The applicability of SRTM in [¹⁸F]fallypride PET investigations: impact of scan durations

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The high-affinity radioligand [¹⁸F]fallypride (FP) is frequently used for quantification of striatal/ extrastriatal D_{2/3} receptors and the receptor occupancies of antipsychotics (APs). Its 110 minutes half-life allows long scan durations. However, the optimum scan duration is a matter of debate. This investigation focuses on scan-duration-related effects on simplified reference tissue model (SRTM) results and the time point of transient equilibrium in a large sample of dynamic FP positron emission tomography (PET) scans. Fifty drug-free and 50 AP-treated subjects underwent FP-PET scans (180 minutes scan duration). The binding potential (BP_{ND}) of the putamen, thalamus, and temporal cortex were calculated using the SRTM and the transient equilibrium model. Furthermore, receptor occupancies were calculated for AP-treated patients. Transient equilibrium in the unblocked putamen occurred after 121 ± 29.6 minutes. The transient equilibrium occurred much earlier in the extrastriatal regions or under AP treatment. Stepwise scan shortening caused BP_{ND} underestimations of 0.58% for the first 10-minute reduction (putamen, SRTM), finally reaching 5.76% after 1 hour scan-time reduction. We observed preferential extrastriatal AP binding irrespective of the analytical method. [18F]fallypride scan durations of 180 minutes reliably reach equilibrium even in D_{2/3}-receptor-rich regions. Moderate reductions in FP scan durations only caused small changes to SRTM results even in receptor-rich regions. Apparently, the D_{2/3} receptor occupancy results of APs, especially preferential extrastriatal binding observations, are not relevantly biased by inappropriate scan durations.

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Introduction

The high-affinity ligand [¹⁸F]fallypride (FP) is one of the most frequently used ligands for $D_{2/3}$ receptor imaging (Mukherjee *et al*, 1995; Vernaleken *et al*, 2008*a*). Because of its high affinity and long half-life, this ligand has proven valuable for the quantification of striatal and extrastriatal $D_{2/3}$ receptors during one

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single positron emission tomography (PET) scan. However, for the use in clinical studies, some more properties should be considered. Shorter, more practical scan durations and the applicability of noninvasive reference region methods can increase the patients' acceptance of such scientific investigations and prevent relevant bias in patient inclusion. The simplified reference tissue model (SRTM; Lammertsma and Hume, 1996) is frequently used in FP-PET investigations. It primarily depends on the existence of a suitable reference region devoid of $D_{2/3}$ receptors. Although there have been some evidence of very low cerebellar $D_{2/3}$ receptor concentrations, this structure appears to be the most appropriate reference region. Kinetic constraints for the application of the SRTM

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as expressed by Buchert and Thiele (2008) can be considered to be fulfilled on the basis of the kinetic parameters reported by Siessmeier *et al* (2005). An early comparison between the two-tissue compartment model (2TCM) and two reference region methods (SRTM, and the transient equilibrium method; TEM) (Farde *et al*, 1989; Ito *et al*, 1998) showed high and statistical significant intermethod correlations (Siessmeier *et al*, 2005).

As depicted earlier, it is of major interest to find the best compromise between practicality and model-related constraints. A systematic evaluation of scan-time effects on the outcome parameters of FP-PET scans has never been performed. Therefore, we conducted a *post hoc* analysis including 100 subjects, of which 50 were under antipsychotic (AP) medication and 50 were free of any neuropsychiatric medication. In respect to medication-free subjects, our study had three goals. (1) We searched for the existence and consequently for the corresponding time point of estimated transient equilibrium conditions. (2) The impact of different scan durations on the SRTM results was depicted by artificial and stepwise removal of later time frame data.

The subgroup analysis of 50 AP-treated patients focuses on the impact of scan durations on $D_{2/3}$ receptor occupancy results. In several studies, FP-PET was used to compare striatal and extrastriatal $D_{2/3}$ receptor occupancies caused by second-generation antipsychotics (Vernaleken *et al*, 2008*a*). However, it has been hypothesized that inaccurate estimation of binding potentials (BP_{ND}) in regions with high $D_{2/3}$ receptor densities (caused by insufficient scan durations) can potentially influence some of the conclusions (e.g., preferential extrastriatal binding) drawn from these results (Olsson and Farde, 2001). Therefore, we also assessed the impact of scan duration on the misinterpretation of AP-induced $D_{2/3}$ receptor occupancies.

