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Fast diffusion kurtosis imaging (DKI) with Inherent CORrelation-based Normalization (ICON) enhances automatic segmentation of heterogeneous diffusion MRI lesion in acute stroke

Iris Yuwen Zhou^{1,‡}, Yingkun Guo^{1,2,‡}, Takahiro Igarashi¹, Yu Wang^{1,3}, Emiri Mandeville⁴, Suk-Tak Chan¹, Lingyi Wen^{1,2}, Mark Vangel¹, Eng H Lo⁴, Xunming Ji³, and Phillip Zhe Sun^{1,2,3,4,*}

¹Athinoula A. Martinos Center for Biomedical Imaging, Department of Radiology, Massachusetts General Hospital and Harvard Medical School, Charlestown, MA, USA

²Department of Radiology, West China Second University Hospital, Sichuan University, Chengdu, Sichuan, China

³China-America joint neuroscience institute, Xuanwu Hospital, Capital Medical University, Beijing

⁴Neuroprotection Research Laboratory, Department of Radiology and Neurology, Massachusetts General Hospital and Harvard Medical School, Charlestown, MA, USA

Abstract

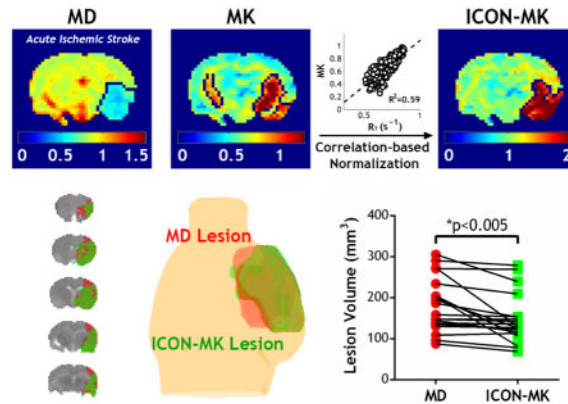
Diffusion kurtosis imaging (DKI) has been shown to augment DWI for defining irreversible ischemic injury. However, the complexity of cerebral structure/composition makes kurtosis map heterogeneous, limiting the specificity of kurtosis hyperintensity to acute ischemia. We proposed an Inherent CORrelation-based Normalization (ICON) analysis to suppress the intrinsic kurtosis heterogeneity for improved characterization of heterogeneous ischemic tissue injury. Fast DKI and relaxation measurements were performed on normal (N=10) and stroke rats following MCAO (N=20). We evaluated the correlations between mean kurtosis (MK), mean diffusivity (MD), fractional anisotropy (FA) derived from fast DKI sequence and relaxation rates of R_1 and R_2 , and found highly significant correlation between MK and R_1 ($P < 0.001$). We showed that ICON analysis suppressed the intrinsic kurtosis heterogeneity in the normal cerebral tissue, enabling automated tissue segmentation in an animal stroke model. We found significantly different kurtosis and diffusivity lesion volumes, being 147 ± 59 and 180 ± 66 mm³, respectively ($P = 0.003$, Paired-t test). The ratio of kurtosis to diffusivity lesion volume was $84 \pm 19\%$ ($P < 0.001$, One-sample t-test). We found relaxation normalized MK (RNMK) but not MD values significantly different between kurtosis and diffusivity lesions ($P < 0.001$, ANOVA). Our study showed that fast DKI with ICON analysis provides a promising means for demarcating heterogeneous DWI stroke lesion.

Graphical Abstract

*Corresponding author: Phillip Zhe Sun, Ph.D. (pzhesun@mgh.harvard.edu), 149 13th Street, Rm 2301, Charlestown, MA 02129, United States, Athinoula A. Martinos Center for Biomedical Imaging, Department of Radiology, MGH and Harvard Medical School, Phone: 617-726-4060, Fax: 617-726-7422.

[‡]These authors contributed equally.

