Review Article: Modern Trends in Imaging VII

Magnetic Resonance Microscopy

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1. Introduction

One of the driving forces for recent developments in magnetic resonance microscopy (MRM) has been the explosion of genetically engineered mouse models. These models have opened avenues for understanding the basis of human diseases, as well as the relationship between genotype and phenotype. In addition, the environment, which can also modulate the phenotype, as well as gene expression patterns, can be well controlled in rodents. It has therefore become essential to develop and improve upon existing noninvasive techniques that can follow the longitudinal progression of disease, as well as the effects of putative treatments.

Among imaging techniques, magnetic resonance imaging (MRI) is particularly well suited to phenotype rodent models of disease because of its unparalleled rich soft tissue contrast. Much progress has been shown in high-field magnetic resonance instrumentation, including gradient and radiofrequency (RF) coils, increased computing power and efficient software processing of large arrays, as well as animal and specimen preparation techniques. All these have allowed the acquisition of magnetic resonance images with spatial resolutions high enough to characterize changes in the volume and microstructure of mouse body organs. The acquisition of MR images at resolutions higher than 300 microns (µm) [65] has been defined as magnetic resonance microscopy (for

Because of its three-dimensional (3D) nature, MRM allows appreciating the rodent anatomy in fixed specimens free of spatial distortions inherent to optical histology [61, 130]. In contrast with traditional histology, the hydration of tissues is preserved in MRM, and both the whole body and then isolated organs can be imaged (at different resolutions to increase the level of observed detail), since MRM is nondestructive. While the inherently digital MRM images can be virtually resliced in any arbitrary plane, the spatial relationships and connections among anatomical structures, as well as their shapes remain unaltered. From a structural, qualitative perspective, MRM can exploit a plethora of contrast mechanisms to differentiate tissue types. In addition, quantitative data can complement the global picture one can obtain from MR. Quantitative volumetric measures and shape parameters can be used to characterize normal strains of mice, models of disease at distinct time points in the evolution of that disease, or treated mice versus controls. Atlases are being developed and made available online, to provide baseline estimates for the variability within a given population, as well to help phenotype mouse models of disease.

comprehensive treatment see [14]). However, it has now become routine to acquire images at resolution of, or higher than 100 µm [35, 108]. To resolve the same kind of anatomical detail in a mouse as in a human, MRM operates with much smaller voxel sizes (1 nl for 100 µm-sized cube, or 10 pl for a 22 µm-sized cube) compared to clinical imaging (1-µ, for a 1mm-sized cube); differences which translate into 4–5 orders of magnitude weaker signal coming from one voxel. To compensate for this signal loss, it has been necessary to develop specific hardware and software for image formation, reconstruction, and analysis.

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Perhaps the most important attribute of MRM in small animals is the ability to image the live animal longitudinally. Functional measures of the heart, lungs, and brain complement the picture of the anatomy. Besides normal physiology in the adult rodent, MRM has the potential to illustrate changes occurring with development and aging, as well as to characterize abnormal lesions from ischemia, to cancer, or the migration of stem cells. Because of its nondestructive nature, and the recent availability of multimodal probes, MRM can be now used in conjunction with other in vivo imaging modalities such as SPECT, PET, or CT, providing different probes of anatomy and function. One can perform nondestructive, volumetric analysis studies using MR and still have the possibility of performing optical histology studies afterwards.

Together, these attributes of MR make it a powerful method for 3D, spatially coherent, high-resolution imaging of fixed specimens, an invaluable tool for the study of living tissue, while holding great promise for functional measures (from blood flow, perfusion, to gas dynamics and brain function) and even molecular information [43, 106].

Among its numerous applications, MRM has been used intensively in the past 10 years to phenotype small animal models of human disease—to assess structural [62] and functional [13] phenotypes in the adult and developing mouse [22, 91], as well as for drug development [9], in toxicology [66], and genetics [5, 6]. This chapter starts by explaining the basic aspects of image formation in MRM, with an emphasis on the differences from clinical MRI configuration and demands. Mechanisms for contrast manipulation in MR and the benefits of contrast-enhancing agents are described. Given these techniques, also covered are the length of MR scans required to achieve sufficient resolution, sensitivity demands, and special protocols for animal-handling and specimen preparation. We will illustrate recent advances in imaging rodent anatomy, follow with functional aspects that are some of the main strengths of live animal imaging made possible by MR, and end by highlighting recent trends in MRM, including imaging genetics.

1.1. Basic principles of image formation and hardware requirements

While MRM may appear to be a natural extension of clinical MRI to small animal imaging, the

transition has been accompanied by significant challenges. The major problem is that small animal samples require smaller voxels to achieve comparable anatomic resolution in small animals. Since fewer protons occupy these smaller voxels, the signal is substantially weaker. To address this problem, most small animal imaging studies are performed at higher magnetic fields (4–16 T) than clinical studies (usually 1-3 T), to benefit from increased magnetization. Other steps towards improving the signal-to-noise ratio (SNR) include using more efficient detectors for the MR signal i.e., RF coils, as well gradients that are stronger (500-10000 mT/m versus 10-50 mT/m gradient strength) and faster (1000-10000 Tm⁻¹s⁻¹ versus 10–200 Tm⁻¹s⁻¹ slew rate) [54, 64]. Strong gradients with high slew rate, high duty cycle, and high power amplifiers are critical for achieving fast imaging scans of samples that occupy small volumes $22 \times 11 \times 11 \text{ mm}^3$ ($\sim 2.7 \text{ cc}$) for a mouse brain compared to $20 \times 20 \times 20 \text{ cm}^3$ ($\sim 8000 \text{ cc}$) for a human brain. The use of higher magnetic fields introduces important physical effects that are not as prominent in the clinical domain, e.g., susceptibility variations at the interface of tissue and air, which may result in signal drop and distortions. Relaxation parameters also change when operating at higher fields. Biological motion from the heart (300–600 beats/minute (bpm) in the mouse versus 60-12 bpm in humans), and respiration (breathing rates from 150-170 breaths per minute in mice versus 20 breaths/minute in humans) pose additional challenges to achieving the highest resolution.

The principles of image formation, as well as basic hardware components, are shared by MRM with its clinical parent, MRI, and are well described in [42, 110]. The NMR signal from a source, such as a small animal, positioned inside a magnetic field is generated following excitation by a transmitter coil, detected by a receiver coil, and reconstructed usually by means of a Fourier transform [29, 68]. The origin of the signal is most often the motion of nuclear spin magnetic moment characteristic to hydrogen atoms in water (110 mol/l) and lipid components of the animal's body. The magnetic spins precess (Fig. 1a) about the local magnetic field (B₀, defining the z axis) at the Larmor frequency (ω_L), proportional to the local magnetic field strength (B₀), and a constant called the gyromagnetic ratio (γ) :

$$\omega_L = \gamma B_0$$

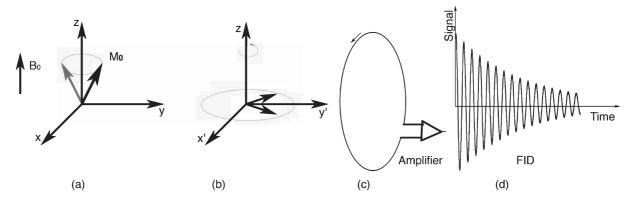


Fig. 1. (a) The source of signal in MR is the proton magnetization, which precesses around the static magnetic field at the Larmor frequency. (b) In a reference frame rotating at this Larmor frequency M_0 appears to be stationary. In the 90-FID pulse sequence, the net magnetization is rotated down into the x'y' plane with a 90° RF pulse. (c) A changing magnetic flux induces a current in the receiver coil. (d) The signal recorded in the laboratory frame following the RF pulse constitutes the FID.

