



Computational Phenotyping from EHR data and Medical Ontologies for Predictive Analytics

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How to get started?





If you use MIMIC data or code in your work, please cite the following publication:

MIMIC-III, a freely accessible critical care database. Johnson AEW,
Pollard TJ, Shen L, Lehman L, Feng M, Ghassemi M, Moody B, Szolovits P,
Celi LA, and Mark RG. Scientific Data (2016). DOI: 10.1038/sdata.2016.35.

Available from: http://www.nature.com/articles/sdata201635



- Critical Care Units
- 2001 2012
- 38,597 adult patients
- 53,423 distinct hospital admissions
- Age (med) = 65.8
- In-hospital mortality = 11.5%
- LOS @ICU (med) = 2.1d
- LOS @HOS (med) = 6.9d

• . . .



EHR Data Analytics: Plug-and-Play?



Electronic Health Records (EHR):



Patient demographics



Providing opportunities for predictive analytics (mortality, next diagnosis, length of stay, ...)



Medication prescriptions (ATC)



Heterogeneous data types



Diagnoses (ICD-10)



Laboratory tests (LOINC)

Complex (different sources, different codes, ...)

Missing, noisy, biased (collection process, reimbursement process, ...)

Hripcsak, George, and David J. Albers. "Next-generation phenotyping of electronic health records." Journal of the American Medical Informatics Association 20.1 (2012): 117-121.

Computational Phenotyping



Suppose you want to identify diabetes patients.

Searching by diagnosis codes is not good enough.

Toy examples:

Diabetes Diagnoses?

Diabetes Medications?

High blood glucose?

Case patient?

Yes Probably Yes Yes No Not

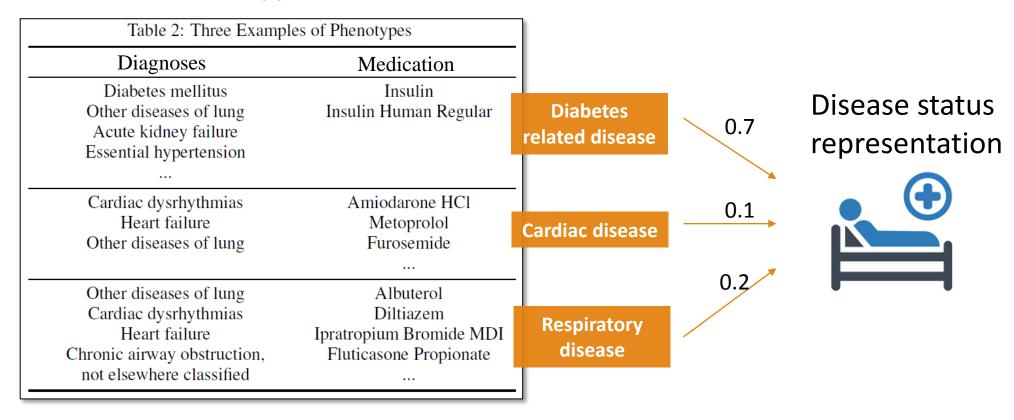
Instead, use <u>the combination of diagnoses, medications, procedures,</u> <u>laboratory tests, etc.</u> to identify patients with certain conditions.

Phenotypes (observable properties)

Computational Phenotyping



Phenotypes



Hripcsak, George, and David J. Albers. "Next-generation phenotyping of electronic health records." Journal of the American Medical Informatics Association 20.1 (2012): 117-121.

Computational Phenotyping

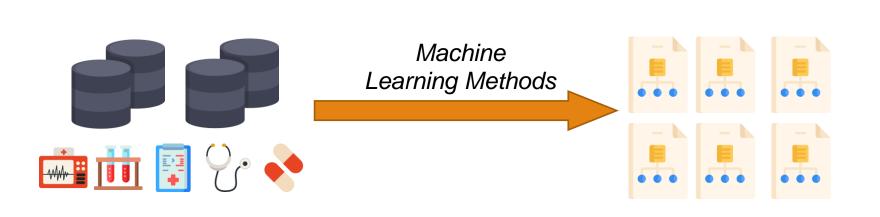


Phenotypes:

The combination of <u>clinically meaningful items</u> (e.g. diagnoses and medications) that <u>reveals the true disease status</u>.

Computational Phenotyping:

The process of automatically discovering meaningful phenotypes from the raw EHR data.



Machine Learning Methods

Natural Language Processing (NLP)

Deep Learning

Matrix Factorization

Tensor Factorization

^[1] Kirby, Jacqueline C., et al. "PheKB: a catalog and workflow for creating electronic phenotype algorithms for transportability." Journal of the American Medical Informatics Association 23.6 (2016): 1046-1052.

^[2] Ho, Joyce C., et al. "Limestone: High-throughput candidate phenotype generation via tensor factorization." Journal of biomedical informatics 52 (2014): 199-211.

