



Quantitative preclinical PET imaging: opportunities and challenges

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PET imaging of metabolism involves many choices, from hardware settings, software options to animal handling considerations. How to decide what settings or conditions to use is not straightforward, as the experimental design is dependent on the particular science being investigated. There is no single answer, yet there are factors that are common to all experiments that are the subject of this review. From physics to physiology, there are many factors to consider, each of which can have a significant impact upon measurements of metabolism *in vivo*. This review examines the most common factors related to all types of quantitative PET imaging.

Keywords: positron emission tomography (PET), quantification, sensitivity, resolution, SUV, small animal, preclinical, animal handling

INTRODUCTION

Biological discovery has moved at an accelerated pace in recent years, with a significant focus on the transition from *in vitro* to *in vivo* models. With animal models widely used in the basic and preclinical sciences, finding ways to conduct animal experiments more accurately and efficiently becomes a key factor in the success and timeliness of research. As a result, there has been a considerable increase in the need to adapt clinical imaging methods, as well as for novel preclinical molecular imaging technologies for biomedical research. Molecular imaging technologies use molecular probes or interactions to visualize *in vivo* processes. Many of the traditional clinical medical imaging technologies, such as ultrasound, computed tomography (CT), magnetic resonance imaging (MRI), single-photon emission computed tomography (SPECT), and positron emission tomography (PET) have been adapted for imaging small laboratory animal models [1].

Animal models of human disease are an important research tool in understanding disease processes and developing novel treatment strategies. Imaging technologies offer the possibility to study biochemical processes in a non-invasive manner *in vivo*. The biggest advantage of *in vivo* imaging is that it provides a bridge from preclinical research to human application in the clinic, enabling similar and sometimes identical experiments to be carried out across species. In preclinical research, most often the mouse is the experimental model of choice, due to many factors, including fast breeding cycle and decreased housing and maintenance costs. A great advantage of mice is the relatively high genetic homology with man and highly developed methodology for genetic manipulation. This has led to the development of wide range of transgenic laboratory animals with customized genetic expression. The most efficient way of utilizing such models is through longitudinal studies of the same animals over an extended period of time. Non-invasive imaging technologies have

proven to be extremely valuable tools in performing such studies and have created the recent surge in small animal imaging [2]. Moreover, the possibility to use nude or severe combined immunodeficiency (SCID) mice with xenografts of human tissue make the immunodeficient mouse model particularly attractive for oncology research. In some disciplines, primarily in neuroscience, rats are more favorable as animal models. The larger brain size means greater stereotactic accuracy in models involving surgical or other invasive procedures directed toward specific brain regions (e.g., Parkinson rat model) and better identification of structures in images that have limited resolution.

Small animal PET is increasingly used to assess efficacy of therapeutic interventions and to validate the application of both current and novel radiotracers for translation to humans [3]. The strength of PET imaging is that it offers quantifiable measures of the underlying biology. The quantitative units given in either percent injected dose per gram tissue (%ID/g) or in standardized uptake values (SUV) are usually extracted from defined regions (or volumes) of interest drawn around an organ. Another more sophisticated method is the use of kinetic modeling to quantify the kinetics of the radiotracer in tissue [4, 5]. All these analytical methods rely on the fact that PET delivers quantitative and reproducible images. This review will look at all the factors that influence the accuracy of quantitative small animal PET imaging.

SCANNER SPECIFIC FACTORS

A variety of dedicated small animal PET scanners have been developed starting from the 1997 and are now commercially available from different vendors. The requirements on a PET detector are (a) high detection efficiency, (b) high spatial resolution, (c) low dead time, (d) good timing resolution, and (e) good energy resolution. These requirements led to the development of new scintillation materials such as LSO or LYSO crystals [6, 7], which

are now widely used in preclinical systems because of their high and fast light output. Most of the commercial preclinical scanners are based on small individual scintillators coupled to photomultipliers such as the microPET R4 and P4 scanners [8, 9], the microPET Focus family [10, 11], the Inveon scanner [12], the nanoPET/CT scanner [13], the Triumph®II –PET/SPECT/CT system [14], the Super Argus PET/CT [15], the ClearPET scanner [16] or the Genisys4 system [17, 18]. In contrast to these scanners, which are based on individual small scintillator pixels, the ALBIRA scanner is made of larger monolithic scintillators, where the point of interaction is estimated from the scintillation light spread into the position sensitive photomultiplier [19, 20]. This scanner offers the advantage of reduced dead areas (higher packing fraction), due to the missing edges between the crystals. A completely different scanner design was realized in the quadHIDAC small animal PET [21], where the detectors are formed by multiwire proportional chambers and lead converter plates.

In the following sections, several scanner specific factors that have an influence on quantitative accuracy are described.

