Clustering Discrete State Trajectories of Varying Lengths: Health Care Utilization Patterns

Laura Hatfield, PhD

Associate Professor, Harvard Medical School

ISPOR May 22, 2018

Baltimore, MD

Thank you to my collaborators

Brianna Heggeseth
- Williams College -> Macalester College

Megan Schuler
- RAND Boston

Nina Joyce
- Brown University
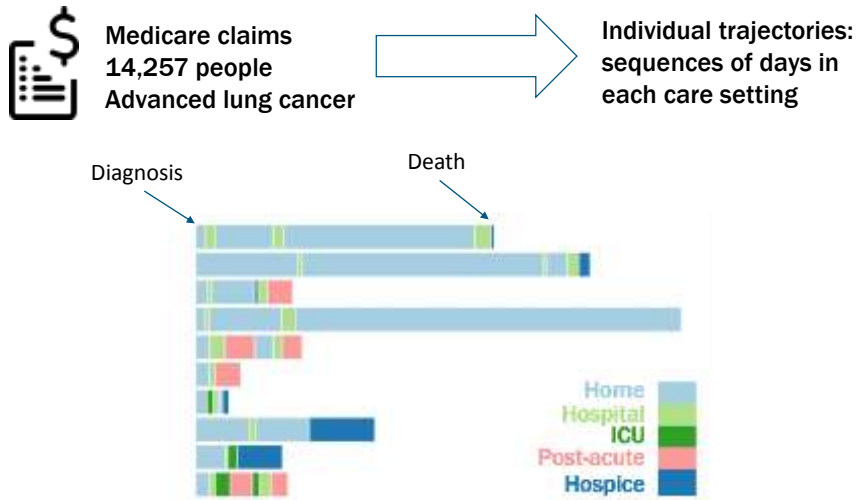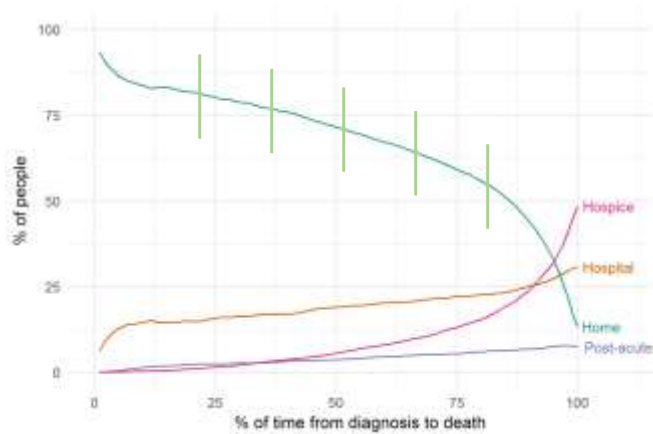
Haiden Huskamp
- Harvard Medical School

Elizabeth Lamont
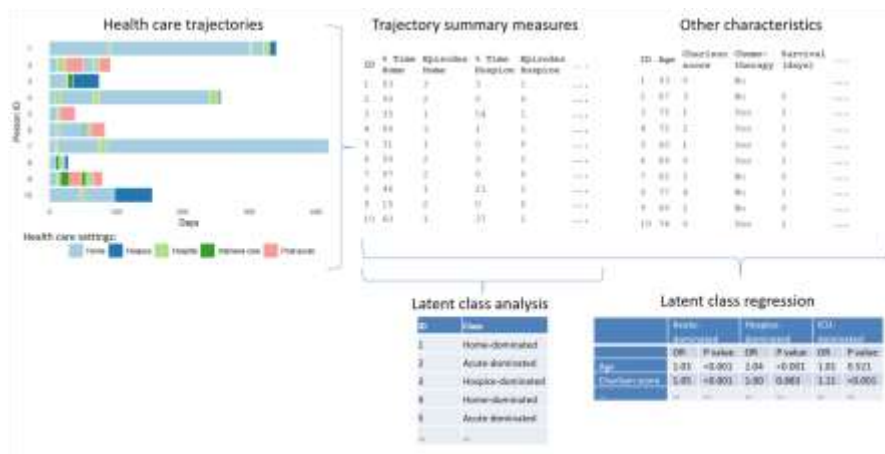- Harvard Medical School

