

# Interactive visual analytics for medical data: application to COVID-19 clinical information during the first wave

José M. Enguita<sup>1</sup>, Diego García<sup>1</sup>, María D. Chiara<sup>2</sup>, Nuria Valdés<sup>3</sup>, Ana González<sup>1</sup>, Abel A. Cuadrado<sup>1</sup>, Ignacio Díaz<sup>1</sup>

<sup>1</sup>University of Oviedo - Dept of Electrical Engineering  
Edificio Torres Quevedo, módulo 2, Campus de Gijón 33204 - SPAIN

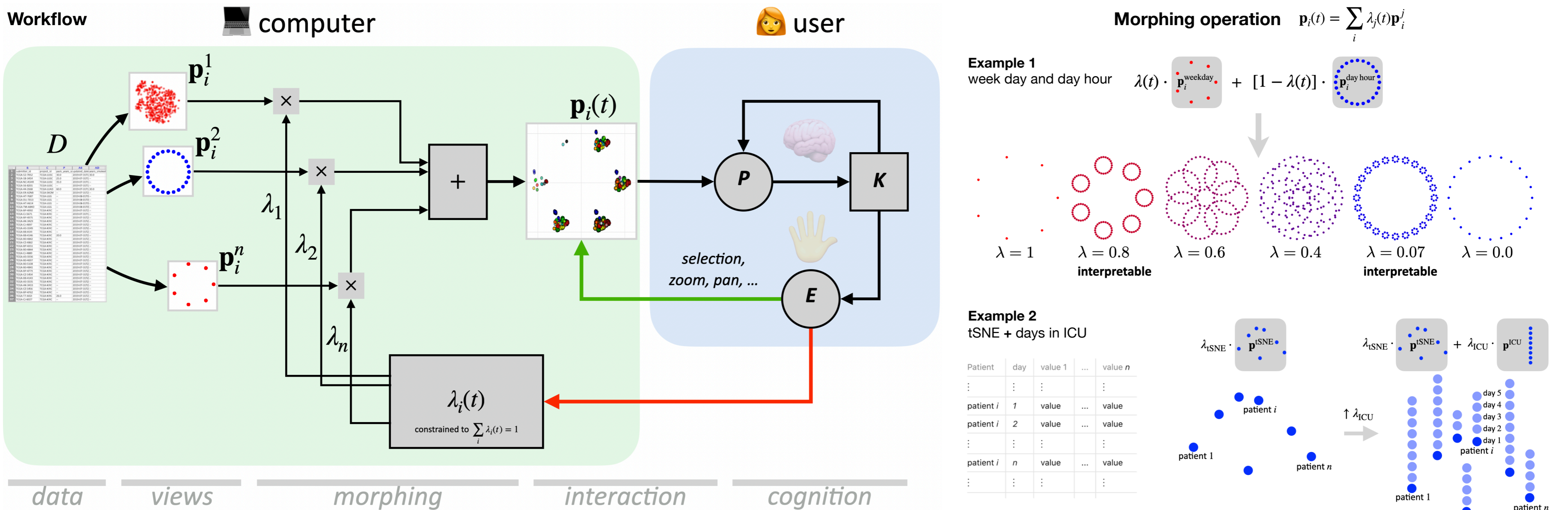
<sup>2</sup>Institute of Sanitary Research of the Principado de Asturias  
Hospital Universitario Central de Asturias, Oviedo 33011 - SPAIN

<sup>3</sup>Department of Internal Medicine, Section of Endocrinology and Nutrition  
Hospital Universitario de Cabueñes, Gijón 33204 - SPAIN

## Abstract

Biomedical data recorded as a result of clinical practice are often multi-domain —involving lab measurements, medication, patient attributes, logistic information—, and also highly unstructured, with high rates of missing data and asynchronously sampled measurements. In this scenario, we need tools capable of providing a broad picture prior to more detailed analyses. We present here a visual analytics approach that uses the morphing projections technique to combine the visualization of a *t*-SNE projection of clinical time series, with views of other clinical or patient's information. The proposed approach is demonstrated on an application case study of COVID-19 clinical information taken during the first wave.

## Methods



**Timeseries: dynamic time warping (DTW)**  
Medical evolution, drug administration

$d_{ij}^a$  DTW distance between timeseries  $i$  and  $j$  for measurement  $a$

Overall distance  $d_{ij}^{\text{overall}} = \sqrt{\sum_{k=1}^5 (d_{ij}^k)^2}$  DTW distance

