Towards precision medicine in mental health from a life-span perspective

Session 4: Developing and Advancing Phenotyping and Biomarker Discovery to Enable Patient Stratification

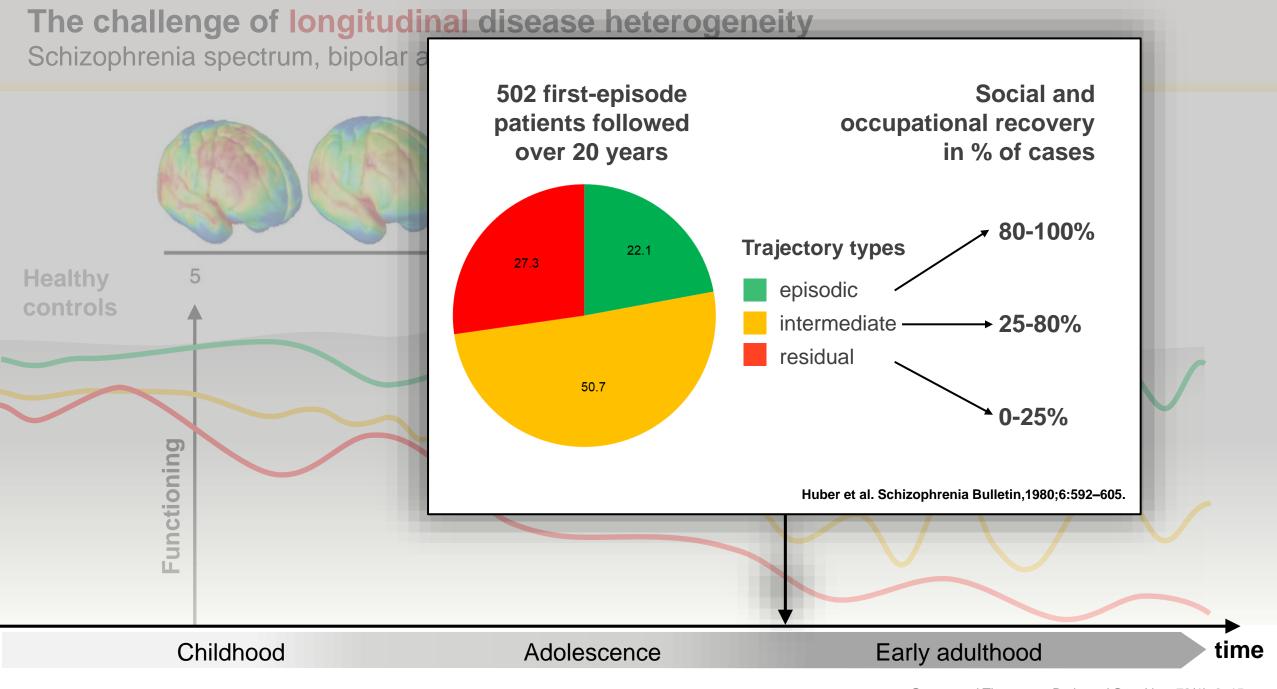
From Molecular Insights to Patient Stratification for Neurological and Psychiatric Disorders: A Workshop National Academy of Sciences, Engineering and Medicine

