

Clinical Data Modelling with Deep Neural Networks: Challenges and Solutions

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Why Deep Learning in Healthcare?

Data

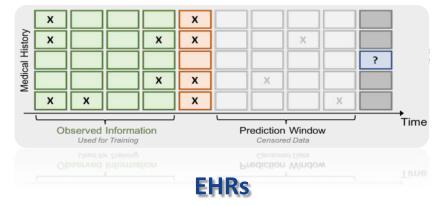
Medical Images



Text

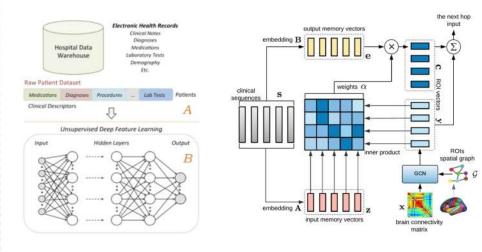


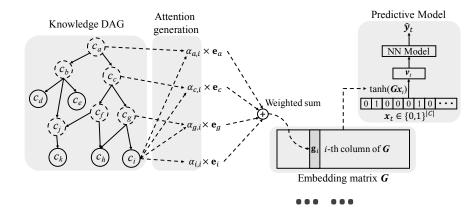
roundstudy different analysis





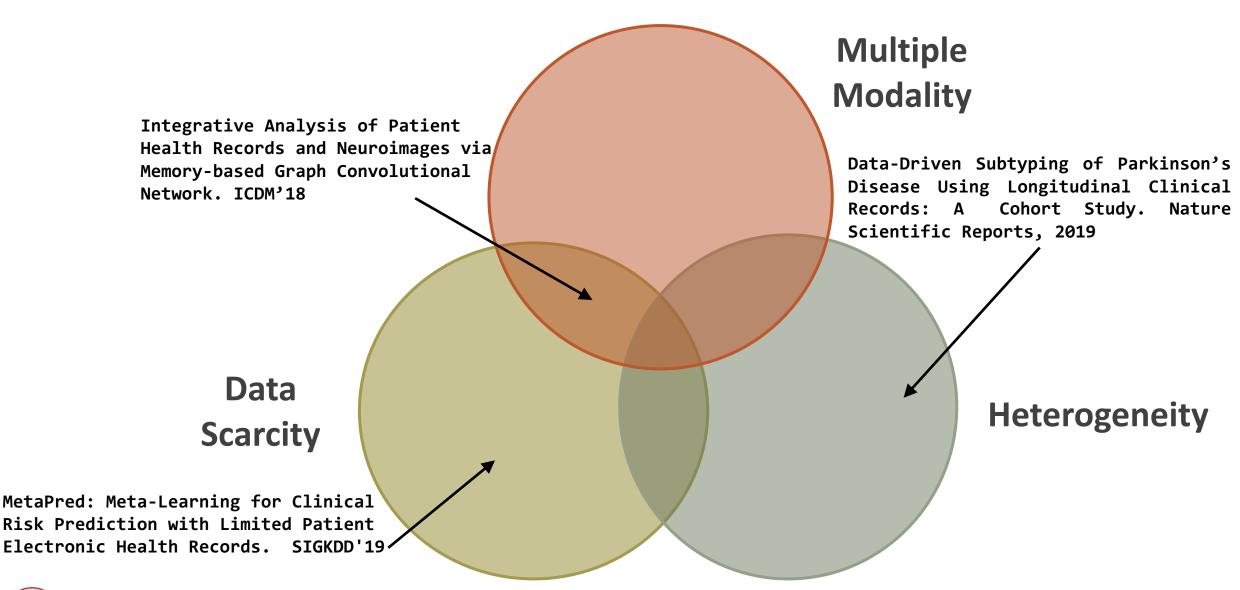
Algorithms







Obstacles





Outline



Part 1: Disease Subtyping on Clinical Times Series

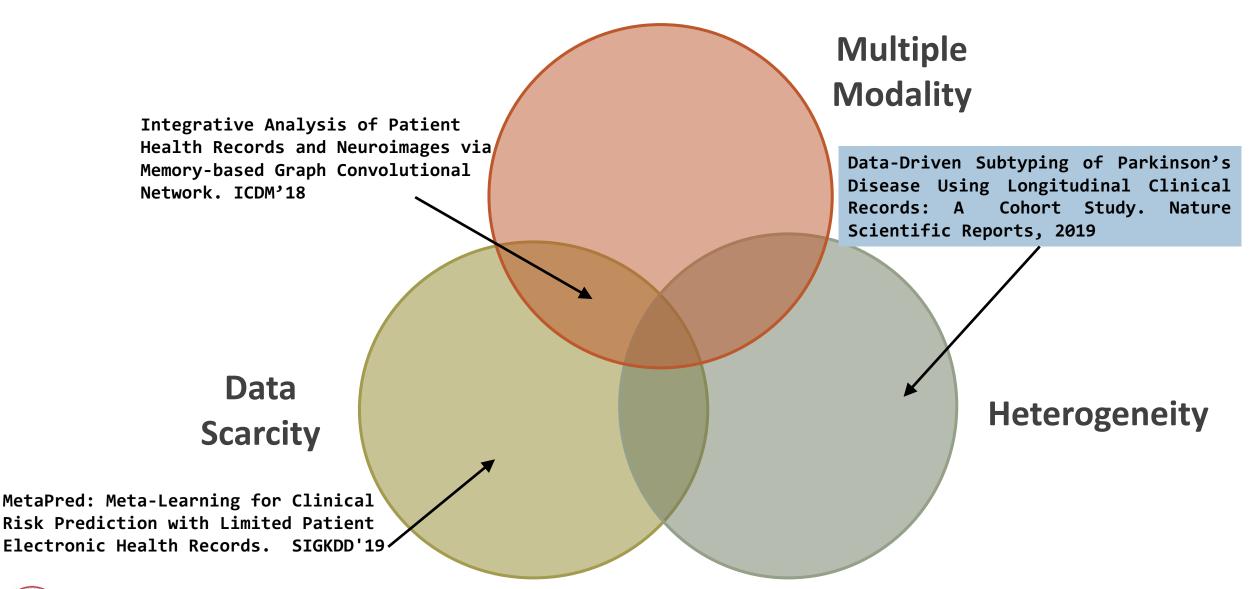


Part 2: Integrative Disease Analysis via Multi-Modality



Part3: Meta-Learning on Limited Clinical Resources

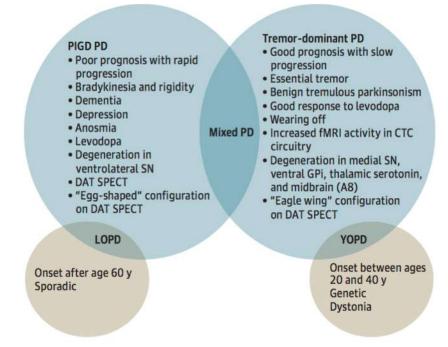
Obstacles





<u>Background</u>: Parkinson's Disease (PD) has been demonstrated heterogeneous in clinical representation and disease progression. Therefore, identifying subtypes with similar characteristics is an important task to study the disease.

- The disease associates with clinical factors of motor, non-motor, and other variables, so that there is no widely accepted consensus on the criteria for patient groups.
- Data-driven approaches of clustering methodologies can identify subtypes without a priori hypothesis about disease knowledge.
- Recurrent neural networks are successful in many sequential learning tasks, and may allow us to find more PD progression patterns among clusters.



Challenges

- ❖ Parkinson's Disease (PD) is clinically *heterogeneous* associated with a broad spectrum of clinical variable factors;
- * How to identify *disease progression* biomarkers, so that we can provide a better population for modifying drug trials.

Solution

- We first concatenated the multi-source records according to their occurring timestamps to form a temporal sequence for each patient;
- A deep learning model **LSTM** is trained to encode the record sequences into a series of standardized embeddings.

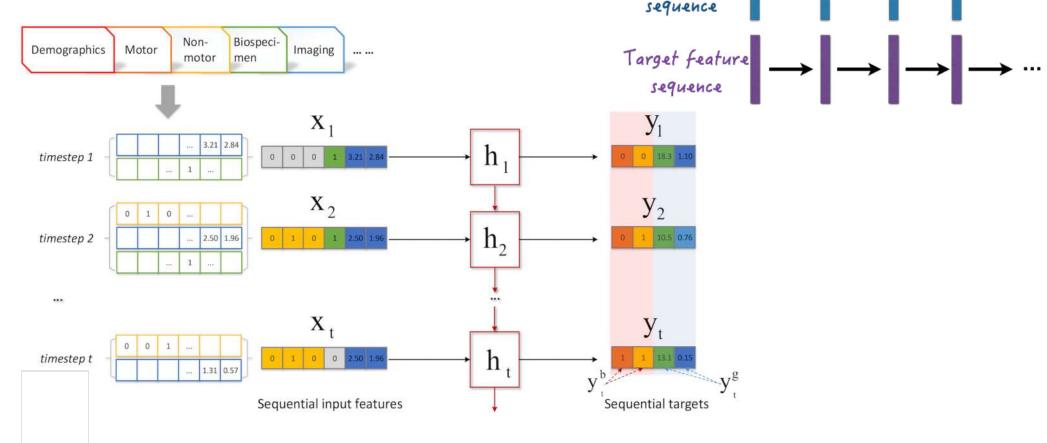


Demographics Motor Non- Biospeci- Imaging

Features

	Target Clinical Variables			
1	Clinical Diagnosis			
2	Demographics			
3	Motor symptoms: MDS-UPDRS scores*			
4	Cognitive Accessments: MoCA*			
-	Cognitive Categorization: Normal Cognition; Mild Cognitive			
5	Impairment; Dementia			
6	Other nonmotor variable: REM Sleep Disorder			
7	Biospecimen: Lumber Puncture Sample Collection			
0	Biospecimen: Laboratory Procedures containing DNA, RNA, Urine,			
8	Plasma, & Serum samples			
9	Imaging Results: DaTScan Striatal Binding Ratio			
10	Imaging Results: Magnetic Resonance Imaging			

Ref: Fereshtehnejad, Seyed-Mohammad, Silvia Rios Romenets, Julius BM Anang, Véronique Latreille, Jean-François Gagnon, and Ronald B. Postuma. "New clinical subtypes of Parkinson disease and their longitudinal progression: a prospective cohort comparison with other phenotypes." JAMA neurology 72, no. 8 (2015): 863-873.



