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Impact of framing scheme optimization and smoking status on binding potential analysis in dynamic PET with [^{11}C]ABP688

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Abstract

Background For positron emission tomography (PET) ligands, such as [^{11}C]ABP688, to be able to provide more evidence about the glutamatergic hypothesis in schizophrenia (SZ), quantification bias during dynamic PET studies and its propagation into the estimated values of non-displaceable binding potential (BP_{ND}) must be addressed. This would enable more accurate quantification during bolus + infusion (BI) neuroreceptor studies and further our understanding of neurological diseases. Previous studies have shown BP_{ND} -related biases can often occur due to overestimated cerebellum activity (reference region). This work investigates whether an alternative framing scheme can minimize quantification biases propagated into BP_{ND} , whether confounders, such as smoking status, need to be controlled for during the study, and what the consequences for the data interpretation following analysis are. A group of healthy controls (HC) and a group of SZ patients (balanced and unbalanced number of smokers) were investigated with [^{11}C]ABP688 and a BI protocol. Possible differences in BP_{ND} quantification as a function of smoking status were tested with constant 5 min ('Const 5 min') and constant true counts ('Const Trues') framing schemes. In order to find biomarkers for SZ, the differences in smoking effects were compared between groups. The normalized BP_{ND} and the balanced number of smokers and non-smokers for both framing schemes were evaluated.

Results When applying F-tests to the 'Const 5 min' framing scheme, effect sizes (η^2p) and brain regions which showed significant effects fluctuated considerably with $F = 50.106 \pm 54.948$ (9.389 to 112.607), P -values 0.005 to < 0.001 and $\eta^2p = 0.514 \pm 0.282$ (0.238 to 0.801). Conversely, when the 'Const Trues' framing scheme was applied, the results showed much smaller fluctuations with $F = 78.038 \pm 8.975$ (86.450 to 68.590), $P < 0.001$ for all conditions and $\eta^2p = 0.730 \pm 0.017$ (0.742 to 0.710), and regions with significant effects were more robustly reproduced. Further, differences, which would indicate false positive identifications between HC and SZ groups in five brain regions when using the 'Const 5 min' framing scheme, were not observed with the 'Const Trues' framing.

Conclusions Based on an [^{11}C]ABP688 PET study in SZ patients, the results show that non-consistent BP_{ND} outcomes can be propagated by the framing scheme and that potential bias can be minimized using 'Const Trues' framing.

Keywords PET, Quantification bias, Non-displaceable binding potential, Framing scheme, mGluR5, Smoking status, Schizophrenia

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Background

Data from dynamic positron emission tomography (PET) acquisitions are often acquired in list-mode with high temporal granularity and are subsequently framed during the reconstruction process. This means that the acquired data are framed into certain time intervals and grouped in a sequence of images (dynamic image stack). In addition to the acquisition of the prompt events for each frame, it is also necessary to correct for several physical effects, such as randoms, scattered events, attenuation, and dead time. However, some of these corrections become more challenging at low count levels. This can become especially relevant during bolus + infusion (BI) studies with ^{11}C radiotracers, where count rates become low toward the end of the acquisition leading to reconstructed frames with low counts. In the frame-based iterative reconstruction of the dynamic PET data, the reconstruction is performed independently for each frame. Once the dynamic frames are reconstructed, the activity of each voxel or selected volume of interest (VOI) can be obtained for each frame, resulting in a time activity curve (TAC) for each voxel or VOI.

The maximum likelihood-expectation maximization (MLEM) and the 3D ordinary poisson ordered-subset (OS) EM (3D OP-OSEM) reconstruction methods tend to be biased in regions with low activity concentrations and at low count rates. Furthermore, they tend to converge rather slowly for these regions. Thus, a very high number of iterations paired with long reconstruction times would be necessary to eliminate the positive bias [1]. Furthermore, VOIs can differ in convergence according to their shape, volume, and activity concentration; TACs with higher time resolution have shorter time intervals and, consequently, higher noise levels and quantification biases [1, 2]. This can be significant for BI studies as the time needed to reach equilibrium means that the count rate decreases during the study. Some of the reconstruction and image correction methods currently available may not sufficiently regularize noise and may introduce high variance during scatter correction at low counts [3, 4]. However, alternative reconstruction methods are emerging that aim to improve the quantification accuracy of the dynamic PET data [5–8]. Quantification bias for low counts during dynamic PET neuroreceptor studies and its propagation into the often-used outcome parameter non-displaceable binding potential (BP_{ND}) is an important factor that must be addressed if accurate quantification is to be achieved. Previous studies have already shown BP_{ND} -related biases of up to -15% due to overestimation of the cerebellum activity concentration (usually used as reference region for several neuroreceptor radiotracers and due to the fact that it shows low

activity concentration and therefore low count rates during the acquisition) [2, 9–11].

As count rates get lower over the course of the BI study protocol, longer time frames are required to control image noise. Hence, one can use time frames of the same length throughout the protocol, or, since the frames after the tracer reaches equilibrium are the most important, non-constant intervals (increasing over the acquisition time) can be used until the tracer reaches equilibrium. At this point, the time frame is then fixed at a constant interval for the duration of the acquisition time (same time interval in the frames from equilibrium to the end of the acquisition time course). However, this choice is a trade-off between frame temporal resolution, counts in the frame (statistical noise), and quantification accuracy. Moreover, for multimodal studies with simultaneous PET/computed tomography (CT) or PET/magnetic resonance (MR), there is no consensus on how a framing scheme should be planned in order to facilitate enough temporal resolution to detect changes in BP_{ND} . When the simple ratio method for analysis/quantification is the method of choice, reducing the bias caused by the low count rates, particularly in the reference region (e.g., cerebellum), is imperative to ensure a high level of accuracy.

The quantification bias described above may not only disturb the accurate detection of changes by stimulation or other challenges of the neuroreceptor system but may also impact biological confounders introduced by the investigated subjects.

To demonstrate these factors relating to quantification bias, patients with schizophrenia (SZ) and healthy controls (HC) underwent PET investigations with [^{11}C]ABP688 [12], an allosteric antagonist for the metabotropic glutamatergic type 5 receptor (mGluR5), which may provide more evidence to elucidate the glutamatergic hypothesis in SZ.

