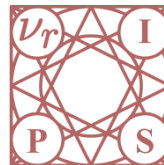




Attentive State-Space Modeling of Disease Progression

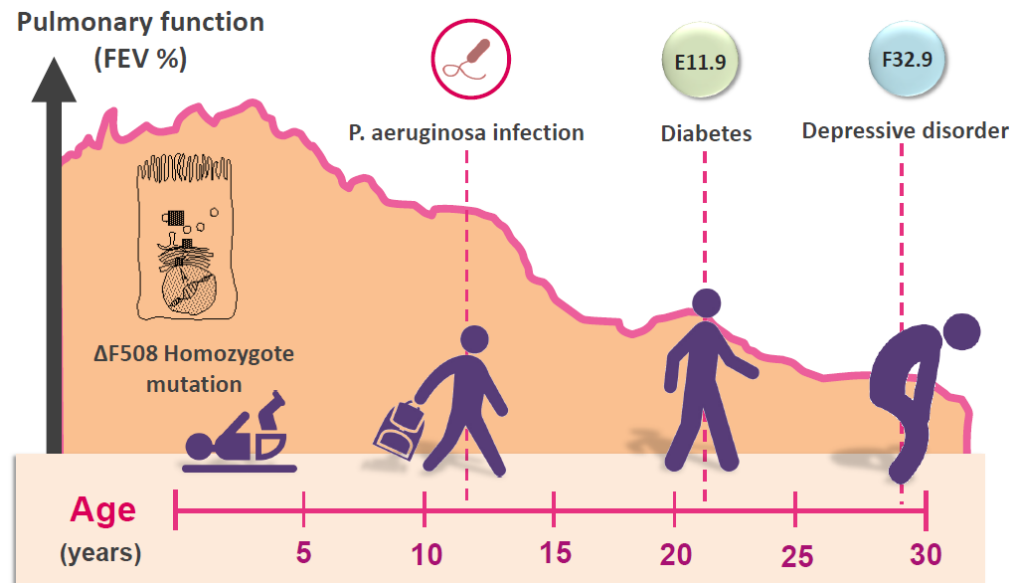
Ahmed M. Alaa
Mihaela van der Schaar



Disease Trajectory Modeling

- Cross-sectional modeling: Simple prognostic questions (how long will a patient *survive*?) are not enough!
- Longitudinal modeling: addresses a more adequate question: *what will happen to the patient in future?*

Inform patient carers
about the patient's
future needs and
**interactions with health
services**