Health care utilization trajectories

Medicare claims
14,257 people
Advanced lung cancer

→ Individual trajectories:
sequences of days in
each care setting

Diagnosis   Death

Home
Hospital
ICU
Post-acute
Hospice

Can we use clustering to **discover** and **illustrate** variation in experiences?
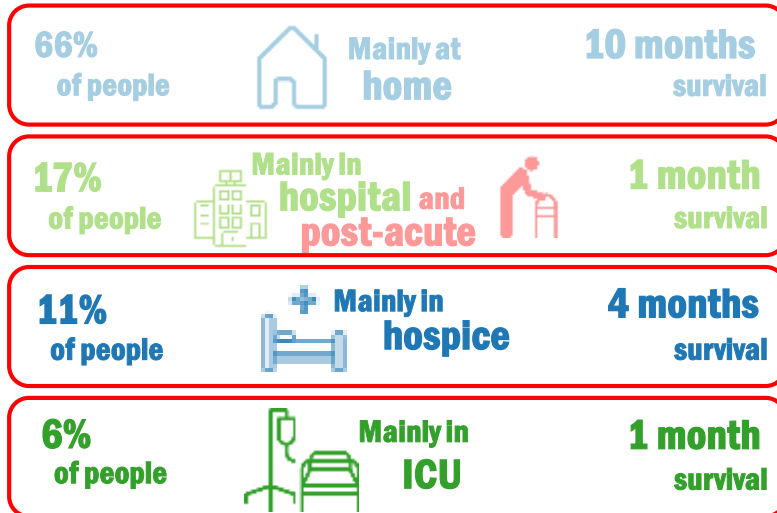
Feature extraction + LCA

Latent class analysis

For response pattern **y** and class c_k

$$\Pr(\mathbf{Y} = \boldsymbol{y}) = \sum_{k=1}^{K} \Pr(C = c_k) \prod_{j=1}^{J} \Pr(Y_j = y_j \mid C = c_k)$$
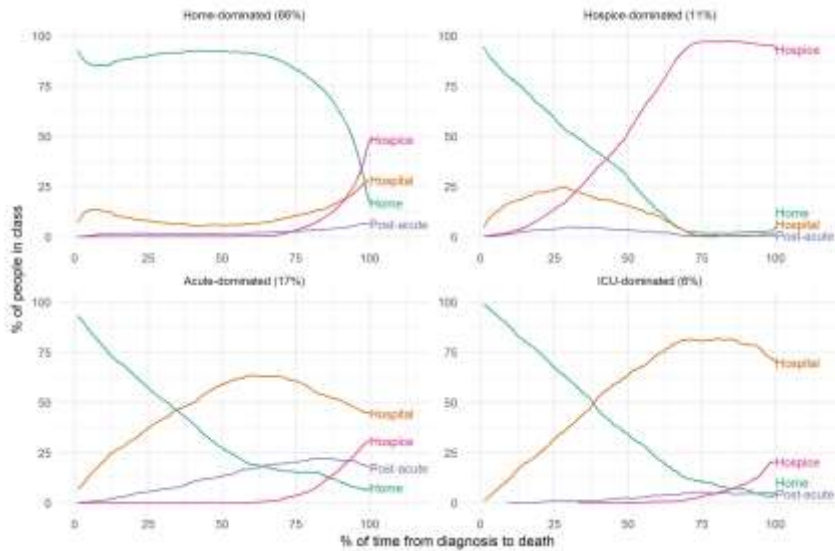
The class indicators are missing data.

Four distinct classes

66% of people	**Mainly at home**	**10 months** survival
17% of people	**Mainly in hospital** and **post-acute**	**1 month** survival
11% of people	**Mainly in hospice**	**4 months** survival
6% of people	**Mainly in ICU**	**1 month** survival

Source: Schuler et al (2017) Health Affairs. doi: 10.1377/hlthaff.2017.0448

Classes have distinct trajectories

Source: Schuler et al (2017) Health Affairs. doi: 10.1377/hlthaff.2017.0448

Remaining methods gaps

Limitations of feature extraction + LCA

| Discards ordering information |
| Requires good feature selection |
| Sensitive to choice of features |

