

FASTGPR: DIVIDE-AND-CONQUER TECHNIQUE IN NEUROIMAGING DATA SHORTENS TRAINING TIME AND IMPROVES ACCURACY



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Introduction

- •Gaussian process regression (GPR) has shown great potential for studying healthy aging and disease via brain-age prediction (BAP) using structural MRI[1].
- A big drawback of GPR is the training complexity which is an O(N^3) operation (N=number of data points).
- The need for expansive datasets and the **high dimensionality of MRI data**, renders the training of GPR impractical with conventional computing resources.
- We investigated whether a divide-and-conquer approach can be used together with the GPR model.

Material

- •**T1w MRI** scans of healthy subjects from IXI [2], eNKI [3], CamCAN [4] (each n>500, **total N=1810**, 18-88 age range).
- •All analyses were run on an Apple M2 pro (12-core) processor with 32GB RAM, under the same conditions.

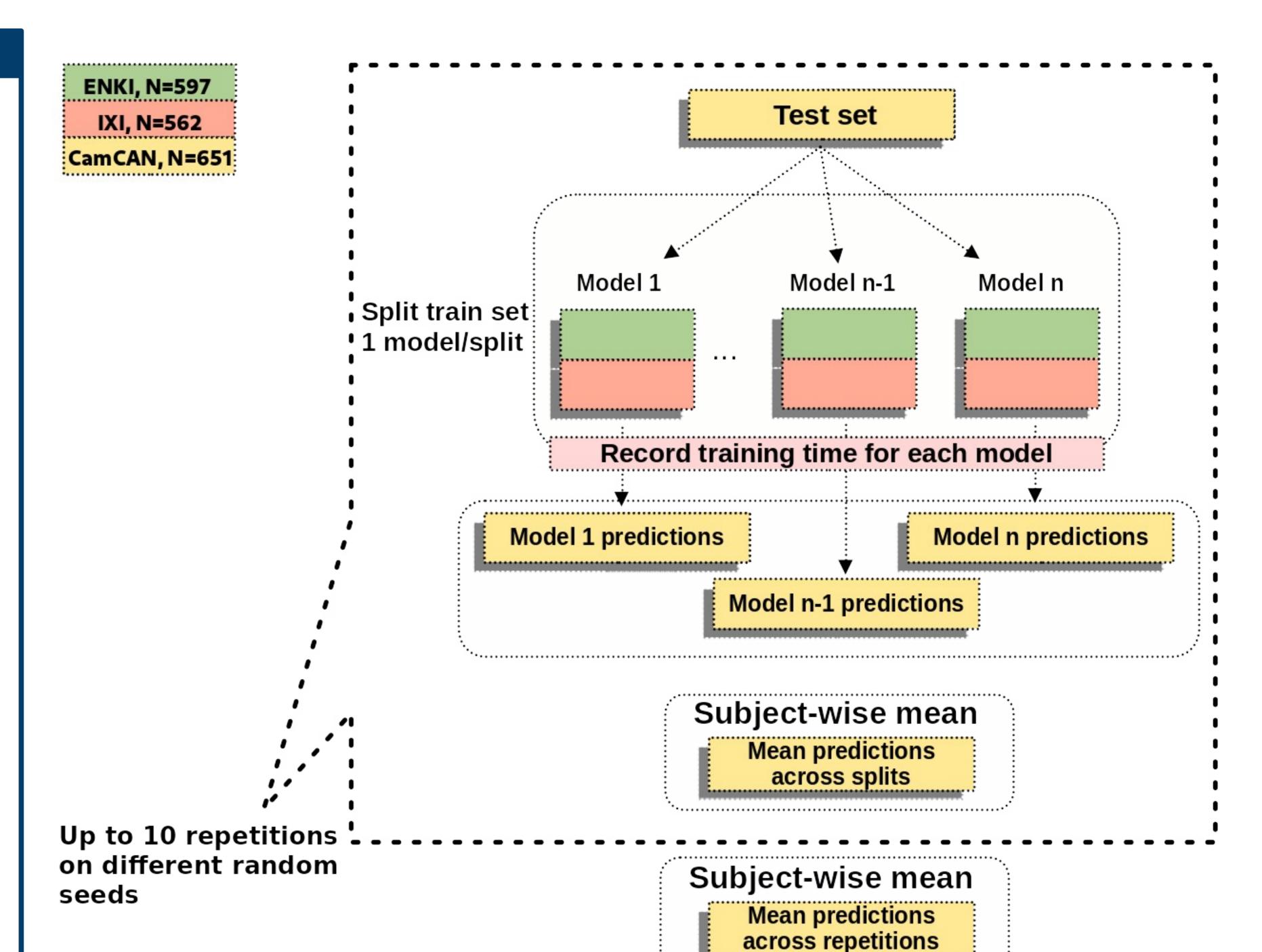