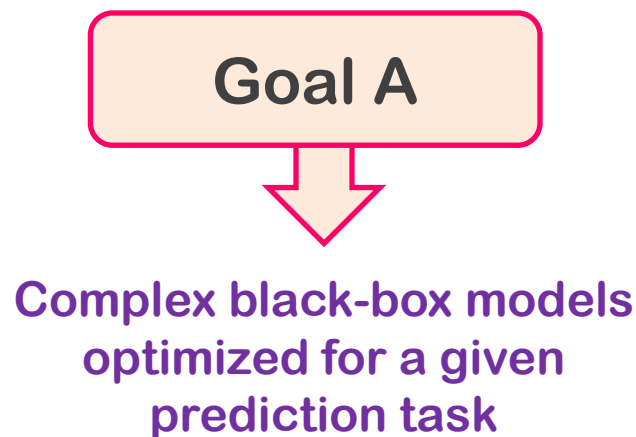


Two Central Goals of Longitudinal Models

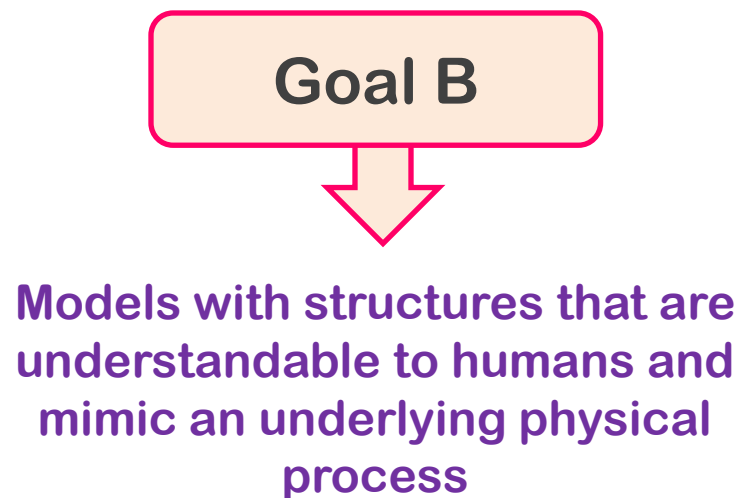
- Goal A: **Predicting** individual-level disease trajectories
 - Early diagnosis
 - What are the risks of mortality, comorbidities, complications, infections in the future?
- Goal B: **Understanding** disease progression mechanisms.
 - Underlying latent structure of disease evolution
 - Causal pathways and comorbidity networks
 - Patients' subgroup analysis
 - Refined phenotypes and ontologies

Accuracy-interpretability Trade-off

- **Goals A and B** have very **different** (and **conflicting**) modeling requirements!



Example: Recurrent neural networks (RNNs)

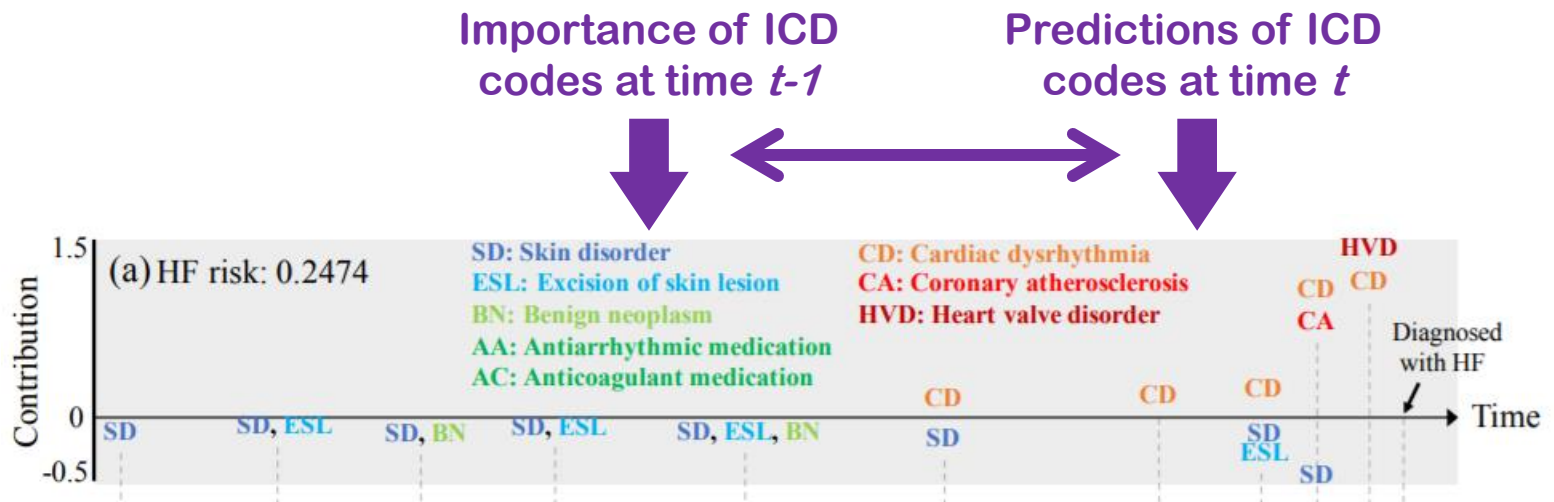


Example: Hidden Markov model (HMM)

Different Forms of Interpretability

● Form A: Instance-wise variable importance

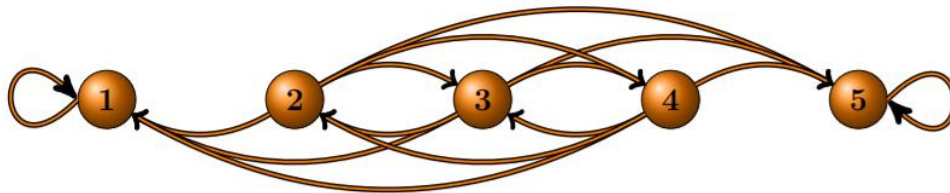
- For every patient: identify important variables for future predictions at every time step based on the patient's history.
- Useful for transparency and actionability but not sufficient (no knowledge extraction).
- Example RETAIN [E. Choi, 2017].



