



# Screening platform of Marseille-Luminy

# Biochemical assays and technical developments

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# The screening platform of Marseille-Luminy (PCML)







**Databases** 

**Chemical Libraries** 

Radioactive or fluorescent screening
HTRF

#### **Robotics**







Scintillation

Fluorescence



Results

Data analysis
- % inhibition
- IC<sub>50</sub>

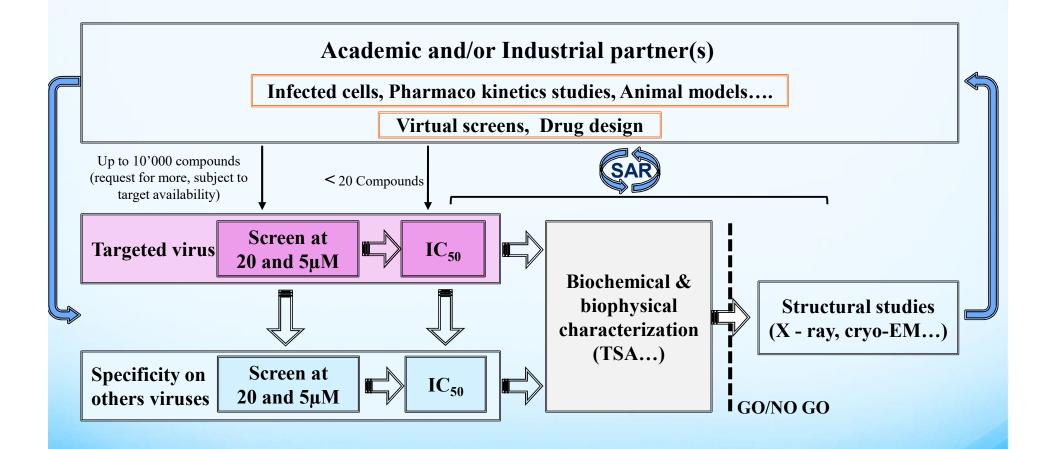
### **PCML Library in 2025**

About 76,000 molecules from various origins are in stock:

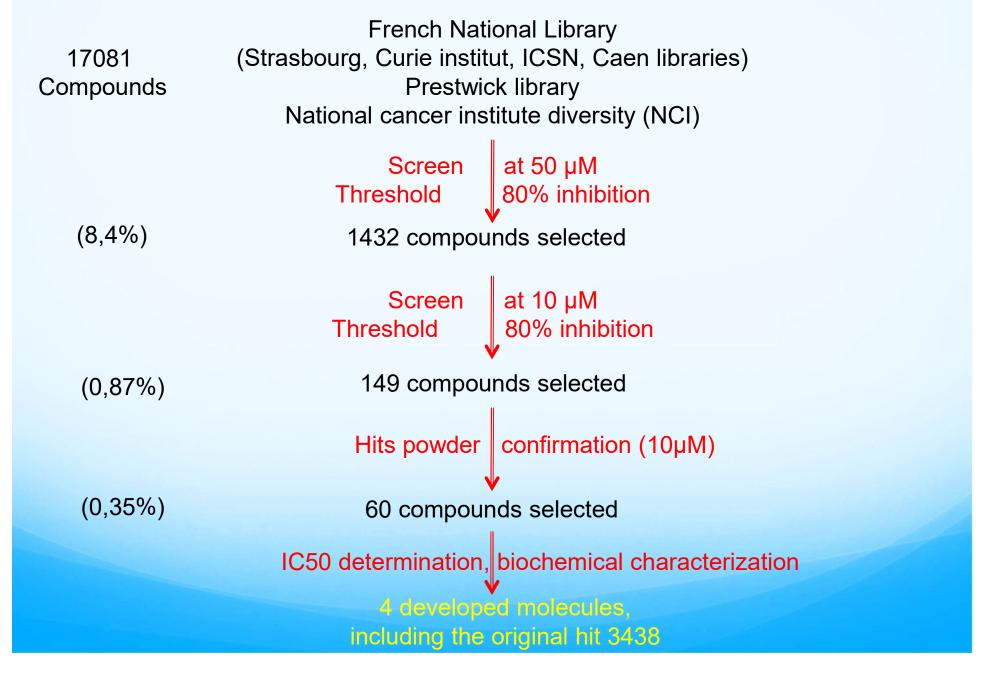
- National cancer institute diversity library (NCI) (2000)
- Chembridge<sup>™</sup> "Diversity Set" library (30,000)
- The "Chimiothèque Nationale Essentielle" (CNE), a representative subset of the Chimiothèque Nationale" (1040)
- The "Chimiothèque Nationale" (21,000) divided into :
  - Strasbourg Library (5000)
  - Curie Institute Library (8000), chemicals (3000) and natural extracts (5000) library, Gif-sur-Yvette ICSN
  - Caen Library (400)
- Active sightTM fragment based library (360)
- Compounds developed in-house by our chemists team (400) or molecules from collaborative studies and commercial providers (>2000) are regularly added to this base.
- Libraries focus on the inhibition of protein-protein or protein-peptide interactions: 2P2I<sub>3D</sub> (1664), Life chemical rule of four (4300), subset ChemDiv Eccentric (515), PPICHem (10 314)
- Prestwick library (2240): chemical (1520), natural (320), pyridazine (400)



