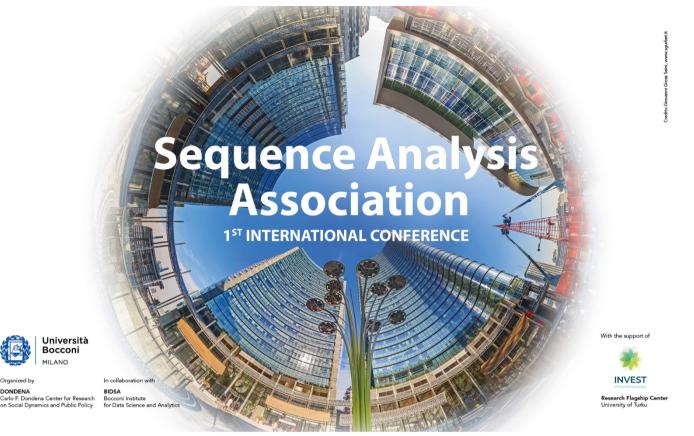
Sequence Analysis Association



The Sequence Analysis Association (https://sequenceanalysis.org) is pleased to announce its first International Conference on June 10-12, 2020 at Bocconi University, Milan, (Italy) organized by DONDENA Centre for Research on Social Dynamics and Public Policy (www.dondena.unibocconi.it) with the support of BIDSA (Bocconi Institute for Data Science and Analytics; www.bidsa.unibocconi.eu) and of INVEST Research Flagship Center (University of Turku, Finland; https://sites.utu.fi/invest/)

The aim of the conference is to bring together scholars applying or developing Sequence Analysis and other innovative methods for analysing longitudinal data in social, economic, managerial, political, health, or environmental sciences.

Since the pioneering work of Andrew Abbott over three decades ago, with the recent releases of versatile software packages, Sequence Analysis has become an increasingly popular exploratory method in the social sciences. The conference will focus not only on the development of Sequence Analysis methodology itself but also on its relations to and integration with other methodological frameworks including qualitative analysis, event history analysis, Markov-based models, and other longitudinal stochastic approaches. The aim is to discuss how to further develop Sequence Analysis and how different methodological approaches can complement one another.

We are very pleased to announce that our **keynote speakers** will be

Andrew Abbott (University of Chicago),

<u>Anette Fasang</u> (WZB Berlin Social Science Center, and Humboldt University, Berlin), <u>Emilio Zagheni</u> (Max Planck Institute for Demographic Research).

All submissions connected with Sequence Analysis or related methods in the social sciences are welcome, especially applications of innovative methodology, new methodological developments, method comparisons, and theoretical discussions linking substantive theory with methodological choices. Proposals across all scientific disciplines and application domains are welcome. The **deadline for the submission** of original papers or extended abstract (of at least two pages not including references) is **January 13th 2020**.

For information, registration and submission, visit the conference website: <u>https://www.unibocconi.eu/saaconference2020</u> To receive updates and relevant information <u>register</u> to the SAA and to our newsletter, or follow us on twitter (@SeqAnalysisAssn).

We would be grateful if you would forward the link to everyone interested.

The organizing committee: Raffaella Piccarreta, Matthias Studer, Emanuela Struffolino, Satu Helske, Brendan Halpin, Tim Liao, Gilbert Ritschard, Francesco Billari.