Different Forms of Interpretability

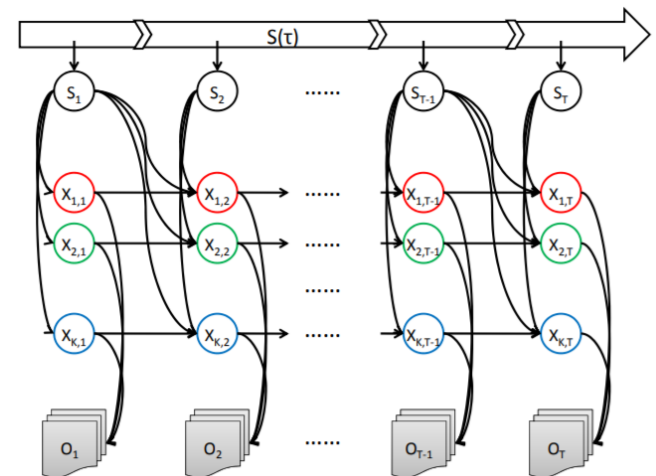
- **Form B: Discovering latent structures in disease progression**
 - Learn a hidden representation of the disease: identify disease states and manifestations of states.

Population-level representation
of disease states



Generative models

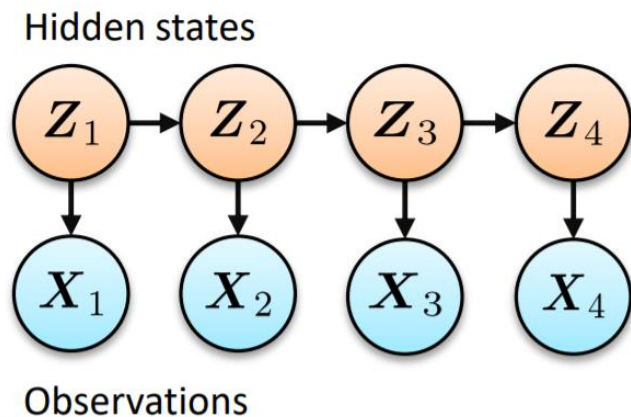
Segmentation and annotation of
individual-level trajectories



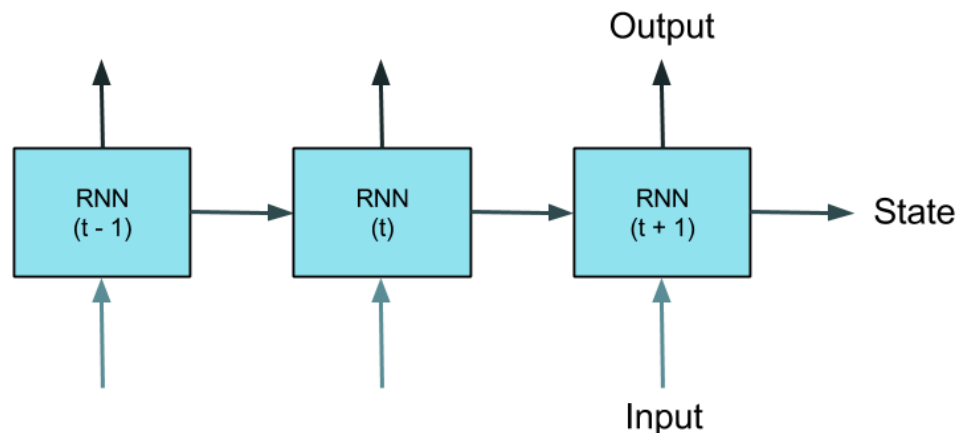
Attentive State-space Modeling

- We want **Form A** and **Form B** + **accurate predictions**
- Our contribution: attentive state-space models
- Main idea:

Maintain probabilistic structure of an HMM

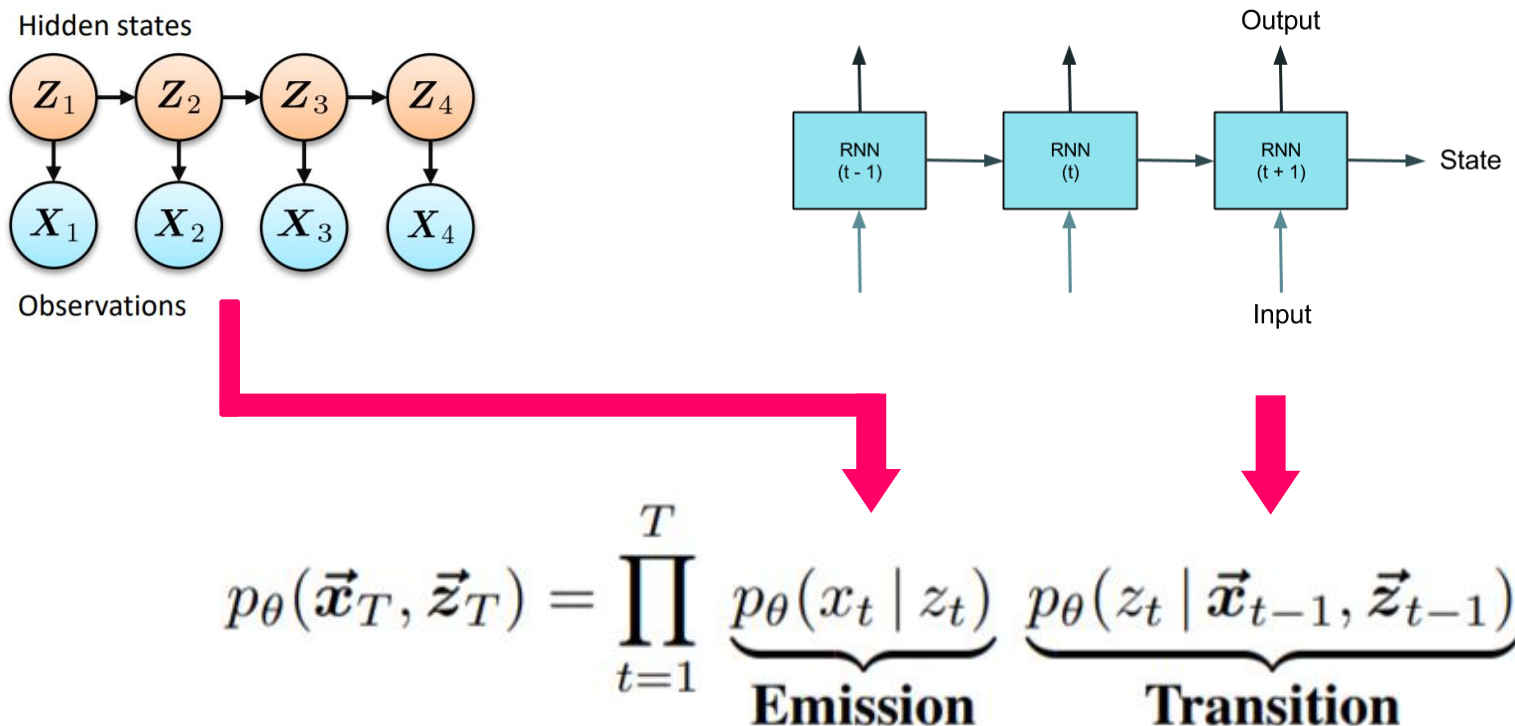


But use an RNN to model the state dynamics



Attentive State-space Modeling

● Attentive state-space models



Contextual Attention

- Use attention mechanism to keep state dynamics interpretable
- Observable variables: (X_1, \dots, X_t)
- Disease states: (Z_1, \dots, Z_t)

