

Published in final edited form as:

*J Magn Reson Imaging*. 2006 November ; 24(5): 1011–1017. doi:10.1002/jmri.20751.

## In Vivo Measurement of Plaque Burden in a Mouse Model of Alzheimer's Disease

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### Abstract

**Purpose**—To demonstrate an MRI method for directly visualizing amyloid- $\beta$  (A $\beta$ ) plaques in the APP/PS1 transgenic (tg) mouse brain in vivo, and show that  $T_{1\rho}$  relaxation rate increases progressively with Alzheimer's disease (AD)-related pathology in the tg mouse brain.

**Materials and Methods**—We obtained in vivo MR images of a mouse model of AD (APP/PS1) that overexpresses human amyloid precursor protein, and measured  $T_{1\rho}$  via quantitative relaxometric maps.

**Results**—A significant decrease in  $T_{1\rho}$  was observed in the cortex and hippocampus of 12- and 18-month-old animals compared to their age-matched controls. There was also a correlation between changes in  $T_{1\rho}$  and the age of the animals.

**Conclusion**— $T_{1\rho}$  relaxometry may be a sensitive method for noninvasively determining AD-related pathology in APP/PS1 mice.

### Keywords

Alzheimer's disease; amyloid- $\beta$ ;  $T_{1\rho}$ ; spin-lock imaging; MRI

Alzheimer's disease (AD) is the most common form of dementia in the elderly (1). Over 4 million people are affected by AD in the United States alone, and as the aging population increases, this number is expected to double by 2025 (2). The classic clinical symptoms of AD include memory loss and confusion. The neuropathological features are neurofibrillary tangles (NFTs) formed by paired helical filaments composed of tau protein, senile plaques (SPs) resulting from A $\beta$  deposits, and neuron loss in the limbic and neocortical regions (3,4). Neuronal loss associated with early AD eventually leads to brain dysfunction and atrophy. Volumetric assessment methods are the current state of the art in magnetic resonance imaging (MRI) of AD in humans and in animal models of AD pathologies. These methods measure gross morphological changes and track disease progression, but do not provide information about biochemical changes. Conventional MRI of the brain, which relies on contrast generated by the variation of  $T_1$  and  $T_2$  or  $T_2^*$  relaxation times of water in tissue, has proven inadequate for observing authentic SPs and NFTs in vivo. This is primarily due

to the high resolution and thin section ( $\sim 10 \mu\text{m}$ ) images required to visualize SPs, which currently are not achievable with MRI. However, interactions between macromolecules and bulk water, and changes in macromolecular content can be indirectly quantified by spatially mapping MR relaxation times. In these relaxometric maps, any change in relaxation times due to the presence of SPs will be reflected over an entire volume-averaged pixel, thereby providing an indirect method of detection without the need for very-high-resolution images.

Intriguing data suggesting that plaques can be detected in brain tissue specimens by  $T_2^*$ -weighted MRI have been reported (5); however, another study using similar methods failed to confirm this (6). Both studies were performed on formalin-fixed brain tissue specimens, and the MR characteristics of fixed tissue differ from those in vivo.  $T_2$ -weighted MRI was employed to visualize plaques in a transgenic (tg) mouse model of AD ex vivo (7,8). Recent work has shown the possibility of detecting A $\beta$  deposits in vivo in the APP/PS1 tg mouse model of AD using  $T_2$ - and  $T_2^*$ -weighted MRI (9,10). The presence of iron in these plaques also exaggerates the actual size of plaques in  $T_2^*$ -weighted images, which makes the accurate quantification of plaque density particularly challenging (9). Also, although very-high-resolution  $T_2^*$ -weighted MRI ( $<100 \mu\text{m}$ ) has been used to image plaque formations in human AD-affected tissue specimens (5), other experiments have contradicted such findings by demonstrating that plaques are *not* visible in  $T_2^*$  images (6). Previous studies have demonstrated the efficacy of gadolinium and magnetically labeled A $\beta$ 1–40 for visualizing A $\beta$  deposits in an APP/PS1 tg model ex vivo (11) and in vivo (12). However, given the neurotoxicity of A $\beta$  and the requirement for mannitol injection for the in vivo experiments, these methods may have limited utility in a clinical setting. Further, in vivo relaxometric measurements established differences in  $T_2$  on the order of a few milliseconds (13) between these mice and age-matched controls. Although these were meticulous experiments, Campeau et al (14) previously demonstrated that  $T_2$  did not significantly differ between AD patients and age-matched normal subjects, which suggests that  $T_2$ -weighted MRI may have limited clinical utility.

