

Predicting Cardiovascular Disease Risk Using Functional Connectivity and Structural Morphology Metrics

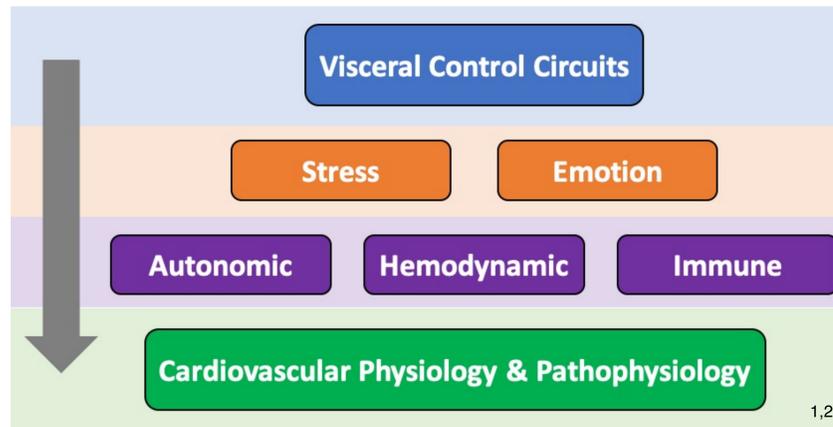
Amy Isabella Sentis^{1,2,3} (aisentis@cmu.edu), Javier Rasero⁵, Peter J. Gianaros^{3,4}, Timothy Verstynen^{3,5}

1. Medical Scientist Training Program, University of Pittsburgh & Carnegie Mellon University
2. Program in Neural Computation, Neuroscience Institute, Carnegie Mellon University
3. Center for the Neural Basis of Cognition, Carnegie Mellon University

4. Psychology Department, University of Pittsburgh
5. Psychology Department, Carnegie Mellon University



Visceral control circuits



- Neurocardiology^{3,4} focuses on identifying brain-based markers of CVD risk that can add to conventional clinical markers⁵.
- Hypothesis: a multimodal approach will generate a brain-based biomarker that reliably predicts a vascular marker of CVD risk, specifically carotid artery intima media thickness (IMT).

Methods

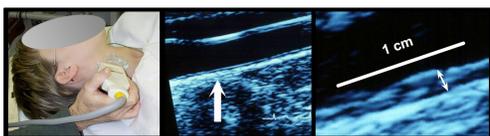
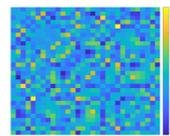
Participants:

- Neuroimaging and demographic data from 324 participants from the Pittsburgh Imaging Project were included in our analyses:
 - Ages 30-51 (mean: 40)
 - Female = 49%, male = 51%



Data:

- 3 Tesla whole-body scanner, 12-channel phased-array head coil
- T1-weighted MPRAGE acquisition:
 - 7 minute 17 second scan
 - FOV = 256x208mm, matrix size = 256x208, TR = 2100ms, inversion time = 1100ms, TE = 3.31ms, FA = 8° (192 slices, 1mm thickness, no gap)
 - cortical surface area, cortical thickness and subcortical volume extracted using FreeSurfer v6
- Resting-state functional connectivity (FC) acquisition:
 - 5 minute scan, eyes open
 - FOV = 205x205mm, matrix size = 64x64, TR = 2000ms, TE = 28ms, and FA = 90° (39 slices interleaved inferior-to-superior for each of 150 volumes, 3mm thickness, no gap)
 - Pearson correlation to estimate edges for FC
- IMT acquisition: Carotid artery mean IMT was measured bilaterally at distal common carotid artery, carotid artery bulb, and internal carotid artery then averaged



Methods

Multimodal stacking:

- Five-fold cross-validation prediction stacking algorithm, see Figure 1:
 - Step 1: train individual support vector regression models on each of the input data sources: resting-state FC, cortical surface area, cortical thickness, subcortical volumes
 - Result: four predictions of IMT for each participant, one from each data source
 - Step 2: stack SVR predictions as input into random forest model to give one final prediction for each participant

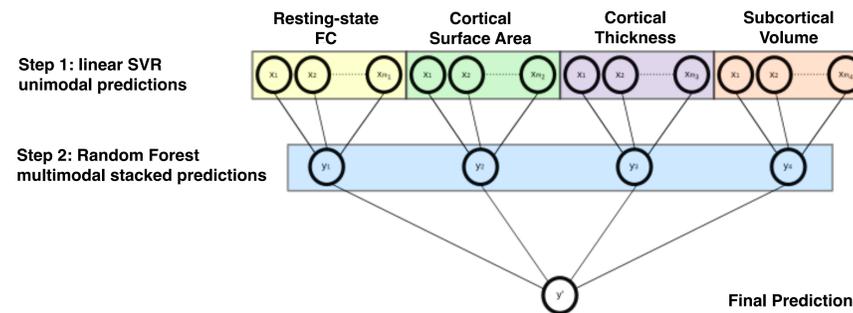


Figure 1: Prediction stacking model schematic, with linear SVR used in the unimodal predictions and random forest used in the multimodal prediction. FC = functional connectivity. SVR = support vector regression.