^[3] Yang, Kai, et al. "TaGiTeD: Predictive Task Guided Tensor Decomposition for Representation Learning from Electronic Health Records." AAAI. 2017.



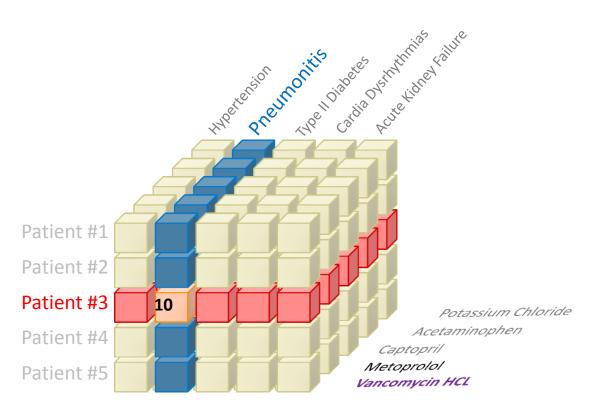
Hidden Interaction Tensor Factorization [IJCAI-18]

for Joint Learning of Phenotypes and Diagnosis-Medication Correspondence

Yin, Kejing, et al. "Joint Learning of Phenotypes and Diagnosis-Medication Correspondence via Hidden Interaction." Proceedings of the Twenty-Seventh International Joint Conference on Artificial Intelligence. 2018.

Tensor Factorization for Phenotyping





Patient #3 is prescribed with Vancomycin HCL for ten times in response to Pneumonitis.

^[1] Ho, Joyce C., et al. "Limestone: High-throughput candidate phenotype generation via tensor factorization." Journal of biomedical informatics 52 (2014): 199-211.

^[2] Ho, Joyce C., Joydeep Ghosh, and Jimeng Sun. "Marble: high-throughput phenotyping from electronic health records via sparse nonnegative tensor factorization." *Proceedings of the 20th ACM SIGKDD international conference on Knowledge discovery and data mining.* ACM, 2014.

^[3] Wang, Yichen, et al. "Rubik: Knowledge guided tensor factorization and completion for health data analytics." Proceedings of the 21th ACM SIGKDD International Conference on Knowledge Discovery and Data Mining. ACM, 2015.

^[4] Kim, Yejin, et al. "Discriminative and distinct phenotyping by constrained tensor factorization." Scientific reports 7.1 (2017): 1114.

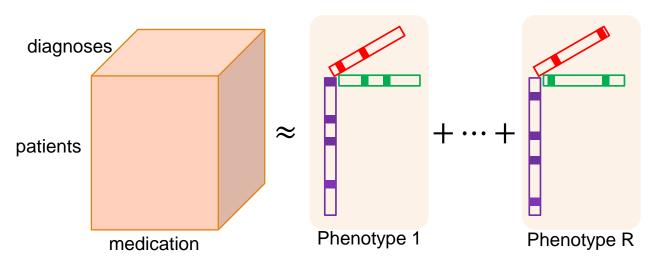
^[5] Yang, Kai, et al. "TaGiTeD: Predictive Task Guided Tensor Decomposition for Representation Learning from Electronic Health Records." AAAI. 2017.

^[6] Henderson, Jette, et al. "Granite: Diversified, Sparse Tensor Factorization for Electronic Health Record-Based Phenotyping." 2017 IEEE International Conference on Healthcare Informatics (ICHI), 2017.

Tensor Factorization for Phenotyping



Non-negative CP factorization for computational phenotyping:



Approximation with sum of R rank-one tensors:
$$\mathcal{X} \approx \hat{\mathcal{X}} = \sum_{r=1}^{R} a_r \circ b_r \circ c_r$$

Minimize the reconstruction error: min $\operatorname{Error}(\mathcal{X}, \mathcal{X})$

Interaction patterns are captured by the rank-one tensors.

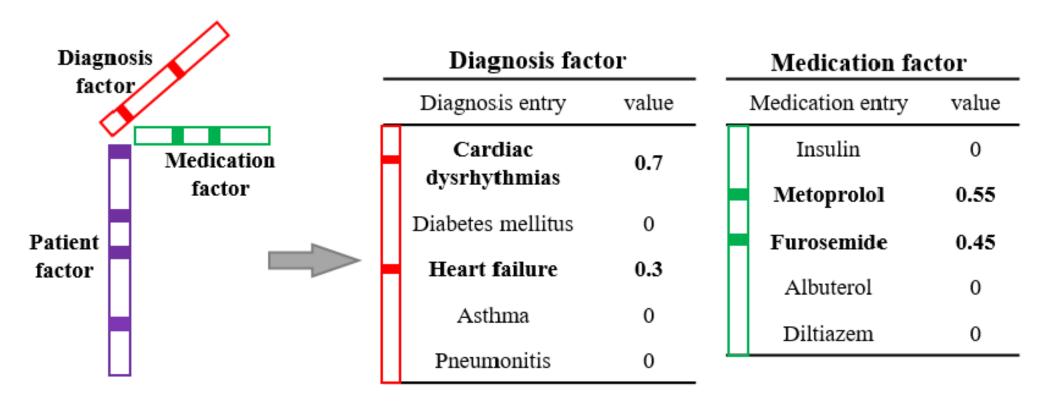
^[1] Kolda, T. G., & Bader, B. W. (2008). Tensor Decompositions and Applications. SIAM Review, 51(3)

