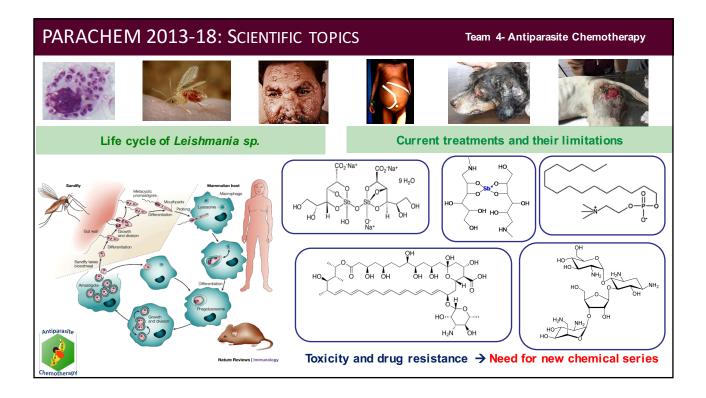
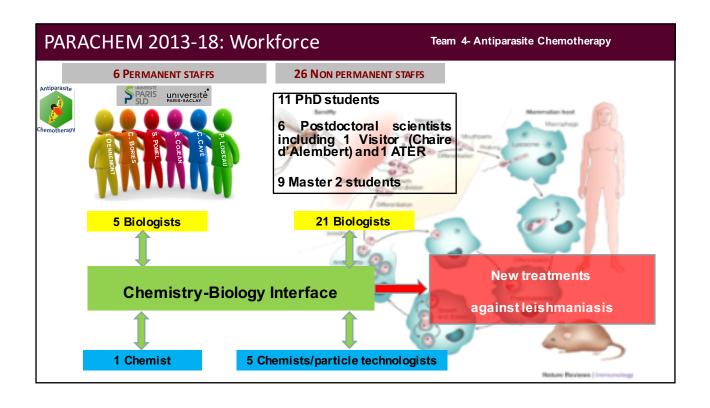
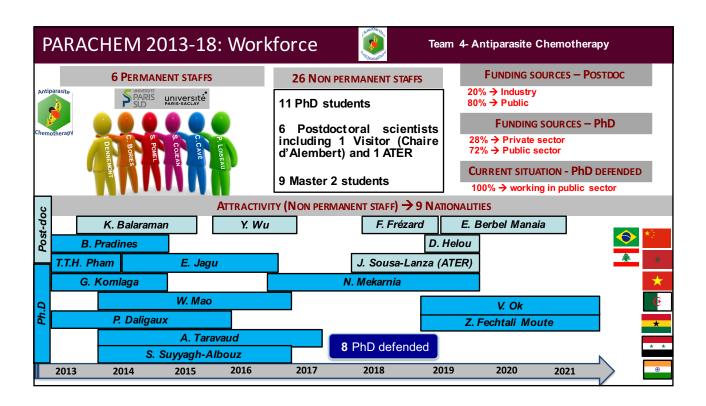
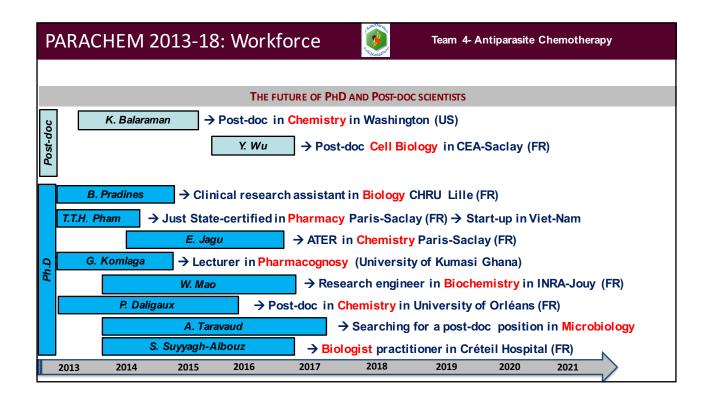
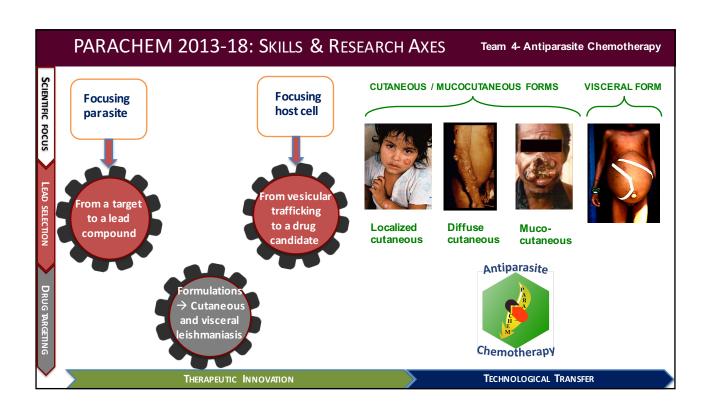
# Presentation of the scientific activities and upcoming research projects Antiparasite Chemotherapy





