Abstract Submission Form - PET is Wonderful 2020

Title The negative relationship between brain metabolism and its network dynamics: stability requires more energy

Max. 250 characters inc. spaces

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spaces

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Body of abstract (1500 characters inc. spaces)

The brain's metabolic consumption, imaged by 18F-FDG PET, is partially explained by the functional connectivity (FC) architecture emerging from resting state fMRI (rs-fMRI) studies. Correlations between 18F-FDG and static FC were reported, but the role of FC temporal variability has never been investigated.

Simultaneous 18F-FDG and rs-fMRI data were acquired in 28 healthy subjects (Riedl *et al., J.Neurosci.,* 2014; Aiello *et al., NeuroImage*, 2015), and standard uptake value relative (SUVR) to brain's global mean was extracted from 18F-FDG data.

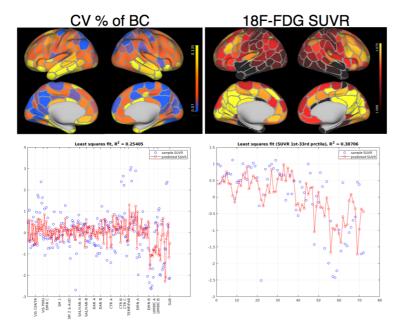
Sliding window time-varying FC was computed from rs-fMRI (window size: 30 TRs, step: 1 TR, TR=2s); strength (STR), betweenness centrality (BC), clustering coefficient (CC), local efficiency (LE) were calculated for each window and their node-wise coefficient of variation (CV%, i.e. temporal median absolute deviation divided by median) were computed.

Spearman's correlation and multiple linear regression were performed on group median values.

There were statistically significant anti-correlations between SUVR and CVs% of STR (rho = -0.33, p < 0.01), BC (rho = -0.42, p < 0.01), CC (rho = -0.19, p < 0.01), LE (rho = -0.25, p < 0.01), implying that the slower the FC dynamics, the higher the glucose consumption.

The multivariate model explained \sim 25% of the SUVR variance (R² = 0.25), with FC variability accounting for metabolic variance mainly in low SUVR nodes (R² = 0.39).

These findings require further investigation using 18F-FDG absolute quantification.



Top: Group-wise CV% of BC (*left*) and SUVR (*right*) sampled on the Schaefer atlas' cortical parcels: regions with high SUVR (*yellow*) tend to have low FC variability (*blue*).

Bottom: Agreement between SUVR (blue dots) and prediction (red line and dots) by multivariate model with CVs% as predictors for a) the whole brain (left) b) nodes belonging to the lower third of the SUVR distribution (right).

Figure legend max. 400 characters inc. spaces

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