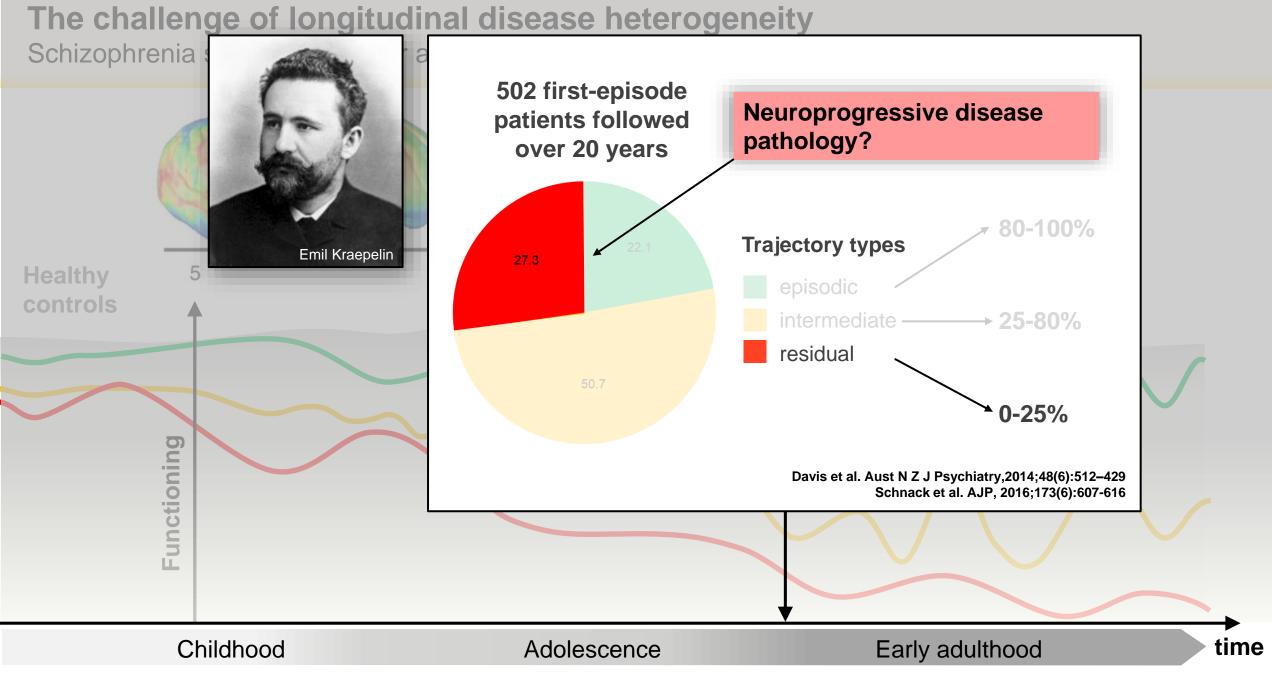
10/6/2021

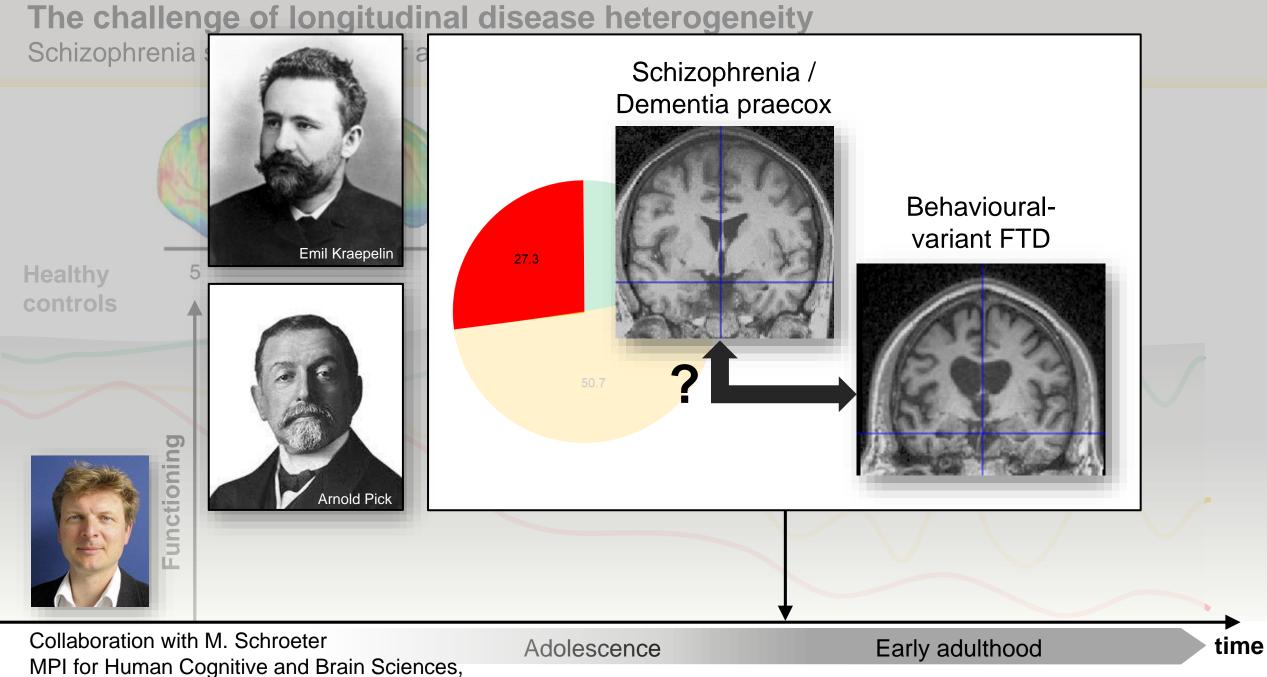


Nikolaos Koutsouleris

Professor for Precision Psychiatry
Ludwig-Maximilian-University | King's College London
Max-Planck Fellowship Group for Precision Psychiatry