Results: data sources

Coefficient of determination for unimodal and multimodal model predictions

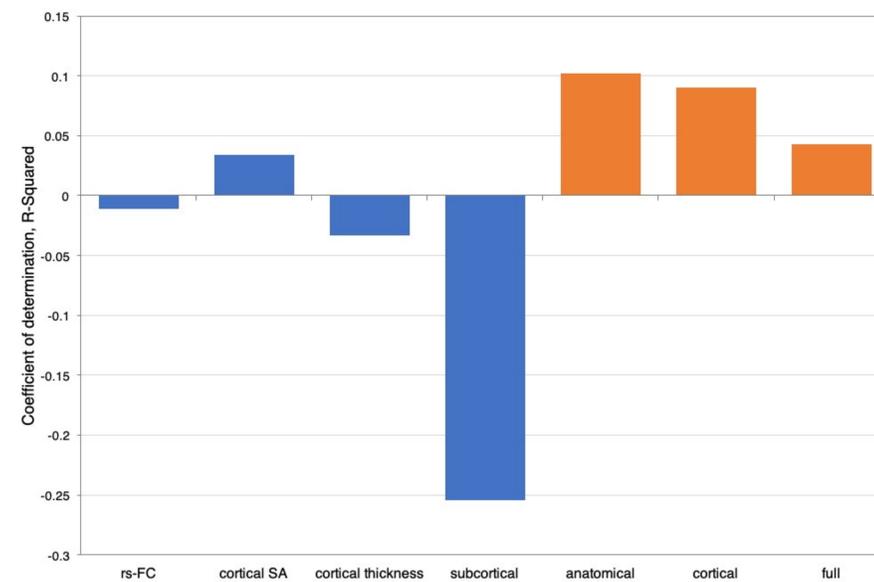


Figure 2: Coefficient of determination, r-squared, for single data sources from step 1 (blue) and best three candidate models from step 2 (orange): anatomical (cortical surface area, cortical thickness), cortical (resting-state FC, cortical surface area, cortical thickness), full (resting-state FC, cortical surface area, cortical thickness, subcortical volume). rs-FC = resting-state functional connectivity, SA = surface area.

Results: IMT

Correlation between observed IMT and predicted IMT

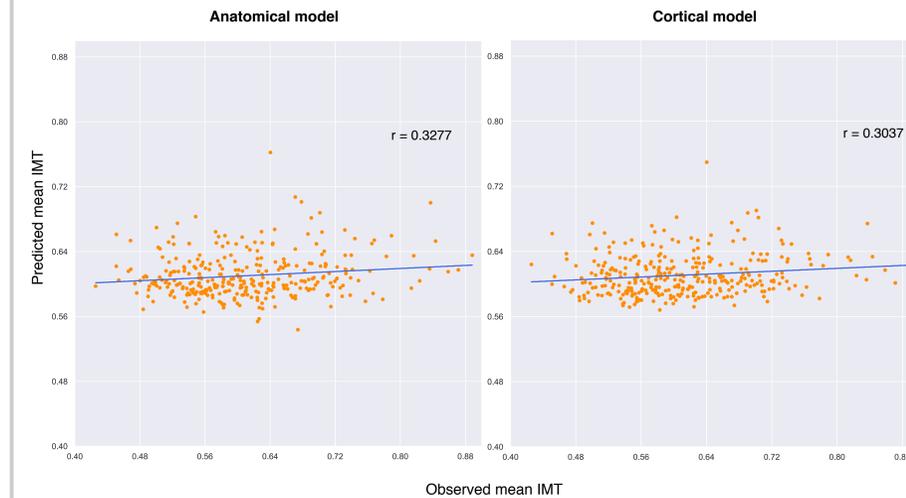


Figure 3: Correlation between observed and predicted mean IMT for the two best performing models: anatomical model, including cortical surface area and cortical thickness, R-squared = 0.1017, $p < 0.01$ (left), cortical model, including resting-state FC, cortical surface area and cortical thickness, R-squared = 0.0982, $p < 0.01$ (right).

Summary and future directions

Summary:

- Our anatomical and cortical models performed most strongly, accounting for 10.1% and 8.9% of the variance in mean IMT, respectively.
- This work builds on growing neuroimaging evidence by showing that functional and structural features of neural circuits may complement and add to the utility of conventional risk factors for predicting CVD.

Future Directions:

- We will perform interaction analyses using covariates such as gender and age to determine if these biomarkers behave differently in different demographic subsets.
- We will evaluate whether limiting resting-state FC features to those within the visceral control circuits that influence physiology will improve model performance.

References

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