## Meeting abstract

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**Metabolism of (R)-[<sup>11</sup>C]verapamil in epilepsy patients** Aiman Abrahim<sup>1,5</sup>, Gert Luurtsema<sup>6</sup>, Martin Bauer<sup>1</sup>, Rudolf Karch<sup>2</sup>, Christoph Baumgartner<sup>3</sup>, Kurt Kletter<sup>4</sup>, Markus Müller<sup>1</sup> and Oliver Langer<sup>\*1,5</sup>

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from 13th Scientific Symposium of the Austrian Pharmacological Society (APHAR). Joint Meeting with the Austrian Society of Toxicology (ASTOX) and the Hungarian Society for Experimental and Clinical Pharmacology (MFT) Vienna, Austria. 22–24 November 2007

Published: 14 November 2007 BMC Pharmacology 2007, **7**(Suppl 2):A23 doi:10.1186/1471-2210-7-S2-A23

This abstract is available from: http://www.biomedcentral.com/1471-2210/7/S2/A23

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### Introduction

(*R*)-[<sup>11</sup>C]Verapamil (VPM) is a new positron emission tomography (PET) tracer to measure P-glycoprotein (Pgp)-mediated transport at the blood-brain barrier (BBB). Owing to the lack of a suitable reference region in brain that is devoid of P-gp, a metabolite-corrected arterial input function is required for quantitative analysis of VPM PET data [1]. The aim of this study was to compare metabolism of VPM in epilepsy patients and healthy volunteers.

### Methods

Selected arterial blood samples from 9 patients, who underwent VPM PET, were analyzed for radiolabeled metabolites by a previously described combined solidphase extraction/HPLC assay [2].

### Results

VPM metabolism was significantly faster in patients as compared to healthy volunteers [1] (unchanged VPM at 60 min after injection:  $26.1 \pm 6.4$  vs.  $49.0 \pm 13.4\%$ , p < 0.05, *t*-test).

### Conclusion

Faster metabolism of VPM in epilepsy patients may be caused by CyP450 enzyme induction by antiepileptic drugs. Based on these data caution is warranted when using an averaged arterial input function derived from healthy volunteers for the analysis of patient data. As radiolabeled metabolites of VPM are known to cross the BBB [1], different kinetic modeling parameters obtained in patients and healthy volunteers might be at least partly attributed to different rates of tracer metabolism rather than to differences in cerebral P-gp activity.

#### References

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