

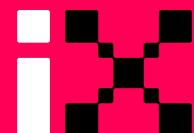
Revolutionizing Connectomics using Generative, federated and holistic learning

Islem Rekik

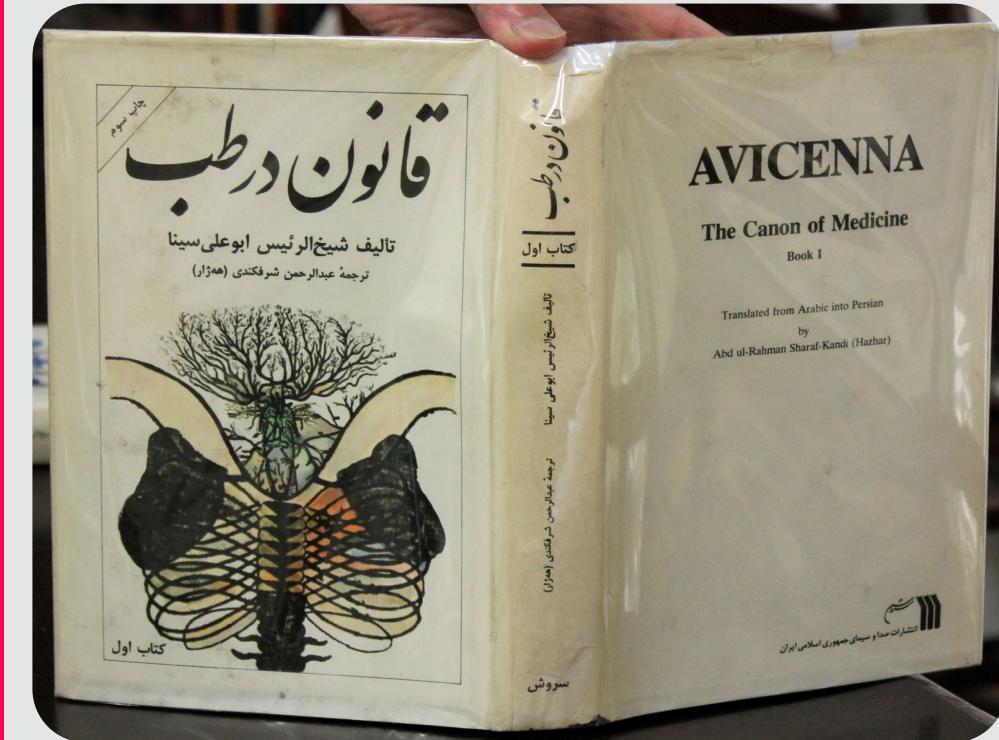
i.rekik@imperial.ac.uk

<https://basira-lab.com/>

Imperial College
London



"The physician should not treat the disease but the patient who is suffering from it."
Avicenna (980–1037)





FORBES > INNOVATION > ENTERPRISE TECH

The 10 Biggest Trends Revolutionizing Healthcare In 2024

Bernard Marr Contributor ◉

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Oct 3, 2023, 03:15am EDT

Facebook icon A longer-living population, the emergence of transformative technologies with applications across the healthcare spectrum, and continued global economic uncertainty. These are the key societal drivers that will impact healthcare in 2024.



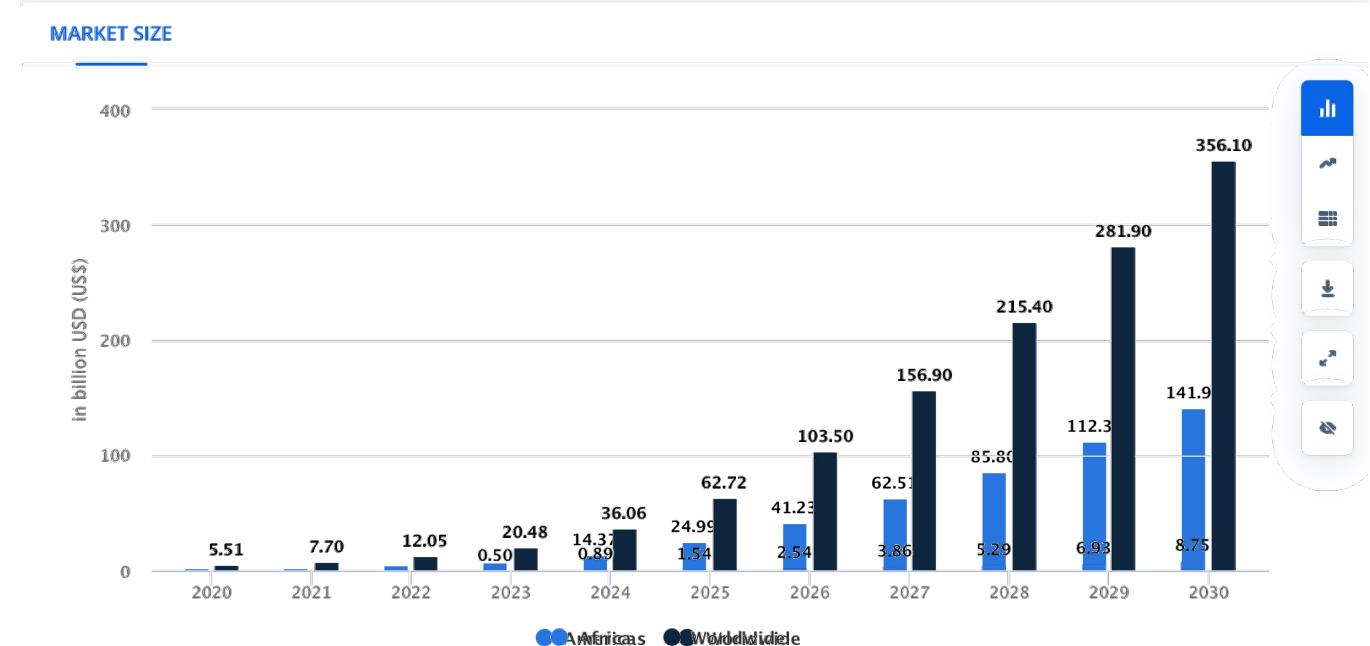
The 10 Biggest Trends Revolutionizing Healthcare In 2024 ADOBE STOCK

Personalized Medicine

Generative AI In Healthcare

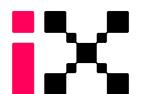


Generative AI market size worldwide



Notes: Data was converted from local currencies using average exchange rates of the respective year.

Source: Statista Market Insights



Three trends in the AI-powered healthcare revolution

Generative learning

Generate *multidimensional* brains

Generate population brain templates

Federated learning

Federate to generate the future of the brain

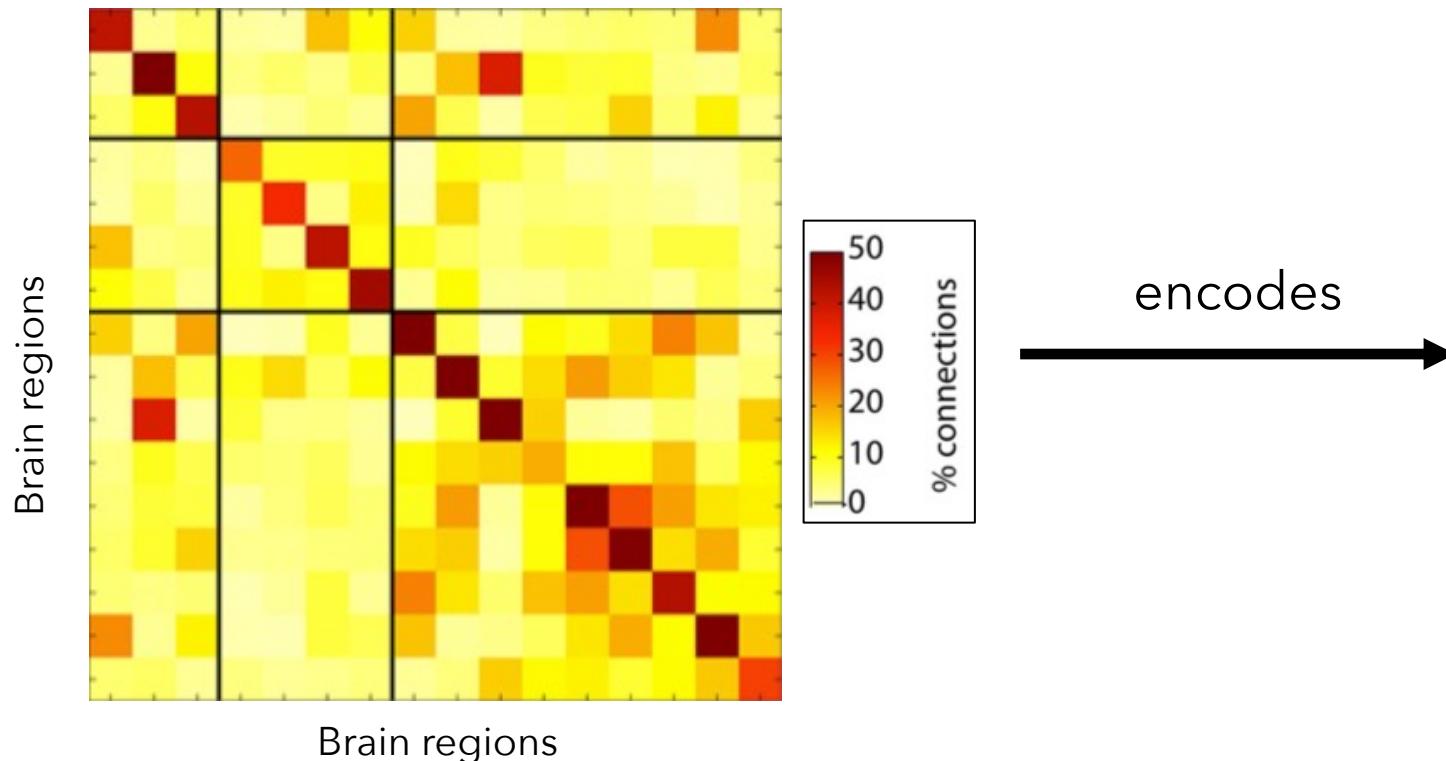
Federate heterogeneous diagnostic tasks

Holistic learning

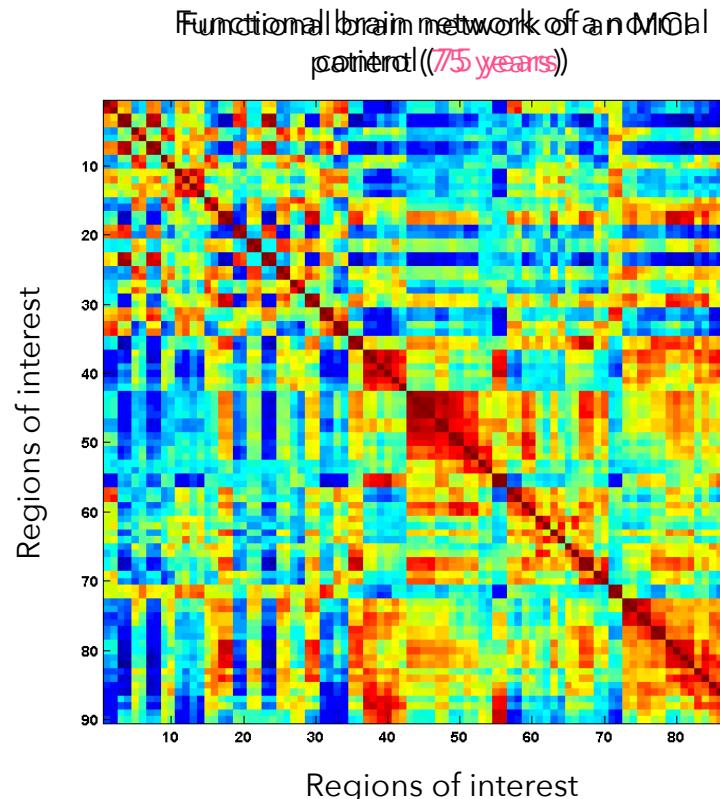
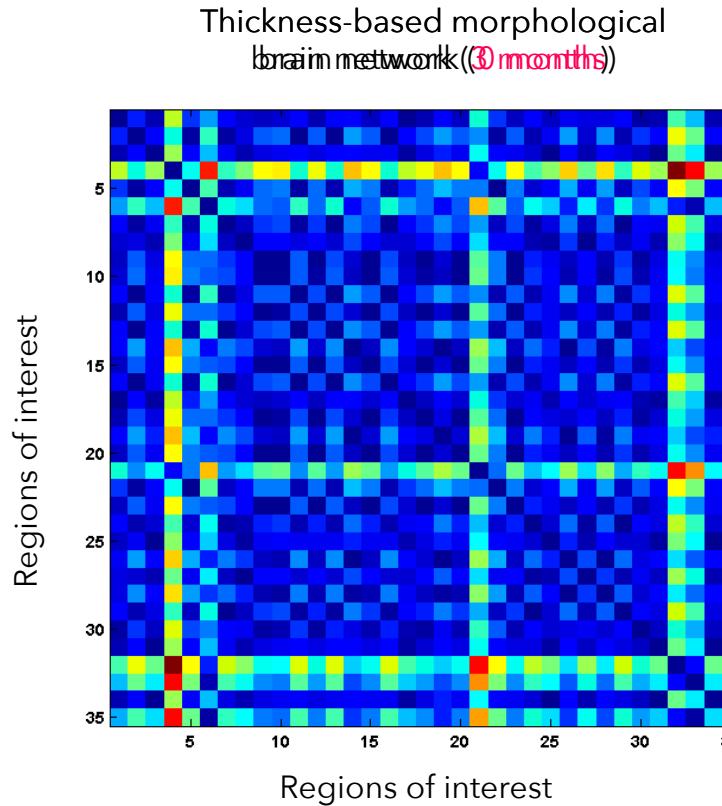
Insights into the future of an inclusive and holistic AI in medicine



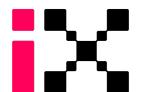
EXAMPLE (2): BRAIN GENERATION



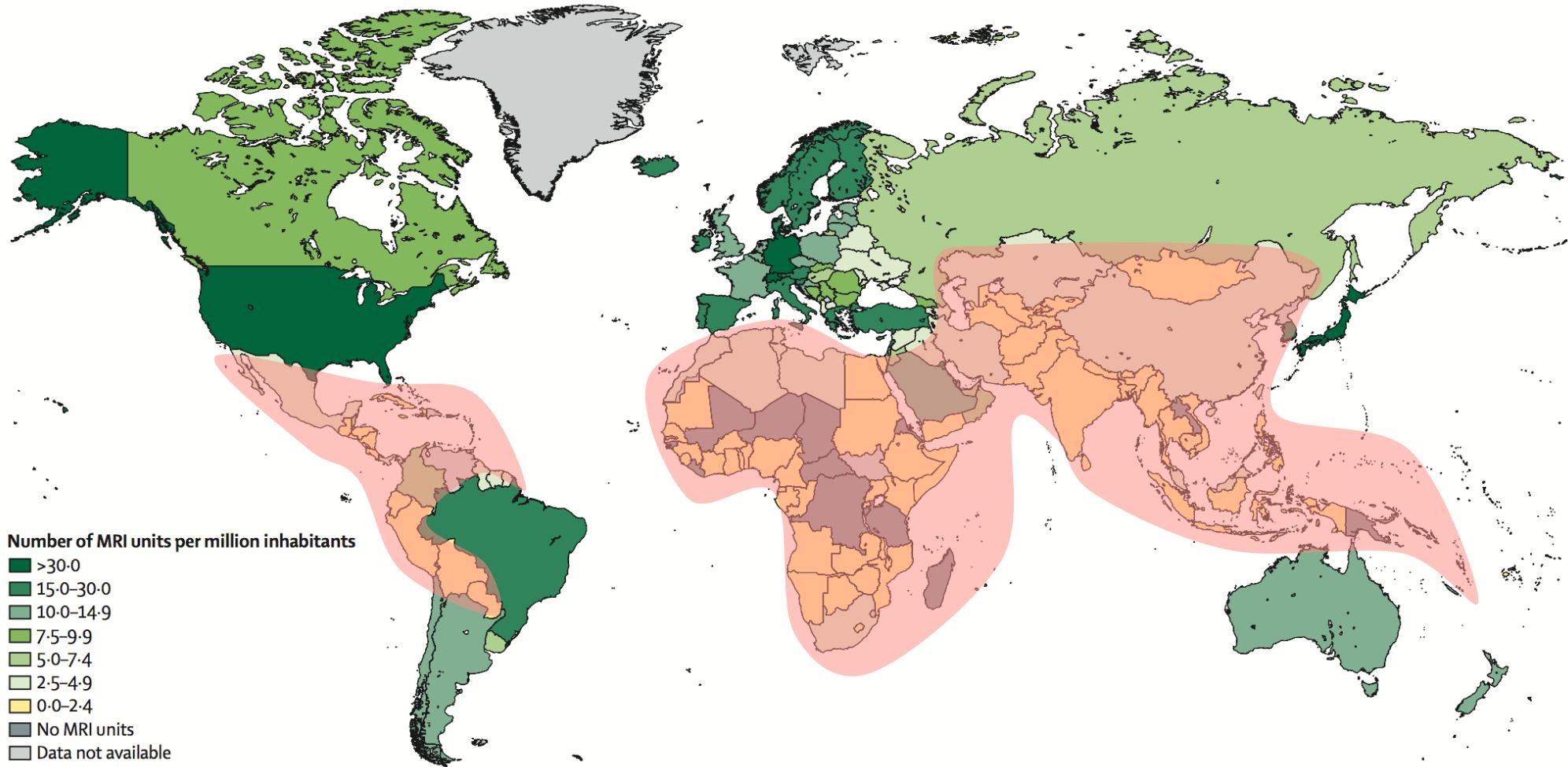
FROM THE CRADLE TO THE GRAVE



A mechanistic understanding of the **human connectome** is not only important for understanding **normal brain function**, but will also help tackle the even **more complex pathological brain**.



Mind the medical imaging gap!



Estimates of the number of MRI units per million inhabitants. The map was produced by the International Atomic Energy Agency (Vienna, Austria).¹



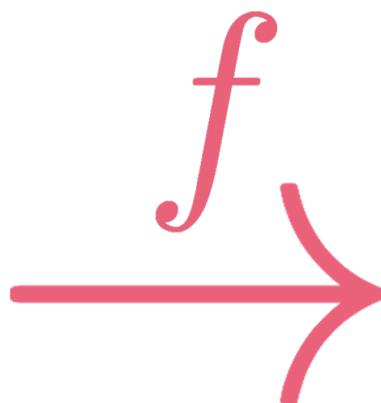
¹Hricak, Hedvig, et al. "Medical imaging and nuclear medicine: a Lancet Oncology Commission." **The Lancet Oncology** 22.4 (2021): e136-e172.

Can we diagnose early and better with **minimal resources (limited data)** and **at lower costs**?

