



Artificial Intelligence in Medical Imaging for Precision Medicine

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Talk overview

- Medical imaging in Precision medicine
- AI Applications in imaging for precision medicine
 - *Individual subject-level*
 - *Population-level demographic factors-driven*
- Challenges in AI tool development
 - *Data/label limitations*
 - *Domain shifts across centres and scanners*
 - *Data sharing and privacy*
- Opportunities/future directions



What is precision medicine?

- Personalized care
 - subject's medical history / lifestyle factors
 - Disease biomarkers
 - Genetic factors
- Patient complains about mood changes and gait disturbances
 - Could be symptoms of many disorders
 - Knowing additional information could help (e.g., hypertension, lesions in specific regions, smoking/stress etc.)



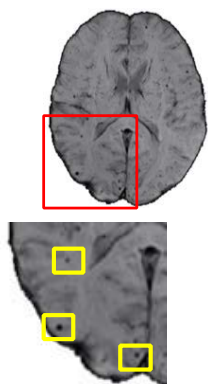
Medical imaging in precision medicine

- Early diagnosis
 - Routine scans, *incidental findings*
- Differential diagnosis
 - Differentiating diseases with similar symptoms
- Screening and treatment planning
 - *Longitudinal analysis*, response to drug administration
- Monitoring and predictive analysis
 - Disease severity prediction, *pre- vs post-surgery analysis*
- Data-driven and hybrid decision making
 - Using scans and *imparting domain (clinical) knowledge/factors*
- Further understanding of disease conditions
 - Large-scale epidemiological studies, *population-level modelling*



AI application in imaging for precision medicine

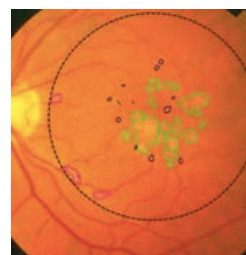
Early diagnosis



Detection of cerebral microbleeds

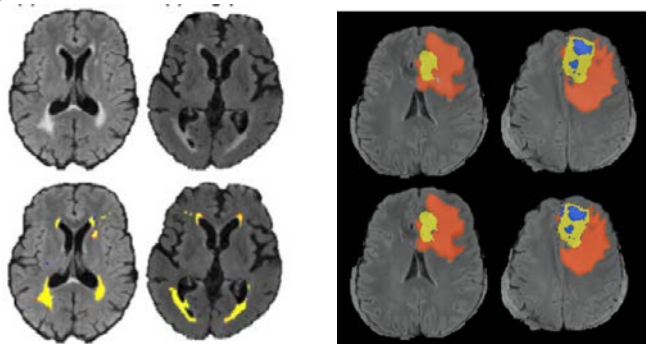
Early sign of neurodegeneration of various cerebrovascular diseases

Screening & monitoring



Screening diabetic retinopathy and macular degeneration

Scoring based on early, intermediate and late signs



Extraction of imaging biomarkers for diseases

Segmentation of white matter hyperintensities and brain tumor lesions – estimate the lesion load

Useful to investigate their distribution and rating to get more information on their incidence.