Materials and methods

This investigation is a *post hoc* analysis of several PET studies using FP as the radioligand. Some patient and control subject data are part of previously published reports about the extent of $D_{2/3}$ receptor occupancies under AP treatment with clozapine, ziprasidone, and quetiapine (Gründer *et al*, 2006; Vernaleken *et al*, 2008b, 2010). All clinical studies were approved by the ethics committee of the medical faculty of the RWTH Aachen University (Aachen, Germany), the ethics committee of the medical chamber of Rheinland-Pfalz (Mainz, Germany), and the German radiation safety authority.

Subjects

Subjects included healthy volunteers (n=27) and patients with various neuropsychiatric disorders (n=73). All subjects participating in the study gave written informed consent. They all underwent physical and mental state

examinations. Patients suffering from neurologic or relevant somatic diseases were not included, nor were patients reporting illegal drug consumption in the last 6 months, except two patients with Tourette's syndrome who ingested THC-containing products up to 6 weeks before the PET scan. Drug screening was performed at the beginning of the inpatient stay and 1 day before the scan. All PET studies were performed at the PET centers of the University of Mainz and RWTH Aachen University, Germany.

A total of 100 subjects were included (age: 18 to 64 years, mean \pm s.d.: 33.4 \pm 10.4; 78 \Diamond + 22 \heartsuit). At the time of the PET scan, 50 subjects were free of any centrally acting substances and drugs directly acting on dopaminergic systems for at least 6 months (age: 18 to 58 years, mean \pm s.d.: 31.0 ± 10.5 ; 40 δ + 10 \Im). This unblocked group consisted of 21 patients suffering from schizophrenia (according to the Diagnostic and Statistical Manual of Mental Disorders (DSM-IV); 10 subjects were drug naive), two patients suffering from Tourette's syndrome (scanned under propofol anesthesia) (Vernaleken et al, 2009), and 27 healthy subjects. The remaining 50 subjects (age: 18 to 64 years, mean \pm s.d.: 35.7 \pm 9.9; 38 \Diamond + 12 \heartsuit) were patients undergoing stable AP treatment (100 to 500 mg clozapine, 80 to 200 mg ziprasidone, 200 to 1,300 mg quetiapine, or 5 to 20 mg olanzapine). In this group, the following variable diagnoses were made: schizophrenia (n=38), schizoaffective disorder (n = 9), transient psychosis (n = 2), cyclothymic disorder (n = 1), and personality disorder (n = 1).

Radiochemistry

[¹⁸F]Fallypride was synthesized using a high-yield modification of the method for [¹⁸F]DMFP synthesis (Hamacher *et al*, 1986; Gründer *et al*, 2003) by using the tosylated precursor ((*S*)-*N*-[(1-allyl)-2-pyrrolidinyl)methyl]-5-(3-toluenesulfonyloxy-propyl)-2-ethoxybenzamide. Quality control before injection included determination of chemical and radiochemical purity, specific activity, pH, and absence of pyrogens.

Positron Emission Tomography Data Acquisition

Images were acquired on a Siemens ECAT EXACT wholebody PET scanner in 3D mode (field-of-view, 16.2 cm; 47 planes; full width at half maximum axial, 6.0 mm; in-plane, 6.0 mm). Data acquisition consisted of a series of 39 time frames $(3 \times 20 \text{ seconds}, 3 \times 1, 3 \times 2, 3 \times 3, 21 \times 5, 2 \times 8, \text{ and}$ 4×10 minutes), resulting in a total scanning time of 180 minutes. A 15-minute transmission scan using a ⁶⁸Ge source was performed before each study for subsequent attenuation correction. Head movement was minimized by means of a vacuum mask. A mean of $218 \pm 33.9 \text{ MBq}$ (mean ± s.d.; range: 101 to 264 MBq) FP was injected intravenously as a bolus into a cubital vein over ~ 30 seconds (unmedicated subjects: 215 ± 28.6 MBq, range: 149 to 251 MBq; treated patients: 221 ± 38.7, range: 101 to 264 MBq). There was no statistical significant difference between the two groups (Δ : 6 MBq; P = 0.45; unpaired two-sided *t*-test). The average specific activity at the time point of injection was $316 \pm 825 \, \text{GBq}/\mu \text{mol}$. There was no correlation between binding potentials and specific activities (medication-free subjects), irrespective of the volume of interest (VOI) analyzed (putamen, r=0.137; thalamus, r=0.231; inferior temporal cortex (GTi), r=0.146; Pearson's correlation).