Kurtosis augments DWI for defining irreversible ischemic injury. However, the complexity of cerebral structure/composition makes kurtosis map heterogeneous, limiting the specificity of kurtosis hyperintensity to acute ischemia. With strong correlation found between mean kurtosis and R_1 , we proposed an Inherent CORrelation-based Normalization (ICON) approach to mitigate the kurtosis heterogeneity in normal brain with substantially reduced scan time. We further demonstrated that this approach enabled automatic lesion segmentation and enhanced stratification of heterogeneous DWI lesion, aiding the translation of fast DKI to the acute stroke setting.



Keywords

Acute Stroke; Diffusion kurtosis imaging (DKI); Mean diffusivity (MD); Inherent CORrelation-based Normalization (ICON); Relaxation normalized mean kurtosis (RNMK); K-means Clustering

1. INTRODUCTION

Diffusion-weighted imaging (DWI) is sensitive to early ischemic tissue injury, and can detect ischemic lesion within minutes of the ischemic insult. Given its advantage in early detection of acute stroke, DWI has become one of the most widely used MRI techniques for stroke imaging (1–4). Various mismatch paradigms, including perfusion/diffusion, angiogram/diffusion and clinical/diffusion where the NIH Stroke Scale (NIHSS) used as a clinical indicator of infarction volume, have been proposed to guide stroke treatment beyond the standard therapeutic window (5–8). However, it has been recognized that ischemic tissue damage within the DWI lesion is heterogeneous (9–15). A portion of the DWI lesion represents injury that is reversible when reperfusion is prompt and effective, and the extent of DWI lesion reversibility is independently associated with improved outcome (16–19). As such, the underlying assumption that the DWI lesion identifies the irreversibly injured ischemic core is oversimplified, and advanced imaging techniques are urgently needed to refine diffusion stroke MRI (20).

Diffusion kurtosis imaging (DKI) has been introduced as a method to quantify non-Gaussian diffusion and has been applied to a number of neurological disorders including Parkinson's disease, Huntington's disease, traumatic brain injury and stroke (21–31). It has been shown that there is significant mismatch between mean diffusivity (MD) and mean kurtosis (MK) lesions in an animal model of acute stroke (32). In addition, diffusivity lesion without

kurtosis abnormality responds favorably to early reperfusion, where little change is found in the kurtosis lesion, suggesting that the kurtosis abnormality captures the more severely damaged ischemic core and hence, stratifies the conventional DWI lesion. However, unlike the relatively homogeneous trace diffusion image, the complexity of cerebral structure and composition leads to a heterogeneous kurtosis map, in which the specificity of kurtosis abnormality to ischemia is compromised, making automatic lesion segmentation somewhat difficult. A means to minimize the intrinsic cerebral tissue kurtosis variation would thus enhance the conspicuity of the ischemic kurtosis lesion.

The conventional DKI protocol requires collecting DWI images with multiple b-values along at least fifteen directions, resulting in relatively long acquisition time (21). Recently, a fast DKI protocol has been introduced by Hansen et al. for the rapid mapping of MD and MK (33,34). The diffusivity and kurtosis maps obtained with the fast DKI agreed well with those estimated from conventional DKI using the diffusion tensor model (MD_{tensor} and MK_{tensor}) in normal brain (33), in glioma (35) and importantly in acute stroke brain (33,34,36). We have further shown that kurtosis is one of the most sensitive indices in capturing acute ischemic lesion (37). Herein we developed a relaxation-normalized fast DKI approach to refine acute ischemic lesion stratification. Briefly, we evaluated the correlation between cerebral MK and multiple MRI indices including MD, fractional anisotropy (FA), and relaxation (R_1 , R_2). There were significant correlation between MK and R_1 , MK and R_2 , and MK and FA, with R_1 showing the highest correlation coefficient. Because MD and FA may alter substantially during acute stroke while the change in relaxation rate is relatively small, we proposed an Inherent CORrelation-based Normalization (ICON) analysis and validated it in suppressing the intrinsic kurtosis heterogeneity. We demonstrated that the ICON approach enabled automatic ischemic lesion segmentation during acute ischemic stroke, superior to the MK-based lesion analysis. In summary, our study developed relaxation-normalized fast kurtosis imaging for characterizing heterogeneous ischemic tissue injury, aiding the translation of fast diffusion kurtosis imaging to the acute stroke setting.

