

EPTRI General Assembly and Scientific Meeting

Warsaw, 28-29 July, 2026

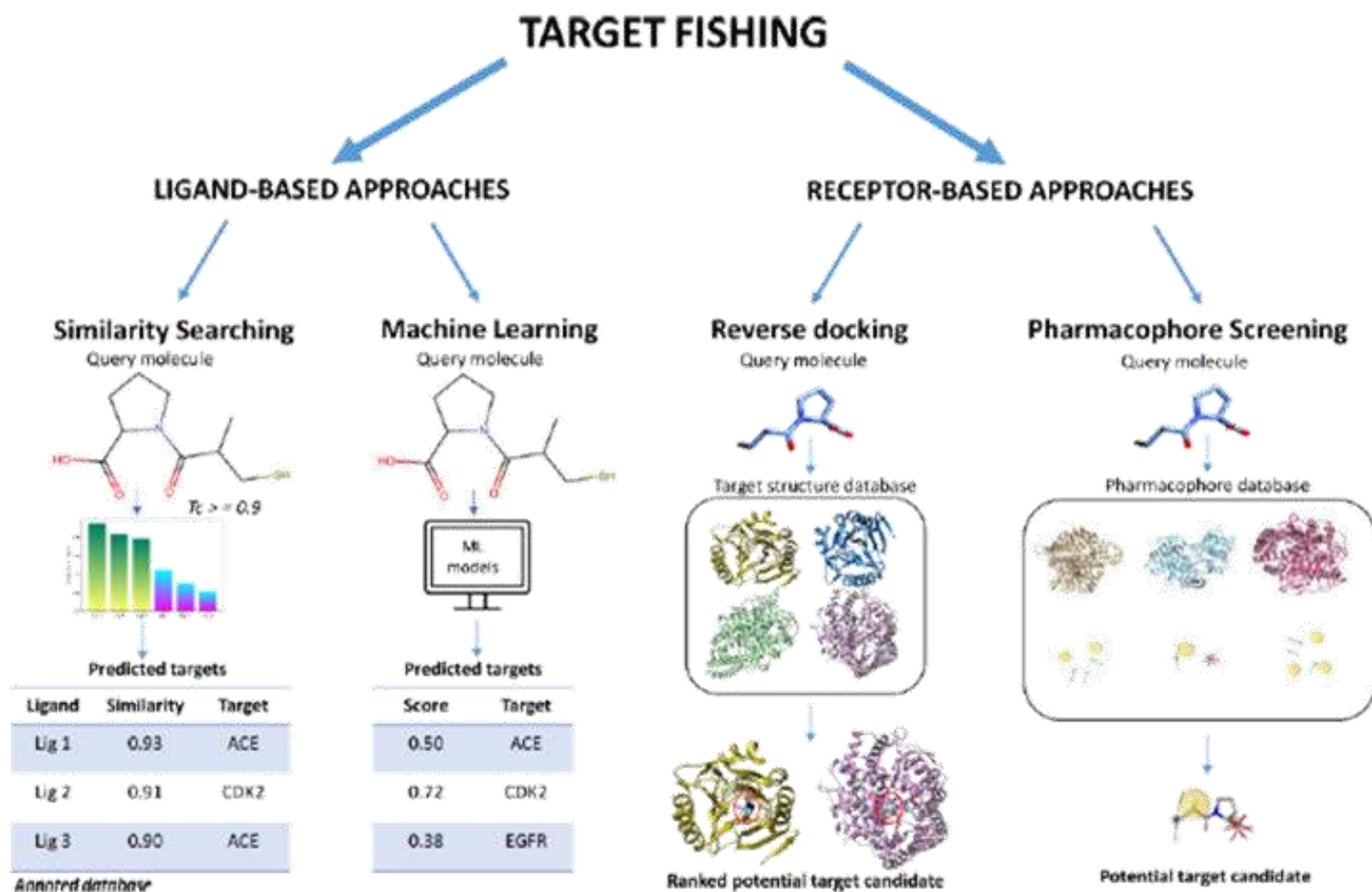
AI-Based Digital Platforms for Early Drug Repurposing Studies in Paediatric Diseases

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Bari, Italy**



**UNIVERSITÀ
DEGLI STUDI DI BARI
ALDO MORO**



In silico target fishing, aiming at identifying possible protein targets for a query molecule, has a wide variety of applications in Drug Discovery.

Figure 1. Overview of the main target fishing approaches.

Prometheus Lab



PLATO

POLYPHARMACOLOGY
PLATFORM PREDICTION

SCAN ME



TIRESIA

TOXICOLOGY INTELLIGENCE
AND REGULATORY
EVALUATIONS FOR
SCIENTIFIC AND INDUSTRY
APPLICATIONS

SCAN ME



CIRCE

CANNABINOID ITERATIVE
REVALUATION FOR
CLASSIFICATION AND
EXPLAINABILITY

SCAN ME



TISBE

TIRESIA IMPROVED BY
STRUCTURE-BASED
EXPLAINABILITY

<< JUST SUBMITTED >>

PLATO (Polypharmacology pLATform predictiOn)

Free and user-friendly web platform for target fishing and bioactivity prediction

PLATO r34 - Polypharmacology pLATform predictiOn

A predictive drug discovery web platform for efficient target fishing and bioactivity profiling of small molecules

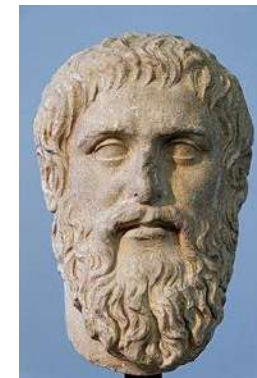
HOME

PREDICTION

DATA

ABOUT US

RELEASE



SCAN ME



Open Access

Communication



PLATO: A Predictive Drug Discovery Web Platform for Efficient Target Fishing and Bioactivity Profiling of Small Molecules

by  Fulvio Ciriaco,  Nicola Gambacorta,  Daniela Trisciuzzi and  Orazio Nicolotti

Int. J. Mol. Sci. **2022**, *23*(9), 5245; <https://doi.org/10.3390/ijms23095245> - 08 May 2022

<https://prometheus.farmacia.uniba.it/plato/>

PLATO r35 - Polypharmacology pLATform predictiOn

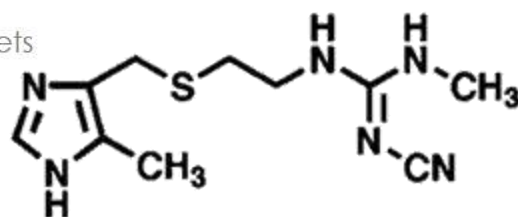
A predictive drug discovery web platform for efficient target fishing and bioactivity profiling of small molecules

HOME PREDICTION DATA ABOUT US RELEASE

release 35 - 2024-12-17 >

- Based on ChEMBL v.35
- Targets: 6297 of which 3008 from Homo sapiens
- Compounds: 634116
- Activities: 1123506
of which :
 - 876553 regarding human targets
 - EC50: 92253
 - IC50: 669945
 - Kd: 31956
 - Ki: 329352

A ligand-based screening of a large collection of bioactive chemicals (small molecules) on **protein targets** implementing two optimized **multifingerprint similarity-based algorithms**.



cimetidine



molecular fingerprint

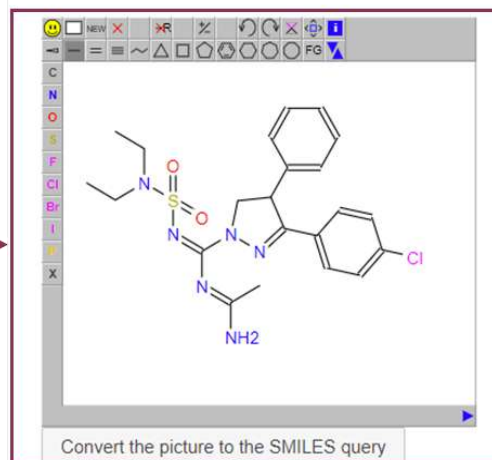
bit string or feature set representations of molecular structure and properties

PLATO - Polypharmacology pLATform predictiOn

A predictive drug discovery web platform for efficient target fishing and bioactivity profiling of small molecules

HOME PREDICTION DATA ABOUT US

Draw your molecule or insert your SMILES and enjoy your prediction!