An important confounder for [^{11}C]ABP688 studies is the smoking status of the subject, as smoking has been shown to cause a global reduction in mGluR5 binding, as reported [13]. Furthermore, 60–90% of SZ patients are smokers [14], and it is known these individuals smoke more heavily and experience more difficulties in quitting smoking than smokers without SZ [15]. Further bias may arise from the fact that smokers might metabolize the radioligand faster in other organs (i.e., liver) than non-smokers, possibly influencing the uptake of [^{11}C]ABP688 in the brain during BI protocol. As this might affect the count rate, the quantification bias and consequently the BP_{ND} results could also be impacted.

Other confounders may be gender and age. Although no significant effects of this kind were found in a study with healthy subjects [16], SZ patients presented different patterns between males and females [17]. Therefore,

to eliminate this potential confounding factor, only male SZ patients (and HC) took part in this study.

The aim of this work was to evaluate the impact of the chosen framing scheme on BP_{ND} quantification and to show if an optimized frame scheme can minimize quantification bias. In addition, using the example of a SZ study with [^{11}C]ABP688 as the radiotracer, we demonstrated the need to control confounders, such as smoking status, and the potential risk of misinterpreting the data due to the aforementioned influences on BP_{ND} quantification.

Methods

Subjects

Seventeen male subjects were studied for each group (SZ patients and HC) and were matched for age (SZ = 38.4 ± 10.6 and HC = 38.5 ± 11.4), level of smoking addiction, and education. All SZ patients were diagnosed with F20.0 paranoid SZ, except one patient who was diagnosed with F20.3 undifferentiated SZ. The patients in the SZ group had an illness duration of 19 ± 11 years. Subjects were instructed not to drink coffee or alcohol and not to take any medicine within 24 h before the measurement (with the exception of daily medication).

The study was approved by the Ethics Committee of the Medical Faculty at the RWTH Aachen University and the German Federal Office for Radiation Protection (Bundesamt für Strahlenschutz). Patients were recruited from Uniklinik RWTH Aachen and complied with the diagnostic criteria according to DSM-IV [18]. Informed consent was obtained for all subjects.

Three groups of SZ patients and HC were selected to represent different conditions for the analysis, as shown below:

Majority of smokers in the groups (unbalanced sample)

The effects of the smoking status on BP_{ND} quantification were evaluated in HC (a.1 and a.2) and SZ patients (b.1 and b.2) individually. Each group consists of nine smokers (S) and six non-smokers (nS), and therefore has a majority of smokers. To evaluate this effect, 2 nS were randomly excluded from each group of 17 SZ and 17 HC twice-represented by study cases called: A-2, A-3, B-2, and B-3. This unbalance mirrors the usual condition and the difficulty in finding nS SZ patients for such studies since smoking addiction is a known and relevant comorbidity of SZ. We tested the influence that the unbalance in the number of S in the analyzed groups (in two different cases for HC and SZ group selection) can have on BP_{ND} between and within the groups.

Equivalent number of smokers and non-smokers in the groups (balanced sample)

The effects of the smoking status on BP_{ND} quantification were individually evaluated in the HC and SZ groups again, but this time with the same number of S and nS in each group (8 S and 8 nS)-represented by case study A-1 and B-1) excluding one S per group (HC and SZ). As mentioned previously, the majority of published studies report a numerical balance of smoking status in SZ and HC, but usually with a majority of S in each group.

SZ versus HC-Equivalent number of smokers in the groups (balanced sample)

Balanced by smoking status, the samples were compared within and between group effects on BP_{ND} . This analysis focused on potential differences in BP_{ND} caused by SZ.

HC versus SZ-Normalized BP_{ND} equivalent number of smokers in the groups (balanced sample)

Balanced by smoking status, the samples were compared, but this time normalized by differences in smoking status estimated from the temporal cortex regions. The focus of this analysis was on potential differences in the normalized BP_{ND} caused by SZ. The temporal cortex was chosen as the reference region for the smoking status as it shows a robust significant difference between S and nS, which was observed during this study in both groups for all kinds of evaluations of smoking effects on BP_{ND} .

PET/MR acquisition and PET BI protocol

Data were acquired with a hybrid 3 T MR-BrainPET insert [19] as reported in [12]. The PET data were acquired over 65 min in list-mode while applying a BI protocol (50% of the [^{11}C]ABP688 activity was applied to the subject as a bolus and 50% was infused, not exceeding 600 MBq of injected activity per subject). PET and MR were synchronized during the acquisition so that the first 5–6 min of the MR resting data are acquired during a time where the equilibrium of [^{11}C]ABP688 is reached (30 min after the bolus injection), thus enabling the evaluation of the BP_{ND} baseline for each subject. Smokers were not allowed to smoke in the two hours prior to the radiotracer injection.

The MR data acquisition started simultaneously with the PET bolus injection and the PET list-mode acquisition. Before the radiotracer reached equilibrium, a longitudinal relaxation time (T_1) MPAGE image was acquired for structural information with the following parameters: repetition time (TR)=2250 ms; echo time (TE)=3.03 ms; field of view (FOV)= $256 \times 256 \times 176$ mm³, matrix size=256 × 256 pixels, flip angle=90°, 176 sagittal slices with 1 mm slice thickness, a generalized

autocalibrating partially parallel acquisition (GRAPPA) factor of 2, and 70 auto-calibration signal lines. The structural MR image was also used for the PET attenuation correction based on the template method [20]. At about 30 min after the PET bolus injection, functional MR images (fMRI) were acquired using an effective transverse relaxation time (T_2^*) weighted echo planar image (EPI) sequences with the following parameters: TR=2200 ms, TE=30 ms, matrix size: 64×64 pixels with 36 slices and $3 \times 3 \times 3.75$ mm³ voxel size. During the fMRI resting state (RS) scan, 180 functional volumes (approximately 5–6 min) were recorded for each subject. During this measurement, subjects were asked to lie down and keep their eyes closed without falling asleep. Only the time corresponding to the RS acquisition was chosen for the BP_{ND} analysis in this work.

PET image reconstruction and BP_{ND} estimation

To compare the influence of the framing on the BP_{ND} bias, the data were reconstructed with two different framing schemes. In the first reconstruction, the list-mode data were sorted into 5 min frames ('Const 5 min'). In the second reconstruction, the frames were reconstructed according to a method already presented [11, 21] with a constant number of true coincidences in each frame ('Const Trues'), which minimizes and keeps the bias constant. For the 'Const Trues' scheme, PET true counts per frame based on this scheme were equivalent between subjects and also synchronized according to the fMRI RS acquisition moment. The subject with the lowest average of true counts registered a total of 9.35×10^6 counts in the last 5 min of the acquisition. This value was taken as the fixed reference count per frame, and three interval time fractions from the subject where this low limit occurred were computed to give 0.267, 0.324, and 0.408 for the acquisition intervals. The intervals were computed for each subject according to the total length of the fMRI intervals and these fractions. The three frames within the time interval were then grown symmetrically around the center of these three intervals until the reference count (9.35×10^6) was reached. Frames without matched counts and/or were not synchronized with fMRI were discarded for complete acquisition. In this way, all seventeen subjects had the same number of counts during the RS moments, and the reconstruction bias was equal for all time intervals.