“data”

X

- Affordable to acquire
- Accessible
- Low-middle quality

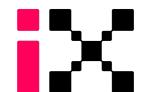
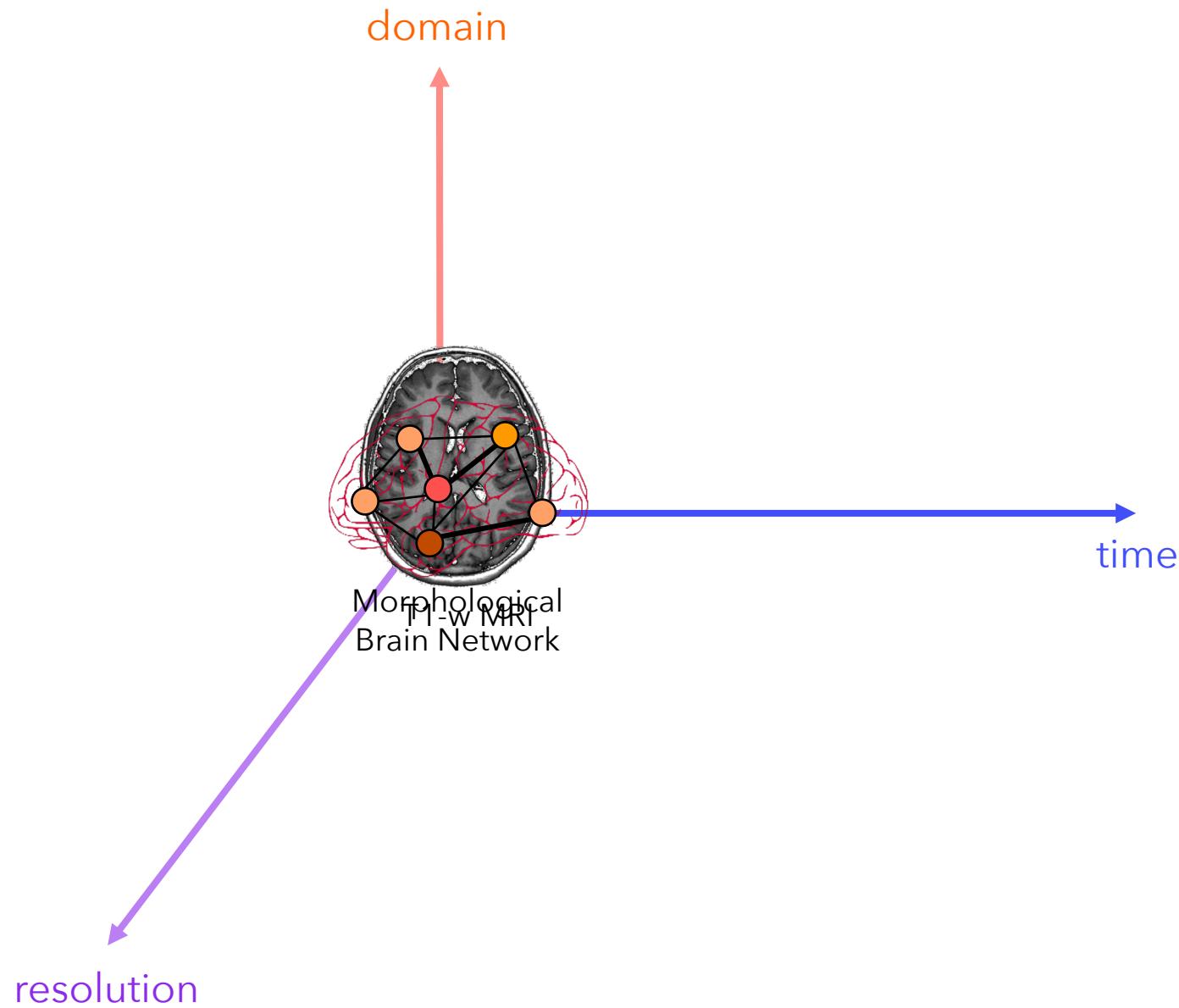


“no-data”

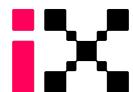
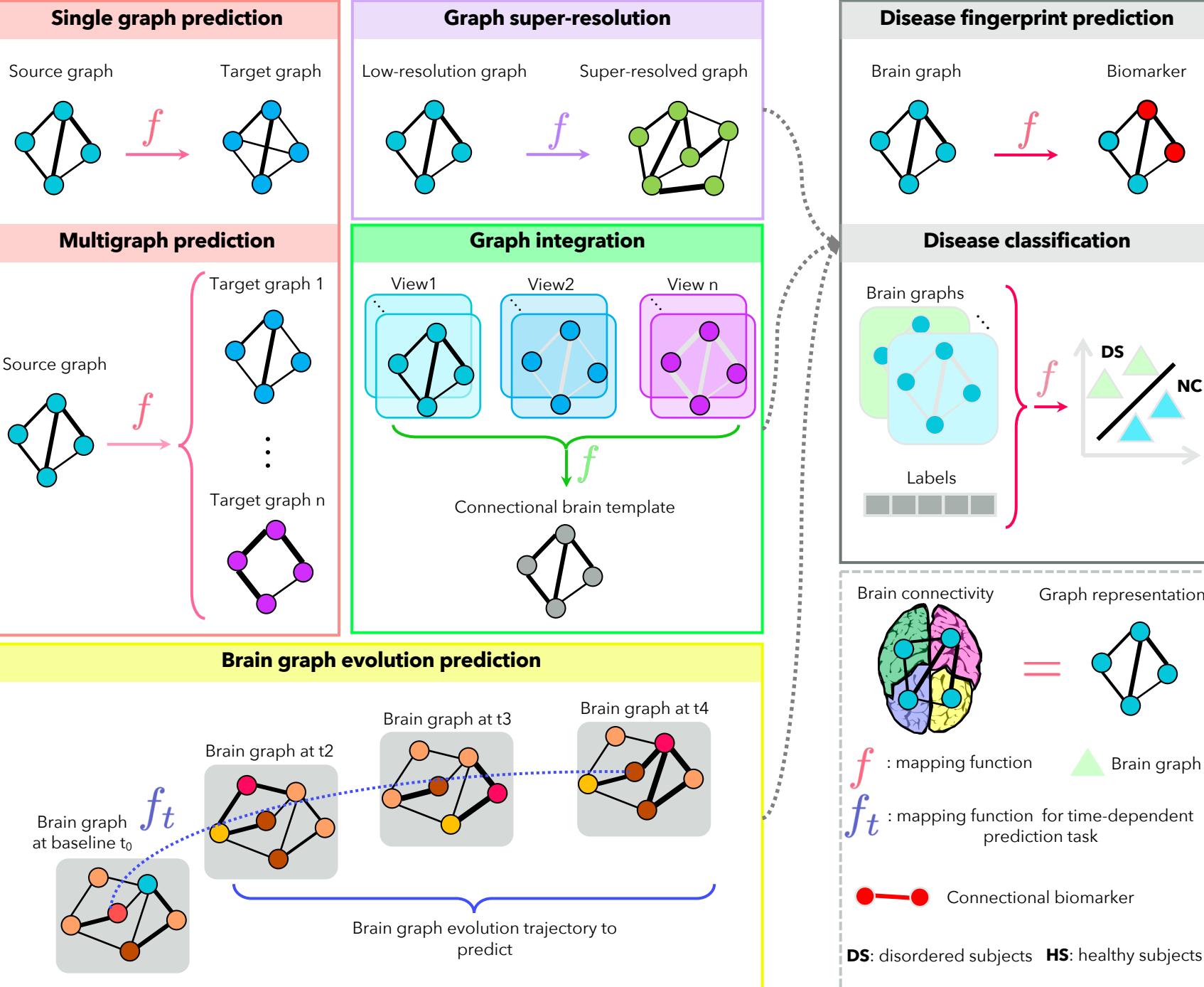
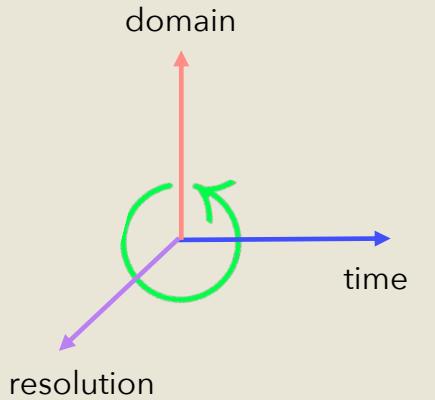
X̂

- Costly to acquire
- Inaccessible in many countries
- High quality

Pioneering and developing **70+ predictive/generative learning** works in **network neuroscience (2019-2023)**



Predictive intelligence dimensions



Graph neural networks in network neuroscience

arXiv:2106.03535v1 [cs.LG] 7 Jun 2021

A-1 Single graph prediction

B- Super-resolution graph prediction

E- Disease fingerprint prediction

Source graph

Source graph

Brain graph at baseline

Graph Neural Networks in Network Neuroscience

In Geometric Deep Learning (GDL), one of the most popular learning methods is the Graph Neural Network (GNN), which applies convolutional layers to learn the topological structure of the input graph. GNN has recently been used for the analysis of different types of the human connectome, such as structural, functional, and morphological networks derived respectively from Diffusion Tensor Imaging (DTI), functional magnetic resonance imaging (fMRI), and T1-w MRI data. Such a nascent field has achieved significant performance improvements over traditional deep neural networks in the early diagnosis of brain disorders. Thus, we list here all MICCAI papers of 4 years (2017–2020), some journals, and IPMI (2017–2019) papers that are embedded into multiple tasks related to the disease diagnosis:

- Predicting one or multiple modalities from a single or different modalities (e.g., data synthesis)
- Predicting high-resolution data from low-resolution data
- Predicting the brain evolution trajectory
- Multiple network integration and fusion
- Computer-aided prognostic methods (e.g., for brain diseases)
- Biomarker Identification
- Extra works related to machine learning

If you like to update the file by adding the unlisted open-source articles, feel free to open an issue or submit a pull requests. You can also directly contact Alaa Bessadok at alaa.bessadok@gmail.com. All contributions are very welcome!

arXiv link

The full paper is downloadable at <https://arxiv.org/pdf/2106.03535.pdf>

GitHub link: <https://github.com/basiralab/GNNs-in-Network-Neuroscience>

Marker

Region

NC

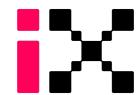
Presentation

in graph

Independent

subjects

.....	2
CE?	4
.....	4
.....	6
.....	6
.....	12
.....	13
.....	17
.....	17
.....	21



Reproducible, inclusive and open science



20 GitHub code video demo (since 2020)

BASIRA Lab GitHub

BASIRA Lab - 1 / 20

2 GitHub demo of MultiGraphGAN for Brain... 3:08
BASIRA Lab

3 GitHub code of GSR-Net for Graph Super-Resolution... 3:48
BASIRA Lab

4 GitHub code of RESNets (Goktas et al., PRIME 2020) 3:36
BASIRA Lab

5 GitHub code of graph GAN (gGAN) (Gurler et al, PRIME...) 3:36
BASIRA Lab

6 GitHub Code of Supervised Diffusion of Biological Graph... 5:09



65+ GitHub codes

Pinned

DGL Public Deep Graph-Based Learning Course (ICL, Computing)
Jupyter Notebook ★ 12 2 215 28

RegGNN Public Regression Graph Neural Network (regGNN) for cognitive score prediction.
Python ★ 68 18

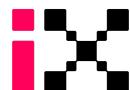
FALCON Public FALCON: Feature-Label Constrained Graph Net Collapse for Memory Efficient GNN
C 66 1

GNNs-in-Network-Neuroscience Public A review of papers proposing novel GNN methods with application to brain connectivity published in 2017-2020.
215 28

survey-multigraph-integration-methods Public Comparative survey of multigraph integration methods
Jupyter Notebook ★ 10 3

RISE-MICCAI/Journal-Club Public The RISE Journal Club aims to create a friendly environment to discuss the latest state-of-the-art papers in the areas of medical image analysis, AI and computer vision. The moderators will briefly...
66 1

<https://github.com/basiralab>



Reproducible, inclusive and open science

Strongly Topology-preserving GNNs for Brain Graph Super-resolution

Pragya Singh and Islem Rekik



BASIRA Lab, Imperial-X and Department of Computing,
Imperial College London, London, UK

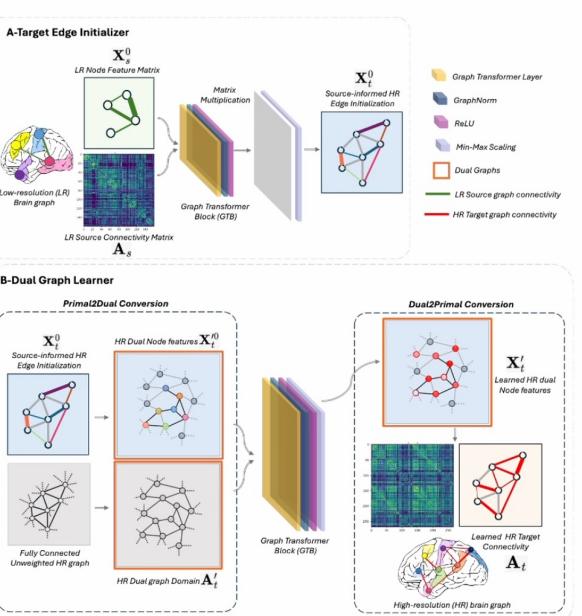
This work is part of my research project at BASIRA lab:

<https://basira-lab.com>

GitHub code: <https://github.com/basiralab/STP-GSR>

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▶ ◀ ↻ 0:01 / 7:25 • Introduction >



PRIME-MICCAI

MICCAI 2024
MOROCCO

50+ oral paper presentations

BASIRA research publications

BASIRA Lab - 2 / 53



1



Topology-aware Graph Neural Network | Distinguished* MSc |...

BASIRA Lab

2



Topology-Preserving GNNs for Brain Graph Super-Resolution |...

BASIRA Lab

3



Generative Hypergraph Neural Networks for Data Fusion | **Or...

BASIRA Lab

4



Diversity-Based Hyperedge Sampling in GNN Learning |...

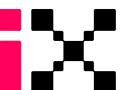
BASIRA Lab

5



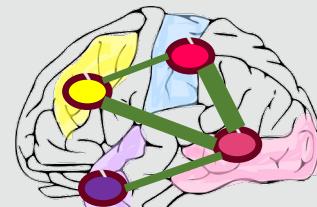
A Universal Federation of a Mixture of Highly Heterogeneou...

BASIRA Lab

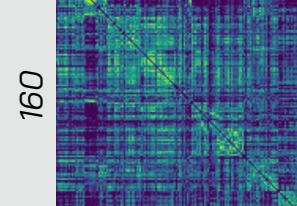


Cross-resolution prediction

**Low Resolution (LR)
Brain Graph**



LR Connectivity Matrix

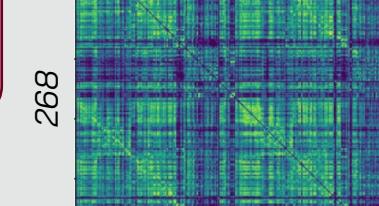
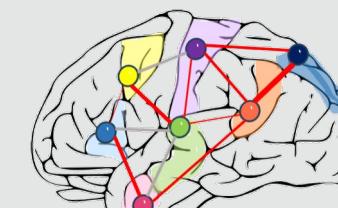


Dosenbach
parcellated

Generate HR Brain Graphs
from LR Brain Graphs,
using **GNN Models**



**High Resolution (HR)
Brain Graph**



Shen parcellated

**HR Connectivity
Matrix**

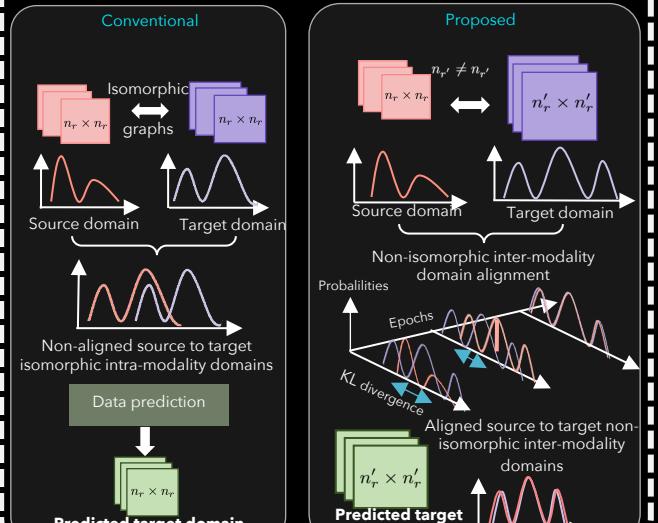


Non-isomorphic Inter-modality Graph Alignment and Synthesis for Holistic Brain Mapping

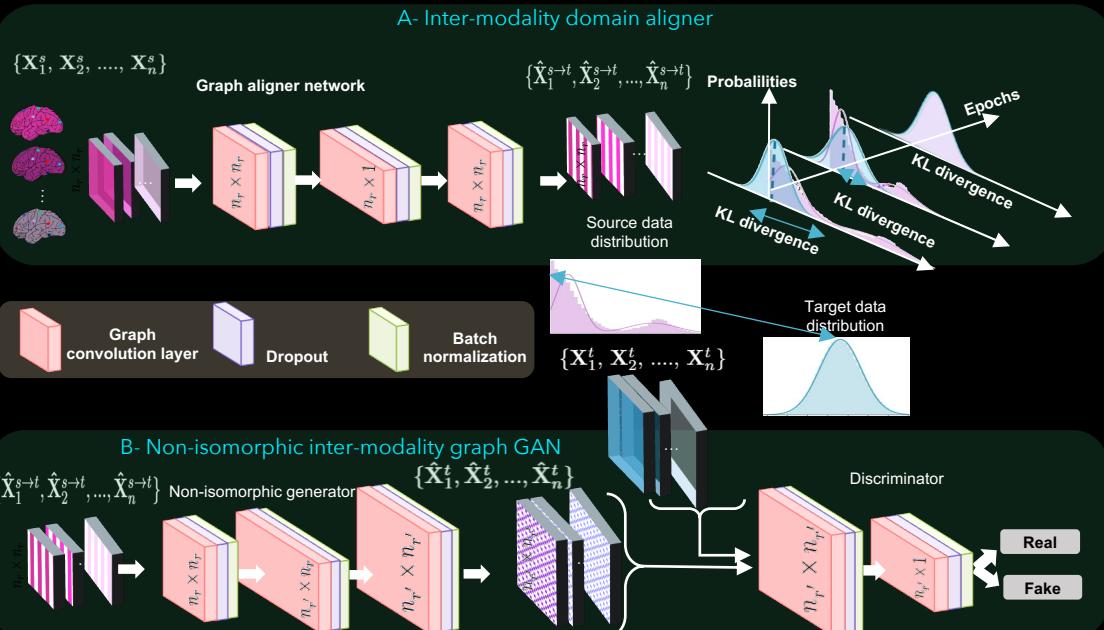
Islem Mhiri , Ahmed Nebli , Mohamed Ali Mahjoub and Islem Rekik

Problem statement

How to predict inter-modality non-isomorphic brain graphs from a base graph ?