V Sundaresan et al. NeuroImage, 2019a

V Sundaresan et al. ISMRM, 2020

V Sundaresan et al. Medical Image Analysis, 2021

V Sundaresan et al. OMIA, MICCAI, 2015

V Sundaresan et al. IEEE EMBC, 2015

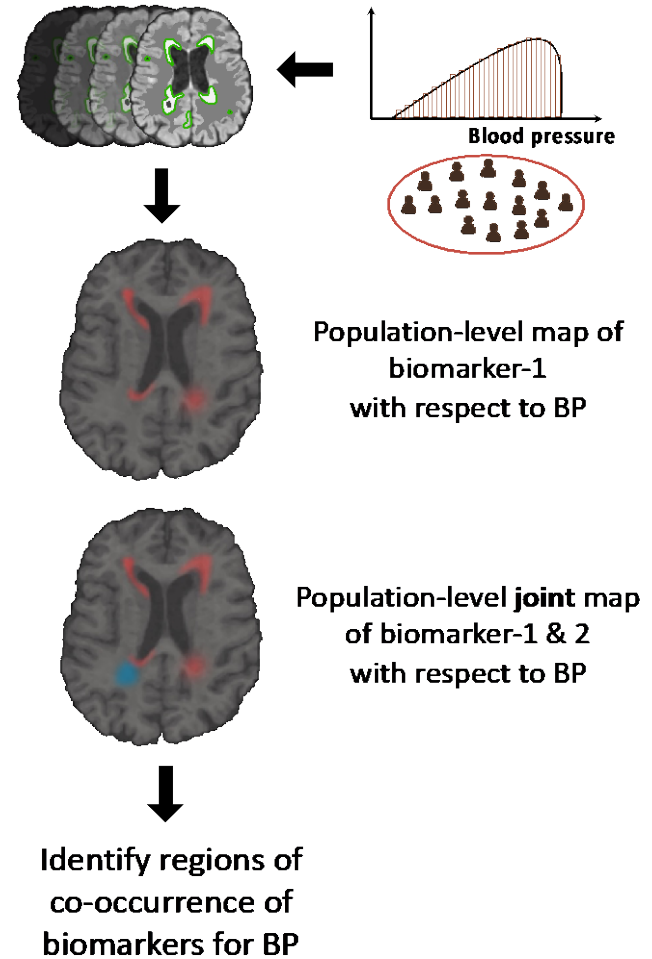
V Sundaresan et al, OHBM, 2018, 2019



Demographic factors-driven
precision diagnosis

Modelling the distribution
of anomalies within a
population

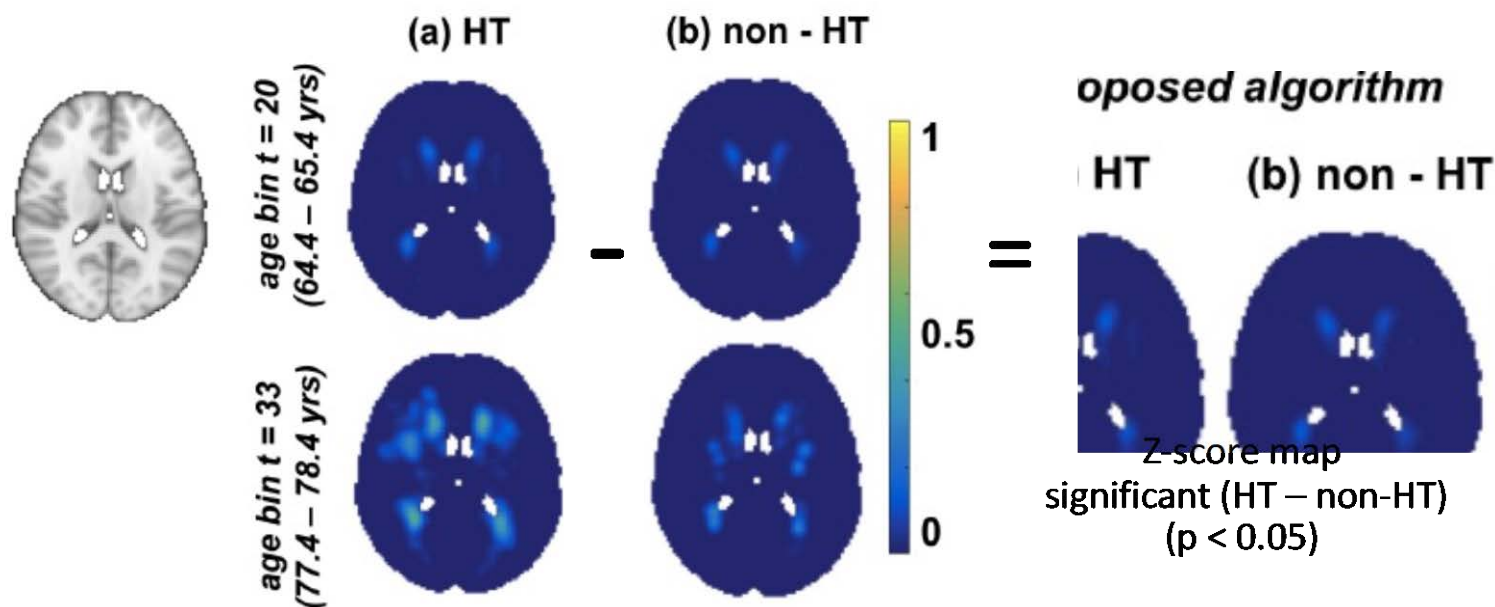
Useful in obtaining a
population-level prior for
specific diseases





Modelling white matter lesion distribution within a population

biobank^{uk} Largest brain imaging database, currently ~40,000 subjects scanned, with the target of 100,000 subjects. Big data - total database size: 500 TB.