For both reconstructions, corrections for attenuation [20], random and scattered coincidences, and dead time were applied. The 3D OP-OSEM [2] reconstruction was performed with 32 iterations and two subsets in a matrix of 256×256 pixels and 153 slices with a 1.25 mm³ isotropic voxel size. Post-processing was performed by filtering with a 2.5 mm 3D Gaussian filter. Head motion

correction was performed with a multi-frame acquisition method (MAF) implemented in-house [22].

The frame(s) corresponding to the fMRI RS (starting around 30 min after the PET bolus injection and about 5 min in duration) were used for computing the BP_{ND} using the simple ratio method shown below (Eq. 1).

$$BP_{ND} = \left(\frac{C_T}{C_R} \right) - 1, \quad (1)$$

where C_T is the activity concentration in the target regions and C_R is the activity concentration in the reference region (cerebellum gray matter (GM)) after the radioligand reaches equilibrium and during fMRI RS.

The activity concentration was extracted using the Pmod v.3.9 PNEURO tool, and T₁ MPRAGE images were used as the anatomical reference. The VOIs were applied in the subject's PET space using the Hammer atlas [23]. The activity concentrations were extracted for 17 regions and the cerebellum GM, which was the reference region [17, 24–26]. The VOIs were: whole-brain GM, frontal left, frontal right, orbitofrontal cortex middle, parietal left, parietal right, temporal left, temporal right, temporal middle, primary auditory, anterior cingulate (ACC), posterior cingulate (PCC), caudate, putamen, thalamus, motor cortex and white matter.

Statistical analysis

Statistical analysis was conducted with the Statistical Package v.25.0 (IBM SPSS Inc., Chicago, IL). The significance level was set to $P=0.05$ for all tests, and all tests were two-tailed. Standard error (SE) bars were computed according to [27], which was implemented as a tool in the SPSS package. The SE bars were added by the SPSS toolbox to the BP_{ND} values calculated from PET image data. All analyzed brain regions were considered as a repeated variable (within-subjects) in the general linear model and, when appropriate, as between-subjects variables for the analysis of smoking status, SZ, and HC. Corrections for multiple comparisons were applied using the Bonferroni method, and the significance level was adjusted accordingly ($P=0.003$). To evaluate the effect sizes of smoking status, the results from F-tests were also reported for each brain region in a pairwise comparison and when $P<0.05$ for significant effects of smoking per region was reached.

For between-subject (HC and SZ) analyses, the regions from each group were compared using the general linear model, and the same significance level was assumed. Effect sizes were reported based on the partial eta squared (η^2p), where 0.01 represents a small, 0.06 represents a medium, and 0.14 or higher represents a large effect size.

Despite having only 16 subjects per group (when groups were balanced with respect to the smoking status), the group size was considered sufficient based on available literature and standard power analysis. Previous comparable studies comprised the unique [^{11}C] ABP688 PET study in SZ ($n=15$ per group) at the time of the design of the study [17] and a ketamine study (one group of $n=10$) [28]. Based on the sample size used in this study, between-subject effects were detected with a statistical power of around 80%, assuming repeated BP_{ND} measures for all analyzed brain regions for two groups with small to medium-size effects and $P=0.05$. The estimations of the power were evaluated using GPower v.3.1 software.

Results

Balanced and unbalanced groups according to smoking status in BP_{ND} analysis

The effect sizes (η^2p) were computed for smoking in each of the HC and SZ groups with both ‘Const 5 min’ and ‘Const Trues’ framing schemes, with standardized SE, and with an equivalent number of S and nS subjects. Table 1 shows a summary of the results for the different conditions of the groups arranged according to smoking status and framing schemes.

Figure two shows all regions evaluated for BP_{ND} changes due to smoking status. The results are arranged according to the applied framing scheme and for the different test cases reported in Table 1. The regions that showed significant effects are marked by an asterisk on top of the green bars, which reports the effect size (η^2p).

Table 1 Summary of all tested cases and correspondent results for smoking effects on BP_{ND}

Framing scheme	Smoking status groups	F-test	P-value	η^2p^*
(a) Const 5 min	A-1 16 HC (8 S and 8 nS) 16 SZ (8 S and 8 nS)	9.389	0.005	0.238
	A-2 15 HC (9 S and 6 nS) 15 SZ (9 S and 6 nS)	28.323	< 0.001	0.503
	A-3 15 HC (9 S and 6 nS) 15 SZ (9 S and 6 nS)	112.607	< 0.001	0.801
	B-1 16 HC (8 S and 8 nS) 16 SZ (8 S and 8 nS)	86.450	< 0.001	0.742
	B-2 15 HC (9 S and 6 nS) 15 SZ (9 S and 6 nS)	79.076	< 0.001	0.739
	B-3 15 HC (9 S and 6 nS) 15 SZ (9 S and 6 nS)	68.590	< 0.001	0.710

* Effect size

Table 1 Analysis for both framing schemes with balanced (cases A-1 and B-1) and unbalanced groups (cases A-2, A-3, B-2 and B-3) with regard to smoking statuses in the subgroups. The results for the effect of smoking on BP_{ND} were obtained with the general linear model

The effect of smoking status per brain region was evaluated in a pairwise comparison. When unbalanced groups were analyzed, we observed an increased significance and larger effect size (as expected). Certainly, this did not dramatically change the conclusions about the smoking status affecting BP_{ND}, but it is important to know how the significance and effects of a variable can influence the study in general and, in particular, how it influences the results in the involved brain regions. For case A-2, a pairwise comparison between regions with both groups—HC and SZ, we observed a slightly changed scenario. In A-3, we found an even larger effect size. This time, the regions that showed significant effects in the comparison were again different for other group compositions (Fig. 1 a and b). However, when the data were analyzed with the ‘Const Trues’ framing scheme (Fig. 1 d to f), we only observed a minor impact on smoking effect sizes between tested cases. Moreover, the F-tests were different to the corresponding results obtained with the ‘Const 5 min’ framing scheme when varying the composition of the groups. Furthermore, when using ‘Const Trues’ framing, the effect sizes and probabilities were reproduced more consistently in all three scenarios, B-1, B-2, and B-3.

