

Effect of blood flow changes on quantification of [¹⁸F]flutemetamol studies

Fiona Heeman¹, Maqsood Yaqub¹, Isadora Lopes Alves¹, Kerstin Heurling², Juan Domingo Gispert³, Christopher Foley⁴, Ronald Boellaard¹, Adriaan A. Lammertsma¹, on behalf of the AMYPAD Consortium

¹Department of Radiology & Nuclear Medicine, Amsterdam Neuroscience, VU University Medical Center ²Wallenberg Centre for Molecular and Translational Medicine and the Department of Psychiatry and Neurochemistry, University of Gothenburg ³Barcelonaβeta Brain Research Center, Pasqual Maragall Foundation ⁴GE Healthcare

Introduction

- Cerebral blood flow (CBF) decreases with age and additional reductions occur in patients suffering from Alzheimer's Disease (AD)
- Reductions in CBF of up to 30% have been reported^{1,2}
- Longitudinal assessment of amyloid load may be biased because of sensitivity to changes in CBF³

Purpose

The purpose of this simulation study was to assess the effect of CBF reductions on quantification of [¹⁸F]flutemetamol scans

Methods

- Clinical input⁴
 - Kinetic parameters for a range of amyloid load levels, based on dynamic 110 minutes time activity curves (TACs) validated previously (Table 1)
- Simulated CBF reductions
 - Tracer delivery reduction K_1 (0 to -25%, while keeping K_1/k_2 ratio constant)
 - Global: reduction in cortical (target) and cerebellar grey matter region (reference) (Fig.1)
 - Local: reduction in cortical region only
- Amyloid load estimation
 - SUVR (90-110 minutes post injection)
 - Simplified reference tissue model⁵: non displaceable binding potential (BP_{ND}) and distribution volume ratio ($DVR = BP_{ND} + 1$)

Table 1 Pharmacokinetic parameters for simulating plasma TACs using reversible two-tissue compartment model

Amyloid level	BP_{ND}	DVR	k_3	V_B
I	0.380	1.020	0.008	0.05
II	0.635	1.208	0.013	0.05
III	0.890	1.397	0.018	0.05
IV	1.145	1.585	0.023	0.05
V	1.400	1.774	0.028	0.05
Reference	0.350	1.000	0.018	0.05

BP = Binding potential of target tissue, DVR = Distribution Volume Ratio. [¹⁸F]flutemetamol target tissue: $K_1=0.248$, $k_2=0.08$, reference tissue: $K_1=0.32$, $k_2=0.103$.