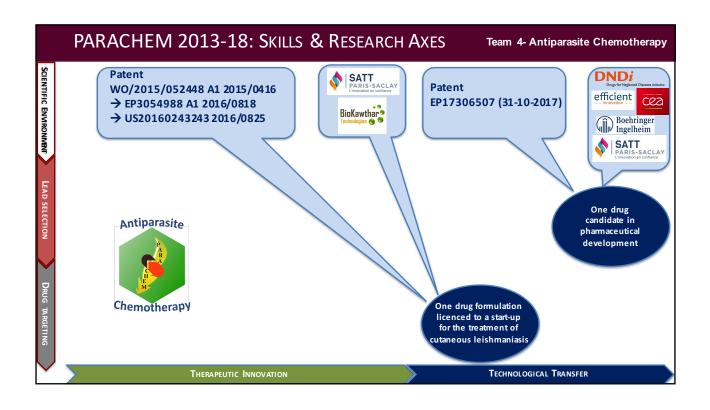


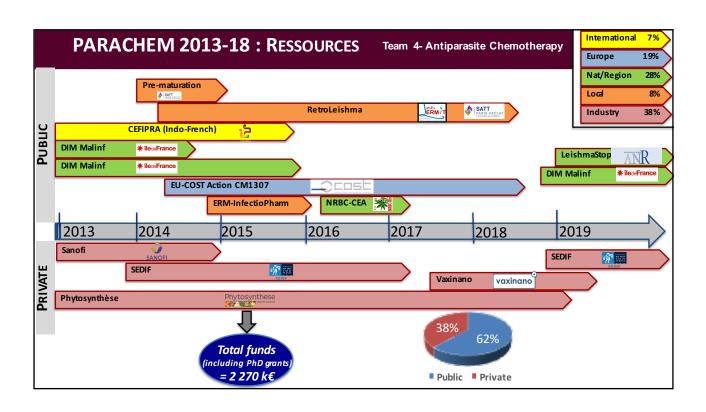


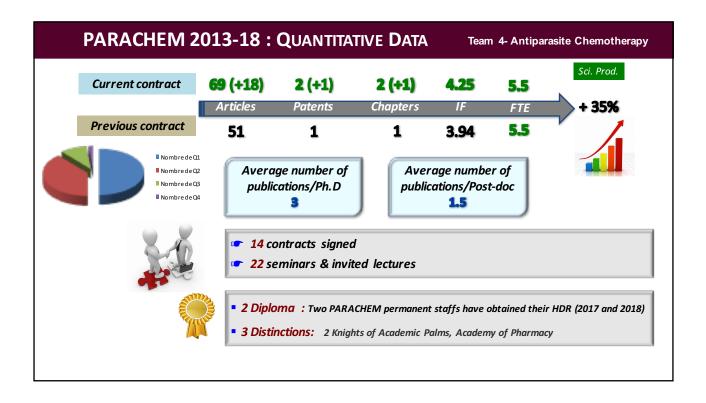


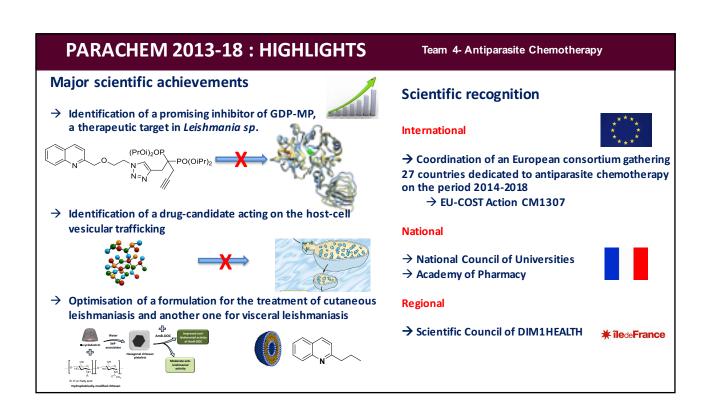


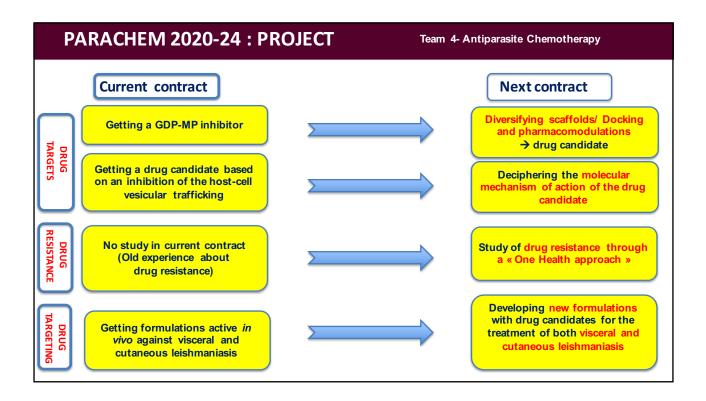