Balanced groups for the evaluation of the effect of smoking status on BP_{ND} per group (HC and SZ individually)

Another important question addressed in this study was whether smoking status affects the BP_{ND} in each group. As discussed in the Introduction, SZ patients find smoking cessation more difficult and usually smoke more heavily compared to non-SZ groups. It is, therefore, important to verify whether smoking status affects the BP_{ND} metrics in both groups unequally. Table 2 shows the regions affected by smoking status in HC (8 S and 8 nS) and SZ (8 S and 8 nS), and in the dependence of the framing scheme.

It is evident that the SZ group showed more prominent effects, and more brain regions were affected by smoking status than the HC group. It must be expected that the outcome of the results is strongly influenced by the SZ group when smoking status was evaluated with both groups merged in the analysis. Interestingly, the temporal cortex showed significant differences for all methods and comparisons, indicating this mGluR5-rich region is heavily impacted by smoking habits. Figure 2 shows the regional average of the BP_{ND} for a) HC and b) SZ groups according to the smoking status and for the ‘Const 5 min’ and ‘Const Trues’ framing schemes. It should be noted that the ‘Const Trues’ scheme led to more brain regions with significant differences when comparing groups with

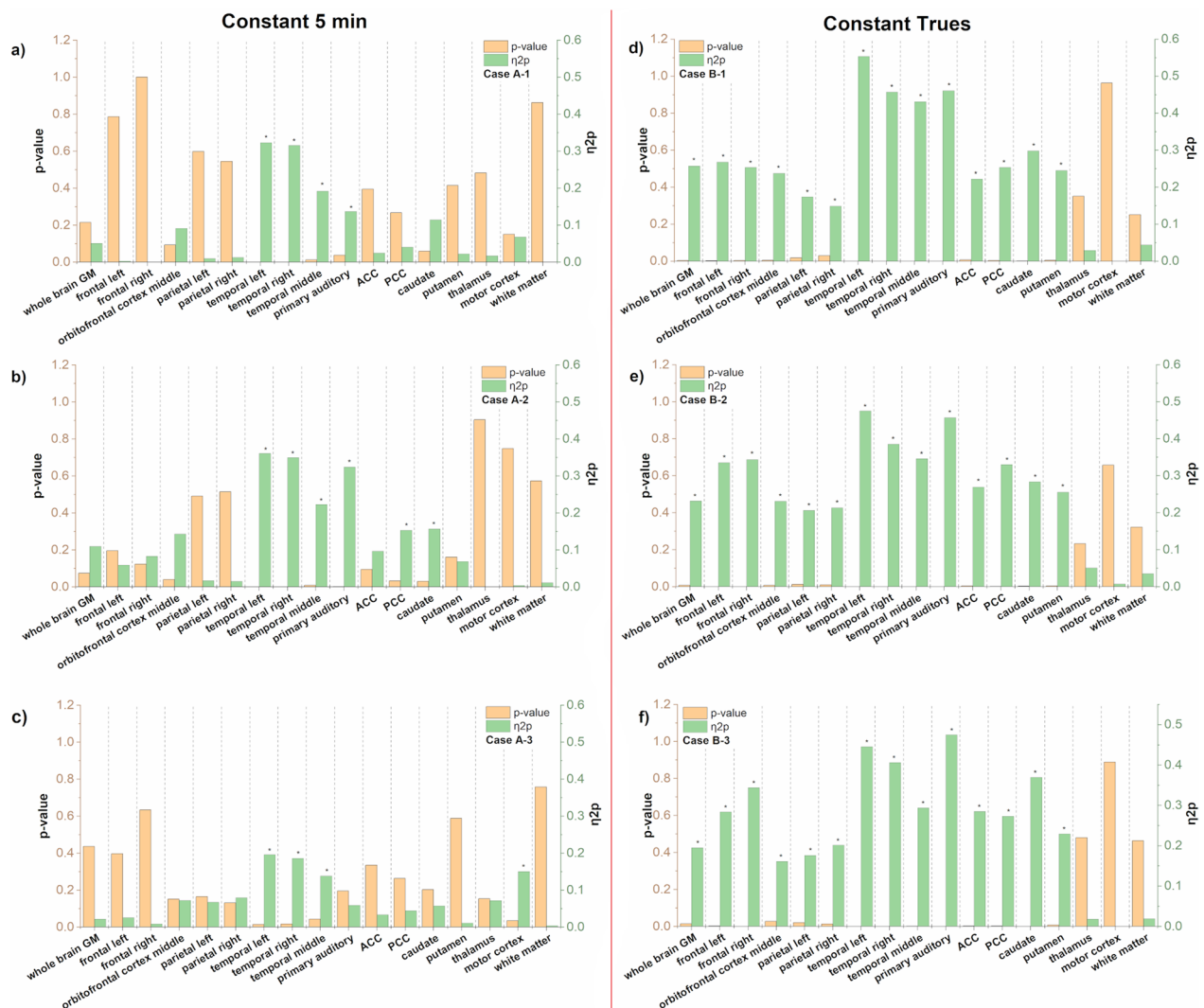


Fig. 1 Regions depicted by significance and effect size according to test cases reported in Table 1. **a–c** Framed with ‘Const 5 min’ scheme **d–f** framed with ‘Const Trues’ scheme. **a** and **d** Cases A-1 and B-1 balanced samples by the number of S and nS, **b** and **e** Cases A-2 and B-2 unbalanced samples by the number of S and nS, **c** and **f** Cases A-3 and B-3 additional case of unbalanced samples by the number of S and nS. Here, 2 nS were randomly removed, and one additional S per group was added to simulate changes in the groups differing to cases A-2 and B-2 obtained with the same random method but differing in the nS removed from each group

different smoking statuses than in the case where the ‘Const 5 min’ scheme was applied.

Balanced groups and BP_{ND} normalized to the temporal cortex: HC versus SZ comparison

In order to find biomarkers for SZ, the different effects of smoking status have to be considered for the comparison between subjects. As the temporal cortex always showed a significant effect of the smoking status on BP_{ND} , independent of the framing scheme or balance of S and nS subjects in the groups, we normalized all other brain regions with respect to the average BP_{ND} of the three

analyzed temporal cortex regions. In addition, we also compared the results for ‘Const 5 min’ and ‘Const Trues’ framing schemes. Table 3 provides a comparison of the groups for all investigated brain regions and both framing schemes.