SPATIAL RESOLUTION

In PET, spatial resolution is defined as the distance where two infinitely small point sources can still be distinguished from each other. Typical preclinical PET scanners exhibit a spatial resolution in the order of 1.5–2.5 mm measured at full-width at half maximum (FWHM) of the point spread function in the central field of view (cFOV). Spatial resolution is critically dependent on the radial offset, which was recently demonstrated in a study where the performance evaluation of several preclinical scanners was performed according to the NEMA NU 4-2008 standards using a ^{22}Na point source [22]. At a radial offset of 5 mm the radial spatial resolution for most of the studied scanners was around 1.6 mm FWHM whereas this value increased to around 2.5 mm FWHM at 25 mm radial offset. It was also demonstrated that scanners with a larger bore size suffer less from this radial elongation effect. In **Table 1** commercially available small animal PET scanners and the respective spatial resolution values are given.

By reducing the scintillator size from 5 mm (human scanners) to around 1.5 mm, the spatial resolution significantly improved. Still, when looking at the size difference between mice and humans, which is about 2000, a three-fold decrease in the spatial resolution is insufficient to create an equal sampling in mice as in humans. Thus, it still remains difficult to image small structure sizes of areas such as the mouse brain. This led to the development of scanners with even smaller scintillator sizes of only 0.975 mm [26, 27]. Smaller crystals means fewer counts per crystal, thus noise and signal strength become significant factors with this approach. Another issue in PET is that the scanner detects the point of annihilation and not the point of positron emission. The positron range in tissue is a function of the kinetic energy with which it is emitted from the nucleus, which is dependent on the specific isotope [28]. Any residual kinetic energy at the time of annihilation creates gamma radiation emitted at less than 180 degrees, resulting in non-collinearity and loss of accurate placement, which is also related to the size of the field of view. These factors create a fundamental limitation of PET spatial resolution; however there is still room to improve upon the current

PET system designs. Different resolution modeling approaches have been proposed to account for these physical effects [29, 30] yet they are only available in their respective academic institutions and are not commercially available. An example of how the positron range affects image quality is given in **Figure 1**. Here, a mini-Derenzo phantom consisting of hollow rods with 2.5, 2.0, 1.5, 1.25, 1.0, and 0.8 mm diameter are arranged in six segments, where the center-to-center distance is equal to two-times the diameter. The phantom was filled with either a ^{18}F -solution or a ^{68}Ga -solution. The kinetic energies of the emitted positrons are 633 keV (^{18}F) and 1.89 MeV (^{68}Ga), leading to a clear difference in mean positron range of around 0.5 mm and 3.1 mm and therefore different spatial resolutions [31].

SENSITIVITY

The sensitivity in PET is usually defined as the percentage between emitted and measured annihilation photons. The sensitivity of a PET scanner is influenced by the detector material, the dead space or packing fraction, the crystal length and the solid-angle coverage of the detector ring. Scanners with longer axial FOVs and smaller bore diameters usually exhibit the highest sensitivity. This can also be seen in **Table 1**, where the Inveon scanner with the rather long axial FOV of 127 mm (161 mm ring diameter) has a reported sensitivity of 6.72%, whereas the Genisys4 scanner with the smaller opening of 50 mm combined with a 94 mm axial FOV exhibits an even higher sensitivity of 14%.

The sensitivity plays an important role when imaging receptor binding probes, where low numbers of receptors are present and low occupancy rates are needed to avoid pharmacological effects, thus a low injectable mass of the tracer is required. Depending on the specific activity and the radiochemical yield, this can lead to rather low radioactivity concentrations in the injectable volume of 100 μL (some kBq/mL) and thus a high sensitivity is needed.

There is also a tradeoff between sensitivity and resolution. Smaller crystals enable better resolution; however this means there are fewer detected events in each individual crystal, which leads to noisy data and loss of image contrast.

SCATTER

In human PET scans, scatter is a dominant factor with scatter fractions of up to 36% in septaless full 3D-mode scanners [32]. In small animal PET scanners, the scatter fraction is usually in the range of $\sim 8\%$ for a mouse and $\sim 20\%$ for a rat scan [22]. One way to increase the throughput of small animal imaging facilities and also to optimize utilization of a radiochemical synthesis batch (especially for C-11) is to image multiple animals at once [33]. Several centers are now imaging 2 or up to 8 mice in one scan [34, 35]. In a recent simulation study [36] it was demonstrated that the scatter fraction increased by 25% or 64%, when imaging three or five subjects (MOBY phantom) simultaneously in comparison to one subject, respectively. This clearly demonstrates that scatter correction in preclinical imaging is important when imaging multiple subjects at the same time.

Scatter and also attenuation correction algorithms rely on the availability of structural information, which can be obtained from transmission scans using positron ($^{68}\text{Ge}/^{68}\text{Ga}$) or single photon (^{57}Co) emitters or from MR or CT scans. The noise level in

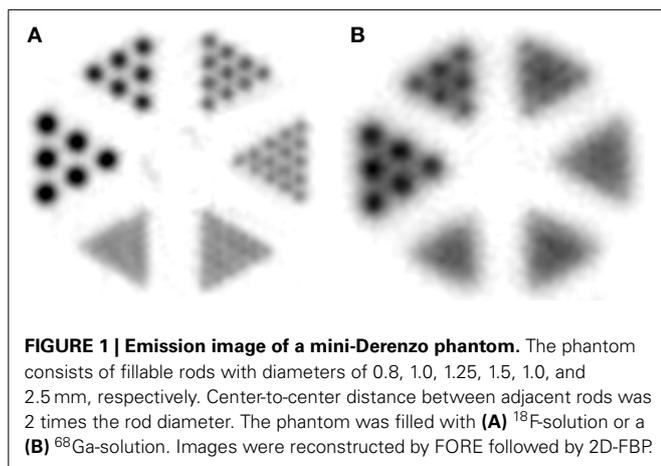
Table 1 | Characteristics of preclinical PET scanners.

