



Development of a third party-accessible, enabling platform for efficient preclinical drug discovery

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WP1 – Overall Goal

Enabling platform for the delivery of **optimised leads** at an **accelerated pace**



Lead optimisation is traditionally considered the place where academic drug discovery projects go to die a slow and painful death

- **Deployment of an actual platform of general applicability**
- **Identification of optimised leads**



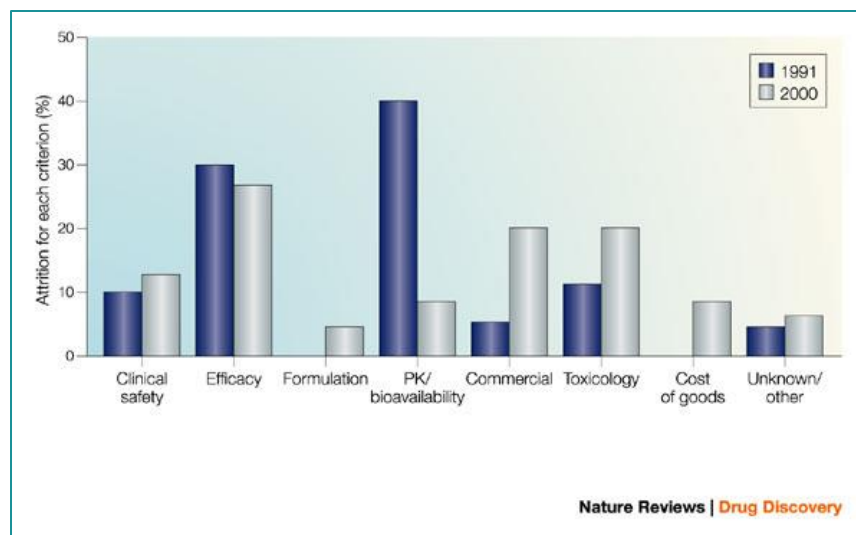
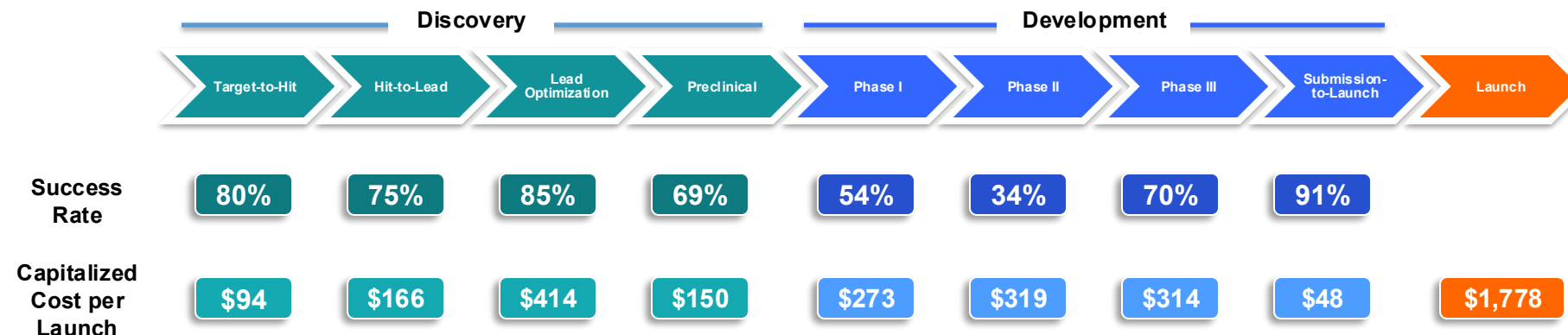
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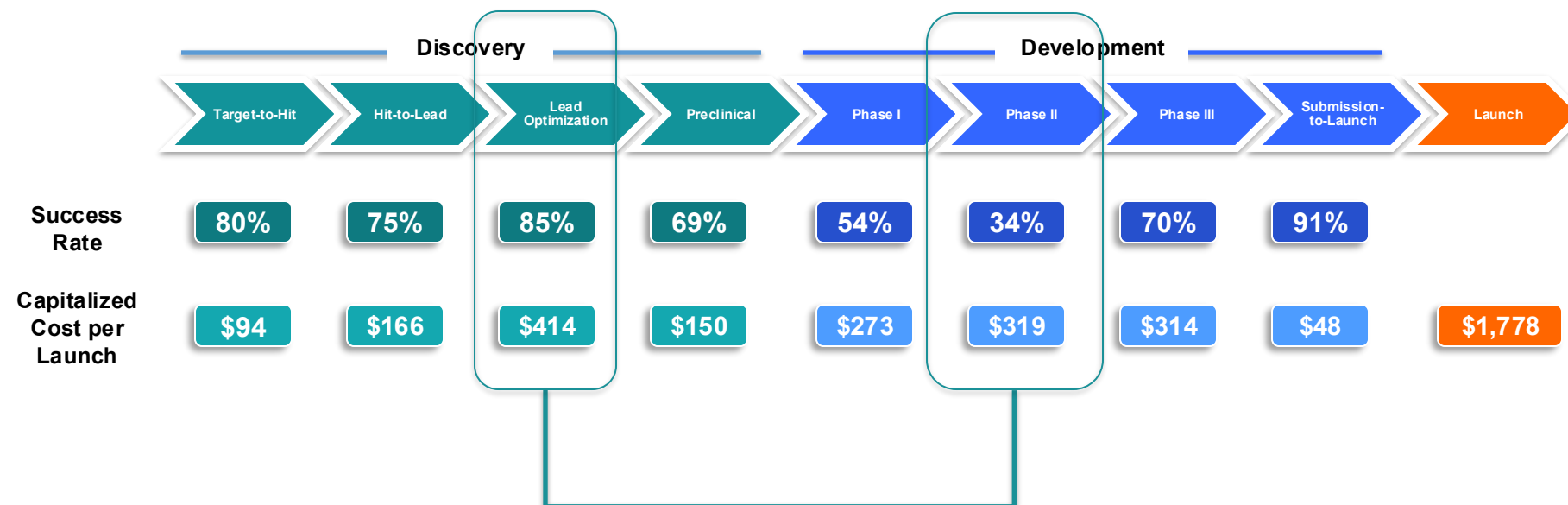
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**Intervening in the discovery phase is
the best way to reduce attrition rates
down the line**