^[2] Chi, Eric C., and Tamara G. Kolda. On tensors, sparsity, and nonnegative factorizations. SIAM Journal on Matrix Analysis and Applications 33.4 (2012): 1272-1299.

Tensor Factorization for Phenotyping



Phenotype extraction from rank-one tensor:



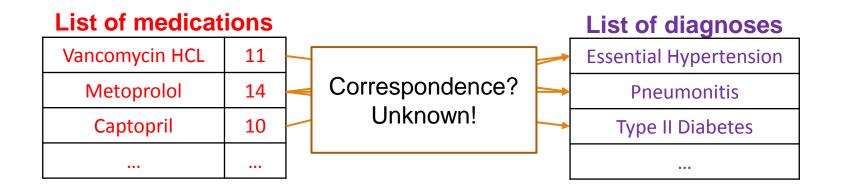
^[1] Kolda, T. G., & Bader, B. W. (2008). Tensor Decompositions and Applications. SIAM Review, 51(3)

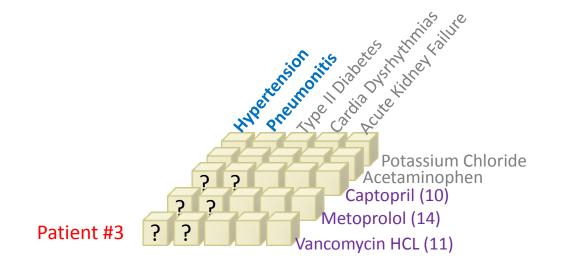
^[2] Chi, Eric C., and Tamara G. Kolda. On tensors, sparsity, and nonnegative factorizations. SIAM Journal on Matrix Analysis and Applications 33.4 (2012): 1272-1299.

Research Challenge



Interaction information is often missing in the records.





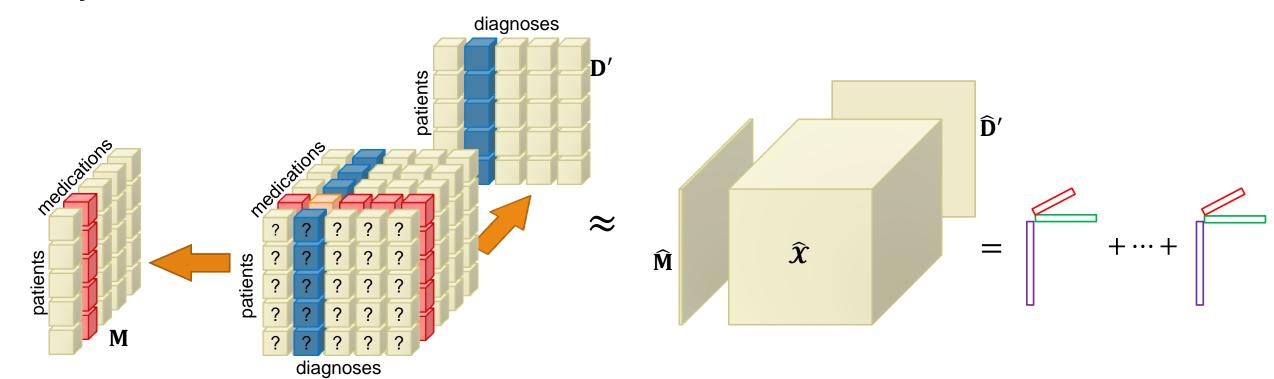
How to fill in the entries?

How to factorize the tensor when we do not observe it?