	Inveon Siemens	Mosaic HP Philips	ClearPET Raytest	Argus Sedecal	Genisys4 Sofie biosciences	NanoPET/CT Mediso	ALBIRA Bruker
Detector material	LSO	LYSO	LYSO/LuYAP	LYSO/GSO LYSO/GSO	BGO	LYSO	LYSO monolithic
Crystal dimension (mm ²)	1.51 × 1.51 × 10	2 × 2 × 10	2 × 2 × 10+ 2 × 2 × 10	1.45 × 1.45 × 7 (LYSO); 1.45 × 1.45 × 8 (GSO)	1.8 × 1.8 × 7	1.12 × 1.12 × 13	40 × 40 × 10
Ring diameter (mm)	161	197	135–225	118	50+	181	111
Axial FOV (mm)	127	119	110	48	94	94.8	40
Energy window (keV)	350–625	385–665	250–650	250–700	150–650	250–750	350–650
Peak detection efficiency (%)	6.72	2.83	3.03	4.32	14	7.7	2
Transaxial FWHM resolution at 5 mm (mm)	1.64	2.34	2.02	1.66	1.4*	~1.6	1.55*
Reference	[12]	[23]	[24]	[25]	[18]	[13]	[19]

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+ Corresponds to the opening of the PET scanner and not the ring diameter, as the scanner is built up of two flat-panel type detector heads.

*Spatial resolution measured in the central FOV



MR and CT scans is considerably lower than nuclear medicine based methods, leading to more accurate estimations of scatter and attenuation corrections.

ATTENUATION CORRECTION

Attenuation of photons absorbed when passing through the subject and surrounding material (e.g., imaging chamber, bed) is usually measured using a transmission scan. In small animals, the amount of tissue through which the 511 keV photons pass

is much smaller as compared to humans, but nevertheless there is a certain percentage that is lost. For example, photons passing through the rat's torso are attenuated by ~40% [37]. A common method is to perform a transmission scan before or after the emission scan using an external source, which is rotating around the subject (rod or point source) [38]. A conventional approach is to use a ⁶⁸Ge/⁶⁸Ga source working in coincidence mode, where the detection of the two annihilation photons and the location of the source form a line of response (LOR). For small animal PET systems, the use of a single photon emitter source such as ⁵⁷Co, that emits photons at a lower energy (122 keV) as compared to the annihilation radiation has shown to be better suited for attenuation correction (in the absence of a CT). Here the LOR is defined by the known source location and the photons detected on the opposing detectors. The advantage of a singles-based approach is the higher detected photon flux and hence better signal-to-noise ratio, however this method requires accurate knowledge of the moving source location. As the energy of the emitted ⁵⁷Co photons is lower than 511 keV, the attenuation sinogram values are scaled up to 511 keV for correction [37]. When using a CT for attenuation correction, a similar energy scaling approach is required [38].

Another method for attenuation correction is based on the segmentation of the PET data, the so called transmissionless attenuation correction [39]. Here a manual or automatic delineation defines the body edges directly on the emission image, and by assuming a uniform distribution of attenuation coefficients

inside the subject an attenuation map is created [40, 41]. The advantages are that less time is required and no additional dose is given to the animals (from the CT or transmission scan). The disadvantages are that they are less accurate in delineating the subject (especially for radiotracers that are not distributed uniformly in the body), provide a less detailed attenuation map and they do not include the surrounding material like the imaging chamber. Similar to the rapid switch to PET/CT in clinical systems, most preclinical systems now utilize both metabolic (PET or SPECT) and anatomical (MR or CT) multimodality imaging as the standard protocol, using the anatomical information to provide scatter and attenuation corrections.

PARTIAL VOLUME EFFECT

Due to the limited spatial resolution of currently available preclinical PET scanners in comparison to the actual size of structures from rodents (e.g., regions in the brain), partial volume effects (PVE) are present in the acquired image data. PVE leads to an underestimation of the activity concentration in the PET images and spillover into surrounding tissues that is observed in objects smaller than three times the spatial resolution. In addition, image sampling where different types of tissue and thus radiotracer concentrations are included in one voxel also contributes to PVE [42–44]. There are several methods for partial volume correction based either on the count recovery models such as the recovery coefficient [45, 46] or the contrast recovery coefficient method [47–49] or they are based on anatomical data (derived from MR imaging) such as the geometric transfer matrix method [50].