The Larmor frequency for ¹H changes from 64 MHz at 1.5 T to 400 MHz at 9.4 T. A collection of protons in a magnetic field will all precess at this same frequency. Since the protons are precessing together, their coherent motion generates a physical property of the tissue, the magnetization M_0 , which is given by the sum over all individual protons. This magnetization is the true source of the signal. Consider a rotating frame of reference, with angular velocity corresponding to the Larmor frequency (Fig. 1b). Using this reference frame helps us understand the generation of the NMR signal from M_0 . Since the reference frame is rotating at the Larmor frequency, M₀ will appear stationary and aligned with the magnetic field. If one applies an oscillating magnetic field at the Larmor frequency, when resonance conditions are met), one can efficiently move M₀. In the 90-free induction decay (FID) pulse sequence (Fig. 1b), the net magnetization was rotated down into the x'y' plane with a 90° RF pulse. M₀ now lies along the y' axis, in the rotating reference frame. In the laboratory frame, which is stationary, the changing magnetic flux induces a current in the surrounding coil (Fig. 1c), which is sent to an amplifier. The recorded signal constitutes the FID (Fig. 1d). As the magnetization continues to precess in the x'y' plane, the individual protons see slightly different fields because of their local molecular environment. As a consequence of these differences, a gradual loss of coherence occurs, hence the decay.

A schematic representation of hardware required for forming an MR image is shown in Fig. 2. Briefly, a biological sample is placed within the center of a (usually superconducting) magnet, which generates a strong, but static field. Superconducting magnets need to be cooled using cryogens, thus decreasing the power requirements for maintaining the magnetic field. Magnets are also shielded from magnetic fields originating from outside the sample, either passively, using a Faraday cage, or actively compensating for the presence of external magnetic fields.

An RF pulse is sent from the pulse waveform generator (transmitter, frequency synthesizer) to an RF amplifier, and finally to the transmitter coil, where it excites the biological sample. This transmit coil must produce a uniform field B_1 perpendicular to B_0 (the static magnetic field), and therefore have relatively large dimensions.

The signal coming from on-resonance spins, as they return to equilibrium, is collected by the receiver coils. The receiver coil has to be sensitive to the relatively low detected signal. It is usually of small dimensions, just large enough to be wrapped around the imaged sample for a volume coil, or a surface coil placed in close proximity to the area of interest can be used. The disadvantage when using such a coil is that the high sensitivity and high signal at the sample surface falls off with increasing depth. Improvements in *coil performance* help improve sensitivity, spatial resolution [54], potentially reducing acquisition time. The improvement may come from developing coils that are 1) smaller, tight-fitting, or implanted; 2) circularly polarized (quadrature coils, giving $\sqrt{2}$ improvement in

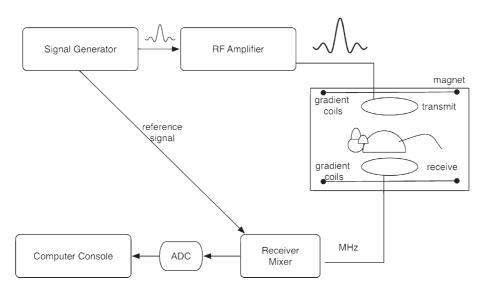


Fig. 2. The basic elements of the MR system include an RF pulse generator, whose signal is amplified before being sent to the transmitter coil. The same signal is used to synchronize the acquisition of the MR signal coming from the receiver coil. After unmixing and passing through the analog-to-digital converter, the real and imaginary components are used to reconstruct and visualize the MR image either by the same console computer used to control the RF and the gradients, or offline.

SNR); 3) birdcage, or 4) cryogenic or superconducting coils with negligible resistance at low temperatures (3–10 fold improvement in SNR). The use of phased arrays of multiple receiver coils operating independently provides similar SNR as single detectors, over larger field of views.

The signal is digitized by the main computer, which also controls the RF transmit and receive, as well as the gradients. The role of the gradient fields is to create a spatially varying magnetic field, which allows the spatial encoding of the signal, i.e. to determine the location of the signal source according to small variations in the resonant frequency and phase. Three pairs of gradient coils, generating orthogonal and linearly varying magnitudes of magnetic field, are used to encode the spatially varying signal.

In essence, the MR signal is sent from the receiver coil to the receiver mixer, where it is amplified by a low-noise amplifier before being sent to the analog-to-digital converter (ADC). A splitter provides imaginary and real components, corresponding to the x' and y' components of the magnetization vector. These are used to provide magnitude and phase information for each location in the spatial frequency domain, or k-space.

An image is typically generated through the repetition of a sequence of RF and gradient waveforms (pulse sequence) that encode the image. The signals from the repeated sequence are collected by the computer and transferred to software that performs the image reconstruction.

1.2. Contrast mechanisms

The signal in MRI is governed by multiple parameters such as the proton density, their polarization, relaxation rates (R1, R2, and R2*; relaxation rates are the inverse of relaxation times T_1 , T_2 , T_2 *), diffusion properties of water and water exchange dynamics. This multivariate signal dependence is the source of the rich soft tissue contrast, which can be tuned by choosing appropriate acquisition parameters.

Since the samples in MRM are small (a mouse is almost 3000-times smaller than a human, per weight), the voxels required are small compared with clinical imaging. Also, the signal coming from these smaller voxels is much weaker. To achieve microscopic resolution, one must somehow increase the sensitivity. One method to do so is to operate at higher magnetic field. But when this is done, it is important to be cognizant of the changes in physics induced by the higher magnetic field.

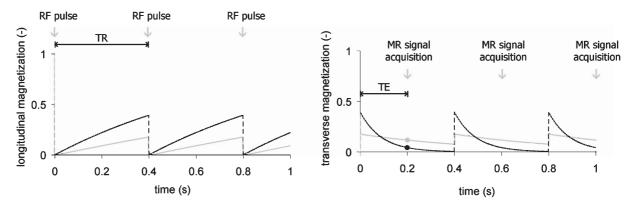


Fig. 3. Differences in the spin-lattice (T_1) and spin-spin (T_2) relaxation times of tissue compartments can be emphasized by intelligent manipulation of acquisition parameters, such as echo time (TE) and repetition time (TR). In the left panel, the longitudinal magnetization of white matter (black) with $T_1 = 0.8$ ms recovers faster than cerebrospinal fluid with $T_1 = 2$ s (gray). The differences are seen using a relatively short repetition time $(0.4 \, s)$. In the right panel, the signal from liver (black), which has a short $T_2 = 40$ ms, diminishes faster than the signal from fat, with a longer T_2 of 60 ms (gray). These differences are captured at an echo time of 20 ms. This figure has been adapted from [25]. It is reprinted with permission from the ILAR Journal, Institute for Laboratory Animal Research, National Research Council, Washington, DC (www.nationalacademies.org/ilar).