Image Processing

Images were reconstructed with filtered back-projection using a Hanning filter (filter width: 4 mm). Although in most scans only moderate head movement was observed, a motion correction was performed (Flexible Image Matching and Fusion Tool (PFUS) implemented in PMOD V3.1, Zürich, Switzerland) using a sequential registration procedure beginning with frame 10 while assuming no relevant dislocation during the first 10 minutes. To check the quantity of misregistration, we checked the position of striatal VOI as well as the appropriate match of the brain surface contours by visual inspection. Rare misregistrations were corrected by manual registration. Thirteen scans had to be interrupted for two 5-minute frames mainly because of the patient's desire to urinate. Repositioning was performed using three marks, indicating the previous head position; this was followed by registration of the postinterruption frames to the previous frames. For spatial normalization, the early frames (usually frames 3 to 8) of the dynamics were averaged and registered to an Montreal Neurological Institute (MNI) template using the warp transformation process of the Automated Image Registration package (Woods et al, 1992) (implemented in MEDx software v3.43; Medical Numerics, Germantown, MD, USA); the corresponding registration parameters were then applied to the entire dynamic scan. To extract the timeactivity curves (TACs), templates of polygonal VOIs were applied to the spatially normalized dynamic recordings. This VOI template was chosen in several previous FP-PET studies (Vernaleken et al, 2008b, 2009, 2010); it focuses on regions showing a sufficient extent of $D_{2/3}$ receptor availability for FP binding and stable statistics. Timeactivity curves were calculated for the following VOIs: the cerebellum, putamen, GTi, and thalamus. Volumes of interest for the caudate nucleus, amygdalae, and other cortical and midbrain regions were also calculated in the original studies. However, these VOIs were omitted in the present post hoc analysis because the present choice of VOIs focuses on the representation of brain regions with high, moderate, and low $D_{2/3}$ receptor densities.

Data Analysis

The general quantification procedures followed the SRTM as described by Lammertsma and Hume (1996). For the calculation of $BP_{\rm ND}$ values by the SRTM, we used the unfitted TACs. The TEM (Farde *et al*, 1989) was applied under the following assumptions: the tissue radioligand activity in the nondisplaceable compartment is the same in the reference and receptor-containing tissues and the tissue radioligand activity bound to the receptors $C_{\rm S}(t)$ equals the difference of $C_{\rm T}(t)-C_{\rm T}(t)$ ($C_{\rm T}$ is the tissue radioligand activity in the receptor-containing region of interest and $C_{\rm T}$ is the tissue radioligand activity of [¹⁸F]FP in the reference tissue (cerebellum)). When $C_{\rm S}(t)$ is plotted and a curve is fitted to these data, $dC_{\rm S}(t)/dt = 0$ at the peak of this curve (the point of 'peak equilibrium'); thus, $C_{\rm S}(t)/C_{\rm T}(t) = BP_{\rm ND} = k_3/k_4$. The TEM provides the time point of the transient equilibrium, which is one of the main variables for this *post hoc* analysis.

For the TEM, the TACs of the cerebellum (as a reference region), putamen, thalamus, and GTi were fitted to predefined functions. The putamen TACs were fitted to a standard Höhrl function $(f(x) = A \times (B^X) \times X^C)$. The TACs of the thalamus, GTi, and cerebellum were fitted to another exponential function: $(f(x) = A \times EXP \quad (B/X + C \times X) +$ $D \times \ln(X)$). These equations provide suitable approximations of the TACs for blocked and unblocked conditions. In only a few cases, the fits of very early time points (<4 minutes) yielded implausible results. However, this did not disrupt the assessment of equilibrium. Parameter approximation for TAC fitting was performed by LabFit software V7.2.42 (Silva and Silva, 1999–2008). The fitted cerebellum curve was subtracted from the corresponding individual target-VOI curves. Afterwards, the local maximum of this curve in the limits of 0 < x < 180 was determined. For TACs that showed no maximum during the 3-hour PET scan, the $C_{\rm S}(t)/C'_{\rm T}(t)$ ratio was determined at t = 180 minutes (n = 3).

To visualize characteristic TAC activity, data were transformed to standard uptake values (SUV = C_T [Bq/g] × BW [kg]/A [MBq] to account for body weight (BW) and injected activity (A).

Statistics

The $BP_{\rm ND}$ values obtained from different analytical methods were compared using the paired two-sided *t*-test. In all analyses, the two-tailed level of statistical significance was set at P = 0.05. Pearson's correlations and linear regression analysis were performed using SPSS software (v14.0; SPSS Inc., Chicago, IL, USA), to characterize the possible impact of the analytical method on $BP_{\rm ND}$ values.

Simulations of reduced scan durations (60 minutes reduction in 10 minutes intervals) were performed on all putamen and thalamus TACs of unblocked subjects. These SRTM results were compared with the individual 180-minute data.

To detect possible influences on receptor occupancy values, the extent of $D_{2/3}$ receptor binding under AP treatment was calculated according to the following equation:

$$ext{Occupancy}[\%] = \left(1 - \frac{BP_{ ext{Drug}}}{BP_{ ext{Control}}}\right) imes 100$$

The 50 medication-free subjects (age matched) served as a control group; we did not perform intraindividual pre/post analyses.