1a) Draw your molecule using sketch and convert in SMILES format

1b) Input query molecule as SMILES

2) Choose your output format

Choose your output format

- ☐ json data
- ☒ pdf report

Get a response

4) Launch prediction

3) Choose your prediction method

Choose the prediction method

- ☐ Bioactivity profiling
- ☒ Target fishing

Results for query in less than 20 s

Workflow

Outputs

Fish for putative protein drug targets

A)

Input query in
SMILES format

PAGE I

smiles: CCN(CC)S(=O)(=O)N=C(C)/N=C(C)/N(C)C(=O)N1CCCC1
C1C2CC(C)C(C)C2=N1



B)

PAGE II

Target	score	reliable
Cannabinoid CB1 receptor:Homo sapiens	9.89	yes
Cannabinoid CB2 receptor:Homo sapiens	9.89	yes
Cannabinoid CB1 receptor:Mus musculus	9.60	yes
Multidrug resistance-associated protein 4:Homo sapiens	6.95	yes
Cannabinoid CB2 receptor:Mus musculus	6.44	yes
Serotonin 6 (5-HT6) receptor:Homo sapiens	2.09	no
Monoamine oxidase A:Homo sapiens	1.52	no
Monoamine oxidase B:Homo sapiens	1.52	no
Kinesin-like protein 1:Homo sapiens	0.76	no
Dihydrofolate reductase:Plasmodium falciparum K1	0.73	no

List of
predicted
targets

Score
associated
to each
predicted
target (0-13)

Reliability
of each
target
prediction
(YES/NO)

C)

OTHER PAGES

id:	57
should id:	CHEMBL225
name:	Cannabinoid CB1 receptor:Homo sapiens
similarity analysis	
structure	adjcat formalcat rdcat pattern syscat tlcat fp2 pubchem cdlcat graph substructure topological kekulacat
CHEMBL225 Ki:200nM	0.78
CHEMBL225 Ki:150nM	0.85 0.82 0.76 0.66
CHEMBL225 Ki:100nM	0.98 0.91 0.92
CHEMBL225 Ki:50nM	0.88
CHEMBL225 Ki:30nM	0.97 0.97 0.92

Target Information

- Target ChEMBL ID
- Hyperlink to ChEMBL Target Report Card
- Target name

13 Molecular Fingerprints

Most similar ChEMBL compounds
to the query

- Hyperlink to ChEMBL Compound Report Card
- Experimental activity value
- Tanimoto value of the fingerprint

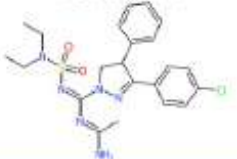
Outputs

Compute bioactivity values for small molecules

A)

Input query in SMILES format

PAGE I



B)

List of predicted targets

PAGE II

Target	IC50	EC50	Ki	Kd	σ_F
Cannabinoid CB1 receptor:Homo sapiens	65.6nM	171nM	104nM	4.68nM	1.3: Ki
Cannabinoid CB2 receptor:Homo sapiens	356nM	50.3nM	91.5nM	7.46nM	1.3: Ki
Cannabinoid CB1 receptor:Mus musculus	127nM	595nM	56.2nM	67.5nM	1.6: Ki
Multidrug resistance-associated protein 4:Homo sapiens	20.0 μ M		6.45 μ M		2: Ki
Cannabinoid CB2 receptor:Mus musculus	620nM	42.4nM	64.1nM		2: Ki
Cannabinoid CB1 receptor:Rattus norvegicus	30.1nM	87.5nM	87.7nM	19.0nM	2.5: IC50
Phosphodiesterase 5A:Homo sapiens	57.3nM	29.1nM	12.5nM	503nM	2.8: IC50
p53-binding protein Mdm2:Homo sapiens	15.5nM	2.93 μ M	154nM	404nM	2.8: IC50

Predicted activity values expressed as IC50, EC50, Ki, and Kd

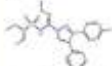
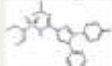
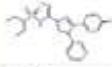
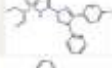
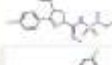
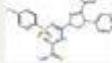
σ_p
the
variance for
the best
predicted
activity type

9

OTHER PAGES

uid: 85
 chemical: C185108219
 name: Cannabidiol CB1 receptor/TrkB agonist

similarity analysis

structure global match	chembl id	r	EC50	EC50	Ki	Kd
		0.12: 0.0	45.6nM	173nM	104nM	4.0nM
	CHEMBL137207	0.001			200nM	3.10nM
	CHEMBL180596	0.001			60.0nM	1.00nM
	CHEMBL198065	0.003			141nM	31.0nM
	CHEMBL196450	0.003			30.0nM	2.51nM
	CHEMBL264390	0.045			117nM	6.51nM
	CHEMBL2133801	0.024	20.0nM			

Target Information

- Target ChEMBL ID
- Hyperlink to ChEMBL Target Report Card
- Target name

Bioactivity prediction results

Most similar ChEMBL compounds
to the query

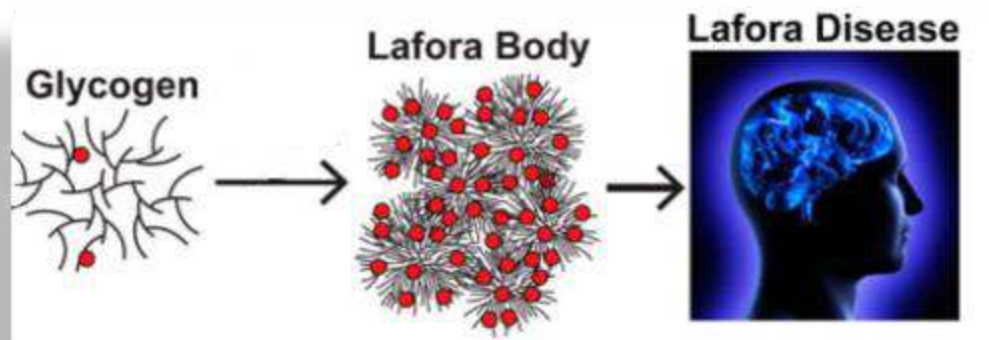
- Hyperlink to ChEMBL Compound Report Card
- τ value expressing biological activity precision
- Experimental activity values expressed as IC_{50} , EC_{50} , K_d and

Lafora disease (LD)

Fatal and ultra-rare **autosomal recessive progressive myoclonus epilepsy**, characterized by a rapid **psychomotor deterioration**

Symptoms typically begin during **late childhood or adolescence**: typically, 8-19 years (peak onset 14–16 years)

The hallmark of the disease is the presence of **insoluble forms of glycogen named Lafora bodies** in the brain and peripheral tissues



LD is caused by **loss-of-function (mutations)** of:

- *EPM2A* encoding for **laforin** (glycan phosphatase)
- *EPM2B* encoding for **malin** (E3 ubiquitin ligase)

In the absence of a functional laforin-malin complex, structurally abnormal glycogen becomes insoluble and accumulates as **Lafora bodies** which drive disease progression in the brain

Clinical progression of Lafora disease

Teenage onset with myoclonic and generalized convulsive seizures

Diminishing response to anti-seizure medication

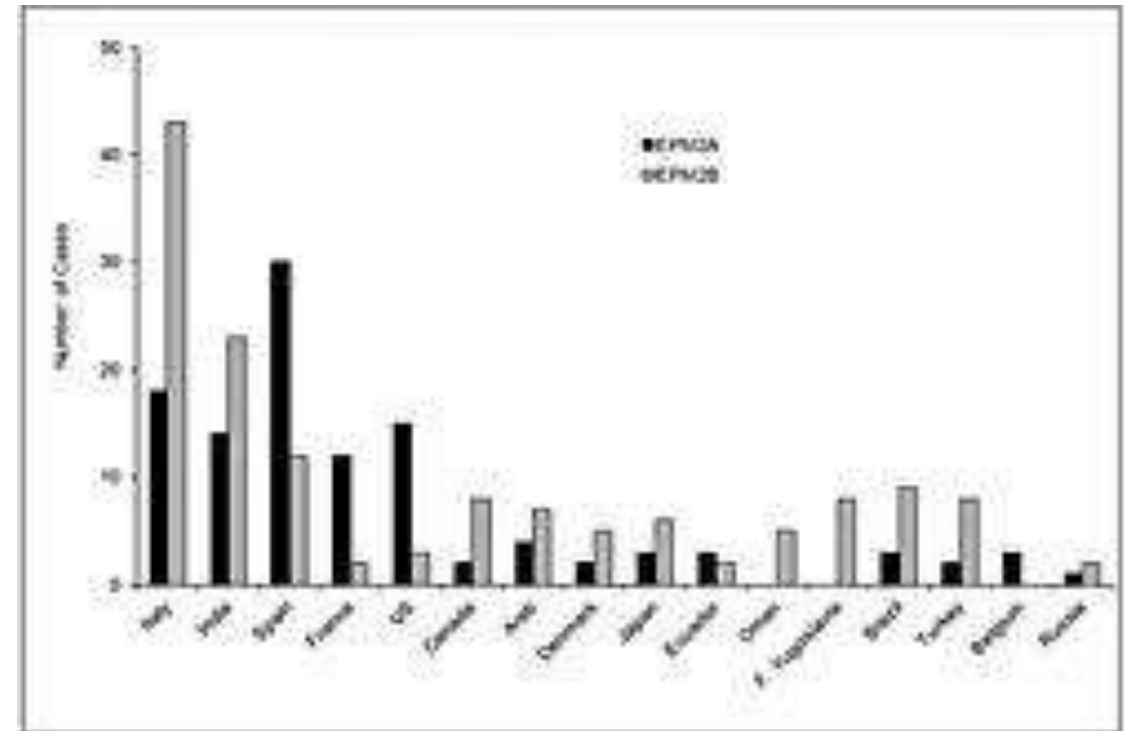
Intractable seizures, near-constant myoclonus, epileptic hallucinations and absence attacks

