

Introduction to sequence analysis for social sciences

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Assignment

Use the data provided to illustrate life course trajectories of a sample of individuals from the German Family Panel (pairfam) data. Data come from [Raab and Struffolino \(2022\)](#). Additional details and code can be found in the book and in the companion github page

Prepare a 5 page max report in which you:

1. Import the data in R and describe the distribution of life states using relevant sequence analysis tools, including tables, indexes and figures.
2. Calculate life course dissimilarities in the data.
3. Use a clustering methods to derive typical trajectories and describe groups

4. (Optional) Describe group differences in the the distribution of life course clusters

Data Source

The data come from the German Family Panel (pairfam), release 10.0 Brüderl et al. (2019). A description of the study can be found in Huinink et al. (2011).

We gratefully acknowledge the permission of the pairfam team to share a reduced version of their data to illustrate all techniques presented in the book with real-world survey data.

If you are interested in using the complete data sets please turn to: <https://www.pairfam.de/en/>

Online references

- **Sequence Analysis Association (SAA)** <https://sequenceanalysis.org>
- **TraMineR R package** <http://traminer.unige.ch>
- **Sequence Analysis (2022) Marcel Raab and Emanuela Struffolino**, Sage <https://sa-book.github.io>

Suggested Readings

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23. R Piccarreta and F.C Billari. Clustering work and family trajectories by using a divisive algorithm. Journal of the Royal Statistical Society: Series A (Statistics in Society), 170(4):1061–1078, 2007.
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