For ‘Const 5 min’ framing, five regions showed significant differences, which was, however, not the case when applying the ‘Const Trues’ scheme. This shows how the uncontrolled bias of constant time framing can affect the results and subsequent interpretations. In this case, the false positive outcome would indicate an important significant difference between HC versus SZ in mGluR5

Table 2 Results for smoking status effects on BP_{ND} in different brain regions separated into HC and SZ groups and with framing schemes applied

Group	Framing scheme	Affected brain region	F-test	P-value	η^2p^*
HC	Const 5 min	Temporal right	5.867	0.030	0.295
		Motor cortex	4.677	0.048	0.250
SZ	Const 5 min	Temporal left	10.693	0.006	0.433
		Temporal right	7.551	0.016	0.350
		Temporal middle	5.039	0.041	0.265
		Caudate	4.849	0.045	0.257
		Temporal left	11.086	0.005	0.442
HC	Const Trues	Temporal right	9.471	0.008	0.404
		Temporal middle	7.010	0.019	0.334
		Whole brain GM	7.019	0.019	0.334
SZ	Const Trues	Frontal left	12.911	0.003	0.480
		Frontal right	18.371	<0.001	0.568
		Parietal left	7.233	0.018	0.341
		Parietal right	13.766	0.002	0.496
		Temporal left	32.300	<0.001	0.698
		Temporal right	20.491	<0.001	0.594
		Temporal middle	16.253	0.001	0.537
		Primary auditory	35.933	<0.001	0.720
		ACC	8.220	0.012	0.370
		PCC	18.291	<0.001	0.566
		Caudate	10.589	0.006	0.431
		Putamen	9.192	0.009	0.396

*Effect size

Table 2 Groups balanced by smoking status, age, gender and education for the analysis of smoking status effects on BP_{ND} in HC and SZ grouped by framing scheme methods. The results from a pairwise analysis between subgroups (S and nS) for brain regions with significant differences reported

binding. No significant differences between the groups were observed when ‘Const Trues’ framing was used for bias minimization together with variance reduction in the data. In this case, the frontal left cortex was only significant before the Bonferroni correction. Thus, bias reduction and the control of confounders in the groups is highly relevant for the analysis of [¹¹C]ABP688 PET in SZ studies.

Discussion

In this work, we have demonstrated the influence of the framing scheme on reconstruction bias in a PET *in vivo* study with [¹¹C]ABP688 for the first time. We have also shown how smoking status affects the BP_{ND} differently between regions by comparing SZ and HC groups. Considering that smoking addiction is a relevant comorbidity of SZ [29–35], the presented work confirms common limitations of studies on SZ that face the challenge of finding non-smokers for group comparisons with balanced groups demographics, and further shows the influence smoking status has on the data analysis in situations

with a majority number of smokers in the study cohort. One of the limitations of this study was the subgroup number, with only eight smokers and non-smokers in each group (HC and SZ). It would be interesting to validate our findings by applying the same kind of analysis to a larger sample size of balanced and unbalanced numbers of smokers in the groups.

For the first time, an inter-subject comparison between framing schemes was performed that showed how framing scheme-dependent bias propagates into the BP_{ND}.

Further, we demonstrated relevant changes in the results when our newly proposed ‘Const Trues’ framing scheme was used instead of the previous ‘Const 5 min’ framing scheme. As already shown in a previous publication [11], the ‘Const Trues’ framing scheme minimizes the reconstruction bias and keeps it constant during the BP_{ND} quantification. When applying the ‘Const 5 min’ framing scheme, the results varied more in terms of statistical significance, effect sizes, and brain regions affected by smoking status. This shows the potential of the ‘Const Trues’ framing scheme to minimize count fluctuations, to reduce the bias, and to obtain more consistent standard errors and variability.

Moreover, we also demonstrated the importance of evaluating the effects of smoking status on BP_{ND} in SZ and HC groups individually and showed that mGluR5 availability in SZ patients is more affected by tobacco than in HC when imaged with PET [¹¹C]ABP688. Furthermore, potential false positive results, even when considering normalization for the different effects of smoking status over groups, were also presented, again showing the limitations caused by uncontrolled framing-dependent bias in the image reconstruction. When the comparison between SZ and HC groups was performed with normalized BP_{ND} (i.e., accounting for different smoking effects in both groups), false positive results were observed in the case of the ‘Const 5 min’ scheme but were not reproduced for the ‘Const Trues’ framing scheme. This means the strategy of normalizing the BP_{ND} by the region most affected by smoking status, i.e., the temporal lobe regions, was not useable as a biomarker for SZ, which significantly could differentiate the groups. However, it is possible that our basic assessment of smoking addiction (cigarettes consumption per day and years of consumption) unintentionally brought more variability into the analysis, and owing to the lack of a complete Fagerstrom scale from all participants and only a rudimentary dichotomous variable (S or nS) in the analysis, this may represent a limitation of the study. In light of this, it would be interesting to perform a deeper evaluation of tobacco addiction in an [¹¹C]ABP688 study of SZ compared to HC in terms of the effects on BP_{ND}. In addition to PET conventional quantification analysis, new methods are becoming available to aid the