Partial volume correction is an active area of research as most of the approaches are either not applicable for irregular structures (hippocampus region in the brain) or they have not been validated against the true activity values. It has also been shown that the recovery coefficient is strongly dependent on the count statistics, which is a problem for dynamic studies, where the count rate changes from time frame to time frame [51]. Together with the fact that the spatial resolution of the scanner and thus the PVE is not uniform in the field of view and that PVE is dependent on the radionuclide and reconstruction algorithm, this makes PVE correction an interesting but complex field of research.

RECONSTRUCTION ALGORITHM

For PET image reconstruction, several rebinning and reconstruction algorithms are available. In commonly used preclinical PET systems, the measured coincidences are stored in list mode format. Then the “raw data” is either converted into 2D sinograms by applying single-slice rebinning [52], multi-slice rebinning [53, 54] or Fourier rebinning [55] or into 3D sinograms. The depth of interaction (DOI) in which an annihilation event occurs inside the crystals can have a significant impact on image resolution, particularly for small ring systems, thus the positioning of oblique angle events is important. In single-slice rebinning (SSRB) the oblique coincidence lines are assigned to a particular 2D sinogram associated with the transaxial slice lying midway axially between the detectors. In multi-slice rebinning (MSRB) a more sophisticated rebinning algorithm is applied, where an oblique coincidence line contributes to the sinogram of all slices which it traverses. MSRB is more accurate than SSRB, but is less

stable in the presence of noise [55]. In Fourier rebinning (FORE) the oblique coincidences are binned to a transaxial slice using the frequency–distance relationship of the data in Fourier space. The reconstruction of the stack of sinograms is then performed using conventional 2D algorithms. The advantage of rebinning and further 2D reconstruction is that it speeds up reconstruction time.

The conventional 2D image reconstruction algorithms are filtered-backprojection (FBP) or 2D ordered-subsets expectation-maximization (OSEM). Apart from the 2D algorithms, fully 3D algorithms such as 3D-OSEM or 3D maximum a posteriori (MAP) [56, 57] or a combination of both can also be applied. The 3D reconstruction algorithms can either work iterative or non-iterative and process oblique coincidence lines without making geometrical approximations. The advantages of the 3D algorithms are that they provide images with better spatial resolution and thus an improved recovery rate as compared to FBP [58, 59], with the assumption that the noise level is the same. Iterative methods can also take into account the geometric and spatial characteristics of the scanner to improve resolution (P matrix). Iterative reconstruction also avoids mispositioning of activity outside the animals, creating images without streaking artifacts commonly seen around the bladder with F-18 labeled probes. Iterative reconstruction can create noise and artifacts from insufficient or too many iterations, thus knowing when to stop becomes a problem, one that is often specific to the imaging probe and scanner settings used to collect the data. Another example is for MAP reconstruction, where the spatial resolution is dependent on the choice of the β hyperparameter (parameter which controls the weight of the penalty term). By choosing a small β , one can obtain better spatial resolution at the cost of image noise and artifacts.

The advantage of FBP over MAP or OSEM is the possibility to define offsets in the x- and y-direction during reconstruction. Thus when imaging multiple animals, small voxel sizes can still be achieved just by using a high zoom and the respective offset values equivalent to the midpoint-position of the animals in the field of view. During MAP and OSEM reconstruction, an offset definition is not feasible and therefore an individual image generation for each animal cannot be achieved. The multiple animals can only be constructed using a rather low zoom thus leading to a larger voxel size. To exemplify this for the Inveon PET scanner (Siemens), for a single mouse scan the usual method for reconstruction is the use of an $128 \times 128 \times 159$ matrix combined with a zoom of 2 leading to a reduced radial field of view of 4.8 cm and a voxel size of $0.38 \times 0.38 \times 0.80 \text{ mm}^3$. For a multiple mouse scan (4 mice) the whole field of view is needed and therefore a zoom of 1 can be used with the $128 \times 128 \times 159$ matrix yielding a voxel size of $0.86 \times 0.86 \times 0.80 \text{ mm}^3$ [35]. This increased voxel size will lead to a lower spatial resolution and therefore a higher partial volume effect.

ANIMAL SPECIFIC FACTORS

As the animal is our “measurement system” in preclinical imaging, several animal specific factors that affect image quantification are discussed below. Overall, the biological variability even in the same strains of animals with the same sex and age is by far larger (~15–20%) than all the scanner related factors. Therefore, great

care should be undertaken to optimize all factors that lead to large standard deviations in animal research such as temperature, diet, anesthesia, injectable volumes and many others [60, 61].

ANESTHESIA

In small animal PET, anesthesia is usually necessary to render the subjects motionless for imaging and to minimize the stress and discomfort associated with restraining procedures. In addition, invasive surgical procedures are often required to allow administration of pharmacological agents or sampling of blood. As a result, a large number of PET imaging studies in rodents are performed under general anesthesia. For all studies it is assumed that anesthesia-induced perturbation must be minimized by using as low doses of anesthetic as possible. The anesthetic method is thus adjusted to the procedure to ensure the necessary level of anesthesia and also compatibility with the purpose of the research. In most cases, the method of anesthesia that was chosen needs to be reversible such that the subject recovers to normal activity for future studies. The most common anesthesia method in PET is the use of an inhalant anesthetic such as isoflurane mixed in oxygen or air. By constantly adapting the isoflurane level to the depth of anesthesia and using a humidifier, it is possible to achieve quite long anesthesia periods of up to several hours without adverse complications. Another method well suited for short imaging protocols is the use of injectable anesthetics such as ketamine or xylazine.