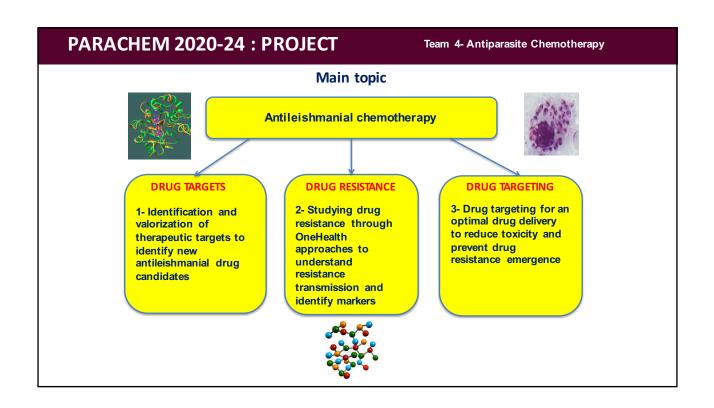


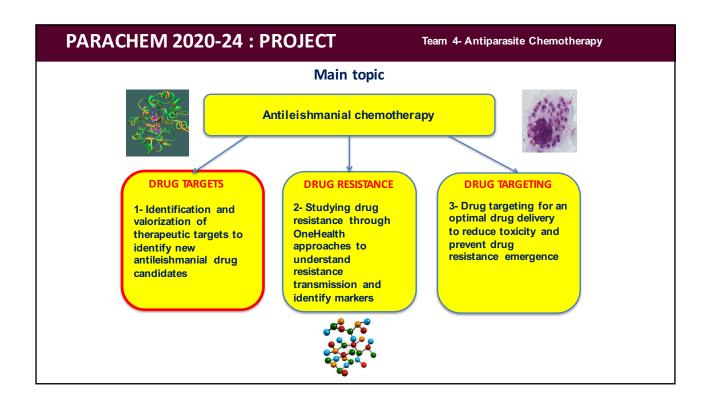


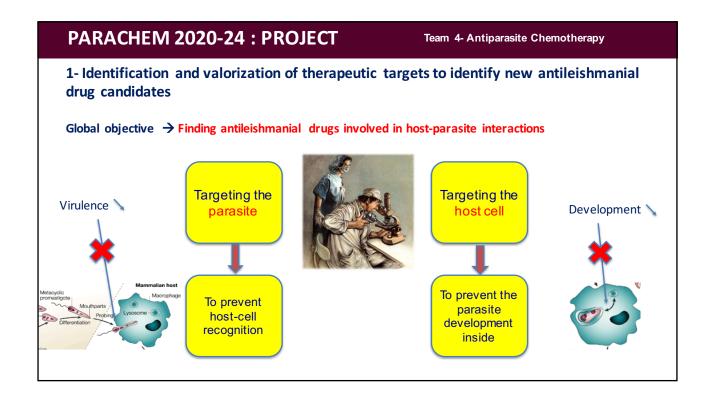


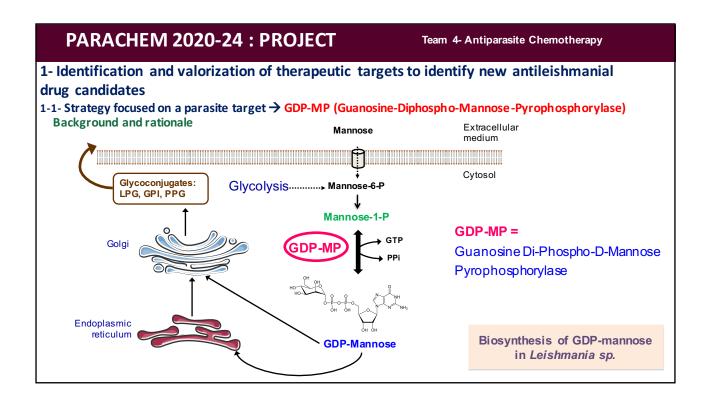


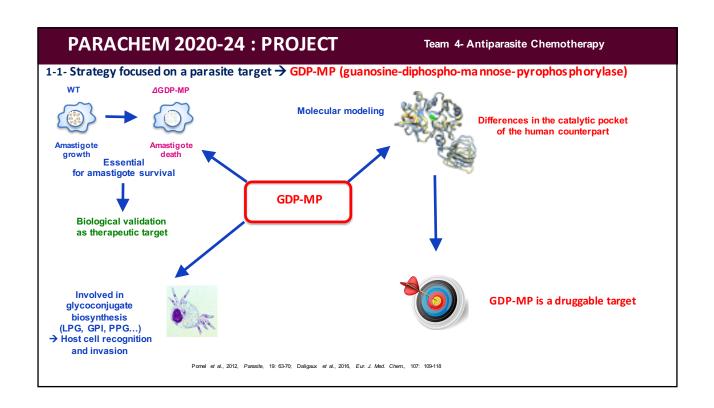


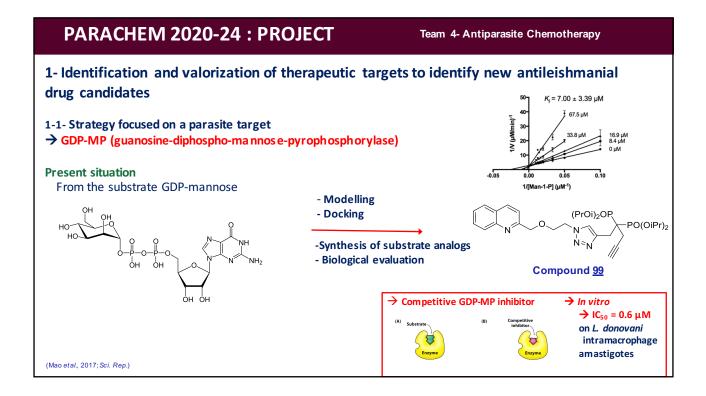


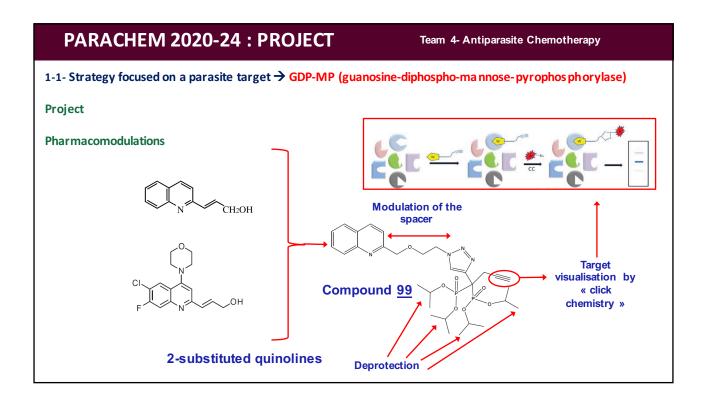












## PARACHEM 2020-24 : PROJECT

Team 4- Antiparasite Chemotherapy

1-1- Strategy focused on a parasite target → GDP-MP (guanosine-diphospho-mannose-pyrophosphorylase)
Project