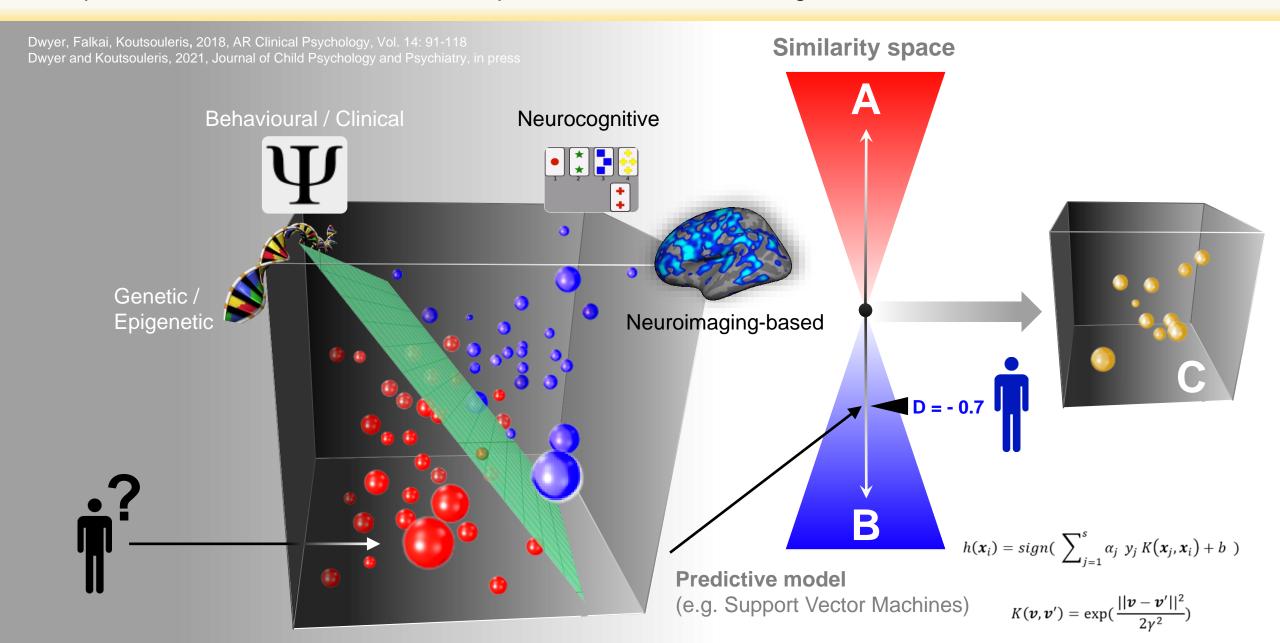




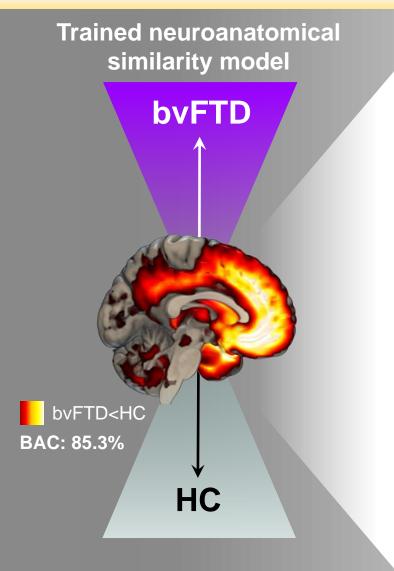


MPI for Human Cognitive and Brain Sciences, Leipzig

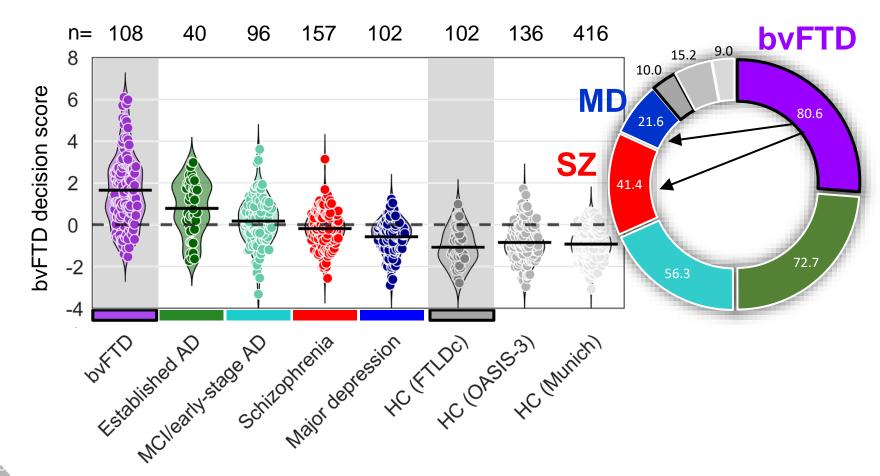
Comparative neuroscience based on supervised machine learning