An alternate contrast mechanism is  $T_{1\rho}$ , or  $T_{1\rho}$ , the spin lattice relaxation time constant in the rotating frame, which determines the decay of the transverse magnetization in the presence of a “spin-lock” radiofrequency (RF) field. In biological tissues  $T_{1\rho}$  is dependent on the macromolecular composition and provides contrast, unlike conventional  $T_1/T_2$ -based methods. While quantifying  $T_2^*$  with MRI can be prone to errors from susceptibility-induced signal losses, and  $T_2$ -weighted MR signal is degraded by diffusion,  $T_{1\rho}$ -weighted MR images are less prone to such losses and have a greater dynamic range of values as recently demonstrated in the human brain (15). Therefore, quantitative  $T_{1\rho}$  MRI is an attractive alternative strategy for detecting and quantifying A $\beta$  deposits, especially since it probes slow molecular interactions and the dynamics of water molecules in tissues based on variations in protein content (16).

The development of novel diagnostic techniques for AD requires the study of an animal model in vivo before these techniques can be translated to clinical utility. For example, if the disease process can be detected in an in vivo animal model of AD with MRI, and this is validated by histological studies, a paradigm for visualizing the presence of AD pathology can be established. Transgenic (tg) mice (APP/PS1) expressing the pathogenic familial AD-associated form of amyloid precursor protein (APP-Swedish/Tg2576) and presenilin-1 (PS1, P246L/wt) were shown to develop a large number of fibrillary A $\beta$  SP-like deposits in the brain, especially in the cerebral cortex and hippocampus (17,18).

In the present work we demonstrate an MRI method for directly visualizing A $\beta$  plaques in the APP/PS1 tg mouse brain in vivo, and show that the  $T_{1\rho}$  relaxation rate increases progressively with AD-related pathology in the tg mouse brain.

## MATERIALS AND METHODS

### Animal Preparation

The institutional animal care and use committee granted approval for all animal experiments discussed in this work. Two APP/PS1 tg mice and two age-matched controls in three age groups (6, 12, and 18 months) were imaged. The mice were initially anesthetized with intraperitoneal injection of 0.1 cc of ketamine/ acepromazine. The mice were then mounted on a bed and connected to an isoflurane (Abbott Labs, Abbott Park, IL, USA) inhalation anesthesia apparatus. A flow rate of 1 liter/minute with 1.5% isoflurane in air was maintained during the experiments. The bed was then placed into an in-house-built, 3-cm-diameter birdcage coil, and the entire apparatus was placed inside the center of the magnet bore. Throughout the experiments each animal's heart rate was monitored by ECG, and body temperature was monitored by rectal thermometer.

### Imaging Protocol

MRI was performed on a 4.7 Tesla horizontal-bore imaging spectrometer connected to a Varian console (Varian Inc., Palo Alto, CA, USA) with a 12-cm, actively-shielded gradient set capable of 25 G/cm amplitude (Magnex Scientific, Abingdon, UK). A two-dimensional (2D) gradient-echo localizer imaging sequence was run in three orthogonal planes through the center of the gradient sets to facilitate reproducible slice planning. The position of each slice relative to the rostral tip of the brain was noted, and extreme care was taken to ensure that the head and body of the animal lay flat on the bed to facilitate subsequent matching with histological sections.

### High-Resolution MRI

Initially, two 18-month-old tg mice and two age-matched controls underwent high-resolution MRI.  $T_{1\rho}$ -weighted images were acquired with a 2D gradient-echo-based  $T_{1\rho}$  pulse sequence (19). Typical imaging parameters were TE/TR = 2 msec/500 msec, flip angle = 45°, FOV = 2 cm × 2 cm, thickness = 130  $\mu$ m, matrix size = 256 × 256, in-plane pixel size = 78  $\mu$ m × 78  $\mu$ m. The time of spin-lock (TSL) was fixed at 15 msec, and the spin-lock field ( $\gamma B_1$ ) was 500 Hz. The total imaging time was a little under three hours.

### $T_{1\rho}$ Mapping

A series  $T_{1\rho}$ -weighted images were acquired from each animal with a 2D spin-echo-based  $T_{1\rho}$  pulse sequence (20) with six equally spaced TSLs from 10 to 60 msec. These images were 250  $\mu$ m-thick pixel size was 156  $\mu$ m × 156  $\mu$ m and were used to generate  $T_{1\rho}$  maps by fitting signal intensities as function of TSL by linear regression. In the fitting routine, pixels whose intensities correlated poorly (i.e.,  $R^2 < 0.95$ ) with TSL were set to zero. This process guaranteed that noisy pixels were rejected from the maps. Other imaging parameters were TE/TR = 7 msec/2000 msec, FOV = 2 cm × 2 cm, and a spin-locking field of 500 Hz. The total imaging time was about three hours for an entire data set of six TSLs.

