




# Event2vec, a python package for medical concept embeddings study

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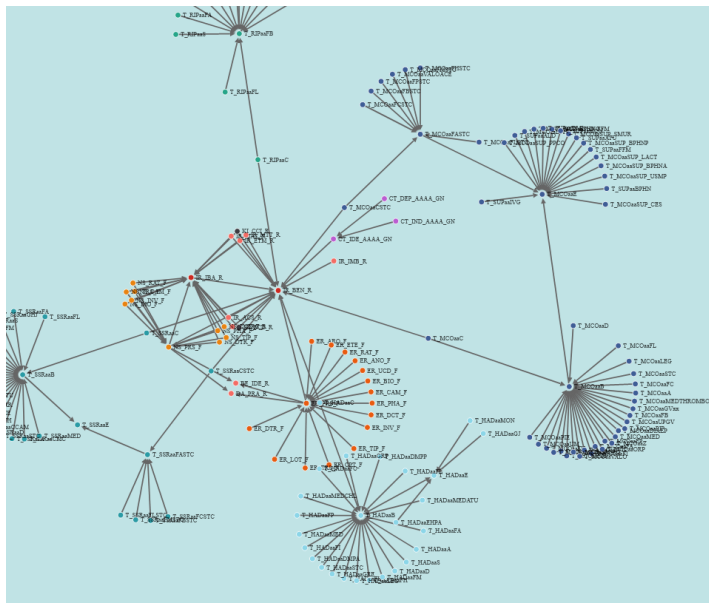
# I. Context and motivations

## II. Medical concept embeddings from structured events

## III. Qualitative results

## IV. Empirical evaluations

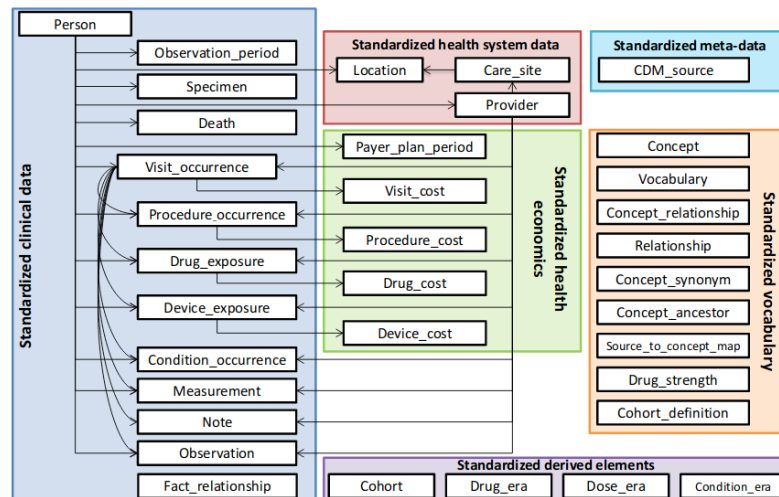
# Large observational structured databases



**Medico-administrative database (claims) :**  
ex. [SNDS](#)  
Care consumptions, reimbursements

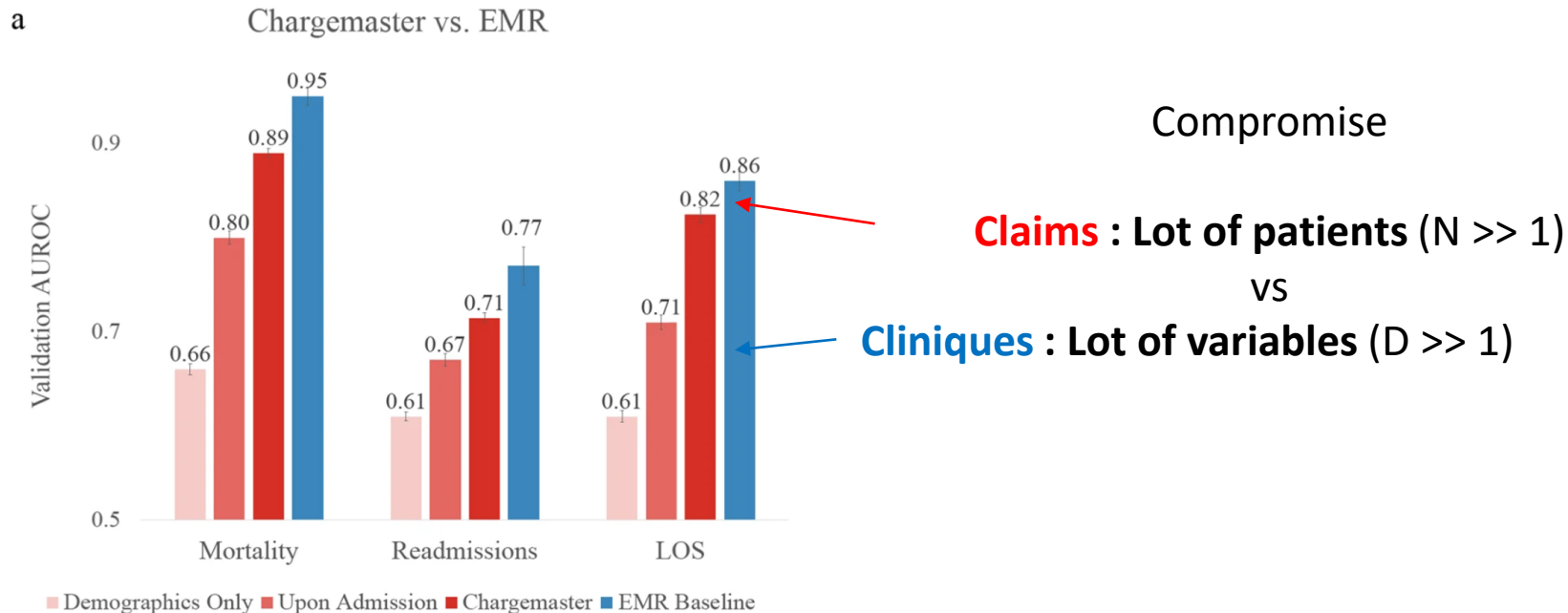


## CDM Version 5 Key Domains



**Electronic Health Records (EHR/EMR):**  
ex. [APHP](#) data warehouse  
Detailed clinical variables, medical reports, ...

# Despite the lack of precise endpoints, claims contain information



(Beaulieu-jones et al, 2021)

Performances of predictive models taking as inputs **claims (chargemaster)** or **Electronic Medical Records (EMR)**

# Patient trajectories : timestamped collection of tokens

Multiple applications of ML in healthcare consider a **triplet event format**

(Rajkomar et al., 2018; Beam et al., 2019; Bacry et al., 2020; Chazard et al., 2022)

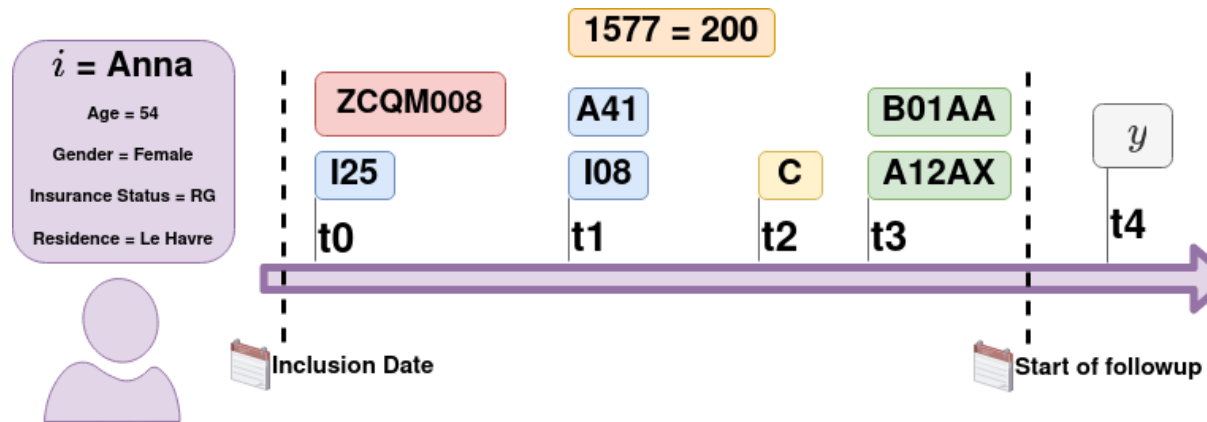
## 👍 Advantages

- **Simple**
- **Sequential**
- **Comparability** of all type of healthcare information