Comparative neuroscience strategies crossing the borders between psychiatry and neurology



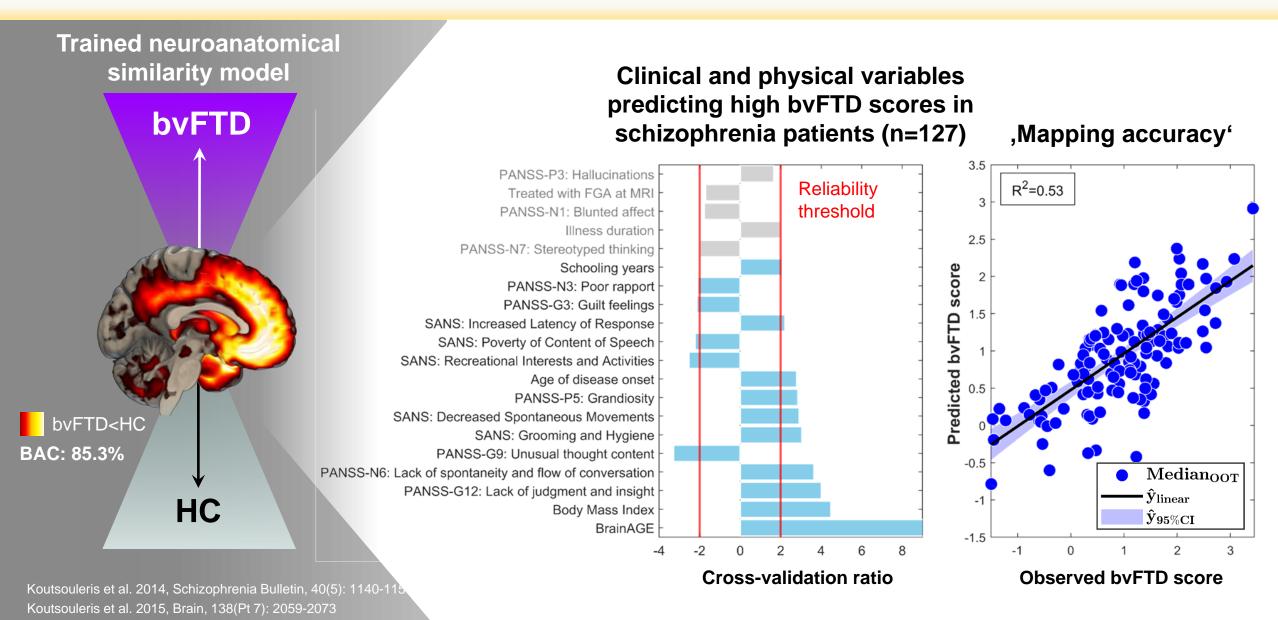
sMRI-based bvFTD classifier applied across different diagnostic cohorts