Attention weights

Patient context

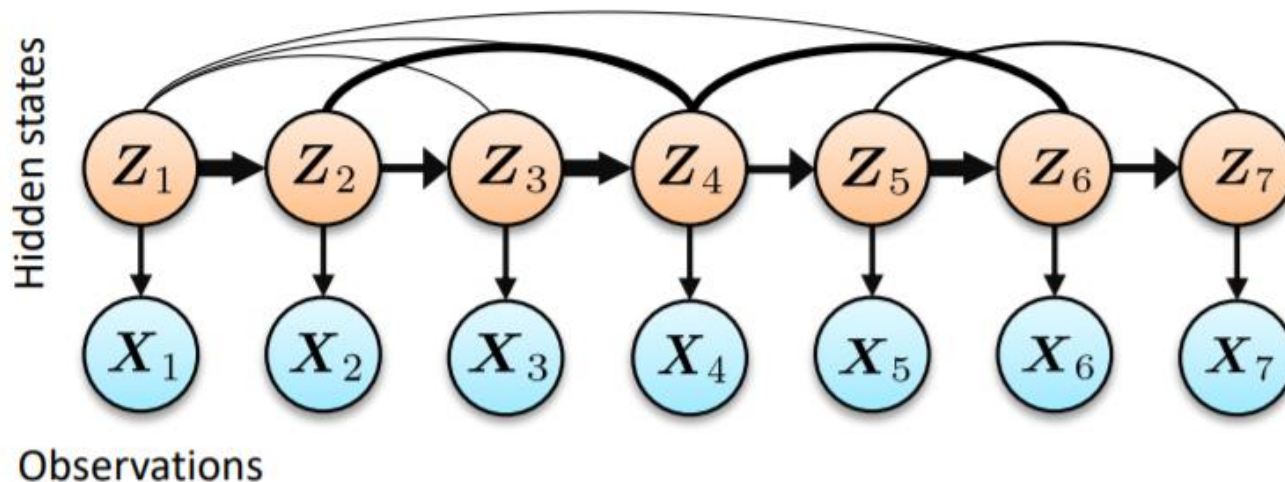
$$(\alpha_1^t, \dots, \alpha_t^t) = \text{RNN}(X_1, \dots, X_t)$$

Baseline Markov chain

$$P(Z_m = z \mid \mathcal{F}_{t_{m-1}}) = P(Z_m = z \mid \underbrace{\{(Z_k = z_k, \alpha_k^m, \Delta_k)\}_{k=1}^{m-1}}_{\text{Sufficient statistics}}) = \sum_{k=1}^{m-1} \underbrace{\alpha_k^m}_{\text{Attention weights}} \underbrace{(e^{\Delta_k \Lambda})_{z, z_k}}_{\text{Transition matrix}}$$

Contextual Attention

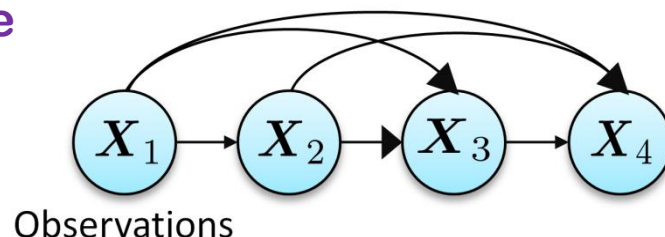
- The attentive state-space model can be interpreted as a **soft version of a variable-order Markov model**
- Memory is shaped by the current patient's context (clinical events, treatments, etc)



Comparison with Existing Models

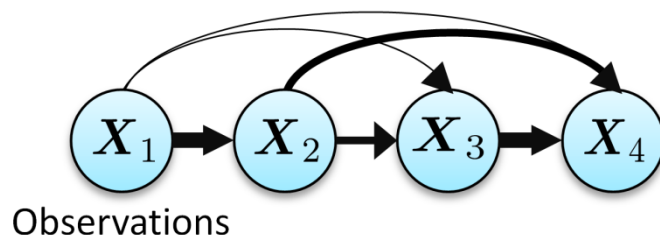
- Difference with RNNs, HMMs, HSMMs, Joint models.