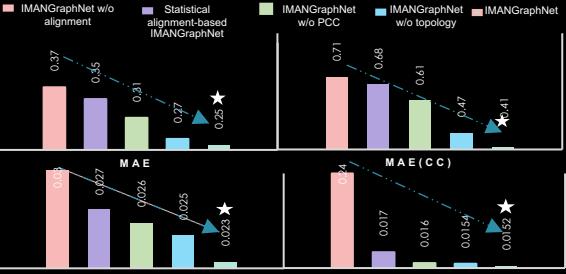
Proposed IMANGraphNet Method



Results

Dataset evaluation
SLIM¹ dataset: 150 functional connectomes (160 × 160)
150 morphological connectomes (35 × 35)

Prediction results using different evaluation metrics



This work *is accepted* at IPMI 2021 and selected for an *oral presentation*.
GitHub code: <https://github.com/basiralab/IMANGraphNet>



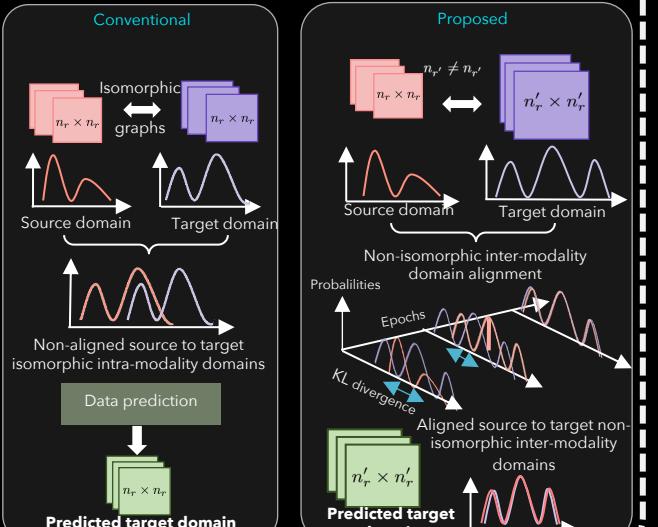


Non-isomorphic Inter-modality Graph Alignment and Synthesis for Holistic Brain Mapping

Islem Mhiri , Ahmed Nebli , Mohamed Ali Mahjoub and Islem Rekik

Problem statement

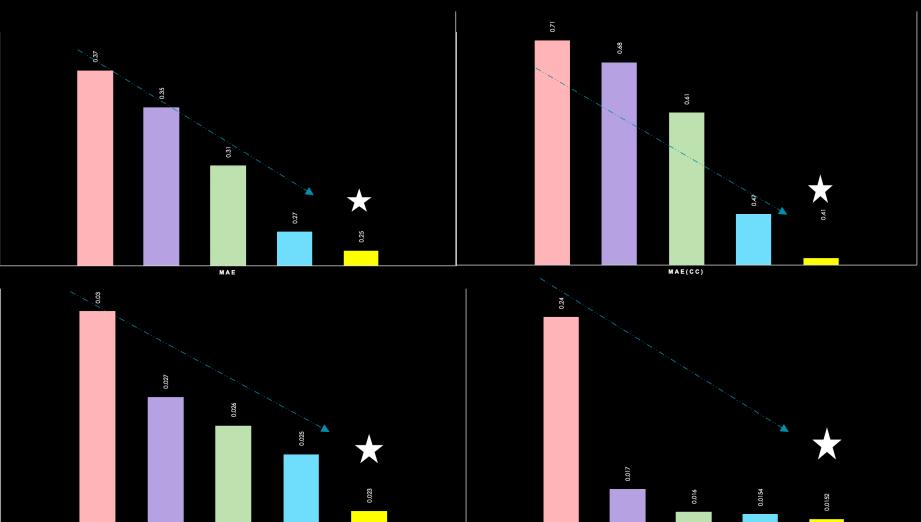
How to predict inter-modality non-isomorphic brain graphs from a base graph ?



Bar plots

Legend:

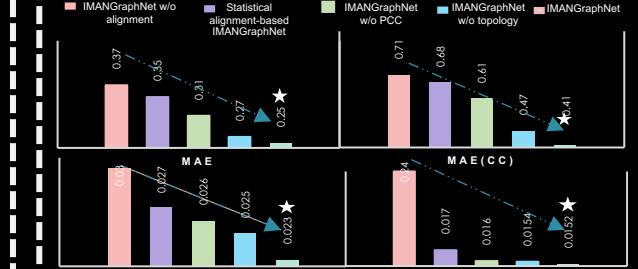
- IMANGraphNet w/o alignment
- Statistical alignment-based IMANGraphNet
- IMANGraphNet w/o PCC
- IMANGraphNet w/o topology
- IMANGraphNet



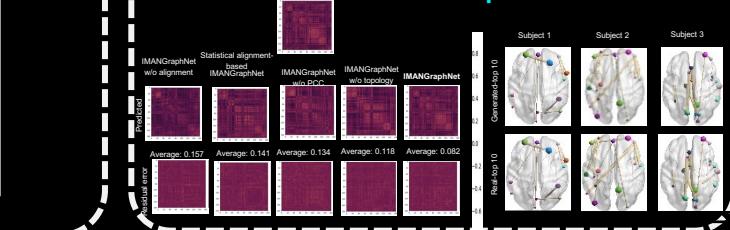
Results

Dataset evaluation
SLIM¹ dataset: 150 functional connectomes (160 x 160)
150 morphological connectomes (35 x 35)

Prediction results using different evaluation metrics



Clinical inspection



This work *is accepted* at IPMI 2021 and selected for an *oral presentation*.
GitHub code: <https://github.com/basiralab/IMANGraphNet>



Non-isomorphic Inter-modality Graph Alignment and Synthesis for Holistic Brain Mapping

Ialem Mhiri^{1,2} (PhD, Presenter), Ahmed Nebli^{1,2}, Mohamed Ali Mahjoub² and Ialem Rekik¹



¹BASIRA lab, Faculty of Computer and Informatics, Istanbul Technical University, Istanbul, Turkey

²University of Sousse, National Engineering School of Sousse (ENISo), Sousse, Tunisia



This work is accepted at "Information Processing in Medical Imaging" (Oral, IPMI 2021), Bornholm, Denmark.
<https://basira-lab.com>
GitHub code: [https://github.com/basiricalab/IMANGraphNet](https://github.com/basiralab/IMANGraphNet)



README.md

IMANGraphNet

IMANGraphNet (Non-isomorphic Inter-modality Graph Alignment and Synthesis for Holistic Brain Mapping), coded up in Python by Ialem Mhiri. Please contact islemmhiri1993@gmail.com for inquiries. Thanks.

This repository provides the official PyTorch implementation of the following paper:

A) Inter-modality domain aligner

B) Non-isomorphic inter-modality graph GAN

Non-isomorphic Inter-modality Graph Alignment and Synthesis for Holistic Brain Mapping [Ialem Mhiri]^{1,2}, [Ahmed Nebli]^{1,2}, [Mohamed Ali Mahjoub]², [Ialem Rekik]¹(<https://basira-lab.com>)

Releases

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Packages

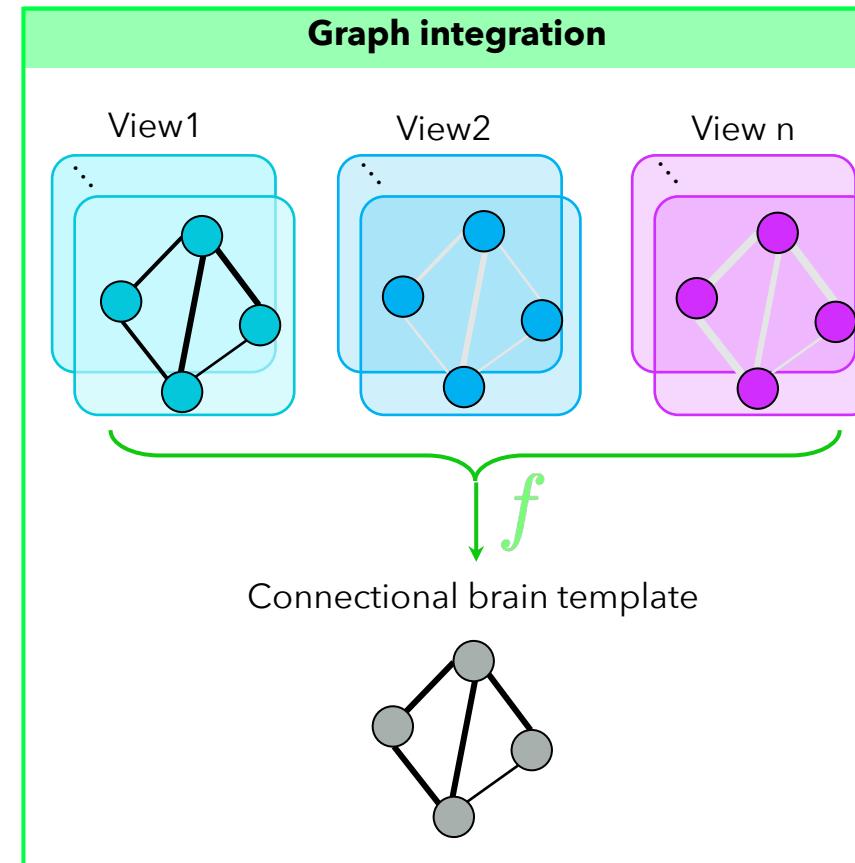
No packages published
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Contributors 3

- basiricalab BASIRA LAB
- YCT1 Yekta Can Tursun
- IalemRekik

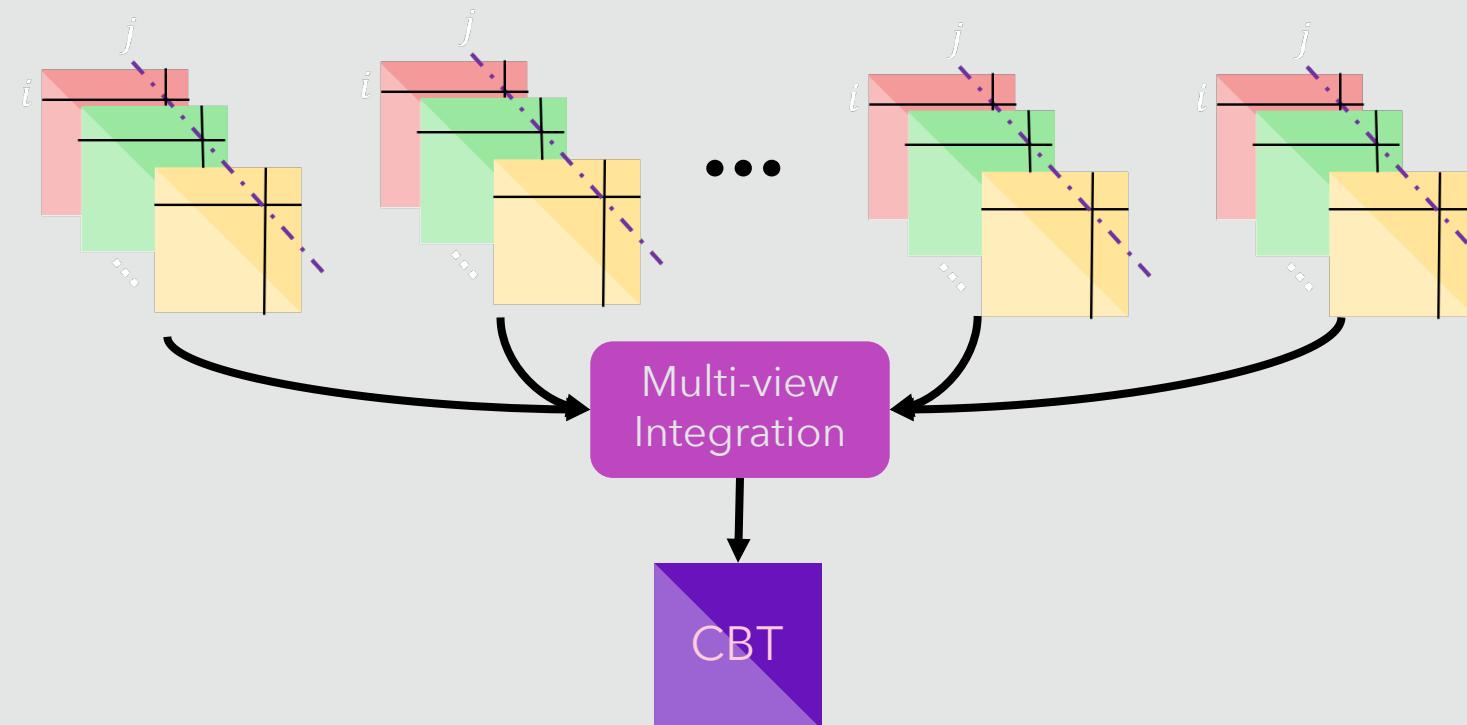
Languages

- Python 100.0%



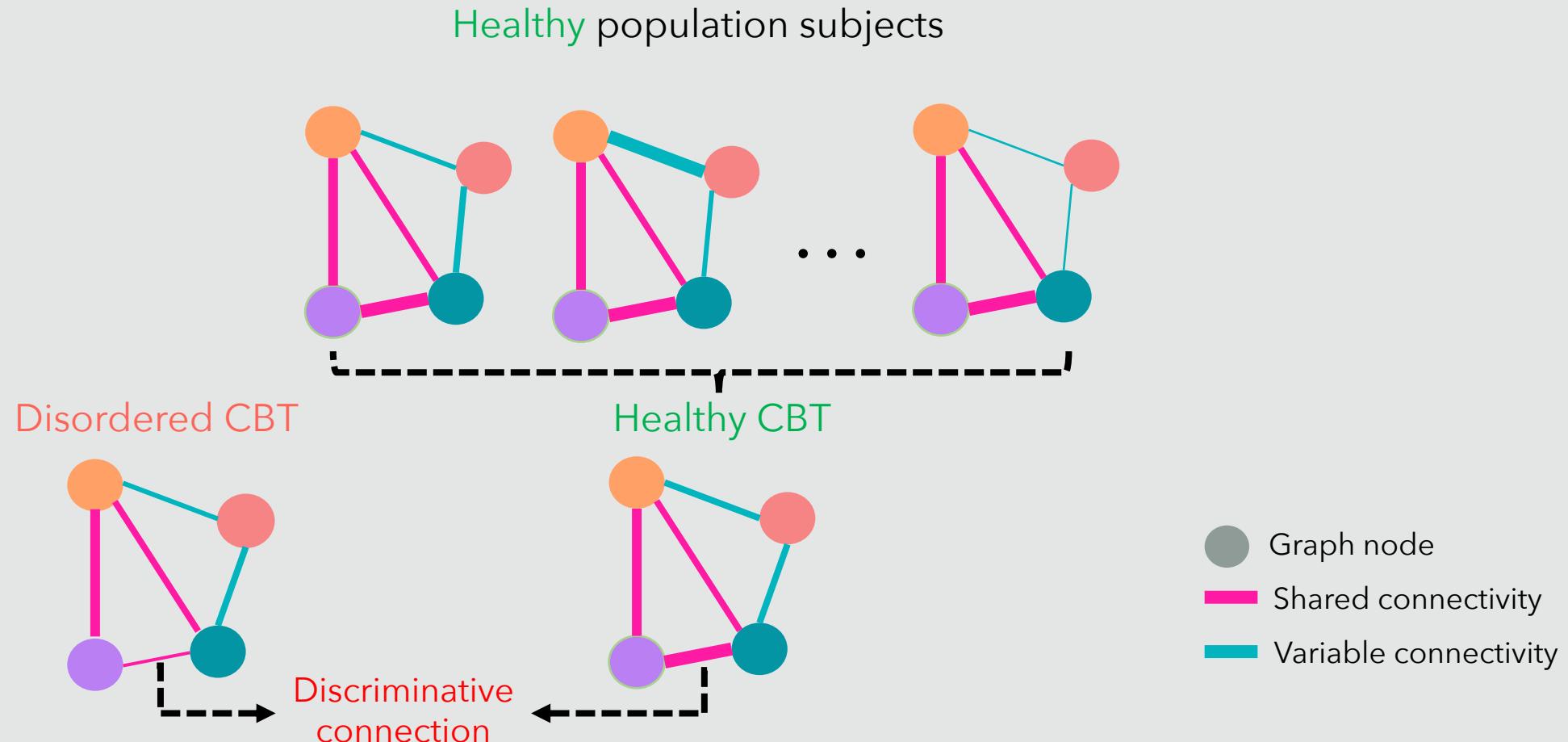
Problem statement

Integrating a population of multi-view brain networks to estimate a connectional brain template (CBT)



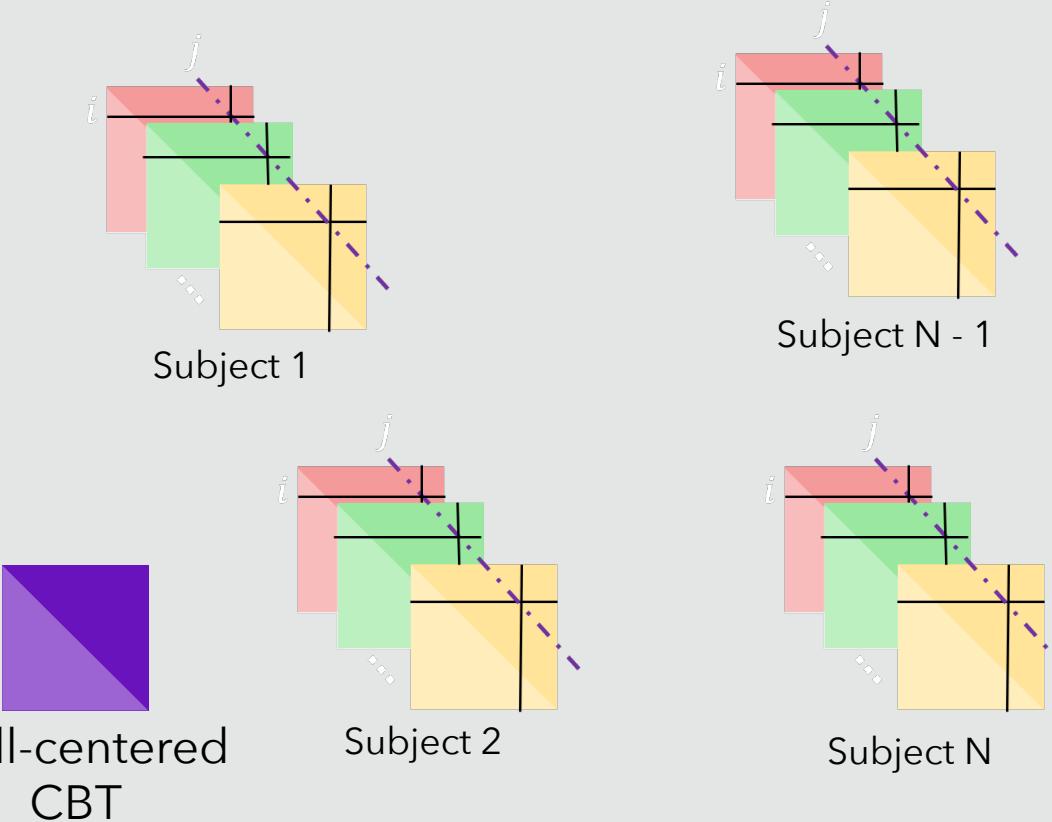
Why CBTs are powerful?

CBTs are powerful tools for group comparison studies and discovering the integral signature of an anomaly.