Hidden Interaction Tensor Factorization



Key Idea



Interaction tensor X: NOT observed



■ Diagnosis-Medication Correspondence

Table 1: Top Five Corresponding Medications for Three Diagnoses Inferred by HITF and Rubik

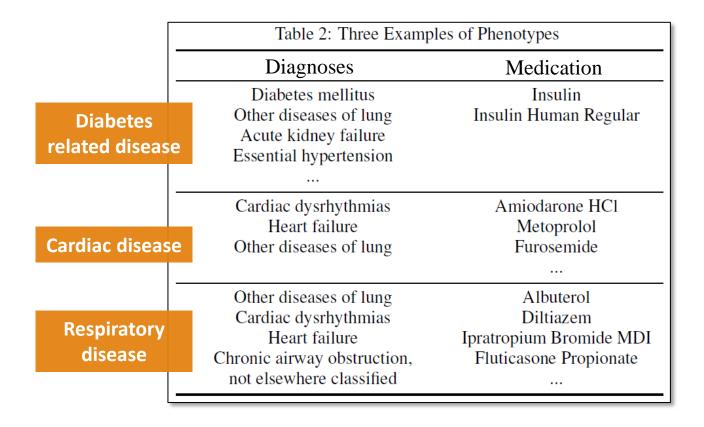
Cardiac dysrhythmias(39.0%)		Diabetes mellitus(25.3%)		Asthma(5.5%)	
HITF	Rubik	HITF	Rubik	HITF	Rubik
Furosemide(0.08)	Potassium Chloride(0.08)	Insulin(0.64)	Insulin(0.09)	Albuterol 0.083% Neb Soln(0.46)	Potassium Chloride(0.08)
Potassium Chloride(0.07)	Insulin(0.06)	Relevant drug identified by HITF gets much higher weight		Ipratropium Bromide Neb(0.39)	Insulin(0.06)
Metoprolol(0.06)	unrelated Furosemide(0.06)			Furosemide(0.08)	Furosemide(0.05)
Amiodarone HCl(0.05)	Magnesium Sulfate(0.04)	Furosemide(0.03)	Magnesium Sulfate(0.03)	He Relevan	Magnesium t drugs 0.04)
Heparin Sodium(0.04)	Acetaminophen(0.03)	Atorvastatin(0.03)	Acetaminophen(0.03)	inferred on	ly by HITF phen(0.03)

Evaluated by a clinician:

"There is qualitative superiority of HITF method over the Rubik method."



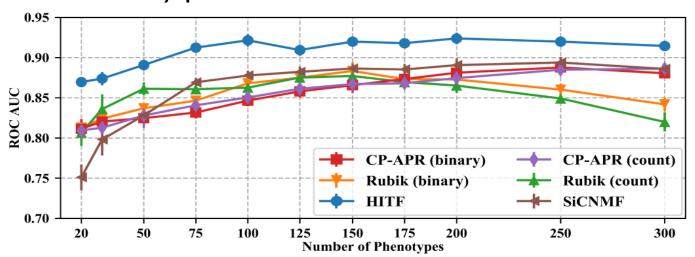
Clinical relevance of the Phenotypes



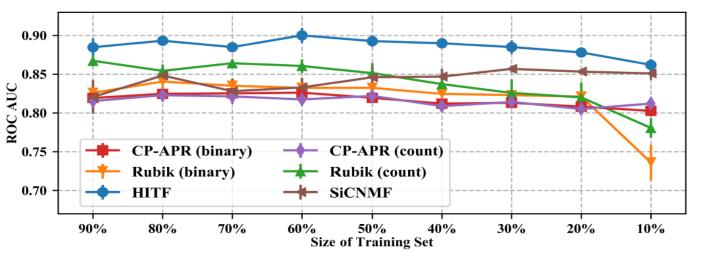
According to the clinician, phenotypes inferred by HITF are clinically relevant.







- HITF outperforms all baselines consistently in terms of mortality prediction task.
- More robust against small size of training set.



Patients can be effectively represented by phenotypes derived using HITF.



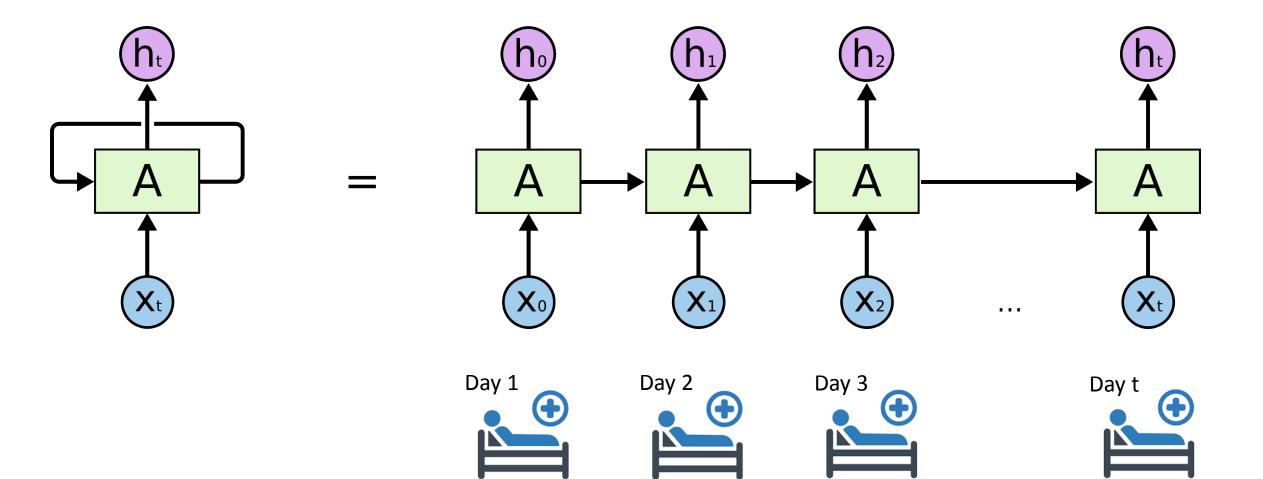
Collective Non-negative Tensor Factorization [AAAI-19]

