

# Reduced Regional Integration in the Default Network of Patients with Very Mild Alzheimer's Disease Detected by Bootstrapping Rest-fMRI Data





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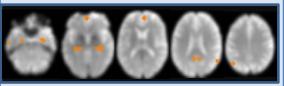
Alzheimer's Disease (AD) is the most prevalent cause of progressive dementia in older adults. Early Alzheimer's disease (AD) is associated with prominent changes of the default network (DN). AD has impact on inter-regional but also on intra-regional neuronal communication processes i.e. on brain integration at distinct levels. Based on the intuition that blood oxygenation level dependent (BOLD) signal variations of a given region might reflect functional aspects of regional integration, we proposed a novel marker BV(r) (Regional BOLD Variation) for the characterization of regional integration changes within the DN of patients with amnestic mild cognitive impairment (aMCI), a risk state for AD.

#### 1. Subjects and Data Acquisition

- resting state fMRI of 24 patients with amnestic MCI (aMCI) and 16 healthy controls
- subjects were instructed to keep their eyes closed, not to think of anything in particular, and not to fall asleep;

### 1. Selection of Regions of Interest

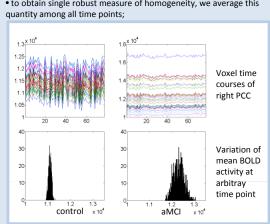
• 10 ROIs within the default network



- medial temporal gyri, hippocampi,
- left superior frontal gyrus, hippocampi.
- right superior frontal gyrus,
- · left and right posterior cingulate cortex, right superior marginal
- · left superior marginal gyrus;
- coordinates of ROI-centers derived from ICA (Sorg et al. 2007);

## 2. Evaluation of Regional BOLD Variation

- bootstrapping the BOLD activity allows detecting minor variations in regional integration;
- for each time point the mean BOLD activity is computed for a large number of samples with replacement;
- width of the 95% confidence interval of all bootstrapping samples is the base of our marker:
- to obtain single robust measure of homogeneity, we average this



# 3. Results · Significant group differences ( α = 95%, Mann-Whitney-U, Westfall-Young Correction) aMCI MTG R MTG R MTG R 00 PCC R · which can not be explained by grey matter loss; 0.4 VBM(PCC 0.35 • only slight correlation to grey 0.3 matter loss in hippocampi; 0.25 -> differences of functional type 300 400 100 200 BV(PCC R) • most pronounced difference in **BOLD Variation in** right posterior cingulate cortex is a good test to identify aMCI; (area under ROC curve 0.81) False positive rate (1-specificity)

#### 4. Conclusion

- regional integration in rest-fMRI captured by bootstrapping plays an important role for aMCI identification:
- significant regions within the DN correspond well to previous findings (Grecius et al., 2004,
- on the data of our study, our novel marker performs superior than previously proposed measures for regional integration (Li et al., 2002, Zang et al., 2004, Xu et al., 2008);
- comparison with VBM demonstrates that the differences are functional in nature;
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