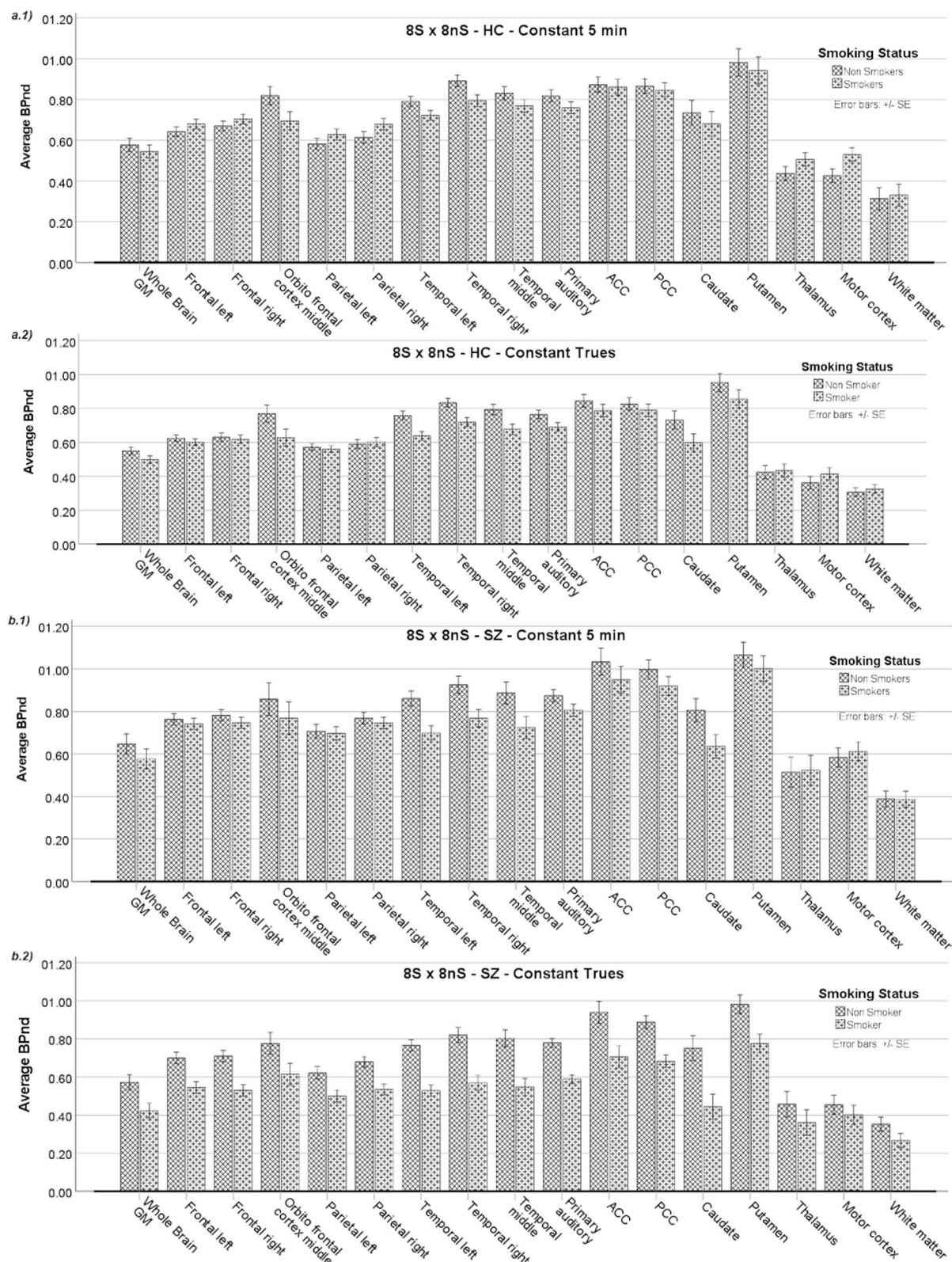


Fig. 2 Smoking effects on BP_{ND} for HC and SZ groups showed individually for both framing schemes. The Average BP_{ND} for balanced subgroups according to smoking status is shown in: **a.1** For HC and 'Const 5 min' framing scheme, **a.2** For HC and 'Const Trues' framing scheme, **b.1** For SZ and 'Const 5 min' and **b.2** For SZ and 'Const Trues' framing scheme

Table 3 Results of the normalized BP_{ND} in the comparison between the HC and SZ groups separated by framing scheme

Brain region	Constant 5 min framing			Constant true counts framing		
	F test	P-value**	η^2p^*	F test	P-value**	η^2p^*
Whole Brain GM	1.401	0.247	0.048	0.326	0.572	0.012
Frontal left	12.036	0.002**	0.301	6.445	0.017	0.187
Frontal right	10.679	0.003**	0.276	2.469	0.127	0.081
Orbitofrontal cortex middle	0.710	0.407	0.025	2.556	0.121	0.084
Parietal left	10.712	0.003**	0.277	4.091	0.053	0.127
Parietal right	16.636	< 0.001**	0.373	5.205	0.030	0.157
Primary auditory	1.072	0.309	0.037	0.910	0.348	0.031
ACC	4.525	0.042	0.139	3.690	0.065	0.116
PCC	4.708	0.039	0.144	3.112	0.089	0.100
Caudate	0.044	0.835	0.002	0.253	0.619	0.009
Putamen	1.632	0.212	0.055	1.561	0.222	0.053
Thalamus	1.351	0.255	0.046	0.271	0.607	0.010
Motor cortex	10.829	0.003**	0.279	4.127	0.052	0.128
White matter	1.366	0.252	0.047	0.278	0.602	0.010

*Effect size, **Significance $P \leq 0.003$ after Bonferroni correction are highlighted in bold italic

Table 3 Samples balanced by smoking status, age, gender and education for the analysis of normalized BP_{ND} based on different framing schemes. All brain regions with statistical analysis results were reported from a pairwise analysis between groups

search for biomarkers in SZ. For example, radiomics is a method that has been tested in neuro-oncology [36] and is particularly interesting as it enables a great deal of additional information to be gathered during the analysis while simultaneously looking for the PET binding metrics and other feature-based metrics [37].

When ‘Const Trues’ framing was chosen, variations in the statistical significance, smoking effects, and effect size were minimized, and the regions affected by smoking status were reproduced when both groups were merged during the analysis. Once more, when evaluating HC and SZ groups individually, the effects of smoking status are more prevalent in a higher number of brain regions in SZ groups compared to HC groups.

We hypothesize that, due to the nature of the metabolism pathway, especially for subjects who smoke, the increased metabolism by other organs, such as the liver, leads to a reduced activity concentration of the radiotracer in the brain cortex (noted in the blood curves during the metabolite’s correction-data not shown). This might contribute to lower activity in the cortex and, in this case, a reduction in mGluR5 receptor binding. This would lead to a reduced count rate and reduced amount of events detected in the reconstructed image, thus causing increased variability in the image frame and bias in the BP_{ND} quantification. However, the results suggest this effect was minimized by the proposed ‘Const Trues’ framing scheme because the scheme keeps constant true events within the frames and, therefore, keeps the bias

constant for the entire acquisition time course and inter-between-subjects frames. It is clear that other mechanisms could be responsible for this interplay of radiotracer metabolism, brain uptake and/or mGluR5 availability since the relationship between nicotine addiction and the functional role of mGluR5 down-regulation remains unclear, as already reported [13, 38]. Interestingly, smoking-related differences in BP_{ND} and the number of brain regions involved were strongly observed in SZ but were not observed in the corresponding HC group.

In light of these results, we strongly recommend the evaluation of the impact on smoking status in samples of SZ patients and HC groups prior to group analysis for other variables considered in the study. Similarly, in order to minimize biased results from the analysis in PET BI studies, we recommend an evaluation of the quantitative bias for the BP_{ND} prior to defining the framing scheme.

