# Disease Knowledge Transfer across Neurodegenerative Diseases

Răzvan V. Marinescu<sup>1</sup>, Neil P. Oxtoby<sup>1</sup>, Alexandra L. Young<sup>1</sup>, Arman Eshaghi<sup>2</sup>, Peter A. Wijeratne<sup>1</sup>, Daniel C. Alexander<sup>1</sup>

Centre for Medical Image Computing, University College London

<sup>2</sup>Queen Square MS Centre, UCL Institute of Neurology

(2)

(3)

# Introduction

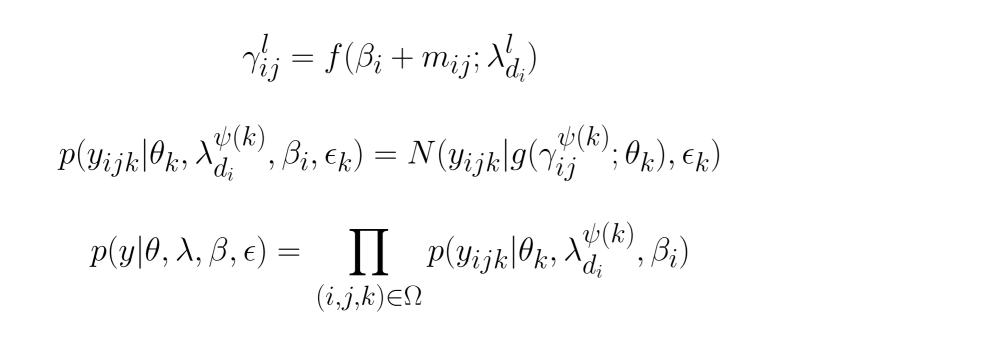
- Recently disease progression models work well on large, multimodal datasets
- Datasets on rare neurodegenerative diseases (e.g., Posterior Cortical Atrophy) are limited and cross-sectional.

## Aim

Develop a transfer learning model to transfer knowledge between diseases

# Method

• We propose Disease Knowledge Transfer:



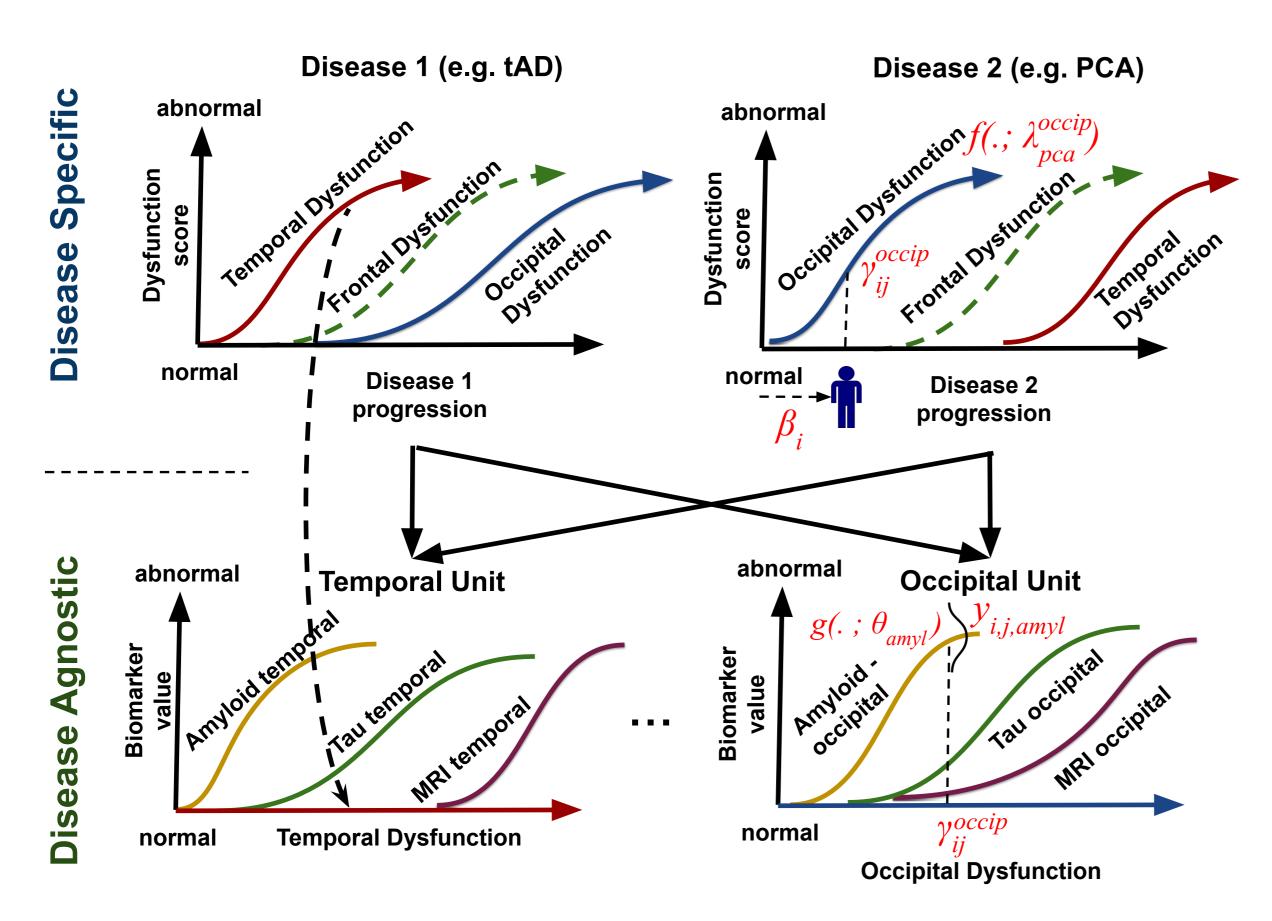


Figure 1

#### References

- 1. Fontejin et al., Neuroimg., 2012
- 3. Villemagne et al., Lancet Neurol., 2013
- 2. Young et al., Nat. Comms., 2018
- 4. Marinescu et al., IPMI, 2017