Results

The estimated time of equilibrium from averaged SUV curves was 120.6 ± 29.6 (76.1 to > 180) minutes (mean \pm s.d. (range)) in the putamen (Figure 1) as

| Scan | duration | of [1 | ⁸ F]fal | lypride | scans |
|--------|------------|-------|--------------------|---------|-------|
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well as 29.2 ± 8.8 (14.6 to 54.7) and 29.4 ± 14.9 (7.7 to 46.1) minutes (mean \pm s.d. (range)) in the thalamus and GTi, respectively. The corresponding time points under AP treatment were in the putamen 77.7 \pm 31.1 (29.5 to 167.2) minutes, in the thalamus 18.4 \pm 9.3 (3.8 to 39.7) minutes, and in the GTi 28.7 \pm 15.9 (6.7 to 67.0) minutes (mean \pm s.d. (range)). The fits of the specific binding curves for the putamen failed to reach a maximum in only two cases of medication-free subjects; their estimated time points of transient equilibrium (calculated by extrapolation of the fitted TACs) were 190 and 205 minutes after tracer injection in the putamen.

The mean $BP_{\rm ND}$ values (SRTM) of medication-free subjects ranged were the following: 22.2 ± 4.9 in the putamen, 2.11 ± 0.57 in the thalamus, and 0.77 ± 0.30 in the GTi (mean \pm s.d.). The TEM results were slightly but significantly different in the putamen



Figure 1 Averaged standard uptake value (SUV) curves of the putamen volume of interest (VOI) (unmedicated subjects: n = 50; red, mean total activity; blue, mean specifically bound activity) and mean cerebellar VOI (green). The average of estimated time points of equilibrium was 121 minutes after injection. The SUV curves (specifically bound) representing the 75% and 25% percentiles of the unmedicated subgroup (borders of highlighted confidence interval; green area) show their maximum before the 180-minute limit.

and thalamus but not in the GTi (for detailed information see Table 1). The $BP_{\rm ND}$ values calculated with the TEM and SRTM corresponded considerably (linear regression analysis for putamen: slope = 0.852, $r^2 = 0.80$, P < 0.0001; thalamus: slope = 0.965, $r^2 = 0.94$, P < 0.0001; GTi: slope = 0.841, $r^2 = 0.87$, P < 0.0001). The comparison of SRTM and TEM results have been performed by a Bland–Altman plot; it depicts higher relative intermethod differences at lower binding potentials (Figure 2). We observed a negative correlation between age and $BP_{\rm ND}$ values of unmedicated subjects (putamen:



Figure 2 A comparison of binding potential (BP_{ND}) values according to the simplified reference tissue model (SRTM) and transient equilibrium method (TEM) for 100 subjects (50 second-generation antipsychotic (SGA)-treated patients and 50 drug-free subjects) and for all regions (the putamen, thalamus, and inferior temporal cortex (GTi); observations: n = 300). Data are presented as Bland–Altman plots, showing the individual mean values (TEM and SRTM) on the abscissa as well as the individual ratios of SRTM/TEM on the ordinate. Because of the large range of outcome values (approximately a factor of 10²), the plot is scaled logarithmically. The mean ratio of SRTM/TEM is 1.073 (solid line). The plot depicts higher relative differences at lower values of binding potentials.

Table 1 Simplified reference tissue model (SRTM) and transient equilibrium model (TEM) results of $D_{2/3}$ receptor availability measured by [¹⁸F]fallypride PET

| Region | | $[^{18}F]$ Fallypride binding (BP _{ND}) | | | | | | | | |
|--------------|----------------|---|---------|-----------------|------------------|---------|-----------------|----------------|---------|--|
| | M | Medication free | | | Treated patients | | | All subjects | | |
| | SRTM | TEM | Р | SRTM | TEM | Р | SRTM | TEM | Р | |
| Putamen | 22.2 ± 4.9 | 23.4 ± 5.2 | 0.001 | 12.6 ± 4.9 | 13.3 ± 5.5 | 0.001 | 17.4 ± 6.9 | 18.3 ± 7.3 | < 0.001 | |
| Thalamus | 2.11 ± 0.57 | 1.89 ± 0.59 | < 0.001 | 1.02 ± 0.46 | 0.81 ± 0.46 | < 0.001 | 1.56 ± 0.79 | 1.35 ± 0.78 | < 0.001 | |
| Temp. cortex | 0.77 ± 0.30 | 0.78 ± 0.33 | 0.831 | 0.35 ± 0.15 | 0.36 ± 0.17 | 0.197 | 0.56 ± 0.32 | 0.57 ± 0.33 | 0.342 | |
| All regions | 4.7 ± 6.3 | 4.8 ± 6.8 | 0.008 | 8.4 ± 10.2 | 8.7 ± 10.8 | 0.018 | 6.5 ± 8.7 | 6.7 ± 9.2 | < 0.001 | |

BP, binding potential; PET, positron emission tomography.