Conclusions

This study shows how the framing scheme impacts reconstruction bias by demonstrating how bias propagation affects BP_{ND} metrics, consequently leading to inconsistent outcomes in a PET BI study on SZ with [¹¹C]ABP688. In this preliminary investigation, potential false positive findings were minimized by applying the proposed ‘Const Trues’ framing scheme. Furthermore, the effect of additional confounding factors, such as smoking status in SZ, can be mitigated with this alternative framing scheme.

Abbreviations

PET	Positron Emission Tomography
[¹¹ C]ABP688	(11C)-methyl iodide with the sodium salt of desmethyl-ABP688 (3-(6-methyl-pyridin-2-ylethynyl)-cyclohex-2-enone oxime)
SZ	Schizophrenia
BP _{ND}	Non-Displaceable Binding Potential
BI	Bolus + Infusion
HC	Healthy Controls
Const 5 min	Constant 5 min
Const Trues	Constant True Counts
η ² p	Effect Size
mGluR5	Metabotropic Glutamatergic Type 5 Receptor
VOI	Volume of interest
TAC	Time Activity Curve
MLEM	Maximum Likelihood-Expectation Maximization
3D OP-OSEM	Ordinary Poisson-Ordered Subset-Expectation Maximization
CT	Computed Tomography
MR	Magnetic Resonance
F20.0	Paranoid Schizophrenia Diagnose Classification
F20.3	Undifferentiated Schizophrenia Diagnose Classification
S	Smokers
nS	Non-Smokers
FOV	Field of View
MPRAGE	Magnetization Prepared-Rapid Gradient Echo
T ₁	Longitudinal Relaxation Time
TR	Repetition Time
TE	Echo Time
GRAPPA	Generalized Autocalibrating Partially Parallel Acquisition
EPI	Echo Planar Image
fMRI	Functional MR Images
T ₂ [*]	Effective Transverse Relaxation Time
RS	Resting State
MAF	Multi-frame Acquisition Method
C _T	Activity Concentration in the Target Regions
C _R	Activity Concentration in the Reference Region
GM	Gray Matter
ACC	Anterior Cingulate Cortex
PCC	Posterior Cingulate Cortex

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Author contributions

CRB: PET bolus + infusion protocol optimization, data acquisition, constant trues framing scheme validations, PET imaging data analysis, statistical design and analysis, manuscript writing, correction, and revision. JS: PET reconstruction discussions, manuscript revision and correction. AI: script for PET True events extraction, and manuscript revision. LT: MR-PET hardware and data acquisition. HH: PET study design discussions and revision and corrections of the manuscript. NJS: MR-PET hardware and revision of the manuscript. IN: ABP study design and setup, approval ethics and BfS, and revision of the manuscript. CL: constant trues framing scheme, study design and setup, data analysis revision, and manuscript corrections and revision. All authors read and approved the final manuscript.

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Availability of Data and Materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

The study was approved by the Ethics Committee of the Medical Faculty at the RWTH Aachen University and the German Federal Office for Radiation Protection (Bundesamt für Strahlenschutz). Patients were recruited from Uniklinik RWTH Aachen and complied with the diagnostic criteria according to DSM-IV [18]. Informed consent was obtained for all subjects. This study was performed in accordance with the ethical standards as laid down in the 1964 Declaration of Helsinki and its later amendments.

Consent for Publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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References

1. Van Slambrouck K, Stute S, Comtat C, Sibomana M, van Velden FH, Boellaard R, et al. Bias reduction for low-statistics PET: maximum likelihood reconstruction with a modified Poisson distribution. *IEEE Trans Med Imaging*. 2015;34(1):89.
2. van Velden FH, Kloet RW, van Berckel BN, Wolfensberger SP, Lammertsma AA, Boellaard R. Comparison of 3D-OP-OSEM and 3D-FBP reconstruction algorithms for high-resolution research tomograph studies: effects of randoms estimation methods. *Phys Med Biol*. 2008;53(12):46.
3. Cheng J-C, Rahmim A, Blinder S, Camborde M-L, Raywood K, Sossi V. A scatter-corrected list-mode reconstruction and a practical scatter/random approximation technique for dynamic PET imaging. *Phys Med Biol*. 2007;52(8):87.
4. Watson CC. New, faster, image-based scatter correction for 3D PET. *IEEE Trans Nuclear Sci*. 2000;47(4):834.
5. Caldeira L, Herzog H, Ferreira HA, Shah NJ. CHAPTER 14 Parametric Imaging. *Hybrid MR-PET Imaging Syst Methods Appl R Soc Chem*. 2019;8:288–99.
6. Otani T, Hosono M, Kanagaki M, Onishi Y, Matsubara N, Kawabata K, et al. Evaluation and optimization of a new PET reconstruction algorithm, Bayesian penalized likelihood reconstruction, for lung cancer assessment according to lesion size. *AJR Am J Roentgenol*. 2019;213(2):76.
7. Rogasch JM, Suleiman S, Hofheinz F, Bluemel S, Lukas M, Amthauer H, et al. Reconstructed spatial resolution and contrast recovery with Bayesian penalized likelihood reconstruction (Q.Clear) for FDG-PET compared to time-of-flight (TOF) with point spread function (PSF). *EJNMMI Phys*. 2020;7(1):960.
8. Tragardh E, Minarik D, Brolin G, Bitzen U, Olsson B, Oddstig J. Optimization of [(18)F]PSMA-1007 PET-CT using regularized reconstruction in patients with prostate cancer. *EJNMMI Phys*. 2020;7(1):410.
9. Lammertsma AA, Hume SP. Simplified reference tissue model for PET receptor studies. *Neuroimage*. 1996;4(3 Pt 1):61.
10. Varrone A, Sjöholm N, Eriksson L, Gulyas B, Halldin C, Farde L. Advancement in PET quantification using 3D-OP-OSEM point spread function reconstruction with the HRRT. *Eur J Nucl Med Mol Imaging*. 2009;36(10):2100.
11. Brambilla CR, Scheins J, Issa A, Tellmann L, Herzog H, Rota Kops E, et al. Bias evaluation and reduction in 3D OP-OSEM reconstruction in dynamic equilibrium PET studies with 11C-labeled for binding potential analysis. *PLoS One*. 2021;16(1):49.