with RNN regularization for Joint Learning of Static Phenotypes and Dynamic Patient Representation

Yin, Kejing, et al. "Learning Phenotypes and Dynamic Patient Representations via RNN Regularized Collective Non-negative Tensor Factorization." Proceedings of the Thirty-Third AAAI Conference on Artificial Intelligence. 2019.

Recurrent Neural Network

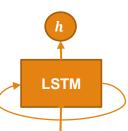




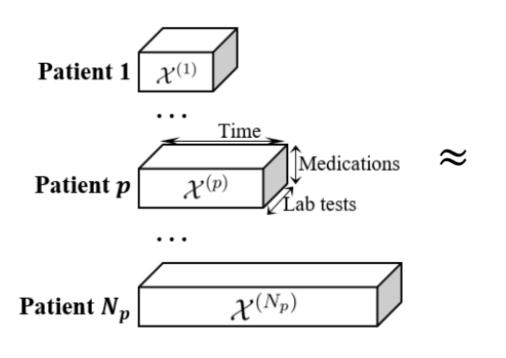
Collective Non-negative Tensor Factorization

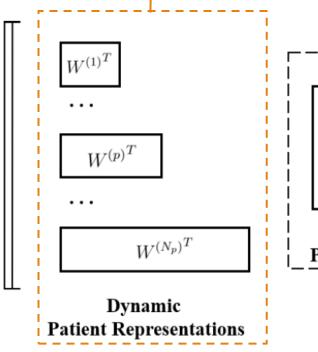


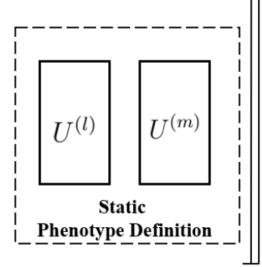
Represent each patient with a temporal tensor



View the <u>temporal representation as a</u> <u>multi-variate time series</u> of the disease states.



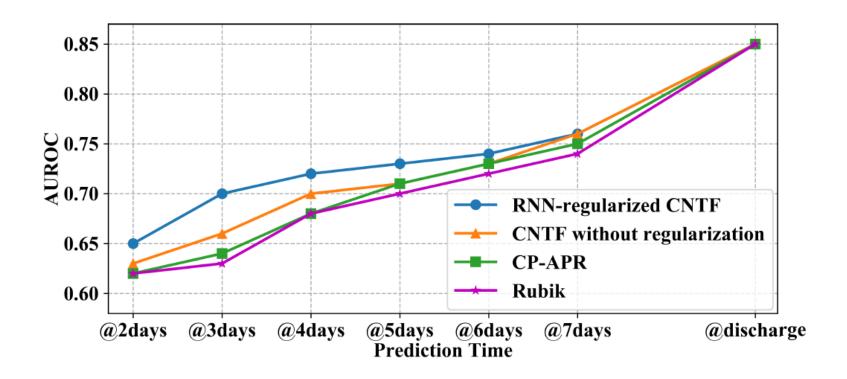




RNN Regularized CNTF



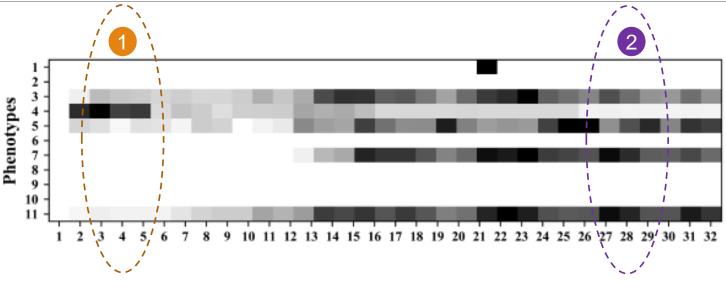
■ Results: Mortality Prediction



Higher prediction rate is resulted

Dynamic Patient Representation





High value for phenotype 4 (Chronic Heart Disease)

2 High value for phenotype 3 (Other Disease of the Lung), phenotype 5 (Cardiac Dysrhythmias), phenotype 7 (Acute Kidney Failure), phenotype 11 (Cardiac Dysrhythmias with Heart Failure)

"Patient <u>admitted with existing condition</u>, chronic heart disease, which is <u>treated unsuccessfully</u>, and <u>eventually developed multiple organ failure</u>." (Supported by reviewing the clinical notes.)