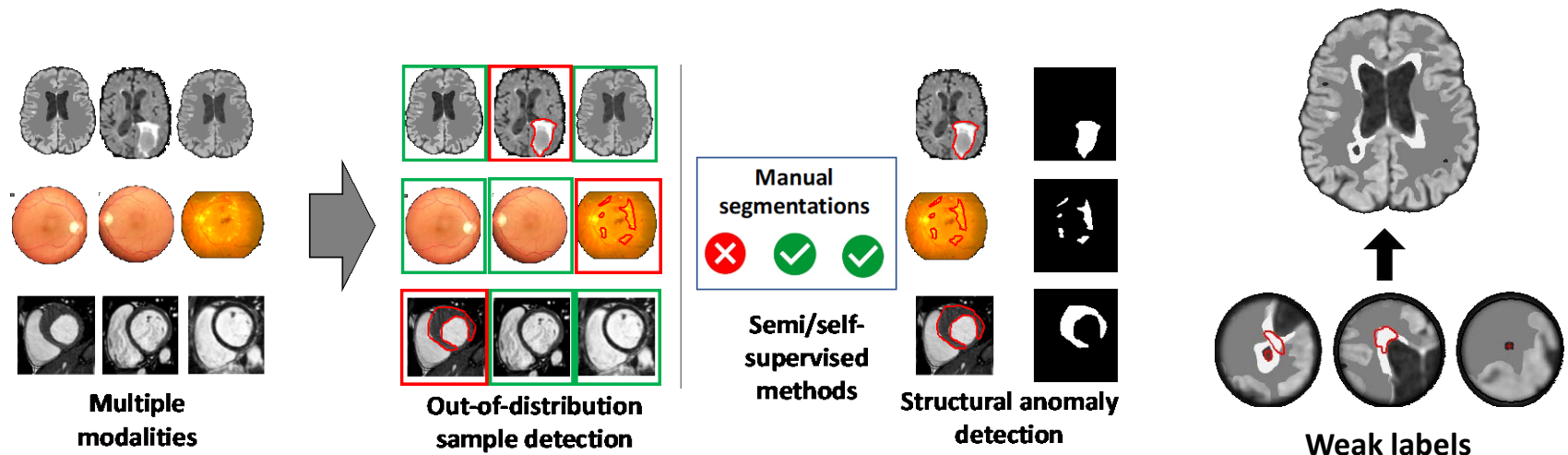


Challenges in AI tool development



Data/label limitations

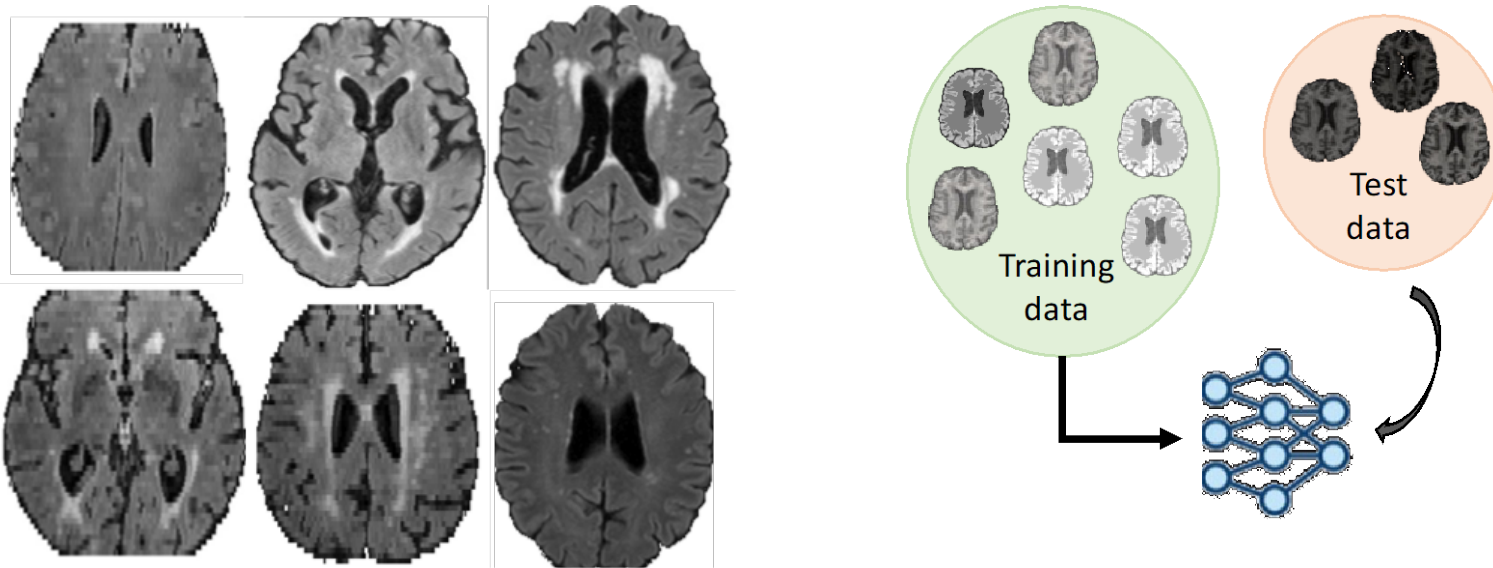
- Medical data is sensitive and difficult to obtain, have to be anonymized
- Manual annotation of medical imaging is time-consuming and expensive
- Self-/semi-supervised techniques