Dementia and a vegetative state in constant myoclonus

Death within 10 years of symptom onset in status epilepticus

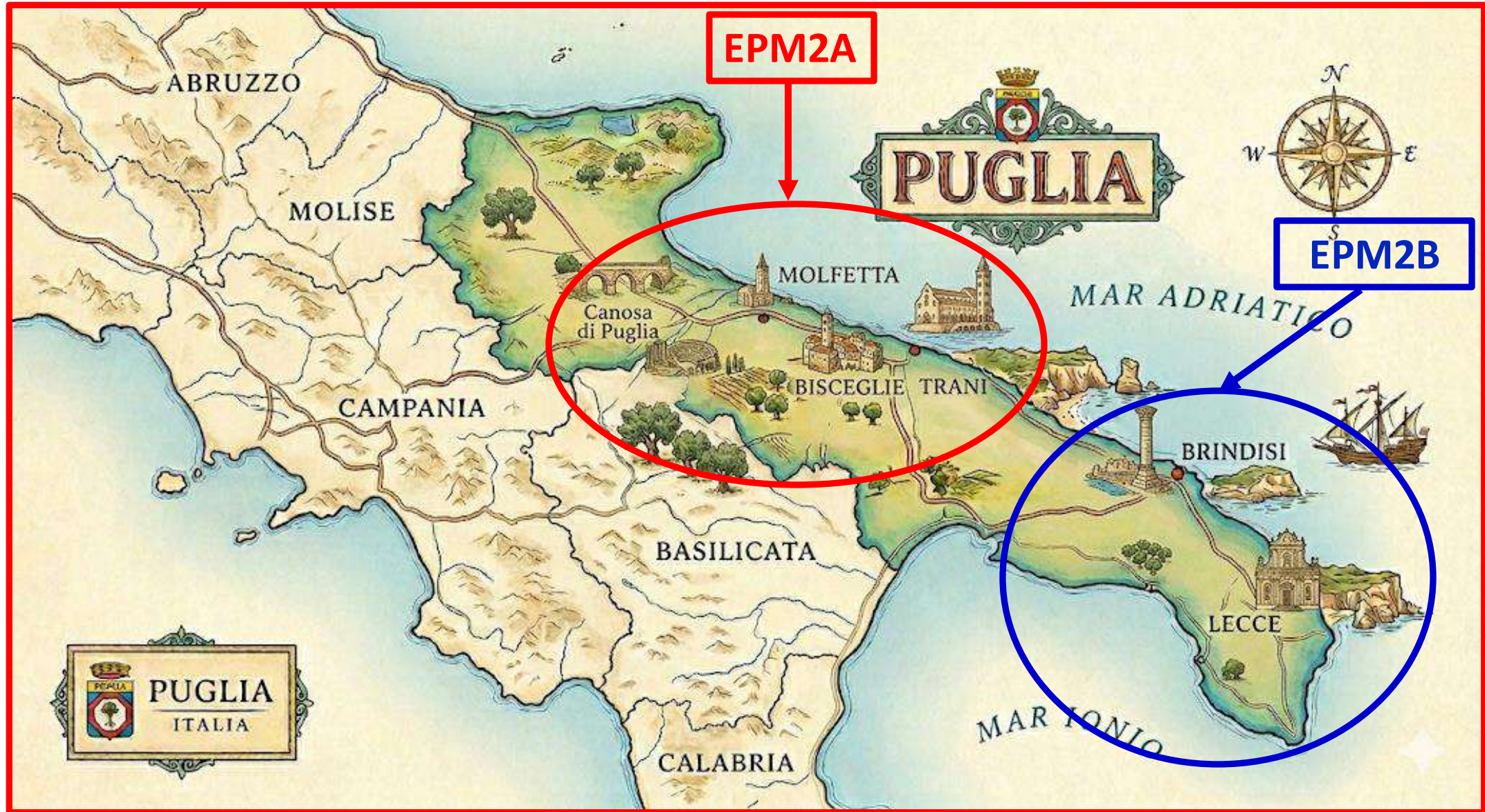
Estimated global prevalence of **fewer than 1 cases per million individuals**

Geographic prevalence: Mediterranean countries (including **Southern Italy/Apulia**), North Africa, the Middle East, and parts of Southern India



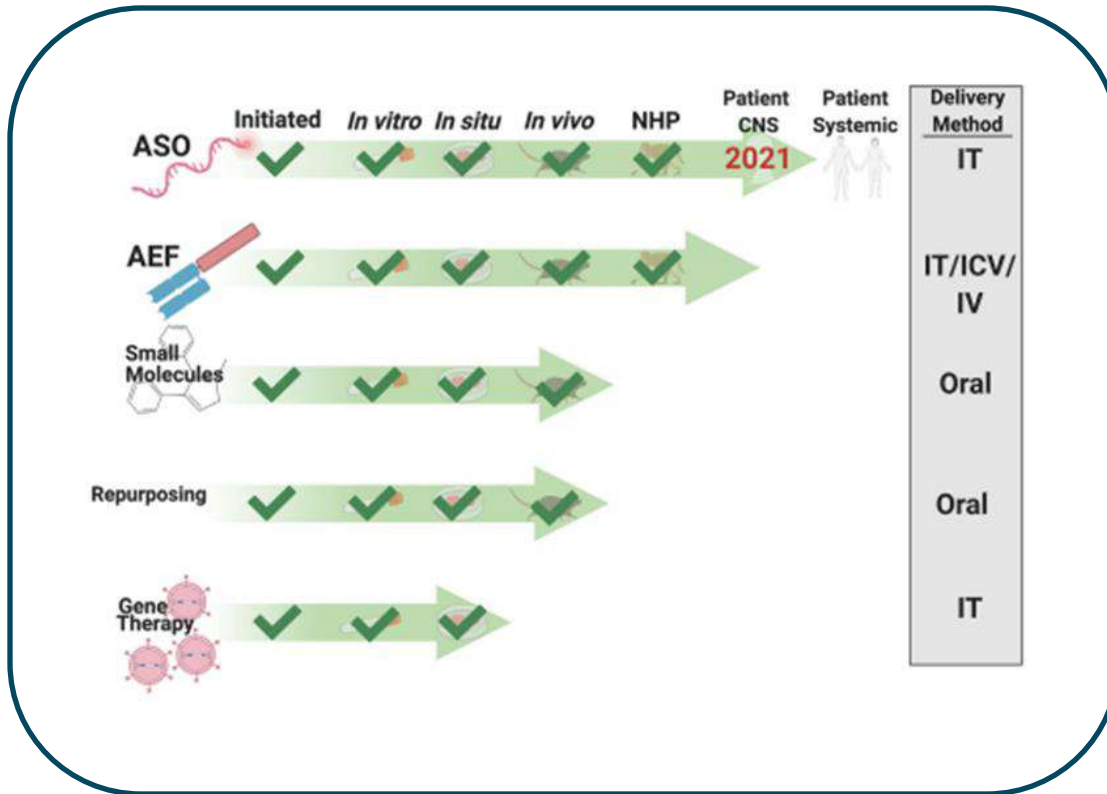
Distribution (2021)

LD in Puglia



There is no effective therapy

Patients are initially treated with anti-seizure medications, to which, after some time, they develop resistance



Metformin, orphan drug since 2016

Drug repurposing in Lafora disease

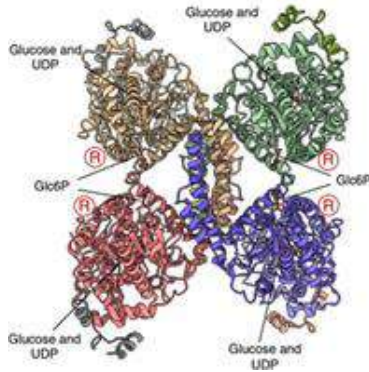
To identify new drug candidates for LD by using *in silico* tools (reverse docking, similarity search) and *in vitro* assays to speed-up (new) therapies

✓ Working hypothesis

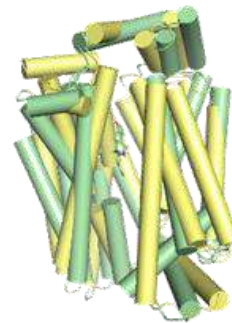
Modulating glucose access into neurons and astrocytes or inhibiting glycogen synthesis, could we reduce glycogen accumulation and Lafora bodies formation?