## 🗨️ Difficulties

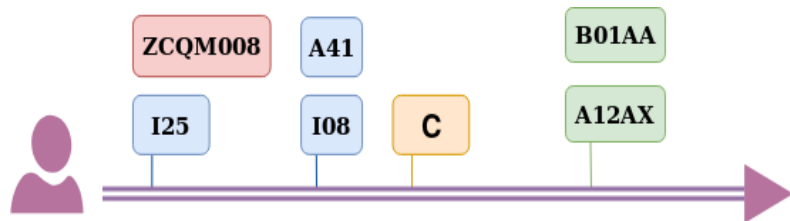
- **High cardinality** of codes
- Choices of **aggregation** for statistical models

$$e = (i, t, c)$$

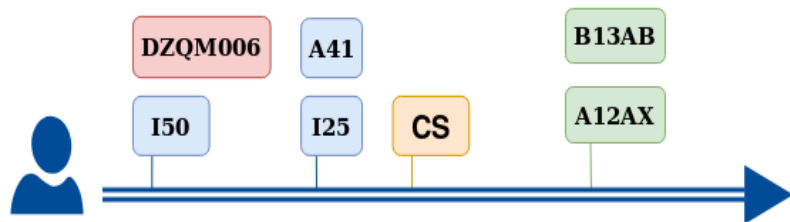


# Patient trajectories: How to derive proximity of

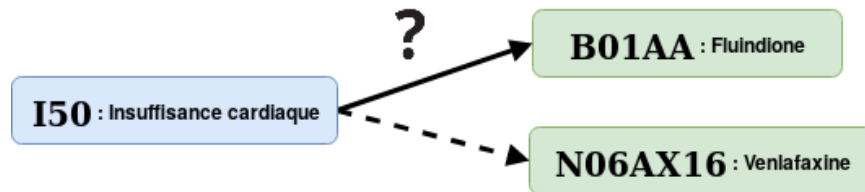
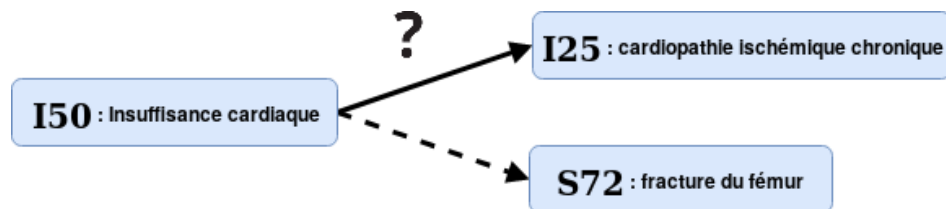
## Trajectories ?



? =



## Concepts ?



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# Medical embeddings of structured data, previous work

## First concept representations algorithms

- [Tran et al., 2015](#): nonnegative restricted boltzmann machines for **suicide-prediction models**
- [Miotto et al., 2016, deepatient](#): Auto Encoder for **78 disease onsets** prediction
- [Nguyen et al. 2016, deepr](#): CNN for deep patient representation and **unplanned readmission**
- [Choi et al., 2016, med2vec](#): MLP for visits and medical codes, for **next visit billing codes** prediction

## Inclusion of time

- [Cai et al., 2019, CBOWA](#): Build a time-aware context window, evaluate on **clustering tasks**
- [Beam et al., 2019, cui2vec](#): Implement context aware svd-ppmi, evaluate on **known associations detection**
- [Xiang et al., 2019](#): extend Beam's algorithm to fastText, applied to **onset prediction of heart failure** (w. LSTM)

## Transformer-based models

- [Rasmy et al., 2021, MedBert](#): Transformers for **heart failure for diabetes patients (DHF)** and **pancreatic cancer prediction**
- [Solares et al., 2020, BEHRT](#): Transformers for **301 diseases predictions** in future visits

## A review paper with benchmarks

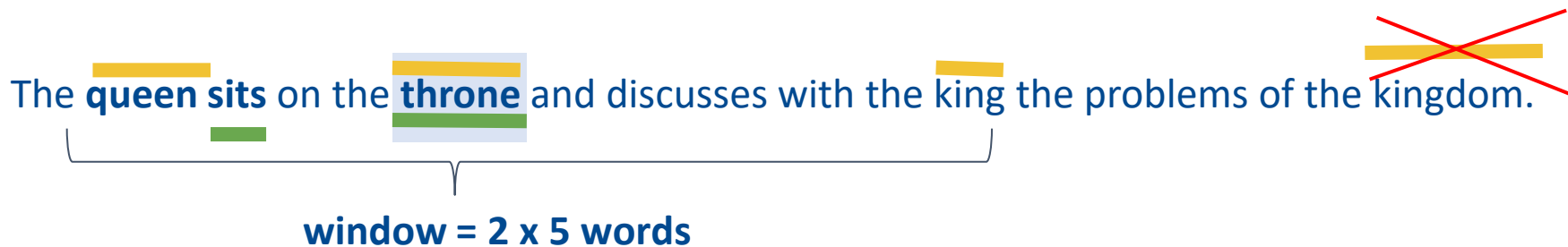
[Solares et al., 2021, Transfer Learning in Electronic Health Records through Clinical Concept Embedding](#)



# Inspiration: back to basics: word2vec in NLP

**Distributional hypothesis** (*Firth, 1957*): Two words are close iff they appear in similar contexts:

*"You shall know a word by the company it keeps"*



**Proximity in the embedding space** is forced by **proximity in the corpus**.

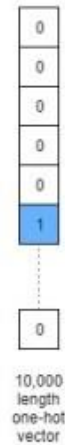
# Focus on a context window approach

- **SGNS (word2vec):** Prediction of the context given a word thanks to a one-layer neural network

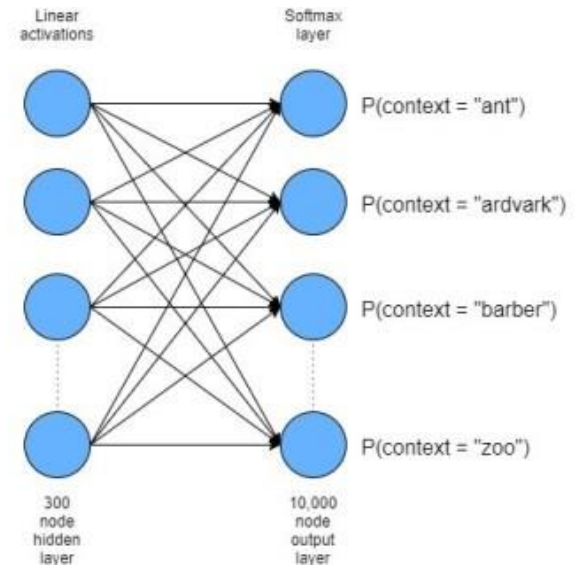
$$\log \sigma(\vec{w} \cdot \vec{c}) + k \cdot \mathbb{E}_{c_N \sim P_D} [\log \sigma(-\vec{w} \cdot \vec{c}_N)]$$

positive example

negative example

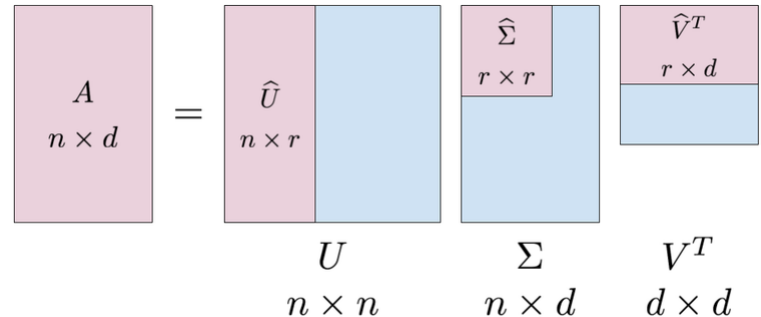


10,000 length one-hot vector

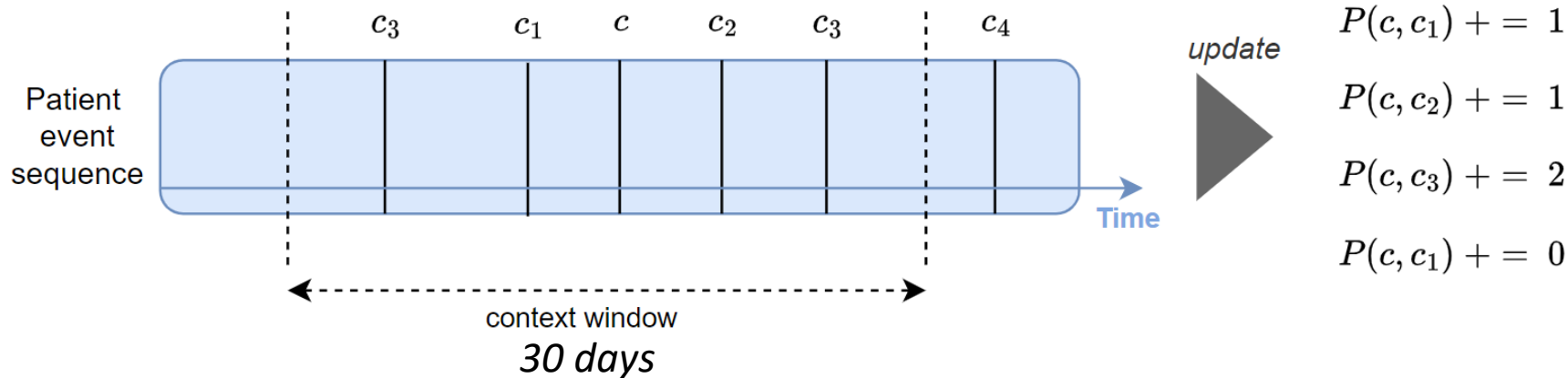


- **SVD-PPMI:** Singular vector decomposition of the transformed word co-occurrence matrix