- Continuous target feature:

$$\frac{1}{2} \sum_{t} \|y_{t}^{g} - W_{g} h_{t}\|_{2}^{2} + \lambda \|W_{g}\|_{F}^{2}$$

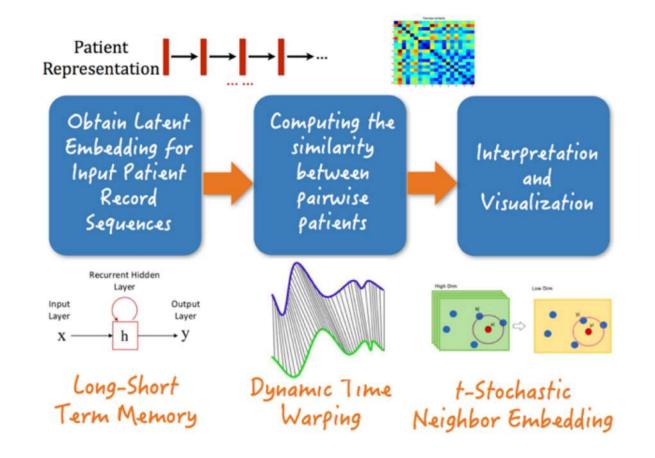
- Binary target feature:

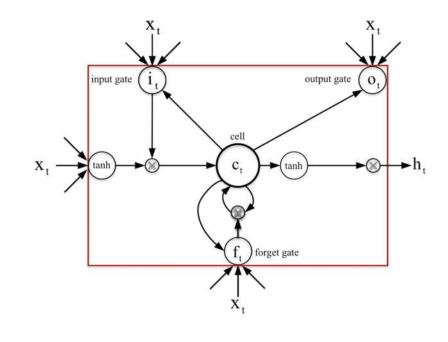
$$\sum_{t} \sum_{j=1}^{m_b} \log(1 + \exp(-y_{t,j}^b(w_{b,j}^T h_t))) + \lambda ||W_b||_F^2$$

Input feature



Method





$$i_{t} = \sigma(W_{i}x_{t} + W_{i}h_{t-1} + b_{i})$$

$$f_{t} = \sigma(W_{f}x_{t} + W_{f}h_{t-1} + b_{f})$$

$$o_{t} = \sigma(W_{o}x_{t} + W_{o}h_{t-1} + b_{o})$$

$$c_{t} = f_{t} \cdot * c_{t-1} + i_{t} \cdot * tanh(W_{c}x_{t} + W_{c}h_{t-1} + b_{c})$$

$$h_{t} = o_{t} \cdot * tanh(v_{c}x_{t} + v_{c}n_{t-1} + b_{c})$$

$$h_{t} = o_{t} \cdot * tanh(c_{t})$$



THE MICHAEL J. FOX FOUNDATION FOR PARKINSON'S RESEARCH

Dataset

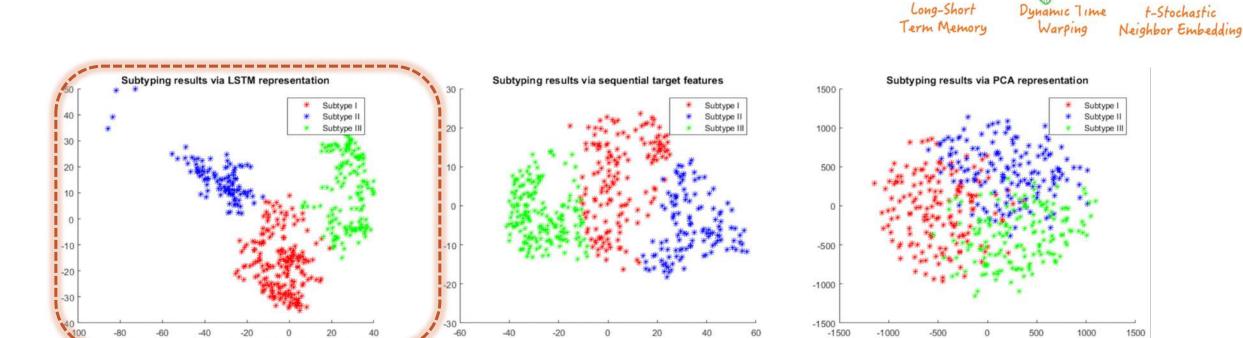
- The patient data were obtained from the Parkinson Progression Marker Initiative (PPMI) study. http://www.ppmi-info.org/
- The de-identified data contained archives of enrolled subjects from June 1, 2010, to June 1, 2016.

Туре	Code	# subjects
Idiopathic PD	1	466
Corticobasal degeneration	4	0
Dementia with Lewy bodies	5	1
Essential tremor	7	2
Multiple system atrophy	11	0
Psychogenic illness	15	1
No PD nor other neurological disorder	17	219
Other neurological disorder(s) (specify)	97	4
		759



Disease Subtyping: Visualization

Results



Comparison with Traditional Clustering Methods



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Obtain Latent

Embedding for

Input Patient

Record

Sequences

Computing the

similarity

between

Pairwise

patients

Interpretation

and

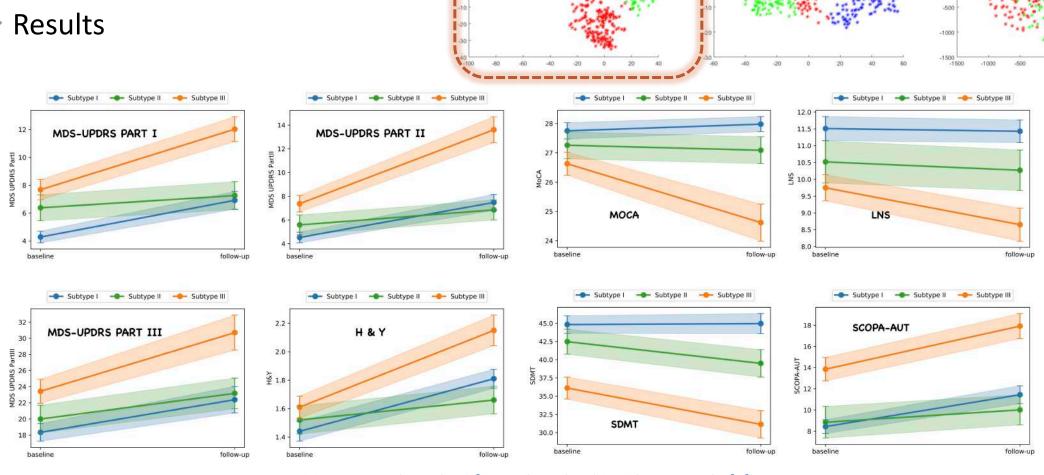
Visualization

Disease Subtyping: Statistical Testing

^a Chi-square test; ^b F-test; ^c One-way ANOVA test; ^d Kruskal-Wallis H-test

	Subtype I	(N = 201)	Subtype I	I (N = 107)	Subtype III	(N=158)	P-V	'alue
Characteristics	Baseline	Follow-up	Baseline	Follow-up	Baseline	Follow-up	Static	Progressi on
Age onset	58.79(9.5)		61.93(9.0)		65.32(8.8)		<0.0001 ^a	
Hoehn and Yahr Stage	1.44(0.5)	1.81(0.4)	1.52(0.5)	1.66(0.5)	1.61(0.5)	2.15(0.6)	<0.0001a	<0.0001a
MDS-UPDRS Part I	4.28(2.9)	6.92(4.5)	6.38(4.8)	7.26(5.2)	7.68(4.6)	12.03(5.7)	<0.0001a	<0.0001a
MDS-UPDRS Part II	4.52(3.2)	7.48(4.8)	5.58(4.4)	6.85(4.4)	7.37(4.4)	13.61(7.0)	<0.0001a	<0.0001a
MDS-UPDRS Part III	18.34(7.9)	22.39(11.8)	19.99(9.0)	23.18(9.9)	23.43(9.5)	30.71(13.9)	0.1146a	<0.0001 ^a
Montreal Cognitive Assessment	27.75(2.0)	27.98(1.8)	27.26(2.4)	27.09(2.4)	26.63(2.5)	24.62(4.0)	<0.0001ª	<0.0001ª
Geriatric Depression Scale	5.11(1.4)	5.2(1.3)	5.2(1.17)	5.31(1.2)	5.47(1.5)	5.96(1.8)	0.0017ª	0.0010 ^a
State Trait Anxiety Inventory	61.84(15.8)	59.52(16.0)	62.14(17.9)	61.89(18.1)	71.0(19.8)	74.25(20.1)	0.0053ª	0.1717ª
DaTScan	1.43(0.5)	1.23(0.5)	1.60 (0.6)	3.05 (0.6)	1.23 (0.5)	0.97 (0.5)	<0.0001°	<0.0001°