Adapted from Paul *et al.*, Nat Rev Drug Discovery 2010. All costs are in million 2008 USD

Kola and Landis, Nat Rev Drug Discovery 2004.



A parametric sensitivity analysis indicates that improving the **lead optimization** step provides the greatest beneficial effect on the progression rate in Phase II and on the **total cost per launch**



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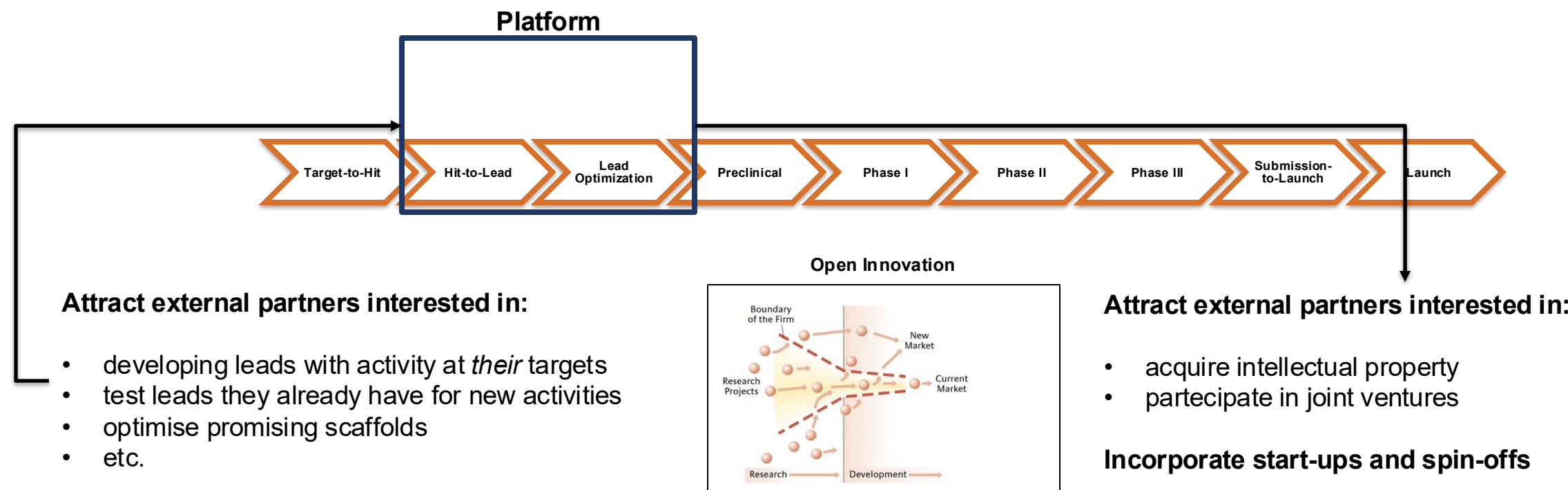
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The Open Innovation Paradigm