The data of patients under stable antipsychotic treatment (clozapine, olanzapine, quetiapine, or ziprasidone; n = 50) and subjects without any D_{2/3} receptorassociated medication (n = 50) are depicted. Tests for significant differences between the SRTM and TEM group results were performed using paired two-tailed *t*-tests. 1961

Scan duration of [¹⁸F]fallypride scans I Vernaleken *et al*



Figure 3 Results of artificial stepwise reduction of scan durations from 180 (0 minutes reduction) to 120 minutes (60 minutes reduction). Binding potential (BP_{ND}) values were calculated for the putamina of 50 subjects without second-generation antipsychotic (SGA) treatment using the simplified reference tissue model (SRTM). (**A**) The mean percentage reductions of BP_{ND} results (circles) and the standard deviations of percent discrepancies (error bars) for every 10-minute reduction in scan time. We found an average BP_{ND} underestimation of 0.58% for the first 10-minute reduction. (**B**) The individual extent of BP_{ND} reduction is linearly correlated with the estimated time point of transient equilibrium. In two subjects (†), the equilibrium could not be observed within 180 minutes; in these two cases, the estimated time points of equilibrium (by extrapolation of the fitted time–activity curves) are depicted.

Table 2 $D_{2/3}$ receptor occupancy results of patients (n = 50) who were under stable antipsychotic treatment (clozapine, olanzapine, quetiapine, or ziprasidone)

| | $D_{\scriptscriptstyle 2/3}$ receptor occupancy (%) | | | | | |
|-----------------|---|-----------------|-------------------------------------|---------------|---------------|--|
| | Putamen | | Thalamus | GTi | | |
| SRTM | 43.7±21.2 | -8.7; P<0.0001 | 52.4 ± 20.3 | -2.8; P=0.178 | 55.2±19.4 | |
| TEM | 43.6 ± 22.3 | -14.5; P<0.0001 | 58.1 ± 21.0 $-10.2; P = 0.0004$ | +4.4; P=0.167 | 53.7 | |
| SRTM versus TEM | +0.17; P=0.83 | | -5.64; P < 0.0001 | | +1.49; P=0.33 | |

Occupancy values have been calculated by comparison to an unmedicated control group (n = 50). Simplified reference tissue model (SRTM) and transient equilibrium model (TEM) results as well as the corresponding differences are depicted. Significant regional differences in drug binding are present for putamen versus inferior temporal cortex (GTi) and for putamen versus thalamus comparisons (paired two-tailed *t*-tests). Italic values separate outcome values from group differences and statistical parameters.

r = -0.43, P = 0.0018; thalamus: r = -0.32, P = 0.025; GTi: r = -0.34, P = 0.016; Pearson's correlation).

In the subgroup of patients receiving AP medications, we congruently found small but significant differences between the TEM and SRTM in the putamen and thalamus but not in the GTi (for detailed information see Table 1). Both methods were strongly correlated with each other (linear regression analysis putamen: slope = 0.878, r^2 = 0.95, P < 0.0001; thalamus: slope = 0.932, r^2 = 0.85, P < 0.0001; GTi: slope = 0.780, r^2 = 0.75, P < 0.0001).

The SRTM simulation of reduced scan durations in the putamen, which was the brain region with the highest D₂ receptor density, revealed an average $BP_{\rm ND}$ underestimation of -0.13 ± 0.21 ($-0.58\% \pm 0.94\%$) for the first 10-minute reduction (180 to 170 minutes) (Figure 3A). This effect increased with ongoing time reductions ($\Delta BP_{\rm ND}$: -0.25 ± 0.25 ($-1.16\% \pm 1.17\%$); 130 to 120 minutes). The summed $\Delta BP_{\rm ND}$ effect of a 1-hour reduction in the scan duration was $-1.28 \pm$ 0.96 ($-5.8\% \pm 4.3\%$). In the extrastriatal regions, we found no effects of reduced scan durations on

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binding potentials (e.g., 1-hour reduction in the thalamus: $-0.01\% \pm 2.37\%$). In the putamen, but not in other regions, the change in $BP_{\rm ND}$ due to a 60-minute scan-time reduction was significantly correlated with the corresponding individual estimation of equilibrium time point (r=0.46, P=0.0008; Pearson's correlation; Figure 3B).