Results



Subtyping results via LSTM representation

Subtyping results via sequential target features

Progression in the Discriminative Variables

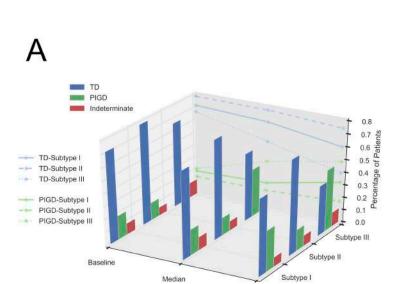


Subtyping results via PCA representation

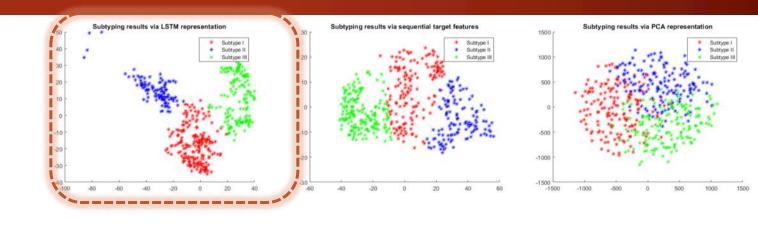
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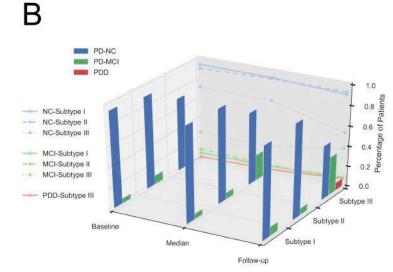
Disease Subtypi	ng _{Subtype I}	Subtype II	Subtype III		Subtype I	Subtype II	Subtype III
ВЈІО -	新 F	j'	i' .	BJLO -	IF.	F	li e
CSF-Abeta 42				DaTScan SBR-CAUDATE LEFT -			
CSF-Total tau -				DaTScan SBR-CAUDATE RIGHT -			14
DaTScan SBR-CAUDATE LEFT -	-			DaTScan SBR-PUTAMEN LEFT -			2
DaTScan SBR-CAUDATE RIGHT -				DaTScan SBR-PUTAMEN RIGHT -			
DaTScan SBR-PUTAMEN LEFT -	•						
DaTScan SBR-PUTAMEN RIGHT -				GDS -			
ESS -				Н&Ү -			14
GDS -				-HVLT-Discrimination Recognition -			
H&Y -				HVLT-Immediate Recall -			
HVLT-Discrimination Recognition -				HVLT-Retention -			
HVLT-Immediate Recall -			,	LNS -			
HVLT-Retention -				-			
LNS -				MCI -			-
MCI -			,	MDS UPDRS Partl -			5
MDS UPDRS Partl -				MDS UPDRS PartII -			
MDS UPDRS PartII -				MDS UPDRS Partill -			
MED USE -				MED USE -			
MoCA -							
RBD -				MoCA -			
Variables with COPA-AUT				RBD -			
p-value < 0.05 are SDM				SDM -			
shown stai		15	5 September 201	SF -		0	Microsoft Research

Results



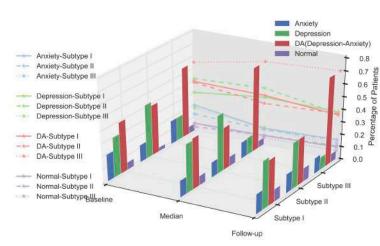
Correlation with Motor Subtypes





Correlation with Cognitive Subtypes

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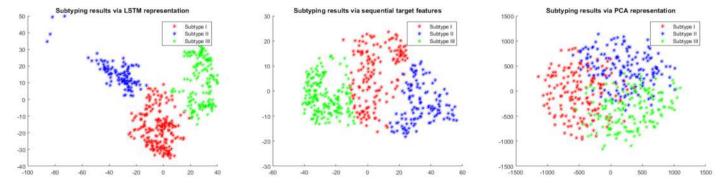


Correlation with Mood Subtypes

Relationship with Conventional PD Subtypes



Interpretation

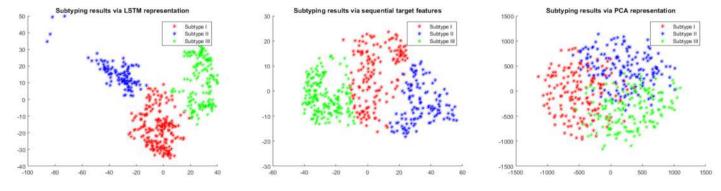


Subtype I (43.1%)	Subtype II (22.9%)	Subtype III (33.9%)
58.79 years at baseline	61.93 years at baseline	65.32 years at baseline
Mild motor symptoms at baseline	Moderate motor symptoms at baseline	Poor motor symptoms at baseline
Mild non-motor symptoms at baseline	Moderate non-motor symptoms at baseline	Poor non-motor symptoms at baseline
Moderate motor decay	Mild motor decay	Severe motor decay

Xi Zhang, Jingyuan Chou, Jian Liang, Cao Xiao, Yize Zhao, Harini Sarva, Claire Henchcliffe, Fei Wang, Data-Driven Subtyping of Parkinson's Disease Using Longitudinal Clinical Records: A Cohort Study. Scientific Reports, Nature, 2018



Summarization



- ✓ This study is an initial attempt on leveraging advanced data analytics for identification of PD subtypes with longitudinal and heterogeneous clinical study data.
- ✓ Our approach has demonstrated strong potentials of identification of comprehensive progressive PD subtypes.

Limitation

✓ the approach the deep learning (LSTM) procedure cannot be straightforwardly interpreted; Also, our study is only conducted on the PPMI cohort.

Source Code: https://github.com/sheryl-ai/Nature-Scientific-Reports

Xi Zhang, Jingyuan Chou, Jian Liang, Cao Xiao, Yize Zhao, Harini Sarva, Claire Henchcliffe, Fei Wang, Data-Driven Subtyping of Parkinson's Disease Using Longitudinal Clinical Records: A Cohort Study. Nature Scientific Reports, volume 9, Article number: 797 (2019).



Outline



Part 1: Disease Subtyping on Clinical Times Series

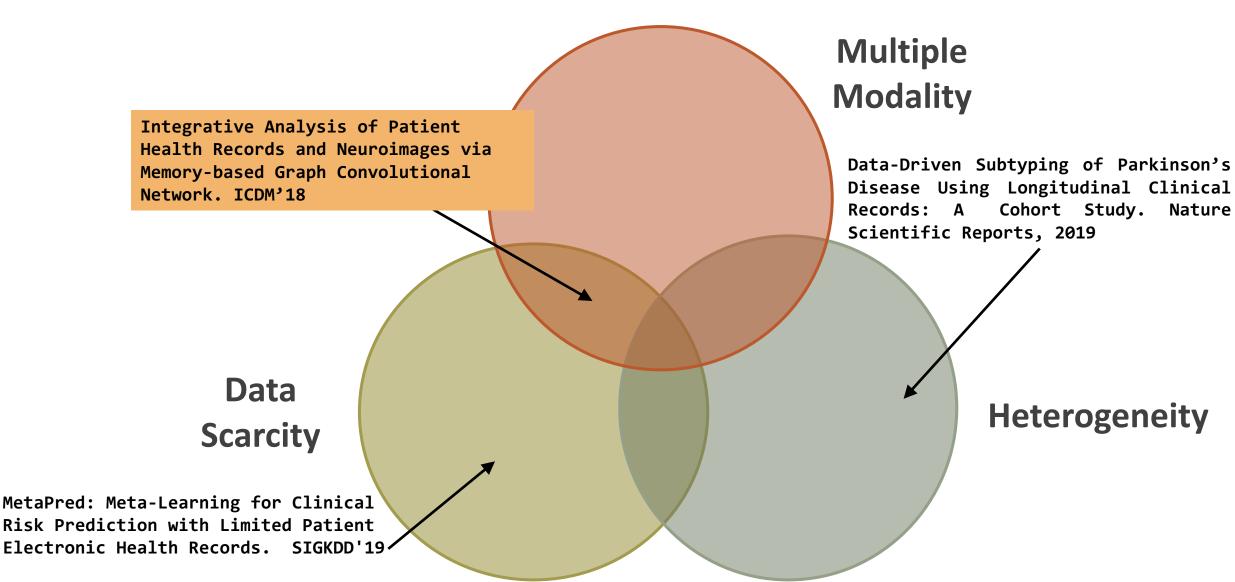


Part 2: Integrative Disease Analysis via Multi-Modality



Part3: Meta-Learning on Limited Clinical Resources

Obstacles





*Background: For complicated diseases such as Parkinson's and Alzheimer's, both patients health records and neuroimaging information are very important for disease understanding.

Goal: Achieving superior classification performance on discriminating patients and controls, with an interpretable learning model based on heterogeneous data structure.
Discriminating

Patients and Health Controls

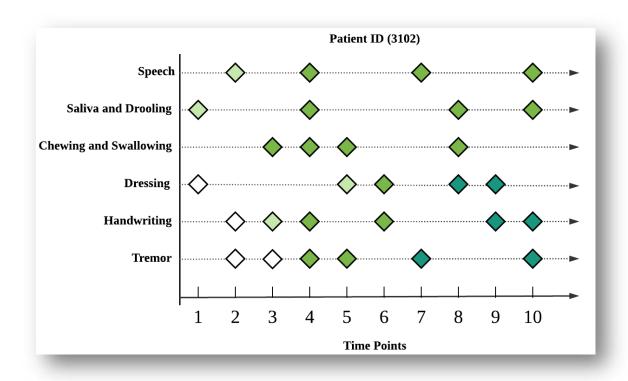
Patients Health Records

Patients Health Records

Patients Mealth Records



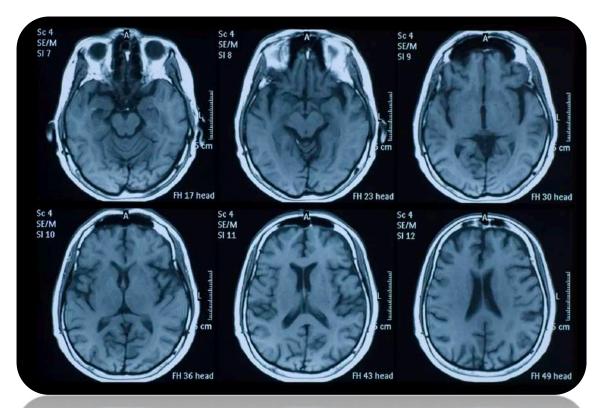
Modality I: Electronic Health Records (Time Series)



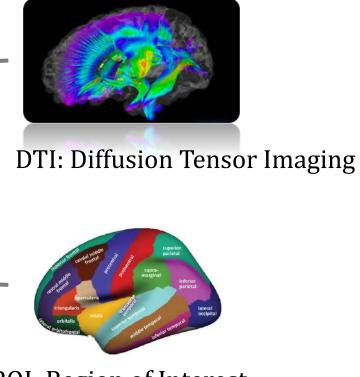
- ✓ Sequential structure
- ✓ Missing Values



Modality II: Neuroimages (Graph Samples)



http://time.com/2860630/mri-scans-can-detect-early-onset-of-parkinsons-study-finds/



ROI: Region of Interest

Desikan-Killiany 84



Challenges

* *Multi-Modality.* The nature of EHR and neuroimage are completely different where EHR data are sequential and a specific brain image is static, i.e., 3-dim tensor or graph.