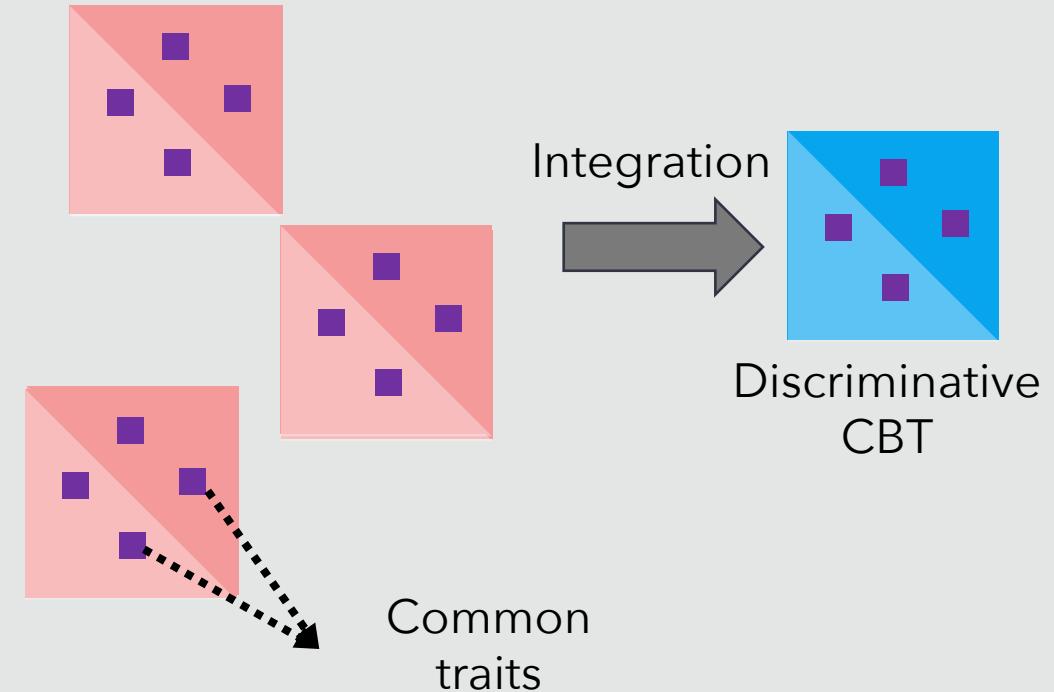
What is a good CBT?

Well-centered → Occupies the center of a population



Discriminative → Captures the unique and common traits of a population

Population samples



Deep Graph Normalizer: A Geometric Deep Learning Approach for Estimating Connectional Brain Templates



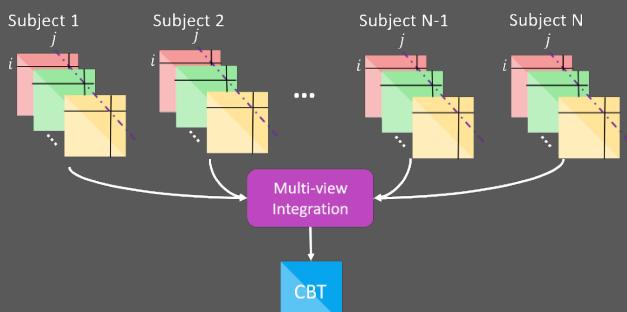
Mustafa Burak Gürbüz and



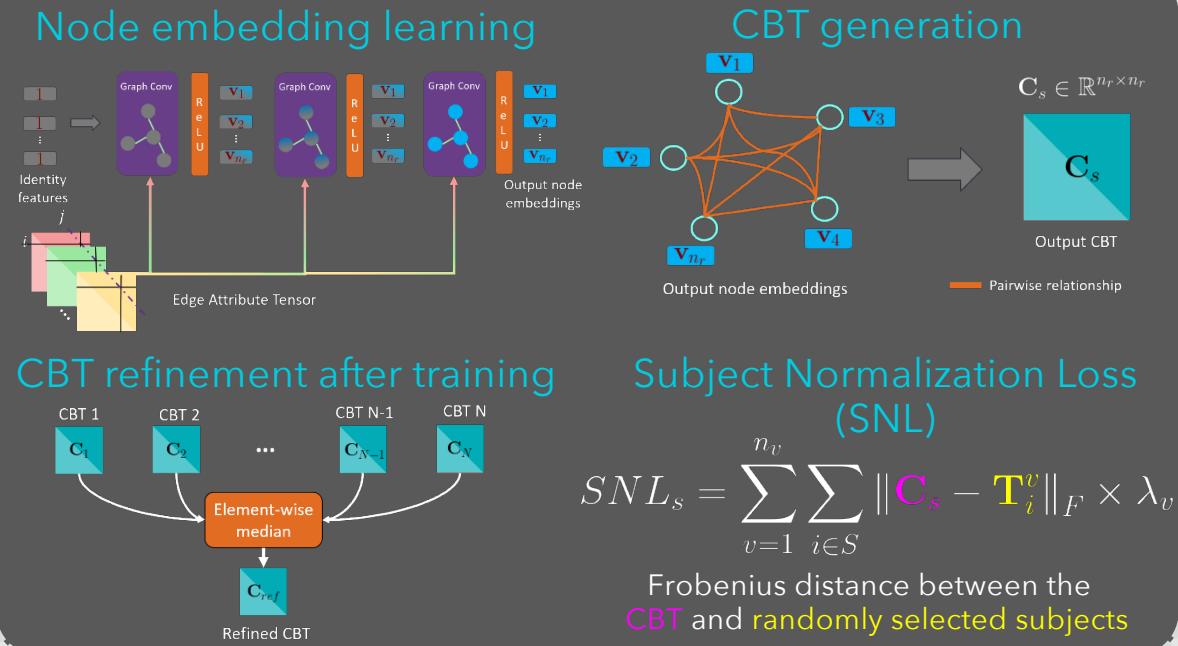
Islem Rekik

Problem statement

How can we **integrate a set of multi-view graph population to estimate a connectional brain template (CBT)?**



Proposed DGN solution



Results

Datasets

AD-LMCI * 77 subjects with 4 different views

NC-ASD ** 310 subject with 6 different views

Centeredness evaluation

Frobenius distance between views of testing subjects and estimated CBT

DGN outperforms netNorm¹ across 40 comparison

Discriminativeness evaluation

The overlap rate between features selected by an MKL*** feature selection and CBT comparison

DGN more reproducible in discriminability than netNorm¹ across 4 classification tasks



This work is accepted at the **MICCAI 2020** conference (**early accept**), Lima, Peru.

Gürbüz, M. B., & Rekik, I. (2021). MGN-Net: A multi-view graph normalizer for integrating heterogeneous biological network populations. *Medical Image Analysis*.

GitHub code: <https://github.com/basiralab/DGN>

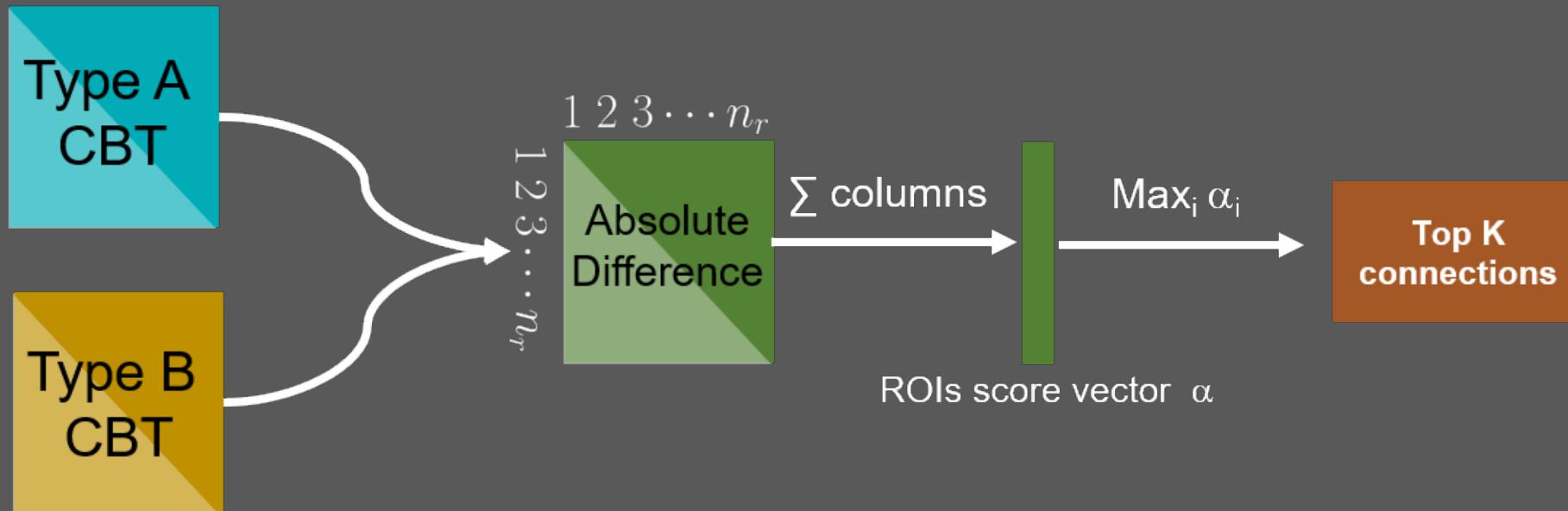
*Dhifallah, S., Rekik, I., "Estimation of connectional brain templates using selective multiview network normalization. Medical Image Analysis" 59 (2019)

AD-LMCI: Alzheimer's Diseases & Late Mild Cognitive Impairment *NC-ASD: Normal Control & Autism Spectrum Disorder ***

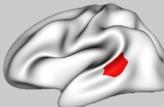
Comparison methods and results (Discriminativeness)

We spot the top 5 most discriminative brain regions where a type-A CBT largely differs from a type-B CBT.

Discriminative feature selection using CBTs



Discovery of most discriminative brain regions for ASD and NC

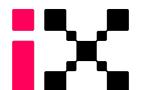
ASD/NC dataset	
Region of Interest	Effect on brain
Insula cortex (left hem.)	 Abnormalities in emotional and affective functions (Yamada et al., 2016)
Superior temporal sulcus (left hem.)	 Social perception impairment (Pelphrey et al., 2009)
Frontal pole (right hem.)	 Atypical right dominant face asymmetry (Hammond et al., 2008)



Yamada, T., et al. "Altered functional organization within the insular cortex in adult males with high- functioning autism spectrum disorder: evidence from connectivity-based parcellation." *Molecular Autism* 7 (2016)

Pelphrey, K., Carter, E., "Brain mechanisms for social perception lessons from autism and typical development" *Annals of the New York Academy of Sciences* 1145 (2009)

Hammond, P., et al., "Face-brain asymmetry in autism spectrum disorders." *Molecular psychiatry* 13 (2008)

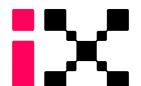


Discovery of most discriminative brain regions for AD and LMCI

AD/LMCI dataset	
Region of Interest	Effect on brain
Temporal pole (left hem.)	 Pathological changes in temporal pole (Arnold et al., 1994)
Entorhinal cortex (right hem.)	 Greater atrophy in the right entorhinal cortex in AD patients (Zhou et al., 2015)



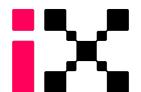
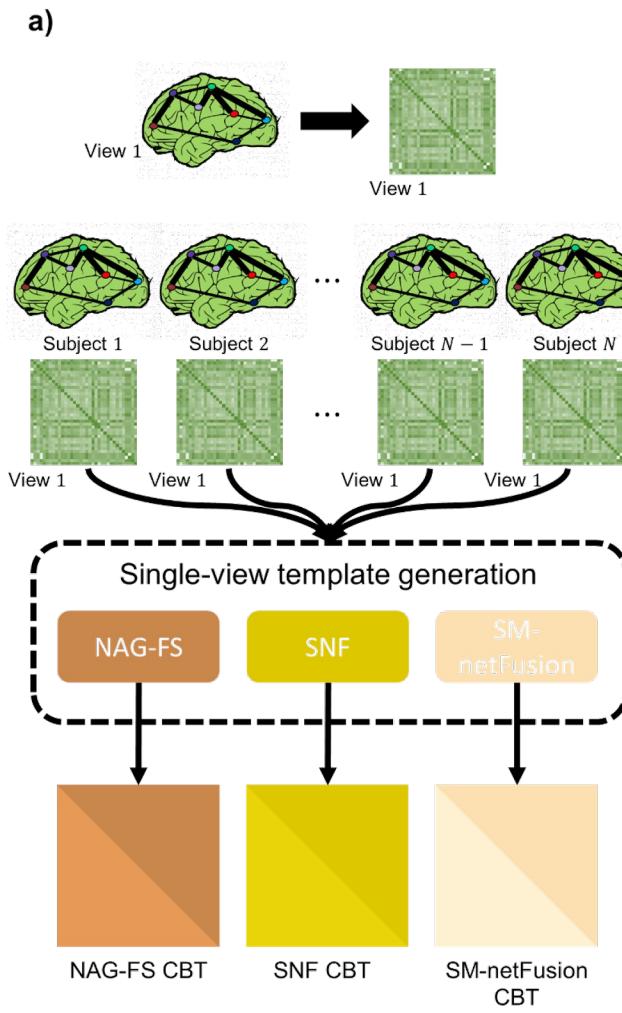
Arnold, S.E., Hyman, B.T., Van Hoesen, G.W., "Neuropathologic Changes of the Temporal Pole in Alzheimer's Disease and Pick's Disease." *Archives of Neurology* 51 (1994)
Zhou, M., et al., "Entorhinal cortex: A good biomarker of mild cognitive impairment and mild Alzheimer's disease." *Reviews in the neurosciences* 27 (2015)

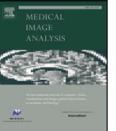




Comparative survey of multigraph integration methods for holistic brain connectivity mapping

Nada Chaari





[basiralab / survey-multigraph-integration-methods](https://github.com/basiralab/survey-multigraph-integration-methods) Public

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main

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<> Code

	basiralab Update README.md	...	on Jan 13	8
	Plots	v0		2 years ago
	Reproducibility of the p...	v0		2 years ago
	Reproducibility of the p...	v0		2 years ago
	data	v0		2 years ago
	Fig1.PNG	v1		2 years ago
	Fig2.PNG	v1		2 years ago
	Fig3.png	v1		2 years ago
	Fig4.PNG	v1		2 years ago
	LICENSE	v0		2 years ago
	README.md	Update README.md		3 months ago

About**Comparative survey of multigraph integration methods**

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Releases

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[Create a new release](#)**Packages**

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[Publish your first package](#)**Survey of multigraph integration methods for holistic brain mapping**Se Camgöz Akdağ^b, Islem Rekik^{a,c,*}^aand Informatics, Istanbul Technical University, Istanbul, Turkey^bIstanbul Technical University, Istanbul, Turkey^cand Innovation Hub, Imperial College London, London, UK**ABSTRACT**

One of the greatest scientific challenges in network neuroscience is to create a representative map of a population of heterogeneous brain networks, which acts as a connectional fingerprint. The connectional brain template (CBT), also named network atlas, presents a powerful tool for capturing the most representative and discriminative traits of a given population while preserving its topological patterns. The idea of a CBT is to integrate a population of heterogeneous brain connectivity networks, derived from different neuroimaging modalities or brain views (e.g., structural and functional), into a unified holistic representation. Here we review current state-of-the-art methods designed to estimate well-centered and representative CBT for populations of single-view and multi-view brain networks. We start by reviewing each CBT learning method, then we introduce the evaluation measures to compare CBT representativeness of populations generated by single-view and multigraph integration methods, separately, based on the following criteria: Centeredness, biomarker-reproducibility, node-level similarity, global-level similarity, and distance-based similarity. We demonstrate that the deep graph normalizer (DGN) method significantly outperforms other multi-graph and all single-view integration methods for estimating CBTs using a variety of healthy and disordered datasets in terms of centeredness, reproducibility (i.e., graph-derived biomarkers reproducibility that disentangle the typical from the atypical connectivity variability), and preserving the topological traits at both local and global graph-levels.

(rsfMRI) encode correlations in functional activity among brain regions, whereas diffusion tensor imaging (DTI) networks provide information concerning structural connections (i.e., white matter fiber paths) between these nodes (Tyan et al., 2017; Jiang et al., 2020; Dadashkarimi et al., 2019). Joining both networks results in two different views of brain connectivity, leading to more insights into the brain as a complex interconnected system.

Understanding how the brain's structural, morphological, and functional levels interlink offers a more comprehensive picture of the brain facets construction (Bassett and Sporns, 2017). For instance, (Farahani et al., 2019) provides an overview of the existing functional and effective connectivity methods used to construct the brain network. However, analyzing such *multi-modal* (also *multi-view*) connectomic dataset together remains challenging due to the large inter-modality variations across different views of connectivity networks and the heterogeneity of brain networks across population samples (Van Essen and Glasser, 2016; Verma et al., 2019). Nonetheless, mapping brain

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Received 30 December 2021; Received in revised form 27 December 2022; Accepted 3 January 2023

Available online 6 January 2023

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<https://github.com/basiralab/survey-multigraph-integration-methods>

Three trends in the AI-powered healthcare revolution

**Generative
learning**

Generate *multidimensional*
brains

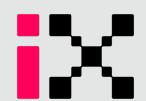
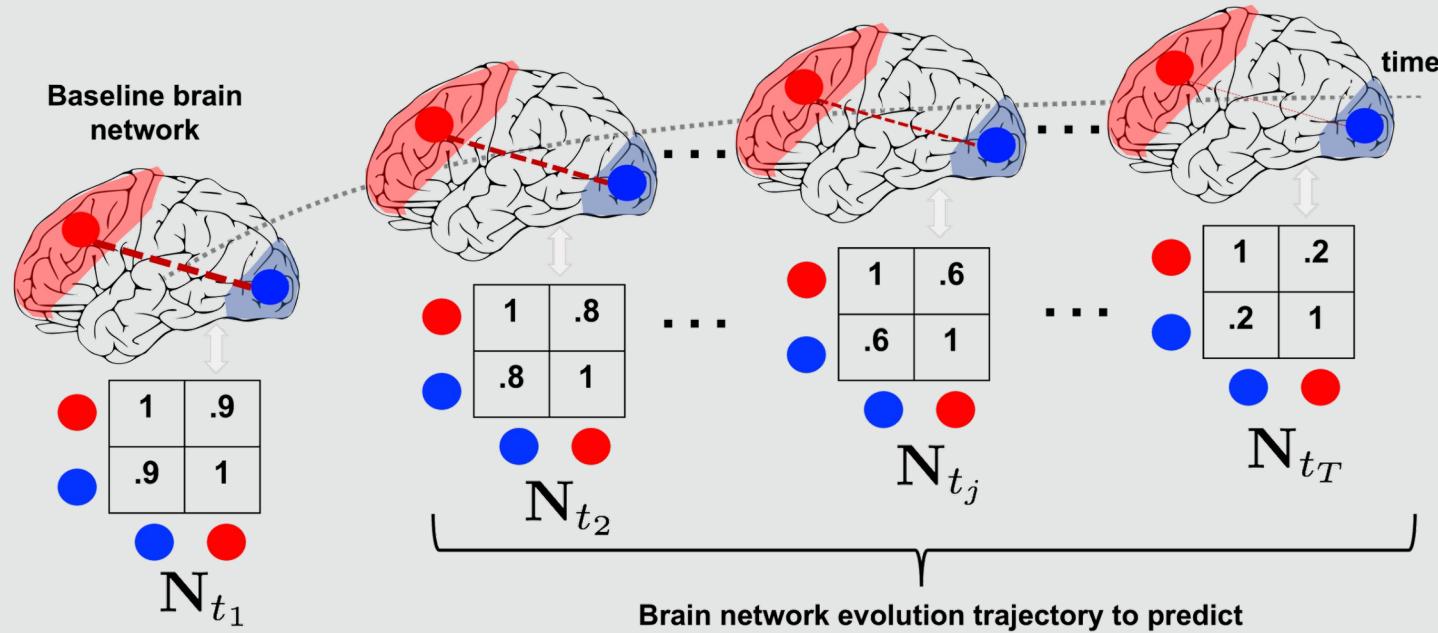
Generate connectional
brain templates