RNN Regularized CNTF



Results: Phenotypes

Our proposed model

Clinically much more meaningful, evaluated by a medical expert.

Phenotype 1	Phenotype 4	Phenotype 9	
Chronic kidney disease	Other forms of chronic ischemic heart disease (0.507)	Other diseases of lung	
(CKD) (0.536)	Cardiac dysrhythmias (0.372) Essential hypertension (0.024)	(0.876)	
RBC (Urine) (0.200)	Hematocrit (Blood) (0.072)	pO2 (Blood Gas) (0.253)	
Osmolality, Measured (Blood) (0.117)	Red Blood Cells (Blood) (0.071)	pCO2 (Blood Gas) (0.237)	
Protein/Creatinine Ratio (Urine) (0.069)	Hemoglobin (Blood) (0.070)	pH (Blood Gas) (0.215)	

Hydromorphone (0.336)

"Phenotype 9 corresponds to the diagnosis Other Disease of the Lung and abnormal laboratory tests pO2, pCO2, pH of the arterial blood gas. Again, this correlates well with the clinical context, where reduced oxygen levels and pH, and elevated carbon dioxide levels all indicate the presence of acute respiratory failure (which is classified under the "other disease of lung" in the ICD-9 coding system)."

"The disease state CKD is indeed associated with elevated RBC in (0.038) urine due to renal tubular necrosis, elevated blood osmolality due to 33) electrolyte retention in the vascular system, and elevated protein loss in the urine leading to an abnormal protein/creatinine ratio."

Baseline: Rubik

Phenotype 1	Phenotype 2	Phenotype 3	
Other diseases of lung (0.045) Septicemia (0.040)	Other diseases of lung (0.040) Acute kidney failure (0.036)	Acute kidney failure (0.039) Other diseases of lung (0.037) Cardiac dysrhythmias (0.033)	
Certain adverse effects not elsewhere classified (0.039)	Certain adverse effects not elsewhere classified (0.032)		
Glucose(Blood) (0.019)	Hematocrit(Blood) (0.017)	Glucose(Blood) (0.018)	
Red Blood Cells(Blood) (0.019)	Red Blood Cells(Blood) (0.017)	Hematocrit(Blood) (0.018)	
Hematocrit(Blood) (0.019)	Glucose(Blood) (0.017)	Red Blood Cells(Blood) (0.018)	
Vancomycin (0.017)	Vancomycin (0.013)	Vancomycin (0.015)	
Insulin (0.015)	Potassium Chloride (0.013)	Potassium Chloride (0.014)	
Potassium Chloride (0.015)	Pantoprazole Sodium (0.012)	Heparin (0.014)	



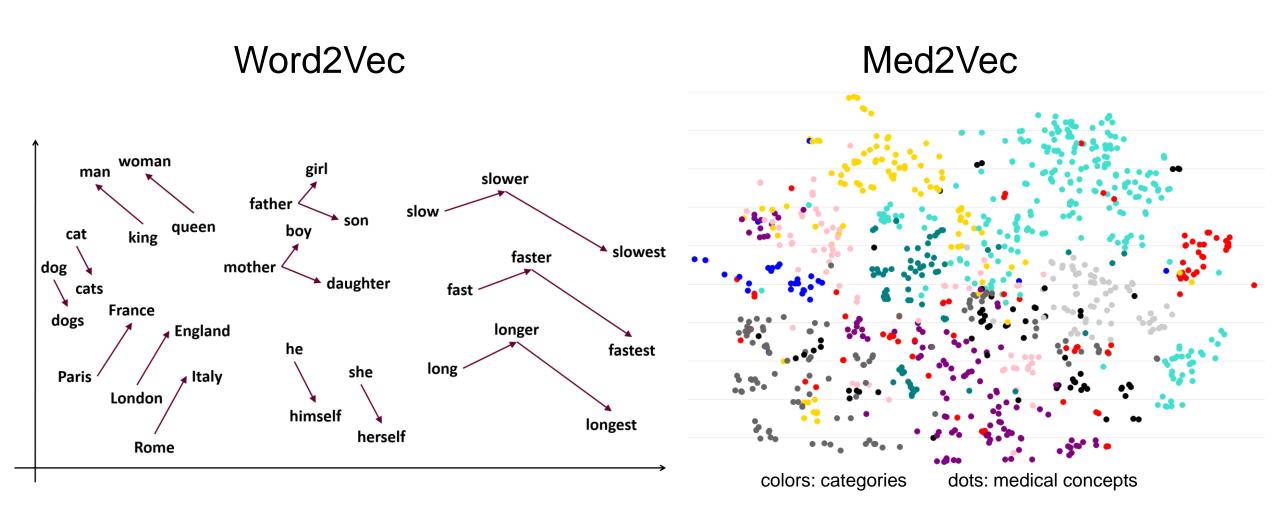
Multiple Ontological Representations (MMORE) [IJCAI-19]

for learning medical concept representations from medical ontologies and EHR

Song, Lihong, et al. "Medical concept embedding with multiple ontological representations ." Proceedings of the Twenty-Eighth International Joint Conference on Artificial Intelligence. 2019.