✓ Drug targets selection

Glycogen synthase (hGYS-1)



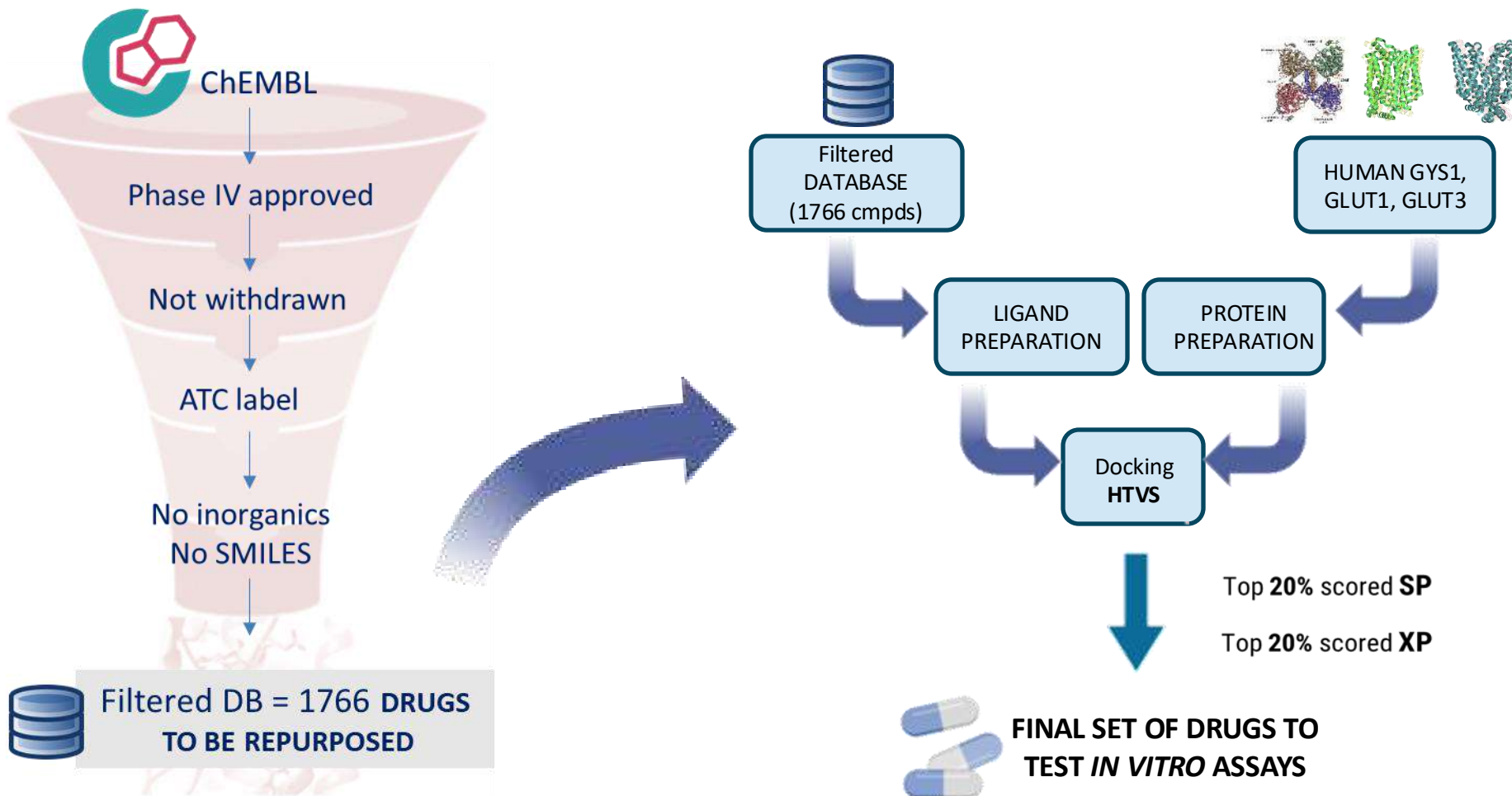
hGLUT1 (astrocytes, BBB)



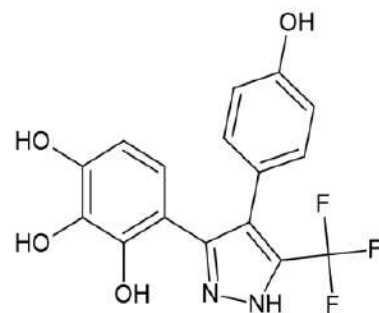
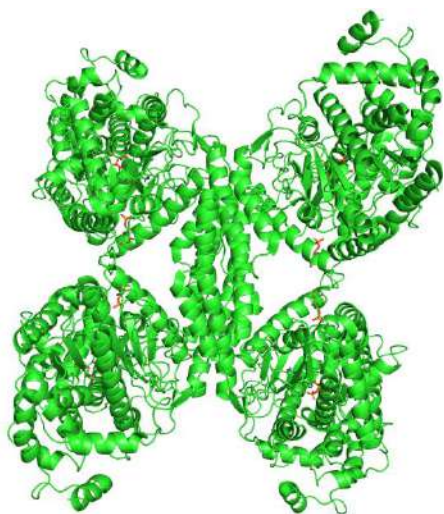
hGLUT3 (neurons)



Docking-based virtual screening of commercial drugs from a filtered ChEMBL database towards hGYS-1 and hGLUT1-3

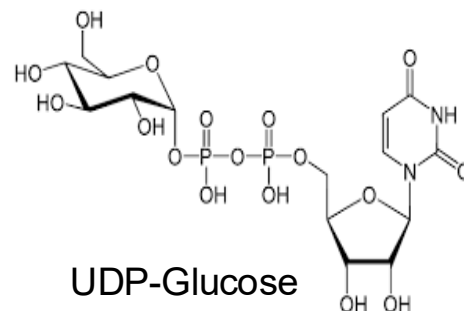


Docking-based virtual screening of filtered ChEMBL database towards hGYS-1



Reference compound

$IC_{50} \sim 2.75 \mu M$

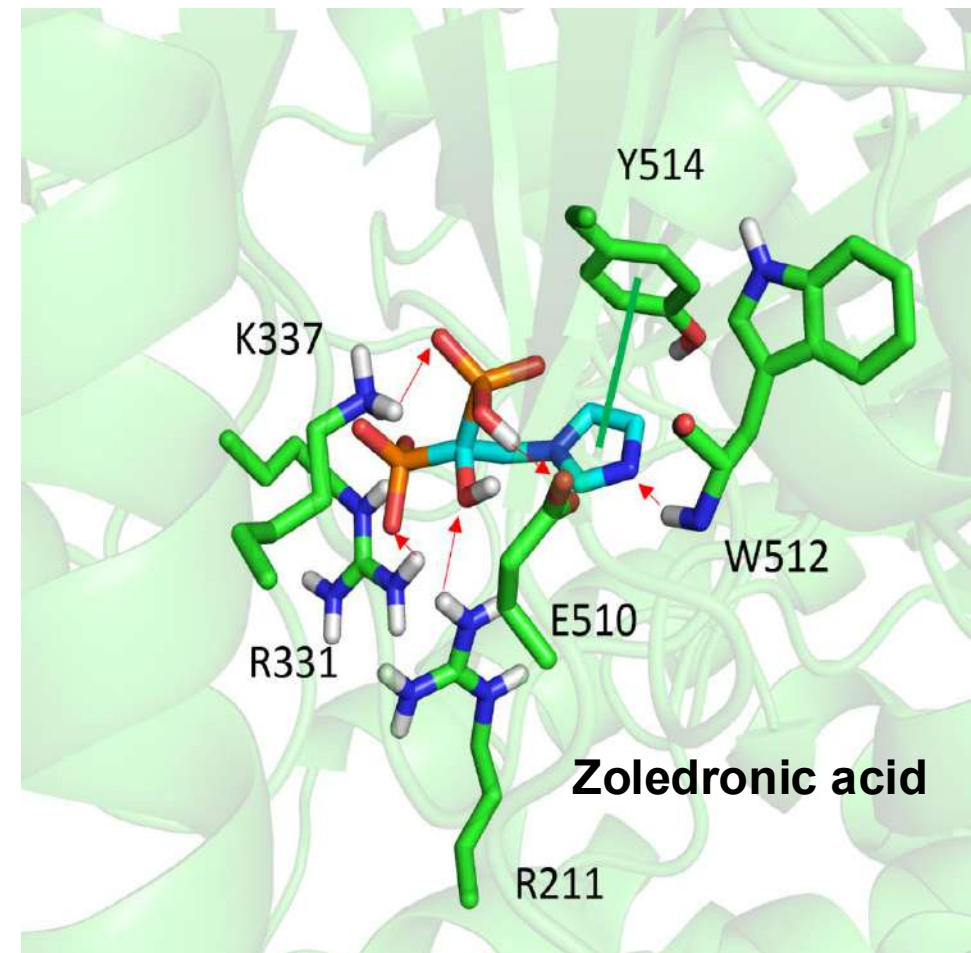


UDP-Glucose

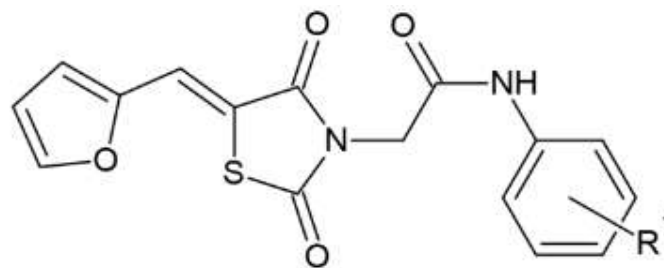
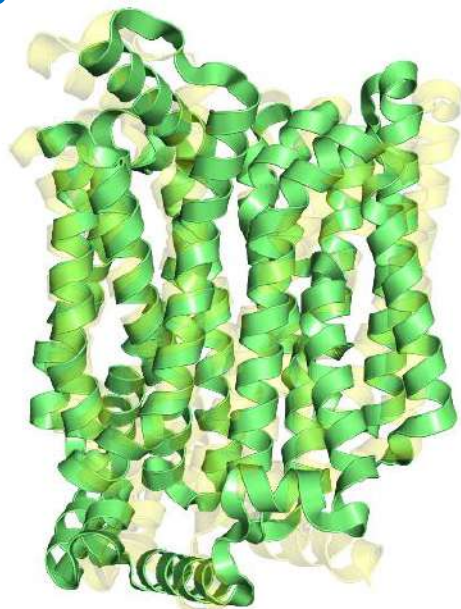
Docking score [UDP-glucose] = -10.118 kcal/mol