12. Régio Brambilla C, Veselinovic T, Rajkumar R, Mauler J, Orth L, Ruch A, et al. mGluR5 receptor availability is associated with lower levels of negative symptoms and better cognition in male patients with chronic schizophrenia. *Hum Brain Mapp*. 2020;6:71.
13. Akkus F, Ametamey SM, Treyer V, Burger C, Johayem A, Umbricht D, et al. Marked global reduction in mGluR5 receptor binding in smokers and ex-smokers determined by [^{11}C]ABP688 positron emission tomography. *Proc Natl Acad Sci U S A*. 2013;110(2):433.
14. Chapman S, Ragg M, McGeechan K. Citation bias in reported smoking prevalence in people with schizophrenia. *Aust N Z J Psychiatry*. 2009;43(3):87.
15. Wing VC, Wass CE, Soh DW, George TP. A review of neurobiological vulnerability factors and treatment implications for comorbid tobacco dependence in schizophrenia. *Ann N Y Acad Sci*. 2012;12:48.
16. DuBois JM, Rousset OG, Rowley J, Porras-Betancourt M, Reader AJ, Labbe A, et al. Characterization of age/sex and the regional distribution of mGluR5 availability in the healthy human brain measured by high-resolution [^{11}C]ABP688 PET. *Eur J Nucl Med Mol Imaging*. 2016;43(1):76.
17. Akkus F, Treyer V, Ametamey SM, Johayem A, Buck A, Hasler G. Metabotropic glutamate receptor 5 neuroimaging in schizophrenia. *Schizophr Res*. 2017;6:183.
18. First MB, Spitzer RL, Gibbon M, Williams JB. Structured clinical interview for DSM-IV-TR axis I disorders, research version. New York: SCID-I/P; 2002.
19. Herzog H, Langen KJ, Weirich C, Rota Kops E, Kaffanke J, Tellmann L, et al. High resolution BrainPET combined with simultaneous MRI. *Nuklearmedizin*. 2011;50(2):84.
20. Rota Kops E, Herzog H, Shah NJ. Comparison template-based with CT-based attenuation correction for hybrid MR/PET scanners. *EJNMMI Phys*. 2014;1(Suppl 1):47.
21. Régio Brambilla C, Veselinović T, Rajkumar R, Mauler J, Matusch A, Ruch A, et al. mGluR5 binding changes during a mismatch negativity task in a multimodal protocol with [^{11}C]ABP688 PET/MR-EEG. *Transl Psychiatry*. 2022;12(1):419.
22. Scheins J, Brambilla CR, Mauler J, Rota KE, Tellmann L, Lerche CW, et al. Image-based Motion Correction for the Siemens Hybrid MR/BrainPET Scanner. Bremen: 57 Jahrestagung der Deutschen Gesellschaft für Nuklearmedizin; 2019.
23. Hammers A, Allom R, Koepp MJ, Free SL, Myers R, Lemieux L, et al. Three-dimensional maximum probability atlas of the human brain, with particular reference to the temporal lobe. *Hum Brain Mapping*. 2003;19(4):84.
24. Akkus F, Mihov Y, Treyer V, Ametamey SM, Johayem A, Senn S, et al. Metabotropic glutamate receptor 5 binding in male patients with alcohol use disorder. *Transl Psychiatry*. 2018;8(1):46.
25. De Lorenzo C, Gallezot JD, Gardus J, Yang J, Planeta B, Nabulsi N, et al. In vivo variation in same-day estimates of metabotropic glutamate receptor subtype 5 binding using [^{11}C]ABP688 and [^{18}F]FPEB. *J Cereb Blood Flow Metab*. 2017;37(8):83.
26. Smart K, Cox SML, Nagano-Saito A, Rosa-Neto P, Leyton M, Benkelfat C. Test-retest variability of [^{11}C]ABP688 estimates of metabotropic glutamate receptor subtype 5 availability in humans. *Synapse*. 2018;6:719.
27. Cousineau D. Representing error bars in within-subject designs in typical software packages. *Quant Methods Psychol*. 2014;3:10.
28. De Lorenzo C, Della GN, Bloch M, Sanacora G, Nabulsi N, Abdallah C, et al. In vivo ketamine-induced changes in [^{11}C]ABP688 binding to metabotropic glutamate receptor subtype 5. *Biol Psychiatry*. 2015;77(3):418.
29. Adolfo AB, Ahn ACG, Tidey JW. Effects of smoking cues on caffeine urges in heavy smokers and caffeine consumers with and without schizophrenia. *Schizophr Res*. 2009;107(2–3):876.
30. Baguelin-Pinaud A, Robert S, Menard JF, Thibaut F. Prenatal exposure to tobacco and risk for schizophrenia: a retrospective epidemiological study. *Compr Psychiatry*. 2010;51(2):547.
31. de Leon J, Diaz FJ. A meta-analysis of worldwide studies demonstrates an association between schizophrenia and tobacco smoking behaviors. *Schizophr Res*. 2005;76(2–3):781.
32. Reed AC, Harris JG, Olincy A. Schizophrenia, smoking status, and performance on the matrices cognitive consensus battery. *Psychiatry Res*. 2016;9:246.
33. Riala K, Hakko H, Isohanni M, Jokelainen J, Weiser M, Rasanen P. Poor premorbid school performance is associated with later cigarette smoking among schizophrenia patients. *Psychiatry Res*. 2005;137(1–2):87.
34. Quigley H, MacCabe JH. The relationship between nicotine and psychosis. *Ther Adv Psychopharmacol*. 2019;8:9.
35. Rabin RA, Sacco KA, George TP. Correlation of prepulse inhibition and Wisconsin Card Sorting Test in schizophrenia and controls: effects of smoking status. *Schizophr Res*. 2009;114(1–3):69.
36. Lohmann P, Kocher M, Ruge MI, Visser-Vandewalle V, Shah NJ, Fink GR, et al. PET/MRI radiomics in patients with brain metastases. *Front Neurol*. 2020;9:11.
37. Carrasco-Poves A, Ruiz-Espana S, Brambilla CR, Neuner I, Rajkumar R, Ramkiran S, et al., editors. Analysis of New Biomarkers for the Study of Schizophrenia Following a Radiomics Approach on MR and PET Imaging. 2022 44th Annual International Conference of the IEEE Engineering in Medicine & Biology Society (EMBC); 2022 11–15 July 2022.
38. Akkus F, Treyer V, Johayem A, Ametamey SM, Mancilla BG, Sovago J, et al. Association of long-term nicotine abstinence with normal metabotropic glutamate receptor-5 binding. *Biol Psychiatry*. 2016;79(6):57.

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