Koutsouleris et al. 2014, Schizophrenia Bulletin, 40(5): 1140-112 Koutsouleris et al. 2015, Brain, 138(Pt 7): 2059-2073 Koutsouleris et al. 2021, under review

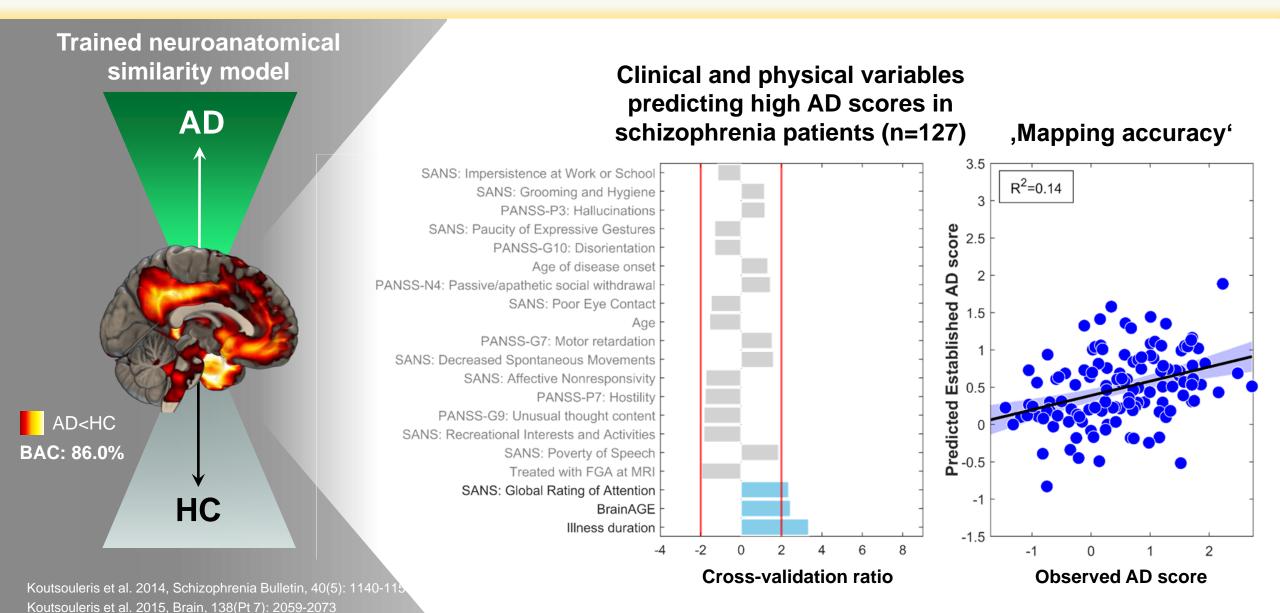
Koutsouleris et al. 2021, under review

Comparative neuroscience strategies revealing shared clinical phenotypes between SCZ and bvFTD