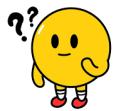
**Federated
learning**

Federate to generate the
future of the brain



Cross-time generation & federation



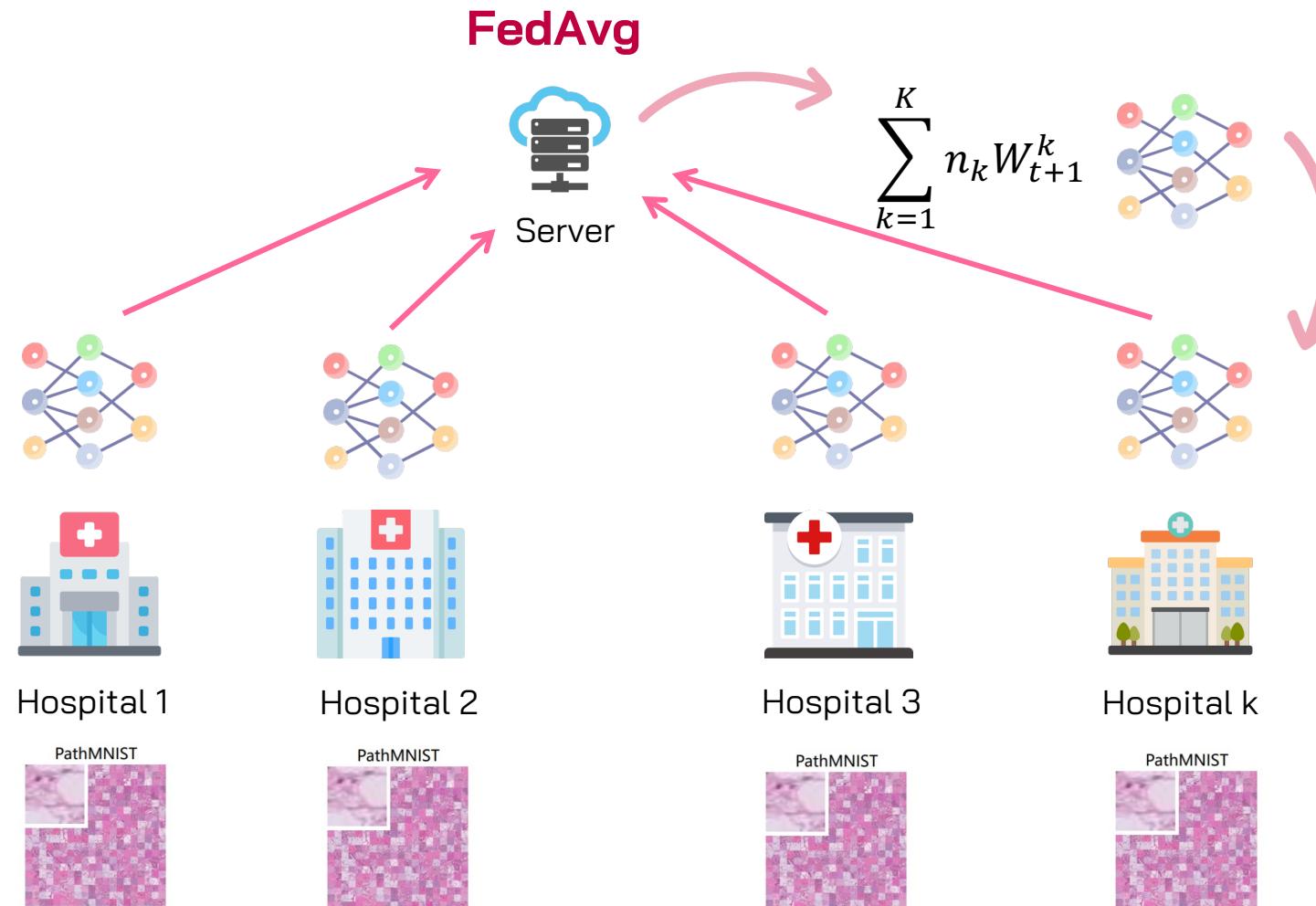


Can we learn better together *without sharing our data?*



Yes, with **federated learning!**

→ A decentralized framework for training global models **without sharing local datasets**, ensuring privacy and security.



New system protects patient data through federated learning

X Share f Share in Share

Publications & research news

25 January 2024

The Big Data Institute along with the collaborators have developed a new, easy-to-use technique for hospitals to contribute to developing artificial intelligence (AI) models, without patient data leaving the hospital's premises.



A scalable federated learning solution for secondary care using low-cost microcomputing: privacy-preserving development and evaluation of a COVID-19 screening test in UK hospitals

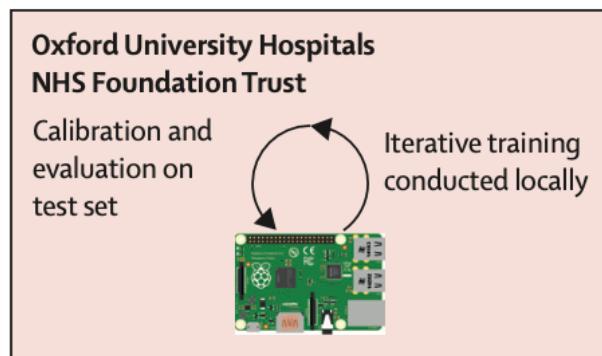
Andrew A S Soltan, Anshul Thakur, Jenny Yang, Anoop Chauhan, Leon G D'Cruz, Phillip Dickson, Marina A Soltan, David R Thickett, David W Eyre, Tingting Zhu, David A Clifton

Summary

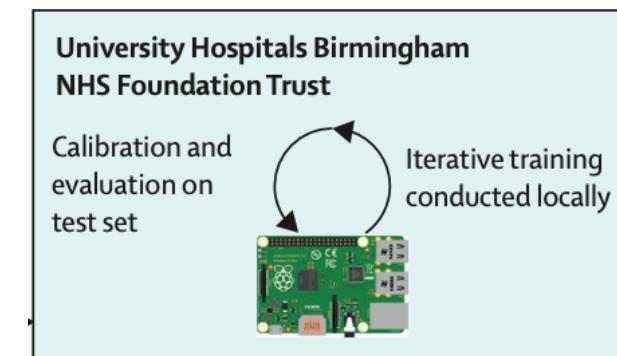
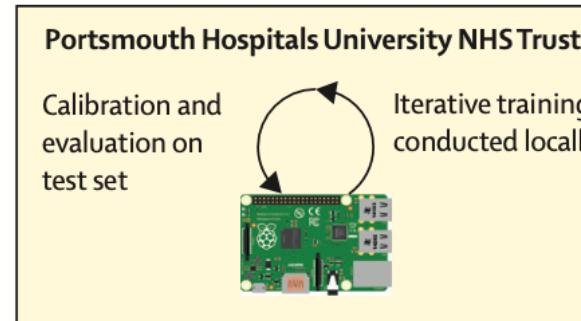
Background Multicentre training could reduce biases in medical artificial intelligence (AI); however, ethical, legal, and technical considerations can constrain the ability of hospitals to share data. Federated learning enables institutions to participate in algorithm development while retaining custody of their data but uptake in hospitals has been limited, possibly as deployment requires specialist software and technical expertise at each site. We previously developed an artificial intelligence-driven screening test for COVID-19 in emergency departments, known as CURIAL-Lab, which uses vital signs and blood tests that are routinely available within 1 h of a patient's arrival. Here we aimed to federate our COVID-19 screening test by developing an easy-to-use embedded system—which we introduce as full-stack federated learning—to train and evaluate machine learning models across four UK hospital groups without centralising patient data.



Federated model development



Updated model parameters, calibrated threshold, and performance metrics from test set evaluation transmitted to coordinating server



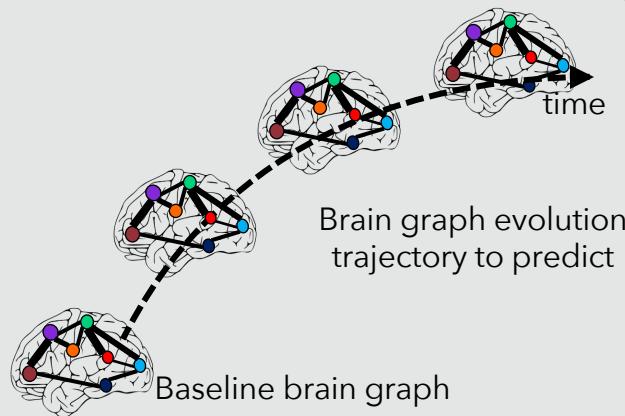
Updated model parameters, calibrated threshold, and performance metrics from test set evaluation transmitted to coordinating server

Federated Time-dependent GNN Learning from Brain Connectivity Data with Missing Timepoints

Zeynep Gurler



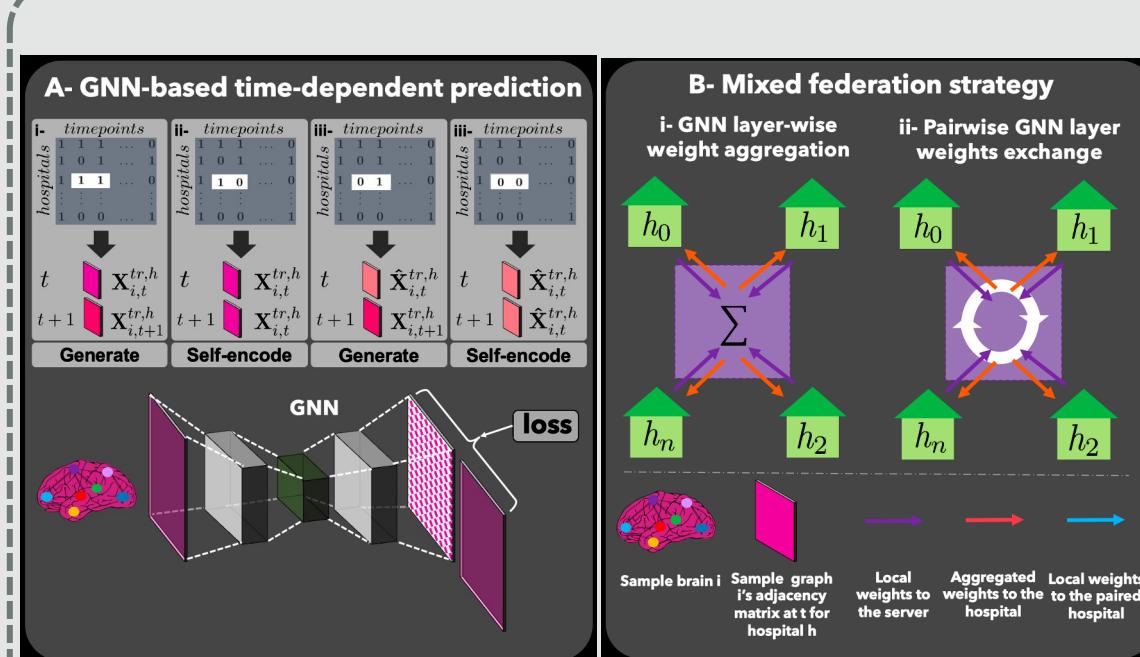
Problem statement



How to predict its evolution trajectory over time in a federated manner?

⌚ Limitation ➔ Absence of federated models for time-dependent graph evolution prediction

Proposed 4D-FED-GNN+



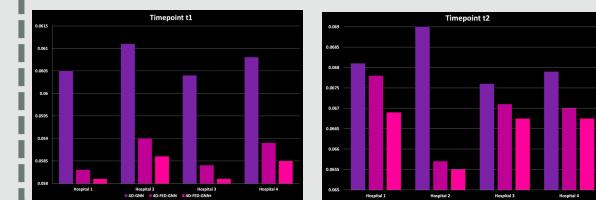
Results

Dataset

- We used 113 subjects from the OASIS-2 longitudinal dataset.
- Each subject was scanned on two or more visits, separated by at least one year.
- We used 5-fold CV for evaluation.
- In each run: 1 fold for testing and remaining 4 each allocated to a local hospital.

Comparison

- a) **4D-GNN**: the vanilla method where hospitals train their local data without federation.
- b) **4D-FED-GNN**: single federation strategy based on GNN layer-wise weight aggregation.



This work is part of my research project at BASIRA lab: <https://basira-lab.com>

This work is accepted at the "PRedictive Intelligence in MEdicine" (PRIME) MICCAI 2022 workshop and the journal of IEEE TMI 2022.

GitHub code: <https://github.com/basiralab/4D-FED-GNN>

¹Tekin, A., et al. "Recurrent brain graph mapper for predicting time-dependent brain graph evolution trajectory." Domain Adaptation and Representation Transfer, and Affordable Healthcare and AI for Resource Diverse Global Health (2021)

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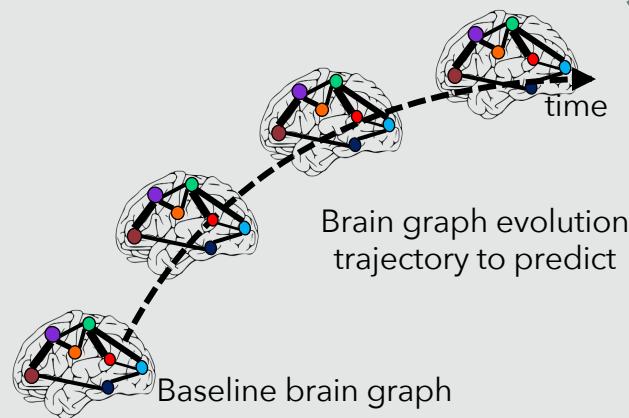


Federated Time-dependent GNN Learning from Brain Connectivity Data with Missing Timepoints

Zeynep Gurler



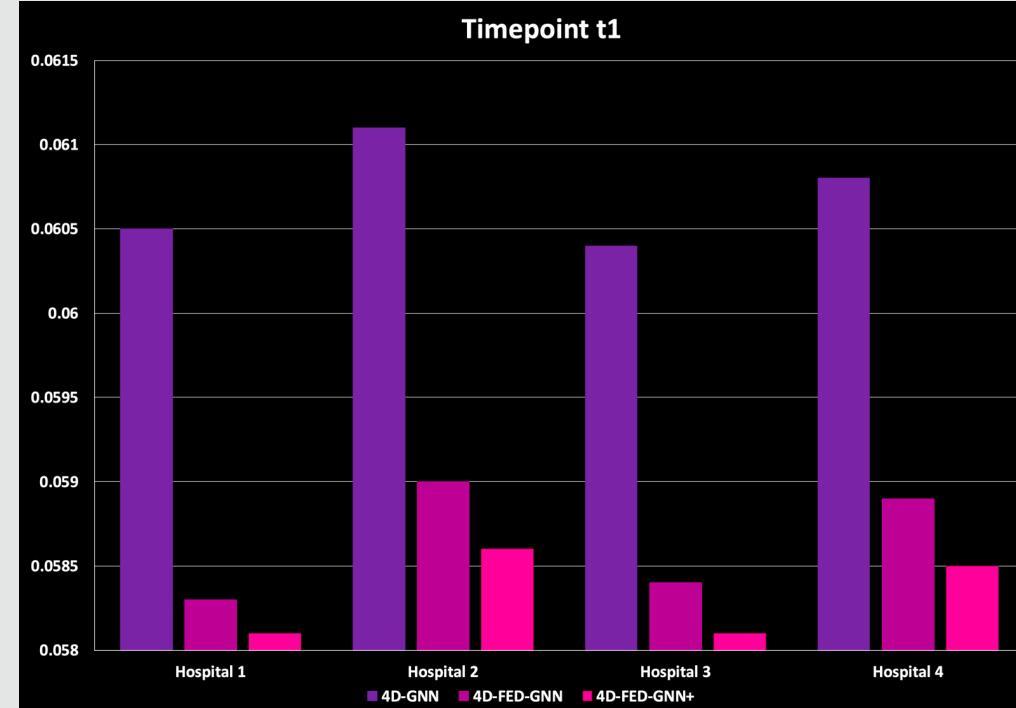
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Bar plots



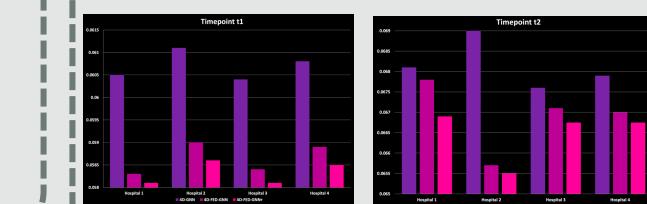
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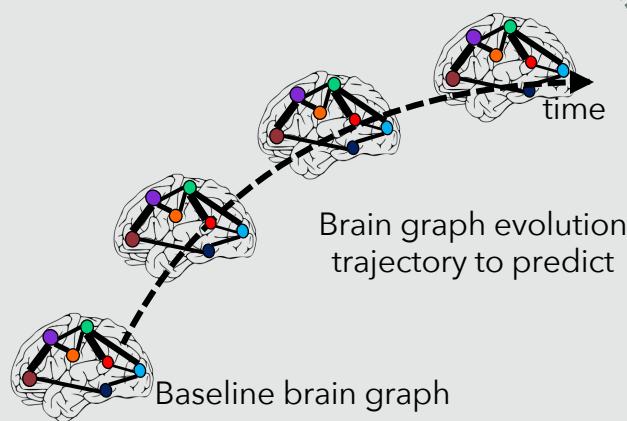


Federated Time-dependent GNN Learning from Brain Connectivity Data with Missing Timepoints

Zeynep Gurler



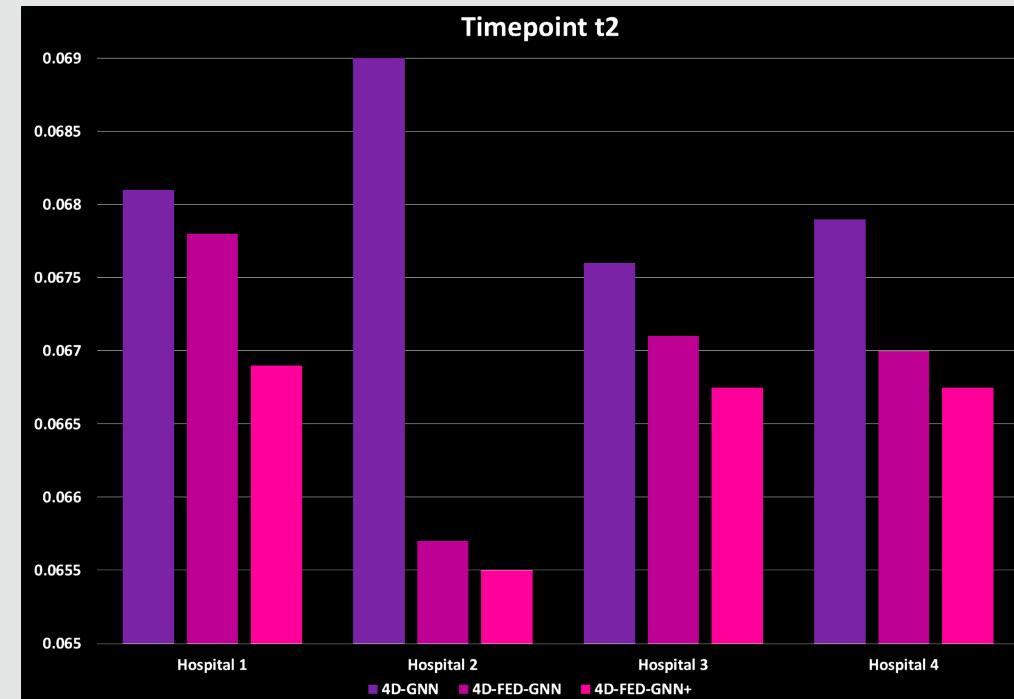
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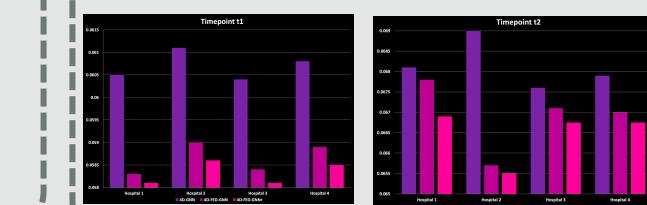
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Federated Time-dependent GNN Learning from Brain Connectivity Data with Missing Timepoints

Zeynep Gurler and Islem Rekik



BASIRA lab, Faculty of Computer and Informatics, Istanbul Technical University, Istanbul, Turkey



This work is accepted as poster presentation at the "PRedictive Intelligence in MEdicine" (PRIME) MICCAI 2022 workshop, Singapore.