The mean regional $D_{2/3}$ receptor occupancies caused by AP treatment are depicted in Table 2. In the thalamus, but not in the putamen or GTi, the SRTM and TEM results showed some significant differences. We found significantly lower $D_{2/3}$ receptor occupancies in the putamen than in the thalamus irrespective of the method used (TEM: -14.5%, P < 0.0001; SRTM: -8.7%, P < 0.0001; two-sided paired *t*-test). The same was true for the GTi (TEM: -10.2%, P = 0.0004; SRTM: -11.5%, P < 0.0001; twosided paired *t*-test). There was no significant difference between the thalamic and cortical occupancies. Based on the simulated data of reduced scan times, a 10-minute reduction of the scan duration would increase these differences (SRTM) to -9.0% (puta-

Discussion

The SRTM approach has frequently been used by different workgroups to quantify the binding potentials using various PET ligands, also including the high-affinity D_{2/3} receptor ligand [¹⁸F]fallypride (Slifstein et al, 2010; Kessler et al, 2009; Siessmeier et al, 2005). Whereas most of these groups accept the SRTM as a suitable model, they use different scan durations and protocols. This retrospective analysis of 100 FP scans reveals some important findings, concerning the sufficient scan duration of FP scans. The estimated time point of the transient equilibrium in untreated subjects (n = 50) and the region with the highest D_2 receptor densities (the putamen) was mainly observed within 3 hours; individual inspection of equilibrium times revealed only 2 out of 100 scans (all two under unblocked conditions) that failed to reach a peak for the binding curve in the putamen during 180 minutes acquisition. However, in most scans, equilibrium occurred much earlier. The mean equilibrium time point in the putamen for unblocked subjects, 121 ± 29.6 minutes, suggests that 180 minutes is adequate for FP scanning in most subjects. The subjects in this study included healthy control subjects, patients suffering from schizophrenia, and subjects undergoing deep brain stimulation to treat Tourette's syndrome. After deep brain stimulation and in schizophrenic patients, $D_{2/3}$ receptor availability was elevated in various brain regions including the basal ganglia (Laruelle, 1998; Vernaleken *et al*, 2009). In particular, deep brain stimulation caused BP_{ND} elevations that exceeded previously measured BP_{ND} values (Vernaleken *et al*, 2009). However, the outliers mentioned above (equilibrium later than 180 minutes) were subjects from the control group. Within the unblocked group, there was no correlation between BP_{ND} and the time to reach equilibrium. This suggests that the present results can be generalized to all subject groups and scan situations. To our knowledge, this is the first systematic publication of equilibrium data for FP.

The main purpose to include the TEM was to estimate the individual time points of transient equilibrium (versus). Though it was not intended to validate the SRTM by the less appropriate TEM, it is noteworthy that there were significant correlations and linear regression slopes of ~0.9 in all regions under both conditions (AP medication and drug free). By definition, such comparisons are only possible when a graphical detection of a transient equilibrium state was possible. A crucial constraint of the TEM is the assumption that the kinetics of nonspecific binding is the same for the reference as well as for the target region. Remarkably, the variability between the two methods was much lower

As a limitation, this investigation did not provide 2TCM data due to the absence of arterial blood sampling. However, in a previous study comparing the SRTM and 2TCM in 10 subjects (also part of the present sample), while there was a high correlation between analytical methods (r=0.95), there was a significant elevation of $BP_{\rm ND}$ values measured by the 2TCM (Siessmeier *et al*, 2005). Similar differences in the extent of $BP_{\rm ND}$ values have been reported in other studies (Slifstein *et al*, 2010). However, the differences between the SRTM and 2TCM are comparable in regions with high and low $D_{2/3}$ receptor binding (10% to 15%).