Yet, one has to be aware of the fact that by using anesthesia one is changing the “measurement system” which is the physiology of the animal. A recent study investigated the effects of anesthesia in brain PET research [62]. They showed that depending on the used anesthetics and animal, radiotracer uptake in brain was either lower (as for [^{18}F]FDG) or binding potentials were higher during isoflurane anesthesia (as for [^{11}C]Raclopride). As isoflurane alters myocardial glucose metabolism it also affects cardiology studies using [^{18}F]FDG [63–66]. Also in oncology, the anesthetic method has a pronounced effect on [^{18}F]FDG or [^{18}F]FLT uptake in tumors [67–69].

In addition to the anesthetic agent, the carrier gas can also have a profound effect on the outcome of a study. Typically, the carrier gas for isoflurane anesthesia is oxygen. For hypoxia studies, where the hypoxic regions are identified and quantified by using hypoxia radiotracer, the breathing conditions have a strong effect on radiotracer retention as shown by Maier et al. [70]. They demonstrated that during the first hour of [^{18}F]FAZA uptake the breathing conditions (air or oxygen) had an effect on tumor uptake. When the mice breathed air, the tumor-to-muscle-ratios were significantly higher as compared to the oxygen breathing protocols.

These factors have led to the development of systems that are capable of imaging conscious animals. One system for conscious rodent imaging is the RatCAP [71, 72], where a miniaturized PET camera is mounted directly on the head of rats. Other solutions for conscious PET imaging are the use of marker-based [73] and also marker-less head motion tracking systems [74]. The marker-less system simplifies motion-compensated imaging by avoiding irritation caused by the attachment of the markers, and also the need to familiarize animals to the markers. Both systems require

a certain period of time in animal handling for successful motion tracking imaging experiments (10 days handling, 12 days training phase using a food reward system to acclimatize the animals to the burrow and scanner environment). Another option is to inject conscious animals and wait until uptake and clearance are essentially complete, then anesthetize and image the animals.

TEMPERATURE

The most important factor when putting an animal under anesthesia is temperature control. Mice and rats quickly equilibrate their internal body temperature to the temperature of whatever surface they are placed upon when under anesthesia [75]. Therefore the animals become hypothermic and can easily die within minutes. Almost every metabolic process is altered by temperature changes, including enzymatic activity, blood flow, and muscle activity. Since the goal is usually to measure normal metabolism, not metabolism under hypothermic conditions, animals need to have sufficient heat support at all stages of the experiment. Thus rodents must be warmed throughout the whole imaging procedure starting around 30 min before induction of the anesthesia till the time when they have fully recovered.

HOUSING CONDITIONS

Often overlooked, the housing conditions for animals can have a significant impact upon physiology and therefore molecular imaging results. Since animals spend nearly all of their time in the vivarium, the conditions are the primary factors that influence the biology under investigation. The bedding amount, type, cage changing frequency, light/dark cycle, room temperature and humidity can affect physiology. In the past several years, caging system use has been shifting from static filter top to individually ventilated cages (IVC). These IVC cages allow for multiple health status and mouse types to be housed adjacent to each other, however they also impose a new source of cold stress upon the animals [76]. These are not small effects, as it was shown that the size of xenograft tumors and FDG uptake can be altered by over 300% based on caging type and ambient temperature [77]. At a minimum, the husbandry conditions need to be reported to enable reproduction of experimental results [78].

FASTING STATE

Because PET images metabolism, measurements can be dependent on the fasting state of the animals. This is particularly true for [^{18}F]FDG studies, where the fasting state has an influence on the study outcome as the endogenous available glucose competes with [^{18}F]FDG and can therefore reduce uptake into the cells [66–68]. The duration of the fast, type of food (high fat, special formulations, etc.) can alter factors such as glucose and amino acid levels, which in turn can alter PET metabolic signals.

POSTURE

The position of the animals inside the scanner is usually determined by the investigators and can affect the image results. Several different imaging chambers are available or provided by vendors that offer the possibility of a defined environment that includes anesthesia supply, heating and some fixation devices such as tooth bars or ear plugs [75]. Apart from using corner posts, where the

limbs can be fixated for a reproducible position, the use of an immobilization bed has also been proposed [79]. The posture of the animal (prone or supine) can have an effect on the outcome of the imaging study has been demonstrated by Richter et al. [80], where they studied regional pulmonary blood flow in rats. As mentioned previously, the position of the animal with respect to the imaging field of view can also change the data, due to factors such as DOI and resolution.