### Histology

After the MR experiments the mice were subjected to histological analysis. The mice were lethally anesthetized and perfused intracardially with phosphate-buffered saline (PBS, 0.1 M, pH 7.4) followed by 10% neutral buffered formalin. Their brains were rapidly removed, immediately frozen on dry ice, and stored overnight at  $-80^\circ$ . Blocks were cut by cryostat in a near serial array of 40- $\mu$ m-thick coronal sections for analysis. Finally, we identified characteristic features of SPs in paraffin sections by subjecting them to histochemistry with Thioflavin-S and immunohistochemistry with anti-A $\beta$  antibodies as described previously (21). Histologic slides were digitally scanned with a Nikon (Nikon Corp., Japan) slide

scanner. Three consecutive immunostained sections were selected and manually overlaid by visual inspection of brain structures using Photo-Shop (Adobe Systems Inc., San Jose, CA, USA) software to generate a 120- $\mu\text{m}$ -thick section for comparison with the corresponding MR image.

### Data Processing

The MR images were transferred to a Macintosh G4 PowerBook computer (Apple Computers Inc., Cupertino, CA, USA). Basic image processing ( $3 \times 3$  smoothing filter, and contrast and brightness adjustments) was performed with custom-written software in the Interactive Data Language (RSI, Boulder, CO, USA). Signal-to-noise ratios (SNRs) were measured as the mean signal intensity in a  $20 \times 20$  pixel region of interest (ROI) within the brain parenchyma in each image divided by the mean intensity of background pixels in a  $20 \times 20$  pixel region outside the animal's anatomy. An increase in plaque density is expected in older animals and in anatomical locations such as the hippocampi and cortex (22). Therefore, we chose ROIs for the measurement of the  $T_{1\rho}$  values in these specific regions (Fig. 1).  $T_{1\rho}$  values were recorded from  $10 \times 10$  pixel ROIs, which were manually selected by a single user (A.B.) in the hippocampi, cortex, and thalami, as indicated in the figure. All 100  $T_{1\rho}$  values in each ROI were tabulated in an Excel spreadsheet (Microsoft Corp.). The data from both animals in each group were averaged at each location. A Student's *t*-test was performed to determine the statistical significance of the difference between the data in each tg group and its corresponding age-matched control group. In a similar manner, the statistical significance of the inter-age difference among the three tg groups was also determined.

## RESULTS

Representative high-resolution ( $78 \mu\text{m} \times 78 \mu\text{m}$  in-plane and  $130\text{-}\mu\text{m}$  slice thickness)  $T_{1\rho}$ -weighted MR images of the tg and control mice are shown in Fig. 2a and f, respectively. Several regions of hypointensity (some indicated by arrows) are evident in the brain of the tg animal (Fig. 2a) corresponding to the location of  $A\beta$  plaques in the histological section (Fig. 2b). This is also seen in the MR images in the middle row of the figure, where the regions indicated by the arrows in Fig. 2a and b are magnified alongside their corresponding histological images (Fig. 2c–e). Analysis of the images revealed that several large plaques were visible in the hippocampus and cortex by MRI. MR image pixels containing such plaques have reduced  $T_{1\rho}$  values because they exhibit lower signal intensity in the  $T_{1\rho}$ -weighted images. However, not all the plaques seen in the histological section could be matched with the MRI section, and some of the plaques visible on histology were too small to be detected in the  $T_{1\rho}$ -weighted images.

Typical in vivo  $T_{1\rho}$ -weighted images that were used to generate  $T_{1\rho}$  maps of the mouse brain are displayed in Fig. 3. The SNRs for the images displayed in this figure were 36:1 for the earliest TSL image (10 msec) and 16:1 for the longest TSL image (60 msec). These images provided  $T_{1\rho}$  relaxometric maps (Fig. 4) in which anatomical structures, such as the hippocampi, corpus callosum, and ventricles, are readily discerned. In Fig. 4 a  $T_{1\rho}$  map of an 18-month-old tg animal and its age-matched control are displayed above their corresponding Thioflavin-stained histologic images. The large diffused plaque aggregates, which are bright green under the fluorescein isothiocyanate filters, are located primarily in the cortex and hippocampi of the animals. Some of these aggregates (indicated with arrows) coincide with regions of decreased  $T_{1\rho}$  values.

The results of  $T_{1\rho}$  measurements are summarized in Table 1 and displayed in Fig. 5. At 6 months, when the tg animals had few plaques, they did not show any significant decrease in relaxation times. However,  $T_{1\rho}$  was reduced in the cortex of the tg animals compared to the

control animals at both 12 months ( $P < 0.05$ ) and 18 months ( $P < 0.01$ ) of age, when plaques are abundant in mice (18). A similar trend was observed in the hippocampus of the 12- and 18-month-old tg animals compared to their age-matched controls, but the difference was smaller ( $P < 0.05$  in both cases). The inter-age difference in relaxation times among the three age groups of tg animals is tabulated in Table 2. Statistically significant ( $P < 0.05$ ) differences were observed between the 12-month-old animals that exhibited an 11% decreased  $T_{1\rho}$  in the cortex than the 6-month-old tg animals and 10% decrease in the hippocampus. The 18-month-old animal had 16% and 8% lower  $T_{1\rho}$  in the cortex and hippocampus, respectively.