Solution

We proposed a novel **Mem**ory-based **G**raph **C**onvolutional **N**etwork (MemGCN) to perform integrative analysis with both patient EHRs and neuroimages, using two major components: Graph Convolution and Memory array.



Utilize 3-dimensional brain coordinates of ROIs

Suppose we have a population of M acquisitions,

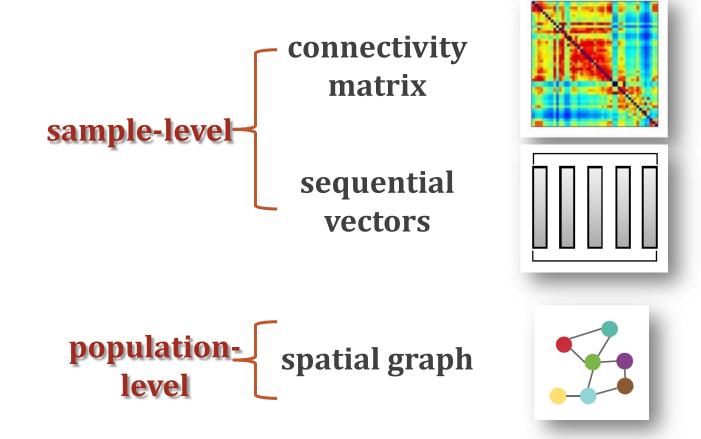
$$\bar{v}_i = \frac{1}{M} (\Sigma_m^M v_{i,m}^x, \Sigma_m^M v_{i,m}^y, \Sigma_m^M v_{i,m}^z), \forall i \in (1, \dots, n).$$

the edges \mathcal{E} can be constructed by

$$w_{ij} = \begin{cases} \exp(-\frac{\|\bar{v}_i - \bar{v}_j\|^2}{2\sigma^2}), & \text{if } i \in \mathcal{N}_j \text{ or } j \in \mathcal{N}_i \\ 0, & \text{otherwise.} \end{cases}$$



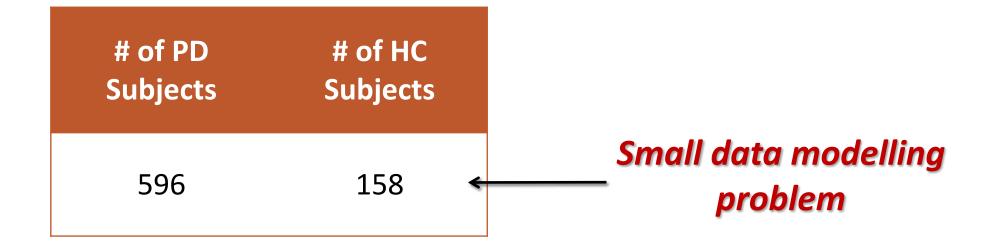
Overview of the input data





How about the amount of training data?





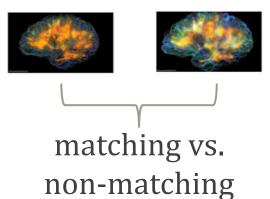


Pairwise-training strategy

# of Matching Samples	# of Non- Matching Samples		
189,713	94,168		

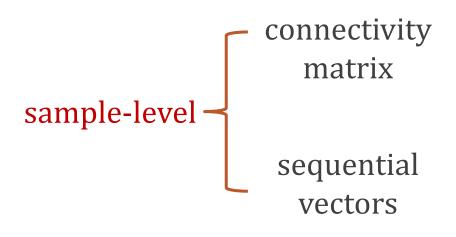


a sample pair

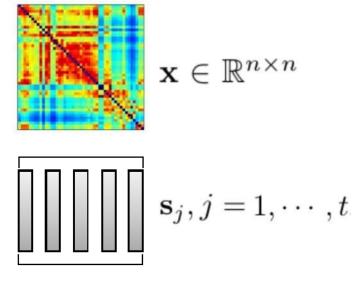


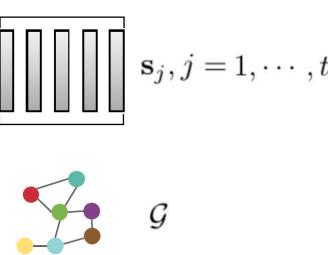
X	\mathbf{y}
PD PD	"same"
PD HC	"different"
HC PD	"different"
HC HC	"same"

The Learning Problem

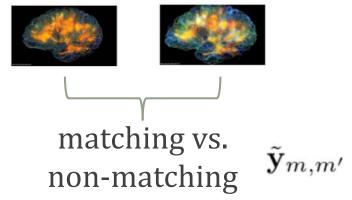


population- spatial graph level



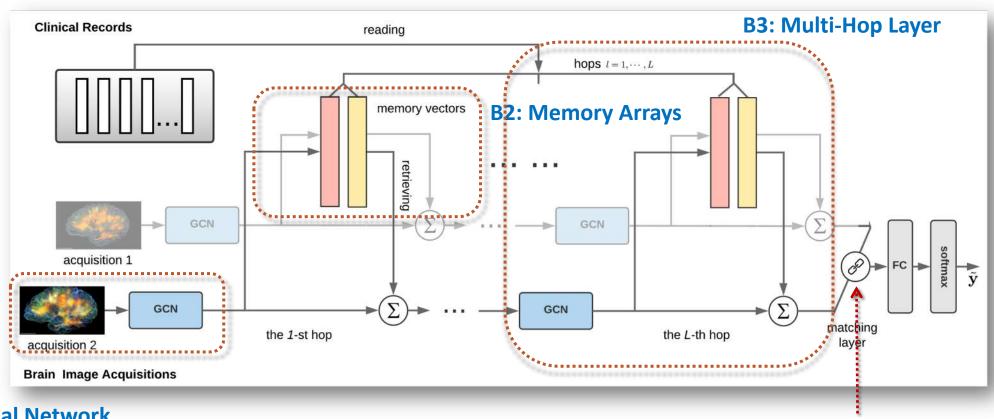








Network Architecture



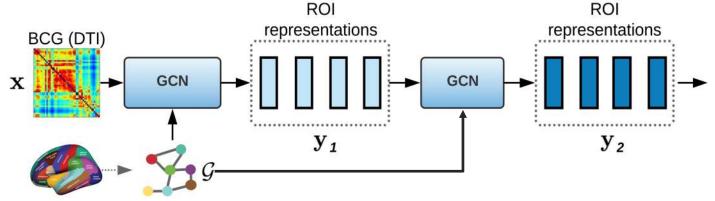
B1: Graph

Convolutional Network

B4: Matching Layer



B1: Graph Convolutional Network (GCN)



ROIs spatial graph)

Graph Fourier Transform

$$\mathbf{x} \star \mathbf{g} = \Phi(\Phi^{\mathrm{T}}\mathbf{x}) \odot (\Phi^{\mathrm{T}}\mathbf{g}) = \Phi g_{\theta}(\mathbf{\Lambda})\Phi^{\mathrm{T}}\mathbf{x}$$
$$= \Phi diag(\hat{q}_{1}, \dots, \hat{q}_{n})\hat{\mathbf{x}}$$

A general operator

$$\mathbf{y}_{m,\,k^{l+1}} = \sum_{k^l=1}^{f_{in}} g_{\theta_{k^l,\,l+1}}(\mathbf{L}) \mathbf{y}_{m,\,k^l} \in \mathbb{R}^n$$

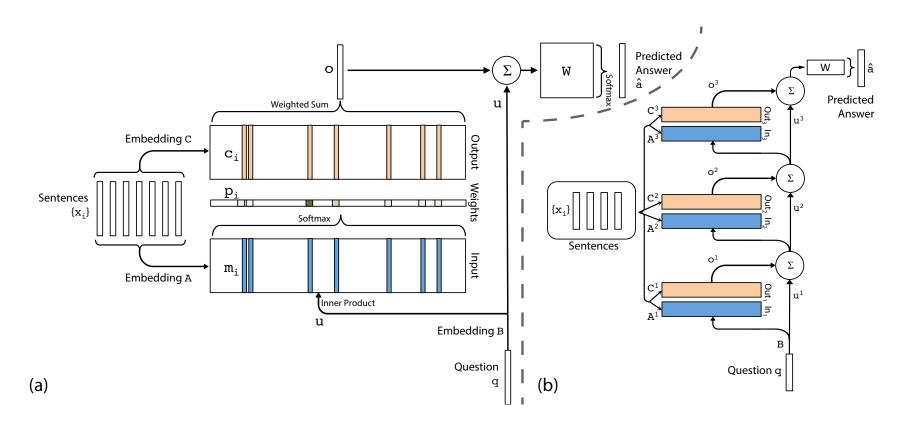
$$g_{\theta}(\boldsymbol{\Delta}) = \sum_{p=0}^{r-1} \theta_p T_p(\tilde{\boldsymbol{\Delta}}) = \sum_{p=0}^{r-1} \theta_p \Phi T_p(\tilde{\boldsymbol{\Lambda}}) \Phi^{\mathrm{T}}$$