*“Open innovation is the use of purposive **inflows** and **outflows** of knowledge to accelerate internal innovation, and expand the markets for external use of innovation.”* Henry Chesbrough



Henry Chesbrough, *Open Innovation. The New Imperative for Creating and Profiting from Technology*. HBR Press 2003

WP1 – Impact

IMPACT

Description

In line with the tenets of **open innovation** (Chesbrough, 2006), we envision a bi-directional flow of people and ideas between UniUrb and companies, that is made possible by an agile and externally accessible platform that enables lead identification and optimisation. Encompassing advanced computational methods, synthetic chemistry skills, in vitro/vivo pharmacology and sound knowledge in key therapeutic areas, the **platform generates** value through:

i) **Unique know-how**, made available to both **big players** and **local SMEs**, limiting the need for **massive infrastructural investments** (for example, for high-throughput wet screening), **shortening time to market**, and, thus, with clear potential for **attracting extramural funds and partnerships**.

ii) **Composition of matter patents**, claiming in house generated compounds active in key therapeutic areas such as oncology. This **IP** is instrumental to **incorporate spin-offs** and attract investments especially within the **emerging framework of portfolio-companies**.

Means of Verification

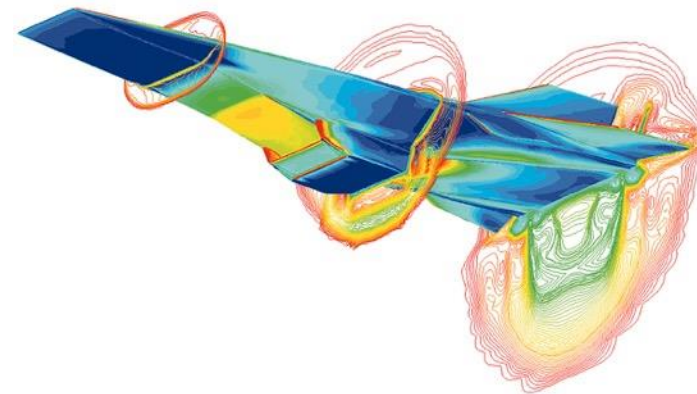
- Number of research contracts involving the platform
- Total amount of extramural funds attracted

- Number of patents granted
- Number of patents out-licensed
- Number of spin-offs incorporated



Of Drugs and Airplanes

“Many **major aerospace projects** require a decade or more of R&D along with over a billion dollars in investment to get a single new product to market.”



Why do not we approach drug discovery the same way?

- Biology is involved
- Limited accuracy of the input
- Multi-disciplinary Effort
- Information is largely incomplete

“All models are wrong, some are useful” G. Box

MISSIONE 4
ISTRUZIONE
RICERCA



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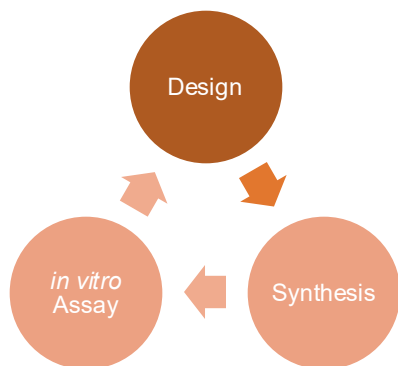


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CARLO BO

WP1 - Development of a third party-accessible, enabling platform for efficient preclinical drug discovery



- Ultra Large Commercial Libraries (10^9)
- DL-generated Compounds (10^7)
- Combinatorial SAR Explorations (10^4)

Compound Prioritization by
Computational Means
(10^2)

Synthesis

Biochemical
or Functional
Assays

Surface
Plasmon
Resonance
(SPR)

MicroScale
Thermophoresis
(MST)

Circular
Dichroism
(CD)

Biophysical Assays

Nuclear
Magnetic
Resonance
(NMR)