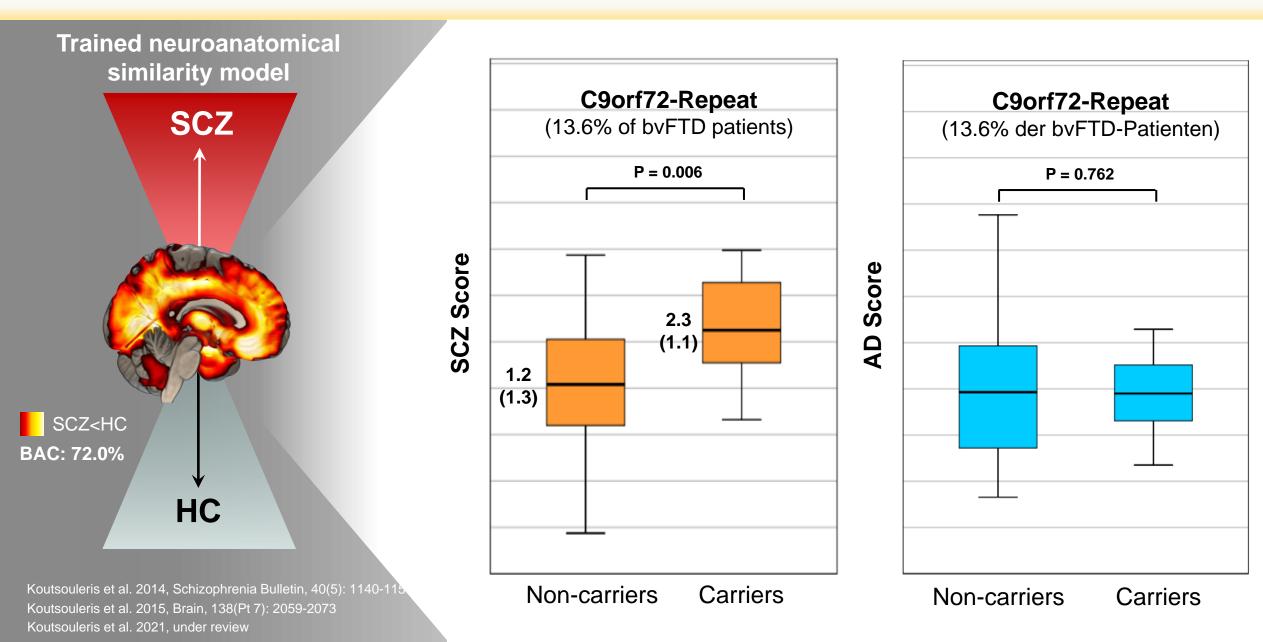


Koutsouleris et al. 2021, under review

Comparative neuroscience strategies showing specific predictability of bvFTD brain patterns in SCZ

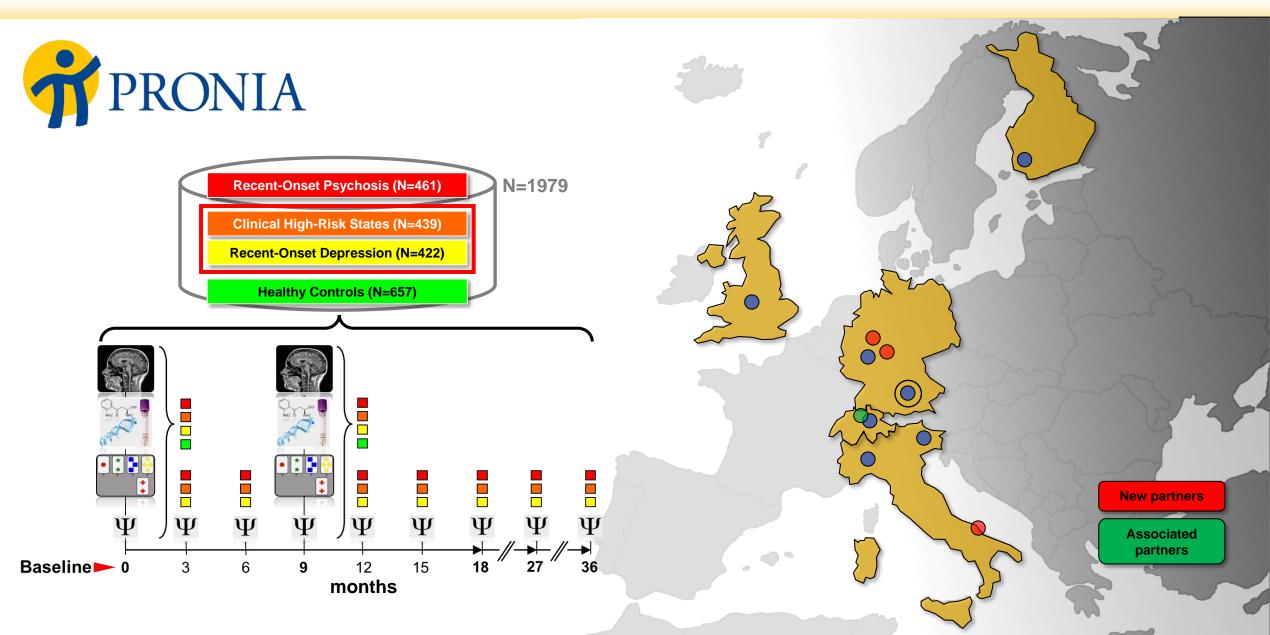


Comparative neuroscience strategies showing C9orf72 association of SCZ pattern expression in bvFTD