MONITORING

Animal Monitoring and supportive care aims to maintain the animal's physiological status as near to normal as possible. The goal is to minimize animal pain and distress by adjusting the depth of anesthesia and to anticipate any impending complication. Supportive care includes monitoring of physiological parameters and analgesia during anesthetic and surgical procedures. Monitoring of vital signs and potential signs of pain should be done throughout the whole animal handling procedure. Monitoring can either be performed visual, manually or by using electronic monitoring systems.

Visual and manual monitoring

Visual monitoring means watching the animal carefully while under anesthesia. By watching the respiration one can note gross change in rate and can classify the character of breathing. The color of mucous membrane and skin is related to the oxygenation state—blue (poor oxygenation), pale (poor blood perfusion). The general behavior of the animal is also monitored visually. Manually one can test the jaw tone just by moving the jaw down. Using palpation, one can feel the heart rate, respiration rate, and body temperature. By pressing mucous membrane or skin, blood flow status can be visually assessed by looking for resumption of color in the skin. Yet, in most preclinical scanners visual and/or manual monitoring is not possible due to the small bore sizes and restricted space inside the scanners.

Electronic monitoring

Nowadays, nearly all the equipment available for human monitoring is also available for small animal monitoring. These include:

- ECG: small electrodes glued to the paws and tail or invasive with needles
- Respiration rate: small pneumatic pillow placed under the abdomen of the animal
- Blood pressure: tail cuff sensor (non-invasive) or positioning of electrode into major artery (invasive)
- Pulse oxymetry: The clip-on-sensors are typically attached to the rat foot or the shaved mouse thigh. Other useful locations include the ankle, paw, wrist, or tail.
- Temperature: rectal probe

Using electronic monitoring is a very reliable tool and also enables monitoring the animals during scans. Some scanners (e.g., μ PET/CT combined systems) have a very long but narrow bore and therefore visual or manual monitoring is not possible. Acceptable limits for all the monitored parameters need to be established and constant monitoring of the anesthetic state

over time is essential. Some monitoring systems have alarm functions that help identify when problems arise, which is helpful during experiments where many demands are present for the investigator's time.

ADMINISTRATION ROUTES

As PET is based on the administration of a radioligand into a subject, the administration itself has to be performed reliably. Animals may be conscious or under anesthesia during the injection, which can influence the measurement as discussed in section Anesthesia. **Table 2** presents administration volumes for the commonly employed routes in the most frequently used species [81]. They are consensus figures based on published literature and internal guidelines. Two sets of values are shown in each column: values on the left are intended as a guide to “good practice” dose volumes for single or multiple dosing; values on the right define the recommended needle size for each administration route in the different animals.

When performing intravenous injections, highly trained personnel are required in order to deliver all the radioactivity reproducibly from the syringe into the blood stream. Otherwise, the error rate would be very high as shown by a recent study by Vines et al, where they evaluated mouse tail vein injections in a qualitative and quantitative approach [82]. They showed that the overall rate of successful intravenous injection was 92% (46 out of 50). If one is only interested in a static image at ~ 45 min post injection, then this error is not as dramatic as the residual activity will be slowly absorbed and transported into the blood stream. For a dynamic study where one is interested in the kinetics of the radiotracer, a paravenous injection severely confounds the results.

The issue with reliable tail vein injection in mice has also led to the development of automatic injection systems, such as the vascular access system (VAS) [83]. In this system, near infrared light, image processing techniques, computer controlled motors and a pressure feedback system was used to insert the needle and to validate the proper location within the tail vein. A study in mice showed that the residual activity inside the vein was around 3%, which was better than the manual injections.

For some tracers (such as FDG), the use of an intraperitoneal injection might be an alternative to avoid cannulation of the tail vein [84]. Especially when performing short static scans at around 45–60 min post injections, the activity concentrations in brain, myocardium, plasma, muscle, liver and also tumors are

Table 2 | Administration routes, volumes and needle sizes considered good practice.

	subcutaneous (mL kg ⁻¹)		intraperitoneal (mL kg ⁻¹)		intravenous (bolus) (mL kg ⁻¹)	
Mouse	10	25G	10	25–27G	5	26G–28G
Rat	10	25G	10	23G–25G	5	25G–27G
Rabbit	10	23–25G	5		2	23G–25G

Table adapted from Diehl et al. [81] and from the German Society for Laboratory Animal Science (GV SOLAS).

comparable [68, 85]. Still, care has to be taken not to inject the substance into the bowel.

In preclinical imaging another important point is the total mass of the injected radioligand. This is usually not a concern in human studies, but is much more important in mice. The injection of too much mass can alter the binding kinetics and therefore the results of the study. For *in vivo* binding studies in rodents, the maximal receptor occupancy to fulfill the tracer kinetic modeling principles should be 10% or even less [86, 87]. It is therefore important to work with high specific radioactivity (i.e., the amount of radioactive molecules vs. unlabeled molecules). A good example of the effect of specific radioactivity was given by Maeda et al. [88], who performed a study with high specific radioactivity of [^{11}C]PIB (~200 GBq/ μmol) in transgenic Alzheimer mice (APP23). Reducing the specific activity to around 20 GBq/ μmol by addition of cold ligand resulted in profound loss of signal-to-noise ratio and reduction in binding potential in the hippocampus and neocortex, along with insensitivity for detection of amyloid plaques.