ChebNet

Chebyshev polynomial



Preliminary: End-to-End Memory Network



Sukhbaatar et al. '15



- *B2: Memory-Augmented GCN (MemGCN)
 - ✓ Clinical Sequences Reading

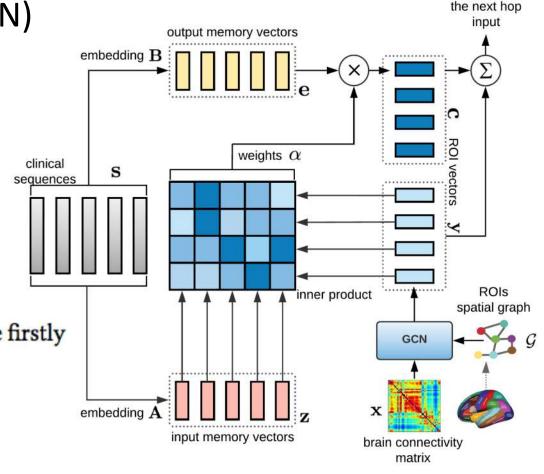
 To embed the sequential vectors $\mathbf{s}_1, \dots \mathbf{s}_t$,

 input memory $\mathbf{z}_j = \mathbf{A}\mathbf{s}_j$ output memory $\mathbf{e}_j = \mathbf{B}\mathbf{s}_j$
 - ✓ Memory Representation Retrieving

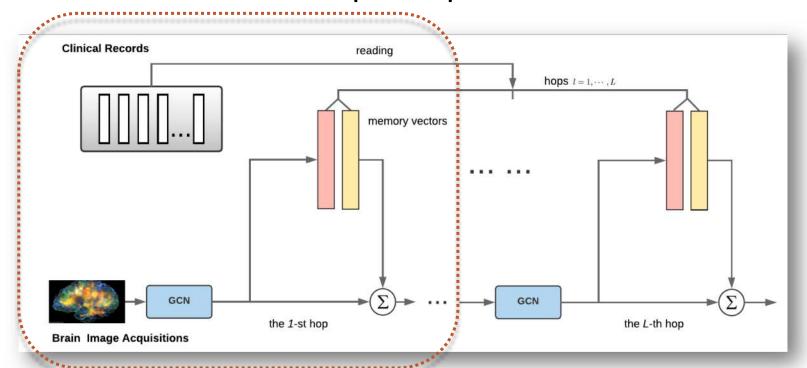
To retrieve memory vectors from the embedding space, we firstly

$$\alpha_{ij} = softmax(\mathbf{y}_i \mathbf{z}_j) = \frac{\exp(\mathbf{y}_i \mathbf{z}_j)}{\sum_{j'=1}^t \exp(\mathbf{y}_i \mathbf{z}_{j'})}$$

$$\mathbf{c}_i = \sum_{j=1}^t \alpha_{ij} \mathbf{e}_i \quad \hat{\mathbf{y}}_i = \mathbf{y}_i + \mathbf{c}_i$$



B3: Extend to multiple hop architecture



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$$\mathbf{A}^1 = \cdots = \mathbf{A}^L$$
 and $\mathbf{B}^1 = \cdots = \mathbf{B}^L$

$$\alpha_{ij}^{l} = \frac{\exp(\mathbf{y}_{i}^{l}\mathbf{z}_{j}^{l})}{\sum_{j'=1}^{t} \exp(\mathbf{y}_{i}^{l}\mathbf{z}_{j'}^{l})}$$
$$\mathbf{c}_{i}^{l} = \sum_{j=1}^{t} \alpha_{ij}^{l}\mathbf{e}_{i}^{l}$$

1-hop MemGCN

the output feature map \hat{y} at the l-th hop can be rewritten as

$$\mathbf{y}^{l+1} = \mathbf{H}\mathbf{y}^l + \mathbf{c}^l, l = 1, \cdots, L$$



- ❖ B4: Matching Layer
 - ✓ Inner Product Matching

$$sim_i(\mathbf{x}_m, \mathbf{x}_{m'}) = (\mathbf{y}_{m,i}^L)^T \mathbf{y}_{m',i}^L, i = 1, \dots, n.$$

✓ Bilinear Matching

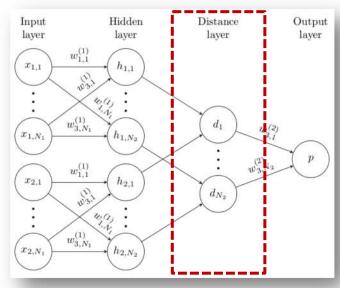
$$sim_{i,j}(\mathbf{x}_m,\mathbf{x}_{m'})=(\mathbf{y}_{m,i}^L)^{\mathrm{T}}\mathbf{M}\mathbf{y}_{m',j}^L,\ i,j=1,\cdots,n.$$

parameter matrix

Siamese-like Network

Impose structure

Learning a metric space

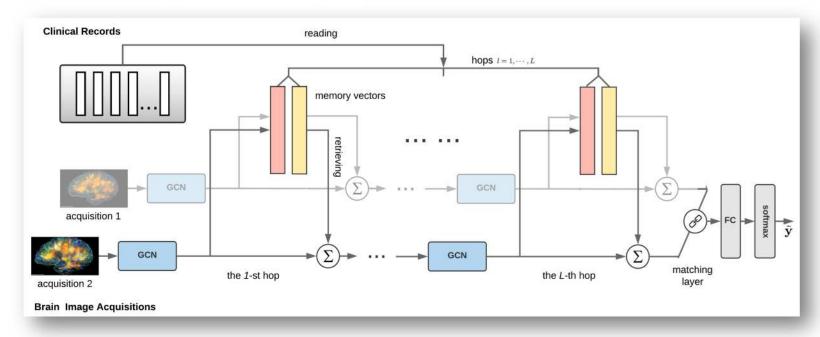


Koch et al. '15

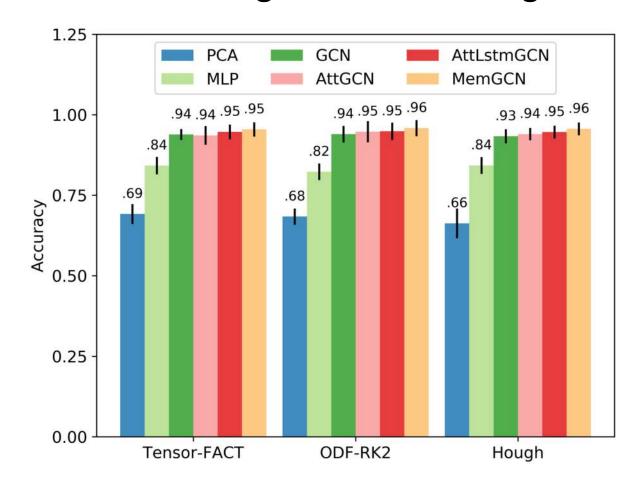
Objective Function (cross entropy, pairwise strategy)

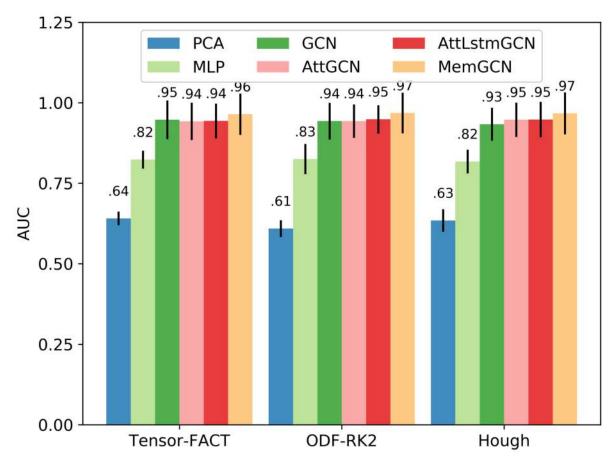
$$\mathcal{L} = \sum_{m,m'}^{N} \tilde{\mathbf{y}}_{m,m'} \log \mathbf{p}_{m,m'} + (1 - \tilde{\mathbf{y}}_{m,m'}) \log(1 - \mathbf{p}_{m,m'}) + \gamma \|\mathbf{\Theta}\|_{2}$$
where $\tilde{\mathbf{y}}_{m,m'}$ denotes the label for sample pair $(\mathbf{x}_{m}, \mathbf{x}_{m'})$

$$\mathbf{p} = softmax(\mathbf{w}_{c}^{\mathsf{T}}\mathbf{r})$$



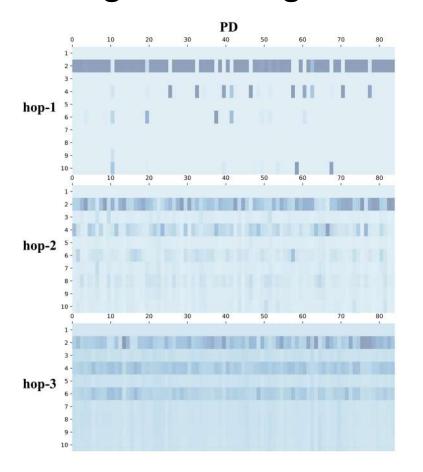
Matching vs Non-Matching Results

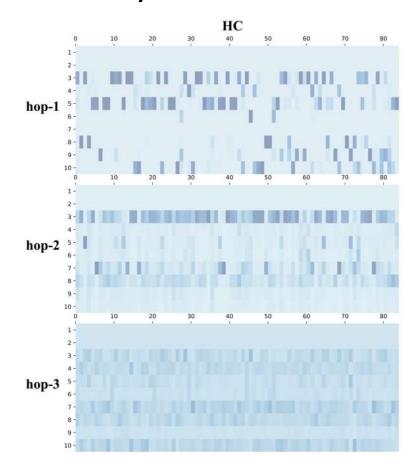


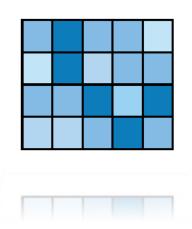




Longitudinal Alignment: Case Study







Visualizations of **attention interaction** matrices for one PD case and one HC case during 3 memory hops.