After the application of a 90° RF pulse that tips the magnetization vector into the transverse plane (see Fig. 1), the longitudinal component of the magnetization recovers at an exponential rate characterized by a tissue characteristic constant, the spin-lattice relaxation time (T_1) . At a time t the longitudinal magnetization is given by:

$$M_z = M_0(1 - e^{-\frac{t}{T_1}})$$

At 1.5 T, typical for clinical systems, T_1 is 0.8 s for brain white matter, but 2 s for cerebrospinal fluid (CSF). As a consequence, white matter recovers its longitudinal magnetization more quickly than CSF [41], as illustrated in Fig. 3. The transverse magnetization, however, decays at a rate governed by the spin-spin relaxation time T_2 , or to be more precise T_2 * (called T-2-star), describing in homogeneities of the local magnetic field.

$$M_{xy} = M_0(1 - e^{-\frac{TR}{T_1}})e^{-\frac{t}{T_2}}$$

The MR image reflects the properties of the transverse magnetization at a time TE (echo time) after application of the RF pulse (B_1 field). As RF pulses are repeated with a periodicity given by TR (repetition time) the intensity of the MR image will be proportional to the value of the starting longitudinal magnetization at TR. Thus, TE and TR interact with T_1 and T_2 to modulate the MR signal.

The endogenous contrast (based on proton density, T_1 , and T_2 differences) can be supplemented with exogenous contrast agents, which affect relaxation properties of water protons to induce dramatic changes in T₁ and T₂ (Fig. 4). Increased signal is obtained due to T₁ shortening by paramagnetic agents, while signal drop is due to strong, but negative susceptibility effect of superparamagnetic agents. These paramagnetic or superparamagnetic compounds are based on the strong magnetic moment of unpaired electrons. Paramagnetic probes are typically complexes of gadolinium (Gd), containing 7 unpaired electrons, in which the metal ion is coordinated by a chelating ligand in order to reduce its toxicity (e.g., GdDTPA or GdDOTA). Superparamagnetic probes are iron oxide nanoparticles, containing hundreds or thousands of iron atoms, exerting a high magnetic moment. Again, biocompatibility is achieved by decorating the iron core with an organic coating, e.g., dextran. The magnitude of the contrast agentinduced changes in proton relaxation times depends on the molar relaxivity of the contrast agents (a measure of their efficiency) and their concentration in tissue. Typically, the method enables the detection of signal changes induced by local concentrations of the contrast agents of $10^{-3} - 10^{-6}$ molar [43]. In general, these changes are not specific to an organ, tissue, or lesion.

The use of blood pool contrast agents [20] allows one to image the vascular system and to evaluate cardiac function [83], and accumulation in

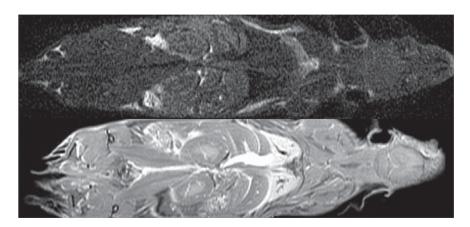


Fig. 4. A five-fold higher signal-to-noise ratio (SNR) per unit scan time can be obtained imaging a specimen prepared with active staining (bottom row: 1:20 Gd-DTPA, 10% formalin) compared to a formalin-fixed specimen (top row: 10% formalin, TR = 100 ms). SNR increased from 10 to 50. Reprinted with permission from [63].

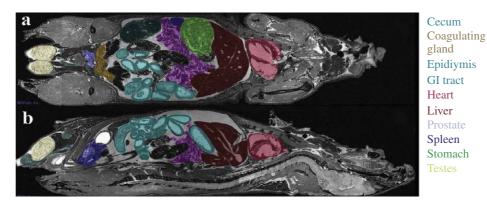


Fig. 5. Whole body imaging of a fixed-perfused mouse specimen at $50 \,\mu m$ resolution allows the identification of major organs, as well as the assessment of microstructural properties. *Reprinted with permission from* [78].

lesions [34]. These are available commercially e.g., gadofosveset (Vasovist, Bayer Schering Pharma, Wedding, GE; MS325, EPIX Medical, Cambridge, MA, USA) or ferumoxtran ultra small superparamagnetic iron oxide (Combidex, AMAG Pharmaceuticals Inc, Cambridge, MA), or are developed by research laboratories [33, 112].

Major organs, such as the liver, heart, or kidneys, can be identified and segmented in the mouse body, imaged at 50 μ m resolution (Fig. 5) on a 7T MR scanner with a GE console. The spatial relationships among organs can readily be appreciated and the slice used to cut through the digital volume can be oriented in any of the three cardinal planes (Fig. 5 shows a horizontal (a), and a sagittal (b) cut), or in an arbitrary plane.

The adaptation from MRI to MRM has required considerable technical developments, including operating at higher magnetic field strengths, improved encoding strategies, and specialized radio-frequency coils [62, 63]. But, these developments have opened a number of fascinating applications for the basic scientist, which we will illustrate in the following sections.

2. Animal preparation for imaging

One of the main advantages of MRM is the capability to image live animals, and to be able to repeat the imaging at multiple time points, e.g., during



Fig. 6. A dedicated cradle helps to quickly and accurately reposition a mouse within the radiofrequency coil (top figure), while allowing anesthesia to be administered by a nose cone, and to monitor physiologic parameters via a respiratory pillow, temperature probe and/or electrodes. (Colours are visible in the online version of the article; http://dx.doi.org/10.3233/ACP-2011-0050)

disease progression to assess the extent of pathological changes, or during treatment to assess its efficacy. However, the animal is located within the small bore of a strong magnet, which makes it challenging for the investigator to supervise its physiological state or to administer contrast agents. An excellent treatise of animal handling for in vivo imaging can be found elsewhere [47], in this section we will cover the basic elements. It is essential that magnet compatible equipment should be used in the proximity of the magnet and a remote physiological system must be used to monitor and adjust body temperature, level of anesthesia, or to trigger the image acquisition. To counteract the effects of motion associated with respiration and to control lung volume, animals can be supported by a ventilator. This way, the position of the lungs is precisely reproduced from one breath to the next [47, 77], and the image acquisition can be triggered as specific phases in the respiratory cycle, where motion is minimal (end expiration or full inspiration). To achieve ventilatorsynchronous acquisition, animals have to be intubated with an endotracheal tube, and closely monitored. ECG monitoring electrodes can be used to achieve cardiac gating to specific intervals during the cardiac cycle.

MR-compatible monitoring systems are commercially available (e.g., BIOPAC MP 150, from BIOPAC Systems, Inc, Galeta, CA) to ensure adequate animal support for the duration of the scan. These can be used in conjunction with specialized cradles that facilitate the accurate positioning/repositioning of the animal. For example, a mouse cradle can accommodate ECG pads, a respiratory pillow, a temperature probe, and a nose cone equipped with an incisor bar, which delivers

gas anesthesia, while ensuring proper positioning [52]. To maintain the animal's normal body temperature, a thermistor attached to the cradle controls the temperature of a heating agent (water or air) flowing through the cold magnet core. Using such a device, the animal can be quickly set up within the coil (Fig. 6), with the head reliably fixed in position.

Since voxel size needs to be small to resolve structures in the rodent body, the signal that comes from the voxel will be proportionally smaller. For example, a voxel size of 20 µl would be sufficient for imaging the human lung, but to obtain comparable details in the mouse chest, the voxel size would be perhaps 0.6 nl [25]. This corresponds to a 33-times smaller signal coming from such a voxel. One way to compensate for such a signal loss is to inject parmagnetic contrast agents, such as Gd chelates (e.g., Magnevist, ProHance, Multihance, etc.). The paramagnetic agents shorten the relaxation times and allow faster MR acquisitions, important when imaging live animals. Contrast agents are usually administered by intravascular (IV) injections. One single injection may suffice, if the contrast agent has a sufficiently long circulation time, or a pump may be used for continuous infusion of small volumes of fluid. Other techniques to obtain higher image contrast increase the number of spins aligned versus those anti-aligned to the applied magnetic field, i.e. the amount of polarization [109]. To deliver hyperpolarized gas to the lungs the flow of air must be rapidly switched to a mixture of hyperpolarized gas and oxygen at specific phases in the breathing cycle [24]. Because of the potential high increase in sensitivity, active efforts are being made to increase polarization

levels and delivery efficiency to image lung perfusion and gas exchange [23, 31], as well as to image blood flow and perfusion in the brain [105, 119], or to estimate oxygenation [114].