## Results

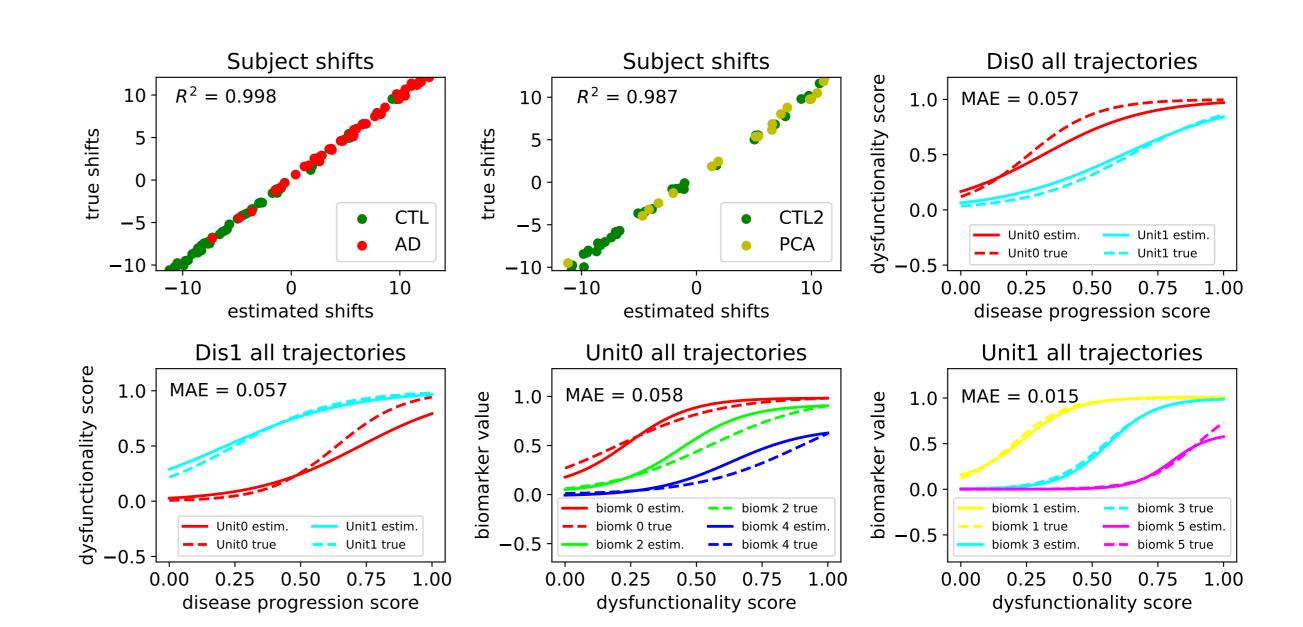
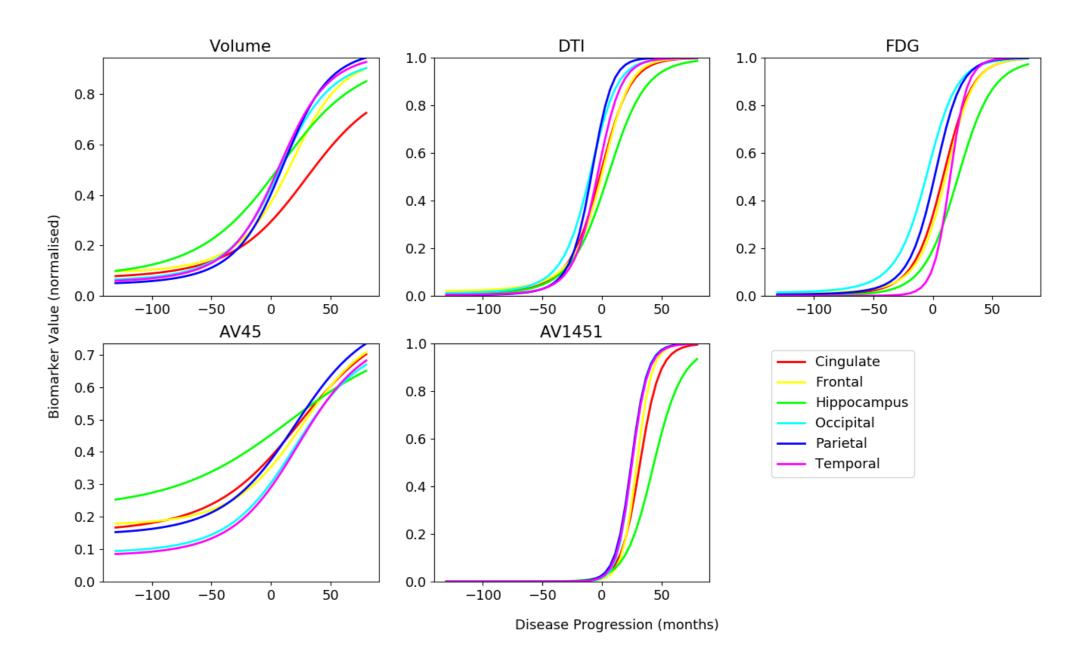


Figure 2: Comparison between true and DKT-estimated subject timeshifts and biomarker trajectories.



**Figure 3:** Estimated trajectories for the PCA cohort. The only data that were available were the MRI volumetric data. The dynamics of the other biomarkers has been inferred by the model using data from typical AD, and taking into account the different spatial distribution of pathology in PCA vs tAD.

## Conclusion

#### Weblinks

- UCL Progression of Neurodegenerative Disease (POND): cmic.cs.ucl.ac.uk/pond/
- UCL Centre for Medical Image Computing: www.ucl.ac.uk/cmic/





