

Evaluating Harmonisation Methods for Multi-Site Psychosis MRI Data Insights from the Psy-ShareD Resource

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Introduction

- MRI is widely used to study brain structure, but variability across scanners, sites, and acquisition protocols can introduce differences in the data, generating site-effects which can lead to biased or inconsistent results and reduce reproducibility [1].
- Another challenge in MRI studies is the limited availability of large, diverse samples due to the high cost and logistical challenges of scanning. This issue is particularly challenging in studies of psychiatric disorders like schizophrenia, where sample sizes are limited [2, 3].
- Harmonisation methods aim to mitigate these differences, enhancing comparability of MRI data [1, 4, 5, 6]. They can be broadly divided in image-level and feature-level.

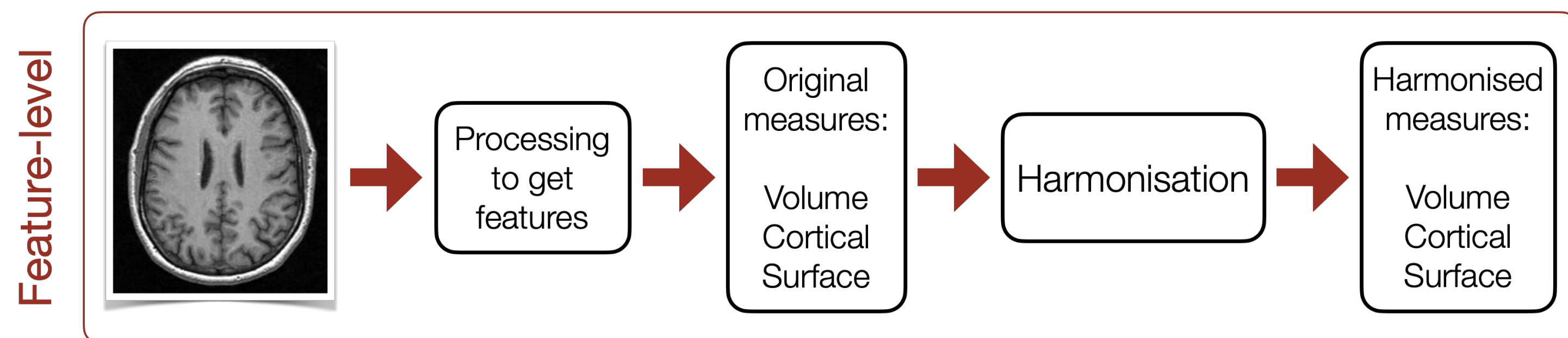
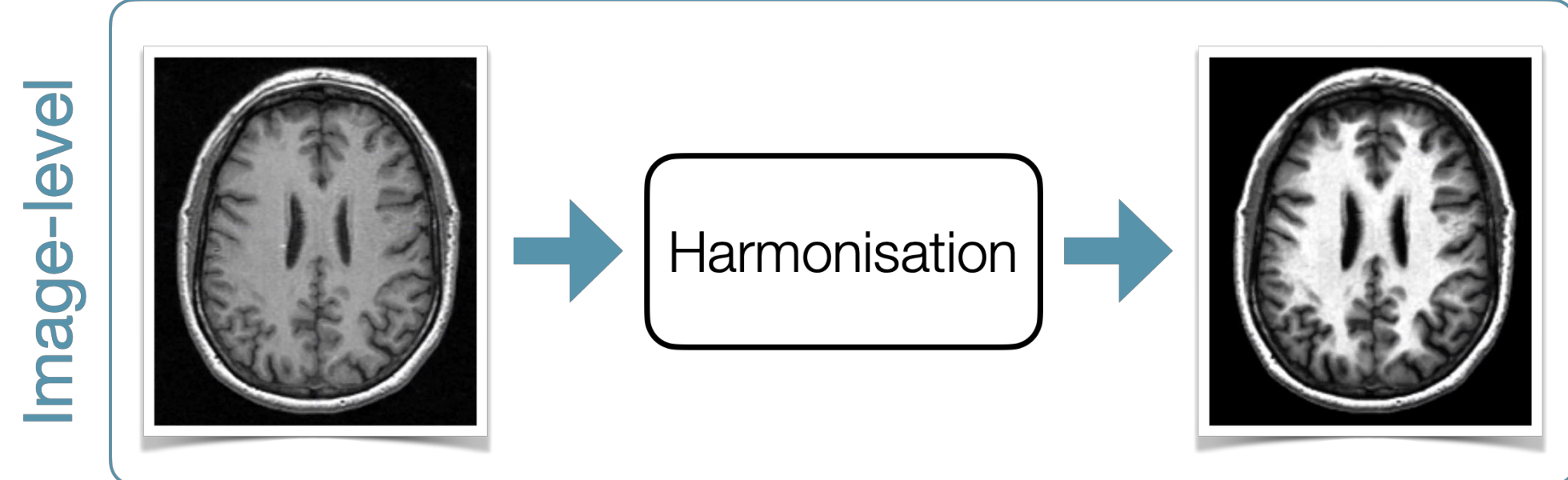
Objective: To compare different harmonisation techniques applied to T1w MRI data from the Psy-ShareD, specifically for individuals with schizophrenia and healthy controls. We assess their ability to preserve relevant biological information and mitigate scanner-related variability.