Compound	docking score (kcal/mol)	ligand efficiency
ZOLEDRONIC ACID	-10,178	-0,636
RISEDRONIC ACID	-9,636	-0,567
ZOLEDRONIC ACID	-9,583	-0,599
RISEDRONIC ACID	-9,255	-0,544
DICUMAROL	-7,012	-0,28



Docking-based virtual screening of filtered ChEMBL database towards hGLUT1

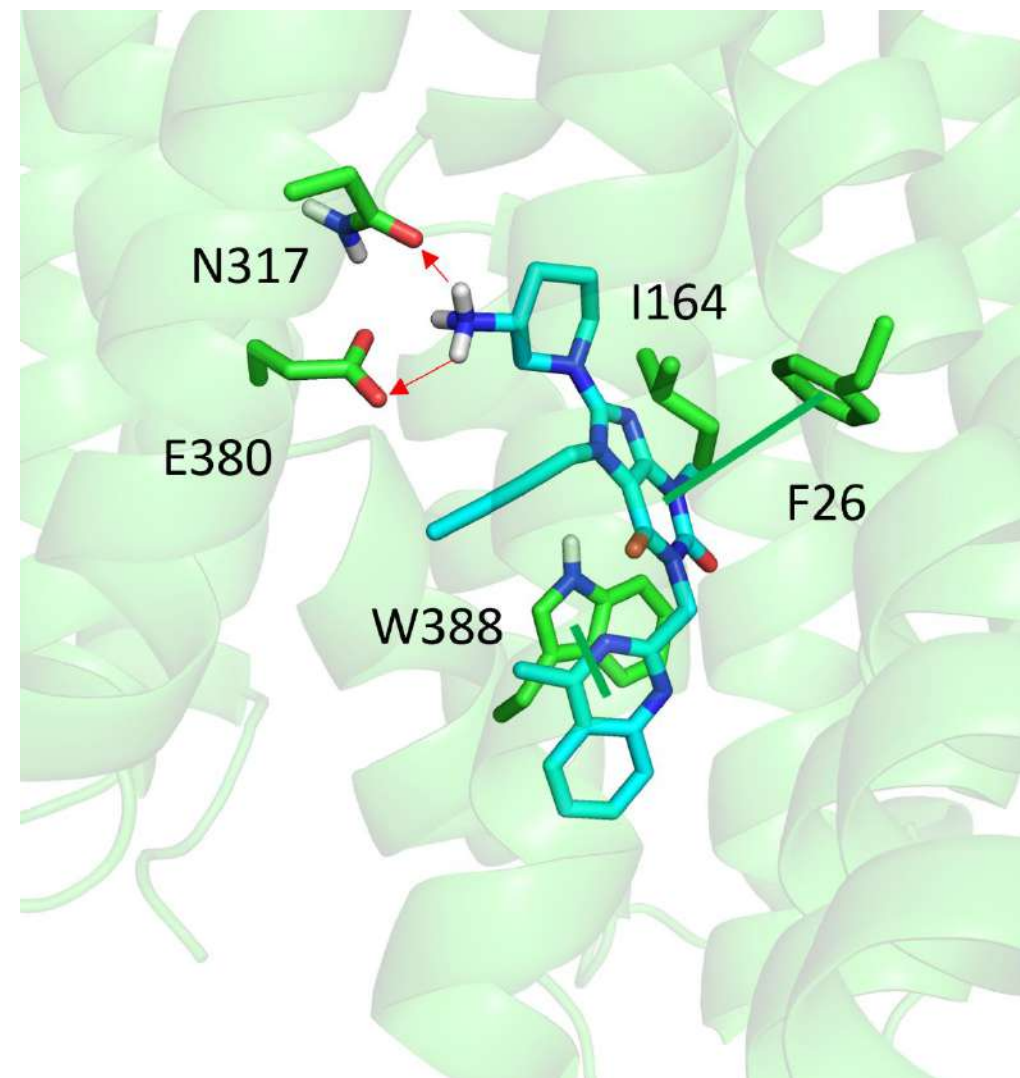


Reference compound
 $IC_{50} \sim 11.4 \mu M$

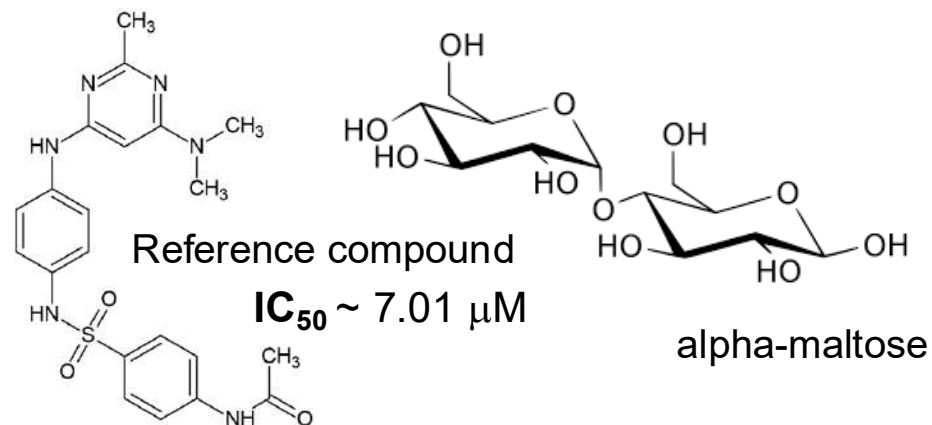
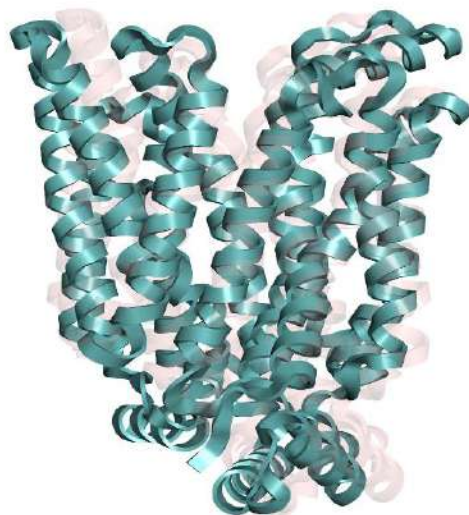
Docking score [Reference compound]= -8.613 kcal/mol



Compound	docking score (kcal/mol)	ligand efficiency
SALMETEROL	-11,134	-0,371
NEBIVOLOL	-10,449	-0,36
LINAGLIPTIN	-8,581	-0,245
THEOPHYLLINE	-8,043	-0,619
ZOLEDRONIC ACID	-6,254	-0,391



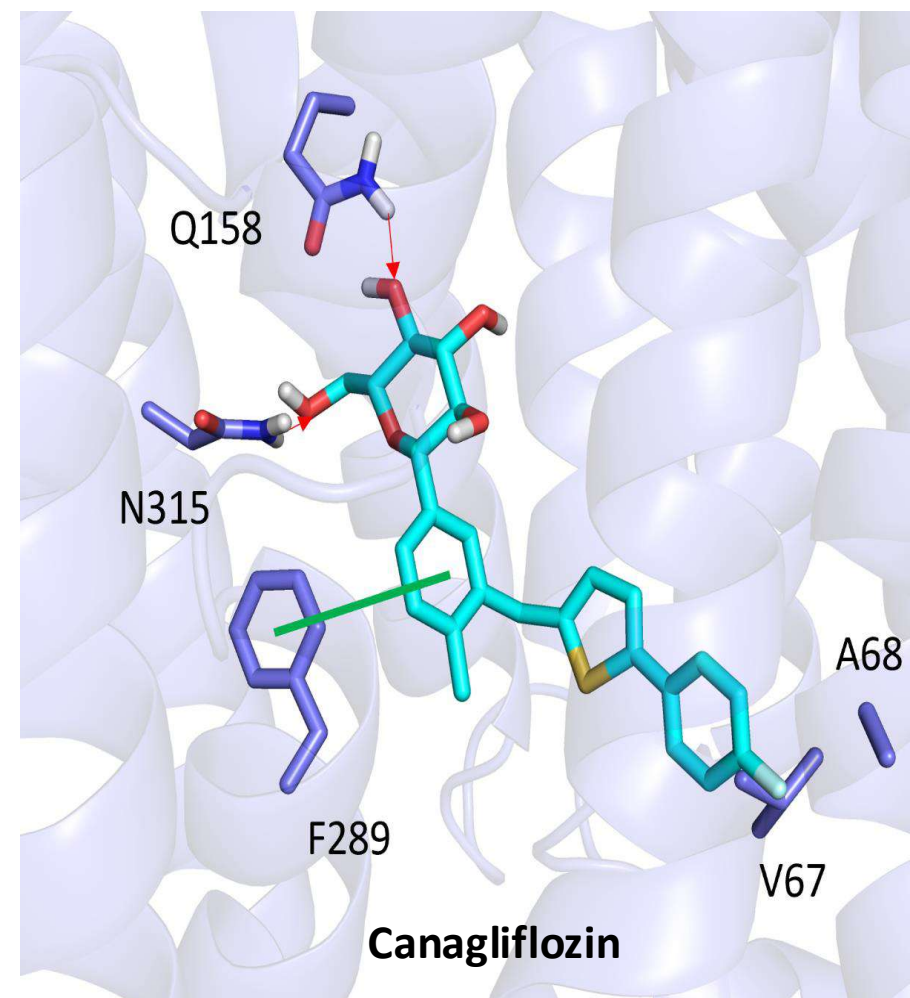
Docking-based virtual screening of filtered ChEMBL database towards hGLUT3



