

Effect of data leakage in brain MRI classification using 2D convolutional neural networks

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In recent years, 2D convolutional neural networks (CNNs) have been extensively used to diagnose neurological diseases from magnetic resonance imaging (MRI) data due to their potential to discern subtle and intricate patterns. Despite the high performances reported in numerous studies, developing CNN models with good generalization abilities is still a challenging task due to possible data leakage introduced during cross-validation (CV). In this study, we quantitatively assessed the effect of a data leakage caused by 3D MRI data splitting based on a 2D slice-level using three 2D CNN models to classify patients with Alzheimer’s disease (AD) and Parkinson’s disease (PD). Our experiments showed that slicelevel CV erroneously boosted the average slice level accuracy on the test set by 30% on Open Access Series of Imaging Studies (OASIS), 29% on Alzheimer’s Disease Neuroimaging Initiative (ADNI), 48% on Parkinson’s Progression Markers Initiative (PPMI) and 55% on a local de-novo PD Versilia dataset. Further tests on a randomly labeled OASIS-derived dataset produced about 96% of (erroneous) accuracy (slice-level split) and 50% accuracy (subject-level split), as expected from a randomized experiment. Overall, the extent of the effect of an erroneous slice-based CV is severe, especially for small datasets.

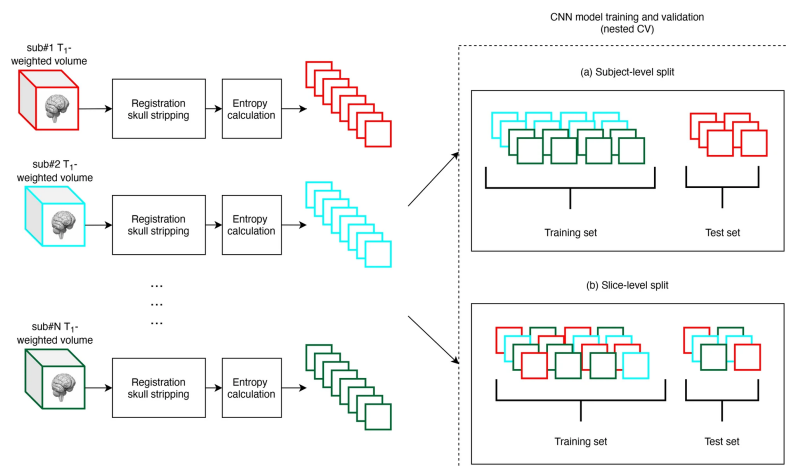


Figure 1: Schematic diagram of the overall T1-weighted MRI data processing and validation scheme. First, a preprocessing stage included co-registration to a standard space, skull-stripping and slices selection based on entropy calculation. Then, CNNs model’s training and validation have been performed on each dataset in a nested CV loop using two different data split strategies: (a) subject-level split, in which all the slices of a subject have been placed either in the training or in the test set, avoiding any form of data leakage; (b) slice-level split, in which all the slices have been pooled together before CV, then split randomly into training and test set [1].

References

[1] E. Yagis *et al.*, *Sci. Rep.* **11**, 22544 (2021). DOI: [10.1038/s41598-021-01681-w](https://doi.org/10.1038/s41598-021-01681-w)