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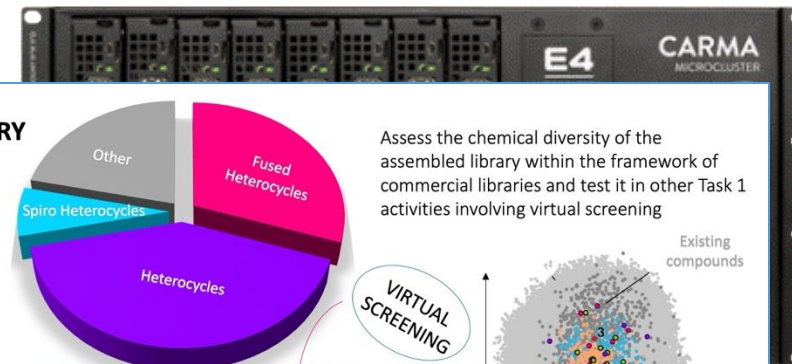


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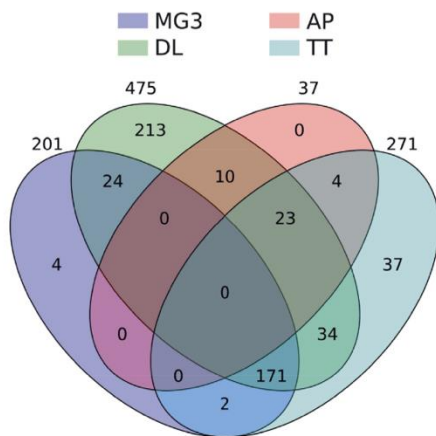


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WP1.1: Computationally-driven, Integrated Infrastructure for Lead Identification and Optimisation

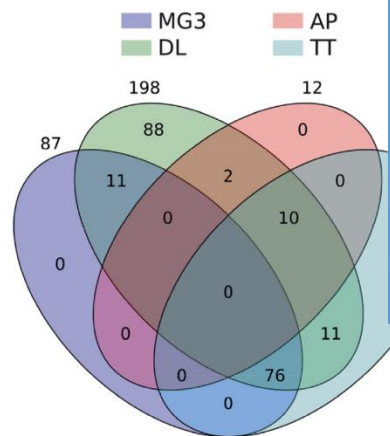


C. Sirocchi, F. Bianucci et al.



(a) Ataxia dataset.

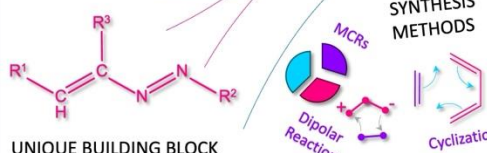
Computer Methods and Programs in Biomedicine



(b) Hypoxia dataset.

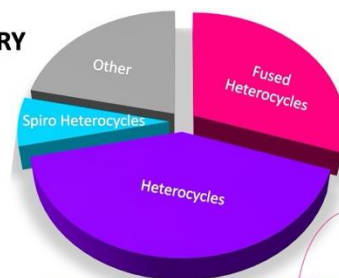
UNIURB PROPRIETARY LIBRARY
MORE THAN 2500 COMPOUNDS

GDB-UniUrb-HetsLab

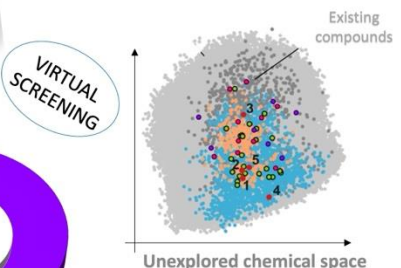


UNIQUE BUILDING BLOCK

This activity aims at compiling a virtual library of novel compounds synthesized at UNIURB over the years, which are: a) available or that b) can be readily synthesized thanks to the proprietary expertise at UNIURB



Assess the chemical diversity of the assembled library within the framework of commercial libraries and test it in other Task 1 activities involving virtual screening



Identify a suitable diversity subset to be tested in targets and/or cell lines relevant to the project



CELL LINES ASSAYS



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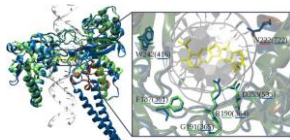
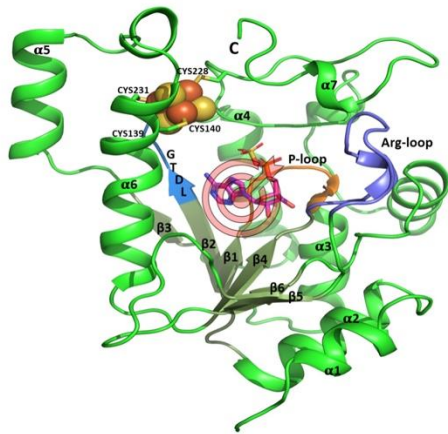


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WP1.1: Computationally-driven, Integrated Infrastructure for Lead Identification and Optimisation

Case study: Selection of potential adenosine 5'-phosphosulfate reductase potential inhibitors



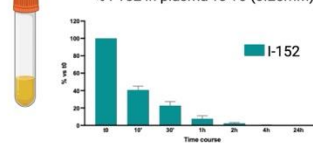
Structural alignment between LiTop1B (PDB 2b9s) and hTop1B crystallized in covalent complex with DNA and in the presence of the inhibitor topotecan.