Docking score [alpha-maltose] = -7.573 kcal/mol



Compound	docking score (kcal/mol)	ligand efficiency
CANAGLIFLOZIN	-12,130	-0,391
GEFITINIB	-10,685	-0,345
CARVEDILOL	-10,050	-0,335
LINAGLIPTIN	-9,400	-0,269
SITAGLIPTIN	-9,040	-0,323



List of approved drugs identified from docking studies

1. TUDCA (Item No. 20277II TUDCA, Cayman)
2. Dapagliflozin (CAS: 004CA11574-50, Cayman)
3. Canagliflozin (CAS: 004CA11575-50, Cayman)
4. Empagliflozin (CAS: 004CA17375-100, Cayman)
5. **Cannabidiol**
6. **Acetazolamide**
7. Gefitinib
8. Nebivolol
9. Sitagliptin
10. Bicalutamide
11. **Carbamazepine**
12. **Levetiracetam**
13. **Stiripentol**
14. **Piracetam**
15. **Etosuccimide**
16. Dicumarol (DMSO-9mM)
17. Zoledronic Acid (stock in water, 1mM)

Starting point for *preclinical* studies

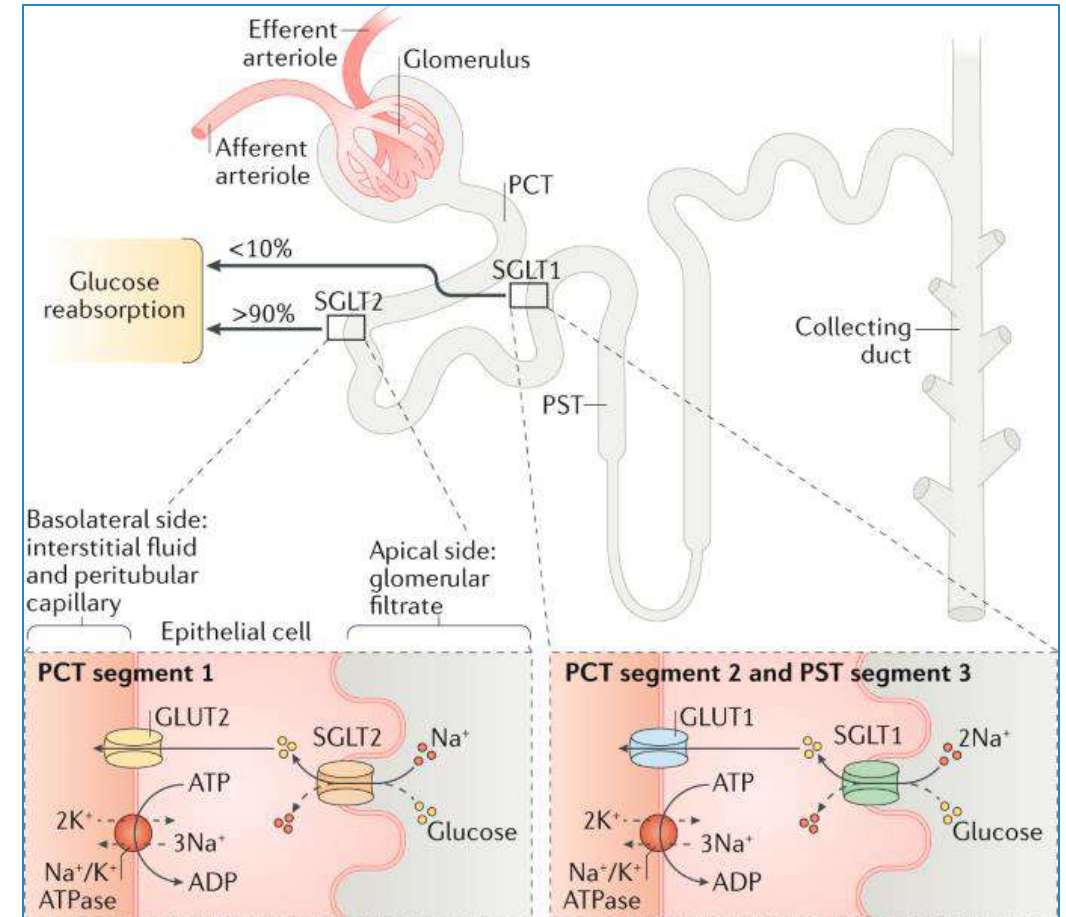


Sodium glucose co-transporters 2 (SGLT2) inhibitors (gliflozins)

In EU, SGLT2i currently authorized to treat DM-2:

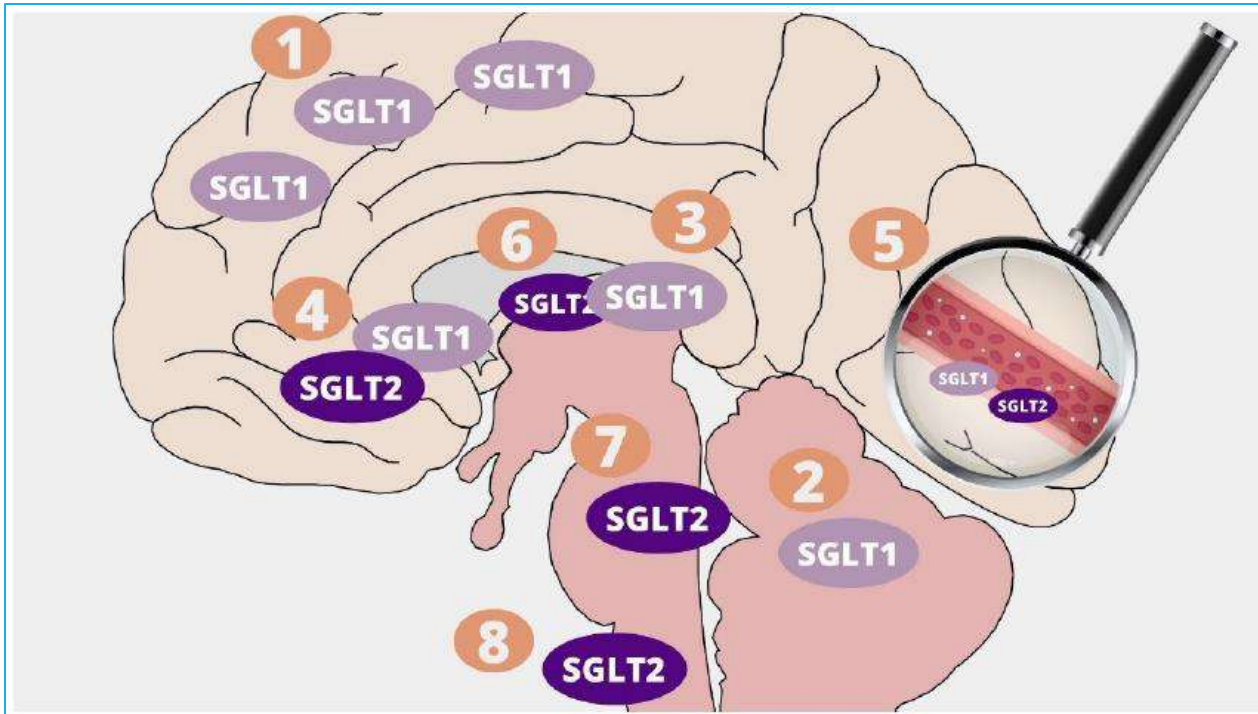
- **dapagliflozin** (*Forxiga* , *Xigduo*)
- **empagliflozin** (*Jardiance*, *Synjardy*, *Glyxambi*)
- **canagliflozin** (*Invokana*, *Vokanamet*)
- **ertugliflozin** (*Steglatro*, *Segluromet*)
- **sotagliflozin** (*Zynquista*)

Mechanism of action: antihyperglycemic drugs, favour glucose excretion from the kidney targeting SGLT2



Hypotheses and evidences supporting the use of gliflozins in LD

Localization of SGLT1/2 in the brain



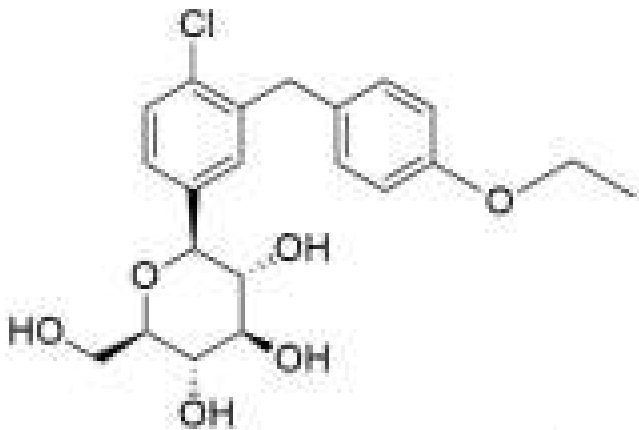
SGLT1 receptor is expressed in brain areas such as hippocampus (CA1, CA3 and the dentate gyrus), cortical and cerebellar neurons, BBB endothelial cells while significant expression of **SGLT2** has been identified in the cerebellum and at BBB endothelial cells.