Observable
models



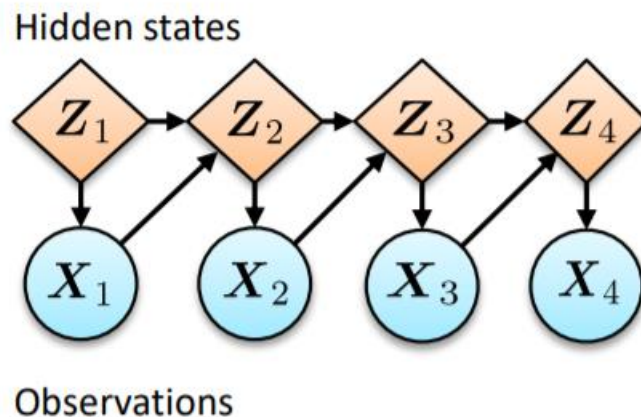
No latent structure

RETAIN



Interpretable predictions,
Uninterpretable latent
structure

RNN

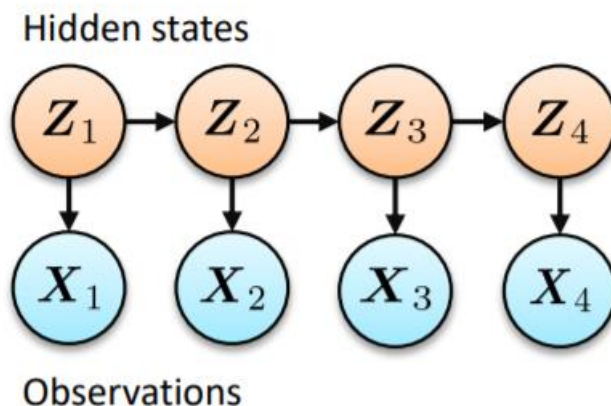


Uninterpretable predictions,
Uninterpretable latent
structure

Comparison with Existing Models

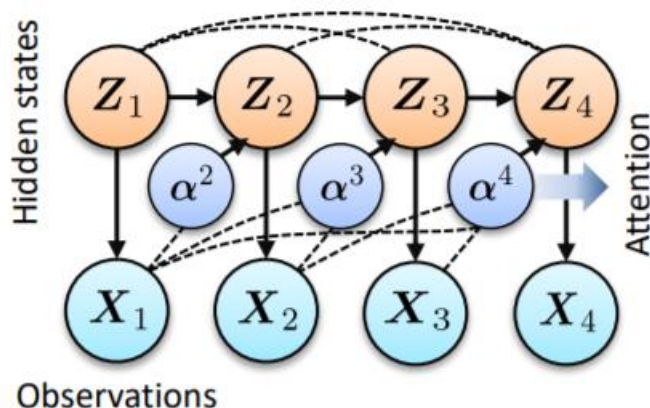
- Difference with RNNs, HMMs, HSMMs, Joint models.

HMM



Memoryless predictions,
Interpretable latent structure

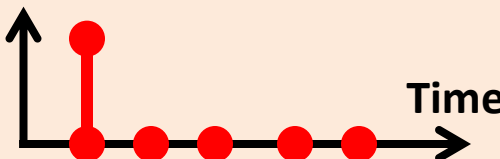
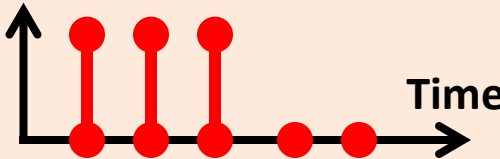
PASS



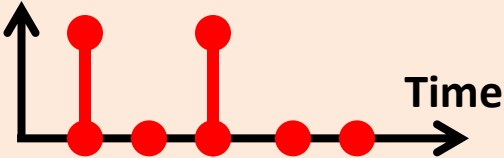
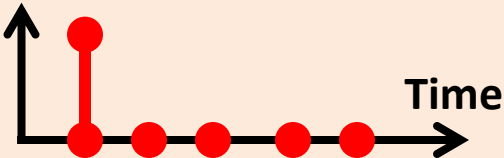
Memoryful and interpretable
predictions,
Interpretable latent structure

Generality of Attentive State Space Model

- Attentive state space models reduce to various classical sequential models for certain settings of attention mechanism.

