



**Offre n°2025-08790**

**Post-Doctoral Research Visit F/M**  
**Postdoctoral fellowship: Metabolic**  
**modelling of the rumen microbiome for**  
**guiding strategies of methane reduction**

**Type de contrat :** Fixed-term contract

**Contrat renouvelable :** Oui

**Niveau de diplôme exigé :** PhD or equivalent

**Fonction :** Post-Doctoral Research Visit

**A propos du centre ou de la direction fonctionnelle**

The Inria center at the University of Bordeaux is one of the nine Inria centers in France and has about twenty research teams.. The Inria centre is a major and recognized player in the field of digital sciences. It is at the heart of a rich R&D and innovation ecosystem: highly innovative SMEs, large industrial groups, competitiveness clusters, research and higher education players, laboratories of excellence, technological research institute...

**Contexte et atouts du poste**

The fellowship is associated to the ANR-funded [H2Rumen project](#), led by [Rafael Muñoz-Tamayo](#).

The fellowship is organised in two parts: this 18-months contract, hosted at Bordeaux (Inria PLEIADE), followed by a 14-month extension at Paris Saclay (INRAE, AgroParisTech).

The postdoc fellow will collaborate with [Clémence Frioux](#), [Simon Labarthe](#) and [David James Sherman](#) from the [PLEIADE](#) team (Inria, INRAE, Bordeaux) and with [Rafael Muñoz-Tamayo](#) from the [MoSAR](#) team (INRAE, AgroParisTech, Université Paris-Saclay). Pleiade hosts researchers in mathematics, computer science and computational biology working on developing approaches for microbial community characterisation.

**Project description.** Ruminants play a significant role in human nutrition and food security. Ruminants can harvest nutrients from forage diets rich in fibres and transform them into human-edible products with high-quality proteins. Feed transformation occurs mainly in the rumen through a metabolic cascade of hydrolytic and fermentative reactions carried out by a complex microbial community (rumen microbiota) constituted by hundreds of species that include bacteria, archaea and eukaryotes. During rumen fermentation, methane ( $\text{CH}_4$ ) is produced and eructed by the animal. This  $\text{CH}_4$  contributes to 44% of greenhouse gas emissions from the livestock sector.

Reducing methane ( $\text{CH}_4$ ) emissions from ruminants is a major challenge for the livestock sector. An optimal  $\text{CH}_4$  mitigation strategy should also induce co-benefits such as enhanced animal productivity and health.  $\text{CH}_4$  is produced during the fermentation of feeds in the rumen. This process is carried out by a complex microbial community (rumen microbiota) and mediated by hydrogen ( $\text{H}_2$ ) in the rumen ecosystem. Our knowledge on the drivers that shape  $\text{H}_2$  flows is still incomplete.  **$\text{H}_2$ Rumen aims to generate fundamental knowledge on  $\text{H}_2$  flows in the rumen ecosystem and to translate this knowledge into predictive mathematical models of rumen fermentation.** Our scientific outputs will be of high value for the optimal design of  $\text{CH}_4$  mitigation strategies with co-benefit for the animal. Our hypothesis is that thermodynamics and microbial interactions jointly control hydrogen transactions in the rumen ecosystem.  $\text{H}_2$ Rumen addresses the following fundamental scientific question: Where does  $\text{H}_2$  go? We will answer this question with an integrative approach that combines *in vitro* experiments, *in silico* modelling and omics approaches. In addition to the impact on ruminant livestock, our methods might be applicable to other ecosystems such as the human gut, engineering bioreactors and fermented food ecosystems.

## Mission confiée

**The postdoc project aims at developing metabolic models of rumen microbiota with capabilities for designing microbial strategies for reducing methane emissions and improving rumen fermentation.** It will contribute to fill the existing gap on how to integrate microbial genomic information into rumen fermentation models [1].

Genome-scale network reconstructions will be done with state-of-the-art methods and pipelines developed in PLEIADE. Metabolic capabilities of rumen microbial communities will be explored using Metage2Metabo [2]. Kinetic models will be

built on the basis of existing models developed at MoSAR [3,4]. Models will be constructed using experimental data produced by our partners at [UMRH](#) and [MoSAR](#).