Understanding longitudinal heterogeneity

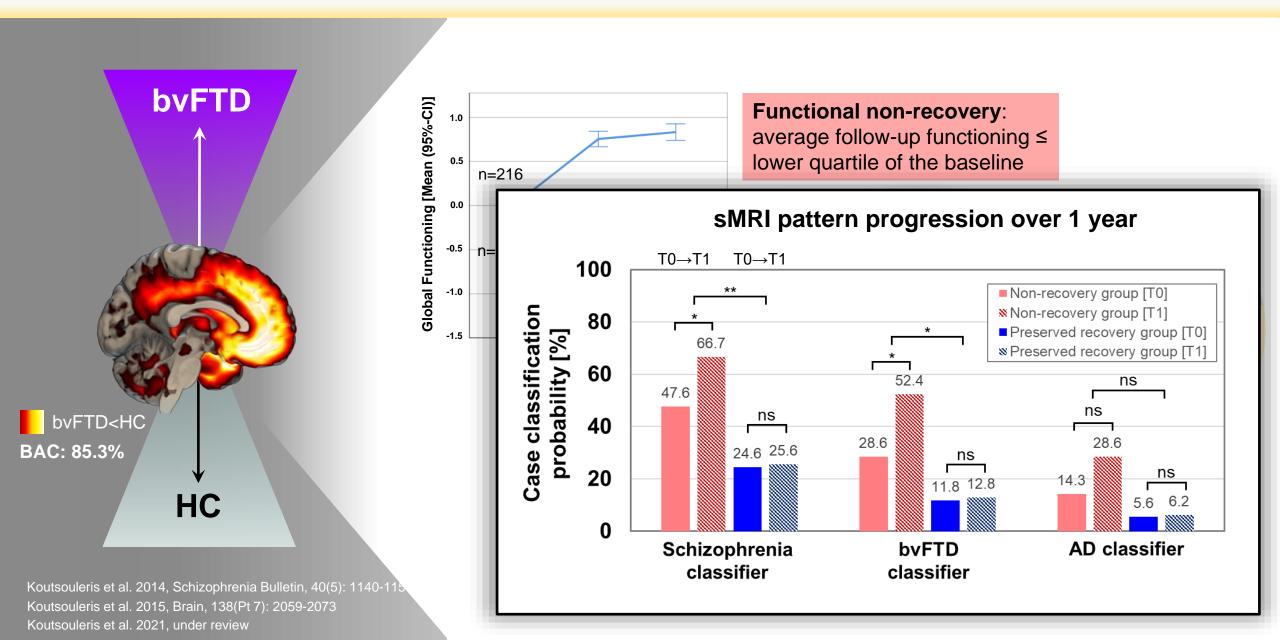
Can we learning something from these cross-sectional analyses for predicting outcomes?



Understanding longitudinal heterogeneity

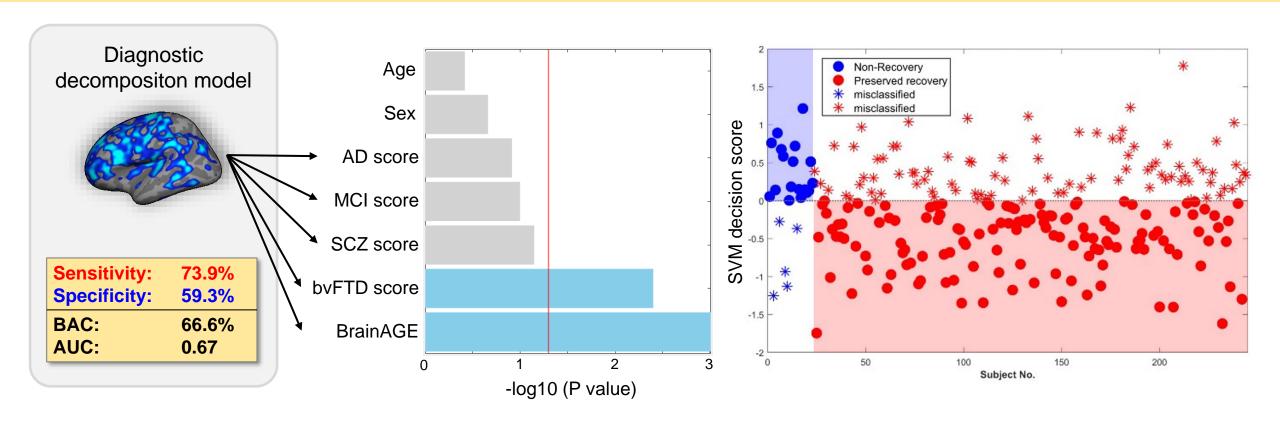
Testing the neuroprogression hypothesis in early-stage disease samples