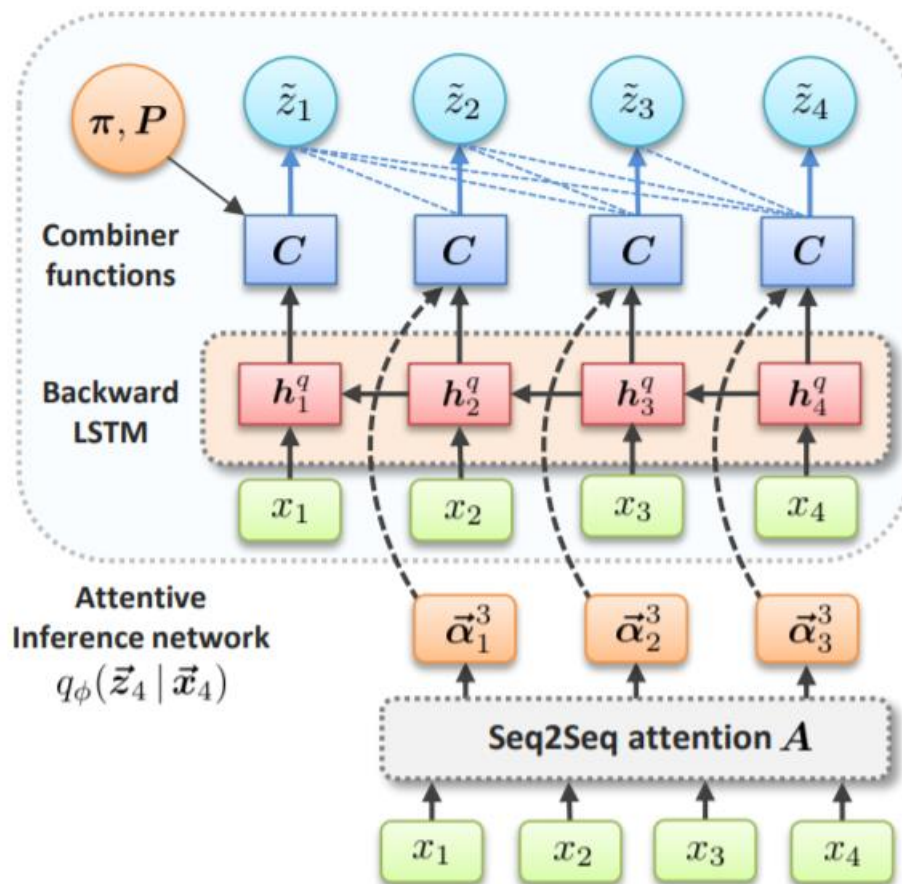
Attention mechanism	Model
<ul style="list-style-type: none">● 1 step back attention  <p>The diagram shows a horizontal axis labeled 'Time' with five red dots representing states. A vertical red line connects the first dot to the second dot, indicating that the state at time t depends on the state at time t-1.</p>	Hidden Markov Model
<ul style="list-style-type: none">● Fixed multi-step attention  <p>The diagram shows a horizontal axis labeled 'Time' with five red dots representing states. Three vertical red lines connect the first, second, and third dots to the fourth dot, indicating that the state at time t depends on states at times t-1, t-2, and t-3.</p>	Higher-order Hidden Markov Model

Generality of Attentive State Space Model

Attention mechanism	Model
<ul style="list-style-type: none">● Phased attention with binarized attention weights 	Variable-order Markov Model (context tree)
<ul style="list-style-type: none">● 1 step back attention with non-linear transitions 	Deep Markov Model

