

PET/MRI: reliability/reproducibility of SUV measurements

ZAIDI, Habib, BURGER, Irene A.

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Habib Zaidi and Irene A. Burger

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6.1 The Standardized Uptake Value (SUV) Metric

Qualitative visual interpretation of PET images is commonly performed to report abnormal tracer uptake in suspected regions. This is usually achieved by comparing observed patterns to expected normal biodistribution. However, the human visual system bears a number of limitations, and as such, this approach intrinsically suffers from intra- and interobserver variability owing to the subjective nature of visual interpretation [1]. Subsequently, guidelines on interpretation of PET images in clinical oncology advocated the adoption of simplified versions of PET metrics including semiquantitative indices, such as the standardized uptake value (SUV) [2].

H. Zaidi (✉)

Division of Nuclear Medicine and Molecular Imaging, Geneva University Hospital,
Geneva, Switzerland

e-mail: Habib.Zaidi@hcuge.ch

I.A. Burger

Department of Nuclear Medicine, University Hospital Zurich, Zurich, Switzerland

e-mail: Irene.Burger@usz.ch

Several methods for measuring PET tracer accumulation were proposed in the literature. The base is to measure the in vivo radioactivity concentration in the suspected malignant lesion (kBq/mL), which is directly linked to the tracer concentration. The two most significant sources of variation of tracer accumulation are the injected dose and the body weight representing whole-body distribution volume [1]. In practice, the SUV is calculated by dividing the decay-corrected activity concentration in the volume of interest (VOI) drawn around the lesion (MBq/mL) by the injected dose (MBq) divided by the body weight (g) [3]:

$$SUV = \frac{\text{Mean ROI concentration (MBq / mL)}}{\text{Injected dose (MBq)} / \text{Body weight (g)}} \times \frac{1}{\text{decay factor}} \quad (6.1)$$

The SUV metric is widely used in the clinic because of its simplicity, ease of use, reproducibility, and compatibility with conventional whole-body PET/CT acquisition protocols, requiring only a static scan as opposed to full kinetic modeling approaches, which require complex dynamic studies and arterial blood sampling. Virtually all commercial and open-source medical image display software platforms offer the option to measure SUVs. However, considerable inconsistencies have been reported among the different software packages used in clinical and research settings as demonstrated in a recent study conducted by the PET technical committee of the Quantitative Imaging Biomarker Alliance initiative [4]. It should also be noted that most packages normalize SUV to patient’s body weight (Eq. (6.1)). However, since adipose tissue is not as metabolically active as other tissues, other variants were suggested, including normalization to lean body mass (SUV_{LBM} or SUL) [5] or body surface area (SUV_{BSA}) [6].

The maximum SUV (SUV_{max}), representing the highest voxel SUV value, and mean SUV (SUV_{mean}), representing the average SUV across all voxels in a defined VOI, are undoubtedly the most widely used semiquantitative metrics (Fig. 6.1). Conversely, SUV_{peak} (Fig. 6.1), defined in PERCIST criteria as representing the SUV_{mean} in a

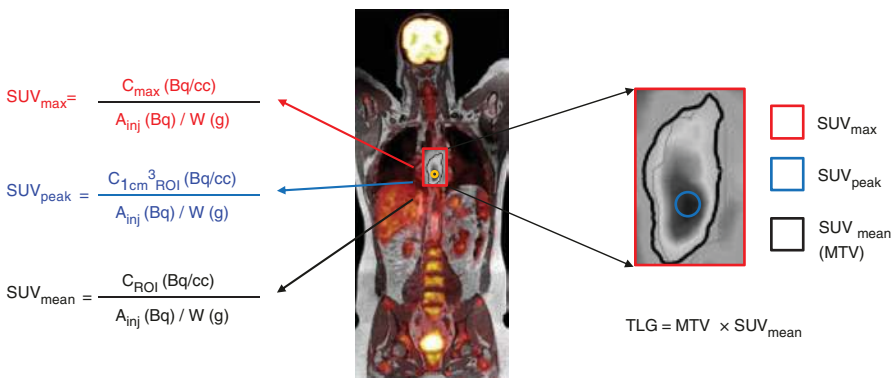


Fig. 6.1 Illustration of the basic foundations of PET quantification and the factors involved in the calculation of first- and second-order image-derived PET metrics used in clinical oncology

spherical VOI (1.2 cm diameter or 1 mL volume) placed over the most active part of a malignant lesion, was advocated as a more robust semiquantitative metric less vulnerable to artifacts [7]. Depending on the defined VOI, a variety of definitions of SUV_{peak} can be envisaged, obviously significantly affecting the resulting values [8].