## DISCUSSION

In conclusion, analyses of  $T_{1\rho}$  values in this study showed that the difference in mean  $T_{1\rho}$  values between 18-month-old tg mice and their age-matched controls was greater than the  $T_2$  difference (3 msec) reported in a similar tg animal model (13). The lower dynamic range of spin-spin relaxation time could also explain why no difference was observed between AD-affected and normal human brain tissue specimens in  $T_2^*$ -weighted images (6).  $T_{1\rho}$  relaxation occurs in the presence of a spin-lock pulse that minimizes the effects of diffusion and susceptibility on the signal, which would explain why  $T_{1\rho}$  values are greater than the reported  $T_2$  values in the same animal. Further, at the spin-lock amplitude of 500 Hz used in the current experiments, the  $T_{1\rho}$  signal was sensitive to relaxation processes that occur in the milliseconds range (e.g., chemical exchange-mediated dipolar relaxation). The greater distribution of  $T_{1\rho}$  values further implies that  $T_{1\rho}$ -weighted MRI can be used to distinguish tissues with similar  $T_2$ . Incidentally, the increased distribution of  $T_{1\rho}$  values compared to  $T_2$  was also observed in the human brain (15).

The observed reduction in  $T_{1\rho}$  in the older tg animals may result not only from increased  $A\beta$  deposition, as demonstrated in Fig. 4a, but also from decreased blood flow and blood volume, which are associated with  $A\beta$  pathology (23,24). More studies are needed to conclusively determine the relevant relaxation processes and the degrees to which these factors affect the  $T_{1\rho}$  relaxation time. To this end, arterial spin-labeled (25) and spin-lock (26) MRI methods can be incorporated into the current protocol to quantify blood flow and perfusion.

The SNR of the images used to calculate  $T_{1\rho}$  maps has to be high to reduce the error in the measurement of  $T_{1\rho}$  via curve-fitting methods. Since noisier data values are introduced into the fitting procedure,  $T_{1\rho}$  maps appear more pixelated because identical tissues would erroneously exhibit different  $T_{1\rho}$  values. In the data presented, the longest-TSL image had an SNR of 16:1; however, higher SNR can be achieved by the use of surface coils for  $T_{1\rho}$ -weighted MRI (27), which is currently being developed for application in the tg animal model.

The MRI slice thickness is three times that of the histologic image, and the MR image's in-plane resolution is  $78 \mu\text{m} \times 78 \mu\text{m}$ . These factors lead to partial voluming and cause mismatching of some large plaques and inadequate resolution to visualize smaller plaques. However, this resolution cannot be achieved on clinical MRI scanners in a practical imaging time. Therefore, it is difficult to translate this technique to direct visualization of plaques in human patients. However, as the  $T_{1\rho}$ -weighted image of the tg mouse brain demonstrates, the presence of plaques reduces  $T_{1\rho}$  relaxation times and could therefore be indirectly quantified even with lower-resolution  $T_{1\rho}$  maps.

The results from this study will form the basis for extending this strategy to a large cohort of animals to study disease progression or response to therapeutic intervention, and eventually

to similar measurements in humans in vivo. However, extending  $T_{1\rho}$  mapping to clinical studies would require additional considerations, such as patient comfort and involuntary motion, which places constraints on the total imaging time. Furthermore, for clinical imaging the imaging protocol must not exceed FDA-mandated energy deposition levels (measured as the specific absorption rate (SAR)) (28). A protocol for mapping  $T_{1\rho}$  in the human brain images with submillimeter resolution in-plane and 2-mm slice thickness can be achieved in two minutes for each TSL by combining a fast spin-echo  $T_{1\rho}$  sequence with a low-SAR sequence (15,29). By employing such a protocol, the total MRI time for five different TSL images that will be needed to calculate  $T_{1\rho}$  maps is ~10 minutes.

An accurate comparison of measurements from histology and MRI necessitates a comparison of data generated with different orientations, slice thicknesses, and resolutions. Ideally, the same slice section thicknesses and orientations should be obtained by both methods for comparison. In the current experiments we achieved this by taking extreme care in aligning the animals inside the scanner and measuring the location of the imaged slice for subsequent histology. However, we will explore the application of several image registration methods in future studies to enable more accurate and reproducible comparisons of MRI and histology. Nonetheless,  $T_{1\rho}$  MRI holds promise for imaging AD to confirm the diagnosis of this disorder and perhaps detect it in its earliest stages without the need to inject tracers or contrast agents.