2. MATERIALS AND METHODS

Animals

All experiments have been approved by the Institutional Animal Care and Use Committee. Adult male Wistar rats (Charles River Laboratory, Wilmington, MA) were anesthetized throughout the surgery and the MRI experiments with 1.5–2.0% isoflurane/air mixture. Animal body temperature was maintained by a circulating warm water jacket positioned around the torso. We imaged ten normal rats (N=10) and twenty three stroke rats following the standard permanent middle cerebral artery occlusion (MCAO, N=23). Briefly, MCAO was induced by inserting a 4–0 silicon-coated nylon suture (Doccol Corp, Sharon, MA) into the lumen of the right internal carotid artery, and then advancing it to block the origin of the middle cerebral artery. Stroke rats were then transferred to the MRI scanner after the surgery and imaged 1.5 to 2 hours after the onset of occlusion. Peripheral oxygen saturation (SpO₂) and heart rate were continuously monitored (Nonin Pulse Oximeter 8600, Plymouth, MN). The SpO₂ was not significantly different between the two groups, being $95 \pm 4\%$ (normal)

vs. $92 \pm 3\%$ (MCAO). Three animals were excluded from analysis due to failed MCAO surgery.

MRI

MRI data were acquired using a 4.7 Tesla small-bore MRI scanner (Bruker Biospec, Billerica, MA). We used a dual RF coil setup, including a 70 mm volume transmitter coil and an actively-decoupled 20 mm surface receiver coil. During MRI, the animals were placed on a plastic cradle with the head fixed with a tooth bar and ear bars to minimize motion. Multi-slice MRI (5 slices, slice thickness/gap = 1.8/0.2 mm, field of view = 20×20 mm², acquisition matrix = 48×48) was acquired with single-shot echo-planar imaging (EPI) (receiver bandwidth = 225 kHz), with the central slice positioned at 2 mm posterior to bregma. Fast DKI was acquired using three b-values: one $b = 0$ s/mm² image, nine $b = 1000$ s/mm² and nine $b = 2500$ s/mm² images along directions of $\hat{n}^{(i)}$, $\hat{n}^{(i+)}$, $\hat{n}^{(i-)}$ (i.e. $\hat{n}^{(i)} = (0,1,1)^T$, $\hat{n}^{(i+)} = (0,1,1)^T$, and $\hat{n}^{(i-)} = (0,1,-1)^T$, and similarly for $i=2$ and 3) (38). Note that the superscript i in $\hat{n}^{(i)}$ labels the position of the “1” while in $\hat{n}^{(i+)}$ and $\hat{n}^{(i-)}$ it labels the position of the “0” (33,34). The other parameters of the fast DKI approach were gradient pulse duration/diffusion time (δ/Δ) = 6/20 ms, repetition time (TR)/echo time (TE) = 2500/36.6 ms, 4 averages, scan time = 3 min 10 s. In addition, T_1 -weighted images were acquired using an inversion recovery sequence, with seven inversion delays ranging from 250 to 3000 ms (TR/TE = 6500/14.8 ms, 4 averages, scan time = 3 min 38 s); T_2 -weighted SE images were obtained with two TEs of 30 and 100 ms (TR = 3250 ms, 4 averages, scan time = 26 s). Perfusion imaging was acquired with amplitude modulated arterial spin labeling MRI (TR/TS/TE = 6500/3250/14.8 ms, 32 averages, scan time = 7 min).