Image resolutions of 100 µm isotropic have become standard in small animal imaging in vivo, but higher resolutions can be obtained from fixed specimens, which can be scanned for long times (sometimes more than 24 hours) to yield high signal-to-noise ratio, or to use multiple contrast mechanisms. To further enhance contrast-to-noise ratio, smaller specimens, such as embryos, can be immersed in a solution of fixative (e.g., Bouin's) and MR contrast agent [89]. For imaging larger specimens a vascular perfusion is done with a mixture of contrast agent (ProHance, Bracco Diagnostics, Princeton, NJ), and fixative (e.g., formalin) to achieve what has been coined as "active staining" [63]. Postnatal aged animals need to be first flushed of blood with a mixture of saline and ProHance, followed by fixation perfusion with formalin and ProHance [90]. The active staining methods allow acquiring MR images of fixed specimens with exquisite contrast and in shorter times compared to those obtained from simply formalin fixed specimens.

3. Anatomical imaging

As shown in the previous section, MRM is well suited to study rodent histology [60–63]. Compared to specimen imaging, live animal imaging is restricted to shorter scans and is complicated by motion. Blood flow, cardiac, respiratory, and peristaltic motions are potential sources of imaging artifacts. One approach to circumvent these problems is to acquire images in a manner that exploits the periodicity of the physiological signal. Other strategies aim to sample faster, switch from Cartesian to radial and spiral acquisition, and/or to correct for motion artifacts. The next sections present examples where specific organs are imaged in the live animal, the associated challenges, and proposed solutions.

3.1. Body imaging

Respiratory motion is one of the foremost challenges in imaging the animal's body *in vivo*. Active mechanical ventilation can be used to synchronize the acquisition with the respiration. However, this

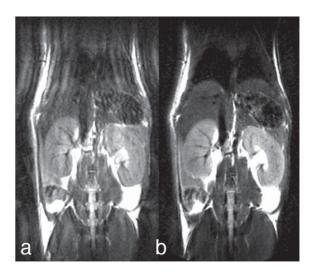
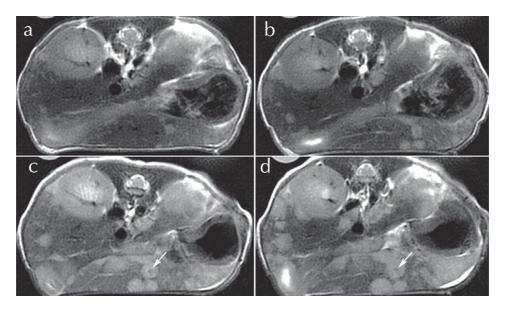


Fig. 7. Effects of motion (ringing around sharp edges) can be greatly decreased relative to Cartesian Fast Spin Echo (a) using PRO-PELLER (b) imaging These images were acquired on a 7T magnet at $195\times195\,\mu m$ resolution in plane, 1 mm slices, TR = 3 s, echo spacing 5 ms, echo train length 24 (lines per blade), in $\sim\!10\,\rm min$. Reproduced with permission from [87].

approach has two drawbacks. It requires intubation of the animal, which significantly reduces the throughput. Also there is some morbidity associated with the intubation, which can limit the number and frequency of scans in a longitudinal study. Alternatively, methods for motion correction similar to those used in humans can be adapted to small animal imaging. One example is the use of PROPELLER (Periodically Rotated Overlapping Parallel Lines with Enhanced Reconstruction) for imaging the mouse body [92]. The method [88] relies on a novel strategy to sample the raw (Fourier) data. Figure 7a shows a coronal image of a free-breathing mouse acquired with a conventional Cartesian sampling strategy. The image is the result of 256 pulses repeated over a period of up to 12 minutes. Since the data are acquired without synchronization with respiration, the inconsistencies of the respiratory state of the animal leads to the ringing artifacts seen in Fig. 7a. The image in Fig. 7b was obtained from the same free-breathing animal using the PROPELLER method. The raw (Fourier) data used to reconstruct the image, has been sampled with "blades"—produced from bursts of data. Each blade is acquired at a different angle. A very low-resolution image can be reconstructed from each of these blades. The phase of the data is used to determine if there are motion differences between the individual blades. When motion irregular-



ities occur, that particular blade of data is rejected. As is evident in Fig. 7b, the motion artifacts are significantly reduced.

A particularly appealing application of PRO-PELLER imaging is characterizing cancer lesions, e.g. metastases from implanted human cell lines of HTN29 colon carcinoma (Pandit, 2010 — personal communication). Careful selection of the acquisition parameters allows obtaining images in which the signal is heavily dependent on differences in tissue spin-spin relaxation (T2) (Fig. 8). Liver metastases are very visible in these T₂-weighted images obtained with a twoshot PROPELLER with TE of 67 ms and TR of 3 s, at 125 µm in-plane resolution, 1 mm between planes, acquired in only 33 minutes on a 7T magnet. Lesions are visible in the ventral liver lobe starting with day 17 after inoculation and they continue to increase in size as illustrated in Fig. 8. Moreover, one can discern a hypointense necrotic core (arrows) surrounded by hyperintense lesion mass.

3.2. Brain

MR has been extraordinarily valuable in imaging the brain because of the superb contrast between gray and white matter. Differences in T_1 provide superb morphologic contrast in the clinic. And differences in T_2

provide the most frequently used method for delineating pathology. Unfortunately at higher magnetic fields used for small animal studies, these mechanisms are less effective. In addition, the use of contrast agents in the brain is hindered by the blood brain barrier (BBB). Several groups have recently described approaches to circumvent this problem by opening the blood brain barrier, which can also be done using ultrasound [16, 51, 55]. This improves both the signal to noise, by reducing the T_1 , and in some cases the contrast to noise. Such images can allow one to distinguish details such as the hippocampal layers, white matter tracts, as well as the organization of the vascular network (Fig. 9). The contrast is much higher than what was obtained without opening the BBB even for a longer TR in previous studies.

We have used this method to detect amyloid plaques in a transgenic mouse model [56]. Opening the blood brain barrier globally reduces the T_1 of the brain parenchyma, permitting the use of a shorter TR, i.e. more rapid scanning. We have and used a multiecho sequence processed in Fourier domain [99] to enhance tissue differentiation on the basis of T_2 . The *in vivo* image shown in Fig. 10 was acquired using a TR of 400 ms, and an echo train length of 8, with spacing of 6.5 ms, at a spatial resolution of $40 \times 52 \times 625 \,\mu\text{m}$, with an acquisition time of less than 2 hours. The plaques appearing hypointense have a shorter (T_2) than



Fig. 9. In Vivo Images of the mouse brain before and after infusion of Gd-DTPA through an opened blood brain barrier. Both images were acquired with the same sequence and parameters: Spoiled gradient recalled image (SPGR), TE = 5 ms, TR = 28 ms, flip angle 30° , BW = 15 kHz, resolution $40 \times 40 \times 200 \,\mu$ m. Courtesy of Nicholas Laucis, Duke Center for In Vivo Microscopy.