Methods

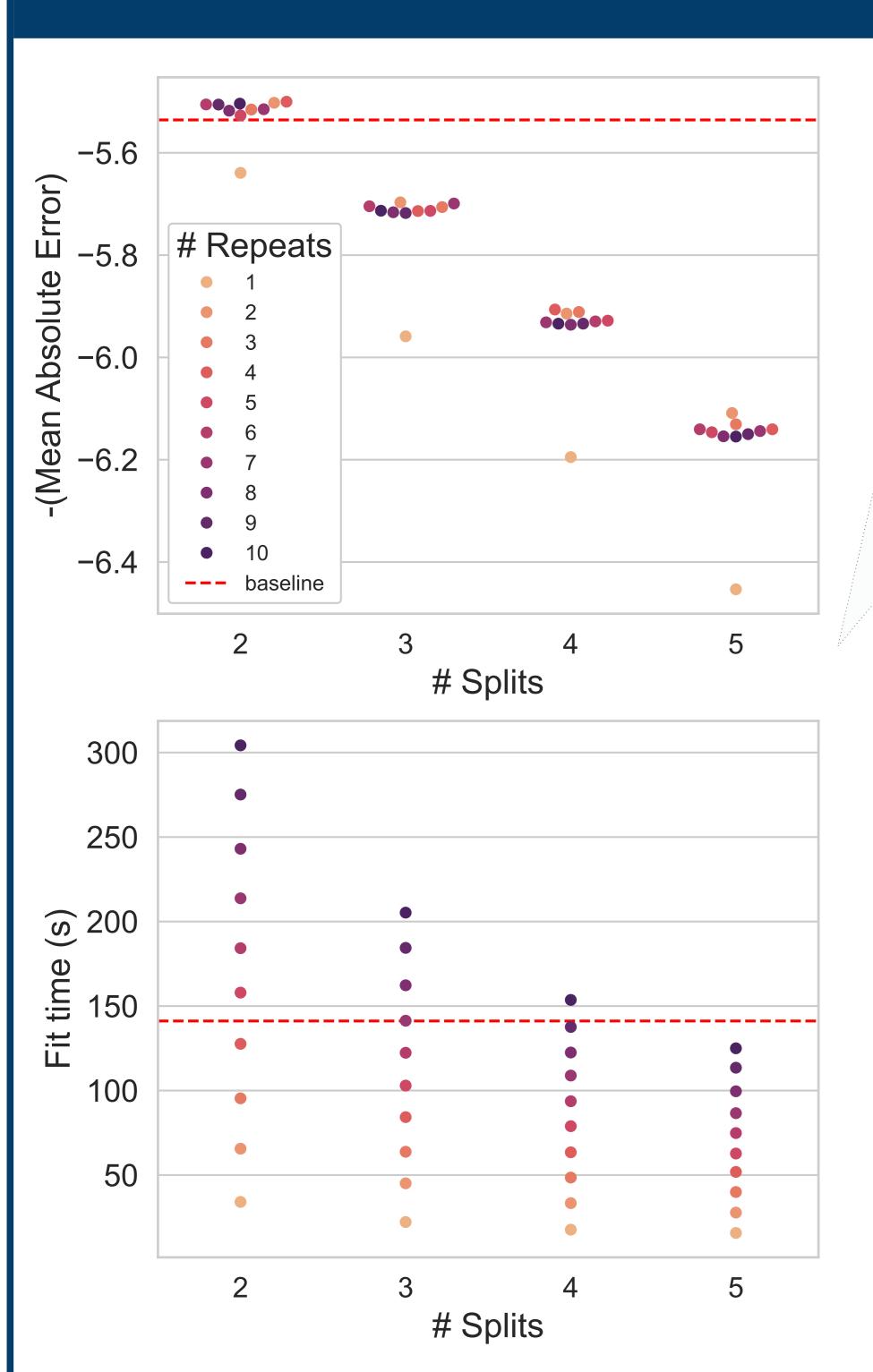
MRI preprocessing: CAT 12.8 [5] \rightarrow linear and non-linear spatial normalization, tissue segmentation and modulation.

238,955 voxels per subject representing voxel-wise gray matter volume.

Smoothing \rightarrow 4 mm FWHM Gaussian kernel, **Resampling** (linear interpolation) \rightarrow 8 mm spatial resolution Finally **n=3747** features per subject.

- Performance estimated in terms of mean absolute error (MAE) using Leave-One-Site-Out.
- Randomly divided training data into non-overlapping subsets while stratifying the splits over age.
- One GPR model was trained on each subset
- •Final prediction was obtained by averaging the predictions of all the models.
- We implemented this process with two-, three-, fourand a five-way split of the training data.
- Repeated up to ten times with distinct random seeds.
 The prediction for a test sample was obtained by averaging predictions across the splits.