Low cerebellar-specific ligand binding may cause a bias in reference tissue-based BP_{ND} calculations. The effects of specific binding in reference tissue on the estimation of occupancy have been studied previously (Asselin et al, 2007; Pinborg et al, 2007). Based on the BP_{ND} calculated by V as V/V_{F+NS} -1, and assuming the regional drug-induced occupancies are same across all brain tissues including the 'reference tissue,' then the apparent occupancy is underestimated as $Occ_{app} = Occ/(1 + (1 - Occ) \times BP_{ND})$ (baseline, cerebellum)), where *Occ* is the true occupancy (see Asselin et al, 2007, Eq. A13). Thus, the underestimation in occupancies increases as $BP_{\rm ND}$ in the reference tissue increases. As depicted in detail by Asselin et al (2007), this effect may cause underestimation of $BP_{\rm ND}$ by >50% in the case of [¹¹C]FLB457. This also confirmed by published [¹²³I]epidepride data (Pinborg et al, 2007). In contrast, for FP, there are no data supporting differences of a similar extent. A comparison of FP and ^{[11}C]FLB457 suggests that FP is less vulnerable to cerebellar receptor binding (Vandehey et al, 2010). These authors argue and illustrate by simulation that in case of [¹¹C]FLB457 a higher plasma clearance and the observed lower k_2 and k_{off} as well as the higher affinity contribute to its higher extent of specific cerebellar binding compared with FP. A relatively low proportion of lipophilic metabolites (>20% after 100 minutes) was reported both for FP and [¹¹C]FLB457, with little effect on parameter estimations (Vandehey et al, 2010). Congruently, for FP, the comparison of 2TCM and SRTM never revealed $BP_{\rm ND}$ differences of much >15% as mentioned previously (Siessmeier et al, 2005; Slifstein et al, 2010). As illustrated by Asselin et al (2007), a possible effect of specific cerebellar FP binding on drugassociated occupancy calculations should be similar for all VOIs, as long as confounding factors of ligand binding (e.g., AP action or stimulation of dopamine release) affect the specific ligand binding in cerebellar and cerebral regions to a similar extent. Finally, another factor may account for miscalculations of outcome parameters using the TEM. We cannot exclude that in different regions of the brain (i.e., reference region versus regions with specific binding) the kinetics of free and unspecifically bound ligands can differ.

Insufficient scan durations, however, would definitely cause spatial selective miscalculations depending on the corresponding regional receptor densities. Previously, Olsson and Farde (2001) mentioned that the high-affinity $D_{2/3}$ receptor ligand, ¹¹C-labeled FLB457, is not suitable for these investigations because of its short half-life. However, concerns over FP also arose because of the potential for failure to reach equilibrium conditions, which would cause an underestimation of $BP_{\rm ND}$ in striatal regions (Vernaleken *et al*, 2010).

Using TEM, we showed that transient equilibrium conditions were fulfilled for nearly all of the scans. The high correlations between the SRTM and TEM in all regions (in the validated presence of a transient equilibrium) do not suggest any bias in SRTM- $BP_{\rm ND}$ calculation caused by improper scan durations. However, for reliable estimations of sufficient scan durations, an analysis of artificially reduced scan durations is necessary. This was recently performed analyses of [¹¹C]raclopride for SRTM and ¹¹C]FLB457 (Ikoma *et al*, 2008). In the present study, we artificially reduced the putamen input values of the SRTM stepwise from 180 to 120 minutes; the effect on BP_{ND} values was surprisingly minimal. The first 10 minutes reduction caused a BP_{ND} reduction in the putamen of only -0.13 (-0.58%). The BP_{ND} difference per 10 minutes time reduction increased with further shortening of the scan time up to -0.25(-1.16%). This increase is expected, because for most of the subjects, the scan duration approximates the time point of equilibrium or even falls below it. Consequently, the error induced by reduced scan duration was strongly associated with the time of equilibrium. The data also suggest that further increase of the scan duration would not cause any relevant change to the BP_{ND} values of high-binding regions (far less than 0.5% per 10 minutes block asymptotically approximating 0%). Thus, our results confirm the significant influence of scan duration on $BP_{\rm ND}$ estimates obtained with the SRTM. However, they also illustrate that the extent of miscalculation is very low for FP scans of 180 minutes, even in the critical subset of subjects that have a later occurrence of transient equilibrium than the mean of 120 minutes. As mentioned previously, shorter scan duration can help to avoid breaks in the scan protocol as well as head movements caused by the subject's increasingly realized inconvenience during the PET scan. Thus, hypothetically, the possibility of scan duration shorter than 180 minutes may be beneficial and needs to be discussed. As depicted by Figure 3A, the scan-time reduction from 180 to 170 minutes causes only little changes in the BP_{ND} measures (-0.58%). After 30 minutes, the average difference raises to -1.05% per 10-minute interval finally leading to -1.12% after an hour of scan-time

reduction. This effect is mainly driven by results of subjects with later time points of transient equilibrium. A further reduction of the scan times would *in realiter* remarkably increase the number of individuals that fail to reach this state of equilibrium (six subjects after reduction to 150 minutes and 21 subjects in case of 2-hour scans). Thus, the data do not suggest a scan duration of < 3 hours.