This work is part of my research project at BASIRA lab:

<https://basira-lab.com/prime-miccai-2022/>

GitHub code: <https://github.com/basiralab/4D-FED-GNN>



4D-FED-GNN code

Federated Time-dependent GNN
Learning from Brain Connectivity Data
with Missing Timepoints

This work is accepted as [poster presentation](#) at
the "PRedictive Intelligence in MEdicine" (PRIME)
MICCAI 2022 workshop, Singapore.



<http://basira-lab.com/prime-miccai22/>
GitHub code: <https://github.com/basiralab/4D-FED-GNN>



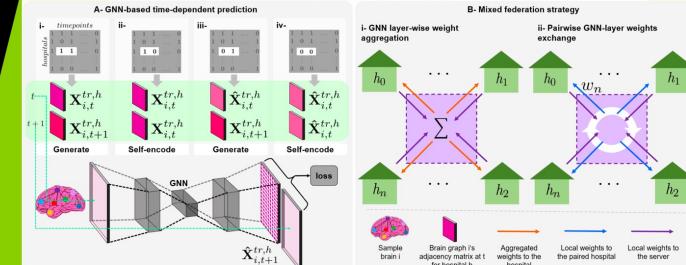
Federated Time-dependent GNN Learning from
Brain Connectivity Data with Missing
Timepoints

Zeynep Gurler and Islem Rekik *

BASIRA Lab, Faculty of Computer and Informatics, Istanbul Technical University,
Istanbul, Turkey (<http://basira-lab.com/>)

Abstract. Predicting changes in brain connectivity between anatomical regions is essential for brain mapping and neural disorder diagnosis across different age spans from a limited data (e.g., single timepoint).

Such learning tasks become more difficult when handling a single dataset with missing timepoints, let alone multiple decentralized datasets collected from different hospitals with varying incomplete acquisitions. In such a non-distributed federated learning (FL) one can learn from decentralized datasets without data sharing. However, to the best of our knowledge, no FL method was designed to predict time-dependent graph data evolution trajectory using non-iid training longitudinal datasets with varying acquisition timepoints. In this paper, we aim to signifi-



Three trends in the AI-powered healthcare revolution

**Generative
learning**

Generate *multidimensional*
brains

**Federated
learning**

Federate to generate the
future of the brain

Generate connectional
brain templates

Federate heterogeneous
diagnostic tasks

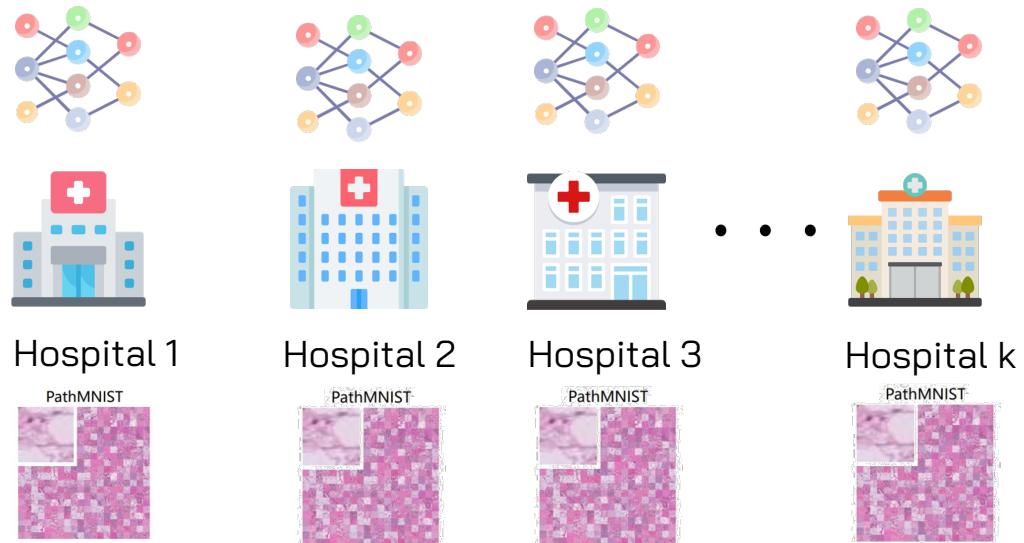




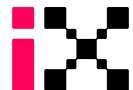
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UniFed: A Universal Federation of a Mixture of Highly Heterogeneous Medical Image Classification Tasks

same dataset types = same learning task (i.e., disease)



This work is accepted at the *MICCAI MLMI 2024* conference (poster presentation), Marrakesh, Morocco.
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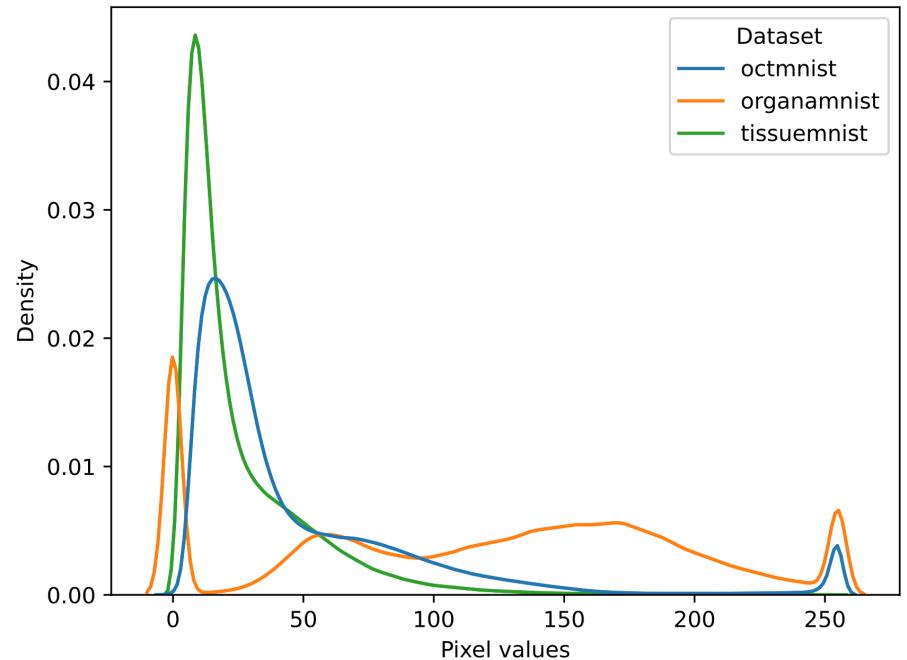
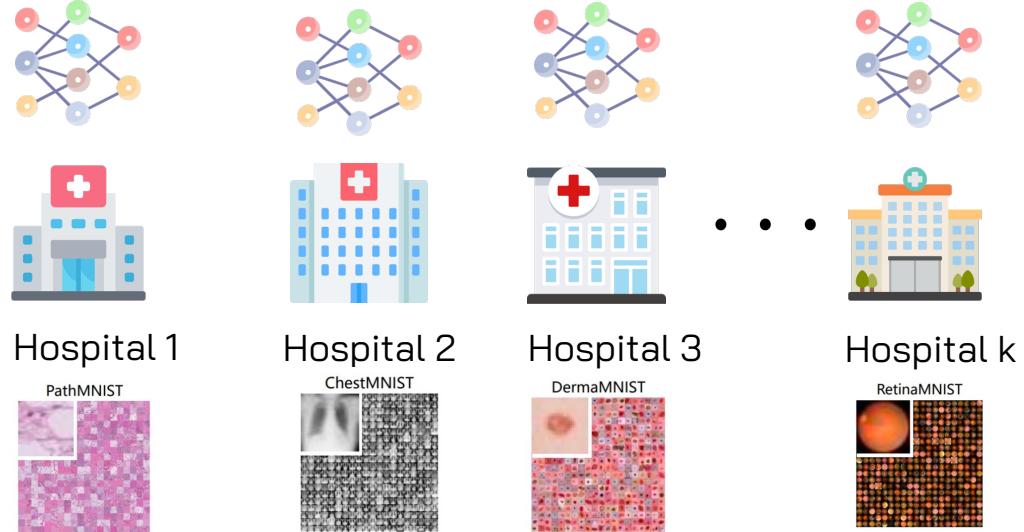




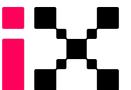
UniFed: A Universal Federation of a Mixture of Highly Heterogeneous Medical Image Classification Tasks

Atefe Hassani

different dataset types = different learning tasks (i.e., diseases)

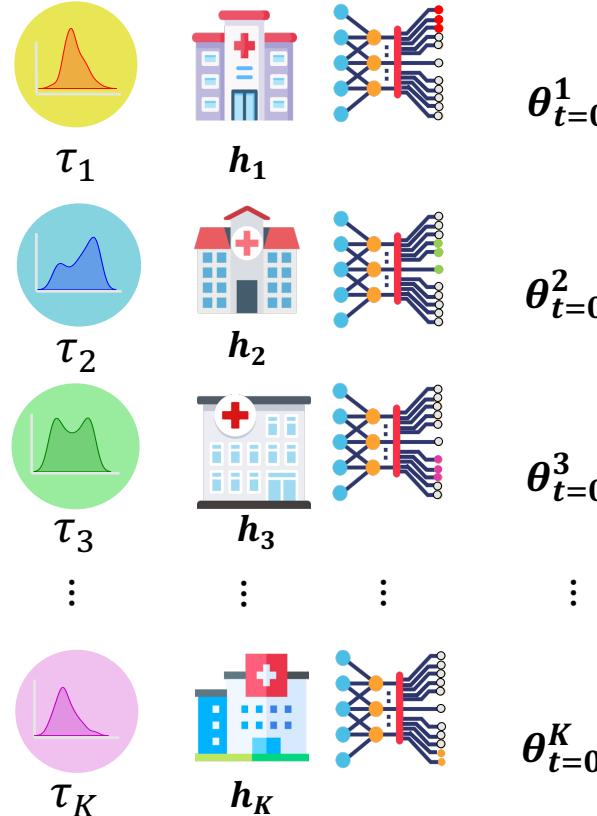


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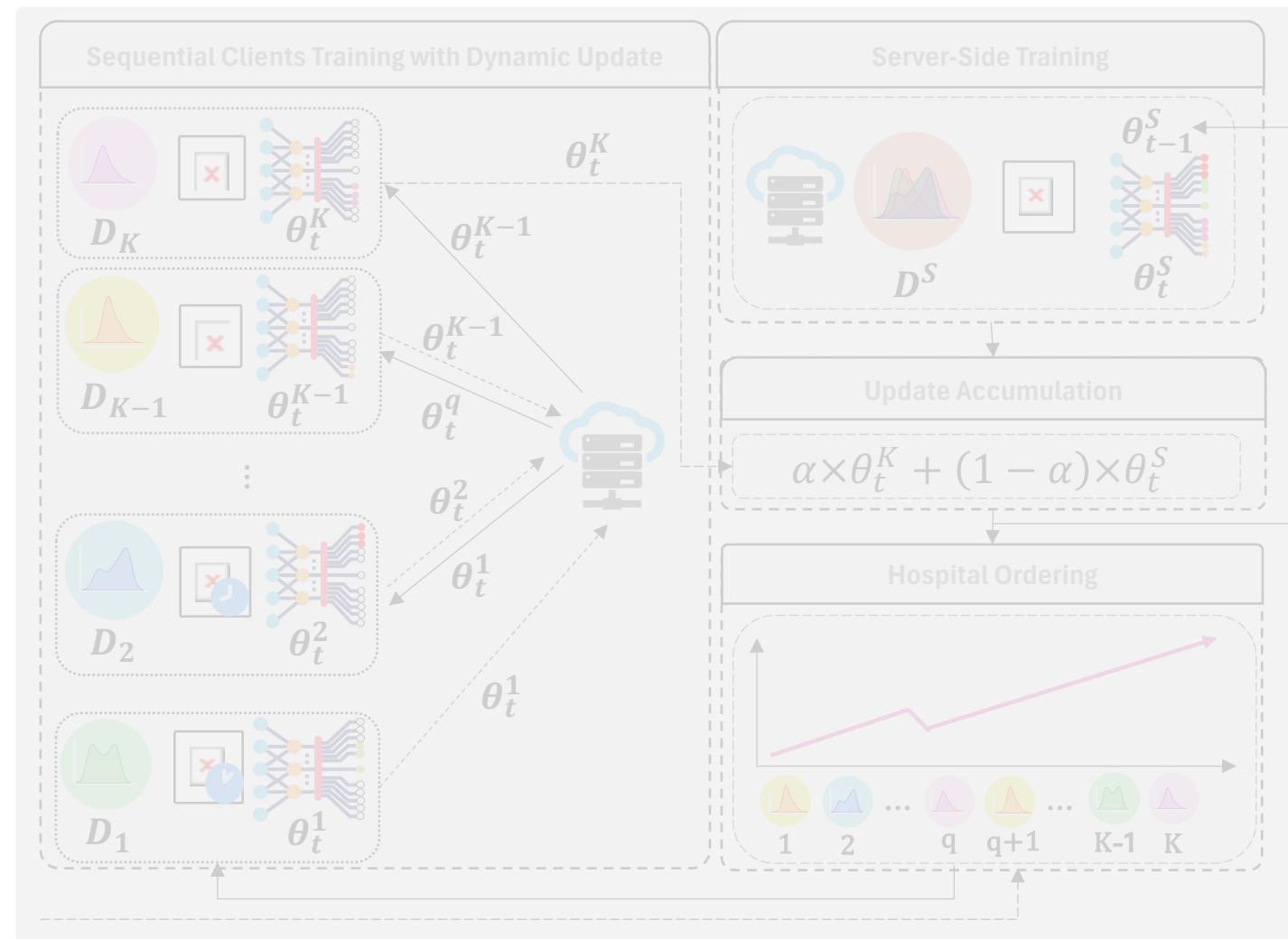
Federated learning

Federate heterogeneous diagnostic tasks



Proposed UniFed learning architecture

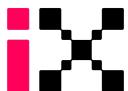
Initial client training



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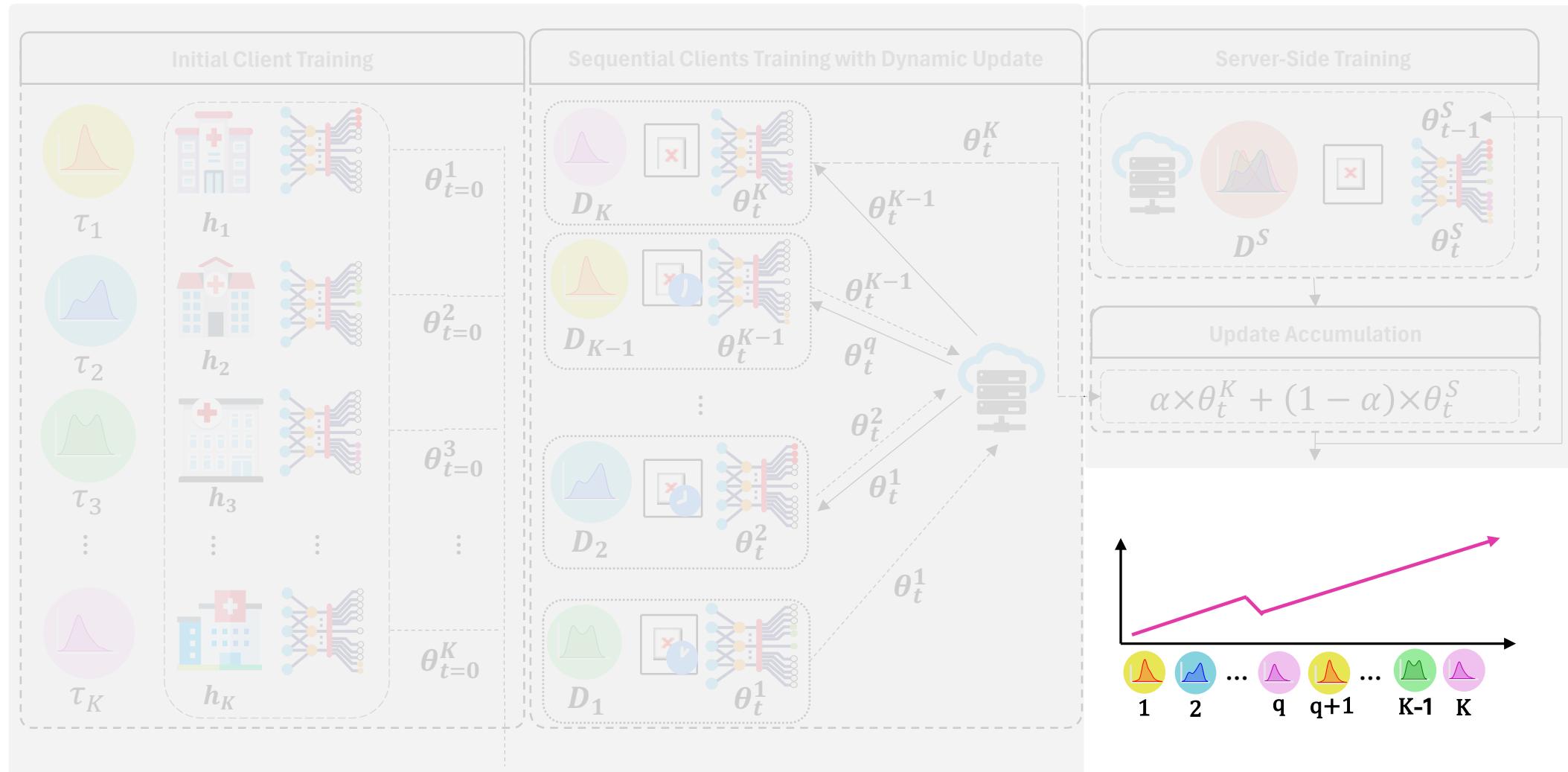