What is PsyShareD?

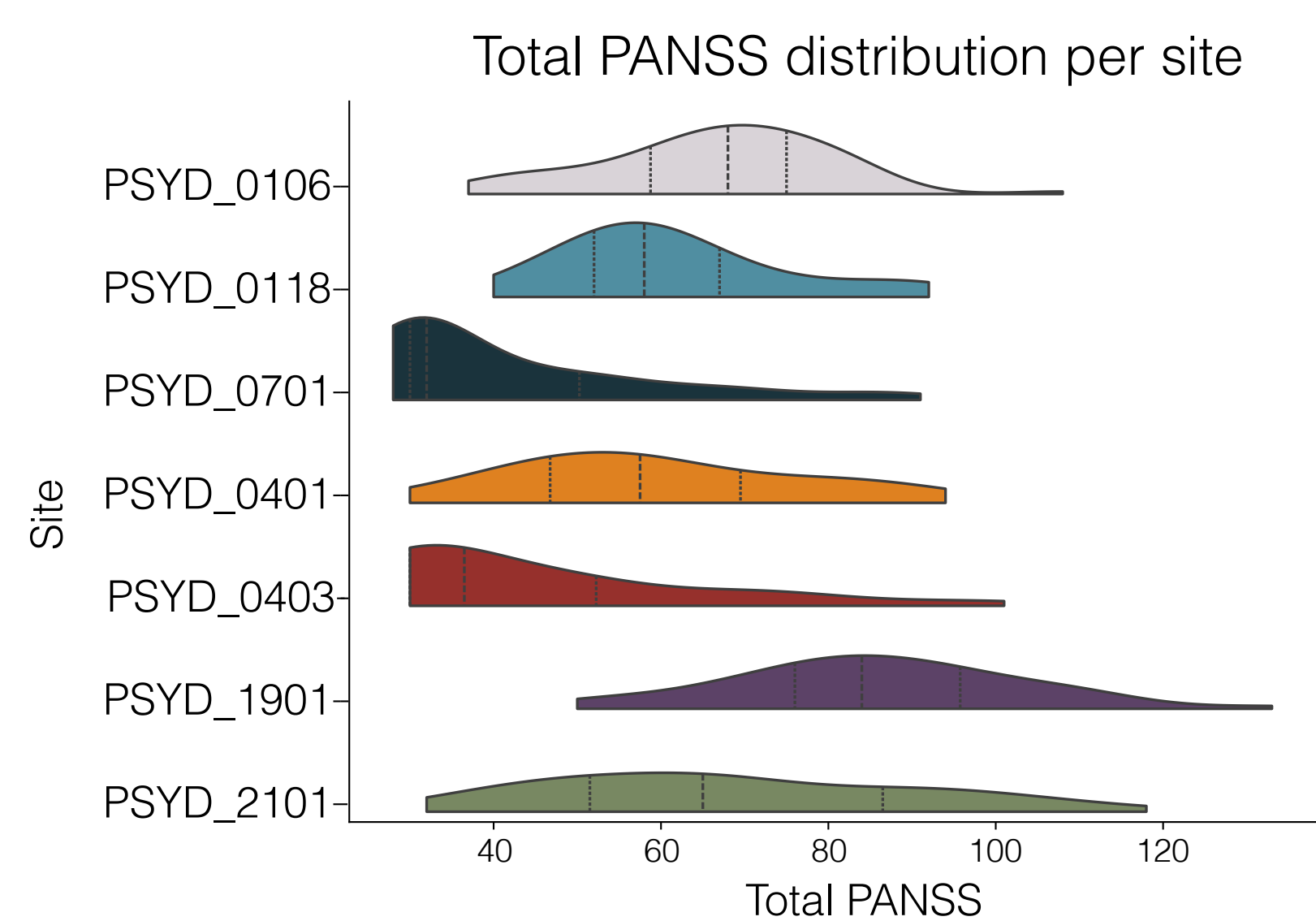
It is a free-to-access database that hosts pre-existing structural MRI data (mainly T1-weighted images) collected from different sites around the world.

