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Identifying return-to-work trajectories among breast cancer survivors using sequence analysis

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Cancérologie eHESP 0 de l'Ouest

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1 Introduction & Objectives

Context

The return-to-work (RTW) process after breast cancer (BC) can be complex. Simple static measures used to assess RTW may ignore this dynamic multistage process that is yet important for targeting interventions aimed at reducing poor RTW outcomes.

Objectives

- (1) Identify RTW trajectories after BC using the sequence analysis method
- (2) Describe their underpinning personal, medical, psychological and occupational factors

Study population



The ELCCA II prospective cohort included 128 BC patients in 2015-2016.

- BC patients were followed at 1 month, 6 months and annually after 1 year following their BC diagnosis
- Medical and sociodemographic characteristics, and quality of life (QoL) were assessed at inclusion and at the follows-up.
- A restropective occupational calendar with a 6-month time scale from diagnosis up to 3 years later was completed by BC patients at their 4 year follow-up (Figure 1)
- BC Patients notified one of the 7 following five occupational situations : "full-time", "part-time", "sick leave", "unemployment", "training courses", "(pre)retirement", "other inactivities"



Exclusion criteria : - BC patients retired or unemployment at the time of diagnostic

Sequence analysis

- Sequence analysis is a quantitative biographical approach
- It generate a sequence for every BC patients based on theirs individual occupational situations every 6 months from their diagnosis up to 3 years after their diagnosis (Figure 2).
- Similarity between BC patients' occupational trajectories were estimated using the optimal matching (OM) distance algorithm¹.
- An agglomerative hierarchical clustering with the Ward's linkage method was used to identify
 patterns of occupational trajectories after BC patients diagnostic (i.e. clusters) based on the
 distance matrix previously elaborated.
- 5 occupational situations were kept in state sequence analysis (Figure 2)

01-021	Sick leave	Sick leave	Sick leave	Part-time	Part-time	Part-time	Full-time
01-020	Sick leave	Full time	Full time	Full time	Full time	Full time	Full time
01-006	Full time	Sick leave	Part-time	Part-time	Full time	Unemployment	Other inactivites
Patient ID	Diagnostic	6_M	12_M	18_M	24_M	30_M	36_M

<u>Figure 2</u>: Example of five BC patients sequences representing their occupational situation from their diagnostic up to 3 years later

¹Gabadinho, A., Ritschard, G., & Studer, M. (2011). Analyzing and Visualizing State Sequences in R with TraMineR. Journal Of Statistical Software, 40(4).

5 Discussion & Conclusion

- The application of sequence analysis highlighted the heterogeneity of the RTW process among BCS.
- It captures trajectories of multiple states and transitions provided a holistic, longitudinal and diachronic approach of RTW.
- Three different patterns of RTW trajectories after breast cancer were identified and underlined a number of key insights not found using conventional static methods.
- However, these results have to be confirmed by using a larger sample of BCS

Results

- 52 BC patients were active at the time of their diagnostic that completed their occupational calendar were included
 3 RTW trajectories' patterns were identified among BC patients by sequence analysis (Figure 3-A).
- 3 R I W trajectories patterns were identified among BC patients by sequence analysis (Figure 3-A).
 Socio-demographic, medical and QoL (QLQ-C30) varied according to RTW trajectories (Figure 3-B).

Figure 3 : Patterns of RTW trajectories among BC patients in the ELCCA II cohort (n= 52).



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