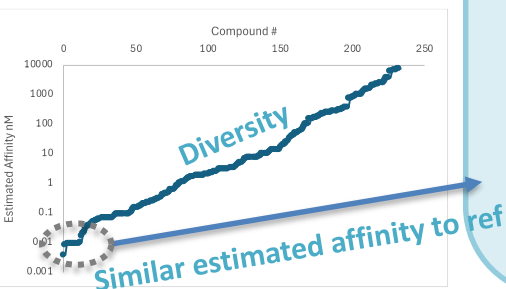
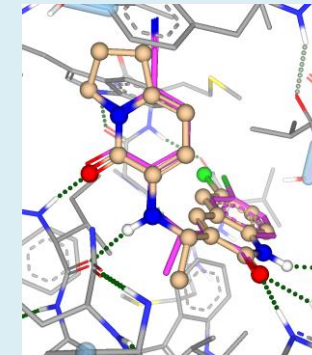
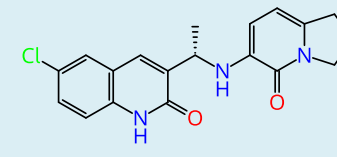
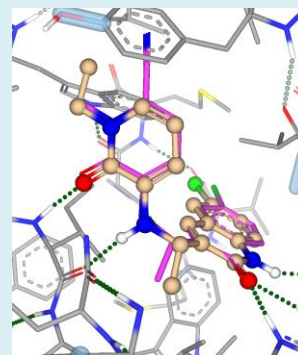
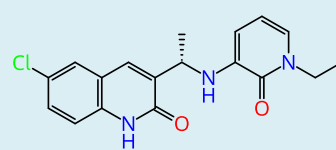
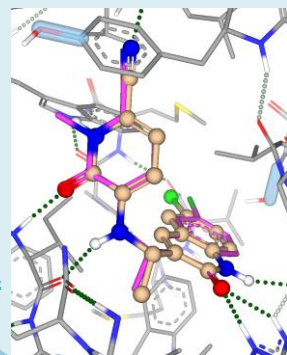
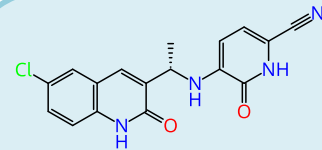


Scoring
Amber
SeeSAR

Ranking (Scoring)