[1] Muñoz-Tamayo R, Davoudkhani M, Fakhri I, Robles-Rodriguez CE, Rubino F, Creevey CJ, et al. Review: Towards the next-generation models of the rumen microbiome for enhancing predictive power and guiding sustainable production strategies. *animal*. 2023;17: 100984. doi:10.1016/J.ANIMAL.2023.100984

[2] Belcour A, Frioux C, Aite M, Bretaudeau A, Hildebrand F, Siegel A. Metage2metabo, microbiota-scale metabolic complementarity for the identification of key species. *Elife*. 2020;9: e61968. doi:10.7554/eLife.61968

[3] Muñoz-Tamayo R, Chagas JC, Ramin M, Krizsan SJ. Modelling the impact of the macroalgae *Asparagopsis taxiformis* on rumen microbial fermentation and methane production. *Peer Community J*. 2021;1: e7. doi:10.24072/PCJOURNAL.11

[4] Fakhri I, Got J, Robles-Rodriguez CE, Siegel A, Forano E, Muñoz-Tamayo R. Dynamic genome-based metabolic modeling of the predominant cellulolytic rumen bacterium *Fibrobacter succinogenes* S85. *mSystems*. 2023;8: e01027-22. doi:10.1128/msystems.01027-22

## Principales activités

The postdoc fellow will be part of the project H2Rumen funded by the French National Agency for Research (ANR). The specific objectives are

- i. To obtain metabolic models of the species constituting a rumen microbial mini-consortium.
- ii. To derive a bag-of-genome model of a complex rumen consortium.
- iii. To develop a community-scale kinetic models of mini and complex consortia accounting for H<sub>2</sub> transactions and thermodynamic control.

The work is organized in two parts. 18 months in PLEIADE to work on the objectives (i,ii) and 14 months at MoSAR to work on the objective (iii).

## Compétences

Technical skills and level required:

- Experience in metabolic modelling with willingness to learn dynamic modelling or viceversa
- Expertise in programming: Python, R or Scilab/Matlab

Languages:

- Proficiency in English

Relational skills:

- Excellent communications skills, willing to discuss with scientists with different backgrounds

## Avantages

- Subsidized meals
- Partial reimbursement of public transport costs
- Leave: 7 weeks of annual leave + 10 extra days off due to RTT (statutory reduction in working hours) + possibility of exceptional leave (sick children, moving home, etc.)
- Possibility of teleworking and flexible organization of working hours
- Professional equipment available (videoconferencing, loan of computer equipment, etc.)
- Social, cultural and sports events and activities
- Access to vocational training
- Social security coverage

## Rémunération

The gross monthly salary will be 2788€ (before social security contributions and monthly withholding tax)

## Informations générales

- **Thème/Domaine** : Computational Biology  
Biologie et santé, Sciences de la vie et de la terre (BAP A)
- **Ville** : Talence
- **Centre Inria** : [Centre Inria de l'université de Bordeaux](#)
- **Date de prise de fonction souhaitée** : 2025-10-01
- **Durée de contrat** : 1 year, 6 months
- **Date limite pour postuler** : 2025-05-31

## Contacts

- **Équipe Inria** : [PLEIADE](#)
- **Recruteur** :  
Frioux Clemence / [clemence.frioux@inria.fr](mailto:clemence.frioux@inria.fr)

## A propos d'Inria

Inria est l'institut national de recherche dédié aux sciences et technologies du numérique. Il emploie 2600 personnes. Ses 215 équipes-projets agiles, en général communes avec des partenaires académiques, impliquent plus de 3900 scientifiques pour relever les défis du numérique, souvent à l'interface d'autres disciplines. L'institut fait appel à de nombreux talents dans plus d'une quarantaine de métiers différents. 900 personnels d'appui à la recherche et à l'innovation contribuent à faire émerger et grandir des projets scientifiques ou entrepreneuriaux qui impactent le monde. Inria travaille avec de nombreuses entreprises et a accompagné la création de plus de 200 start-up. L'institut s'efforce ainsi de répondre aux enjeux de la transformation numérique de la science, de la société et de l'économie.

## L'essentiel pour réussir

We are seeking a researcher with a PhD in applied mathematics, computational biology, or process engineering. The ideal candidate should have expertise in metabolic modelling and a strong willingness to expand their skills in dynamic modelling, or alternatively, experience in dynamic modelling with an eagerness to develop proficiency in metabolic modelling.

In addition to technical expertise, the candidate should have a demonstrated ability to write and publish research articles in peer-reviewed journals. Strong communication skills are essential, as the role involves effectively presenting scientific results to both academic and interdisciplinary audiences.

**Attention:** Les candidatures doivent être déposées en ligne sur le site Inria. Le traitement des candidatures adressées par d'autres canaux n'est pas garanti.

## Consignes pour postuler

If you are interested by this job, please could you apply on website [jobs.inria](https://jobs.inria.fr) with the following documents :

- cv

- cover letter
- recommendation letter

**Sécurité défense :**

Ce poste est susceptible d'être affecté dans une zone à régime restrictif (ZRR), telle que définie dans le décret n°2011-1425 relatif à la protection du potentiel scientifique et technique de la nation (PPST). L'autorisation d'accès à une zone est délivrée par le chef d'établissement, après avis ministériel favorable, tel que défini dans l'arrêté du 03 juillet 2012, relatif à la PPST. Un avis ministériel défavorable pour un poste affecté dans une ZRR aurait pour conséquence l'annulation du recrutement.

**Politique de recrutement :**

Dans le cadre de sa politique diversité, tous les postes Inria sont accessibles aux personnes en situation de handicap.