

^{11}C -PBR28 [3] ($R^2 = 0.93$). Whilst caudate cannot be assumed to be a true reference region, it has been previously shown to have low displaceable fraction of TSPO in rats [5]. In order to explore the association between specific binding and genetics independent of plasma variability, we estimated a pseudo binding potential (BP_{ref}) [4] using grey matter masked caudate as a reference region. The genotypic influence was not observed on the V_T , but was seen on BP_{ref} and particularly evident in cortical regions.

Conclusions: Binding of ^{18}F -PBR111 can be quantified with a 2TC model in healthy subjects. When normal-

ized by a pseudo reference region, there was evidence that the cortical binding was influenced by the status of the rs6971 polymorphism in the TSPO gene.

References

- [1] Owen et al. *J Cereb Blood Flow Metab*, Vol. 32, 2012.
- [2] Owen et al. *J Nucl Med*, Vol. 52, 2011.
- [3] Fujita et al. *NeuroImage*, Vol. 40, 2008.
- [4] Gunn et al. *Synapse*, Vol. 65, 2011.
- [5] Jones et al. *WMIC Meeting*, 2011.

P164. Investigation of noise-induced correlations in dual-biomarker parametric imaging from dynamic [^{11}C]PiB PET

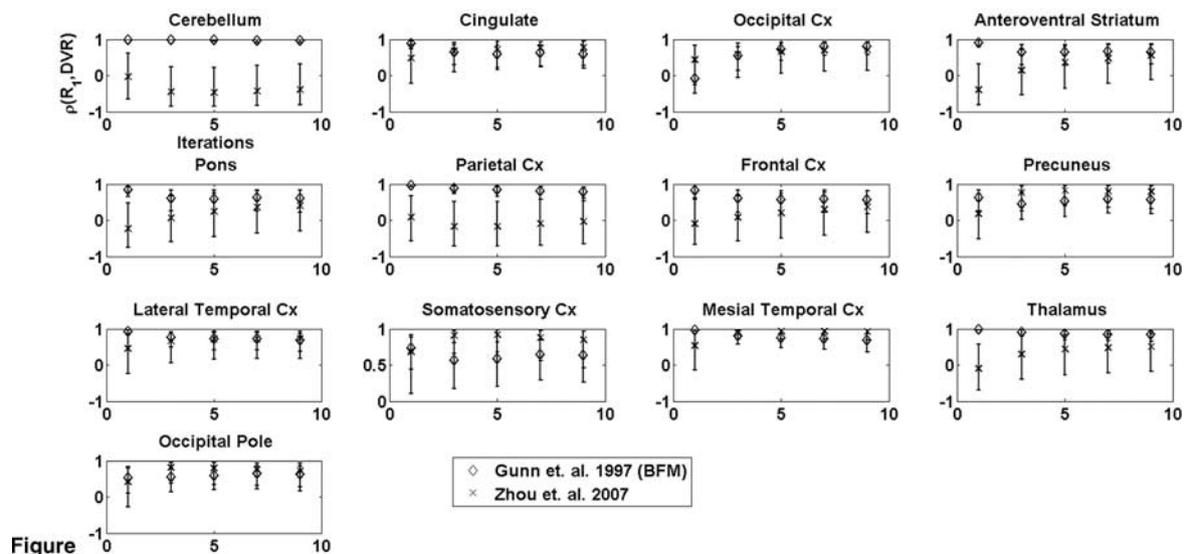
Hassan Mohy-Ud-Din¹, Nicolas A. Karakatsanis¹, Julie C. Price², Yun Zhou¹, Susan M. Resnick³, Christopher J. Endres¹, William E. Klunk², Chester A. Mathis², Dean F. Wong¹ and Arman Rahmim¹

¹Johns Hopkins University, Baltimore, MD, USA; ²University of Pittsburgh, Pittsburgh, PA, USA; ³Intramural Research Program, National Institute on Aging, NIH, Baltimore, MD, USA

Background: To investigate the impact of noise on correlations between R_1 (relative flow) and BP (binding potential) parametric images as obtained from dynamic [^{11}C]PiB PET. **Motivation and Significance:** Generation of R_1 and BP parametric images from dynamic [^{11}C]PiB PET has been proposed as an approach for dual-biomarker imaging of dementia from a single acquisition. This approach has also been considered to evaluate correlations between flow and β -amyloid depositions. We demonstrate that noise-induced correlations pose significant confounds to interpretation.