Last, the residual activity in the syringe also can contribute to errors in quantitation. Due to the small volume (100 μL) and especially with lipophilic radiotracers that stick to the walls, a considerable amount of radioactivity might be present in the syringe and needle—even after a complete injection of the tracer. Therefore, the syringe (and any catheters) should be measured after injection to obtain the accurate amount of injected radioactivity.

BLOOD SAMPLING

In preclinical PET imaging blood sampling can be performed to generate an arterial input function for kinetic modeling, to measure the free fraction of radioligand in plasma and to check for radioactive metabolites. The amount of blood that can be sampled depends on the circulating blood volume present in each species. **Table 3** gives the circulating blood volumes and the recommended maximum blood samples volumes for the most commonly used species. The recommendations are based on published work [81, 89] and on information from “in-house” standard operating procedures. Animal welfare is a prime consideration when blood sampling is approaching limits, however the scientific impact of an animal’s physiological response also must be considered because this may affect data interpretation and validity [90].

There are several sampling sites that can be used to obtain blood samples (mostly at a single time), such as the lateral tarsal (saphenous) vein, the sublingual vein, the lateral tail vein, the retrobulbar plexus or directly by cardiac puncture from the heart. All these sampling methods should be carried out under general anesthesia.

For repeated blood sampling a catheterization of a major artery (e.g., carotid or femoral artery) is recommended [91]. Temporary catheters such as butterfly needles and over-the-needle cannulae can be used in the short term (working day), whereas for long-term use surgical implantation of a biocompatible catheter is required. These methods allow repeated blood sampling with minimal distress and discomfort for the animal.

Table 3 | Circulating blood volume and recommended maximum blood sample volume.

	Rat (250 g rat)	Mouse (25 g mouse)
Blood volume	64 mL/kg (range 50–70) 6.4% BW 16 mL	74 mL /kg (range 70–80) 6.6% BW 1.8 mL
Safe single sample	1.6 mL (10%) max 1.2 mL (7.5%)	0.2 mL (10%) max 0.1 mL (7.5%)
Continuous samples		
Day	0.2 mL	0.02 mL
Week	1.4 mL	0.14 mL

Table adapted from Diehl et al. [81] and from the German Society for Laboratory Animal Science (GV SOLAS). BW, body weight.

One confound for blood sampling is the time it takes to get a sufficient sample and the dispersion that occurs between the sampling site and the area of the animal being investigated. As the sampling site is typically distant from the investigation site, the blood activity may not match well with respect to concentration and time for the target region.

PET DATA QUANTIFICATION

The first steps for obtaining quantitative PET images are to apply the corrections factors that are usually implemented in the reconstruction procedure. In most of the scanners, deadtime and decay correction is already performed during data acquisition. Normalization, attenuation and scatter correction are done in sinogram space. To ensure reproducible data (even after long periods of time) a regular quality control of all these correction factors together with a calibration procedure to convert counts into kBq/cc using appropriate phantoms is advisable [78].

SEMI-QUANTITATIVE VALUES

There are two commonly used semi-quantitative methods used in preclinical PET. The first method is the ratio of the activity in a tissue (given in kBq/cc or mCi/cc) divided by the decay-corrected activity injected into the animal (given in kBq or mCi). This ratio is defined as the percent injected dose per cubic centimeter (or milliliter) in tissue (%ID/cc or %ID/mL). Commonly, a density of 1 g/cc in tissue is assumed and therefore the uptake is often defined as %ID/g. Another method which also includes the weight of the animal is the standardized uptake value (SUV) calculated as the %ID/cc multiplied by the weight (given in g). The resultant number is almost unitless (actually g/cc or g/mL) and is a crude measure of radiotracer uptake into tissue. There are numerous problems with the use of SUV, such as the dependency on body composition (fat vs. muscle), varying radiotracer uptake and imaging durations, endogenous competition and definition of the volume or region of interest (ROI), which are discussed in detail elsewhere [92, 93]. Often substantial residual activity may be present at the injection site and therefore not available for biodistribution, leading to inaccuracies in the injected dose and SUV values. This is

particularly true if the site is the tail and outside of the imaging FOV. Several correction schemes, involving blood glucose level, body surface area, and lean body mass have been proposed in clinical studies to try to improve the accuracy of the SUV method and are now being used at some clinical PET centers [94, 95].

In preclinical PET there are also several methods for the extraction of the SUV from the volume of interest (VOIs). The most commonly used method is the extraction of the mean SUV over a ROI or a VOI. When imaging tumors with necrotic cores, another way of quantification is the definition of a lower threshold such as the mean plus 1 *SD* [96] or the inclusion of voxels exhibiting >75% of maximum radioactivity [97, 98], as a way to exclude areas with low tracer uptake. Sometimes, the use of the max SUV or the “5 hottest voxel” is also reported. The problem in defining the VOI and extracting the SUV is that this value should be a reliable representative measure of radiotracer uptake and thus underlying biology of this specific region.