Rosa-Teijeiro, C. et al. Parasites Vectors 14, 438 (2021).
<https://doi.org/10.1186/s13071-021-04947-4>

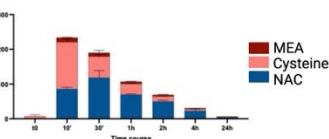
Redox modulating agent: I-152



% I-152 in plasma vs T0 (0.25mM)

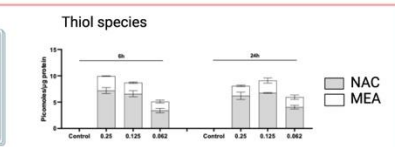
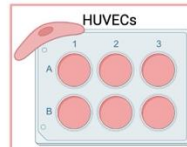
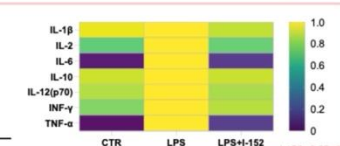
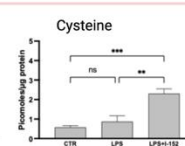
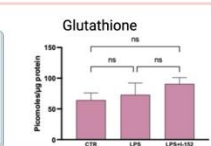
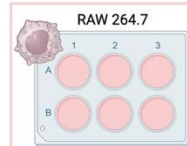
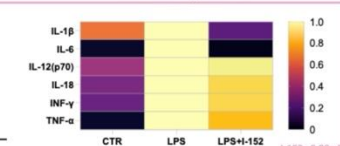
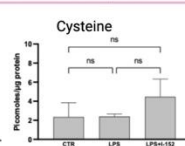
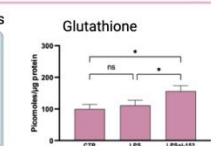
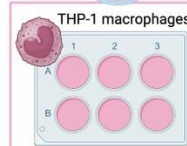


Thiol species in plasma



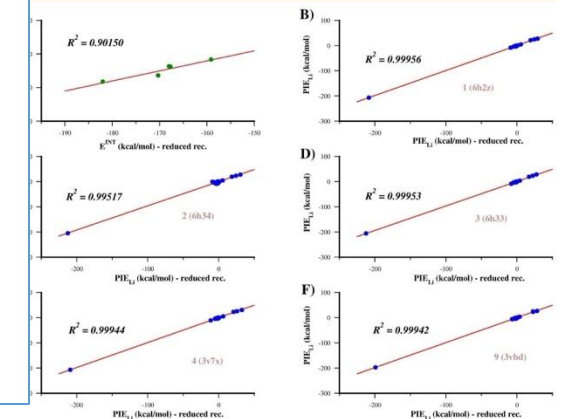
Intracellular thiol species

Secreted cytokines



binding affinity by using the y of hCA II inhibitors

The E_{INT} computed considering the *core model* and the *entire receptor* are in fair agreement, with $R^2 = 0.90$, as well as the single PIE between ligands and each residue ($R^2 \sim 0.999$). Thus, the core model of LR complexes will be adopted in the next steps of our project to compute the E_{INT} and the binding energies (ΔE_{FMO}) at higher level of theory, with the final aim to build a QM-based scoring function for prediction of ligand-metalloenzyme binding affinity.

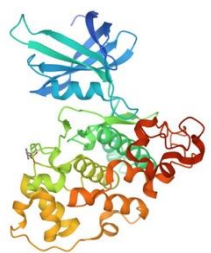


hCA II inhibitors

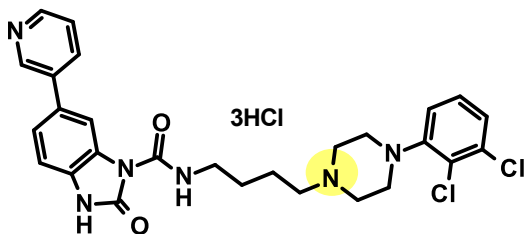
In vitro infection model



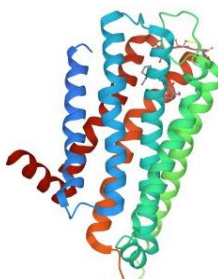
WP1.2: Development, validation and application of tools for rational polypharmacology