Image and Data Analysis

Images were analyzed in MATLAB (MathWorks, Natick, MA). MD and MK maps were obtained by linear combination of log diffusion signals, as described previously (33,34,36,37,39). FA was estimated based on standard diffusion tensor model. In addition, parametric T_1 map was derived using least-squares fitting of the signal as functions of inversion time ($I = I_0 [1 - (1-\eta)e^{-TI/T_1}]$), where η is the inversion efficiency, TI is the inversion time, and I_0 is equilibrium signal. T_2 map was calculated as

$T_2 = \frac{TE_2 - TE_1}{\ln(I(TE_1)/I(TE_2))}$, where $I(TE_{1,2})$ are T_2 -weighted signals obtained at different echo times (TE = 30 and 100 ms). The relationship between MK and other MRI indices (T_1 , T_2 , MD and FA) was evaluated using Pearson’s correlation, excluding ventricular area with MD value above $1 \mu\text{m}^2/\text{ms}$. An estimated MK map (MK_{est}) was calculated using the regression coefficients determined from the univariate linear regression analysis between R_1 and MK per pixel (i.e. $MK_{\text{est}} = C_0 + C_1 \cdot R_1$, where C_0 and C_1 are regression coefficients) and then the relaxation-normalized MK (RNMK) map was calculated as $RNMK = MK/MK_{\text{est}}$. Cerebral blood flow (CBF) map was obtained from amplitude modulated arterial spin labeling MRI

(40) and derived as $CBF = \frac{\lambda (I_{\text{ref}} - I_{\text{tag}})}{2\alpha \cdot I_{\text{ref}}} \cdot \frac{e^{w/T_{1a}}}{T_{1w}}$, where I_{tag} is the label image, I_{ref} is the reference image, λ is the brain-blood partition coefficient for water, α is the degree of inversion with transient time correction, w is the post-labeling delay, and T_{1a} is the arterial blood longitudinal relaxation time (41).

Results were expressed as mean \pm standard deviation (S.D.). We evaluated the association between MK and relaxation, MD and FA with univariate and multiple linear regressions, and *P*-values of less than 0.05 were considered statistically significant. The size of ischemic lesions in MD, MK or RNMK maps were automatically defined using a K-means clustering-based algorithm (42). Briefly, a 1D vector of all the pixels within the brain was formed and clustering algorithm used an iterative refinement technique to partition the data points in the vector into two clusters (lesion vs. normal tissue) based on their intensities.

3. RESULTS

Fig. 1 shows parametric brain T_1 , T_2 , MD, FA, and MK maps from a representative normal Wistar rat. Because cerebral spinal fluid (CSF) has significantly higher water content, the ventricle region appears hyperintense in T_1 , T_2 and MD images. In addition, CSF displays hypointensity in FA and MK maps due to little anisotropic/restricted diffusion. Whereas the contrast between striatum or corpus callosum regions and cortical region was relatively small in T_2 and MD maps, it was distinct in T_1 , FA and MK maps. This suggests that kurtosis, a measure of non-Gaussian diffusion, is partially associated with diffusion fractional anisotropy, a widely-used measure of gross white matter microstructure (43–45) along with longitudinal relaxation rate, which has been shown to be associated with semisolid macromolecules and water content (46,47).

We evaluated the relationship between MK and other MRI indices using Pearson's correlation with a Student's *t*-distribution, excluding ventricular regions using an MD threshold of $1 \mu\text{m}^2/\text{ms}$ (Fig. 2a). There was a weak correlation between MK and MD ($R^2=0.11$, $P<0.001$), consistent with previous report suggesting that diffusivity and kurtosis are two distinct diffusion-based indices for characterizing tissue microstructure (48). The pixel-wise analysis also showed a weak correlation between MK and FA ($R^2=0.17$, $P<0.001$), moderate correlation between MK and R_2 ($=1/T_2$, $R^2=0.37$, $P<0.001$) and strong correlation between MK and R_1 ($=1/T_1$, $R^2=0.70$, $P<0.001$). The correlation between MK and R_1 was significantly higher than that between MK and MD, FA or R_2 in normal animals ($N=10$, ANOVA with multiple comparison test, $P<0.001$). Because MD and FA may change substantially during acute stroke, we compared the univariate linear regression of MK and R_1 versus multiple linear regression of MK with R_1 and R_2 in all the normal animals (Fig. 2b). We found that the coefficient of determination (i.e., R^2) was 0.62 ± 0.08 (MK and R_1 , $P<0.001$) and 0.62 ± 0.08 (MK, R_1 and R_2 , $P<0.001$), showing little difference (Two-sample *t*-test, $P=0.92$). Hence, we chose the univariate linear regression of MK and R_1 to correct for intrinsic kurtosis heterogeneity in intact cerebral tissue.