**Dimensionality Reduction (t-SNE)**  
 $d_{ij} \rightarrow [tSNE] \rightarrow p_i^{\text{tSNE}}$

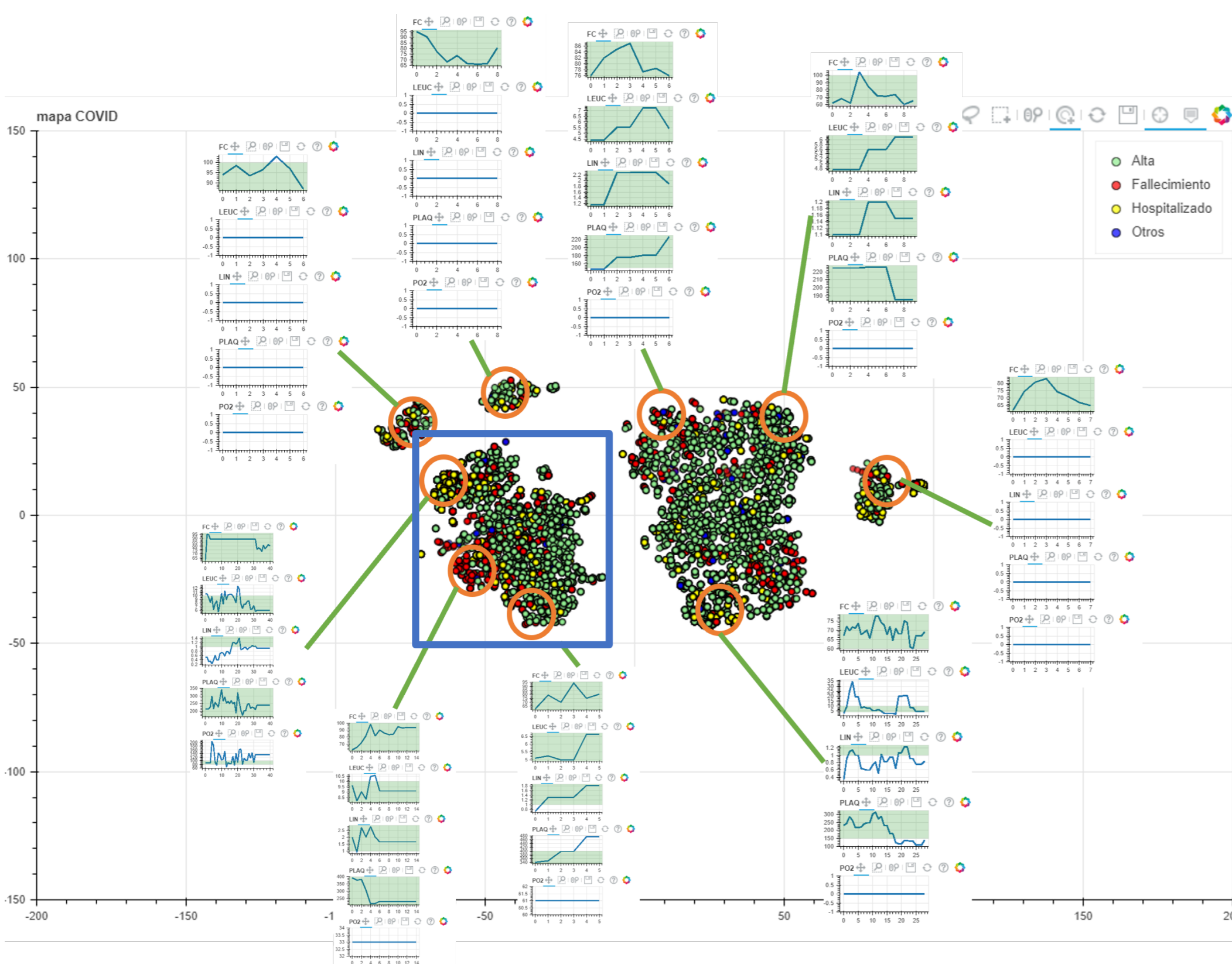
- close projections represent close (small  $d_{ij}$ ) time series behavior
- clusters reveal groups of similar timeseries behavior
- results in a map revealing the temporal behavior of samples

**Typical encodings**

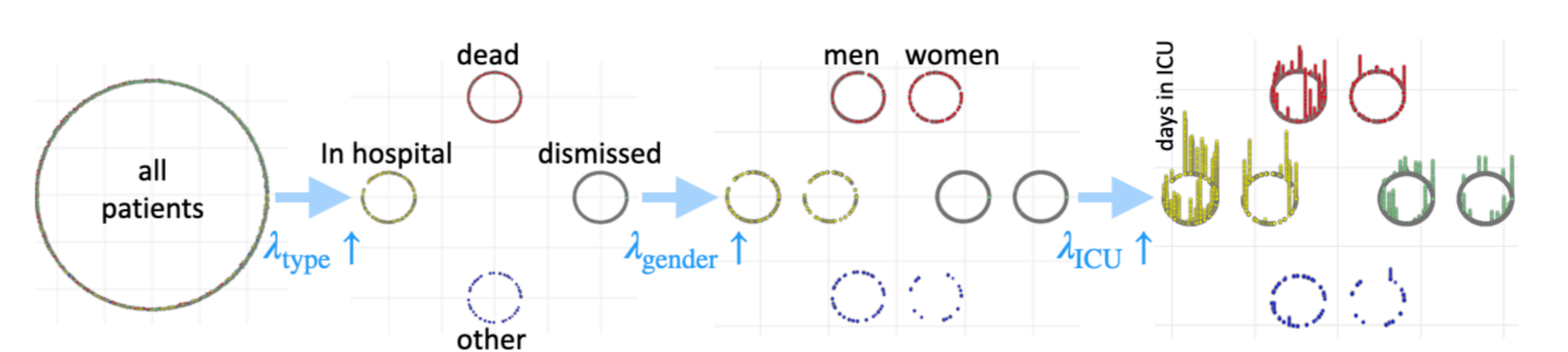
vertical horizontal circular 2D projection geo-location body-location