Pawlos et al., 2021

PLATO (Polypharmacology pLATform predictiOn)

Target fishing

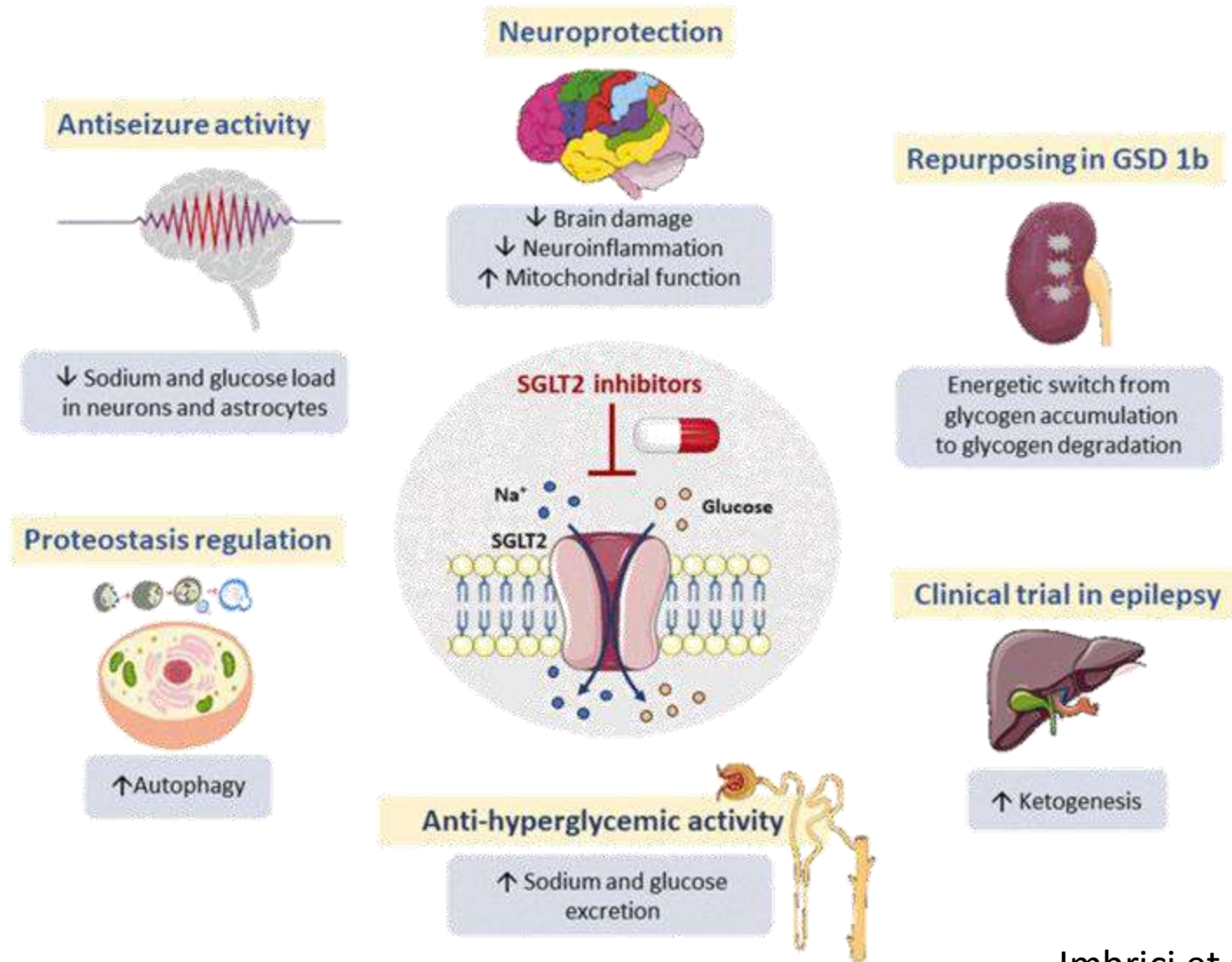
smiles: CCOc3ccc(Cc2cc([C@@H]1O[C@H](CO)[C@@H](O)[C@H](O)[C@H]1O)ccc2Cl)cc3



Dapagliflozin

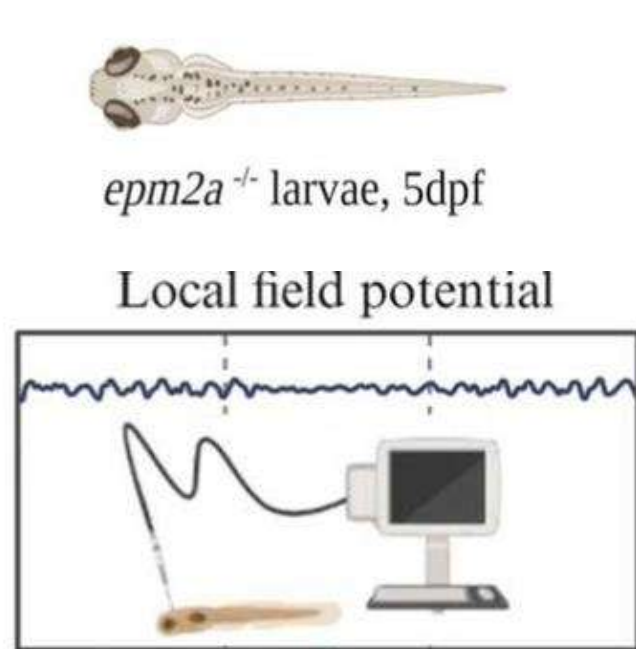
Target	score	reliable
Sodium/glucose cotransporter 2:Rattus norvegicus	13.00	yes
Low affinity sodium-glucose cotransporter:Homo sapiens	13.00	yes
Sodium/glucose cotransporter 2:Homo sapiens	13.00	yes
Sodium/myo-inositol cotransporter 2:Homo sapiens	13.00	yes
Sodium/glucose cotransporter 1:Homo sapiens	13.00	yes
Sodium/glucose cotransporter 2:Mus musculus	10.69	yes
SGLT1 protein:Mus musculus	7.64	yes
Glycogen phosphorylase, muscle form:Oryctolagus cuniculus	5.19	no
Adhesin protein fimH:Escherichia coli K-12	3.70	no
Aldose reductase:Rattus norvegicus	3.65	no
Carbonic anhydrase XII:Homo sapiens	2.88	no
P-selectin:Homo sapiens	2.83	no
Carbonic anhydrase IX:Homo sapiens	2.82	no
Carbonic anhydrase I:Homo sapiens	2.82	no
Leukocyte adhesion molecule-1:Homo sapiens	2.81	no
Tyrosinase:Homo sapiens	2.79	no
Tyrosinase:Agaricus bisporus	2.78	no
Aldose reductase:Homo sapiens	2.76	no
Epoxide hydratase:Homo sapiens	2.44	no
Monoamine oxidase B:Rattus norvegicus	2.34	no
Cytochrome P450 3A4:Homo sapiens	2.30	no
Acetylcholinesterase:Electrophorus electricus	2.28	no
Peroxisome proliferator-activated receptor delta:Homo sapiens	2.27	no
Cyclooxygenase-2:Homo sapiens	2.19	no
Glucose-6-phosphate 1-dehydrogenase:Homo sapiens	2.15	no
Dipeptidyl peptidase IV:Homo sapiens	2.13	no