Methods



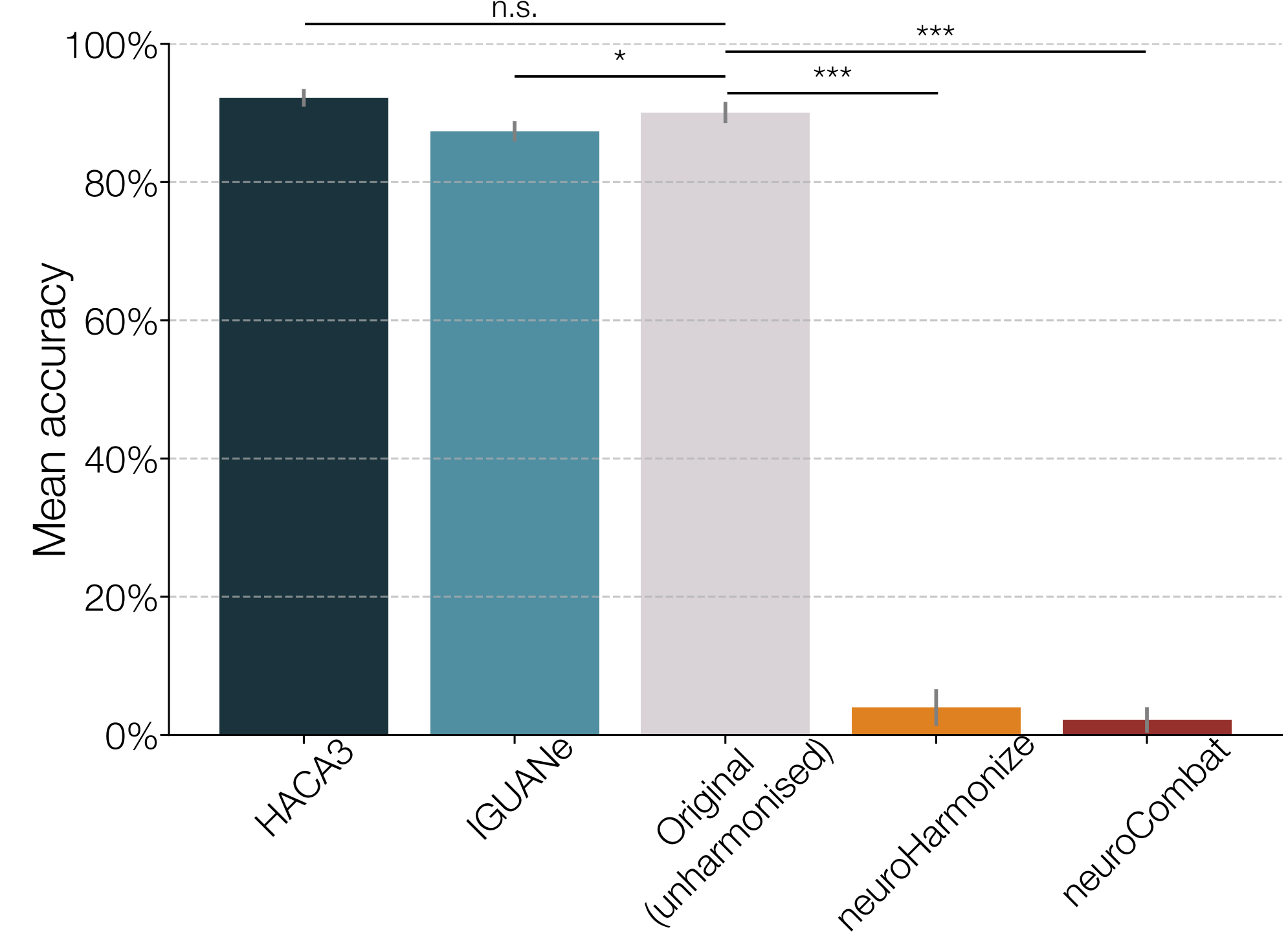
- Image-level harmonisation methods: HACA3 [5] and IGUANE [6]
- Feature-level harmonisation methods: neuroHarmonize [1] and neuroCombat [4]