$$\Phi(w) = \frac{1}{2}(U_d \cdot \sqrt{\Sigma_d} + (V_d \cdot \sqrt{\Sigma_d}))$$






# Adapting word2vec to patient trajectory *(Beam et al., 2019)*




 Build a **time dependant context** for the co-occurrence matrix  $P(c_i, c_j)$

# Why concept embeddings could be interesting ?

## Objectives

-  Predictive and interpolation models (*cf. preceding review slide* →)
-  Treatment effects estimation thanks to G-formula (*Dorie et al., 2018, Wendling et al., 2018*)
-  Vocabulary matching

## Advantages

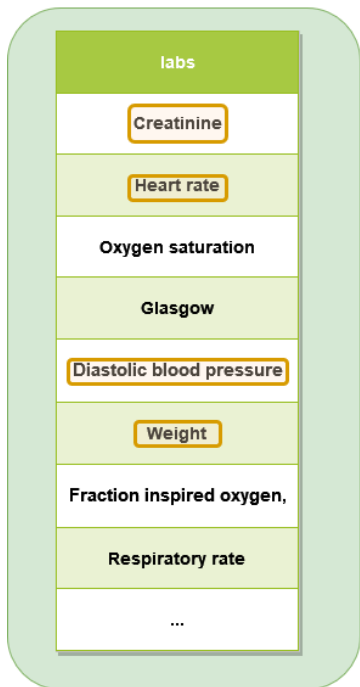
-  **Sharable** aggregated information
- Fewer **Hyper-parameter tuning**
- **Simple implementation** pandas + scipy
- **CPU only** easilly scalable w. distributed backend
- **No softmax** computation

## Difficulties

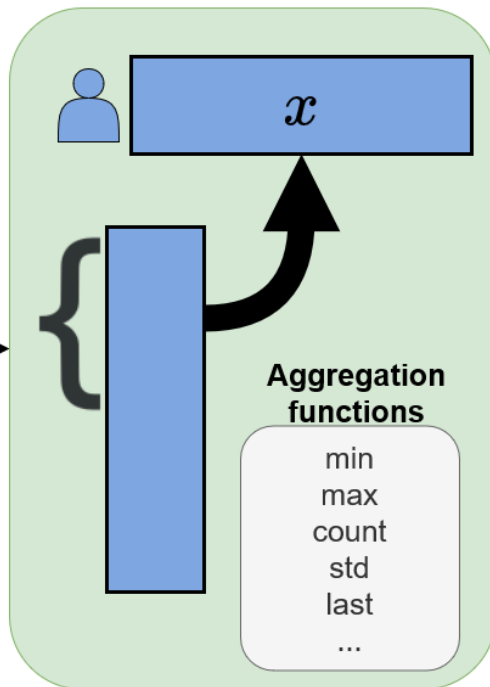
- **Poor in-context** comprehension
- Different **choices of aggregation** for visit modelizations

# Des choix multiples de modélisation

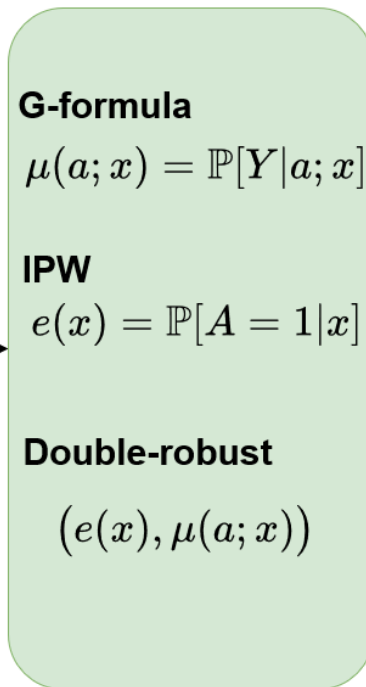
## Variables selection



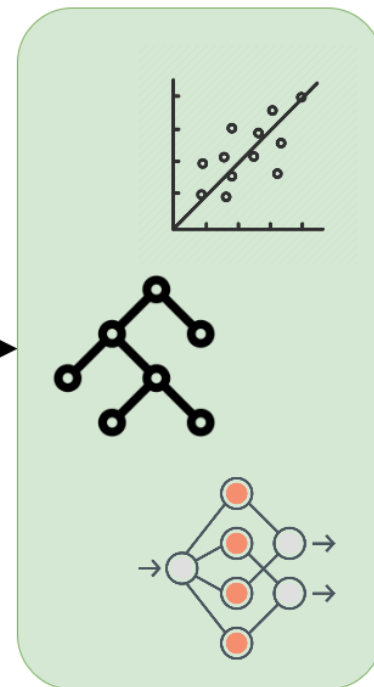
## Feature extraction



## Identification



## Estimation



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**I. Context and motivations**

**II. Medical concept embeddings from structured events**

**III. Demo and qualitative results**

**IV. Empirical evaluations** 

# Event2vec, a package to easily compute concept embeddings

 A python package available on pypi

 A pyspark version for big data (>500m rows)

 Quick start and step by step guides:

[https://straymat.gitlab.io/event2vec/tutorials/0\\_tuto\\_event2vec.html](https://straymat.gitlab.io/event2vec/tutorials/0_tuto_event2vec.html)

## Load events

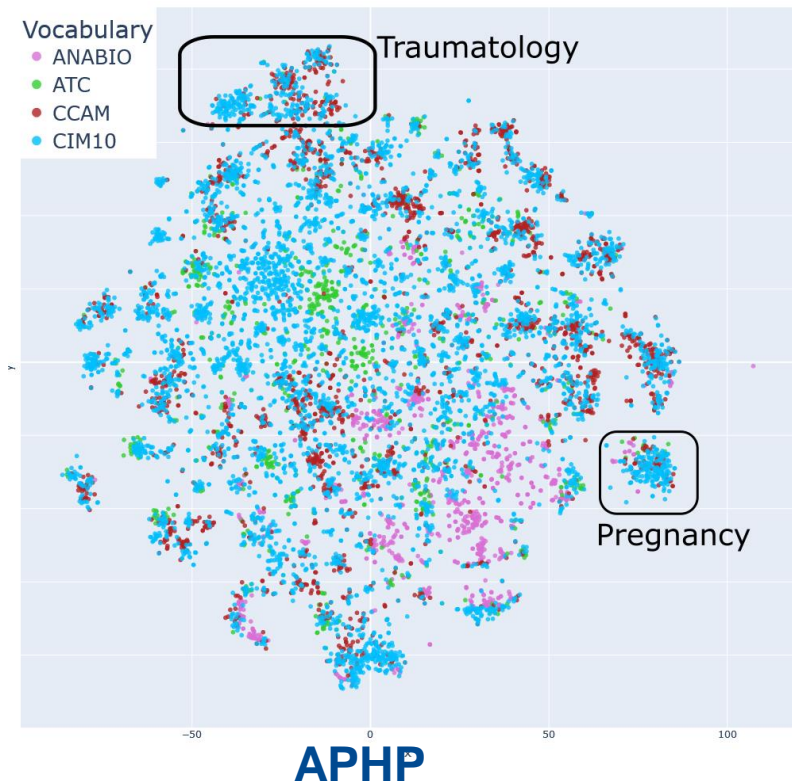
	person_id	start	event_source_concept_id
0	1	2018-11-08 19:24:15	CIM10:N182
4	1	2018-12-20 19:24:15	CCAM:JVJB01
8	2	1993-01-26 07:22:42	CIM10:E12
12	3	2009-04-25 10:14:21	CIM10:N182
9	2	2020-01-26 07:22:42	CIM10:E12