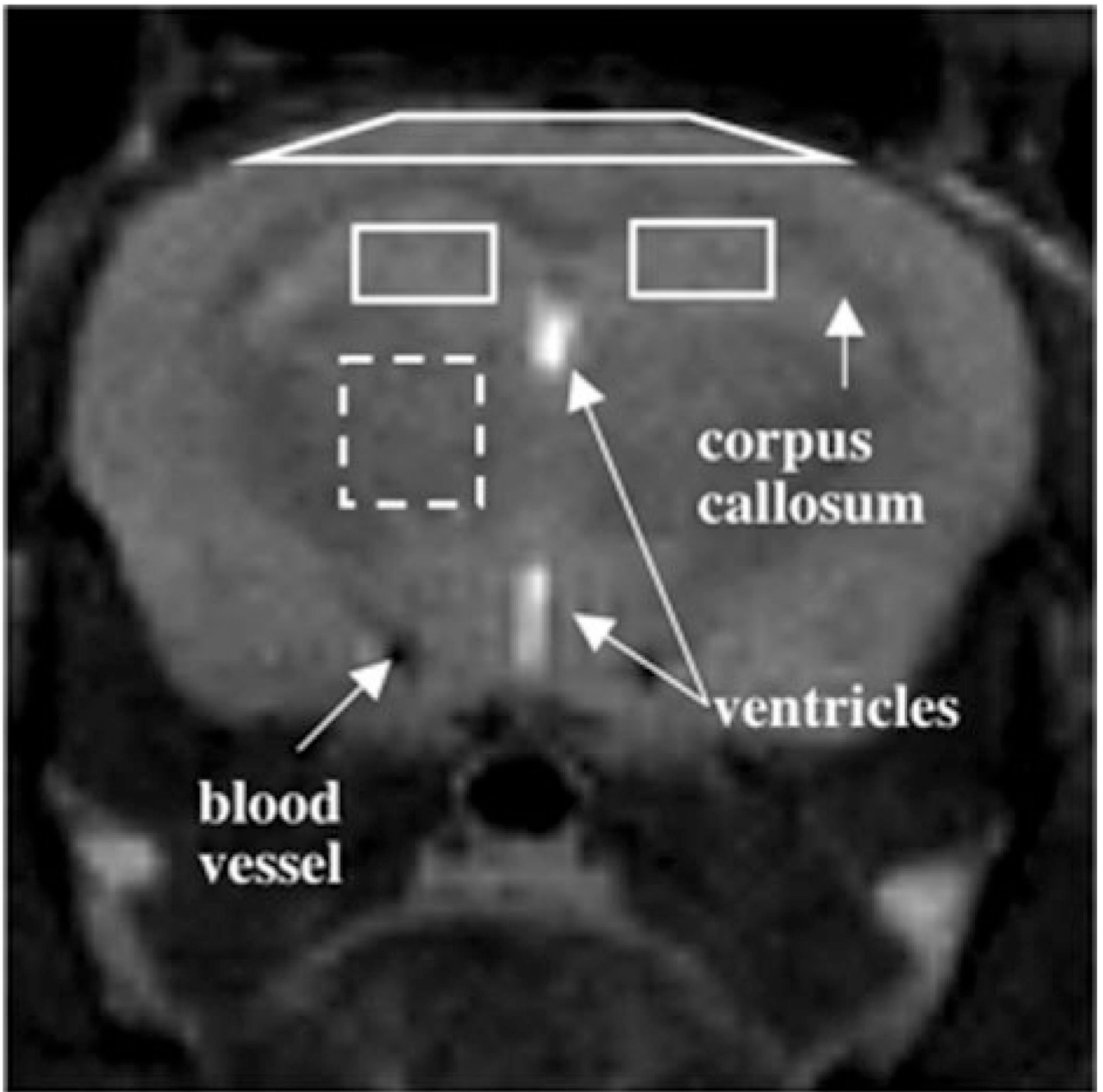
## Acknowledgments

We thank Professor John S. Leigh for his encouragement and support. This work was performed at the MMRRCC, an NIH regional resource (NIH RR02305), and the Center for Neurodegenerative Disease Research, an NIH-funded Alzheimer's Disease Center.