6.2 Limitations of the SUV Metric

Despite its popularity and wide adoption in clinical and research settings, the SUV metric intrinsically bears a number of shortcomings which has limited its adoption in large clinical multicenter trials. The SUV also depends on the time course of the activity concentration in the blood plasma and the time point of acquisition and therefore imposes strict standardization of the uptake time, usually to 60 min with an acceptable range of 55–75 min according to the EANM guidelines 2015 [2]. Also the dependency of SUV on acquisition frame times, reconstruction parameters, physical and physiological factors, and scanner calibrations limits its potential in providing an objective assessment of whole-body PET images across baseline and follow-up studies for robust disease monitoring. Therefore, large multicenter clinical trials require standardization of data acquisition and processing protocols on different scanners for harmonization of PET quantification to enable pooling of data collected at different centers [9, 10].

To overcome the limitations of the SUV metric, alternative measures have been proposed including an uptake time-corrected version of the SUV including an uptake time quotient to normalize to 75 min [11] or simplified population-based blood pool activity-corrected estimations of FDG uptake, both trying to overcome the strong variability of SUV from uptake time and blood pool activity as approximations for more robust PET quantification [12]. The underlying principle for calculation of the glucose influx into the cell (MRGlu) was suggested about 25 years ago [13], but the fact that dynamic acquisition is needed did not render it feasible for whole-body imaging [14] until the advent of whole-body parametric imaging [15], which is now receiving considerable attention by the molecular imaging community. It can be anticipated that the additional information provided by parametric imaging through exploitation of the 4D spatiotemporal nature of the complete list-mode PET data, beyond the currently established semiquantitative SUV metric, might be valuable in whole-body PET imaging for multiparametric assessment of metastatic tumors across multiple beds and for improved reproducibility and evaluation of response to treatment over long periods [15].

Furthermore, volume-based PET metrics such as the total lesion glycolysis (TLG), calculated by multiplying SUV_{mean} by the metabolic tumor volume (MTV) [16], radiomic and texture analysis [17, 18], and parametric imaging have been suggested [14, 15]. The TLG, used to assess global metabolic response of the whole lesion thus providing complementary information to SUV and its variants, was shown to be highly correlated with other PET response parameters and is reproducible [19]. Recent advances in PET image segmentation and delineation of lesion contours [20] combined with progress in partial volume correction techniques have enabled to automate the calculation procedure. More recently, radiomics and texture

analysis emerged as new promising approaches enabling to circumvent the limitations of the above described oversimplified approaches by providing additional features including intratumoral heterogeneity through advanced image processing techniques and knowledge in systems biology [17, 18]. An increasing number of pioneering studies support the underlying assumptions of these hypotheses; however, further research and development efforts using large clinical databases are still required before these approaches can translate to valuable and reliable tools that can be adopted in the clinic.