## Build embeddings

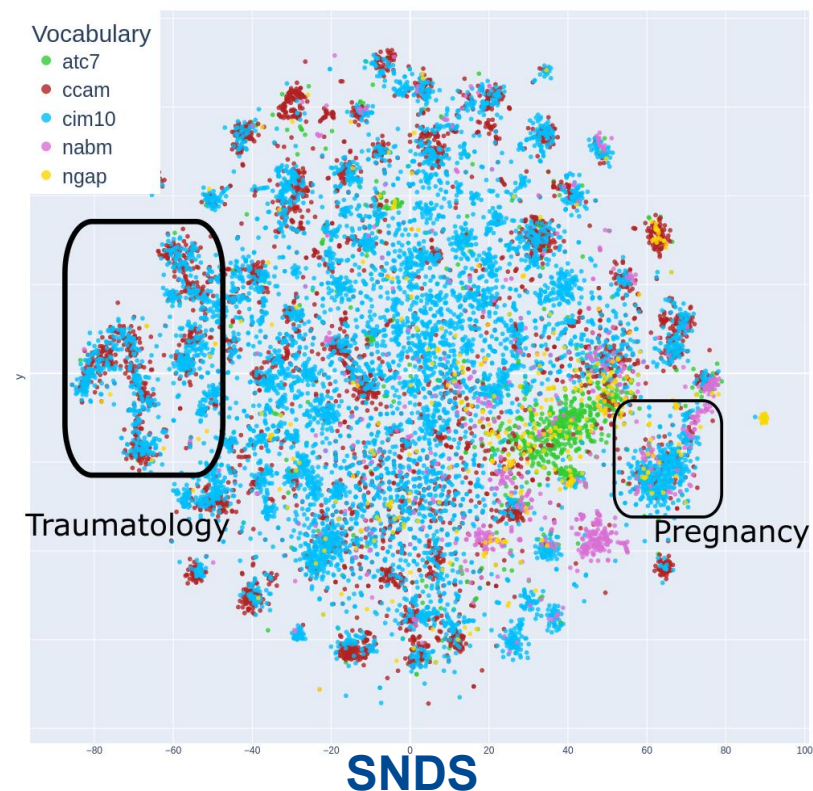
```
alpha = 0.75
k = 1
d = 3

embeddings = event2vec(
    events=events,
    output_dir=output_dir,
    colname_concept="event_source_concept_id",
    window_orientation="center",
    window_radius_in_days=30,
    d=d,
    smoothing_factor=alpha,
    k=k,
    backend="pandas",
)
```

# Qualitative results: <https://straymat.gitlab.io/event2vec/visualizations.html>



**APHP**  
**(200K random patients)**



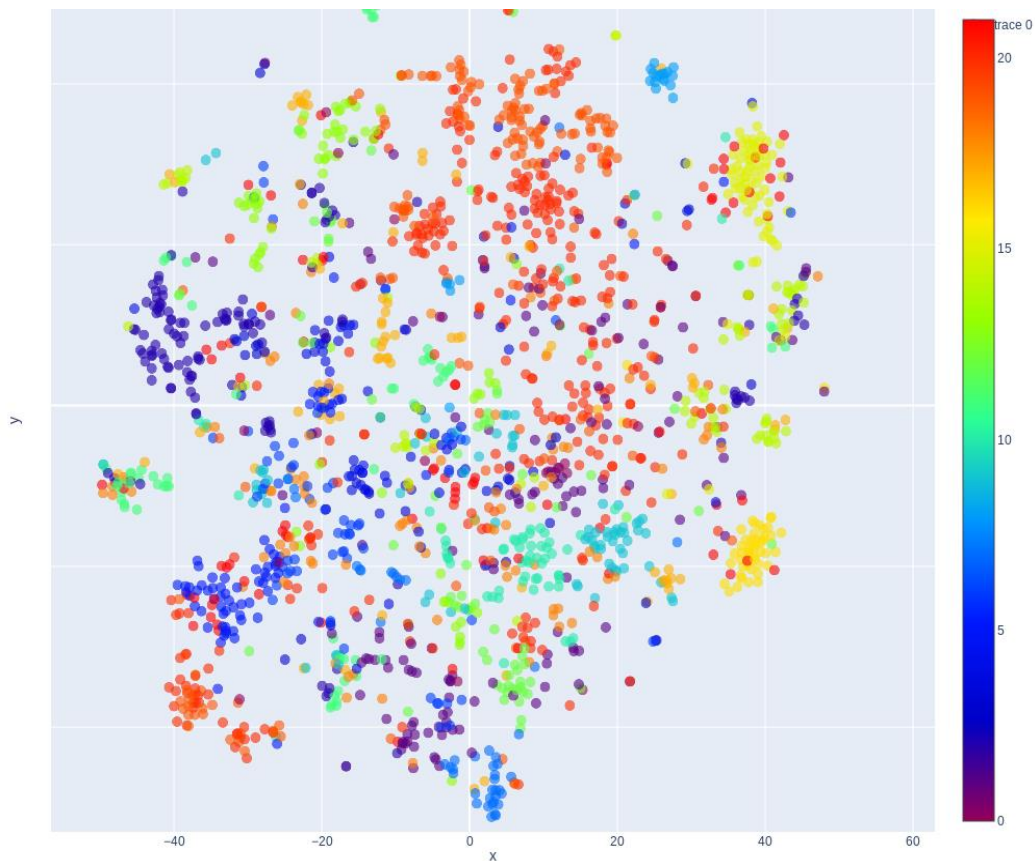
**SNDS**  
**(3M random patients)**



# Qualitative results, Hierarchy reconstruction

CIM10 billing diagnoses  
Third level,  $r=30$  jours

Colored by chapter



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I. Context and motivations

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# Extrinsic evaluation: Compare different models on a downstream task

 **Task:** rehospitalization at 30 days, for planning and outcome modeling (g-estimation)

 **Models = (featurizer, estimator):**

**Compared featurizers:**

**Count vectorizer (+SVD, D=30)**

$[C, C_{decay}]$

**Embeddings fit on train data**

$[C \cdot \Phi_{train}, C_{decay} \cdot \Phi_{train}]$

**SNDS Embeddings (+SVD, D=30)**

$[C \cdot \Phi_{SNDS}, C_{decay} \cdot \Phi_{SNDS}]$

**Compared estimators:**

Random forests, Ridge classifier

*V concept codes*

	0	0	2	0	4
	0	1	0	0	0
	2	0	0	2	0
<i>N patients</i>	0	0	2	0	0
	1	0	0	0	0
	0	1	1	4	1
	0	0	0	1	0

**Sparse count matrix  $C$**

*D dimensions*

	0.251	0.124	0.871
	0.635	0.551	0.487
<i>V concept codes</i>	0.251	0.1687	0.152
	0.530	0.251	0.693
	0.07	0.336	0.714

**Embeddings  $\Phi$**

# Population selection for prediction

**Extraction** of 200,000 random patients from APHP EDS

With **complete hospitalization**

**Study period** 2017-2022 (stable Information System)

**Sufficient horizon for followup** (no right censoring)

**Exclusions:** No children, not deceased during hospitalization

At least one event in **4734 codes** occurring at least 10 times:




**drug exposure administrations:** 663, 027 events

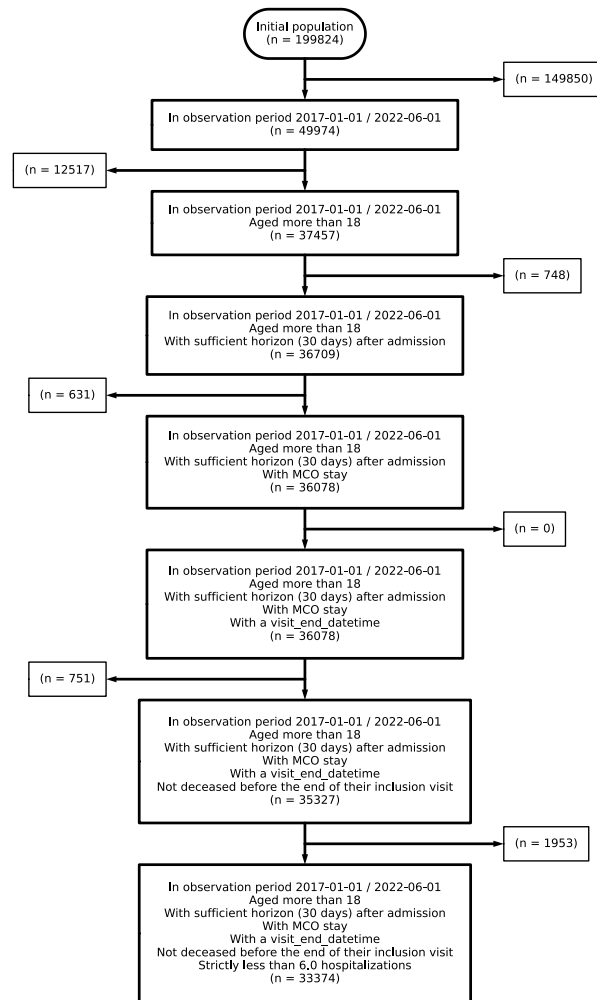


**procedure occurrences:** 222, 770 events

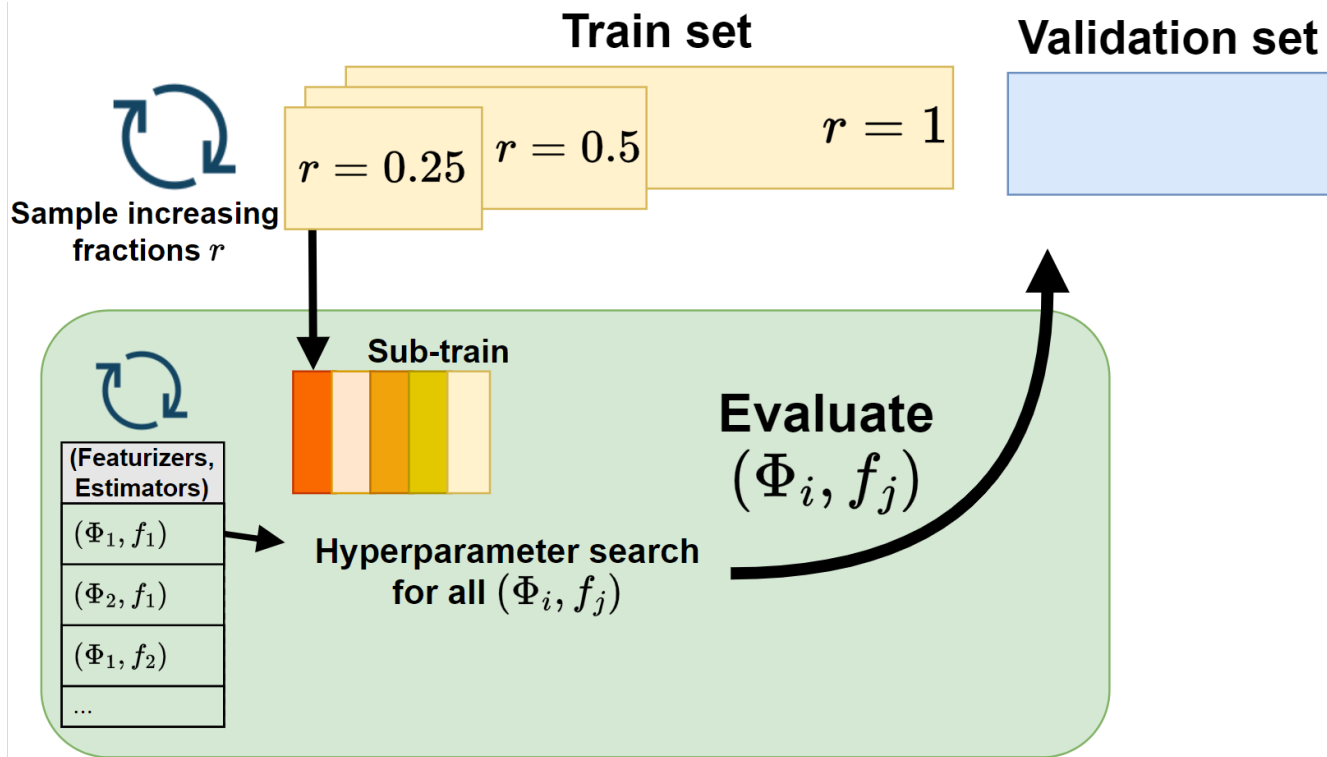


**condition occurrences:** 203, 779 events

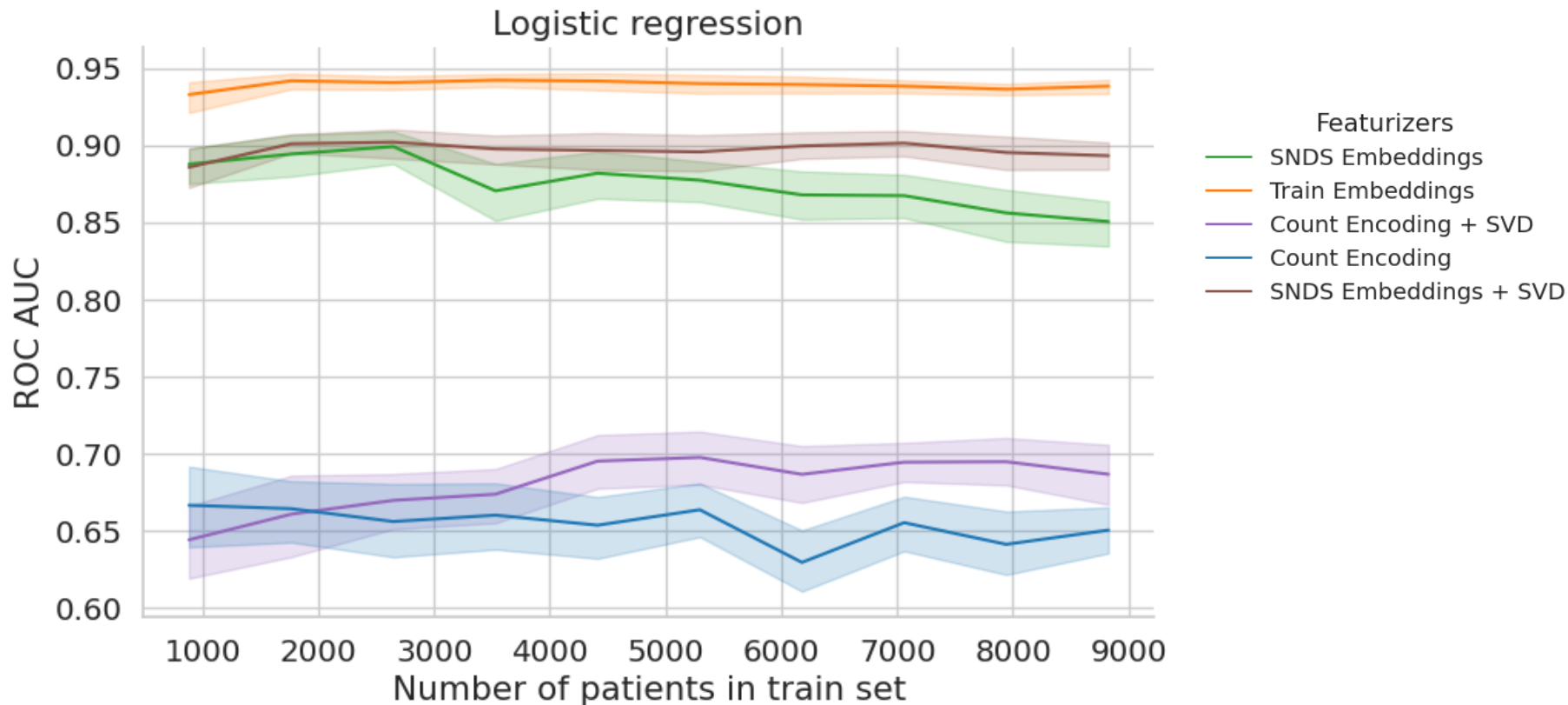
 **25, 063 patients:** mean age = 54.4, female ratio = 54.1%,  
mean LOS > 7 days: 20.80%