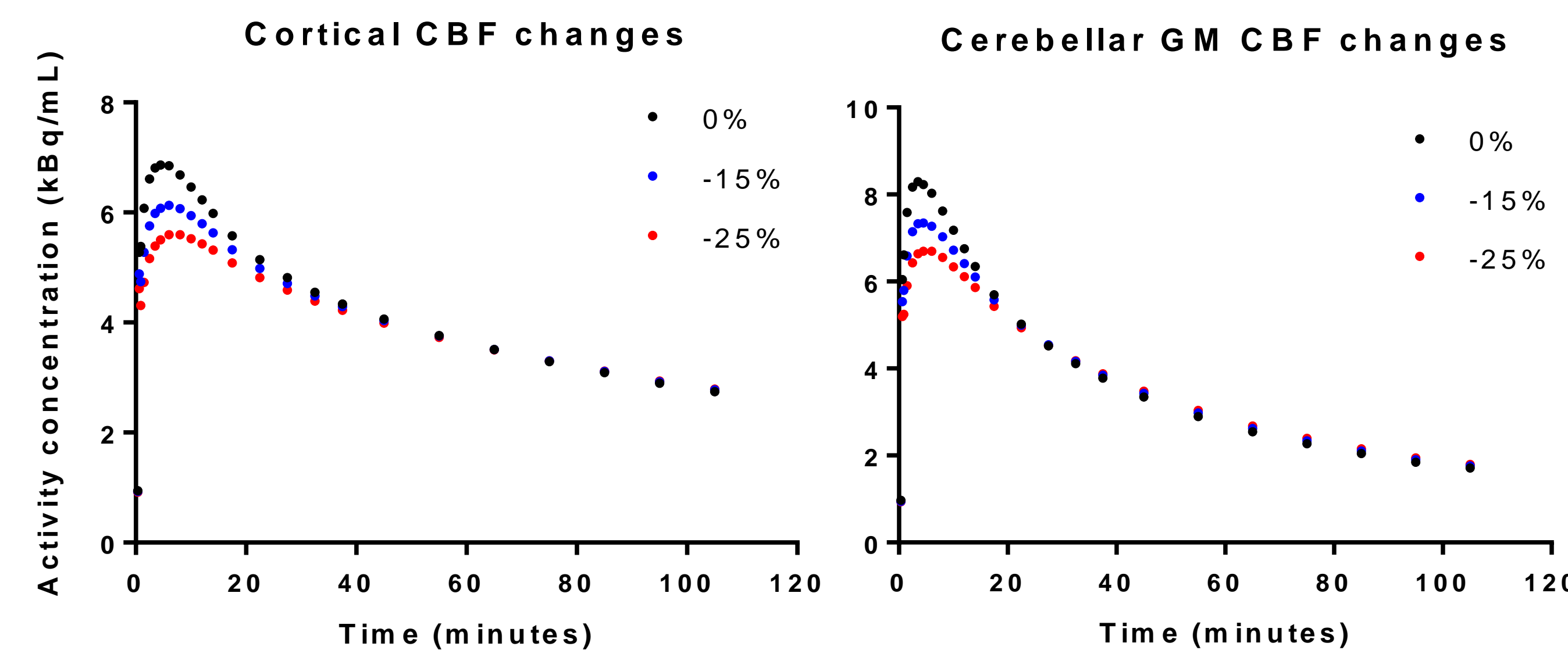


Figure 1. Example TACs showing reductions in tracer delivery (K_1) for a) the cortical region and b) the cerebellar grey matter region. K_1/k_2 ratio was kept constant for both regions.

Results (1)

- Large global CBF reductions resulted in only a small bias in DVR (<3%) for all simulated amyloid levels (Fig. 2a)
- Bias in SUVR was larger (4-16%) and affected by the level of amyloid load (Fig. 2b)

Global CBF change

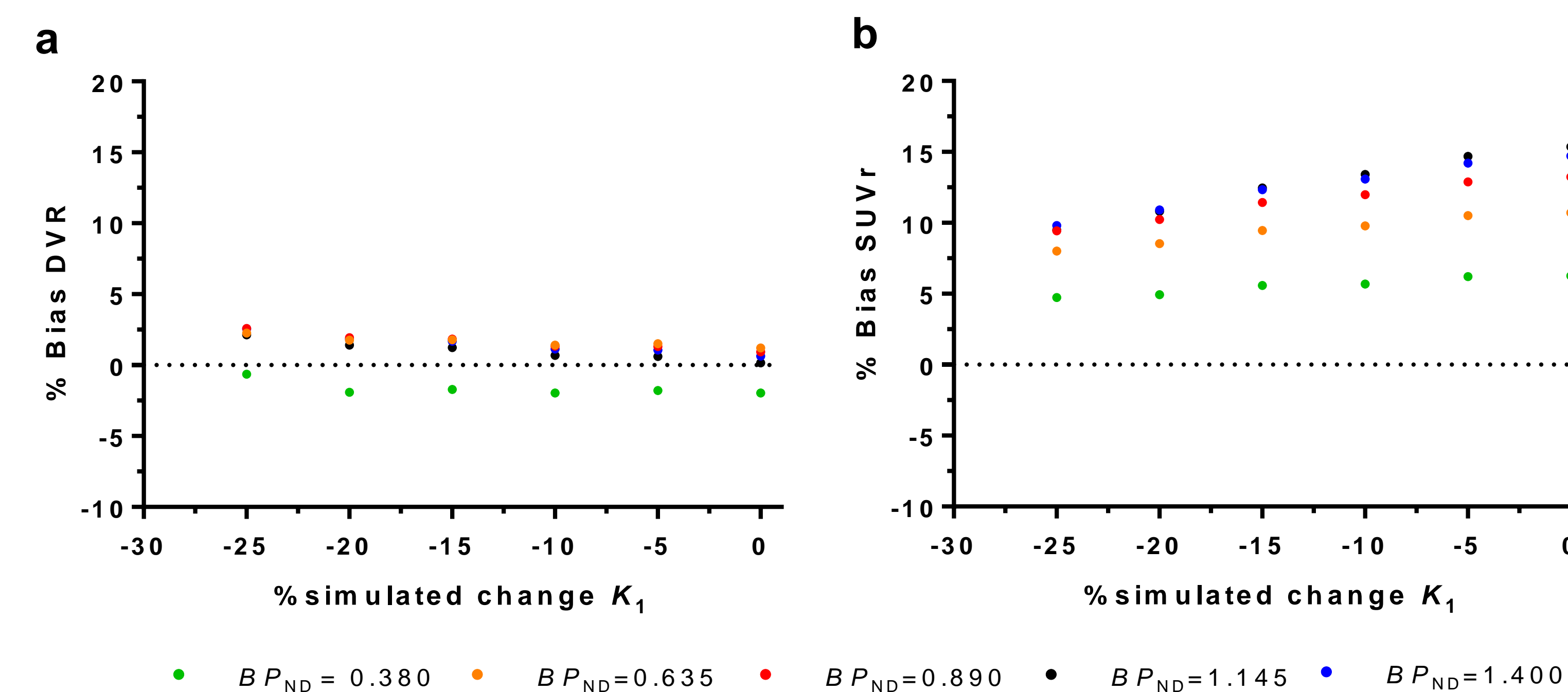


Figure 2. a) Percentage bias in DVR and b) SUVR, as a function of global CBF (K_1) for various levels of amyloid load (BP_{ND}).