6.3 Repeatability of SUV Measurements

A number of studies investigated the reproducibility of multiple PET metrics including SUV_{max} , SUV_{mean} , TLG, and MTV. All publications conclude that PET metrics are reproducible [21], with an almost perfect interobserver agreement for SUV_{max} [22, 23]. Furthermore, numerous studies showed that SUV_{max} has a prognostic value and correlates with progression-free and overall survival [24, 25]. This led to the widespread use of SUV_{max} as predictive and prognostic value for oncology PET assessments supported by the recommendations of the European Association of Nuclear Medicine guidelines 1.0 for PET/CT [26]. Six years later, the updated version 2.0 still recommends reporting SUV_{max} with the addition of SUV_{peak} to quantify tumor activity. SUV_{peak} is using a 3D 1.2 cm diameter (and 1.0 mL volume) spherical VOI positioned such that the average value across all positions within the lesion is maximized [7]. This alternative measure of the highest activity within a tumor was introduced due to the major limitation of SUV_{max} , that is, the high variability introduced by the statistical noise associated with a single voxel analysis. As has been shown in phantom and dynamic patient studies, this statistical noise substantially impairs the repeatability of SUV_{max} [27–29]. Averaging the values of a number of voxels within a given VOI, as proposed by SUV_{peak} [30, 31], can reduce this noise and therefore increases the repeatability without significant reduction of reproducibility. Using SUV_{peak} rather than an averaged number of the hottest voxels in a VOI has the advantage of a standardized volume irrespective of reconstruction methods and voxel sizes. However, the absolute activity drops compared to an averaged number of the hottest voxels in a VOI since not all voxels within SUV_{peak} have a high activity, which might impair the discrimination between high-activity lesions [32].

6.4 Clinical Relevance of the SUV

The use of FDG PET for therapy response assessment and evaluation of tumor aggressiveness is increasing. Most publications using PERCIST 1.0 as the base for PET-based therapy response assessment for solid and nonsolid tumors are based on SUV_{max} or SUV_{peak} [7, 33, 34]. PERCIST 1.0 suggested a cutoff at 30% increase for progressive disease and 30% decrease for partial remission, which is a slight increase in difference compared to the 25% according to the 1999 EORTC recommendation. A study comparing both evaluation systems for response assessment in

metastatic colorectal cancer came to the conclusion that they were equivalent [35]; however, several studies comparing CT-based RECIST with FDG quantification found that PET was superior in predicting histopathological therapy response [36, 37], time to progression [38], or overall survival [39].

Besides the increasing use of FDG PET for therapy response assessment, baseline quantitative PET metrics are also increasingly used to predict outcome. Initially these were predominantly based on SUV_{max} [24, 25, 40]. However, more and more publications come to the conclusion that volume-based PET metrics, such as MTV or TLG, are superior prognostic markers on baseline PET scans compared to SUV_{max} [41–43].

For cardiac FDG-PET evaluation, the absolute SUV values are less important than the relative difference within the cardiac wall using polar maps normalized to the peak activity. Here a direct comparison of the relative activity in the 20 cardiac segments between PET/CT and PET/MR showed an excellent correlation between both modalities that was even slightly improved for PET/MR with time-of-flight (TOF) capability (mean -1.3%) compared to non-TOF PET/MR (mean -2.1%) [44].

Analog to cardiac PET, also quantification of neuro-PET is rather performed based on tumor-to-background (TBR) ratios than absolute SUV values. The first study comparing PET quantification between PET/CT and PET/MR was performed for ^{11}C -methionine PET for gliomas and ^{68}Ga -DOTATOC in meningiomas. The authors concluded that the computed TBR exhibited an excellent accordance between PET/MR and PET/CT systems, with a correlation coefficient of 0.98 and a mean relative error of 7.9% [45].

6.5 Clinical Studies Comparing SUV Measurements Between PET/CT and PET/MRI in Oncology

Since the introduction of fully integrated PET/MR systems, multiple studies have been published comparing quantitative and qualitative results between PET/CT and PET/MR. In most of these studies, a same-day protocol with a single injection was performed. The majority had a significant difference in uptake time between PET/CT (commonly injected 60 min after injection) and the secondary PET/MR scan (with uptake times from 135 min [46] up to 180 min [47]). Table 6.1 summarizes results of studies comparing SUVs estimated on PET/CT and PET/MR images. Despite the sometimes large differences between uptake times, most studies came to the conclusion that there was a high positive correlation for all SUV quantification metrics between both exams and that most of the differences were due to the variance in uptake time between both scans [44–48, 50, 51, 53, 54, 57, 59–61, 64]. In only two studies, some scans were performed on the PET/MR scanner first [52, 58], and in two studies, all PET/MR scans were performed before PET/CT [49, 56]. All other studies had a longer uptake time of 51–120 min (mean 81 min) for the PET/MR scans and showed a slight increase in SUV between both exams. Sher et al. performed 40 scans, 31 on the PET/MR system and 9 on the PET/CT first. They showed that SUV values were always higher on the second scan for malignant lesion further supporting that the observed difference in the other studies is rather due to the increasing uptake over time than different technology.