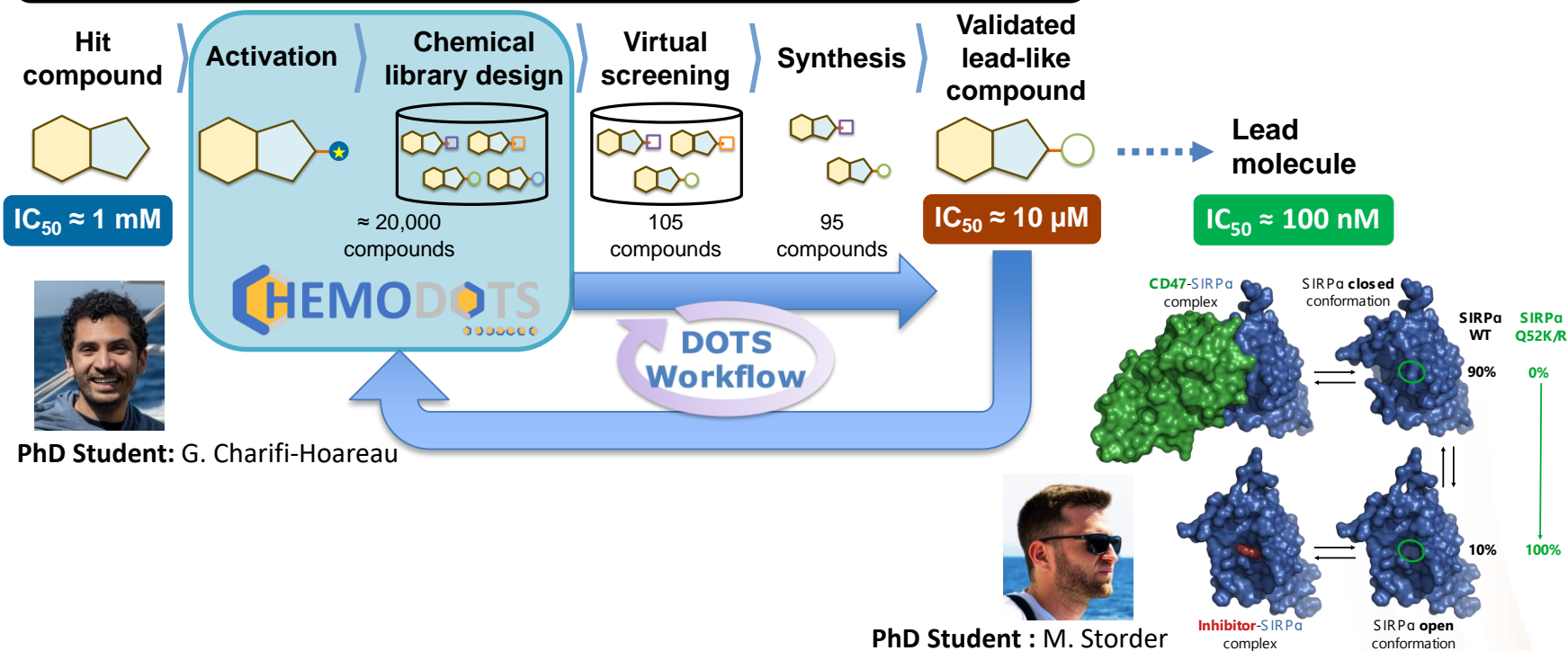


Olutasidenib



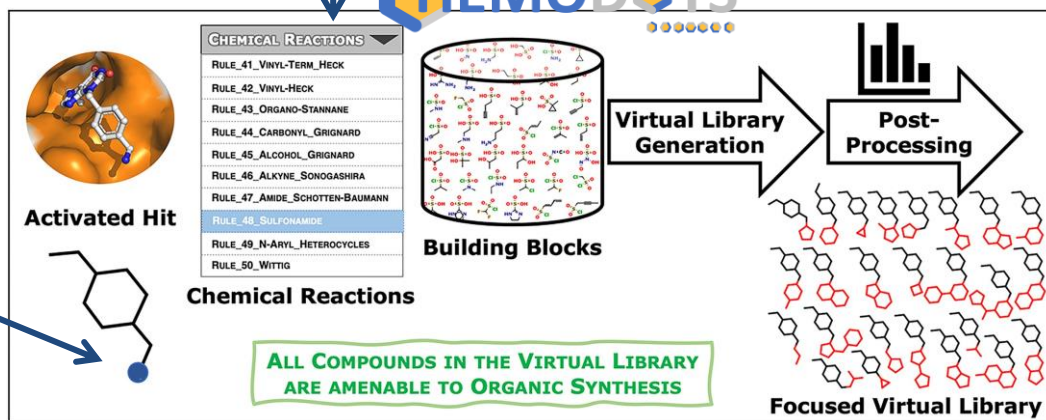
ChemoDOTS ongoing utilization: hit to lead optimization of novel CD47-SIRP α immune checkpoint inhibitors

A **low-affinity hit** identified from screening was refined into a **high-affinity lead-like** inhibitor of SIRP α -CD47 interaction



Continuous improvement and Perspectives

Update with latest robust chemical reactions (e.g. C-C coupling)



Incorporate linking and covalent design strategies



Direct integration in fragment screening libraries

CHEM
18 17 16 15 14 13 12 11 10 9 8 7 6 5 4 3 2 1

NEW!

Ultrafast NMR-based fragment screening
Enhanced Screening Efficiency
Low Sample Concentration
Cost Effective

Fragment Library
450+ COMPOUNDS

Hit-to-lead optimization
utilizing Chemspid's extensive catalog of small molecules

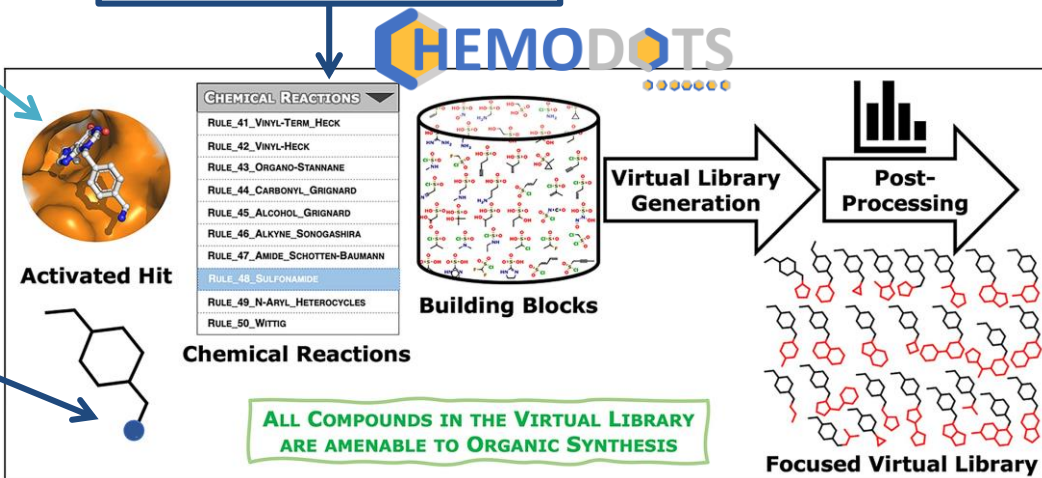
NEXMR

Continuous improvement and Perspectives

Hit ID strategies
CryptoDOTS

Update with latest robust
chemical reactions (e.g. C-C
coupling)

Automated
proposition of
activated fragment



Intelligent library
refinement

Virtual screen

ChemoDOTS
GOLD LIBRARY

Incorporate linking and
covalent design strategies



Direct integration in
fragment screening
libraries



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 Etienne Rebuffet
 Benoit Sicard



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 Camille Montersino



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 Guillaume Andrieu



Aznam Yacoub
 Namarta Vij