Interpretation: Learned Similarity (Region Scores)

Average the learned representations for pairwise sample groups (by inner product)

	Motor		Non-motor		Fusion	
	ROI Name	Score	ROI Name	Score	ROI Name	Score
	Right Thalamus Proper	0.9258	Rh Paracentral	0.8563	Rh Pars Opercularis	0.9344
Identical ROIs	Lh Insula	0.9253	Rh Lingual	0.8180	Rh Lateral Occipital	0.8372
(PD Group)	Right Pallidum	0.9226	Right Pallidum	0.8091	Left Accumbens Area	0.7887
(FD Gloup)	Lh Rostral Middle Frontal	0.9210	Lh Parsorbitalis	0.6554	Rh Parahippocampal	0.7827
	Parahippocampal	0.9206	Left Thalamus Proper	0.6387	Rh Frontalpole	0.7742
	Right Putamen	-0.9134	Left Putamen	-0.7423	Right Thalamus Proper	-0.8960
Discriminative ROIs	Right Accumbens Area	-0.9075	Lh Frontal Pole	-0.5754	Left Caudate	-0.8439
(PD vs. HC Group)	Left Hippocampus	-0.9059	Lh Supramarginal	-0.5731	Lh Paracentral	-0.8227
(FD vs. HC Gloup)	Right VentralDC	-0.9058	Lh Inferior Parietal	-0.5693	Lh Middle Temporal	-0.7865
	Left Caudate	-0.9014	Lh Paracentral	-0.4851	Lh Cuneus	-0.7528

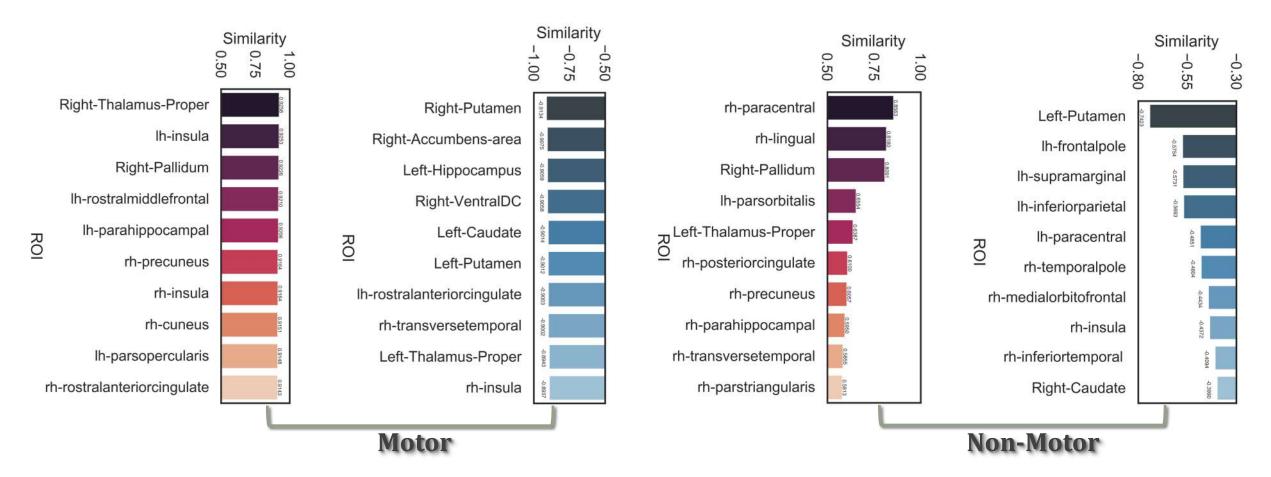
^{*} Lh and Rh are the abbreviations of Left Hemisphere and Right Hemisphere respectively.

Ref: 1. Self-initiated versus externally triggered movements: I. An investigation using measurement of regional cerebral blood flow with PET and movement-related potentials in normal and Parkinson's disease subjects, *Brain*, Volume 118, Issue 4, August 1995, Pages 913–933;

- 2. The functions of the basal ganglia and the paradox of stereotaxic surgery in Parkinson's disease, *Brain*, Volume 117, Issue 4, August 1994, Pages 877–897;
- 3. Cerebral atrophy in Parkinson's disease with and without dementia: a comparison with Alzheimer's disease, dementia with Lewy bodies and controls, *Brain*, Volume 127, Issue 4, April 2004, Pages 791–800.

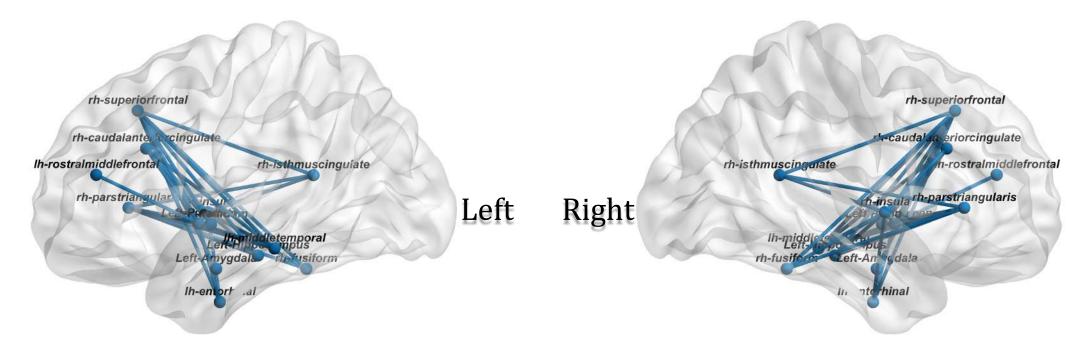


Interpretation: Learned Similarity (Region Scores)



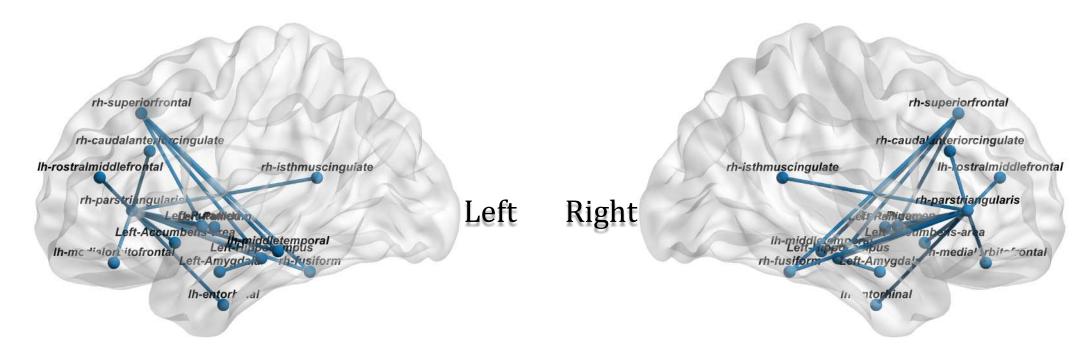


Interpretation: Learned Similarity (Identical Connection)





Interpretation: Learned Similarity (Discriminative Connection)

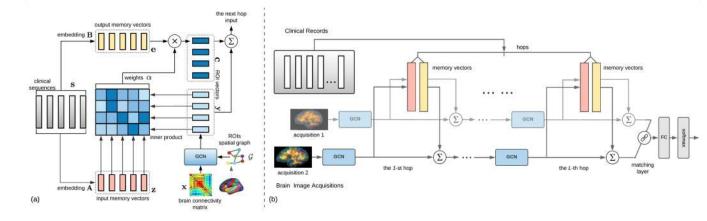


Source Code: https://github.com/sheryl-ai/MemGCN

Xi Zhang, Jingyuan Chou, Fei Wang, Integrative Analysis of Patient Health Records and Neuroimages via Memory-based Graph Convolutional Network. ICDM'18: IEEE International Conference on Data Mining, 2018.



Summary



- ✓ Making a progress in modelling a small cohort data such as PPMI.
- ✓ Interpretable high-level representations extracted from MemGCN are explored.
- ✓ Experiments on classification of Parkinson's Disease demonstrate the superiority of MemGCN.

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Source Code: https://github.com/sheryl-ai/MemGCN

Xi Zhang, Jingyuan Chou, Fei Wang, Integrative Analysis of Patient Health Records and Neuroimages via Memory-based Graph Convolutional Network. ICDM'18: IEEE International Conference on Data Mining, 2018.