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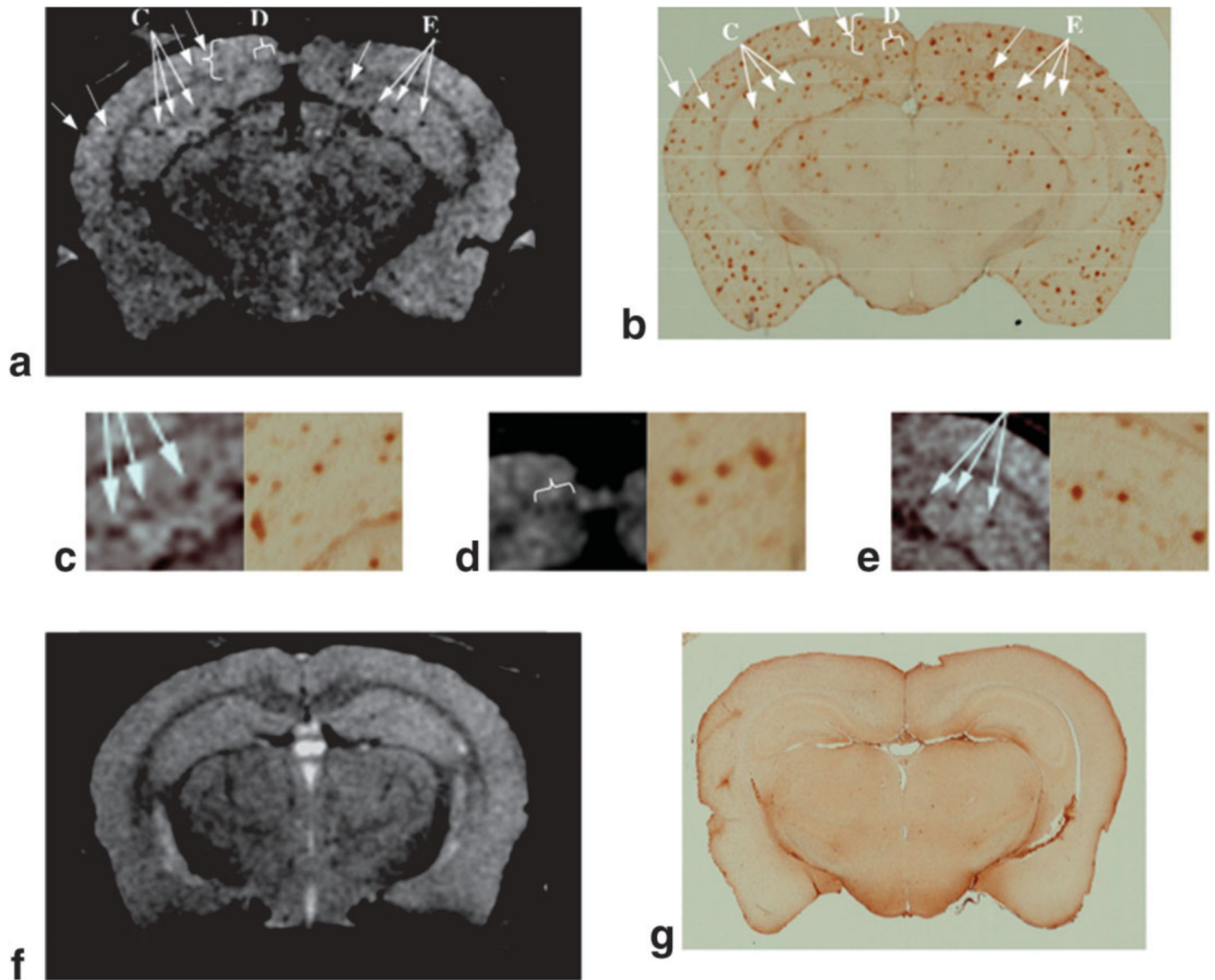
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**Figure 1.**

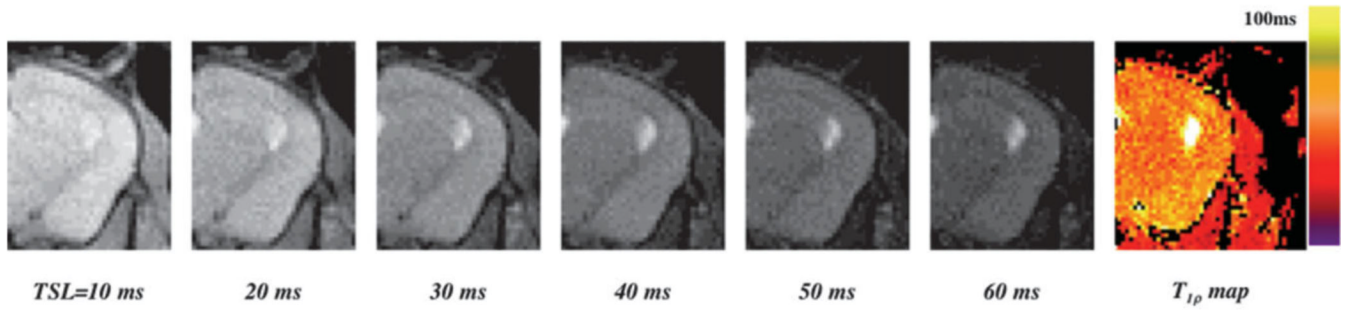
Location of the ROIs used for all data analyses overlaid on a  $T_{1\rho}$ -weighted image. Actual data were collected in the calculated  $T_{1\rho}$  map, an example of which is shown in color in Fig. 3. The trapezoid represents the boundary of the measurements performed in the cortex, the solid squares are located in the hippocampi, and the dashed square is in the thalamus. The total number of distinct pixels and therefore  $T_{1\rho}$  values in each location was 100.