- Data from 7 studies in 5 sites across 3 countries (UK, Italy and Japan)
- 269 healthy controls (101 females)
- 295 individuals with schizophrenia (85 females)

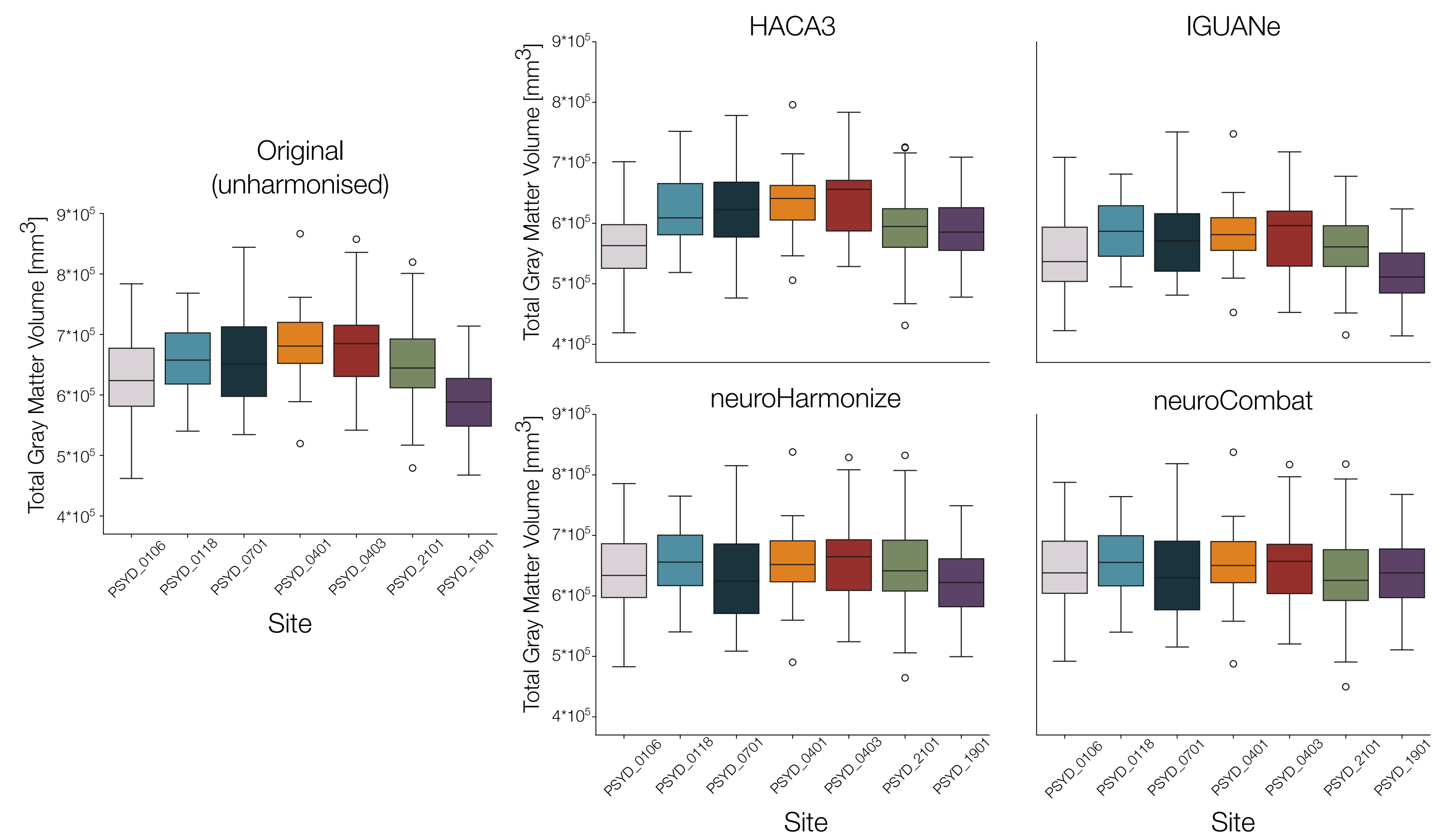


Results

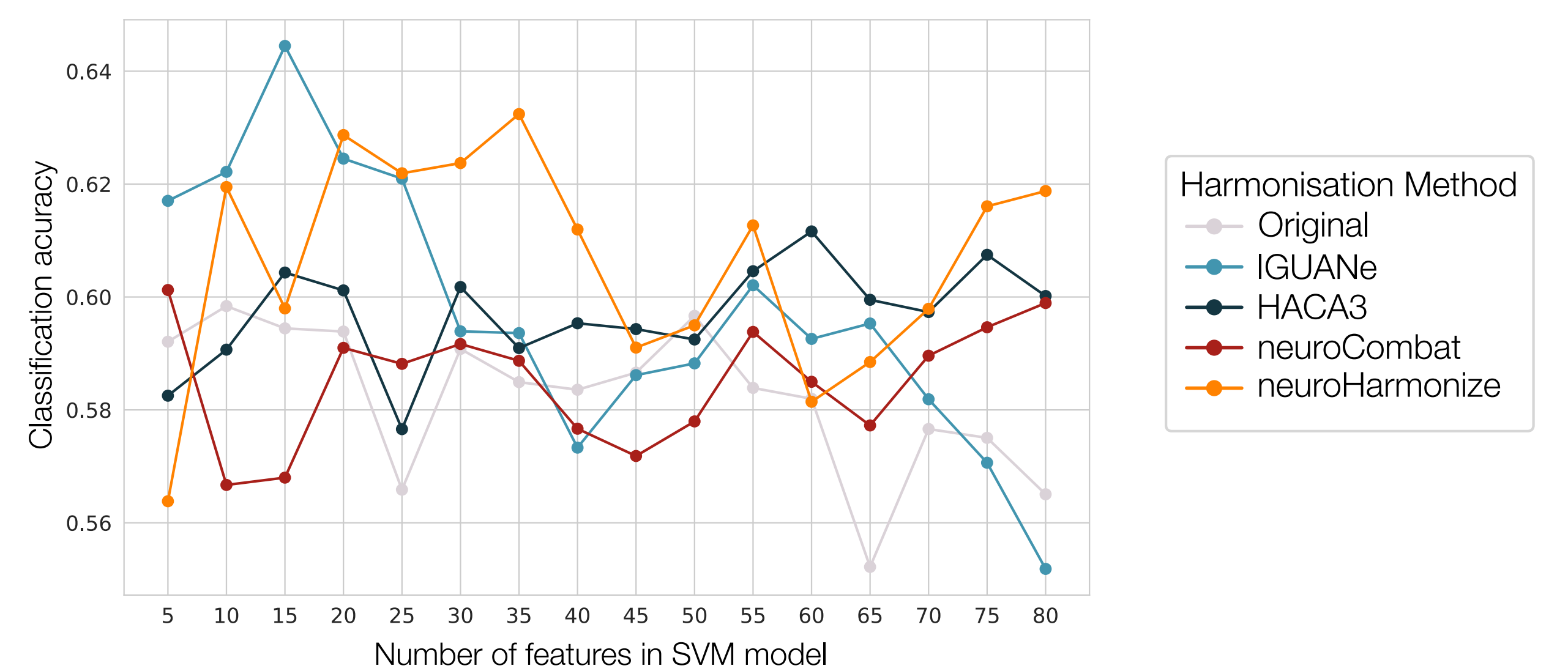
Site classification accuracies per harmonisation method