Federated learning

Federate heterogeneous diagnostic tasks

Proposed UniFed learning architecture

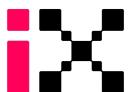
Hospital (client) ordering



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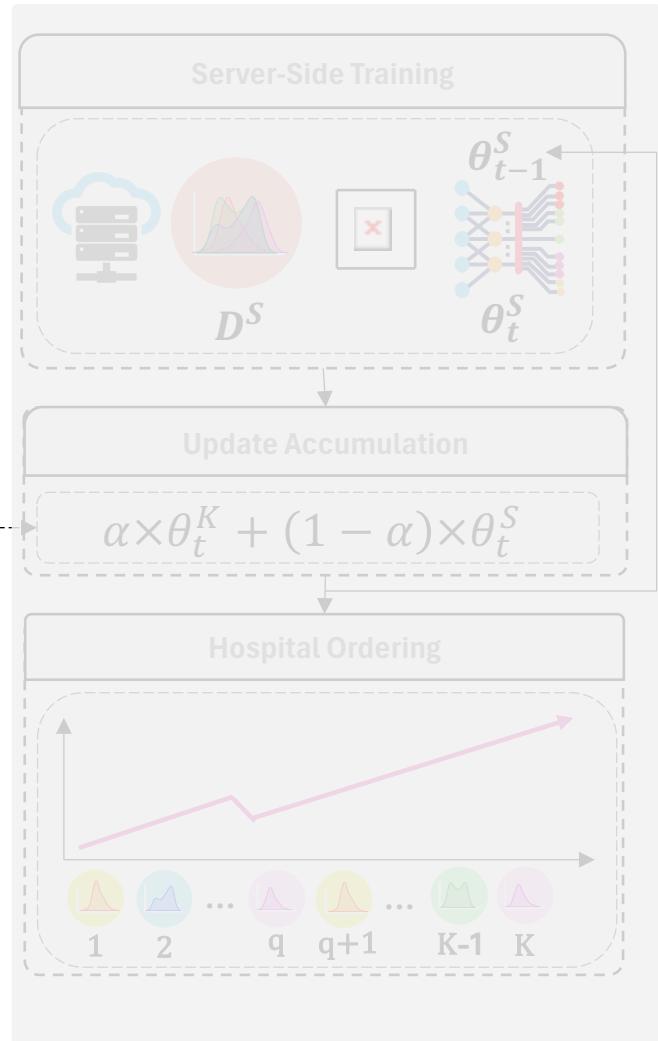
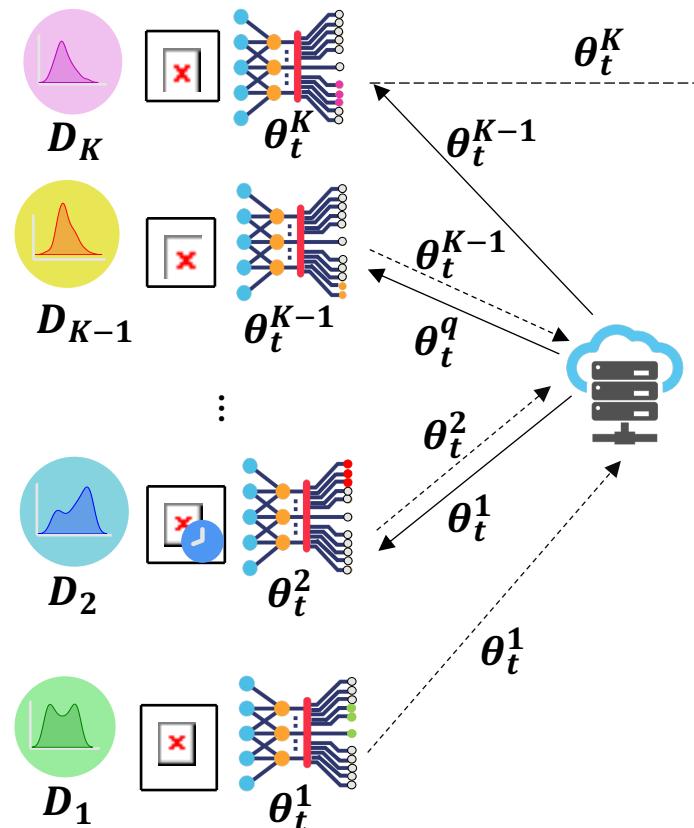
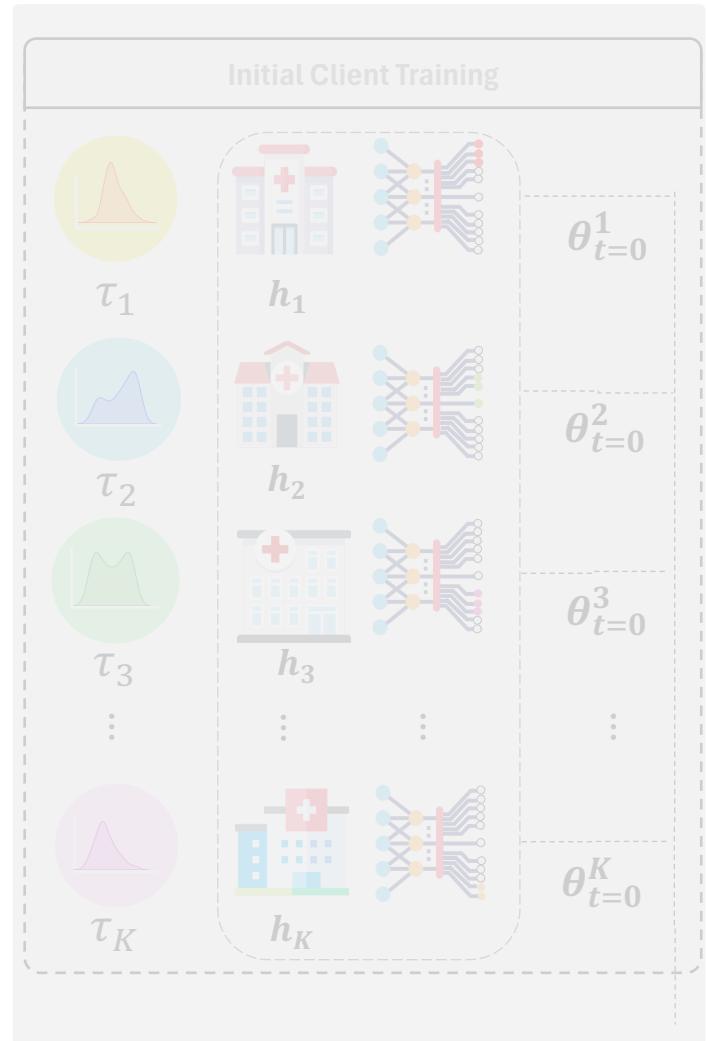


Federated learning

Federate heterogeneous diagnostic tasks

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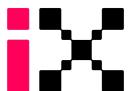
Sequential client training + dynamic update



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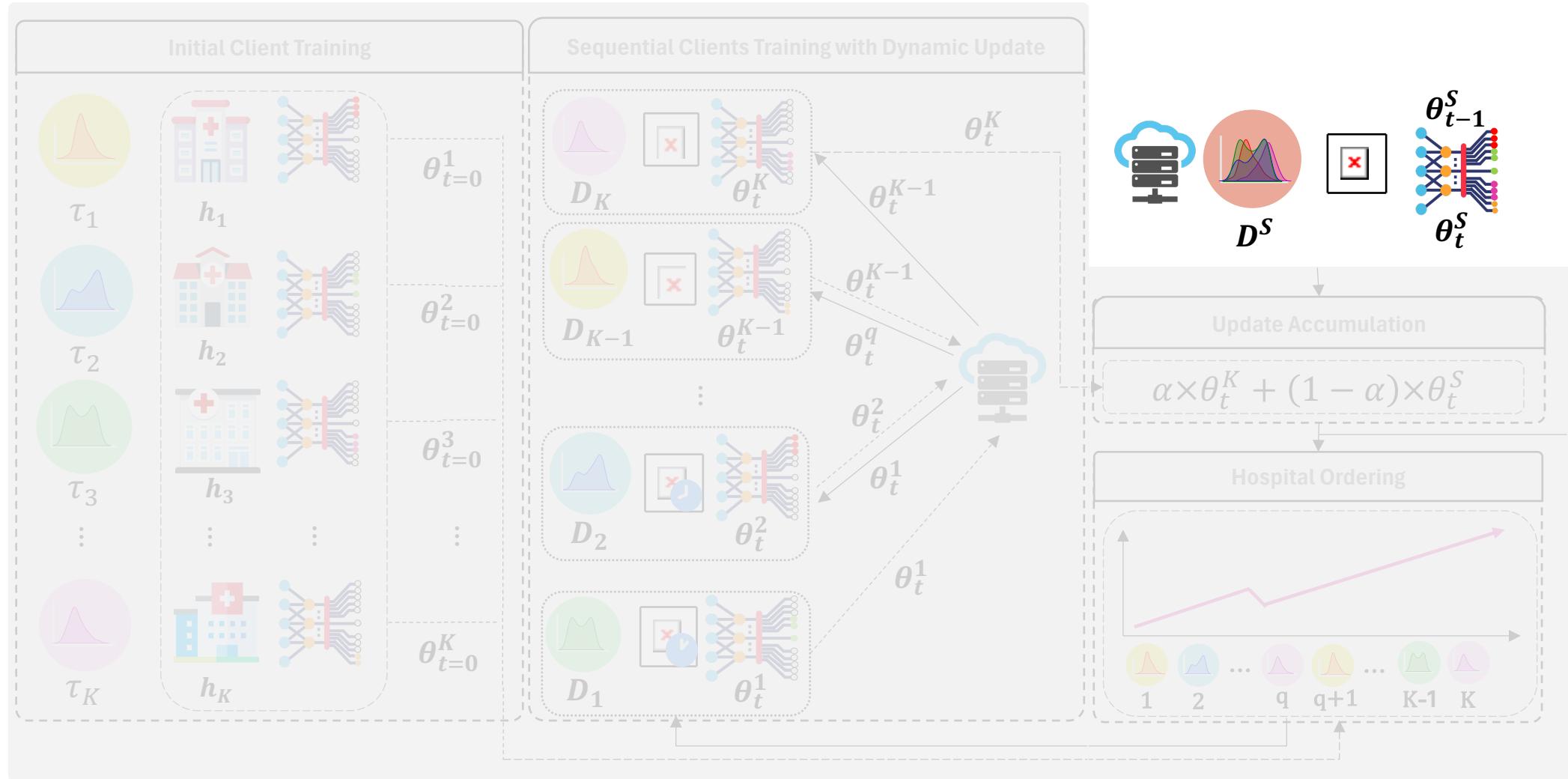


Federated learning

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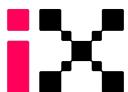
Server-side training



Atefe Hassani



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Federated learning

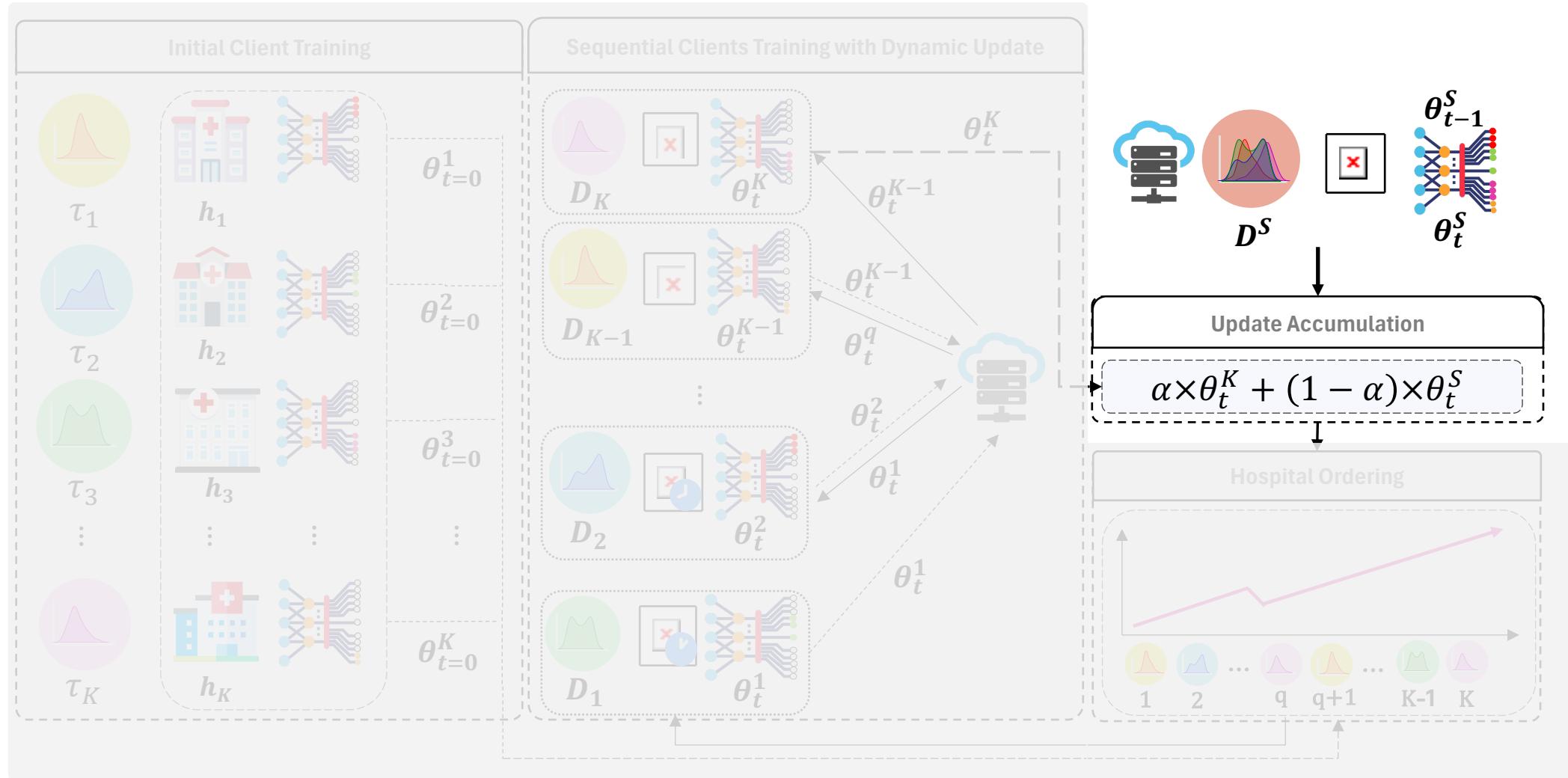
Federate heterogeneous diagnostic tasks

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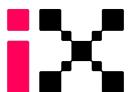
Server-side learning and global model update



Atefe Hassani



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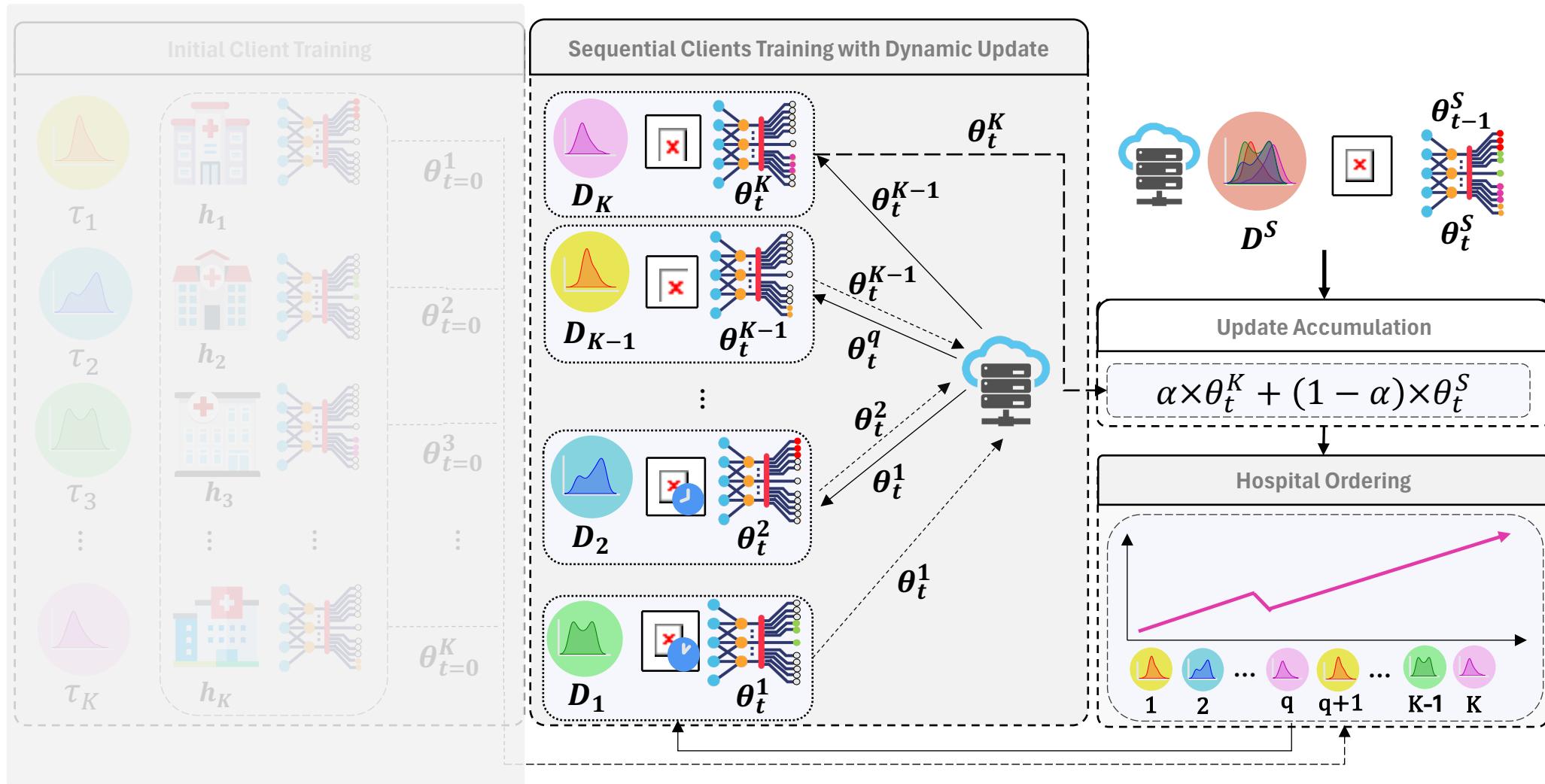


Federated learning

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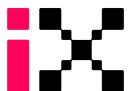
Universal federated learning



Atefe Hassani



This work is accepted at the *MICCAI MLMI 2024* conference (poster presentation), Marrakesh, Morocco.
GitHub code: <https://github.com/basiralab/UniFed>





Atefe Hassani

Table 1. Comparison of model performance (%) in strongly and moderately Non-IID settings.

Data Partition	Method	CNN				VGG11				ResNet18			
		Acc	F1	Sens	Spec	Acc	F1	Sens	Spec	Acc	F1	Sens	Spec
NoFed	NoFed	68.17	54.15	56.42	96.63	77.10	65.90	67.31	97.00	74.94	61.26	63.39	96.94
Strongly Non-IID	FedAvg [1]	38.44	24.88	23.89	96.85	4.90	2.33	1.69	96.18	9.79	6.82	9.13	96.10
	FedProx [32]	37.92	23.37	22.32	96.86	4.06	2.12	1.78	96.17	9.68	5.45	7.28	96.08
	FedSeq [33]	10.73	1.05	0.59	95.73	26.46	17.24	23.42	96.20	22.81	18.60	23.31	96.09
Moderately Non-IID	UniFed	69.37	55.05	55.87	96.75	50.52	37.41	39.18	96.56	46.77	32.45	34.32	96.45
	FedAvg [1]	24.58	22.73	24.24	93.00	6.04	1.83	1.83	94.1	10.10	5.61	6.59	92.58
	FedProx [32]	24.58	22.72	24.22	92.99	5.10	1.31	1.05	93.80	9.37	5.34	5.55	93.46
	FedSeq [33]	3.75	0.91	0.67	94.67	13.75	7.35	8.64	92.74	19.38	19.04	21.46	93.57
UniFed		58.02	55.97	58.36	94.10	46.15	40.26	43.36	93.11	32.40	28.72	31.75	95.3

UniFed outperforms existing federation methods



Atefe Hassani

The computation overhead on MedMNIST dataset with **strongly Non-IID setting** and the communication overhead (local epoch for each hospital × number of hospitals × iteration). The computation overhead shows the total training time in minutes.