Many PET studies using high-affinity ligands have been performed to characterize the $D_{2/3}$ receptorbinding properties of various APs. Some, but not all, investigations of second-generation antipsychotics report increased receptor binding in extrastriatal regions compared with striatal binding (Frankle, 2007). All PET studies describing this preferential extrastriatal binding effect have been performed using FP as the radioligand. In our previous investigations using quetiapine, ziprasidone, and clozapine (Gründer et al, 2006; Vernaleken et al, 2008b, 2010), we found an overall elevation of +26%in $D_{2/3}$ receptor binding in the GTi compared with that in the putamen (55.2% versus 43.7%; GTi versus putamen) using the SRTM. The difference in occupancy values calculated by the TEM was comparable. If the BP_{ND} values of unblocked putamen or caudate nucleus are systematically biased by improper scan durations, this would cause relevant and region-specific errors of D_{2/3} receptor occupancy estimations. The present results do not suggest a relevant methodological bias-at least none caused by insufficient scan durations: reducing scan time by 10 minutes increased the differences only by 0.3%. This small but significant effect is most likely due to the fact that the sensitivity of BP_{ND} calculations regarding scan-time reductions was present only in unblocked striatal regions (later time point of transient equilibrium) but could neither be observed under blocked conditions nor in regions with low $D_{2/3}$ receptor density. On extrapolating a change of 0.58% per 10 minutes block to a scan duration of 4 hours, even after neglecting the asymptotic reduction of this effect, significant preferential extrastriatal binding observation would still remain. Thus, the preferential extrastriatal binding of APs with low or moderate affinity for the $D_{2/3}$ receptor does not appear to be a methodological artifact caused by improper scan duration.

The calculation of the AP-induced $D_{2/3}$ receptor occupancies was based on the comparison with the unmedicated control group but not on individual pre/posttreatment comparisons. Because of the comparatively large interindividual variability of the binding potential under unblocked conditions, this may cause miscalculations of receptor occupancies on the individual level and can affect correlations with other individual measures (e.g., drug plasma level). The mean occupancy levels as well as the average regional differences in drug binding should be unaffected by this procedure.

We furthermore compared the occupancy values obtained by the SRTM and TEM, and calculated the

effect of scan-time reduction on the occupancy values. In the comparison of $D_{2/3}$ receptor occupancy results measured by SRTM versus TEM, it needs to be mentioned that we found significant discrepancies between the TEM and SRTM for the thalamus VOI. The observed 33% reduction in drug binding in the thalamus compared with the putamen using the TEM shrank to a 20% difference when using the SRTM; that was nevertheless statistically significant. This decrease was most likely caused by fitting problems of the blocked thalamus TAC. Inspection of these curves showed that the Hoerl function was unable to perfectly describe this kind of measure. However, in general, the SRTM results are in line with the TEM data. Both the methods suggest preferential extrastriatal binding.

In summary, to the best of our knowledge, this post hoc analysis includes the largest sample of FP data sets thus far. Justifying our chosen time duration for FP-PET scans (180 minutes), we found only a negligible number of scans that exceeded the 180minute limit in respect to transient equilibrium. Neither the time point of equilibrium nor the choice of the reference method had any relevant influence on $BP_{\rm ND}$ or occupancy values. Three hours FP-data acquisition is sufficient to reliably calculate the D_{2/3} receptor availability using the SRTM and TEM. A limitation of this study is the lack of compartment analysis in the absence of arterial input functions in most subjects.

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Disclosure/conflict of interest

Dr Ingo Vernaleken has served on the speakers' bureaus of Bristol-Myers Squibb (New York, NY, USA), Eli Lilly (Indianapolis, IN, USA), and Glaxo-SmithKline (London, UK). Dr Gründer has served as a consultant for Astra Zeneca (London, UK), Bristol-Myers Squibb (New York, NY, USA), Johnson & Johnson (Beerse, Belgium), Lundbeck (Copenhagen, Denmark), Otsuka (Rockville, MD, USA), and Pfizer (New York, NY, USA). He has served on the speakers' bureaus of Astra Zeneca, Bristol-Myers Squibb, Eli Lilly, Janssen Cilag, Otsuka, Pfizer, Servier (Paris, France), and Wyeth (Madison, NJ, USA). He has received grant support from Alkermes (Cambridge, MA, USA), Bristol-Myers Squibb, Eli Lilly, Johnson & Johnson, and Pfizer. Dr Dean F. Wong has received grant support from the NIH, Amgen, Avid, the DANA Foundation, Intracellular, Johnson & Johnson, Otsuka, Lilly, Lundbeck, Merck, Orexigen, Roche, and Sanofi Aventis. Dr Wolfgang Schaefer has served as a consultant for GE Healthcare Buchler (München, Germany), Siemens Medical Solutions (Erlangen, Germany), Philips (Aachen, Germany), and Bristol-Myers Squibb (München, Germany). He has received grant support from Mediso Medical Imaging Systems (Budapest, Hungary).

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