Table 6.1 Summary of results presented in the literature comparing SUVs estimated on PET/CT images to those measurements on PET/MR images

Authors (reference)	Year	Journal	Tracers	Patient number	PET/CT uptake	PET/MRI uptake	Indication	Scanner	Main conclusions
Boss et al. [45]	2010	JNM	¹¹ C-methionine, ⁶⁸ Ga-DOTATOC	10	60 min	90–120 min	Brain tumors	Biograph mMR	Tumor-to-reference tissue ratios with excellent correlation (0.98), with a mean relative error of 7.9%
Drzeżga et al. [48]	2012	JNM	¹⁸ F-FDG	32	86 min	140 min	Cancer general	Biograph mMR	High correlation between mean SUVs on PET/MRI and PET/CT in lesions ($r = 0.93$) and background tissue ($r = 0.92$)
Varoquaux et al. [49]	2014	EJNMMI	¹⁸ F-FDG	32	146 min	85 min	Head and neck	Ingenuity TF PET/MR	Excellent correlation between SUVs on both modalities, with an underestimation on PET/MRI as compared to PET/CT
Kershah et al. [50]	2013	MIB	¹⁸ F-FDG	40	66 min	117 min	Cancer general	Ingenuity TF PET/MR	High correlation for SUV values obtained from PET/MRI compared to those from PET/CT, as the reference standard

Wiesmüller et al. [51]	2013	EJNMMI	¹⁸ F-FDG, ⁸ F-FEC, ⁶⁸ Ga-D-TATE	43	60 min	140 min	Cancer general	Biograph mMR	Differences demonstrated in quantification of tracer uptake between PET/CT and PET/MRI were minor but statistically significant
Schäfer et al. [52]	2014	Radiology	¹⁸ F-FDG	20	60 min/118 min ^a	128 min/63 min ^a	Pediatrics	Biograph mMR	13.1% deviation of SUVs for bone marrow and 5% deviation for other tissues
Afshar-Oromich et al. [53]	2014	EJNMMI	⁶⁸ Ga-PSMA	20	60 min	180 min	Prostate cancer	Biograph mMR	Scatter correction on PET/MRI challenging and direct comparison of SUVs from PET/CT and PET/MRI needs to be conducted carefully
Al-Nabhani et al. [46]	2014	JNM	¹⁸ F-FDG, ⁶⁸ Ga-D-TATE, ¹⁸ F-FECH	50	56 min	135 min	Cancer general	Biograph mMR	PET data on both modalities were similar with significant correlation between mean SUVs

(continued)

Table 6.1 (continued)

Authors (reference)	Year	Journal	Tracers	Patient number	PET/CT uptake	PET/MRI uptake	Indication	Scanner	Main conclusions
Sachpekidis et al. [54]	2015	AJNMMI	^{18}F -FDG	30	60 min	120 min	Multiple myeloma	Biograph mMR	Significant correlation between the two techniques was demonstrated, despite the statistically significant differences in lesional SUV _s between PET/CT and PET/MRI
Queiroz et al. [55]	2015	PLoS One	^{18}F -FDG	75	na	na	Cancer general	SIGNA PET/MR	TOF-PET/MRI device requires significantly less activity to generate PET images with good-to-excellent image quality
Lyons et al. [56]	2015	AJR	^{18}F -FDG	35	108 min	61 min	Pediatrics	Ingenuity TF PET/MR	Good correlation for SUV on both modalities, PET/MRI underestimated SUV compared with PET/CT