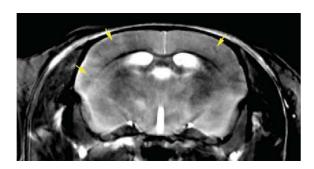


Fig. 10. *In vivo* imaging of the mouse brain after Gd-DTPA delivery in conjunction with ultrasound opening of BBB. (TR = 400 ms, ETL = 8 echoes, echo spacing 6.4 ms, resolution $40 \times 52 \times 625 \,\mu\text{m}$, matrix size of $512 \times 512 \times 32$ slices, BW 62.5 kHz. Total acquisition time is under 2 hours. Note the reduced signal from the plaques (arrows). (Colours are visible in the online version of the article; http://dx.doi.org/10.3233/ACP-2011-0050)

the surrounding tissues. We speculate that this might be due to the presence of iron in the plaque.

Several investigators have used targeted contrast agents for MR visualization of amyloid plaques. These agents can be modified to enhance BBB permeability e.g., using a polyamine modification [93] or delivered in conjunction with opening of the BBB [100]. The targeted agents bind specifically to the amyloid plaques, resulting in enhanced local contrast.

The use of MR to study fixed specimens, i.e. MR histology, was first suggested in 1993 [60, 61]. Since the tissues are fixed, one can use specialized methods

to actively stain the tissues, and longer scanning times to achieve higher spatial resolution than attained in vivo [62, 63]. As with the actively stained in vivo studies, one can exploit several different sources of contrast as shown in Fig. 11. For example, a T₁-weighted scan (Fig. 11a) is useful to accurately determine the borders of gray matter structures, such the superior and inferior coliculi, olfactory bulbs, or hippocampus, as well as its subfields; or to identify cortical layers and white matter tracts 1. A longer T₂-weighted scan (Fig. 11b) adds rich information on thalamic nuclei, such the medial and lateral geniculate. This information from the two scans can be combined to segment the brain compartments [2, 59, 98], and to infer information on their on their shape, size, and location relative to the each other and to the whole brain (Fig. 11c and d).

The volumes of segmented structures can give information on the covariance structure in the CNS of mouse strains that have been well genotyped, e.g. the BXD family. The high estimates of heritability for these structures [5, 6] allows us to believe that it will be possible to map genes responsible for variations in the anatomical phenotype.

3.3. Atlasing efforts

Analogous to efforts in human brain imaging, the mouse neuroimaging community has developed brain atlases, for several mouse strains including C57BL/6,

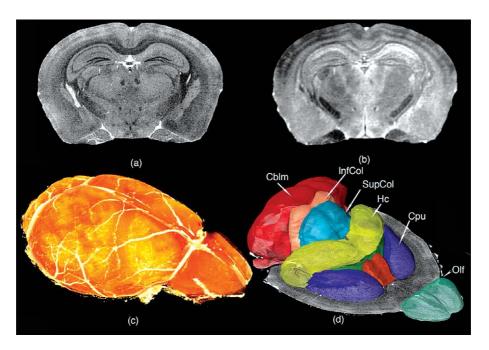


Fig. 11. (a) T_1 -weighted and (b) T_2 -weighted images of the actively stained mouse brain offer rich contrast to differentiate and segment brain regions, and can help determine the genetic basis of their volume variation; (c) Volume rendered T_1 image showing the brain surface; (d) Surface rendering of structures that have been segmented and quantified, including olfactory areas (Olf), striatum (CPu), septal nuclei, fimbria, hippocampus (Hc), superior and inferior colliculi (SupCol, InfCol), and cerebellum (Cblm). Reprinted with permission from [5, 6].

129S1/SvIm [4, 59, 74–76]. The digital imaging task force of the International Neuroinformatics Coordinating Facility (INCF, www.incf.org) is developing tools to facilitate the international sharing of data from the mouse brain [46]. Waxholm Space (WHS), one such tool, is a coordinate system based on multispectral 3D data from three different MRM imaging protocols and conventional Nissl stained histology (Fig. 12). Thirty-seven substructures of the brain have been labeled and the data and visualization tools are freely available on line (http://www.civm.duhs.duke. edu/neuro201001/index.html). Data from 13 additional specimens scanned using the same imaging protocol has allowed construction of an average atlas with T₁T₁-weighted, T₂-weighted, and T₂*-weighted images, where one can distinguish with increased precision gray mater nuclei and white matter bundles, as well as construct an average label and probabilistic atlas.

Such atlases can be used to estimate limits of normal variability within one strain, as well as to make comparisons across strains and characterize populations of mouse models of neurological disorders.

3.4. Diffusion tensor imaging

The diffusion properties water in tissue can be used to complement the information from (T_1) -weighted, T₂-weighted, and T₂*-weighted images of the rodent brain [8, 70, 82], and provide information on the degree of orientation in an image. The richness of the tensor information supports a number of different diffusion-related parameters, and helps characterize the brain microstructure. Figure 13 illustrates a few important parameters derived from the diffusion tensor. FA (fractional anisotropy) reflects how anisotropic the diffusion in one voxel is. AD (axial diffusivity) is the primary eigenvalue, representing diffusivity along the primary axis of the diffusion tensor ellipsoid. RD (radial diffusivity) is the mean of the secondary and tertiary eigenvalues, characterizing diffusivity perpendicular to the primary axis. White matter exhibits higher anisotropy, characterized by the FA, higher AD, and lower RD than the gray matter, since in white matter bundles, in the axon and its surrounding extracellular space, the parallel diffusivity is higher and the perpendicular diffusivity is lower compared to those

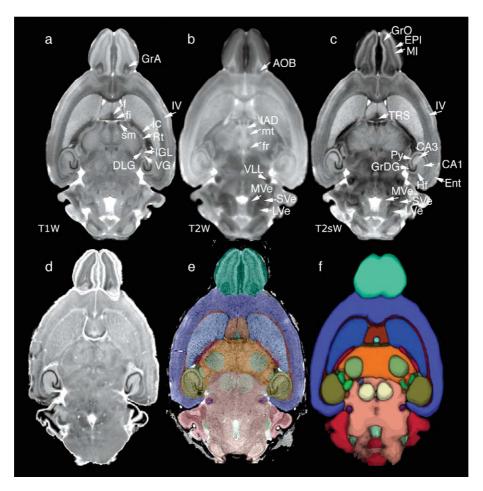


Fig. 12. Multiple specimens imaged with MRM have been used to generate population-averages: (a) T_1 -weighted $(T_1 W)$, (b) T_2 -weighted $(T_2 W)$, and (c) T_2 *-weighted $(T_2 s)$; (d) also a 3D volume of Nissl histology was acquired on one of the specimens; (e) labels defined on the one specimen atlas images can be used to map other mouse brains; (f) labels from multiple specimens have been used to produce a probabilistic atlas that gives insight into volume and shape variability. *Modified and reproduced with permission from* [59].

in the soma. These images, which characterize orientation, can be used to distinguish small white matter bundles, such as the habenular commisure, or among fibers located closely in space such as fimbria, optic tract, stria terminalis, cerebral peduncle; or the cingulum from the rest of the corpus callosum (Fig. 13).

Fiber tracts can be reconstructed based on DTI using tractography algorithms [81], to give insight into the architecture of neural connections. This information, together with parameters extracted from DTI, can be used to characterize connectivity in the normal brain and to identify changes in mouse models of disease involving loss of white matter integrity. DTI atlases [18, 58] help assess the bounds of normal variability in the adult, as well as progression of myelination during

development, or defects in animal models of diseases involving demyelination [32].

