

# Biological pathways discriminating African trypanotolerant and trypanosusceptible cattle breeds

Moana Peylhard, David Berthier, Laurence Flori, Guiguigbaza-Kossigan Dayo, Isabelle Chantal, Sophie Thevenon

#### ▶ To cite this version:

Moana Peylhard, David Berthier, Laurence Flori, Guiguigbaza-Kossigan Dayo, Isabelle Chantal, et al.. Biological pathways discriminating African trypanotolerant and trypanosusceptible cattle breeds. Trypanosomatid parasites: From the field to the lab, Dec 2017, Paris, France. 1 p., 2017. hal-02788332

## HAL Id: hal-02788332 https://hal.inrae.fr/hal-02788332v1

Submitted on 5 Jun 2020

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



### **Individual registration form**

To send via email to <a href="mailto:frederic.bringaud@u-bordeaux.fr">mailto:frederic.bringaud@u-bordeaux.fr</a>; <a href="mailto:lkohl@mnhn.fr">lkohl@mnhn.fr</a>; <a href="mailto:rotureau@pasteur.fr">rotureau@pasteur.fr</a>

Before November 10 <sup>th</sup> 2017

Talk / Poster title: Biological Pathways Discriminating African Trypanotolerant and Trypanosusceptible Cattle Breeds?

#### Authors:

Moana Peylhard<sup>1,2</sup>, David Berthier<sup>1,2</sup>, Laurence Flori<sup>3,4</sup>, Guiguigbaza-Kossigan Dayo<sup>5</sup>, Isabelle Chantal<sup>12</sup>, Sophie Thévenon<sup>1,2</sup>

#### Affiliations:

- 1 CIRAD, UMR INTERTRYP, F-34398 Montpellier, France.
- 2 INTERTRYP, Univ Montpellier, CIRAD, IRD, France
- 3 CIRAD, UMR SELMET, F-34398 Montpellier, France
- 4 SELMET, Univ Montpellier, CIRAD, INRA, SupAgro, Montpellier, France
- 5 CIRDES unité URBIO Bobo-Dioulasso

#### Key words (5 maximum):

Animal African Trypanosomosis, trypanotolerance, RNA-seq, pathways analysis, host\*parasite interactions

#### Abstract (250 words maximum):

Animal African Trypanosomosis (AAT) is a vector-borne disease caused by blood protozoan parasites of the Trypanosoma genus. It represents a major constraint to the development of cattle breeding in the humid and sub-humid zones of Africa because of the high morbidity and mortality it causes. Zebu breeds and European taurine breeds are very susceptible to AAT and they usually die in the absence treatment. On the contrary, some taurine breeds in West Africa have the capacity to tolerate the disease and are called trypanotolerant.

The trypanotolerant phenotype is known to be polygenic and multifactorial, but up to now, its mechanisms remain unknown. In order to decipher the molecular bases of trypanotolerance, we chose to analyse the genes expression of blood cells of susceptible and tolerant cattle during an experimental infection, performed in Burkina Faso, in 40 cattle from five West African breeds with T. congolense. mRNA were extracted from blood, comprising bovine leukocytes and parasites, before and during the infection and were sequenced using a Illumina highSeq2000. We mapped the reads on the bovine and trypanosome genomes, counted the reads on the annotated genes and performed a differential expression analysis. The genes identified as differentially expressed during the infection were then analysed using the Ingenuity Pathway Analysis software in order to identify enriched functional patterns. The functional analyses highlighted upstream regulators and canonical pathways associated with the immune response, the cell proliferation and signaling. Very fine differences in the modulation of the response between trypanotolerant and susceptible cattle were observed.