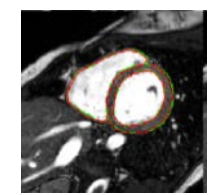
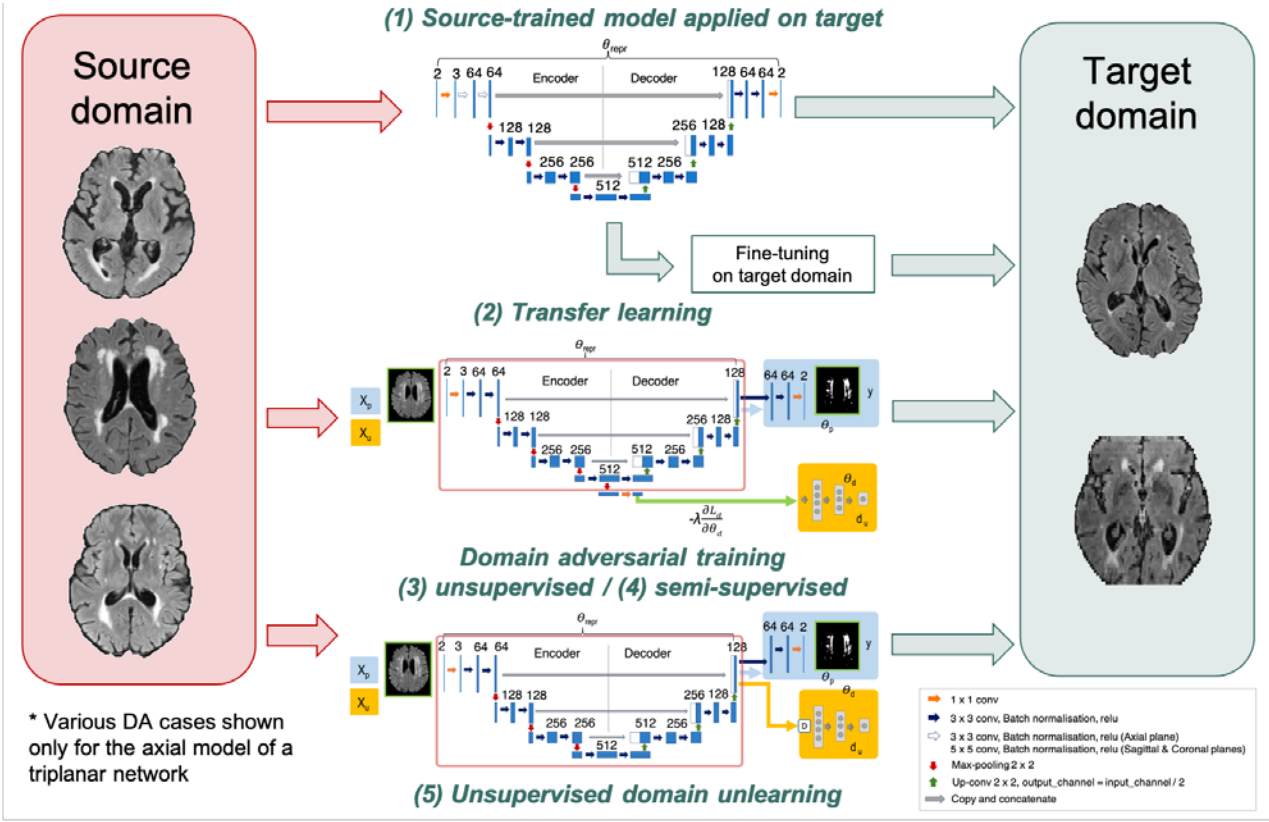
Data domain shifts in multi-centric data

Domain adaptation: Transfers knowledge from source to target domain by leveraging domain invariant features to reduce domain shift.