Outline



Part 1: Disease Subtyping on Clinical Times Series

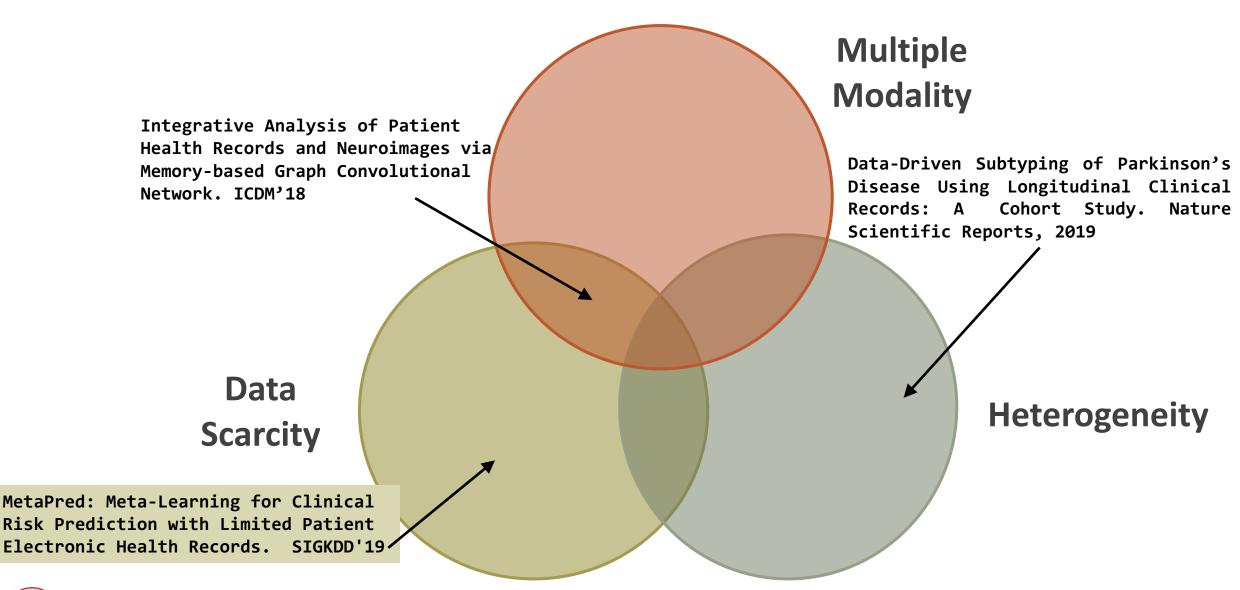


Part 2: Integrative Disease Analysis via Multi-Modality



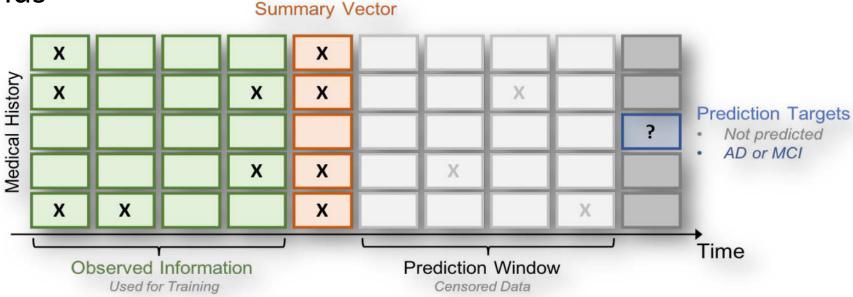
Part3: Meta-Learning on Limited Clinical Resources

Obstacles





Backgrounds



- Patient EHRs: each patient has a sequence of vectors;
- > Predictive models: build for clinical risks, such as in-hospital mortality, hospital readmission, chronic disease onset, condition exacerbation, etc.
 - LR, SVM, k-Nearest Neighbor, Random Forest, MLP;
 - RNN, CNN.



- How about patient samples that are insufficient?
 - > it is expensive and sometimes even impossible for obtaining labelled new samples
 - reusing data on other domain/tasks becomes a feasible strategy
 - transfer learning
 - meta-learning (learning to transfer)



Using the learning experiences from a set of relevant tasks ...



Challenges

- * Data Scarcity: EHRs are suffering sparsity, irregularity, temporality;
- **Label Insufficient:** labelled samples in medicine (patients) are relatively limited, and creates troubles for building an effective predictive model.

Solution

We proposed a **MetaPred**, a model agnostic meta-learning framework for low-resource predictive modelling with patient EHRs.

Motivation

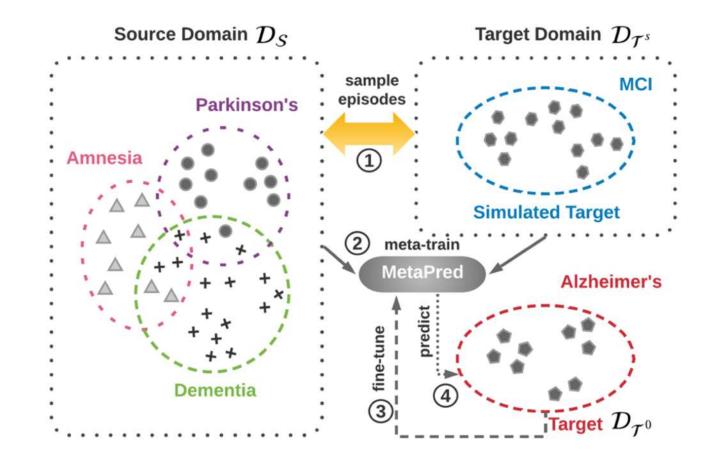
Goal: is to predict the risks of target disease with few labeled patients, which give rise to a low-resource classification.

The idea: is to take advantage of labeled patients from other relevant high-resource domains and design the learning to transfer framework with sources and a simulated target.



Meta-Learning for Clinical Risk Prediction

- Problem Setup
 - 4 steps:
 - ✓ sample episode
 - ✓ meta-train
 - ✓ fine-tune
 - ✓ predict





 $\theta^* \stackrel{\text{meta-learning}}{\sim} \theta_1^*$

Preliminary: MAML

Meta-learning, also known as <u>learning to learn</u>, aims to optimize the objective over a variety of learning tasks \mathcal{T} which are associated with the corresponding datasets $\mathcal{D}_{\mathcal{T}}$.

$$\Theta^* = \arg\min_{\Theta} \mathbb{E}_m \mathbb{E}_{\mathcal{D}_{epi}^m \sim p(\mathcal{D}_{\mathcal{T}})} \mathcal{L}_{\Theta}(\mathcal{D}_{\mathcal{T}})$$

$$\underline{Loss \ function}: \quad \mathcal{L}_{\Theta} = \frac{1}{|\mathcal{D}_{epi}^{te}|} \sum_{(\mathbf{X}_{i}, \mathbf{y}_{i}) \in \mathcal{D}_{epi}^{te}} \mathcal{L}_{\Theta} \left((\mathbf{X}_{i}, \mathbf{y}_{i}); \mathcal{D}_{epi}^{tr} \right)$$

Finn et al. '17



Two-level adaptation

$$\Theta^* = \text{Learner}(\mathcal{T}^s; \text{MetaLearner}(\mathcal{S}^1, \cdots, \mathcal{S}^{K-1}))$$



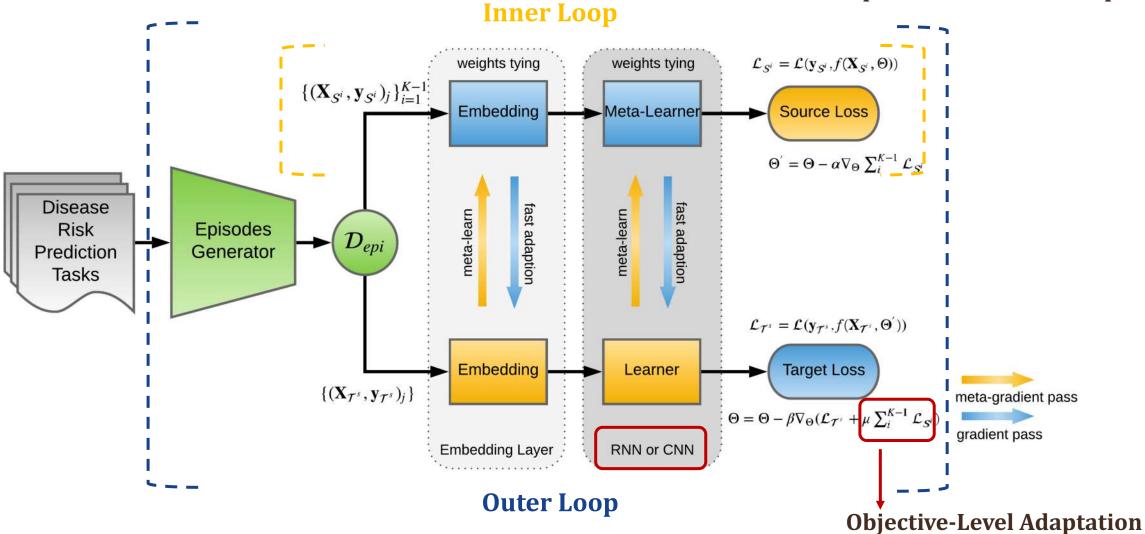
$$\Theta^* = \text{Learner}\left(\mathcal{T}^s, \{\mathcal{S}^i\}_i^{K-1}; \text{MetaLearner}(\{\mathcal{S}^i\}_i^{K-1})\right)$$

- ✓ <u>Parameter-level adaptation</u>: Model-agnostic, typing parameters for source/target domain.
- ✓ *Objective-level adaptation*: bounded target risk via empirical risk of source domains.