#### **Workflow on PCML**



# Screening results on Dengue polymerase



### Production/Purification of enzymes of interest

#### 1) Production

- Bacterial transformation (E.coli)
- pre-cultures and cultures (1 to 10L /protein)



#### 2) Purification

- Bacterial lysis
- 1° step of purification: Affinity chromatography (Cobalt or Nickel) using histidin tag
- 2° step of purification : chromatography by ionic exchange ou size exclusion
- Appropriate storage according stability and future use of the protein
- Yield : from 0,3 mg to 2 mg/L

# **Available proteins on PCML**

18 NS5 from Orthoflavivirus



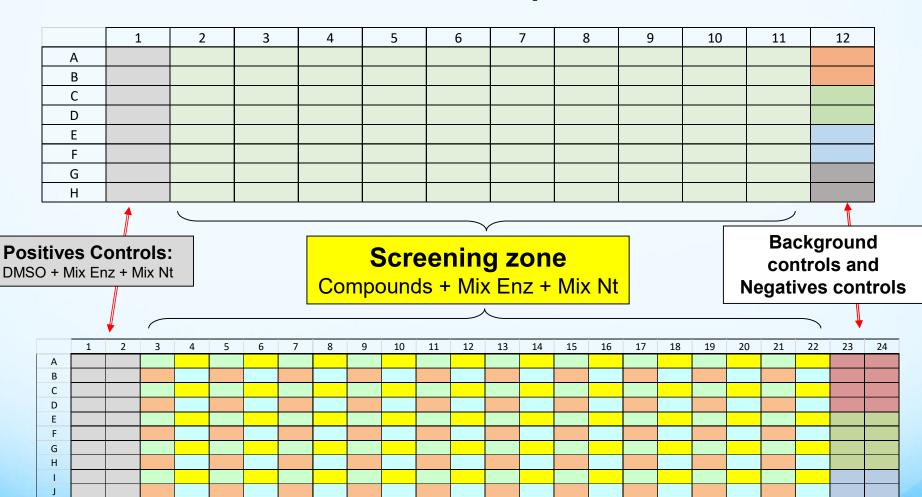
#### **SARS** replication complex:

- nsp12 Cov-1 and Cov-2
- nsp8 Cov-1 and Cov-2
- nsp8L7 Cov-1 and Cov-2
  - nsp7 Cov-1 and Cov-2
    - Nsp13 Cov-2