Method	Computation			Communication		
	CNN	VGG11	ResNet18	CNN	VGG11	ResNet18
NoFed	8.43	23.56	30.58	24000	24000	24000
FedAvg	207.65	251.56	162.02	24,000	24,000	24,000
FedProx	71.26	276.72	244.78	24,000	24,000	24,000
FedSeq	170.58	227.2	169.03	48,000	24,000	24,000
UniFed	98.43	50.41	39.78	30,327	7,765	6,213

UniFed is more affordable to train

UniFed: A Universal Federation of a Mixture of Highly Heterogeneous Medical Image Classification Tasks

Atefe Hassani and Islam Rekik



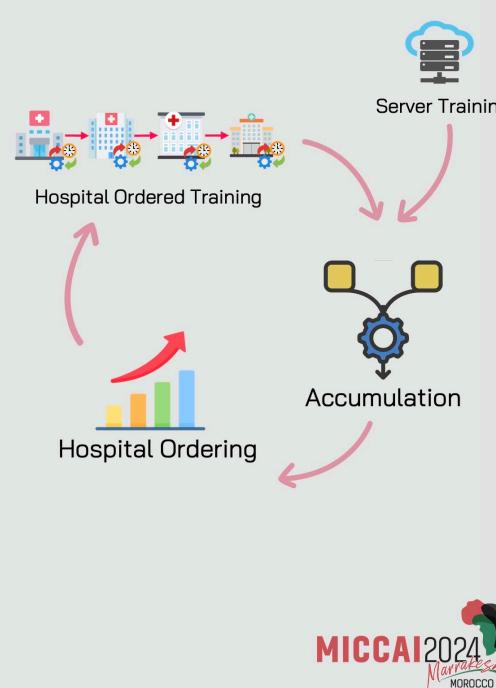
BASIRA Lab, Imperial-X and Department of Computing; Imperial College London, UK

This work is part of my research project at BASIRA lab:

<https://basira-lab.com>

GitHub code: <https://github.com/basiralab/UniFed>

Imperial College London



<https://github.com/basiralab/UniFed>

The GitHub repository page for UniFed is shown. The repository is public and has 1 branch and 0 tags. The last commit was made by HassaniAtefe 3 months ago. The repository contains files such as README.md, figs, models, utils, .gitignore, LICENSE, baseline.ipynb, configs.yaml, main.ipynb, and requirements.txt. The repository has 2 stars, 1 watching, and 0 forks. It includes sections for About, Releases, Packages, Contributors, and Languages.

About
No description, website, or topics provided.

Releases
No releases published
[Create a new release](#)

Packages
No packages published
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Contributors 2

HassaniAtefe Atefe Hassani
basiralab BASIRA LAB

Languages

Jupyter Notebook	63.0%
Python	37.0%

UniFed: A Universal Federation of a Mixture of Highly Heterogeneous Medical Image Classification Tasks

This repository provides the official implementation of our UniFed model accepted to [MLMI-MICCAI-2024](#).

What is UniFed?



Three trends in the AI-powered healthcare revolution

Generative learning

Generate *multidimensional* brains

Generate connectional brain templates

Federated learning

Federate to generate the future of the brain

Federate heterogeneous diagnostic tasks

Holistic learning

Insights into the future of an inclusive and holistic AI in medicine



Avicenna
Ibn Sina (980–1037),

A Persian **polymath** whose work in **medicine, Islamic philosophy, and science** laid foundational principles that influenced medical practice for centuries.

His visionary insights are particularly noted in *The Canon of Medicine*, which became a **standard text in medical schools across Europe and the world** for hundreds of years.

[Mental Health]

"The imagination is the intermediary between the bodily organs and the soul."

[Holistic Healing]

"The physician should not treat the disease but the patient who is suffering from it."



A History of Population Health: the rise & fall of diseases

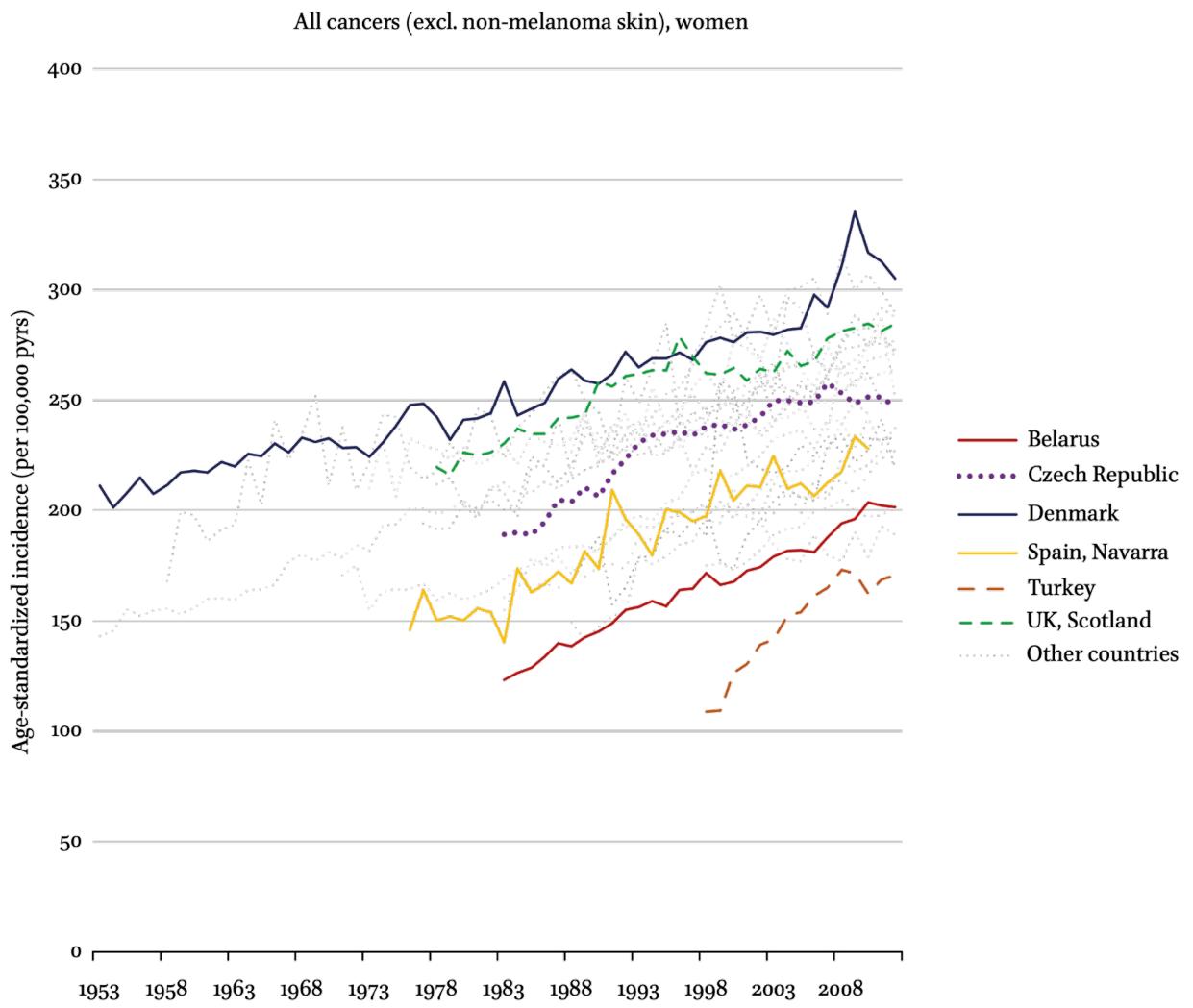


FIGURE 4 Trends in cancer incidence in Europe, 1953–2011

SOURCE OF DATA: NATIONAL OR REGIONAL CANCER REGISTRIES (THROUGH GLOBAL CANCER OBSERVATORY (GCO.IARC.FR; ACCESSED 24/04/2019))



A History of Population Health

Rise and Fall of Disease in Europe

Johan P. Mackenbach



That 'rise-and-fall' is the usual time-pattern in which diseases afflict human populations must have a deeper explanation. Why have, in the course of human history, so many diseases emerged? Why have, on the other hand and with shorter or longer delay, most of these diseases become less common again, or at least become less lethal? And is there a causal link between the disappearance of one disease and the emergence of another?

We need a more **holistic**
approach.

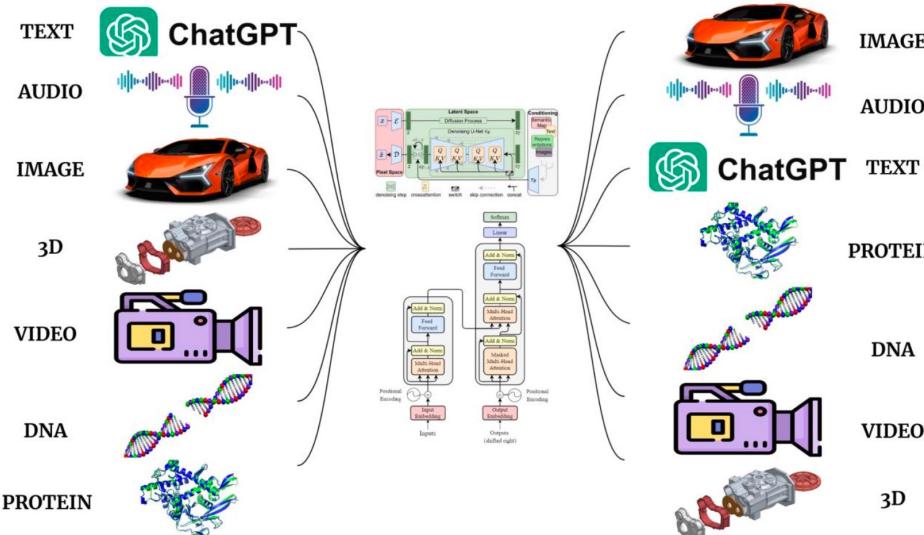
Are we asking the *right*
questions and looking at the
right data?



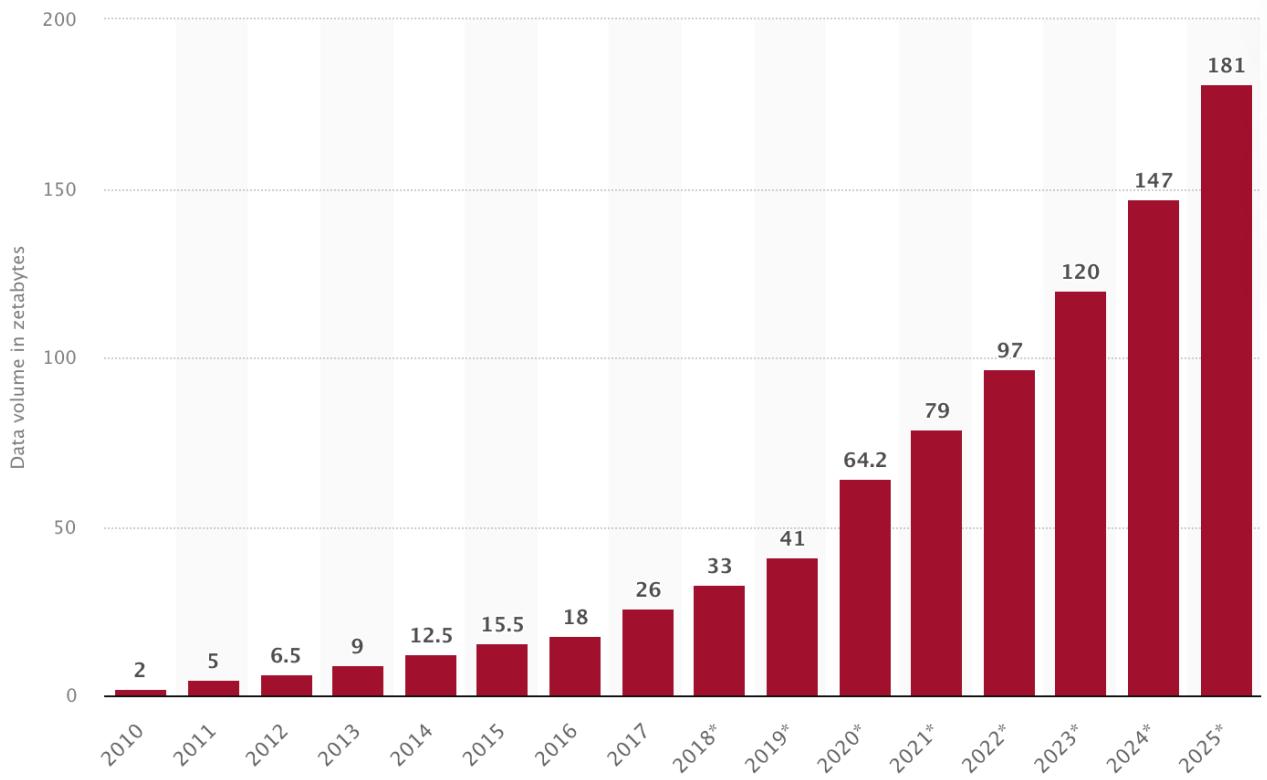
A History of Population Health
Rise and Fall of Disease in Europe
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Generative AI needs data, but how much?



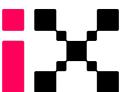
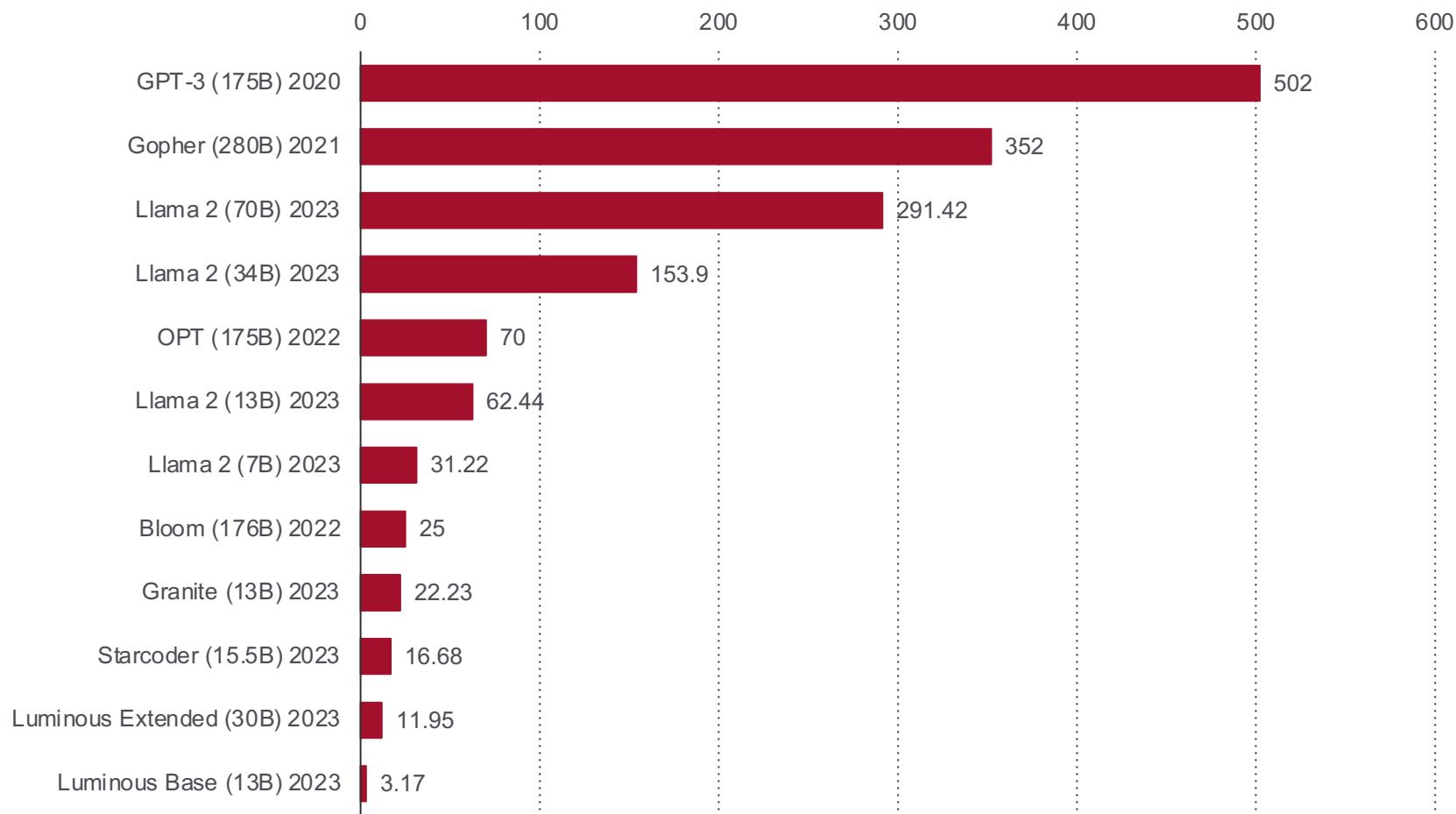
Volume of data/information created, captured, copied, and consumed worldwide from 2010 to 2020, with forecasts from 2021 to 2025 (in zettabytes = 10^{21} bytes)



© Statista 2024



CO₂ equivalent emissions by AI models 2024 in tonnes





AI BOOK CO-WRITING PROJECT

THURS
DAYS

29 MAY 2025

13:00 – 14:00 UK TIME



“**AI: THE GOOD, THE BAD
AND THE GAME CHANGER**”

HOW TO JOIN US?



Collaborative
Creative Projects

APPLICATION LINK: [HTTPS://SHORTURL.AT/YIGFU](https://shorturl.at/yigfu)

DEADLINE: 27 MAY 2025

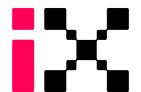


<https://basira-lab.com/>

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WRAP UP

The main trends

Generative learning

- *improve disease diagnosis/prognosis*
- *generate costly medical data from low-cost*
- *enable the forecasting of health/disease evolution (individual, population)*

Federated learning

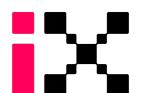
- *promote low-cost and privacy-preserving learning from data*
- *increase AI fairness by learning from diverse datasets*
- *learn better together and without data*

The other side to rethink

Holistic learning

Gen-AI, at which cost?

How to enable a more holistic learning and healing of diseases?

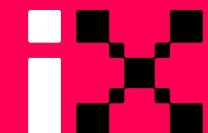


"The requisites of knowledge: a quick mind, zeal for learning, humility, foreign land, a professor's inspiration, and a life of long span."
Juwaini of Nishapur (d.1085)



**Brain And SIgnal Research & Analysis
Lab**

<https://basira-lab.com/>



**Generative
learning**

**Federated
learning**

**Holistic
learning**