Literature data supporting the use of gliflozins in LD

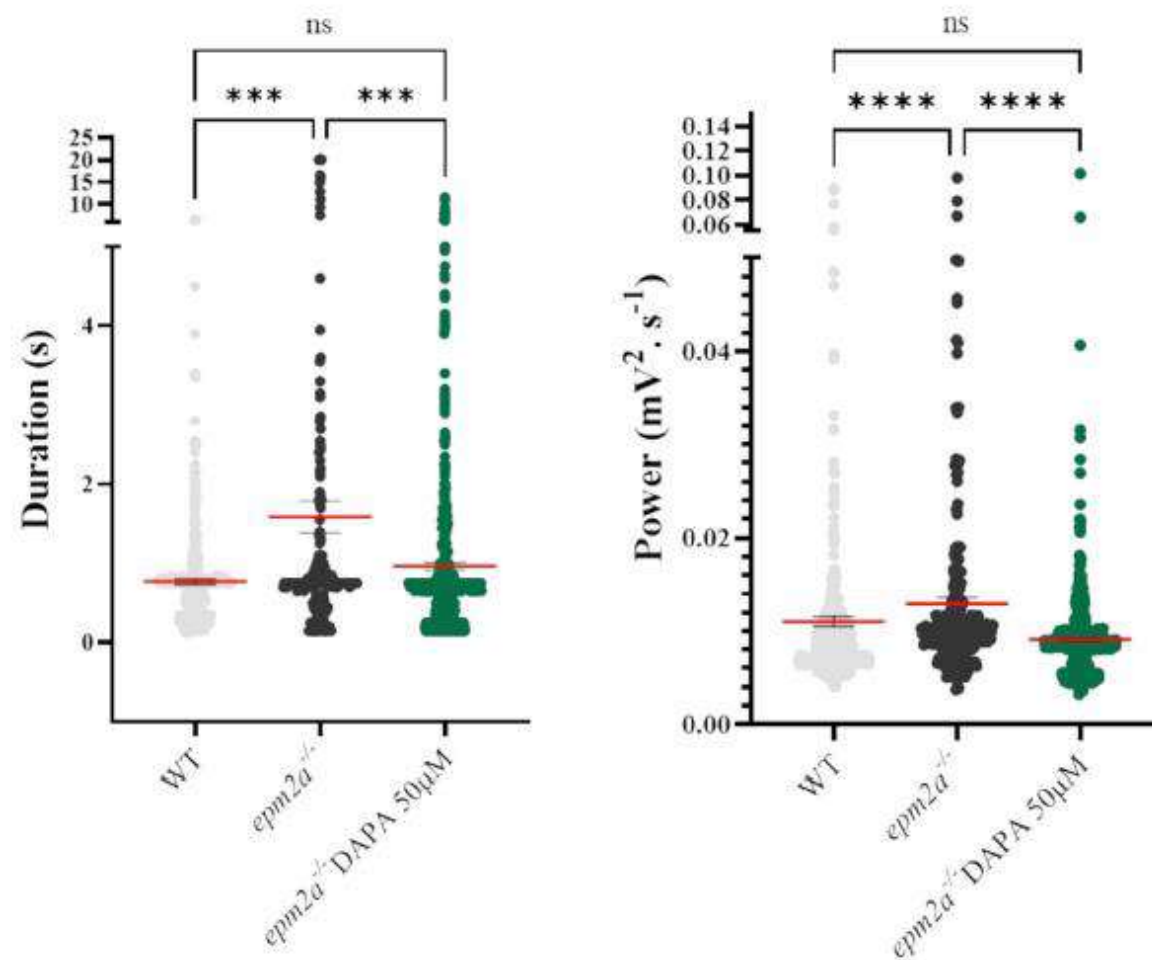


Imbrici et al., *Pharmacol Res* 2024

Dapagliflozin attenuated epileptic phenotype in LD zebrafish model

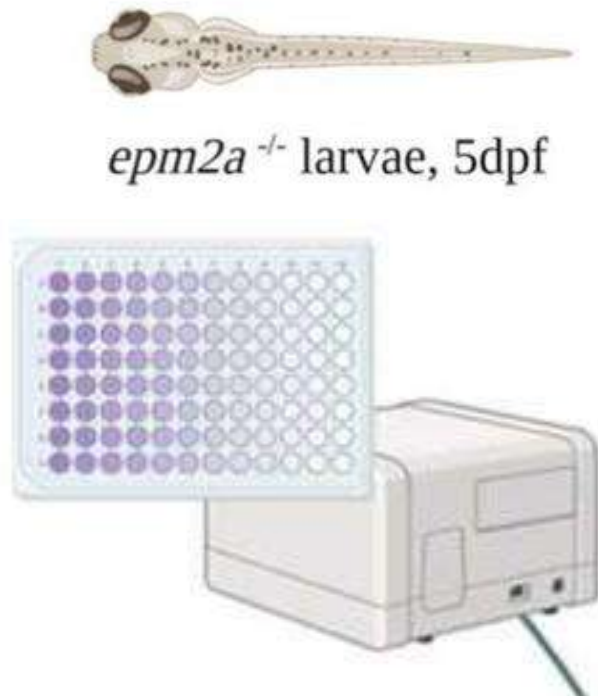


Treatment with DAPA (50 μ M) was able to significantly reduce the **duration and power of seizure-like events**

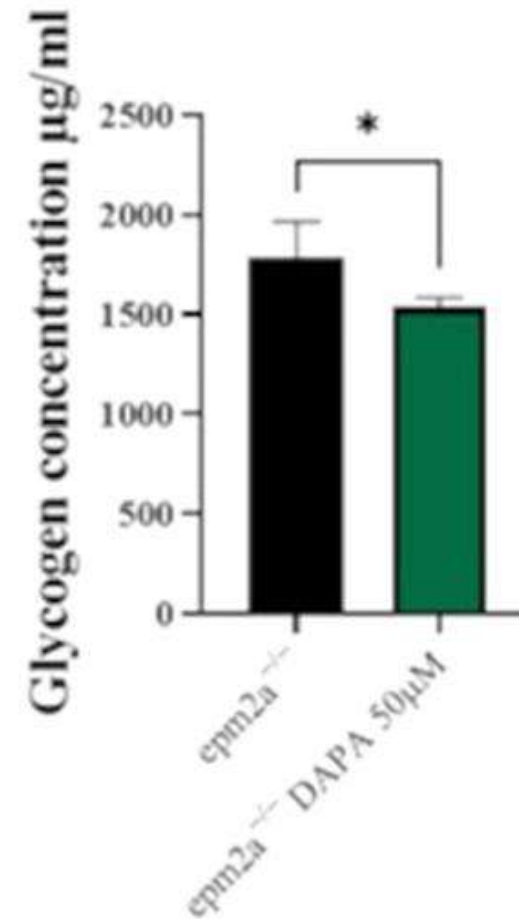


Della Vecchia et al., *Biomed & Pharmacother* 2025

Dapagliflozin slightly reduced glycogen content in LD zebrafish model



Glycogen was quantified in whole LD zebrafish model



Della Vecchia et al., *Biomed & Pharmacother* 2025

ASSESSING THE POTENTIAL OF REPURPOSING EMPAGLIFLOZIN IN LAFORA DISEASE TREATMENT: A STUDY IN PROGRESS

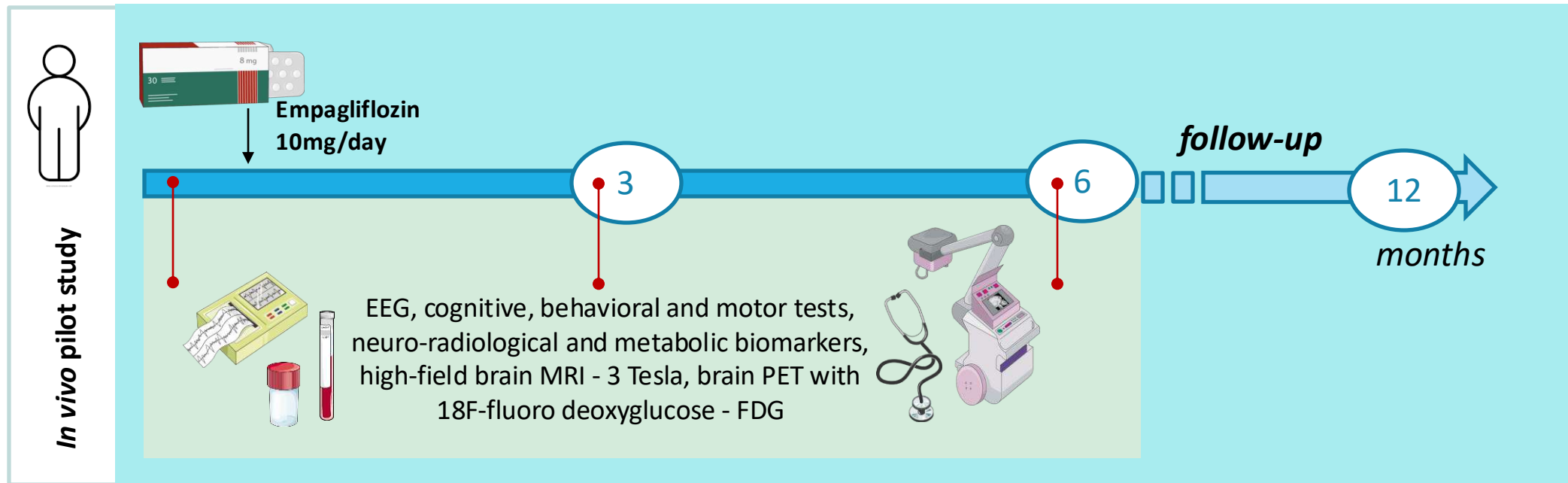
Giuseppe d'Orsi¹, Paola Imbrici², Daniela Trisciuzzi², Nicola Gambacorta^{2,3}, Maria Teresa Di Claudio¹, Cosimo D. Altomare², Antonella Liantonio², Francesca Bisulli⁴, Massimo Carella³ and DEFEAT-LD study group

¹Neurology Unit, Fondazione IRCCS Casa Sollievo della Sofferenza, San Giovanni Rotondo (Fg), Italy.

²Department of Pharmacy - Pharmaceutical Sciences, University of Bari Aldo Moro, Bari, Italy

³Fondazione IRCCS Casa Sollievo della Sofferenza, San Giovanni Rotondo (Fg), Italy

⁴IRCCS Istituto delle Scienze Neurologiche di Bologna, Dep of Biomedical and Neuromotor Sciences, Bologna, Italy



Medicinal Chemistry

Marco Catto
Modesto de Candia
Gabriella la Spada
Caterina Deruvo
Alessandro Gadaleta

Computational Chemistry

Orazio Nicolotti
Nicola Amoroso
Fulvio Ciriaco
Daniela Trisciuzzi
Nicola Gambacorta

Pharmacology

Antonella Liantonio
Paola Imbrici
Annamaria De Luca



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Division of Medical Genetics

Massimo Carella

Orazio Palumbo



Thanks for
your attention