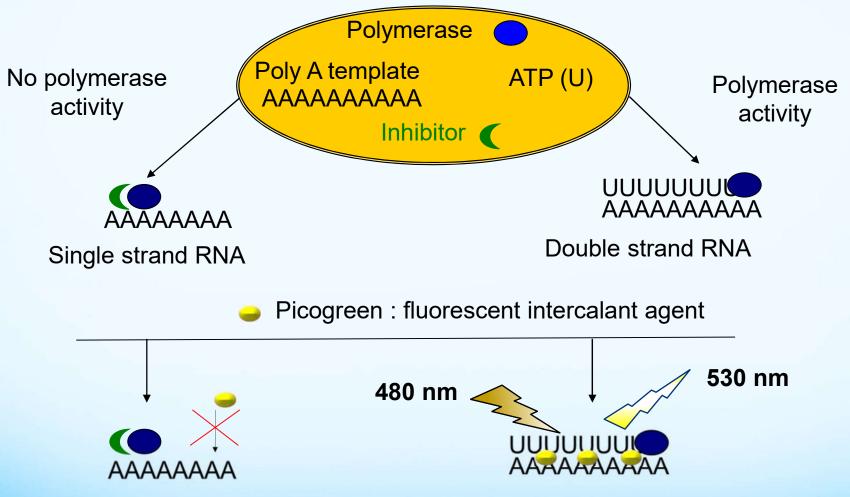
15 Polymerases from Orthoflavivirus

6 Methyltransferases from Orthoflavivirus

# Experimental screening plate pattern 96 and 384 wells plates



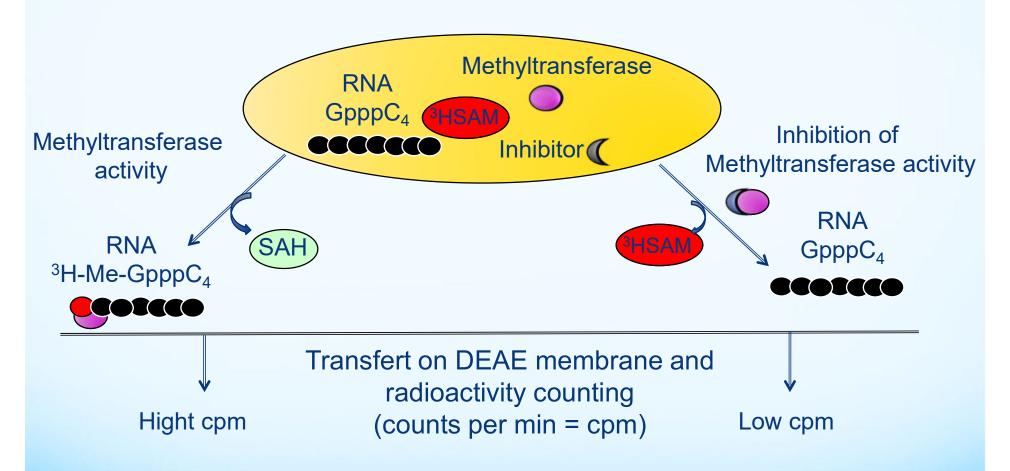
#### A screening polymerase assay based on fluorescence



Detection and quantification (RFU) with a Tecan Safire<sup>2</sup> spectrofluorometer % inhibition – IC50



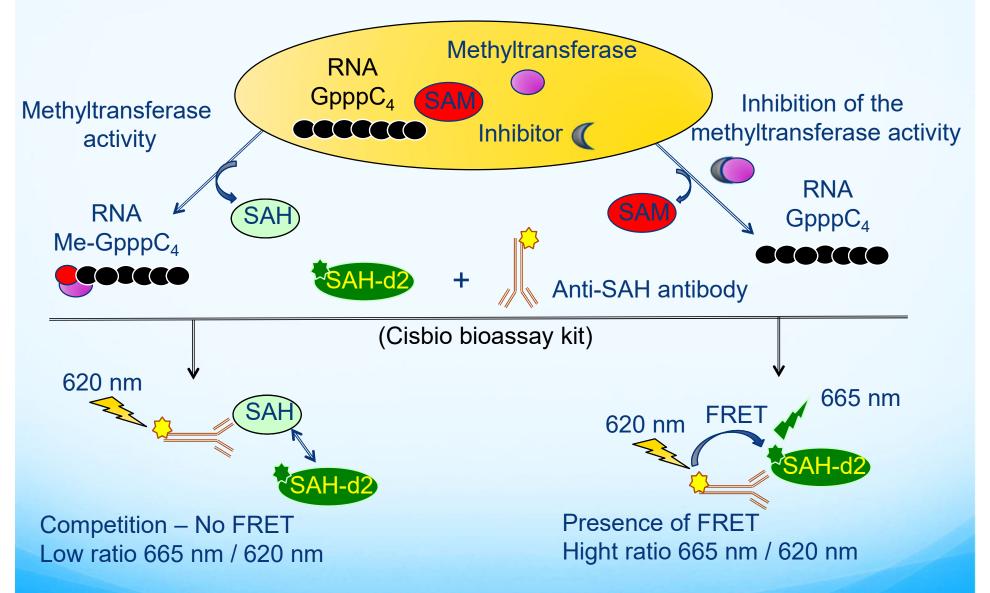
# Determination of methyltransferase activity on radioactive assay



#### <u>Using</u>:

- Enzymes stocks validation
- Inhibitor potency evaluation of compounds (screens, IC50...)

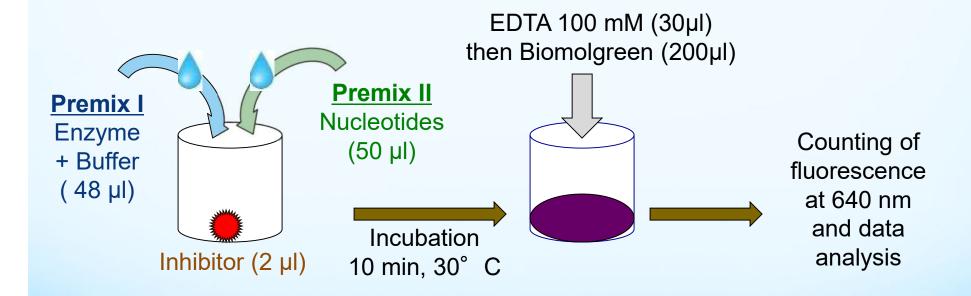
#### Determination of methyltransferase activity on HTRF assay