In vitro and in vivo biological evaluation

→ HTS on human and leishmanial recombinant GDP-MPs







- → Chemical libraries (BioCIS, ICSN, CNE, GSK and WIPO Re:Search BVGH) → New scaffolds
- → In vitro and in vivo evaluation against Leishmania donovani
  - → Hit identification
- → Docking analysis of identified hits



→ GDP-MP functional analysis





to determine its biochemical importance in the main Leishmania species

- → Functional analysis → knockout (CRISPR-Cas9) → analysing the phenotype of the GDP-MP KO
- → Optimizing the strategy of specific inhibitor development

### PARACHEM 2020-24: PROJECT

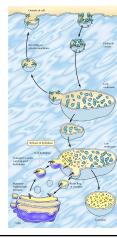
Team 4- Antiparasite Chemotherapy

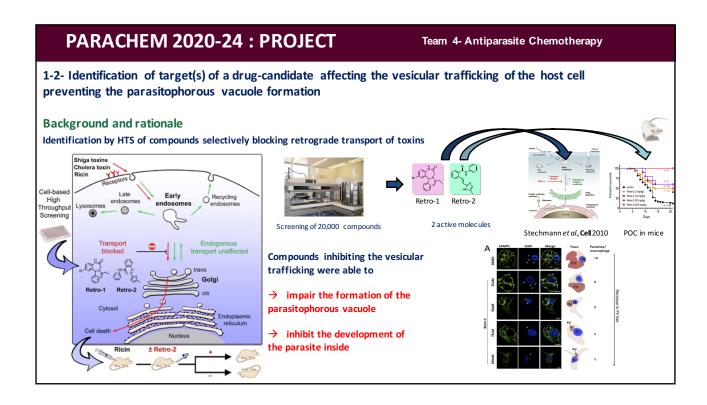
- 1- Identification and valorization of therapeutic targets to identify new antileishmanial drug candidates
- 1-2- Identification of target(s) of a drug-candidate affecting the vesicular trafficking of the host cell preventing the parasitophorous vacuole formation

**Background and rationale** 

Identifying a drug-candidate for the treatment of visceral leishmaniasis having the following characteristics:

- → An original mechanism of action that interferes with vesicle trafficking in host-cell impairing the development of the vacuole in which the parasite proliferates
- → No direct and intrinsic antiparasitic activity on the parasite itself in order to reduce the risk of drug resistance
- $\rightarrow$  No toxicity on the host-cell
- → A suitable druggability for oral or intravenous administration





# PARACHEM 2020-24: PROJECT

Team 4- Antiparasite Chemotherapy

1-2- Identification of target(s) of a drug-candidate affecting the vesicular trafficking of the host cell preventing the parasitophorous vacuole formation

### **Present situation**



Getting an antileishmanial drug-candidate (compound ABMA-2) and one back-up (compound ABMA-3) from a library of 300 compounds selected as affecting vesicular trafficking

### Compound ABMA-2

- → In vitro activity
  - → IC<sub>50</sub> = 40 nM on intramacrophage amastigotes
  - $\rightarrow$  Selectivity Index = CC<sub>50</sub>/IC<sub>50</sub> = 5625
  - → RetroLeishma Index= IC<sub>50</sub> amas axenic/IC<sub>50</sub> intramacrophage amastigotes = 240
- → Active in vivo on the L. infantum/BALB/c mice model at 10 mg/kg/day x 5 days by oral and iv routes (60% reduction of parasite burden)
- → No in vivo toxicity at 100 mg/kg
- → Microsomal stability (human): 46% at 45 min
- $\rightarrow$  PK
  - → Per os at 10 mg/kg: C<sub>max</sub>=80 ng/ml in 15 min
  - → Iv: at 1 mg/kg: Bioavailability 1%
  - $\rightarrow T_{1/2} = 4h$



### PARACHEM 2020-24 : PROJECT

Team 4- Antiparasite Chemotherapy

1-2- Identification of target(s) of a drug-candidate affecting the vesicular trafficking of the host cell preventing the parasitophorous vacuole formation

### **Project**

The goals of the study consist in:



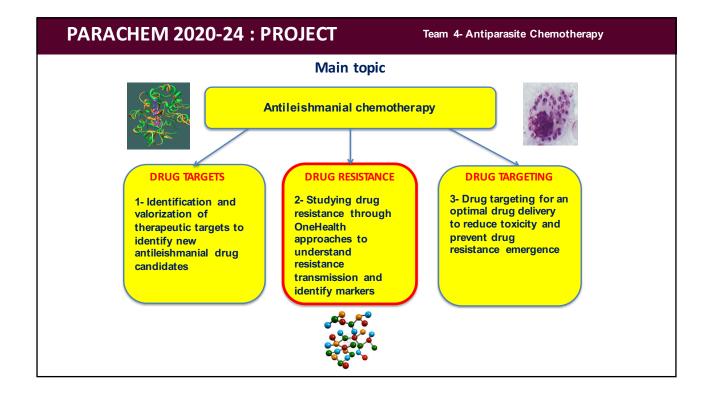




ANR LeishmaStop



- Understanding the basis for cellular protection against Leishmania given by ABMA-2 and ABMA-3 by confocal microscopy and videomicroscopy
  - → Effect on the parasitophorous vacuole development using GFP expressing parasites and markers of different steps of the endo-lysosomal pathway (such as EEA1, Rab7, Lamp1, etc.)
- Identifying drug targets of ABMA-2 and ABMA-3 by proteomic analysis and siRNA libraries
- Identifying the intracellular localization of the targets by confocal microscopy
  - in infected macrophages during cell invasion and intracellular parasite development
    - → If altered localization after treatment → confirmation of involvement in the mechanism of action of the compounds
- Functional analyses of the identified target genes
  - → Knockout using CRISPR-Cas9, or by knockdown using si/shRNA specific of the target genes
  - → Phenotype analysis of these cell lines and as well as their susceptibility to ABMA-2/ABMA-3



# PARACHEM 2020-24: PROJECT

Team 4- Antiparasite Chemotherapy

# 2- Studying drug resistance through OneHealth approaches to understand resistance transmission and identify markers

Background, rationale and objectives

- → Development of drug resistance in the field
  - → Resistance to antimonials
    - → Main molecular mechanisms described from isolates/clones
      - → What about the influence of the successive hosts (mammal and insect) in the transmission of drugresistant parasites ?

### **Project**

→ Developing a OneHealth concept for studying drug resistance parameters by modelling a natural life cycle of cutaneous leishmaniasis

### PARACHEM 2020-24: PROJECT Team 4- Antiparasite Chemotherapy Setting up of a natural life cycle with Leishmania major, its host Meriones shawii analysis Phenotype of selective and the insect vector Phlebotomus papatasi collected in gerbil holes in Algeria pressions from host change (Collaboration with Algeria Pasteur Institute, Dr. Z. Harrat) parasite biological parameters: Institut Pasteur In vitro et in vivo fitness d'Algérie **BioCIS** Virulence Variation of resistance intensity Analyses **Parasites** Stability/reversibility of Resistant? →Virulent? resistance breeding Parasite recovery Molecular analysis of levels of Metacyclic amplification and expression of some Parasite recovery → Infected ? promastigotes molecular markers: Gerbils infection by L. major MRPA (Multidrug Related Protein A) WT and Drug-Resistant MDR1 (Multidrug Resistance 1) TP (Tryparedoxine Peroxidase) **Blood** meal Infected Insects **Blood** meal gerbils → Infected ?

# PARACHEM 2020-24: PROJECT

Team 4- Antiparasite Chemotherapy

2- Studying drug resistance through OneHealth approaches to understand resistance transmission and identify markers

The natural life cycle of *L. major* → Proposed as a predictive model to measure the resistance parameters for any drug candidate in development

- → First application to:
  - → Meglumine antimoniate

- → Next applications
  - → Amphotericin B
  - → Any drug candidate to assess the risk of drug resistance (ABMA-2, ...)

# **PARACHEM 2020-24: PROJECT**

### Team 4- Antiparasite Chemotherapy

### **Conclusion**

- → Main PARACHEM objectives
  - $\rightarrow$  Reinforcing relevant international collaborations
  - → Participating to the development of new antileishmanial drug candidates /formulations