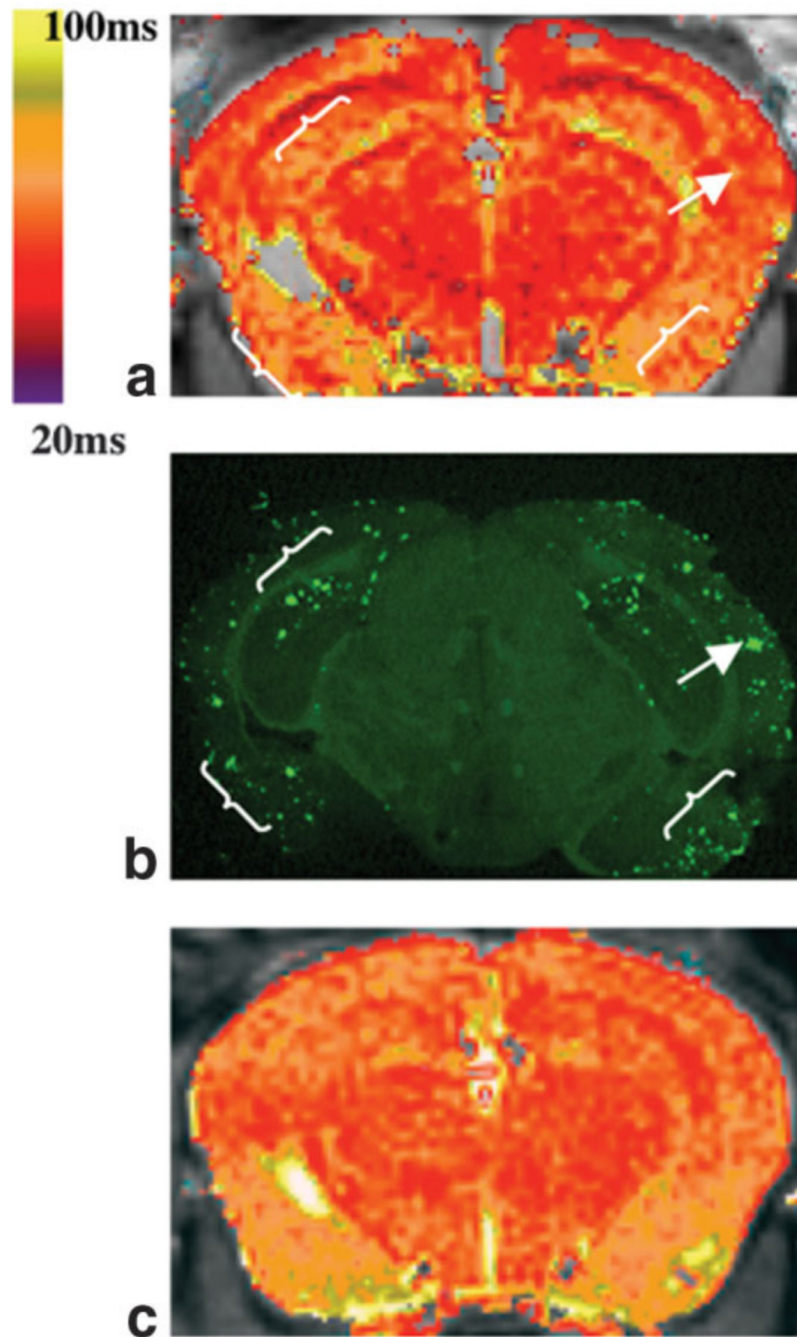
**Figure 2.**

**a:** A  $T_{1\rho}$ -weighted MR image of the APP/PS1 mouse brain shows hypointense regions, some of which are indicated by arrows. **b:** The corresponding histologic image created by combining five 10- $\mu$ m-thick immunostained sections shows abundant A $\beta$  deposits. **c–e:** The higher magnification views correspond to regions indicated by the arrows in part a to show identical clusters of A $\beta$  deposits in the hippocampus and cortex in both MR images and histological sections. The  $T_{1\rho}$ -weighted MR image of an age-matched control (**f**) and corresponding histology (**g**) failed to show SPs.