Vontobel et al. [44]	2015	EJNMMI	¹⁸ F-FDG	23	60 min	106 min	Cardiac	SIGNA PET/MR	PET/MRI with and without TOF showed minimal underestimation of tracer uptake (-2.08 and -1.29%, respectively), compared to PET/CT
Iagaru et al. [57]	2015	Clin Nucl Med	¹⁸ F-FDG	36	74 min	161 min	Cancer general	SIGNA PET/MR	The mean SUV _{max} values were higher in PET/MRI than PET/CT for all lesions
Sher et al. [58]	2016	AJR	¹⁸ F-FDG	40	90 min/132 min ^b	71 min/124 min	Pediatric lymphoma	Ingenuity TF PET/MR	PET/MRI quantification strongly correlated with PET/CT, but the SUVs were not interchangeable
Freitag et al. [47]	2016	EJNMMI	⁶⁸ Ga-PSMA	26	60 min	180 min	Prostate cancer	Biograph mMR	The correlation between PET/MRI and PET/CT SUVs was linear in lymph nodes and bone metastases

(continued)

Table 6.1 (continued)

Authors (reference)	Year	Journal	Tracers	Patient number	PET/CT uptake	PET/MRI uptake	Indication	Scanner	Main conclusions
Sawicki et al. [59]	2016	JNM	^{18}F -FDG	121	61 min	113 min	Pulmonary nodules	Biograph mMR	Lesion size, SUV, and characterization correlate strongly between the two modalities
Atkinson et al. [60]	2016	Abdom Radiol	^{18}F -FDG	18	60 min	na	Lymphoma	Biograph mMR	SUV values were comparable with high positive correlation. Similar lesion detection rate
Xin et al. [61]	2016	Eur J Radiol	^{18}F -FDG	45	64 min	170 min	Abdominal/pelvic	Ingenuity TF PET/MR	High correlation between the two modalities in terms of SUV_{max} and SUV_{mean} in focal lesions
Karlberg et al. [62]	2016	EJNMMI Phys	^{18}F -FDG	Phantom	na	na	na	Biograph mMR	NECKs and sensitivity are higher for PET/MRI. Differences in image quality are most evident for challenging settings. PET/CT performance better owing to TOF capability

Seith et al. [63]	2016	Invest Radiol	¹⁸ F-FDG, ⁶⁸ Ga-DOTATATE, ¹¹ C-choline	66	na	na	Cancer general	Biograph mMR	MRI-based AC is accurate in most tissues (SUV deviations generally <10%). Underestimations in the bone and overestimation close to the lung can be pronounced
Afaq et al. [64]	2017	Clin Nucl Med	¹⁸ F-FDG	68	64 min	136 min	Lymphoma	Biograph mMR	SUV _{max} correlated significantly with a correlation coefficient of 0.84
Law et al. [65]	2017	Clin Nucl Med	¹⁸ F-FDG	100	63 min	116 min	Cancer general	Biograph mMR	Excellent correlation between SUVs in lesions on PET/MRI and PET/CT, across all body regions and in all tumor types studied

^aIn four patients PET/MRI scans were performed before PET/CT

^bFor nine scans PET/CT before PET/MRI

To investigate the influence of MR-based attenuation correction (AC) on absolute SUV, Seith et al. used the same PET dataset from an integrated PET/MR scanner and reconstructed it either with the tissue segmentation-based MRAC μ -map or the CTAC μ -map using a nonrigid registration of the CT to the MR-based μ -map. They showed that MR-based AC is very accurate in most tissues with SUV deviations of generally <10%. A systematic underestimation of SUV was only present in the bones, while some lesions close to the lung were overestimated [63]. These results are in line with earlier published simulation data showing a significant decrease of up to 30% in sclerotic lesion, after subtraction of all voxels with bone density from the CTAC μ -map (Fig. 6.3) [38].

A quantitative phantom study comparing image quality between TOF PET/CT (Siemens Biograph mCT) and non-TOF PET/MR (Siemens Biograph mMR) showed that the spatial resolution was similar for the two systems. Average sensitivity was higher for the mMR (13.3 kcps/MBq) compared to the mCT system (10.0 kcps/MBq), and peak noise equivalent count rate (NECR) was slightly higher for the mMR (196 kcps @ 24.4 kBq/mL) compared to the mCT (186 kcps @ 30.1 kBq/mL). Highest hot contrast for the smallest sphere (10 mm) was achieved with the combination of TOF and PSF on the mCT compared to the mMR. Overall the differences between both systems were mainly due to the TOF possibility on the mCT, which resulted in an overall better image quality, especially for the more challenging settings with higher background activity and small uptake volumes [62]. For the fully integrated PET/MR scanner with TOF, only a few studies were published comparing PET quantification on PET/CT and PET/MR until today [44, 55, 57]. A prospective trial comparing TOF PET/CT (Discovery D 690 PET/CT, GE Healthcare) and TOF PET/MR (SIGNA PET/MR) in 75 patients based on phantom NECR curves investigated the potential dose reduction for PET/MR with equivalent image quality. They concluded that a reduction of FDG activity of slightly more than 50% can be achieved thanks to improvements in detector geometry and technologies [55].