Target 2



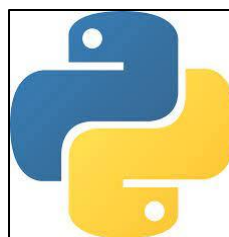
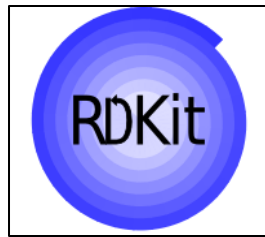
DTDL



Target 1

Manually curated dataset encompassing 158 DTDLs

```
1 from chembl_webresource_client.new_client import new_client
2 import json
3 import os
4 import pandas as pd
5
6 os.chdir('/common/users/vlembo/TESE/PYTHON_SCRIPTS/RETRIEVE_BY_ACTIVITY/TARGET_ACTIVITIES_JSON/')
7
8 target1_id = "CHEMBL1865"
9 target1_name = "HDAC6"
10 target2_id = "CHEMBL1827"
11 target2_name = "PDE-5A"
12
```



Lembo and Bottegoni, **2024** J Med Chem, *just accepted*

- Targets are paired based on pre-existing independent validation at a given pathology. The promise of network pharmacology (Hopkins, Nat Chem Biol **2008**), that the druggable genome would have been greatly expanded by new target combinations, has yet to be fulfilled.
- DTDLs largely exploit known regions of the chemical space that the members of the target pair have in common
- Representative DTDLs never significantly depart from single target prototype compounds

**We have barely scratched the surface
of what D/MTDLs have to offer to
therapy and ultimately human health**



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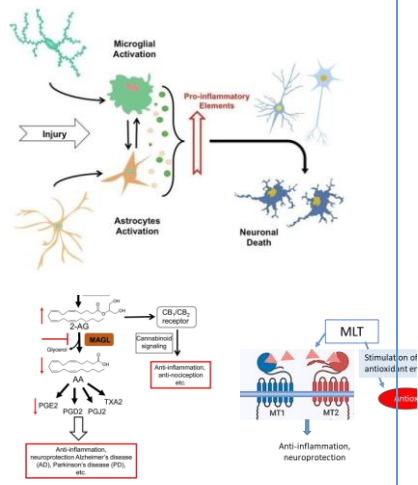
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WP1.2: Development, validation and application of tools for rational polypharmacology

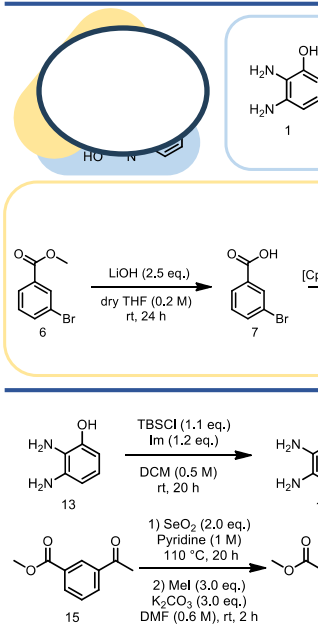
VITALITY – Spoke 8 – WP1 – Task2

Title: Discovery of dual agents targeting monoacylglycerol lipase (MGL) and to promote neuroprotection (Gilberto S. et al.)

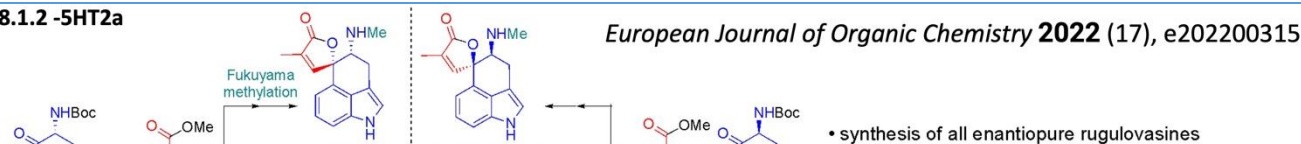
Potential neuroprotection induced by MGL inhibitors and melatonin