KINETIC MODELING

Tracer kinetic modeling that describes mathematically the mechanism of transport and biochemical reactions of the tracer in tissue is a method to extract transfer rate constants, volumes of distribution or binding potentials from the radiotracer to the target [99]. Most of the modeling approaches require taking a series of blood samples from the animal to provide the time course of the tracer delivery (called the input function) which is matched to the PET image of the dynamics of the radiotracer concentration in local tissue over time. By applying model fitting or regression analysis the desired biological information can be obtained. In compartment modeling, the model describes the behavior of the tracer mathematically using a number of compartments, each of which represents distinct anatomic, physiologic, or biochemical stage of the behavior of the tracer. Typically the model can be described using a set of linear first order differential equations.

Several methods for PET tracer kinetic modeling have been introduced and are now applied in clinical and preclinical research. Generally speaking they can be divided into model-driven methods [99–104] and data-driven methods [105–110]. The model-driven methods include compartmental models that require the radioactivity concentration in arterial blood or plasma, the so-called “input function,” and reference tissue models, where the target tissue time-activity curve is expressed as a function of the reference tissue time-activity curve. In contrast to the compartmental models based on the input function, which can be applied to all areas of research, reference tissue models are primarily used in neuroimaging [111]. The requirement for using a reference tissue model is that there must be a region in the brain that is devoid of receptors and representative of the blood time activity values. This region is then used as an indirect input function to the target region. The data-driven methods include graphical analysis or spectral analysis. Graphical analysis methods such as Logan and Patlak plots [105–108, 110] use a transformation of the data such that a linear regression of the transformed data yields the desired parameters. These methods are simple and elegant, yet they may be biased by statistical noise

[112], the definition of the starting point for the linear regression is variable and they also fail to provide information about the underlying compartmental structure. A more detailed description together with the mathematical background can be found elsewhere [113–115].

GUIDANCE ON IMAGING STUDIES

A good description on the methods used in preclinical studies is given by Stout et al. [78]. This paper focuses on the methods descriptions in manuscripts, however it can also be used as a “how to do imaging” guide as well. Quantitative preclinical PET is highly dependent on scanner and animal related parameters. Therefore, it is difficult to define parameters required to perform an adequate small animal PET imaging study that are valid for all studies, as each study design must be optimized based on the biological target, species used, radiopharmaceutical and scanner performance.

As an example, for a mouse [¹⁸F]FDG oncology study the following parameter settings seem to be optimal [68]:

- Static scan after 50–60 min
- 6–8 h fasting before [¹⁸F]FDG injection
- Warm animals before [¹⁸F]FDG injection and during the uptake period
- Isoflurane anesthesia during the injection and uptake period (optional) and during imaging (required)

The details on how the imaging study was performed are extremely important when it comes to replication or expanding on existing work from other laboratories. Standardization of the imaging procedure is particularly essential when pooling data acquired in different labs.

OPPORTUNITIES

Molecular imaging can help in understanding of biological processes behind a particular disease. In central nervous system (CNS) drug development, the use of translational imaging is essential for successful clinical proof-of-concept testing [116]. The potential of CNS PET imaging to translate the finding allows the assessment of target occupancy in preclinical species, before first-in-human safety and tolerability studies, to refine dose ranges for exploration in humans. Hargreaves et al. pointed out that PET imaging has been an important tool for investigating drug–target interactions and the pathophysiology of many diseases such as monoaminergic dysfunction in schizophrenia and depression. Especially studies in non-human primates allow a prediction of the behavior of a drug in humans due to the high genetic homology [117].

Another area of research where PET has a pivotal role is the application of molecular imaging to oncology. Using a variety of radiotracers, different diagnostic targets such as proliferation, hypoxia, cell surface molecules, tumor vasculature and oxygenation can be assessed [118]. An interesting research area is the use of imaging for the assessment of early drug response in “mouse avatars,” where sectioned patient tumor samples are implanted in immunodeficient mice for subsequent use in drug efficacy studies [119]. The avatars allow for each patient to have their own tumor

grow in an *in vivo* system, thereby enabling the identification of a personalized therapeutic concept. This avatar approach is a promising concept and has been shown in a murine lung cancer study that mirrored an ongoing human clinical trial in patients with KRAS-mutant lung cancers [120]. Using FDG-PET in mice and patients, they were able to predict antitumor efficacy of docetaxel, selumetinib or combined selumetinib-docetaxel in treating Kras-mutant lung cancers.

CONCLUSION

PET imaging of metabolism is an excellent research tool that requires careful consideration of both physical and metabolic factors in order to obtain meaningful and reproducible information. There are many different hardware and software options that require appropriate selection of options in order to obtain useful images. The use of animals needs to be carefully considered with respect to their physiological environmental support to obtain meaningful metabolic measurements. With proper attention to the many details involved with PET imaging, reproducible measurement that have translational potential can be obtained.

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