Representation Learning for Medical Concepts



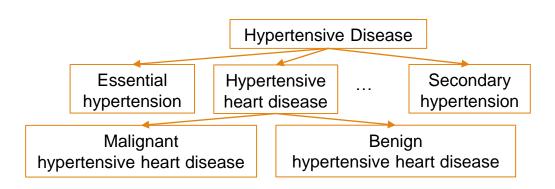


Choi, Edward, et al. "Multi-layer representation learning for medical concepts." Proceedings of the 22nd ACM SIGKDD International Conference on Knowledge Discovery and Data Mining. ACM, 2016.

Research Challenge



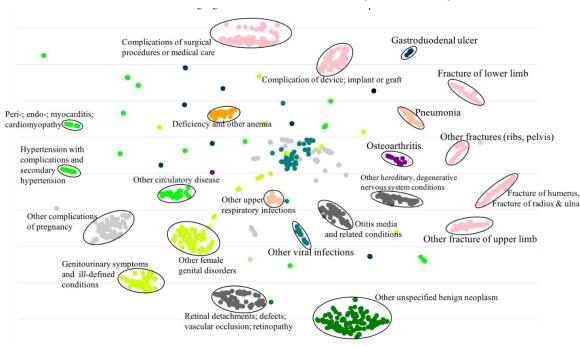
Inconsistency between medical ontologies and EHR



Example: ICD-9 ontology

- Good enough?
- Medical concepts under the same category should co-occur with other concepts in EHR in a similar manner. Correct? *E.g.*, essential hypertension & secondary hypertension.

GRAM model (KDD '17)

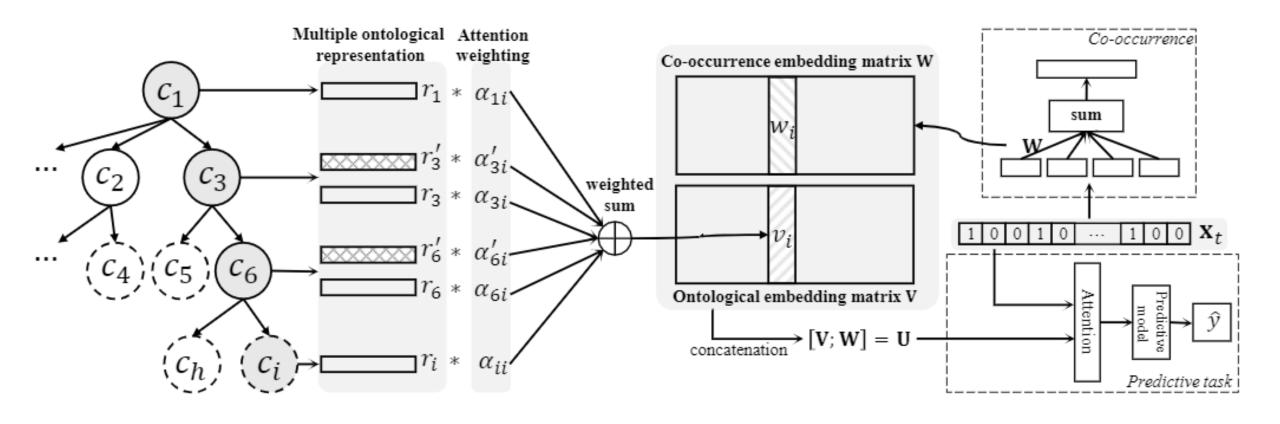


Choi, Edward, et al. "GRAM: Graph-based attention model for healthcare representation learning." Proceedings of the 23rd ACM SIGKDD International Conference on Knowledge Discovery and Data Mining. ACM, 2017.