Methods: For validation, kinetic parameters (K_1 , k_2 , k_3 , k_4) from [^{11}C]PiB studies of normal controls were estimated, averaged and assigned to a mathematical brain phantom, from which time-activity-curves (TACs) were generated using a 2-tissue compart-

mental model (blood-volume fraction was set to 3%). This was followed by realistic simulations of dynamic frames for the geometry of the HRRT scanner (including 20 noise realizations in the sinogram-space of the same subject; i.e. simulating no biological correlations). OSEM reconstructions (1-10 iterations) for each dataset were generated. Parametric Images of R_1 and DVR (=BP+1) were obtained via the Simplified Reference Tissue Model (SRTM) using (1) the Basis Function Method (BFM) of Gunn et al. (1997), and (2) linearized formulation and regression (LR) of Zhou et al. (2007) to estimate the parameters. Noise-Bias trade-off curves were obtained for R_1 and DVR images, demonstrating reduced bias with increasing iterations at the cost of enhanced noise levels, as expected. Next, Pearson



Figure

correlation coefficients were determined for each voxel-pair of (R_1 , DVR) vectors (across the 20 noise realizations) with lower and upper bounds for a 95% confidence interval, and quantitatively evaluated for each OSEM reconstruction. A hypothesis test for no correlation with a probability value (p-value) of 0.05 was also conducted, and the resulting correlation images were analyzed qualitatively and quantitatively for all OSEM iterations.

Results: Specifically, 13 regions-of-interest (ROIs) were considered. Overall, R_1 and DVR showed statistically significant correlations, across the OSEM iterations, for all ROIs using the BFM approach, while this was also

the case for the LR method with the exception of the parietal cortex, pons and occipital pole.

Conclusions: Significant correlations attributed purely to noise were observed between R_1 and DVR parametric images in parametric quantification of dynamic [^{11}C]PiB PET. Caution should be exercised when performing R_1 and DVR analysis, where the detection of biological correlations may be confounded by noise-induced correlations. Future work should explore potential approaches that quantify and account for these associations to provide more accurate estimates of correlations between DVR and R_1 estimates from a single acquisition.

P165. *In vivo* amyloid deposition in the aging brain: methodological considerations for partial volume correction

Olivier Rousset¹, Pierre-Louis Bazin², Aaron Carass³, Christopher Endres⁴, Amith Harsha⁴, Dzung Pham⁵, Susan Resnick⁶ and Dean F. Wong¹

¹Department of Radiology and Radiological Sciences, Division of Nuclear Medicine, Section of High Resolution Brain PET imaging, Johns Hopkins University, Baltimore, MD, USA; ²Max Planck Institute for Human and Cognitive Brain Sciences, Leipzig, Germany; ³Image Analysis and Communication Lab, Johns Hopkins University, Baltimore, MD, USA; ⁴Department of Radiology and Radiological Sciences, Division of Neuroradiology, Johns Hopkins University, Baltimore, MD, USA; ⁵Center for Neuroscience and Regenerative Medicine, Bethesda, MD, USA; ⁶Gerontology Research Center, National Institute on Aging, National Institutes of Health, Baltimore, MD, USA

Background: PET imaging with amyloid tracers such as [^{11}C]PiB is contributing to differential diagnosis

of Alzheimer's disease (AD) by demonstrating the presence of beta-amyloid plaques. While partial

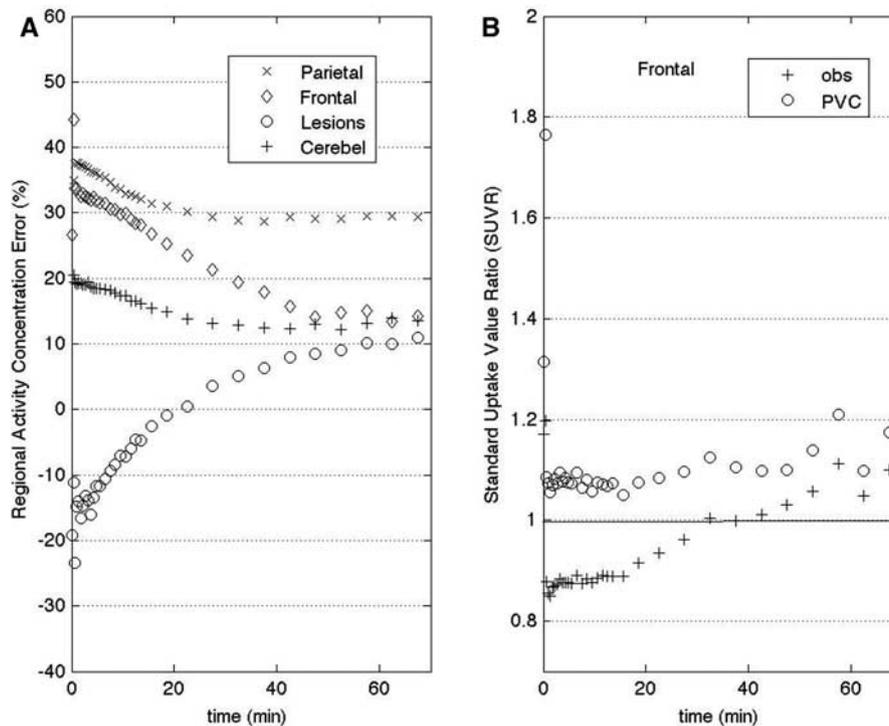


Figure (A) %Error in regional activity concentration in selected regions, and (B) Standard Uptake Value Ratio (cerebellar grey taken as reference region) in a typical PiB study, in the frontal cortex before (+) and after PVC (o).