Learning and Inference

Sequence-to-sequence architecture for the attention mechanism

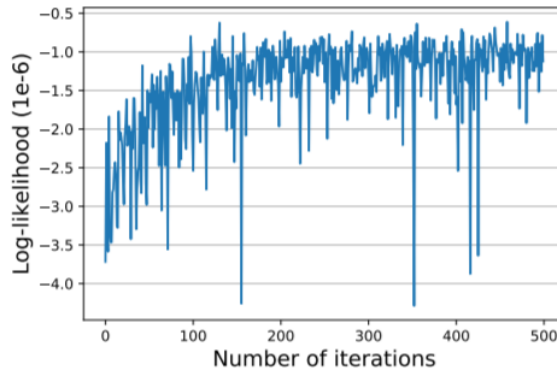


Experiments

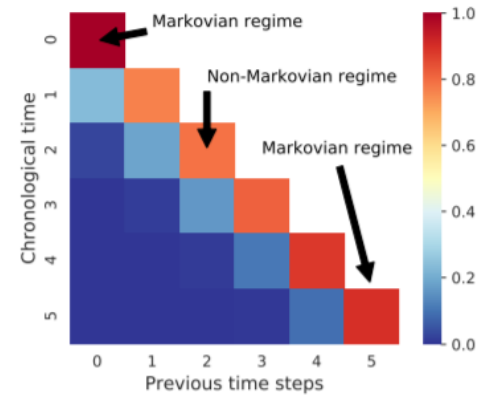
- We used data from a cohort of patients enrolled in the UK CF registry, a database held by the UK CF trust.
- The dataset records annual follow-ups for 10,263 patients over the period from 2008 and 2015, with a total of 60,218 hospital visits.
- Each patient is associated with 90 variables, including information on 36 possible treatments, diagnoses for 31 possible comorbidities and 16 possible infections, in addition to biomarkers and demographic information.

Experiments

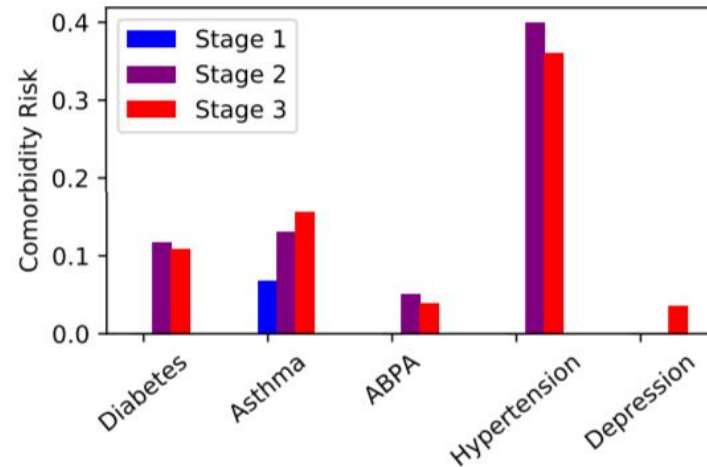
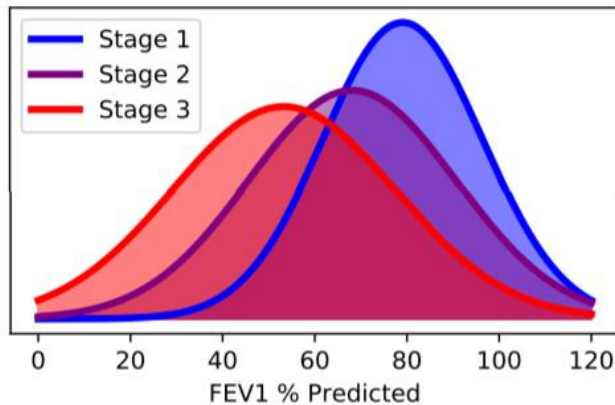
Learning curve



Individual trajectory



CF Phenotypes



Experiments

- Attentive state-space models do not only extract interpretable representations, but improve accuracy of prediction of comorbidities as well.

Model	Diabetes AUC-ROC	ABPA AUC-ROC	Depression AUC-ROC	Pancreatitis AUC-ROC	P. Aeruginosa AUC-ROC
Attentive SS	0.709 ± 0.02	0.787 ± 0.01	0.751 ± 0.03	0.696 ± 0.04	0.680 ± 0.01
HMM	0.625 ± 0.02	0.686 ± 0.03	0.667 ± 0.08	0.625 ± 0.04	0.610 ± 0.02
RNN	0.634 ± 0.03	0.727 ± 0.10	0.575 ± 0.01	0.590 ± 0.06	0.654 ± 0.01
LSTM	0.675 ± 0.03	0.740 ± 0.07	0.609 ± 0.12	0.578 ± 0.05	0.671 ± 0.01
RETAIN	0.610 ± 0.06	0.718 ± 0.05	0.580 ± 0.09	0.600 ± 0.08	0.676 ± 0.02