Fig. 3 compares the routine MK and relaxation-normalized MK maps in a normal Wistar rat. Both the relaxation R_1 (i.e., $1/T_1$) map (Fig. 3a) and the MK map (Fig. 3b) show hyperintensity in striatum and corpus callosum. Fig. 3c shows the estimated MK map (MK_{est}) using the regression coefficients determined from univariate linear regression analysis between R_1 (Fig. 3a) and MK (Fig. 3b) per pixel ($\text{MK}_{\text{est}}=1.42 \cdot R_1 - 0.26$). Fig. 3d shows the proposed relaxation-normalized MK map (i.e. $\text{RNMK}=\text{MK}/\text{MK}_{\text{est}}$), which was substantially more homogeneous than the MK map without relaxation normalization (Fig. 3a). For all the normal animals studied, the coefficient of variation (COV, i.e., S.D./mean)

was $23.6 \pm 2.3\%$ and $12.1 \pm 2.8\%$ for MK and RNMK maps, respectively. This represents $48.6 \pm 10.2\%$ decrease in COV, showing that the ICON analysis removes a substantial portion of kurtosis heterogeneity which is intrinsic to normal cerebral tissue.

We further evaluated the ICON analysis in a representative acute stroke animal within 2 hrs after ischemia. R_1 (Fig. 4a) and R_2 (Fig. 4b) maps show relatively small change during acute ischemic stroke, with a reduction of 1.8 % in R_1 and 0.1% in R_2 in the ischemic lesion. In comparison, substantial reduction in MD map was observed in the ischemic lesion (Fig. 4c). The diffusivity lesion was 42.7 mm^3 for the slice. The MK map is heterogeneous, particularly in the internal capsule and striatum, which leads to overestimated kurtosis lesion (Fig. 4d). While the ipsilateral cortex shows relatively increased kurtosis from that of the contralateral cortex, the K-means clustering algorithm missed part of the cortex lesion but incorrectly included the contralateral internal capsule and ipsilateral striatum to the kurtosis lesion (Fig. 4d, arrows), resulting in kurtosis lesion substantially larger than the diffusivity lesion (54.2 vs. 42.7 mm^3). Fig. 4e demonstrates significant correlation between R_1 and MK indices from the contralateral normal brain ($R^2=0.59$, $P<0.001$). For the contralateral brain, MK and RNMK values were found to be 0.67 ± 0.13 and 1.00 ± 0.12 , respectively. This represents a relative COV decrease of 42%, similar to that found in intact brains. The kurtosis lesion was segmented from RNMK map, using the K-means clustering algorithm (Fig. 4f). Compared to the lesion segmented in Fig. 4d which inaccurately includes part of the contralateral normal tissue, the ICON analysis corrected for such error and resulted in a kurtosis lesion size of 41 mm^3 , which is close to but slightly smaller than the MD lesion (as seen in Fig. 4c and 4f).