3.5. Developmental imaging

Atlases have also been useful in studying development in the mouse embryo and fetus [19, 91]. Images acquired for embryos of age E9.5 with a FOV of 5 mm and a $256 \times 256 \times 256$ matrix resulted in $20 \,\mu m$ isotropic resolution. Larger arrays are however required for imaging larger specimens. To obtain the same resolution for an E19.5 embryo, which requires a FOV of $20 \times 10 \times 10 \, mm$, the image array must be $1024 \times 512 \times 512$. This results in $\sim 4GB$ of

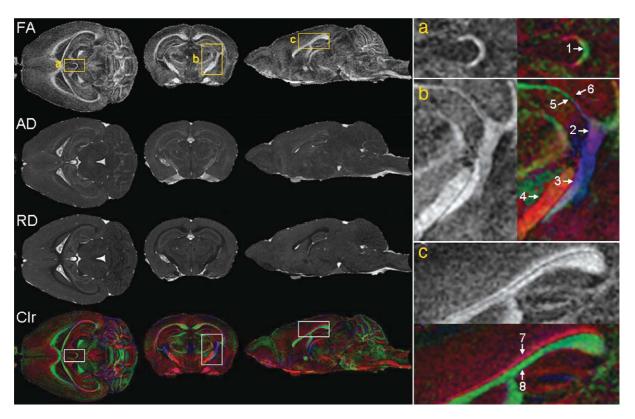


Fig. 13. Diffusion tensor imaging (DTI) provides rich information on white matter structures, connectivity patterns, as well as the integrity of these connections in the mouse brain. Among the parameters calculated from DTI are: FA (fractional anisotropy), AD (axial diffusivity), RD (radial diffusivity), and Clr (color-coded orientation map of the primary eigenvector. Green: left-to-right; red: rostral-to-caudal; blue: dorsal-to-ventral). The right column shows magnified views that correspond to the regions specified by the rectangles (a, b, and c) in the left column. Arrows indicate structures that are better discerned using directional information on the colormap: 1. Habenular commissure; 2. Fimbria; 3. Optic tract; 4. Cerebral peduncle; 5. Thin branch of optic tract adjacent to 6. Thin branch of fimbria; 7. Cingulum (rostro-caudal orientation); 8 Corpus callosum (medial to lateral orientation). *Reproduced with permission from* [58].

raw data, and yields an array of \sim 500 MB for the reconstructed image. The approach requires powerful computational resources, relatively large storage space, and large amounts of memory to visualize the 3D anatomy. To address these problems [91] have shared their imaging protocol and imaging database of the developing mouse, at 16 stages, from E10.5 to postnatal day (PND) 32, as well as 200 associated labels. The protocol starts with actively staining specimens either through immersion or cardiac perfusion under ultrasound guidance. This active staining allows the acquisition of 3D images with 19.5 µm resolution in \sim 3 hours. Figure 14 illustrates the differences in scale encountered in developmental imaging from E10.5 to E18.8, where the crown to rump length varies from 4 mm at E10.5, to about 16.5 mm at E18.8, but this continues to get larger—up to 75 mm crown to tail length in PND32.

A high-throughput version of the embryo imaging protocol has been used to characterize cardiac defects in a mouse strain with a conditional ablation of the Smo receptor gene [91]. In these mutant mice, the outflow tract appears as a singe entity, while in the wild type (WT) control it branches into the aorta and pulmonary trunk (Fig. 15A and B). The interventricular septum appears open in the mutant rather than continuous as in the WT. In the most caudal slices the dorsal mesenchymal protrusion (DMP) is absent in the mutant. 3D surface renderings of the reconstructed heart and associated blood vessels allow for direct observation of changes in the shape and position of the different parts of the



Fig. 14. Mid sagittal slices from volume datasets of developing mouse at stages E10.5–E18.5 illustrate differences in specimen scale occurring in less than 10 days during early development [91].

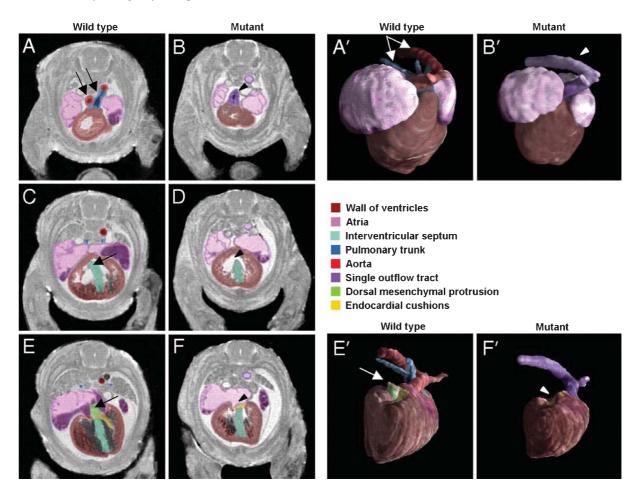


Fig. 15. MRM imaging at $19.5 \,\mu m$ resolution allows discerning cardiac defects in a mouse with conditional ablation of Smo receptor gene. From most cranial to most caudal slices, the arrows point to (A, B) outflow tract abnormalities in the mutant mouse, which fails to branch into the aorta and pulmonary trunk (C, D) an open septum defect (E, F), absence of DMP in mutant mouse. (A', B') present surface rendering of the segmented heart (ventral view) and abnormal outflow tract in the mutant in comparison to the wild type control. (E', F') illustrate the absence of DMP (in a dorsal view of the heart). Reproduced with permission from [91] Copyright 2008 National Academy of Sciences, USA.

anatomy, as segmented based on high-resolution MRM images.

While demands on the resolution and scan time are exacerbated in the case of developmental imaging, because of the small sample size, early detection of abnormal phenotypes is a promising avenue not only for studies of mutant [36] or engineered mice, but also for studies on the effects of the environment, toxic insults, and substances of abuse [37].

4. Functional imaging

While structural imaging, as well as recordings of physiological parameters, have both been used for a long time, they have traditionally provided independent views on the same animal model. MRM has the ability to provide both anatomical imaging, and insights into the physiological status, through estimates of cardiac contractility, blood volume and flow dynamics, neuronal, and respiratory function. Such imaging studies require that the animal must be held under anesthesia for the duration of the scan, which may affect the physiological parameters and make studies such as functional MRI (fMRI) very challenging in small animals. In addition, biological motion tends to exacerbate problems in live animal imaging, compared to clinical imaging. This is because the mouse heart rate is in the range of 500 bpm, while the respiratory rate is around 160 bpm, depending on the activity level of the animal.