POTENTIAL DUAL AGENTS QUINOXALINE AND



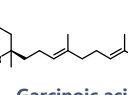
Task 8.1.2 -5HT2a



Garcinoic acid: A natural lead compound advanced to chemical optimization



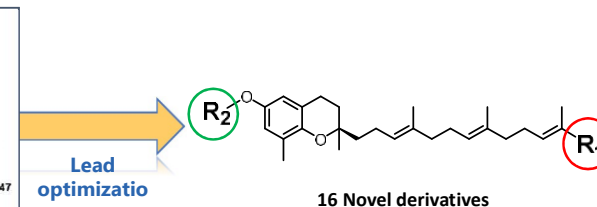
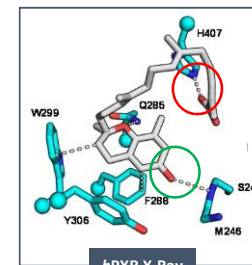
Garcinia kola



Garcinoic acid

Key hot-spots

PXR activity
Off-target (selectivity): PPARα, 5-LO
Metabolism



- Synthesis of novel garcinoic acid analogs (n= 16) (*Gioiello - UniPg*) ✓
- *In vitro* screening of the novel derivatives as PXR ligands (*Tes Pharma*) ✓
- Computational analysis (structure-activity relationship) (*Bottegoni - UniUrb*)
- *In silico* e biophysical PK/PD profiling of best PXR ligands (*Goracci/Macchiarulo - UniPg*)
- *In vivo* validation of best PXR ligands (*Galli/Zelante - UniPg*)
- Scale-up optimization of preclinical candidates (*Gioiello/Vaccaro - UniPg*)



DSS - Colitis



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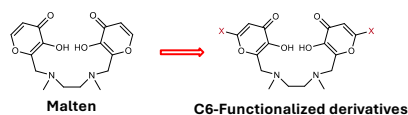
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WP1.3: Test bed for preclinical compounds against innovative targets for metastatic cancer

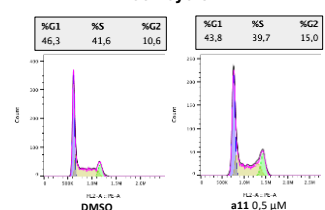
Polyamino-bis-Maltol Compounds



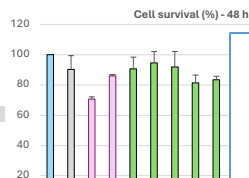
Maltol analyzed at both biological and molecular levels for its antiproliferative activity against cancer cells showed biological activities associated with the modulation of genes having key roles in cell progression and apoptosis. Recently a new set of compounds (polyamino-bis-maltol derivatives) was synthesized by functionalizing the C6 units with different groups.

[*British Journal of Cancer* (2012) 103(2), 231-240]

cell cycle 24 h



The biological activity of **a11** was deepened using the slight accumulation of cells in the G2/M phase of the cell cycle. The percentage of hypodiploid cells starting from the cell cycle activity on U937 cells through the activation of the cell cycle confirm this observation.



The new set of molecules in parallel with

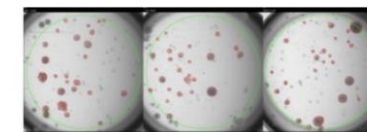
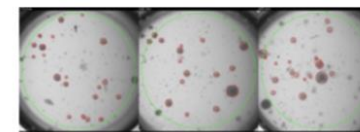
Lead compound optimization o

- Modification of three side chains

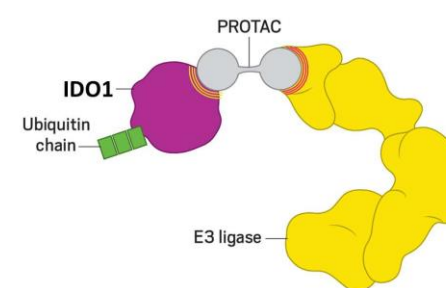
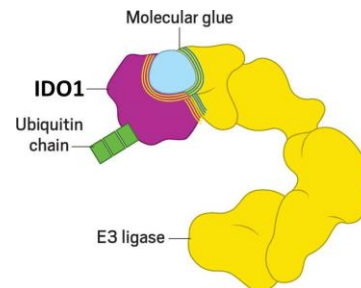
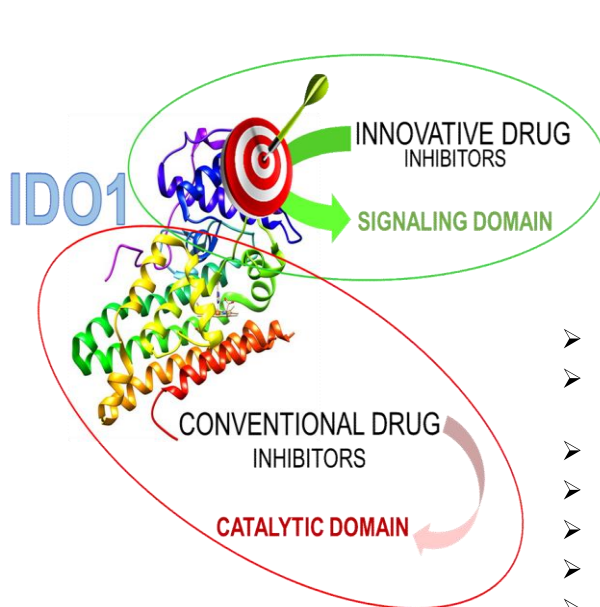
- Docking analysis