MMORE



Key Idea: Multiple representations for each ontological category





Next-admission Diagnosis Prediction

Measure the predictive performance by $Accuracy@k = \frac{\text{# of true positives in the top } k \text{ predictions}}{\text{# of positives}}$

Utilize only the EHR data

Mainly focus on medical ontologies

Consider both the ontologies and the EHR co-occurrence

- Less sensitive to the medications
- Ontologies could serve the role to "regularize" the learned representations

Size of training data are varied to train models

Data	Model	20%	40%	60%	80%
	RETAIN	0.4422	0.4447	0.4449	0.4545
Dx	Med2Vec	0.5064	0.5187	0.5200	0.5290
	GRAM	0.4980	0.5218	0.5409	0.5498
	MMORE	0.5205	0.5426	0.5548	0.5618
	RETAIN	0.4422	0.4447	0.4449	0.4547
Dx &	Med2Vec	0.4920	0.4967	0.4979	0.5110
Rx	GRAM	0.5057	0.5285	0.5426	0.5548
	MMORE	0.5243	0.5498	0.5619	0.5689

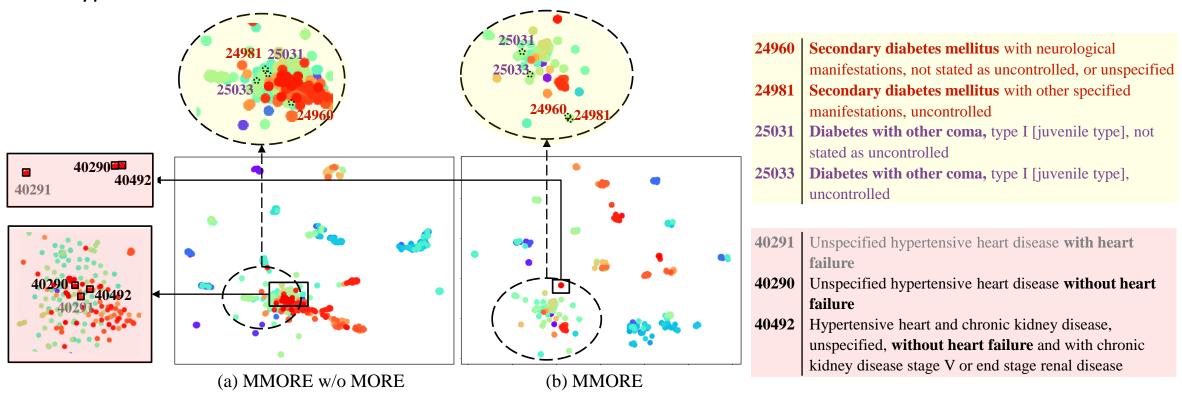
Dx for the diagnosis, Rx for the medication



Case study

Diabetes with neurological manifestations & Diabetes with other manifestations

Hypertensive heart disease with or without heart failure



Learned representations align with both EHR and medical ontologies

Experimental Results: Phenotyping



- Applying Non-negative Matrix Factorization to Attention Matrix
- Basis factors try to group related concepts together (phenotypes)

Phenotype 1

Dx: Atrial fibrillation; Congestive heart failure, NOS; ...

Heart diseases

Rx: Warfarin; Heparin; ...

Phenotype 2

Dx: Cirrhosis of liver w/o mention of alcohol;

Dx: Alcoholic cirrhosis of liver; ...

Rx: Lactulose; Folic acid; ...

Liver diseases

Phenotype 3

Dx: Chronic airway obstruction, NEC;

Dx: Obstructive chronic bronc w/ (acute)

exacerbation; ...

Rx: Ipratropium bromide; Albuterol sulfate; ...

Respiratory diseases

Closing Remarks



- Three ML methods proposed for EHR Data Analytics.
 - Tensor Factorization -> HITF model
 - Tensor Factorization + RNN -> CNTF model
 - Representation Learning + Ontology -> MMORE model

- Future Research Directions:
 - More data modalities (e.g., vital signs)
 - Going beyond categorical ontology (e.g., SNOMED-CT)
 - Continuous-time modelling (from ICU to primary care data)





Thank you!

Q&A



References



- [1] Adoption of Electronic Health Record Systems among U.S. Non-Federal Acute Care Hospitals: 2008-2015.
- [2] Johnson AEW, Pollard TJ, Shen L, Lehman L, Feng M, Ghassemi M, Moody B, Szolovits P, Celi LA, Mark RG. MIMIC-III, a freely accessible critical care database. *Scientific Data*, 2016.
- [3] Hripcsak, George, and David J. Albers. "Next-generation phenotyping of electronic health records." *Journal of the American Medical Informatics Association* 20.1 (2013): 117-121.
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- [8] Jennifer Pacheco and Will Thompson. Northwestern University. Type 2 Diabetes Mellitus. PheKB; 2012 Available from: https://phekb.org/phenotype/18
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- [16] Choi, E., Bahadori, M. T., Searles, E., Coffey, C., Thompson, M., Bost, J. & Sun, J. (2016, August). Multi-layer representation learning for medical concepts. In *Proceedings of the 22nd ACM SIGKDD International Conference on Knowledge Discovery and Data Mining* (pp. 1495-1504). ACM.