Meta-Learning on Limited Clinical Resource: Framework

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Optimization-Level Adaptation



Objective adaptation

Domain Adaptation Bound Ben-David et al. '10

Theorem 1 For a hypothesis h,

$$\epsilon_{T}(h) \leq \epsilon_{S}(h) + d_{1}(\mathcal{D}_{S}, \mathcal{D}_{T})$$

$$+ \min \left\{ E_{\mathcal{D}_{S}} \left[|f_{S}(\mathbf{x}) - f_{T}(\mathbf{x})| \right], E_{\mathcal{D}_{T}} \left[|f_{S}(\mathbf{x}) - f_{T}(\mathbf{x})| \right] \right\}.$$

$$\mathcal{L}_{\mathcal{T}}(f_{\Theta'}) = \mathcal{L}_{\mathcal{T}^s}(f_{\Theta'}) + \mu \sum_{i}^{K-1} \mathcal{L}_{\mathcal{S}^i}(f_{\Theta}) \qquad \text{all the domains share the same feature space}$$

$$= \sum_{\mathcal{D}_{epi}^{\mathcal{T}^{S}}} \mathcal{L}\left(\mathbf{y}_{\mathcal{T}^{S}}, f(\mathbf{X}_{\mathcal{T}^{S}}, \Theta')\right) + \left(\mu \sum_{i}^{K-1} \sum_{\mathcal{D}_{epi}^{S^{i}}} \mathcal{L}\left(\mathbf{y}_{S^{i}}, f(\mathbf{X}_{S^{i}}, \Theta)\right)\right)$$

Parameter adaptation

Feedforward 1:

Embedding → LSTM→ MLP

Feedforward 2:

Embedding → 1d-CNN→ MLP

Obj: Source **Domains**

Obj: Simulated Target Domain

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```
Algorithm 1 MetaPred Training
```

Require: Source domains S^i ; Simulated target domain T^s ;

Require: Hyperparameters α , β , μ ;

1: Initialize model parameter Θ randomly

2: **while** Outer-Loop not done **do**

Sample batch of episodes $\{\mathcal{D}_{epi}\}$ from \mathcal{D}_{S^i} and $\mathcal{D}_{\mathcal{T}^s}$

while Inner-Loop not done do

$$\{(\mathbf{X}_{S^{i}}, \mathbf{y}_{S^{i}})\}_{i=1}^{K-1}, \{(\mathbf{X}_{T^{s}}, \mathbf{y}_{T^{s}})\} = \{\mathcal{D}_{epi}\}$$

5: $\{(\mathbf{X}_{S^{i}}, \mathbf{y}_{S^{i}})\}_{i=1}^{K-1}, \{(\mathbf{X}_{T^{s}}, \mathbf{y}_{T^{s}})\} = \{\mathcal{D}_{epi}\}$ 6: Compute $\mathcal{L}_{S^{i}} = \mathcal{L}(\mathbf{y}_{S^{i}}, f(\mathbf{X}_{S^{i}}, \Theta)), i = 1, \dots, K-1$

Parameter fast adaption with gradient descent:

$$\Theta' = \Theta - \alpha \nabla_{\Theta} \sum_{i}^{K-1} \mathcal{L}_{\mathcal{S}^{i}}$$

end while

10: Compute $\mathcal{L}_{\mathcal{T}^s} = \mathcal{L}(\mathbf{y}_{\mathcal{T}^s}, f(\mathbf{X}_{\mathcal{T}^s}, \Theta'))$

Update $\Theta = \Theta - \beta \nabla_{\Theta} (\mathcal{L}_{\mathcal{T}^s} + \mu \sum_{i=1}^{K-1} \mathcal{L}_{S^i})$ using Adam

12: end while





Disease	ICD-9 Codes
Mild Cognitive Impairment Alzheimer's Disease	331.83, 331.89, 331.9
Parkinson's Disease	331.0, 331.2, 331.6, 331.7 332.* Target :
Dementia	290.*, 291.*, 294.*, 331.82
Amnesia	780.93
Huntington's Disease	333.4
Mechanical Obstructions	331.3, 331.4, 331.5
Frontotemporal Dementia	331.1, 331.11, 331.19

	Domain	Case	Control	# of visit	Ave. # of visit
	MCI	1,965	4,388	161,773	22.24
5	Alzheimer's	1,165	4,628	136,197	20.73
	Parkinson's	1,348	3,588	105,053	20.01
	Dementia	3,438	1,591	98,187	18.06
	Amnesia	2,974	4,215	180,091	21.60



 $^{^{\}ast}$ means that all the codes in this diagnosis group are included.

Prediction Accuracy

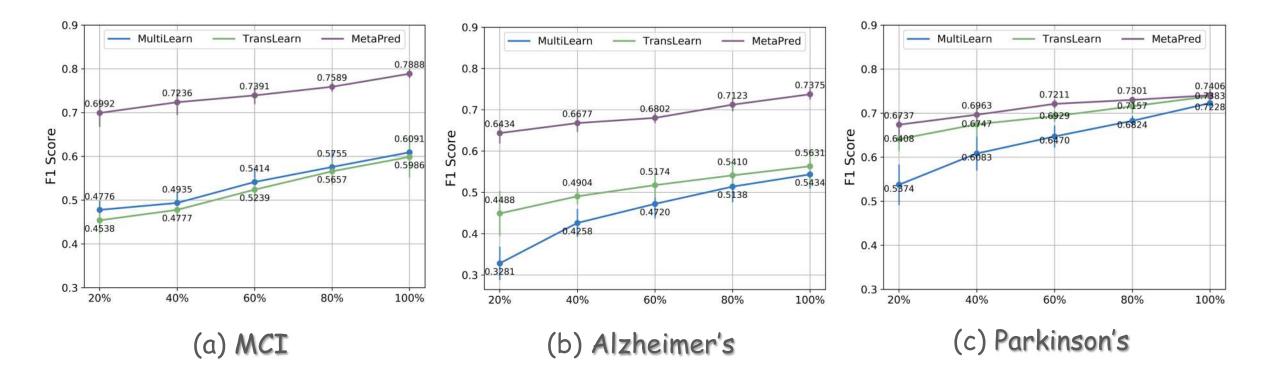


Training Data	Model	M	CI	Alzheimer's Disease		
Training Data	Woder	AUCROC	F1 Score	AUCROC	F1 Score	
Eulles Comencie al	LR	0.5861 (.01)	0.3813 (.02)	0.5369 (.01)	0.2216 (.02)	
	kNN	0.6106 (.01)	0.4540 (.01)	0.6713 (.02)	0.4686 (.03)	
	RF	0.6564 (.01)	0.4998 (.01)	0.6300 (.02)	0.4111 (.04)	
Fully Supervised	MLP	0.6515 (.01)	0.5077 (.01)	0.6639 (.02)	0.4901 (.03)	
	CNN	0.6999 (.01)	0.5816 (.02)	0.6755 (.03)	0.4935 (.04)	
	LSTM	0.6874 (.01)	0.5666 (.02)	0.6902 (.01)	0.5316 (.02)	
Low-Resource	Meta-CNN	0.7624 (.02)	0.6992 (.02)	0.7682 (.01)	0.6434 (.03)	
	Meta-LSTM	0.7876 (.02)	0.7225 (.02)	0.7464 (.02)	0.6170 (.03)	
Enlls Eine Tour 1	Meta-CNN	0.8470 (.01)	0.7888 (.02)	0.8461 (.01)	0.7375 (.01	
Fully Fine-Tuned	Meta-LSTM	0.8477 (.01)	0.7963 (.02)	0.8232 (.01)	0.7364 (.01)	

Outperform fully supervised model

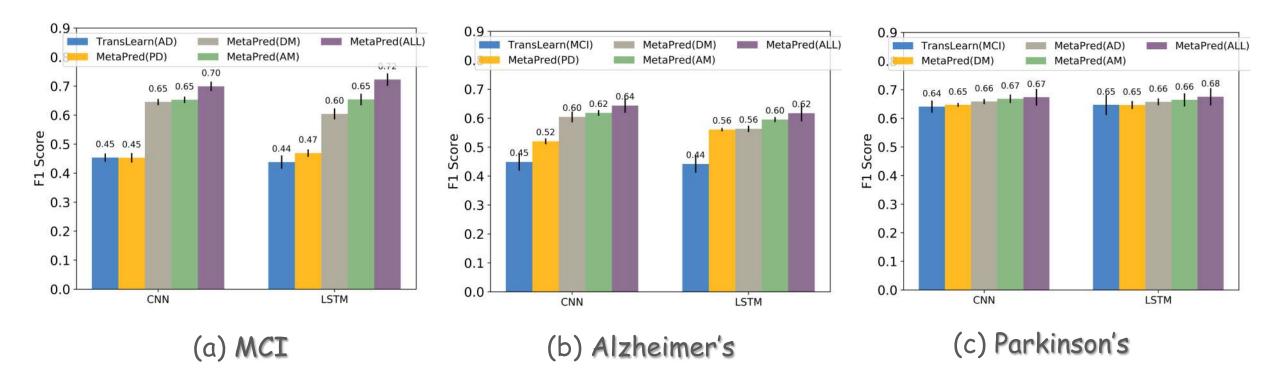


Fine-tuning on Target Domain

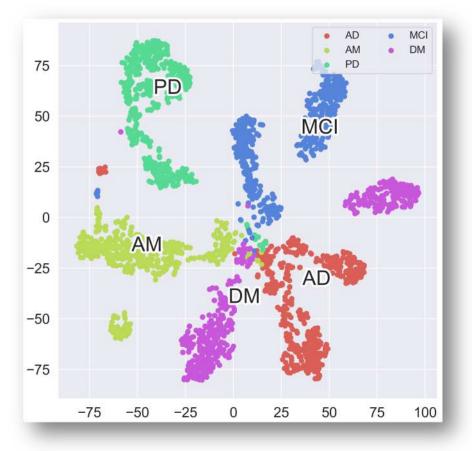




Different combinations of source disease domains







t-SNE Visualization

Source Code: https://github.com/sheryl-ai/MetaPred

Xi Zhang, Fengyi Tang, Hiroko Dodge, Jiayu Zhou, Fei Wang, MetaPred: Meta-Learning for Clinical Risk Prediction with Limited Patient Electronic Health Records. SIGKDD'19: ACM SIGKDD International Conference on Knowledge Discovery and Data Mining, 2019.

Summary

- ✓ Leverages deep predictive modeling with the model agnostic metalearning to exploit the medical records from high-resource domain.
- ✓ Introduce two different kinds of adaptation, which are parameter-level adaptation, objective-level adaptation.
- ✓ Extensive evaluation involving 5 cognitive diseases is conducted on realworld EHR data for risk prediction tasks.

Source Code: https://github.com/sheryl-ai/MetaPred

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Hiroko H. Dodge UMich



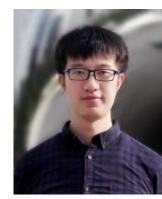
Fei Wang Cornell



Andy Tang MSU



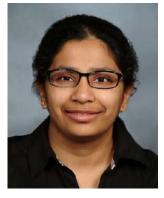
Jiayu Zhou MSU



Jian Liang Tencent



Claire Henchcliffe Cornell



Harini Sarva Cornell



Yize Zhao Yale



Jingyuan Chou UVA



Cao (Danica) Xiao IQVIA

Thank You!

Q & A