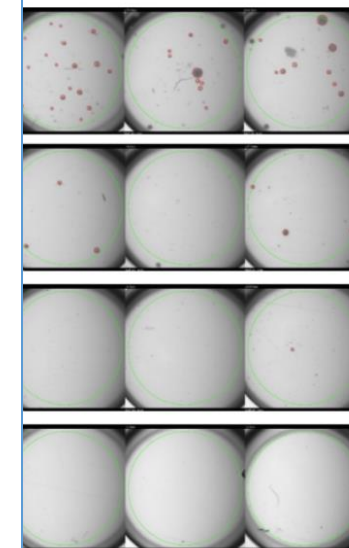
CTR



Design and development of IDO1 protein degraders (PI: Ciriana Orabona)



#36 – UR1286



- Characterization of IDO1-mediated signaling in tumor cells (*Orabona – UniPg*) ✓
- Rational design of small molecules stabilizing IDO1 conformations suitable for the protein degradation (*DSF – UniPg*)
- Synthesis of IDO1 degraders (*UniPg + UniUrb?*)
- *In vitro* screening of IDO1 degraders (*Orabona – UniPg*)
- Biophysical PK/PD profiling of best IDO1 degraders (*Macchiarulo – UniPg*)
- *In vivo* validation of the best IDO1 degraders (*Orabona – UniPg*)
- Scale-up optimization of preclinical candidates (*UniPg + UniUrb?*)

Dose-dependent uptake of borono tryptophan derivatives. Both have been proposed to express CMT1



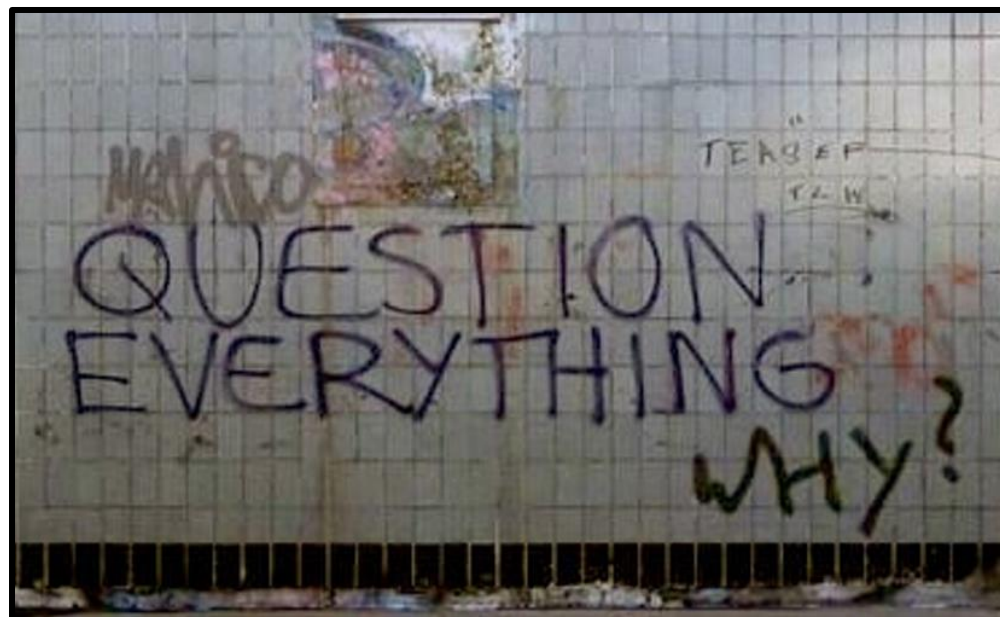
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Grazie per l'attenzione