## Results

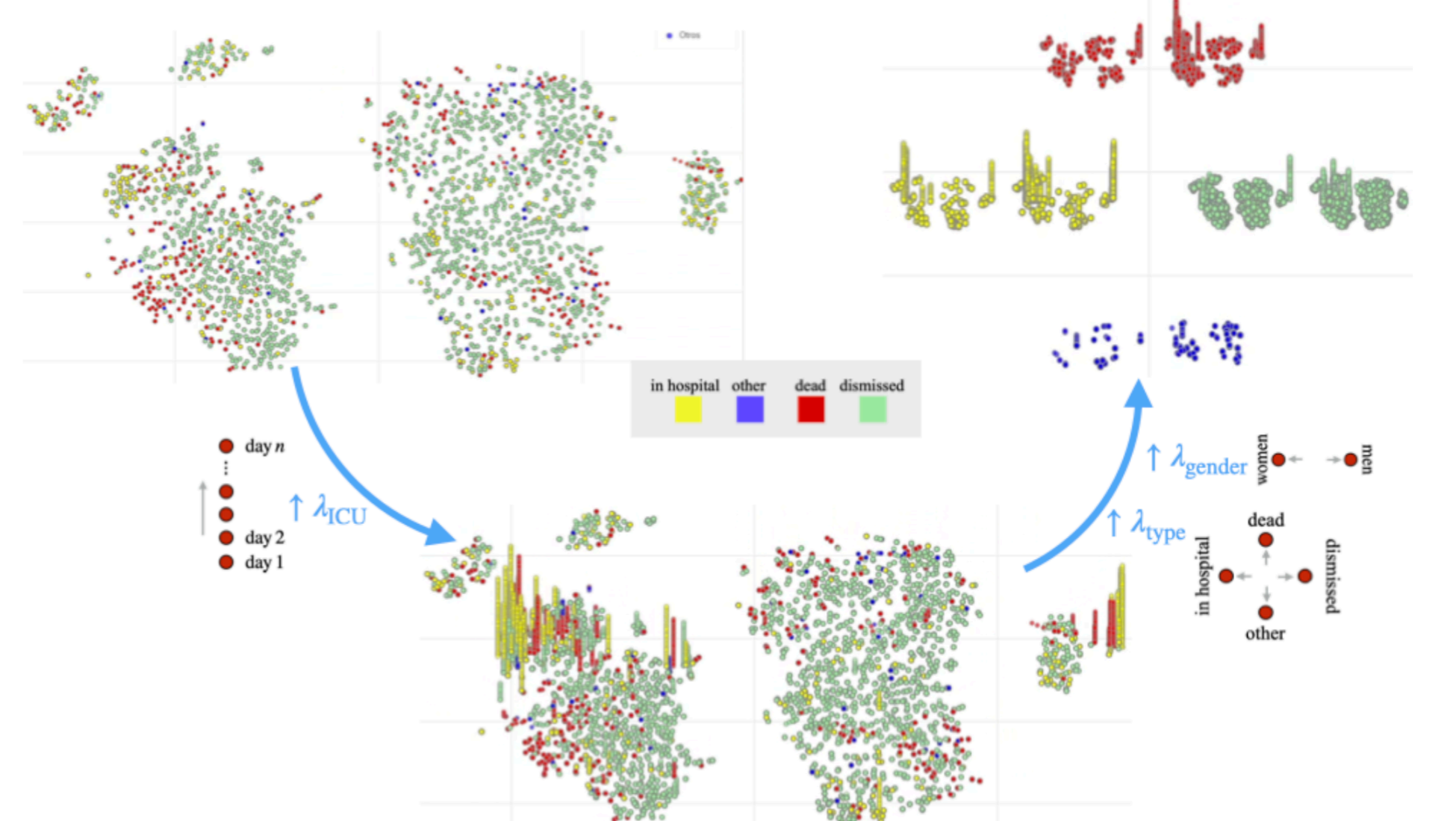
### Analysis by evolution (map of timeseries)



### Stratification of patients (status, sex, days in ICU)



### Combined analysis: evolution + days in ICU + status



## Conclusions

- Visual Analytics approach → synergy between computational intelligence and human-based reasoning
- Allows to combine knowledge from multiple domains: time evolution, clinical information (status, sex, other)
- Fluid interaction allows immediate feedback to the user → iterative question/answer exploration
- Tolerant to highly unstructured multifaceted data
- Smooth transitions foster the visual tracking of items, enhancing the discovery of connections between different domains

## Acknowledgments



This work is part of Grant PID2020-115401GB-I00 funded by MCIN/AEI/ 10.13039/501100011033.

The authors also want to thank HM Hospitales for the dataset "covid data saves lives" used in this paper (<https://www.hmhospitales.com/coronavirus/covid-data-save-lives>)

## Demo video