From promise to practice

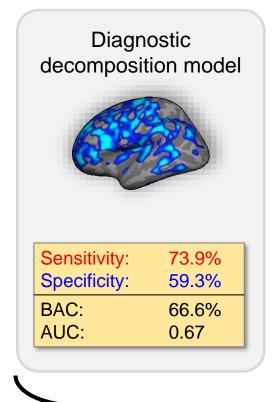
Using diagnostic decompositions to predict 2-years non-recovery courses in CHR and ROD patients

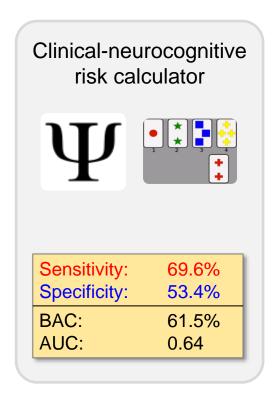


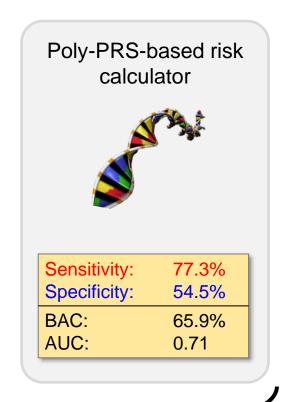


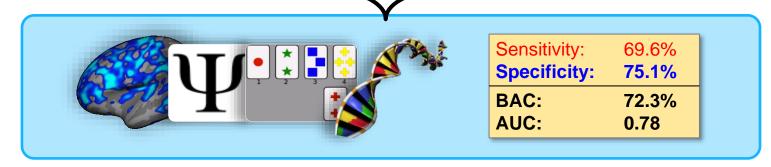
From promise to practice

Using multiple data domains to predict 2-years non-recovery courses in CHR and ROD patients



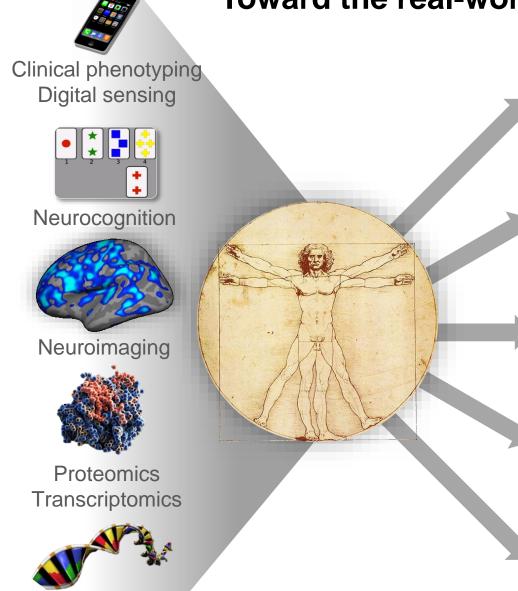








Toward the real-world implementation of biomarkers



Genetics

To assess and compare data domains for their diagnostic, prognostic and theragnostic utility.

To test models across different phenotypes, patient populations and disease gradients/stages.

Based on lessons learned, to iteratively refine clinical applications areas for 'algorithmic augmentation'.

To test clinical effectiveness of model-informed treatments in biomarker-stratified clinical trials.

To improve models and treatments via bedside-tobench and bench-to-beside development cycles.