

JOB OFFER

Postdoctoral position

Federated Statistical Learning for New Generation Meta-Analyses of Large-scale and Secured Biomedical Data (Fed-BioMed)

Project Description.

The project Fed-BioMed focuses on methodological and technical advances towards the development of a novel generation of federated learning methods for the analysis of private and large-scale multi-centric biomedical data. The project has a specific focus on the efficient federation of frameworks robust to data heterogeneity and uncertainty, and tackles the following scientific challenges:

- Methodological. Extending the federated paradigm to novel scalable approaches to probabilistic modeling and prediction from siloed data.
- Technical. Developing our federated learning framework through a self-contained system that can be securely deployed across different centers and collaborators (fedbiomed.gitlabpages.inria.fr).
- Translational. Demonstrating federated learning on two applications: 1) Discovering novel genetic underpinnings of neurological and psychiatric disorders, and 2) Prediction of response to immunotherapy from the analysis of federated lung imaging data.

During the project the candidate will:

- Develop learning methods for federated analysis for private and distributed data;
- Deploy advanced statistical learning methods into a wide range of biomedical/clinical applications;
- Interact with INRIA researchers and engineers, and participate to the scientific life of the team;

Required competences.

Demonstrable experience in some of the following topics (the more the better):

- Statistics, Bayesian Modeling;
- Optimization, Distributed Computing;
- Python and PyTorch/TensorFlow;
- Biomedical Data Analysis;
- Signal Processing;

Strong communication abilities are necessary, as well as motivation in taking responsibilities (e.g. supervision, organization of scientific events).

Hosting Research Group.

[Epione](#) team (Inria), located in the tech park of Sophia Antipolis (France).

The longstanding research activity of our group revolves around the analysis and treatment of biomedical data, with a focus in machine learning, medical imaging, computational anatomy and computational physiology. Over the past twenty years the group developed innovative approaches in image processing, statistical learning and patient-specific biophysical modeling, with translation to the clinical domain, and to the creation of several biotech startups. The group is currently composed by 6 permanent researchers, several postdoc fellows and research engineers, and by more than 20 PhD students.



Offer.

18 months contract
 Salary upon experience
 Starting date: asap

Contact.

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References:

- Yann Fraboni, Richard Vidal, Laetitia Kameni, Marco Lorenzi. Clustered Sampling: Low-Variance and Improved Representativity for Clients Selection in Federated Learning. Proceedings of the 38th International Conference on Machine Learning (ICML 2021).
- Yann Fraboni, Richard Vidal, and Marco Lorenzi. Free-rider Attacks on Model Aggregation in Federated Learning. In AISTATS 2021.
- Irene Balelli, Santiago Silva, and Marco Lorenzi. A Probabilistic Framework for Modeling the Variability Across Federated Datasets of Heterogeneous Multi-View Observations. International Conference of Information processing in Medical Imaging, IPMI 2021.
- Santiago Silva, Boris Gutman, Barbara Bardoni, Paul M Thompson, Andre Altmann, Marco Lorenzi. *Multivariate Learning in Distributed Biomedical Databases: Meta-analysis of Large-scale Brain Imaging Data*. IEEE International Symposium on Biomedical Imaging (ISBI), Venice, 2019.
- Marco Lorenzi, Andre Altmann, Boris Gutman, et al. *Susceptibility of brain atrophy to TRIB3 in Alzheimer's disease: Evidence from functional prioritization in imaging genetics*. Proceedings of the National Academy of Sciences of the United States of America (PNAS). March 20, 2018. 115 (12) 3162-3167.