6.6 Reliability of SUV Measurements in PET/MRI

The challenges faced by quantitative PET/CT imaging have been investigated since the commercial availability of this technology more than 15 years ago, and several professional societies established committees and task groups (e.g., QIBA/RSNA, CQIE/ACRIN, QIN/AAPM, etc.) to support and promote the use of quantitative imaging biomarkers in the context of cancer staging and therapy response assessment. The deployment of hybrid PET/MRI in the clinic poses new challenges and additional difficulties to enable reliable, quantitative imaging biomarkers. The primary challenge is the lack of a robust MRI-guided attenuation correction particularly in whole-body imaging. Furthermore, partial volume and motion correction need to be considered to produce artifact-free and quantitative PET images, with robust and reliable quantitative indices for routine application and advanced tools for clinical and research applications. The bulk of quantitative PET/MRI research to date focused on addressing the challenges of MRI-guided PET attenuation correction. Three categories of MRI-guided attenuation correction techniques have emerged [66]. This

includes (1) segmentation-based approaches, which segment MR images into different tissue classes and assign predefined attenuation coefficients to each class, (2) atlas-based and machine learning techniques in which co-registered MR-CT Atlas pairs are used to derive a pseudo-CT image or to learn a mapping function that predicts the pseudo-CT from actual patient's MRI, and (3) the recently revisited joint emission and attenuation reconstruction algorithms or maximum likelihood reconstruction of attenuation and activity (MLAA), in which the attenuation map is estimated from emission or transmission data. Figure 6.2 shows a representative clinical FDG brain PET study comparing various attenuation correction strategies.

Segmented MRI-guided attenuation correction widely used on commercial PET/MRI systems suffers the lack of bones in the derived attenuation map, which induces underestimation of the SUV in the corresponding regions within or close