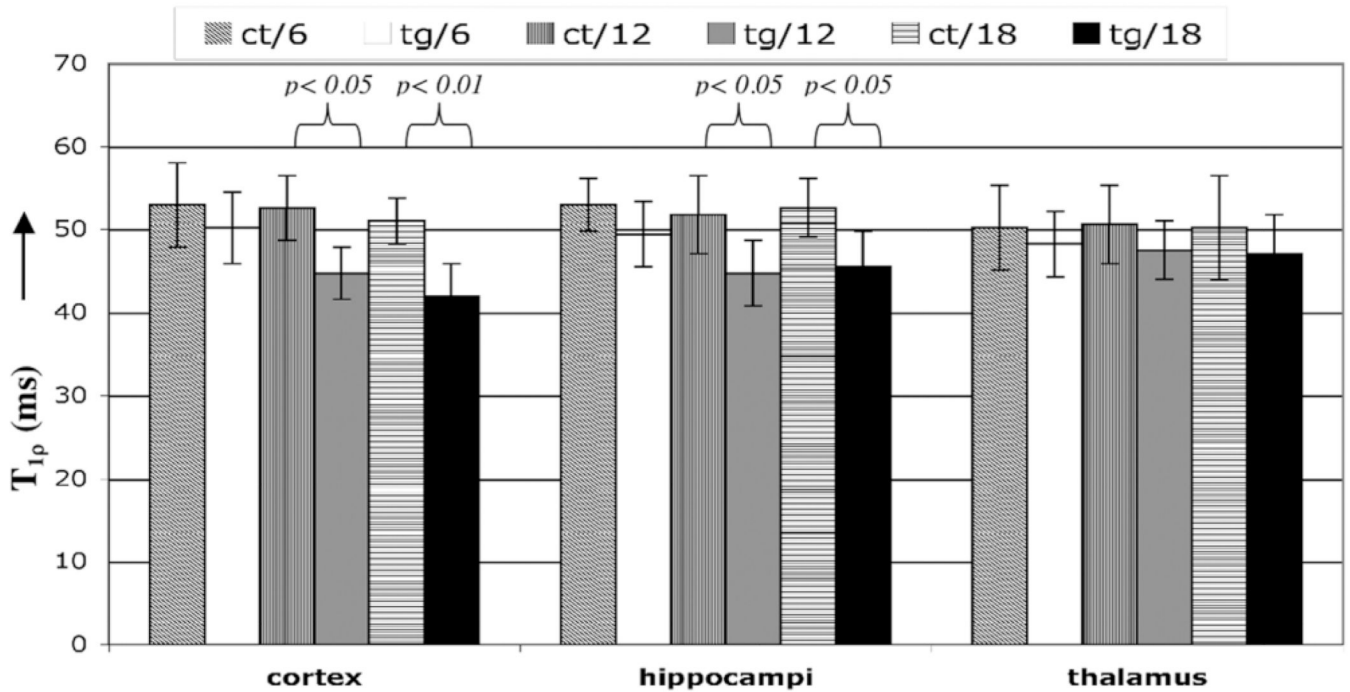
**Figure 3.**

Example of  $T_{1\rho}$  images from a single data set obtained at different durations of the spin-lock pulse (TSL) that was used to calculate the  $T_{1\rho}$  map (rightmost image, shown in color). In such maps, each pixel's intensity is the actual  $T_{1\rho}$  relaxation time constant of that pixel. Values above 100 msec, such as in fluid in the ventricles, are set to yellow, and pixels that did not correlate with TSL in the fitting routine, such as in the skull and background, were set to 0 (transparent).



**Figure 4.**

$T_{1\rho}$  color map overlaid on a grayscale  $T_{1\rho}$ -weighted image (a) and corresponding Thioflavin-S stained histology section (b) of an 18-month-old tg mouse, and a  $T_{1\rho}$  map of an age-matched control (c). During the MRI, identical slices were selected in both animals for meaningful comparisons. Although the histologic section is 40  $\mu\text{m}$  thick and the MRI slice thickness is 250  $\mu\text{m}$ , some of the pixels with low  $T_{1\rho}$  values are in the same location as large A $\beta$  deposits in the histology (indicated in a and b). The color bar-scale on the left indicates the range of  $T_{1\rho}$  values.



**Figure 5.**

$T_{1\rho}$  values averaged from ROIs in both animals, in the locations indicated. The maximum difference was observed between the 18-month-old tg animals and controls (9 msec), and was statistically significant ( $P < 0.01$ ). The difference between the 12-month-old tg mice and controls was significant in both the cortex and hippocampus. The thalamus had reduced  $T_{1\rho}$  values (~2 msec) in all age groups, but the difference was not significant.

**Table 1**Summary of the Results of  $T_{1\rho}$  Measurements

	$T_{1\rho}$ (msec)					
	6 month		12 month		18 month	
	Control (N = 2)	fg (N = 2)	Control (N = 2)	fg (N = 2)	Control (N = 2)	fg (N = 2)
Cortex	53.1 ± 5.1	50.2 ± 4.3	52.6 ± 3.9	44.7 ± 3.1	51 ± 2.8	41.9 ± 3.8
Hippocampus	52.9 ± 3.1	49.4 ± 3.7	51.8 ± 4.7	44.7 ± 3.9	52.5 ± 3.5	45.5 ± 4.3
Thalamus	50.2 ± 5.1	48.2 ± 3.9	50.6 ± 4.3	47.4 ± 3.5	50.2 ± 6.3	47.1 ± 4.7

**Table 2**

Interage Difference Among the Three Age Groups of tg Animals

	$\Delta T_{1\rho}$ (msec)	
	6 month tg	12 month tg
12 month tg	5.5 msec, $P < 0.05$ (cortex) 4.7 msec, $P < 0.05$ (hippocampus)	
18 month tg	8.3 msec, $P < 0.01$ (cortex) 3.9 msec, $P < 0.05$ (hippocampus)	2.8 msec, $P < 0.05$ (cortex) 0.8, not significant (hippocampus)