Comparison of domain adaptation (DA) techniques for neurodegeneration lesion segmentation



Multi-Centre, Multi-Vendor and Multi-Disease Cardiac MR segmentation

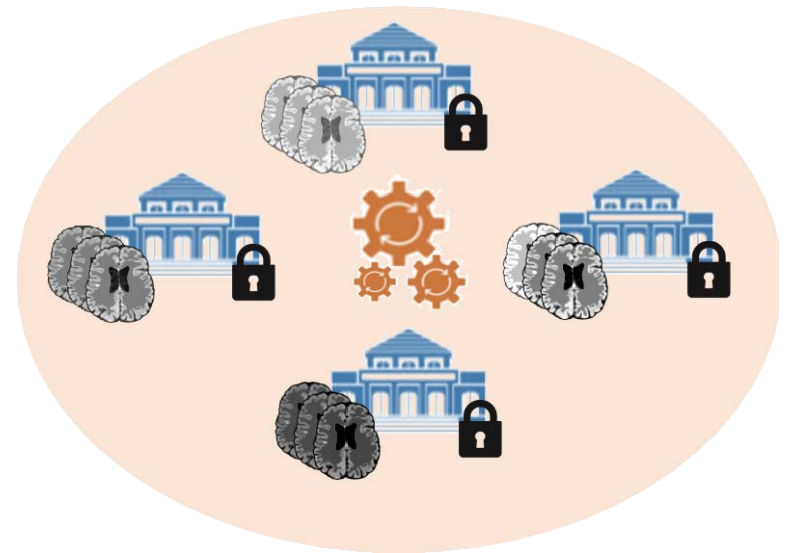


Classification of X-ray images from different domains



Data sharing and privacy protection

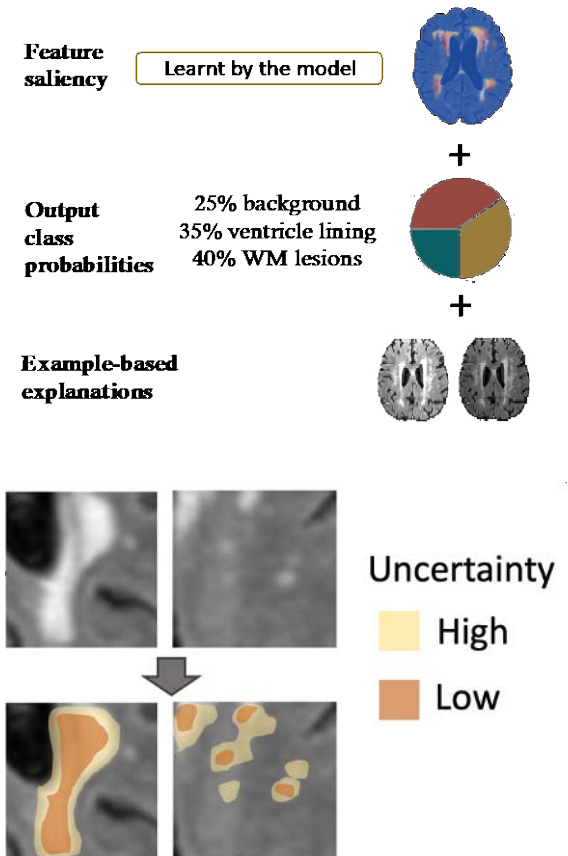
- Federated learning:
 - Preserves data privacy
 - Model training done individual centres rather than sharing data
 - Aggregate model parameters *centrally* from individual models
- Emerging areas:
 - Addition of unseen centres in FL, cross-modal domain adaptation, missing modalities





Opportunities/future directions

- Interpretability and Trust – explainability
 - Including clinicians in the loop
- Out-of-distribution detection
 - Open-world anomaly detection (open-set recognition)
 - Integration within a federated learning set-up
 - Anomaly informed precision diagnosis (incidental findings)
- Accurate extraction of underexplored biomarkers
 - Disentanglement of source (e.g., pathological vs scanner-induced)
 - Characterization and reporting





THANK YOU!

IISc, Bangalore

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