Results & Conclusions

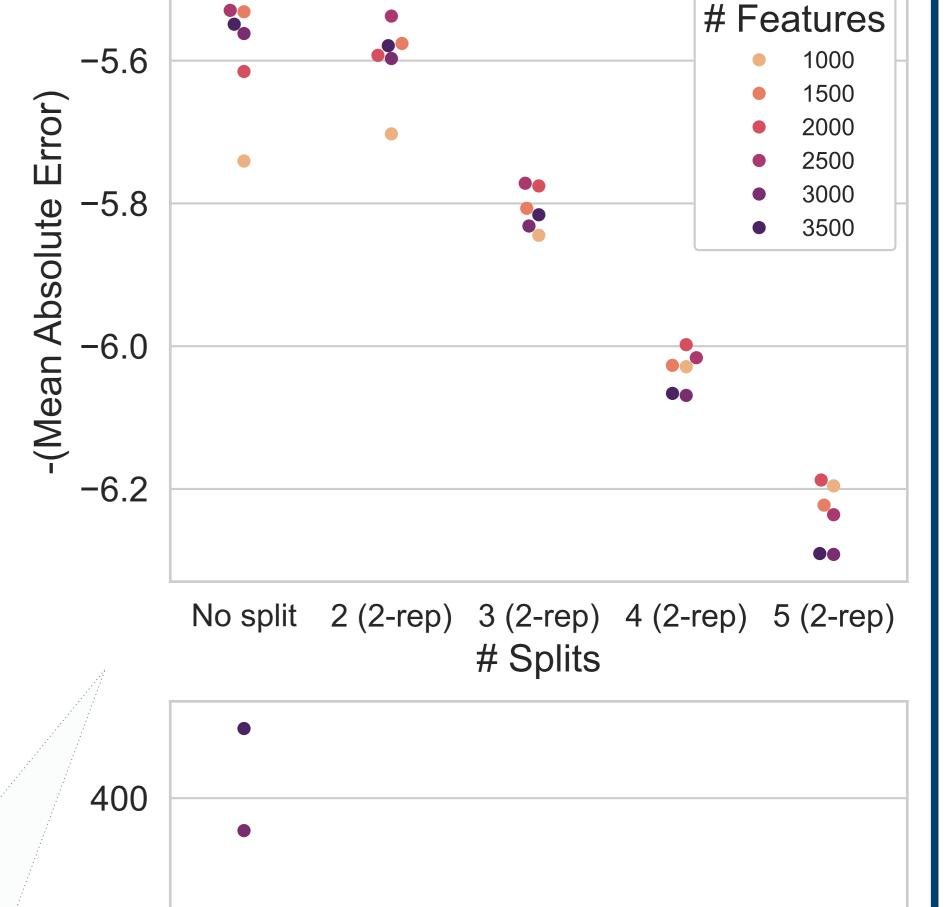
- ** training time is estimated as
- ** no-split model's performance (red line)
- 2-splits MAE=5.63 total compute time=24.2% of the no-split model's time
- The 3-, 4- and 5-split configurations manifest a reduction in training time and a pronounced drop in predictive performance
- 2 repetitions (2 random seeds) of 2-splits MAE=5.5, training time 46.4%

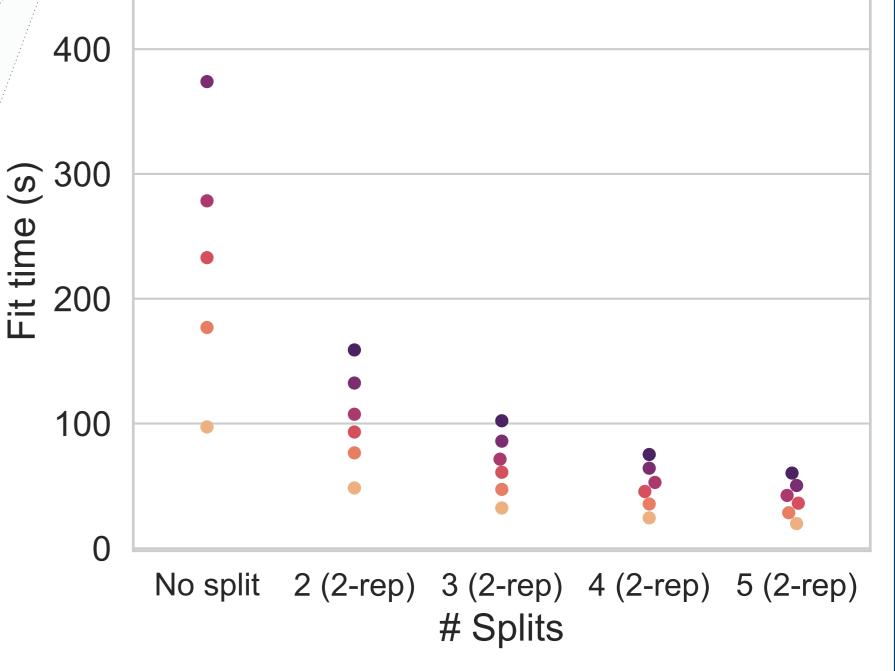
The 2-split configuration with at least two iterations as the optimal trade-off between training time and prediction accuracy when using GPR for brain age prediction.

No-split model, with 1000 features: total training time represented 42.1% of the training time of 2000 features, & 26.4% of the training time of using 3000 features. Similar time reductions were obtained within the split setups

The non-split, 2x2-splits and 2x3-splits models exhibited a less pronounced impact on MAE, except for 1000 features. The 2x4- and 2x5-splits models displayed a decrease in predictive performance for high number of features (3000 and 3500).

Low sample/feature ratio might cause a drop i predictive performance.





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