4.1. Heart

Given the prevalence of heart disease in the modern world, cardiac function has become one of the foremost interests of live animal imaging [28]. Normal as well as mouse models of cardiac dysfunction have received special attention. One major goal has been to obtain high throughput in estimating cardiac function, and associated parameters like cardiac mass, wall thickness, stress, ventricular volumes, cardiac output, and ejection fraction [96, 115]. MR allows one to appreciate infarct size, remodeling after infarction, as well as atherosclerotic plaque. But now one must consider 4 dimensional arrays, i.e. spatial and temporal resolution. The rapid heart rate of the mouse, requires temporal resolution on the order of 10 ms to achieve comparable frame rates in the

mouse as in humans. Bucholz et al. have addressed this challenge using a novel 4D radial sampling method [12]. This method differs from the more traditional encoding method by sampling the raw data in a polar coordinate system. Traditional sampling strategies use a Cartesian sampling strategy where one line of a square (or cubic) array is acquired with each excitation. The radial sampling strategy is particularly useful in reducing artifacts from respiratory motion in the mouse and in providing the rapid sampling required to capture the high temporal resolution. But the approach requires specialized reconstruction algorithms, such as the nonuniform fast Fourier transform (NUFFT) described in [101], Bucholz et al have used the 4D radial approach to acquire 4D arrays with 87 µm isotropic spatial resolution and 10 ms temporal resolution in 31 minutes. Figure 16 shows selected images from one such array at diastole. Since the spatial resolution is isotropic, one can choose any arbitrary plane without loss of spatial resolution, allowing us to discriminate not only the ventricles and the atria, but also the major blood vessels (like aorta, pulmonary artery, left carotid artery), and four valves (tricuspid, mitral, pulmonary, and aortic). These high spatial resolution images reduce the partial volumes effects on volumetric estimates of the cardiac chambers, and allow one to assess abnormal morphological and functional phenotypes, including valve defects or aortic stenosis [12].

Such advances in cardiac imaging allow investigation of mouse models of cardiac dysfunction, and assessing the effects of genotype or environment. Bucholz et al. [13] acquired contrast-enhanced cardiac MRM images, using a high-throughput protocol, which results in 87 µm in plane, 352 µm between planes spatial resolution, and a temporal resolution of 9.6 ms (TE = 300 μ s, TR = 2.4 ms, α = 45°, BW = 125 kHZ). These images allowed identifying significant differences among three mouse strains in left ventricle distolic volumes, end systole volumes and ejection fraction (EF), and a markedly lower left ventricle EF for the transgenic mice (Fig. 17). Such studies illustrate the possibilities offered by MRM for rapid phenotyping of the mouse heart, holding great promise for elucidating the relationship between phentotype and genotype, as well as assessing the influence of the environment on cardiac function. Mouse imaging studies, which can provide rich information on cardiac anatomy and function, are particularly relevant to developing drugs and assessing their efficacy [49].

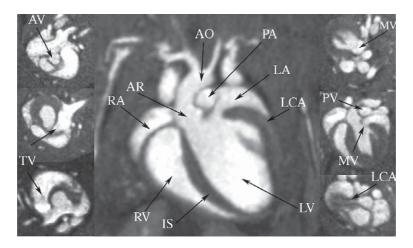


Fig. 16. MRM imaging at resolution of $87 \,\mu m$ allows clear visualization of the four major compartments of the mouse heart: left (LV) and right ventricles (RV), left (LA) and right atria (RA), as well as the intraventricular septum (IS). The major cardiac blood vessels are visible: the left carotid (LCA) feeding the heart, the aorta (AO) ands its root (AR), as well as the pulmonary arteries (PA). In addition, the valves for the aorta (AV) and pulmonary artery (PV), as well as the tricuspid (TV) and mitral valve (MV), are identifiable (insets). Reproduced with permission from [12].

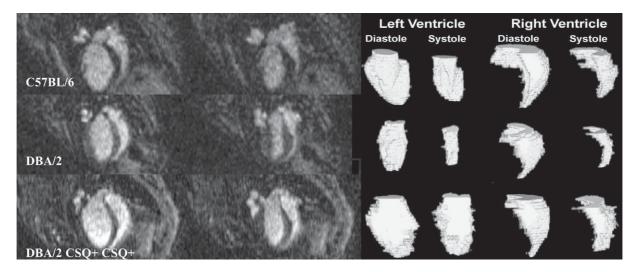


Fig. 17. MRM can be used to identify differences in cardiac function among strains of mice. Here the C57BL/6 was found to be slightly hypertrophic compared to the DBA/2 and DBA/2 CSQ+. The ventricular volumes were used to determine significant differences in left ventricle ejection fraction, smallest for DBA2 CSQ+, and largest for DBA/2, while the right ventricle ejection fraction was significantly higher in DBA/2. Significant differences are also apparent in the right ventricle end systole, larger in DBA2 CSQ+. These differences are also apparent from surface renderings of the segmented heart compartments. *Reproduced with permission from* [13].

4.2. Lung

The challenges for imaging the lungs with MR arise not only from the effects of motion (in the range of 160 breaths/minute in the mouse), but also because the proton density of the lung is lower than that of muscle

(\sim 0.2–0.4 relative to muscle), which correspond to a proportional loss of signal intensity. The many interfaces between the air spaces and tissue lead to rapid signal decay, which can be addressed with ultra-short TE. A more important signal increase can be obtained, however, through enhancement of the nuclear magnetic

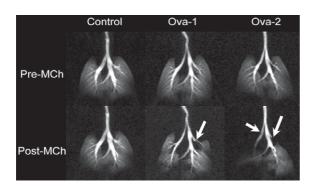


Fig. 18. Gas exchange in the mouse lungs of models of asthma (Ova sensitized) compared to a control animal. Maximum intensity projections based on 3D (radial acquisition) images of 3 He distribution in the lungs before and after administration of $250 \,\mu\text{g/kg}$ methacholine (MCh) challenge. The image were aquired using TR = 5 ms, TE = $132 \,\mu\text{s}$, bandwidth (BW) = $62.5 \,\text{kHz}$, with 20 radial k-space views per breath, interpolated to $256 \times 256 \times 16$ matrix and resolution is $125 \times 125 \times 1000 \,\mu\text{m}^3$. Airways constriction is indicated with white arrows in the Ova-sensitized mice. *Reproduced with permission from* [26].

moment alignment—hyperpolarization [1]. Hyperpolarization (HP) of gas (³He or ¹²⁹Xe) [27] can be used to produce ventilation images. The gas is polarized outside the MR scanner using optical pumping and spin exchange, where the alignment is transferred from laser photons to gas nuclei via an intermediary alkali metal [44]. The hyperpolarized gas

can be captured and transferred to the animal via a dedicated animal ventilator. An immediate application of gas imaging is illustrated in Fig. 18, where a mouse model of asthma, sensitized to ovalbumin (OVA) presents with impaired ventilation patterns compared to normal control after they were both administered a methacholine challenge. Hyperpolarized 3 He distribution can be visualized at resolution of $125 \times 125 \times 100 \,\mu\text{m}$, and in this example was sufficient to allow identification of airway constriction in major bronchi in this mouse model of asthma.

Since ³He availability and price are not practical for large cohort studies, ¹²⁹Xe has received renewed attention and can be used to provide functional information on the lungs [24]. Even more attractive is the fact that ¹²⁹Xe is soluble in fluids and tissue, and exhibits distinct resonance frequencies, separated by large frequency shifts. Because of these characteristics, one can image in the lungs three separate compartments corresponding to peaks of the frequency in the airspace, interstitial space, and blood. Such a compartmental analysis allows the study of gas exchange, for example in mouse models of fibrosis (Fig. 19).