A new distance measure

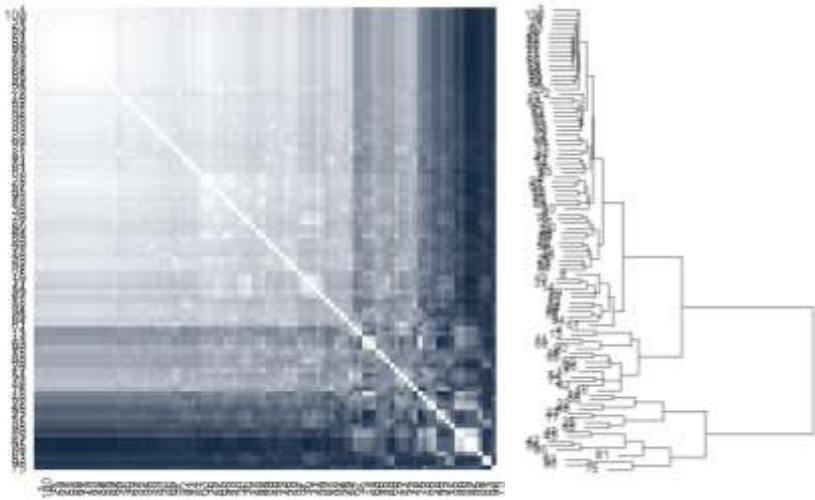| Uses sequence information directly |
| Does not require feature selection by investigator |
| Facilitates standard clustering methods |

Distance is a weighted combination of

1. moving average of discordant days and
2. length difference

$$
\begin{aligned}
d(\boldsymbol{a}, \boldsymbol{b}) \\
= w \frac{1}{K} \sum_{k=1}^{K} \frac{\sum |\boldsymbol{s}(a_t | t \in (k, k+\tau)) - \boldsymbol{s}(b_t | t \in (k, k+\tau))|}{2\tau} \\
+ (1-w) \frac{|l(\boldsymbol{a}) - l(\boldsymbol{b})|}{\max\{l(\boldsymbol{a}), l(\boldsymbol{b})\}}
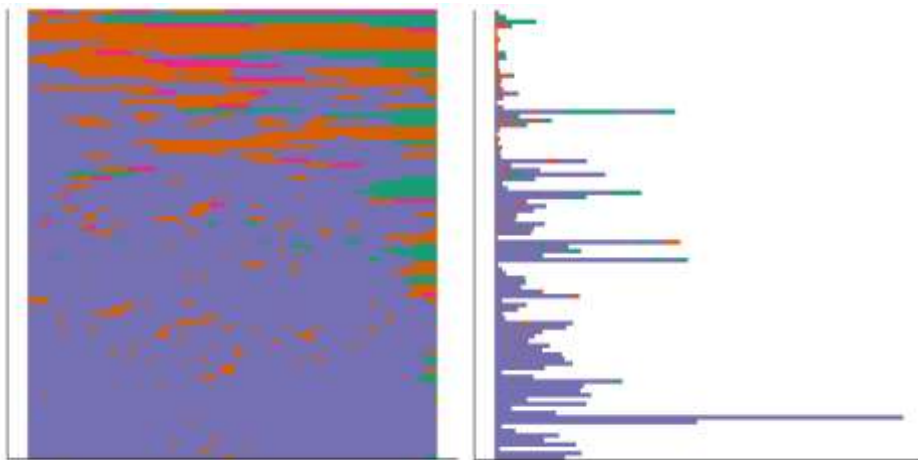\end{aligned}
$$

$\boldsymbol{s}(a_t | t \in (k, k+\tau))$ — Vector number of "days" in each state during time window of width τ
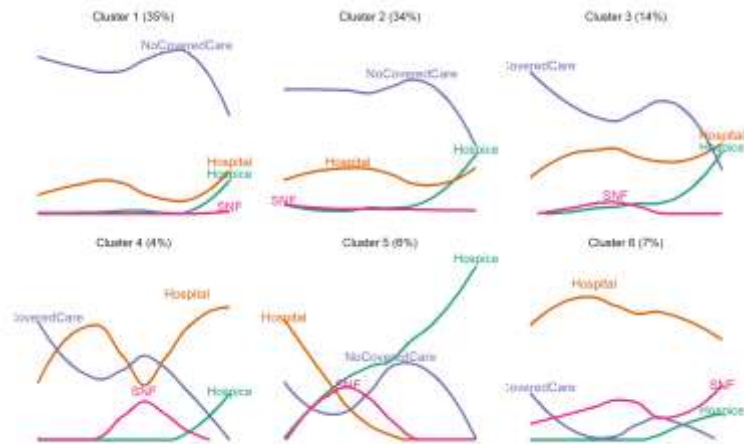
τ, w — Bandwidth and weight tuning parameters

Standardized time   Original time

Conclusions

- Clustering can show variation in longitudinal data
- Feature extraction enables use of LCA clustering
- Custom distance measure enables other clustering methods

hatfield@hcp.med.harvard.edu

Thanks!

@laura_tastic
@HPDSLab

healthpolicydatascience.org