#### **Using:**

- Inhibitor potency evaluation of compounds (screens, IC50)

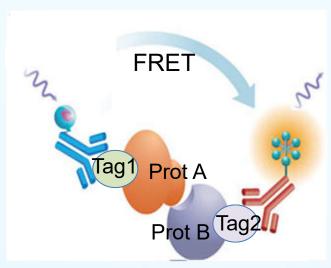
#### **Determination of ATPase activity on fluorescent assay**



#### <u>Using:</u>

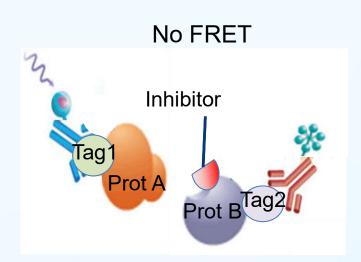
- Inhibitor potency evaluation of compounds (IC50)
- Available on Dengue 3 NS3 and Helicase domain

#### Protein-Protein interactions: screening method by HTRF



Absence of inhibitor

➤ HTRF signal



Presence of inhibitor

➤ no HTRF signal

- Robotized assay (384-wells)
- Avalaible on Dengue NS3 NS5MTase
- Screening on PPICHem library

### Thermalshift assay (TSA)

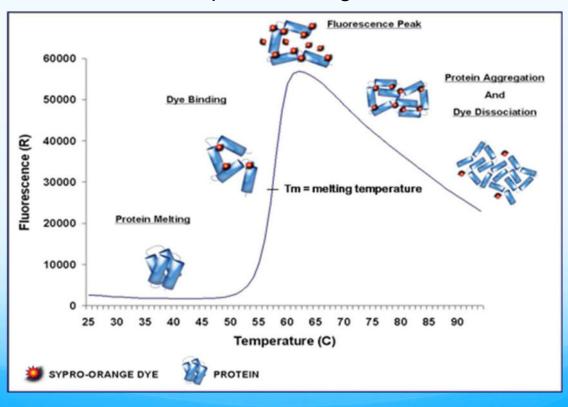
- Determination of the protein of interest Tm (melting temperature) by using a fluorescent probe (Sypro-orange)
- Tm measurement with compound and determination of the variation (ΔTm)

 $\Delta Tm > 0$ : the compound stabilises the protein

 $\Delta$ Tm <0 : the compound destabilises the protein

-> validation of the compound/protein interaction

-> help to cristallogenesis



### Characterisation of mode of action: gel based assay

Specific FAM labelled Dengue Primer/Template

Enzyme: NS5,RdRp or nsp



Inhibitor /





+ NTPs

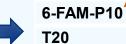


AGU UGU UAG UCU ACG UGG AC

**T20** UCA ACA AUC AGA UGC ACC UG



+ NTPs



AGU UGU UAG U

UCA ACA AUC AGA UGC ACC UG

2 orders of addition availables

Enzyme + P/T 10min 30°C

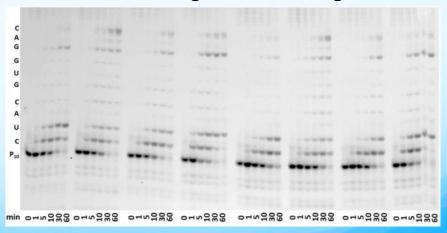
NTPs and Inhibitor X min 30°C

Kinetic (0/1/5/10/30/60min) 30°C

Enzyme + P/T 10min 30°C + Inhibitor X min 30°C + **NTPs** 

**Kinetic** (0/1/5/10/30/60min) 30°C

Migration on Urea gel



D1 NS5 D2 NS5 D3NS5 D4 NS5 D1 NS5 D2 NS5 D3NS5 D4 NS5

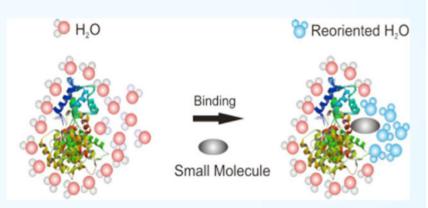
**DMSO** 10 µM Inhibitor

Reading and quantification of the enzymatic activity (Typhoon system)

## Affinity constant (Kd) determination by thermophoresis

#### Principle:

Following of water molecules reorientation in a temperature gradient after binding a ligand to a fluorescent labeled substrate



Substrate: Dengue Polymerase or Dengue NS5 Ligand: Ions, Nucleotides, VHHs or Inhibitors

- 1) Development of the assay:
- Fluorescent labelling conditions
  - Choice of capillary
- Buffer composition and protein concentration
  - Stability and reproductibility