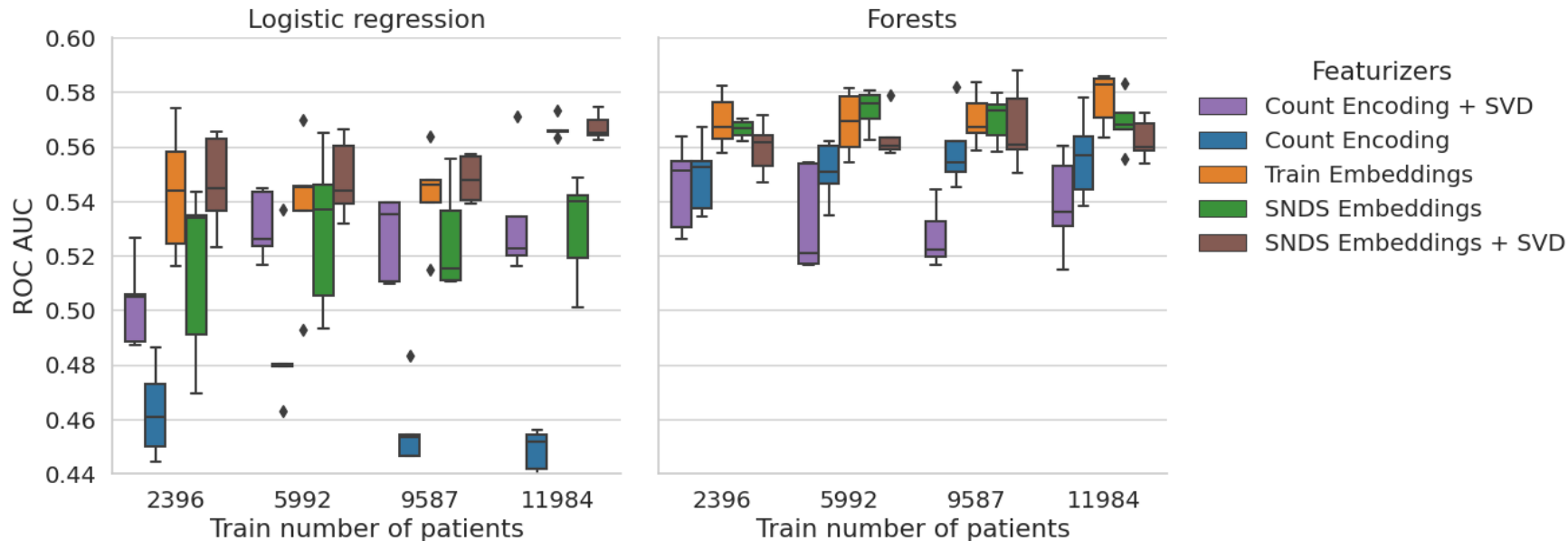
# Selection procedure



# Task: Length Of Stay interpolation : $\leq 7$ days vs. $> 7$ days



# Current results for rehospitalization or death @ 30 days



Overall, **performances are low** (too difficult task ?, badly defined ?)

Logistic regression: **in-domain embeddings are equivalent to SNDS embeddings**

**Forests smooth these differences** (leverages better the missing value mask ?)

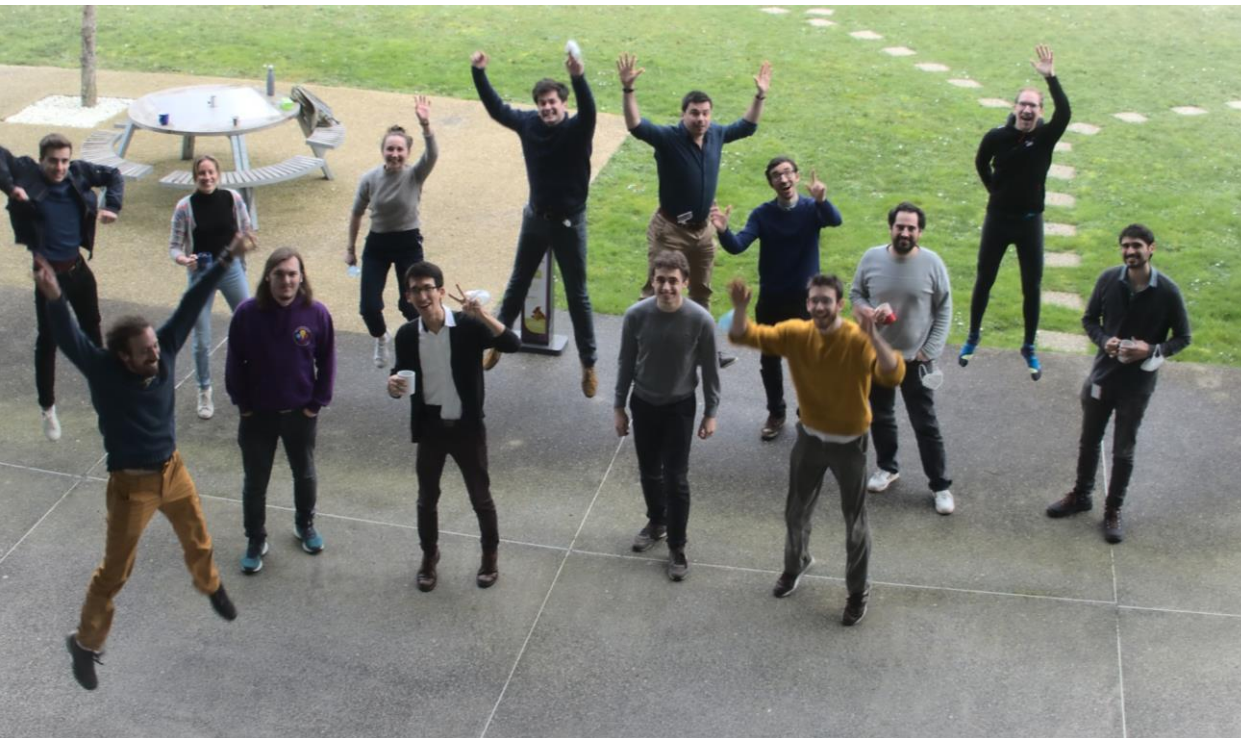
# Further work

- Study **transfer capabilities inside APHP**
- Study **transfer capabilities with international embeddings** such as cui2vec
- Perturbe the learning by dropping some codes
- An **intermediate task** to study predictive performance:  
*mortality prediction ? Disease onset ? Computational phenotyping ?*
- Evaluation for **concept proximity** : eg. [eds-scikit biology concepts as ground truth](#)

## Collaborations ?

- Better **inclusion of temporality** with **transformer-based models**
- Transfer **from APHP to SNDS ?**





## Research axes

### Representation learning for heterogeneous databases

- Learning despite database normalization errors
- Tabular deep learning

### Health and Social Sciences

- Electronic health records
- Epidemiological cohorts
- Educational data mining

### Data-science with statistical learning

- Statistical learning with missing values
- Machine learning for causal inference

### Turn-key machine-learning tools for socio-economic impact

Helping to maintain and grow tools such as [scikit-learn](#), [joblib](#)...

## HAS, mission Data



1. Données produites par la HAS



2. Données observationnelles



3. Connaissances textuelles



4. Organisation





# Supplementary slides

# Skip-gram with Negative Sampling

- Given (word, context) = (w, c) pairs, and random representation in the embedding space:

$$\vec{w} \in \mathbb{R}^{V_w \times d}$$

$$\vec{c} \in \mathbb{R}^{V_c \times d}$$

- The probability of occurrence of a pair (w, c) is given by:

$$\mathbb{P}(D = 1 | w, c) = \sigma(\vec{w} \cdot \vec{c}) = \frac{1}{1 + e^{-\vec{w} \cdot \vec{c}}}$$

- We maximize for a pair:  $\log \sigma(\vec{w} \cdot \vec{c}) + k \cdot \mathbb{E}_{c_N \sim P_D} [\log \sigma(-\vec{w} \cdot \vec{c})]$

- On the whole corpus:

$$P_D(c) = \frac{\#(c)}{|D|}$$

$$l = \sum_{w \in V_w} \sum_{c \in V_c} \#(w, c) (\log \sigma(\vec{w} \cdot \vec{c}) + k \cdot \mathbb{E}_{c_N \sim P_D} [\log \sigma(-\vec{w} \cdot \vec{c})])$$

# SVD-PPMI as the solution of the SGNS objective

Given the **Pointwise Mutual Information matrix**:  $PMI(w, c) = \log \frac{P(w, c)}{P(w)^\alpha P(c)^\alpha}$

Rewrite **Cooccurrence** as:  $PMI(w, c) = \log \frac{\#(w, c) \cdot |D|^{2\alpha - 1}}{\#(w) \cdot \#(c)}$

Enforce **sparsity**:  $PPMI(w, c) = \max(PMI(w, c), 0)$

**Factorization**:  $SPPMI(w, c) = U_d \Sigma_d V_d$

**Dense representations** as singular components:  $\Phi(w) = \frac{1}{2} (U_d \cdot \sqrt{\Sigma_d} + (V_d \cdot \sqrt{\Sigma_d}))$