Results (2)

- Bias in DVR increased slightly for large local CBF reductions (Fig. 3a)
- A local CBF decline resulted in a relatively stable bias in SUVR, however strongly affected by level of amyloid load (Fig. 3b)

Local CBF change

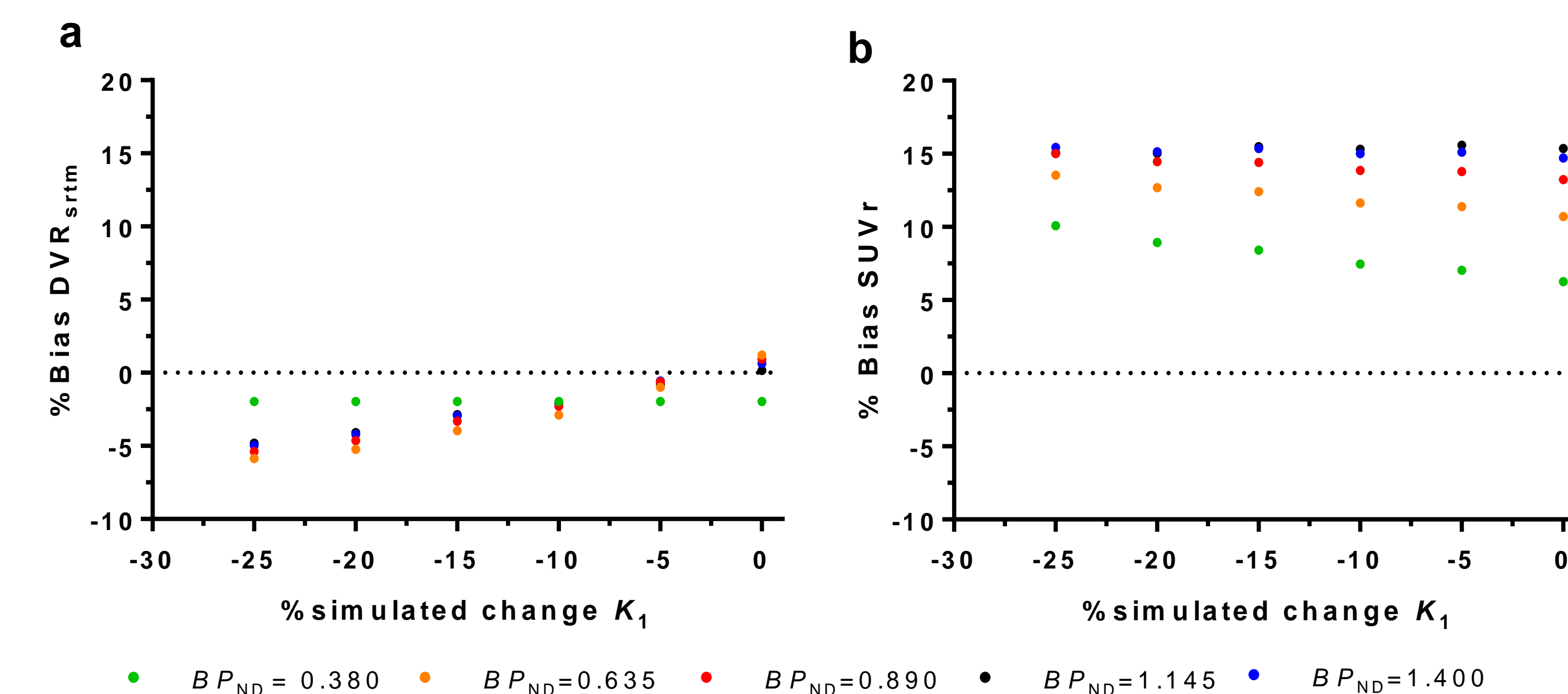


Figure 3. a) Percentage bias in DVR and b) SUVR, as a function of local CBF (K_1) for various levels of amyloid load (BP_{ND}).

Conclusion

- ✓ Changes in derived amyloid load may be affected by reductions in CBF when using SUVR. Hence, for more accurate amyloid load estimates, especially in a longitudinal setting, dynamic scans should be used.

1. Frackowiak (1986) *Prog Brain Res.* 1986;70:69–85, 2. Bremner et al. (2010) *Molecular Imaging and Biology*;13: 759–68. 3. van Berckel et al. *J Nucl Med* 54.9 (2013): 1570-1576. 4. Heurling et al. (2015) *Neuroimage* 121: 184-192, 5. Lammertsma & Hume (1996) *Neuroimage* 4.3 (1996): 153-158.

Contact: f.heeman@vumc.nl