We further tested relaxation-normalized kurtosis analysis for multi-slice segmentation in acute stroke rats. Fig. 5a shows the outlined diffusivity and kurtosis lesions superimposed on a scout scan, in which diffusivity and kurtosis lesions appear in red and green, respectively. Fig. 5b shows substantial kurtosis/diffusivity mismatch. Fig. 5c compares diffusivity and kurtosis lesion volumes in all stroke animals. Specifically, we found kurtosis lesion significantly smaller than that of diffusivity lesion (147 ± 59 vs. $180 \pm 66 \text{ mm}^3$, $P=0.003$, Paired-t test). The ratio of kurtosis to diffusivity lesion volumes was $84 \pm 19\%$ ($P<0.001$, One-sample t-test). Moreover, we compared the CBF, MD, RNMK values in different brain regions including contralateral normal, diffusivity lesion, kurtosis lesion and their mismatch (Table 1). MD was 0.63 ± 0.06 and $0.65 \pm 0.06 \text{ } \mu\text{m}^2/\text{ms}$ in diffusivity and kurtosis lesions (n.s., ANOVA) while their RNMK values were significantly different (1.60 ± 0.27 vs. 1.77 ± 0.21 , $P<0.05$, ANOVA). The kurtosis/diffusivity lesion mismatch had a relatively higher MD value ($p<0.05$, ANOVA) while its RNMK value was significantly lower than that of diffusivity lesion ($p<0.001$, ANOVA) and kurtosis lesion ($p<0.001$, ANOVA). Our results demonstrated that relaxation-normalized kurtosis imaging effectively reduced the intrinsic kurtosis heterogeneity, enabling automatic tissue segmentation of acute stroke DKI.

4. DISCUSSION

Our results demonstrated significant correlation between mean kurtosis and longitudinal relaxation, and that relaxation-normalized fast DKI reduces the intrinsic kurtosis heterogeneity, facilitating automatic segmentation of heterogeneous ischemic tissue. In

water content may affect multiple MRI indices of healthy brain, which may partially explain the correlation among the measurements. Additional MRI indices such as magnetization transfer (MT) and amide proton transfer MRI which contain information about the chemical and physical environments of macromolecular protons in tissue may further enhance the correlation (58–60) and hence improve suppression of intrinsic heterogeneity in kurtosis map and specificity of MK changes to ischemic tissue injury. It is necessary to point out that multi-parametric MRI acquisition beyond the highly correlated T_1 and kurtosis MRI will prolong the scan time, limiting its use during acute stroke imaging. Techniques such as fingerprinting MRI may make the proposed approach more applicable in the emergency settings (61).

Accurate identification of the salvageable ischemic tissue is crucial in the clinical settings, particularly with the recent advancement in embolectomy. In this work, we studied acute ischemic tissue injury within 2 hours from occlusion using a well-established filament MCAO model. This time window is similar to the hyperacute phase of stroke patients, which is the most critical period for effective treatment. Although reasonably reproducible, the filament model induces severe hypoperfusion and, arguably, significantly reduces the salvageable tissue and shortens the treatment window. This limitation may be reasonably addressed by using an embolic stroke model, albeit its moderately higher variability (62). Another advantage of using the embolic model is that it enables evaluation of thrombolytic therapy that mimics recanalization in stroke patients. More importantly, our method can be used to study the irreversible and reversible ischemic lesion in the embolic model with thrombolytic treatment, especially beyond the hyperacute window before significant T_1 change occurs, potentially extending the treatment window. On the other hand, there may be substantial relaxation changes in the ischemic lesion at extended stroke onset time, which can be largely attributable to edema formation. The performance of ICON approach in subacute stroke imaging needs to be investigated in the future. In the current study, no histology was obtained because commonly used Hematoxylin and eosin (H&E) and TTC staining techniques cannot reliably identify acute ischemic tissue injury. Notably, before we can investigate histological changes, we need to develop a reliable lesion segmentation approach to guide histology, which is the goal of our current study. It would be interesting to confirm the tissue classification and reversibility of ischemic brain using more sensitive immunohistological techniques in the future.

In this study, we chose a relatively short diffusion time, which is desirable given that brain T_2 at 4.7 T is shorter than that at 1.5 and 3 T. Recently, Baron et al. found reduced diffusion contrast in stroke patients under short diffusion times (63). Latt et al. showed that DWI acquired with long diffusion time is susceptible to intra- and extra-cellular water exchange, especially in regions with compromised cell membranes (64). In addition, tissue with intact but functionally impaired and nearly impermeable membranes shows significantly reduced ADC. Consequently, long diffusion time may enhance diffusion contrast with elevated heterogeneity, for which the proposed ICON analysis might provide crucial information for tissue characterization. To summarize, the proposed relaxation-normalized fast kurtosis imaging is promising to aid the evaluation of fast DKI and its translation to the acute stroke setting.