## Theoretical arguments in favor of Glove model *(Pennington et al., 2014)*

$$\sum_{i,j} f(X_{ij}) (w_i^T \tilde{w}_j - \log X_{ij})^2,$$

- Offline (like SVD-PPMI)
- Avoid high cost of softmax (computation of normalization functions)
- No cross-entropy error (model poorly long tail distributions)

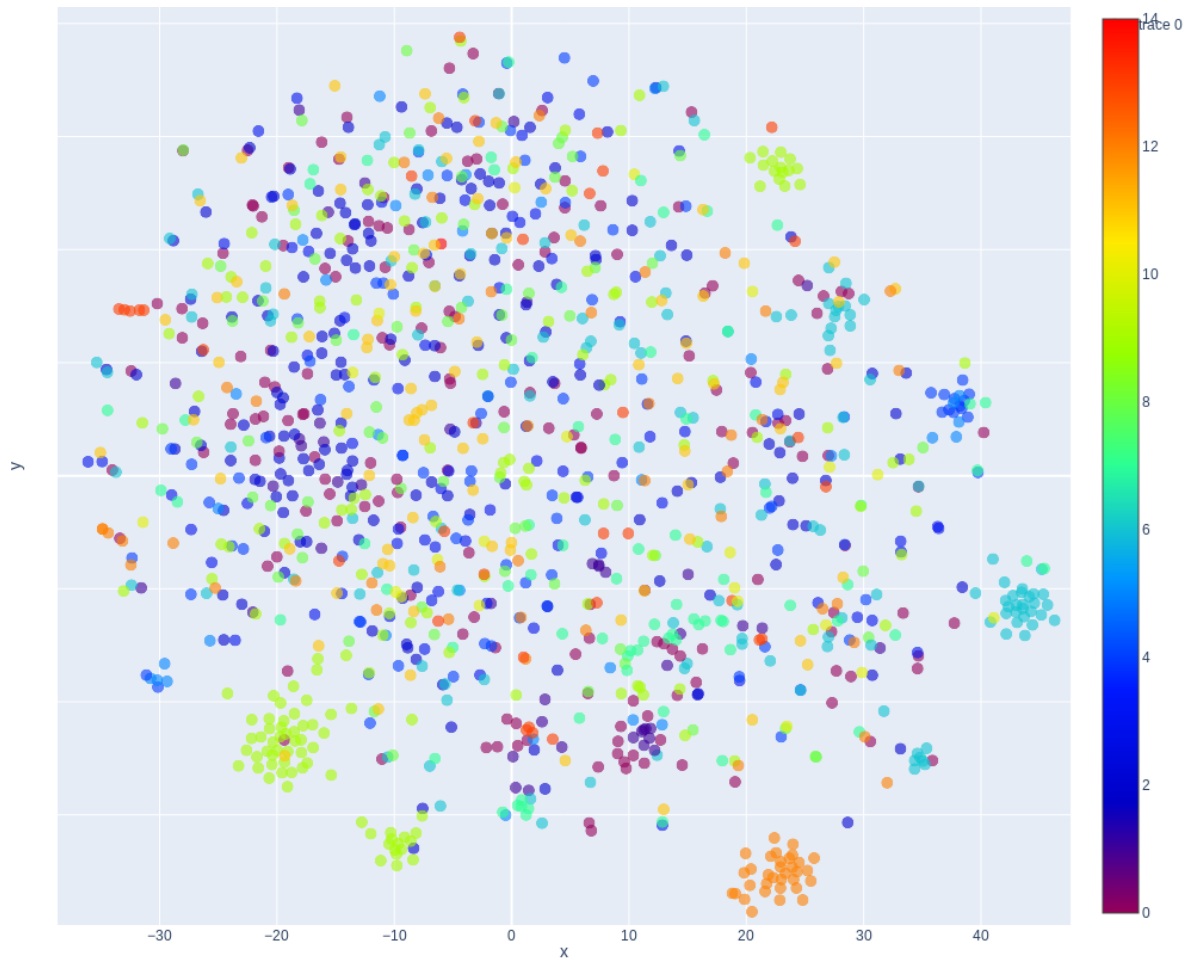
# SNDS, details on data

- **Extraction:** Sample of three millions of patients followed 9 years
- **Sources:** DCIR (assurance maladie), PMSI (hospital billing codes) MCO, MCO\_CE, SSR, SSR\_CE, HAD
- **Events:** CIM10 (diagnostics), CCAM procedures (outpatient, inpatient, city care), city drugs, city biology
- **Granularity of codes:** ATC 7, CIM10 (4 characters), CCAM (7 characters), biology (4 characters) -> **15968 codes**
- **4416 codes** in common with APHP study cohort for rehospitalization@90 days

# Quantitative results

ATC drug codes,  $r=30$  days

Colored by chapter

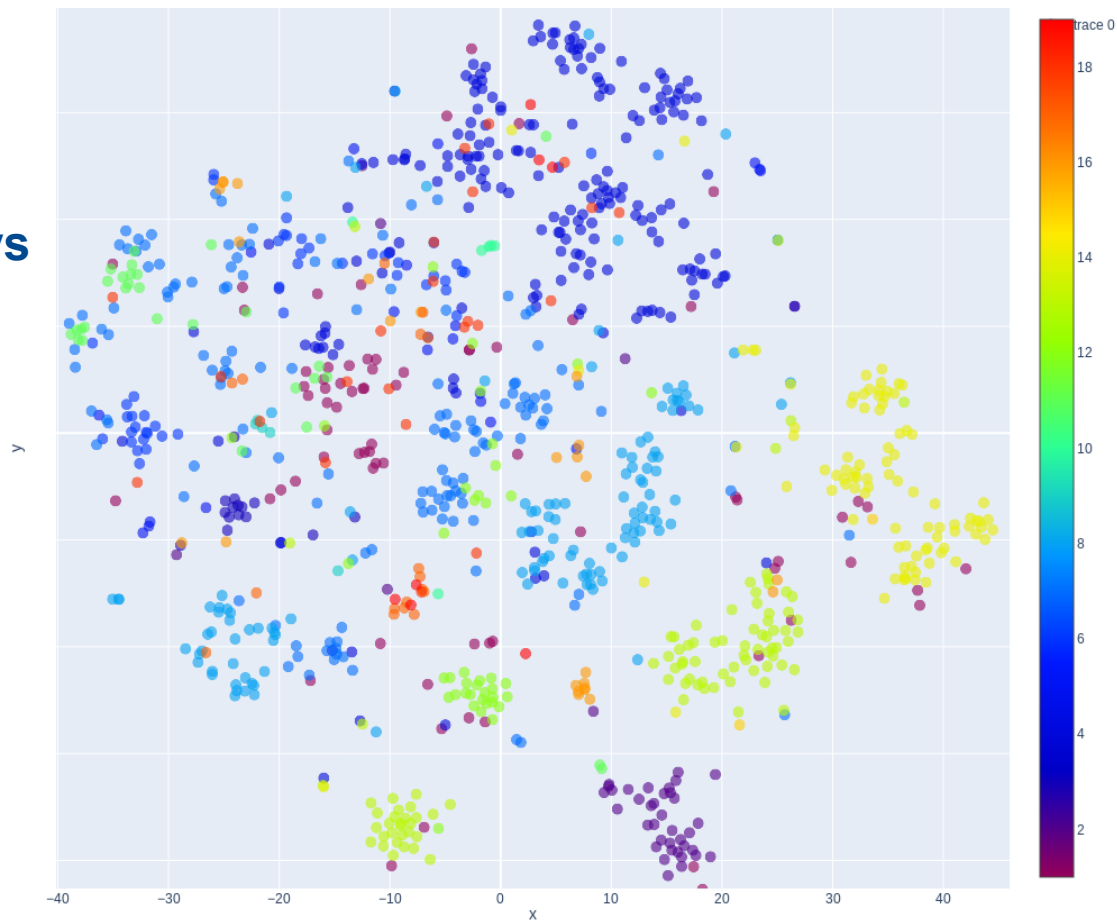




# Qualitative results

CCAM billing procedures  $r=30$  days

Colored by chapter



# Benchmarks for intrinsic evaluation

**Metrics** (Beam et al., 2019):

- **FMI**: quality of clustering
- **Medical Relatedness Measure**: How many close neighbors in the same hierarchical category ?
- **NDF-RT may treat / may prevent**
- **UMLS causative relationship**

granularity	radius	orientation	icd10	ccam	atc
L	30	centered	0.3818	<i>0.489</i>	0.2237
		future	<i>0.3878</i>	0.4821	0.1535
	90	centered	0.3677	0.4873	<b>0.3412</b>
	F	30	centered	0.4533	0.5159
future			0.4465	0.4997	0.1368
90		centered	<b>0.456</b>	<b>0.5306</b>	<i>0.3159</i>
		future	0.4528	0.5106	0.2376

Table 3: MRM computed on the embeddings grouped per terminology hierarchy. The best score is given in bold and the best score per granularity is given in italics.

