

Programme COMPAS 2024

6th-8th November

TEAMS LINK:

https://teams.microsoft.com/l/meetup-join/19%3ameeting_ZmYxMTc5MjAtMWlyNy00ZDM0LWJkM2MtZDg5YmJmMzdY2E2%40thread.v2/0?context=%7b%22Tid%22%3a%22158716cf-46b9-48ca-8c49-c7bb67e575f3%22%2c%22Oid%22%3a%227f728b67-7aa4-4166-8de9-2daa6efec a1f%22%7d

LOCATION: Espace Rabelais, Salle des Thèses, Campus du Saulcy, Metz

Wednesday 6th

First session (30 minutes presentation + 15 minutes discussion):

Philology and stemmatology

12am: *Welcome lunch*

1:30-2 pm: Coffee and welcome talk - *Sophie Robert-Hayek, University of Lorraine*

2:00-2:45 pm: The Stemma of Ben Sira's Hebrew Manuscripts as a Test Case for Computational Stemmatology - *Frédérique Rey, University of Lorraine*

2:45-3:15 pm: The Relevance of Minor Variants for the Validation of Stemmatic Relations – “and” in Isaiah as a Test Case - *Benjamin Ziemer, University of Halle*

3:15-4:00 pm: Genealogical Trees in Biblical Textual Studies Phylogenetic Insights from Kennicott's Collation of the Hebrew Bible - *Luigi Bambaci, EPHE*

4:00-4:30 pm: *Break and informal discussion: 30 minutes*

4:30-5:30 pm: *First keynote: Jean-Baptiste Camps, Ecole des Chartres (45 minutes + 15 minutes discussion)*

Imbalanced evolutionary trees and lost manuscripts

7:30 pm: *Dinner at the Theatris*

Thursday 7th

Second session (30 minutes presentation + 15 minutes discussion):

Computational approaches to stemmatology

9:00-9:45 am: Beyond Singular Readings: Priors on Scribal Habits from Non-substantive Readings - *Joey McCollum, Australian Catholic University*

9:45-10:30 am: Classifying transcriptional changes using Large Language Models - *Robert Turnbull, Australian Catholic University*

10:30-11:15 am: Exploring phylogenetic models for stemmatology applications - *Jacques Istas, Université Grenoble Alpes*

11:15-11:45 am: *Break and informal discussion: 30 minutes*

11:45-12:15 am: The Importance of Metric Selection for the Benchmarking of Stemmatology Algorithms - *Sophie Robert-Hayek, University of Lorraine*

12:30-2:30 pm: *Lunch break*

2:30-3:15 pm: Localization of new textual variants within an existing stemma using neural networks - *Armin Hoenen, independent scholar*

3:15-4:00 pm: Imi Khily in Khanty Folklore: A Phylogenetic Approach to Tale Type Identification in Ob Ugric Narratives - *Anna Grinevich, Institute of Philology*

4:00-4:30 pm: *Break and informal discussion: 30 minutes*

4:30-5:30 pm: *Second Keynote: Philipp Roelli, University of Zurich (45 minutes + 15 minutes discussion)*

*Some Methodological Considerations Concerning Computational Tools in the Reconstruction of the *Stemma codicum*.*

6:00-7:00 pm: *Visit of Metz*

7:30 pm: *Dinner*

Friday 8th

Third session (30 minutes presentation + 15 minutes discussion):

Current trends in phylogeny and applications

9:00 - 10:00 am: *Third Keynote (45 minutes + 15 minutes discussion)*

Models of biological sequence of evolutions, beyond independence between sites - Bastien Boussau, LBBE

10:00 - 11:15 am: *Fourth Keynote: Robin Ryder (1 hour + 15 minutes discussion)*

Bayesian Methods in Historical Linguistics: a Workflow

Phylogenetics of Primate Call Functions: a methodological proposal and case study -

Valentin Thouzeau, Guillaume Dezecache, Philippe Schlenker, Ewan Dunbar, Emmanuel Chemla and Robin Ryder

11:30 - 12:30 am: *Coffee break and round table and discussion around further collaborations*

12:30 am: *Goodbye lunch*