4.3. Brain

While fMRI has become an important tool in human neuroimaging, a major challenge in obtaining func-

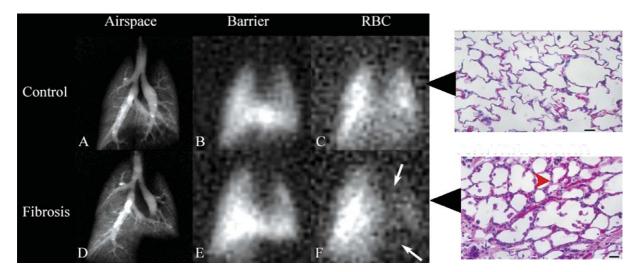


Fig. 19. Gas exchange impairment can be assessed with MR imaging of HP ¹²⁹Xe in the three compartments of the lung: airspace (A, D), tissue barrier (B, E), and red blood cells (C, F). Arrows in F indicate diminished ¹²⁹Xe uptake in red blood cells (RBC) due to inflammation and fibrosis caused by bleomycin. The thickening of blood gas barrier, confirmed by histology slides, slows down the ¹²⁹Xe transfer to gas space. *Adapted and reprinted with permission from* [23]. *Copyright 2006 National Academy of Sciences, USA.* (Colours are visible in the online version of the article; http://dx.doi.org/10.3233/ACP-2011-0050)

tional information from the brain of small animals comes from the fact that anesthesia is known to interfere with brain activation [69]. Imaging awake animals requires control and correction of motion affects, including the difficult task of immobilizing the animal. Even after circumventing these difficulties, the signal recorded is very small. Brain activation was detected by fMRI in the rat somatosensory cortex during electrical stimulation of the forepaws [40], and highly localized in the postero-medial barrel subfield during mechanical stimulation of a single whisker [116]. Photic stimulation was used to elicit brain activation in the mouse brain occipital area containing the visual cortex [53]. Pain-processing and the effects of analgesics were studied by fMRI [107]. In another study, brain activation following intravenous cocaine infusions was characterized in halothane-anesthetized rats using contrast-enhanced fMRI [79]. Intravenous cocaine-induced activation was reflected in increased regional cerebral blood volume (rCBV) throughout the frontal cortex, and partially in the parietal and occipital cortex at cocaine doses greater than 0.1 mg/kg [79].

As an alternative to fMRI, which is coupled to the vascular response, researchers have developed techniques to look into the small animal functional anatomy based on the use of manganese ions or MnCl2 (Manganese Enhanced MRI or MEMRI) [72, 86], a calcium analog that has high uptake in highly active regions of the brain. This technique was shown to identify activated regions of the rodent brain which "light-up" in T_1 -weighted images following auditory [117, 118] or somatosensory stimulation [3, 50, 113], performed after injection manganese chloride [50]. However, the fact that this technique requires general anesthesia and opening of the blood brain barrier, as well as the toxicity of manganese, are issues that need to be considered [30] in order to fully exploit the potential of MEMRI for imaging plasticity, testing behavioral paradigms, or studying the effects of pharmacological agents or substances of abuse on the central nervous system [73].

5. Current trends in MRM

Although present from about the same time as clinical MRI [48], MRM of small animals presents additional challenges and opportunities. Perhaps the most important limiting factor in the translation from human to mouse imaging is that the smaller voxels provide a weaker signal.

One way to improve sensitivity is to increase *coil performance* by developing better copper coils operating at room temperature and customized for specific geometry of samples. Cryogenic probes can increase the sensitivity by a factor of up to 2.9 by operating at low temperature, e.g. 60 K [85], or 30 K [7]. However the cost is still high and temperature must be well controlled to avoid frequency shifts and changes in coil performance. Another approach to improve imaging efficiency is to use phased arrays of small coils that cover a larger field of view [104, 97]. *Multiple animal imaging* can benefit from such developments and contribute to increased throughput scanning of live animals [11, 21, 95] or fixed specimens.

Major efforts are being made to increase the molar relaxivities of contrast agents based on paramagnetic T₁ agents such as Gd and Mn (MnDPDP), or superparamagnetic T2*/T2 agents such as iron oxide nanoparticles. The later category present high sensitivity and can be good candidates for molecular imaging. Contrast enhancement is an ever-developing area of interest, excellently reviewed by [106], providing either higher SNR per unit time overall in the image, or improving the basically low specificity of MR, and more importantly to provide insight into molecular and cellular processes. One approach to enhance the image information content is to use multiple sources of information. For example, anatomical information from high-resolution imaging modalities, such as CT or MR, can be combined with imaging modalities that have great sensitivity but lower resolution, such as PET/SPECT. The first approaches combine imaging modalities through software fusion, while hardware fusion has now become possible. **Software fusion** is a possible answer, but it is a complex process with many degrees of freedom, especially for aligning images of flexible organs that change shape or size. To address these problems, hardware fusion holds great promise, but it is a challenging engineering approach in the case of MR. On one hand, PET/SPECT detectors based on photomultiplier tubes are sensitive to magnetic fields, on the other hand the homogeneity of the magnetic field required for MRM can be perturbed by the presence of PET/SPECT detectors. It has become possible to combine two such techniques into one single system and a low field MRPET system has been proposed [15], as well as a MRSPECT [38]. MR has the potential to improve results currently obtained with imaging modalities of high sensitivity, but lower resolution such as PET and SPECT. Hybrid multimodal imaging techniques aim to combine the benefits of sensitivity with those of resolution and MRM, and can be used in conjunction with other imaging techniques such as CT or fluorescence. *Dual probes* [57] can facilitate the transition from one imaging modality to another, but it is essential that they bind specifically. To achieve this, *molecular probes* are being developed for detecting tumor cell lines, enzyme activity, proteins, or pathogens, for use with MR or joint use with other modalities such as fluorescence [45]. These probes may contain a paramagnetic or superparamagnetic moiety, which improves the MR contrast, and a specific molecular structure. The development of such targeted probes help raise MR to the level of a molecular imaging technique. MR is diverging today from the traditional imaging of water protons, to make use of the enhanced signal, but also the spectral information given by hyperpolarized agents including ³He, ¹²⁹Xe, and ¹³C. These agents are themselves the source of signal, rather than acting on the tissue relaxation parameters. The increase in sensitivity is useful for characterizing lung ventilation patterns and gas exchange [80], brain perfusion alterations in stroke [119], as well as illustrating the metabolic activity in tumors [39]. These techniques are however very expensive, and therefore only accessible to a limited number of research laboratories.

MRM produces in general large image arrays compared to clinical standards, requiring powerful and efficient processing tools from optimized acquisition schemes, efficient computer reconstruction protocols, memory management, and visualization components. Statistical analysis tools are needed to infer quantitative information, which can be used to assess anatomical variability in control populations, and deviations from the normal in mouse models of disease. Quantitative information derived from MRM can be used to explore the relationship between anatomical phenotype and genetic variations [5, 27, 84]. *Imaging genetics* can be used for gene discovery either through genome-wide association studies using imaging data or derived quantitative traits as phenotype, either voxel-wise [102] or using regional phenotypes [94].

Pharmacological research should benefit from the possibility of performing longitudinal studies for drug development, as well as from the ability to image specific processes such as apoptosis or plasticity. One of the challenges when imaging the central nervous system (CNS) is to penetrate the blood brain barrier (BBB) and deliver imaging or pharmacological agents. To

address this problem, chemical [93, 111] and mechanical methods using ultrasound [10, 17, 67] have been developed to disrupt the BBB, as well as cell permeable contrast agents [71]. This may help reduce toxicity, as well as enhance efficacy of treatment for the CNS. Conjugated efforts from multiple disciplines contribute to making MRM one of the foremost imaging modalities for the small animal brain and body.

From its beginnings, MRM has been moving towards efficient high-resolution images approaching the diffusion limit. Moving beyond anatomical imaging, MRM provides growing evidence of its value for studying physiological function of the circulatory system, the heart, as well as neural activity. Molecular MR is still a developing field, along with cellular imaging or imaging genetics, but these are all avenues to watch, because they have the potential to lead towards better understanding of fundamental biological processes.

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