# Population selection for prediction

**Extraction** of 200,000 random patients from APHP EDS

**Study period** 2017-2022 (stable Information System)

**Sufficient horizon for followup** (no right censure)

**Exclusions:** No children, not deceased during hospitalization

At least one event in **4923 codes** occurring at least 10 times:




**drug exposure administrations:** 608, 577 events

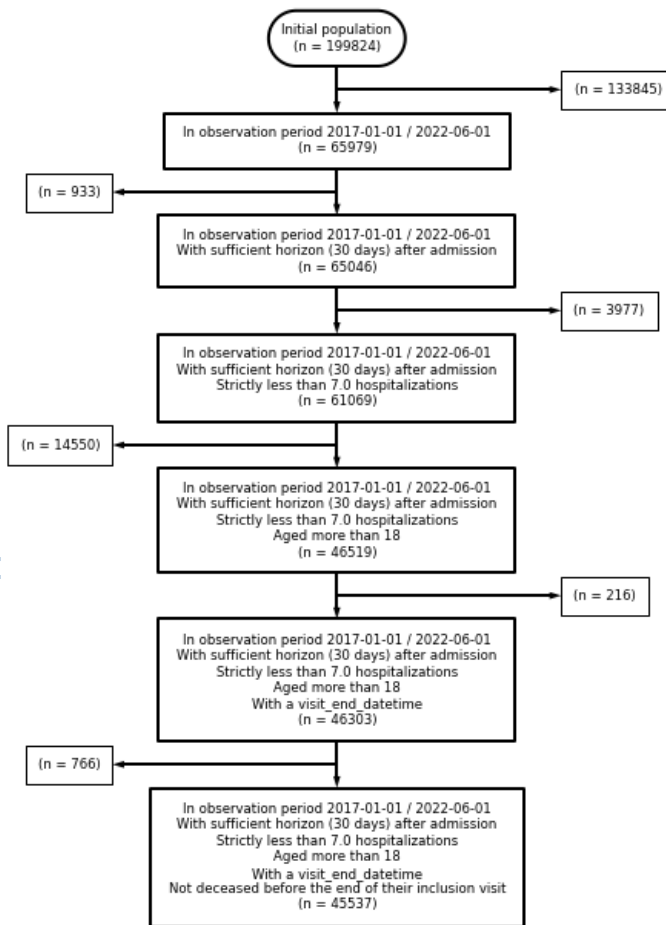


**procedure occurrences:** 252, 668 events



**condition occurrences:** 219, 666 events

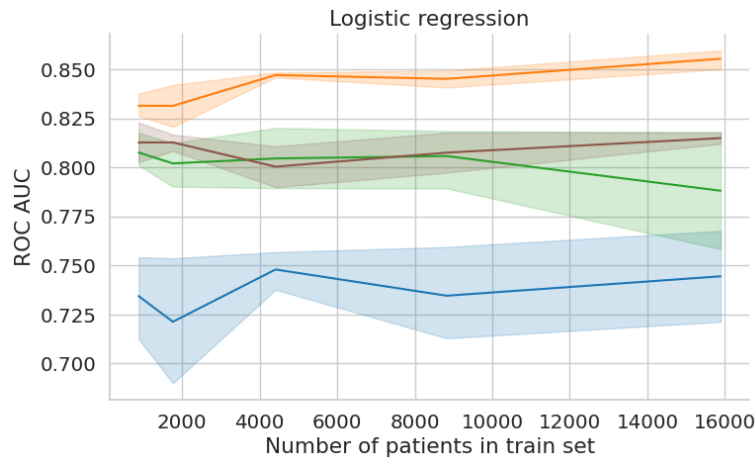
 **34, 063 patients:** mean age = 54.4, female ratio = 55.8%,  
mean rehospitalization@30d=10.5%



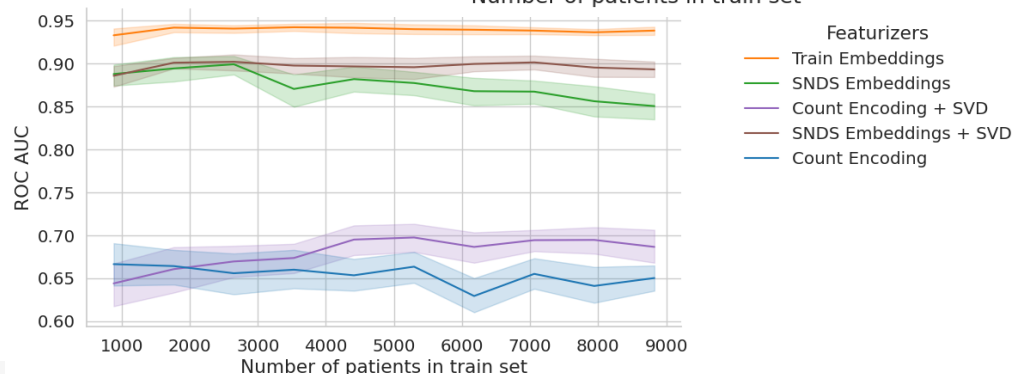
# Task: Length Of Stay interpolation : $\leq 7$ days vs. $> 7$ days

Adding a simple temporality decay seems very efficient.

Wo temporality decay

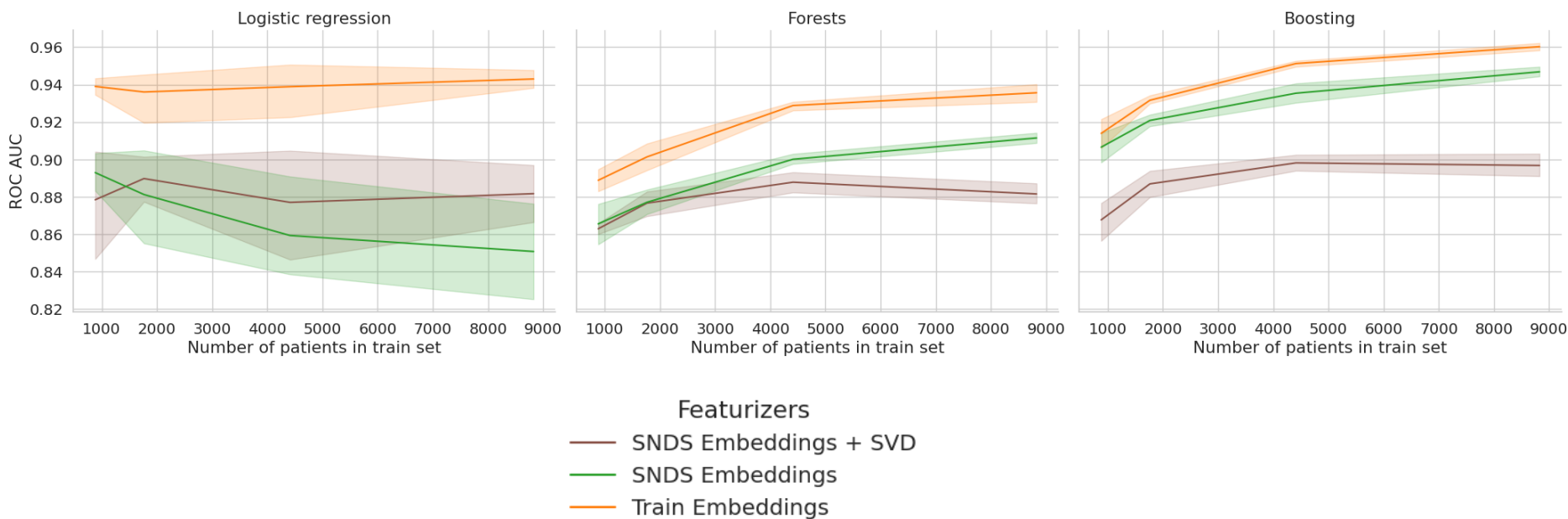


W temporality decay



# Task: Length Of Stay interpolation : $\leq 7$ days vs. $> 7$ days

There is sample gains at least for boosting and forests (no saturation of the task)



# References

## Event format:

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# References

Embeddings for for predictive tasks

Ref. slides 8