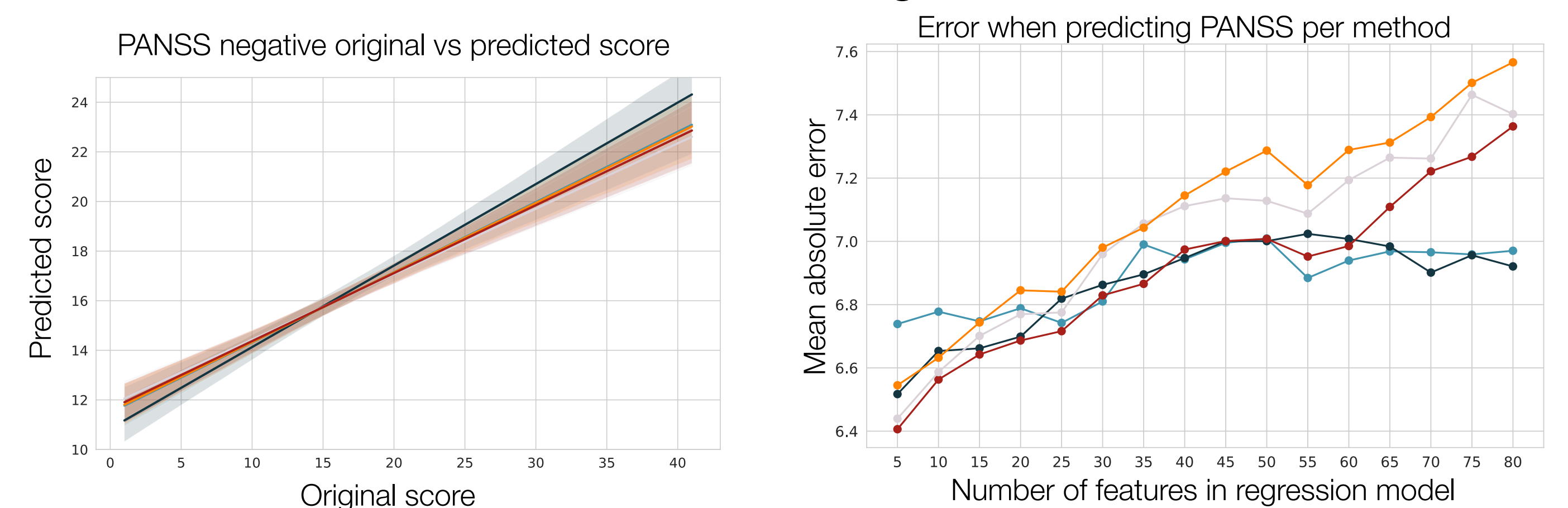
Mean total Gray Matter volume per harmonisation method



Classification accuracy of SVM into schizophrenia vs. control



Prediction of PANSS negative score



Discussion

- Feature-level harmonisation methods effectively removes most site-related variance, as shown by site classification and total GM volume results.
- IGUANE image-level harmonisation significantly reduces site recognisability, though to a lesser extent than feature-based methods, while HACA3 does not reduce it.
- Harmonised images perform similarly to original images in schizophrenia classification and symptom prediction (PANSS negative).
- Reducing site-related variance is important, but it should not come at the cost of removing biologically relevant information. Image-level harmonisation methods, particularly IGUANE, show promising progress in achieving this balance.
- Careful selection of harmonisation methods is crucial and should align with the study's objective.

References: [1] Pomponio et al., (2020) NeuroImage; [2] Evans et al., (2025) Schizophrenia Research; [3] Allen et al., (2025) Human Brain Mapping; [4] Fortin et al., (2018) NeuroImage; [5] Zuo et al., (2023) Computerized Medical Imaging and Graphics; [6] Roca et al., (2025) Medical Image Analysis