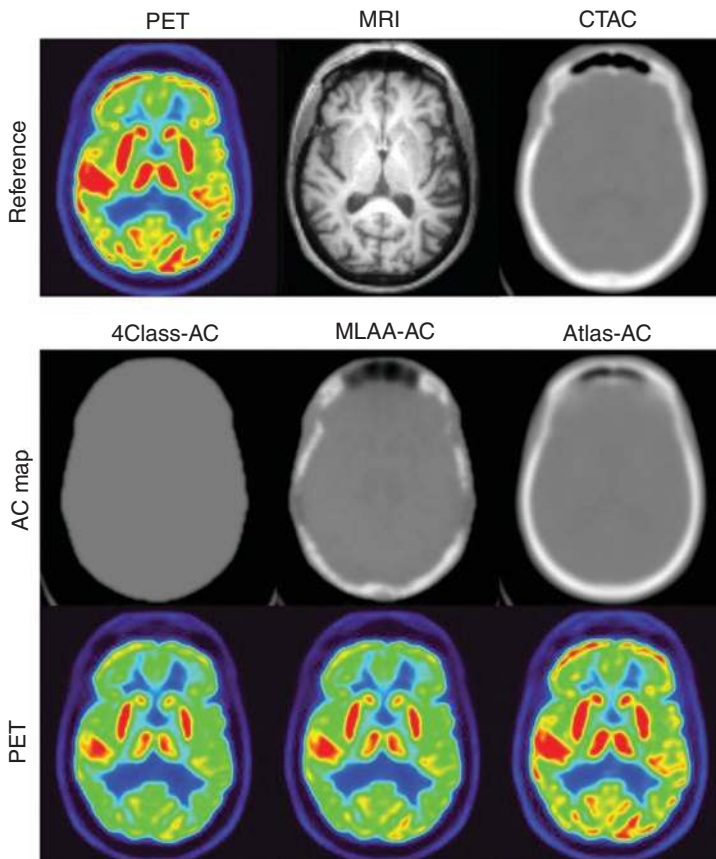


Fig. 6.2 Representative clinical FDG brain PET study showing from left to right: (top row) PET, MR, and CT images used as reference for evaluation, (middle row) attenuation maps derived using different attenuation correction approaches including segmented MRI, Atlas-guided, and maximum likelihood reconstruction of attenuation and activity (MLAA) technique, (bottom row) corresponding reconstructed PET images. Note the limitations of segmented MRI-based approach which ignores the skull and air cavities. The Atlas-based approach better matches the patient's CT image, clearly outperforming the MLAA algorithm

to the bones on PET images (Fig. 6.3) [67, 68]. Likewise, metal artifacts can lead to substantial signal voids and underestimation of SUV. This can be compensated by exploiting TOF reconstruction [69, 70], which was shown to reduce the SUV bias compared to non-TOF-PET emission data [71, 72]. Further improvement is possible with the joint emission-transmission reconstruction algorithm which showed promise in the correction of metal susceptibility artifacts by estimating the high attenuation coefficients of metallic implants [73]. Furthermore, the development of appropriate MR sequences, such as multi-acquisition variable-resonance image combination (MAVRIC) [74], can reduce metal artifacts. An additional complexity arises from the attenuation and scattering of annihilation photons by objects present in the field of view, which may also induce SUV underestimation if not accounted for. This includes patient's bed, MRI radiofrequency body or surface coils, and patient positioning aids [66]. Transmission or CT scanning-based predetermination of attenuation maps for rigid objects (bed, body coils, etc.) and nonrigid registration of templates of flexible objects (surface coils) [75] is currently used to account for the additional attenuation from these items. Joint reconstruction of emission and transmission images was purportedly promoted as potential approach enabling to estimate the attenuation maps of these objects. However, the performance of this approach for recovering objects lacking support from emission data remains to be demonstrated in clinical setting.

6.7 Summary

Objective quantification of PET tracer uptake is gaining significance with the increasing use of PET as a prognostic biomarker for therapy response assessment. Ease of use, availability, as well as excellent reproducibility and correlation with

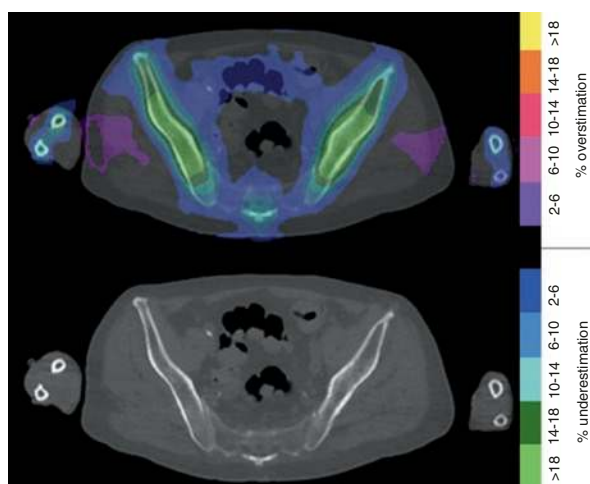


Fig. 6.3 Relative difference between PET images reconstructed with and without considering bony structures on CTAC attenuation map with the corresponding axial CT image

tumor aggressiveness lead to widespread use of SUV_{max} for tumor assessment, despite its well-known limitations, such as high variability with uptake time, blood pool activity, and limited repeatability.

With PET/MRI as a new hybrid modality being now commercially available, additional challenges are faced for reliable PET quantification. Although numerous studies showed an overall high correlation among SUV measurements between PET/CT and PET/MR, impaired attenuation correction due to the missing density of bone and metal implants can substantially affect SUV values within the bones and adjacent structures. More research and development efforts are required to improve MRI-based attenuation correction using machine learning algorithms to create a pseudo-CT or through advanced priors for joint estimation of emission, and attenuation using TOF-PET data to correct the attenuation map is currently ongoing and will further improve reliable PET quantification for PET/MRI.

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