5. CONCLUSION

Our study demonstrated that fast kurtosis imaging with Inherent CORrelation-based Normalization (ICON) reasonably corrects for the intrinsic regional kurtosis variation, enabling automatic segmentation of ischemic kurtosis lesion. We confirmed significant kurtosis/diffusivity lesion mismatch, which is a promising approach to elucidate the diagnostic value of fast DKI in acute stroke prior to its translation to the clinical setting.

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ABBREVIATIONS

ADC	Apparent Diffusion Coefficient
CBF	Cerebral Blood Flow
COV	Coefficient of Variation
CSF	Cerebrospinal Fluid
DKI	Diffusion Kurtosis Imaging
DWI	Diffusion-weighted Imaging
EPI	Echo Planar Imaging
FA	Fractional Anisotropy
MCAO	Middle Cerebral Artery Occlusion
MD	Mean Diffusivity
MK	Mean Kurtosis
TE	Echo Time
TR	Repetition Time

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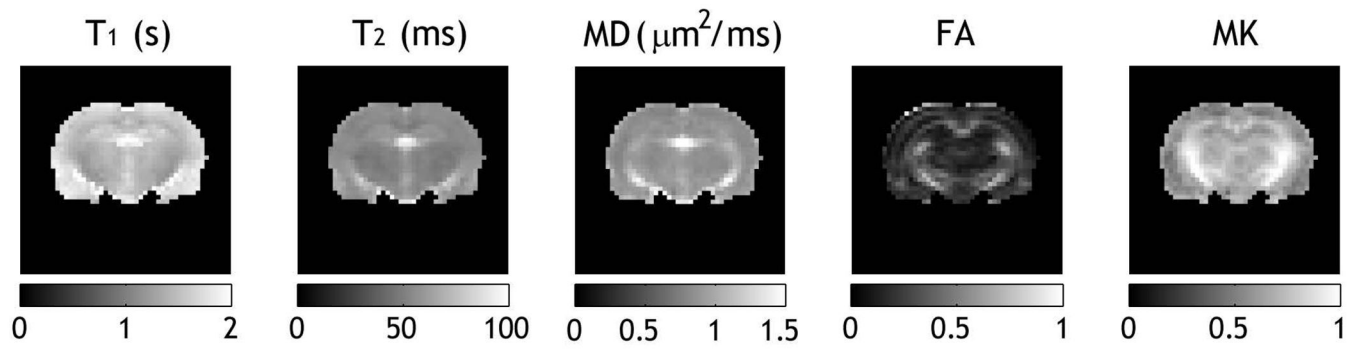


Fig 1.
Comparison of T₁, T₂, mean diffusivity (MD), fractional anisotropy (FA) and mean kurtosis (MK) maps of a representative normal rat.

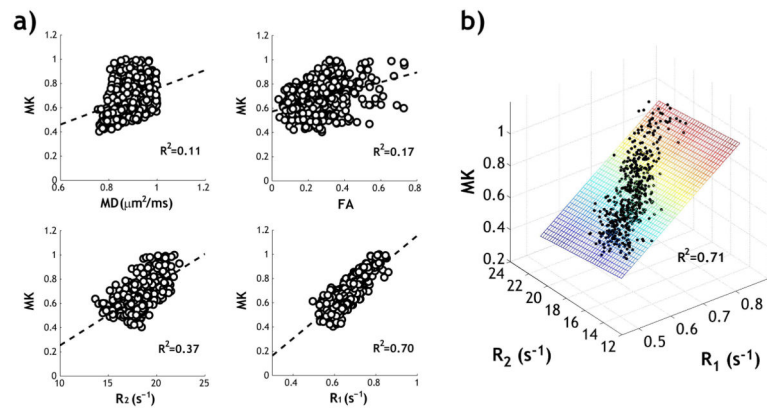


Fig 2.

a) Univariate linear regression analysis between mean kurtosis and multiple MRI indices, including MD, FA, R_2 ($=1/T_2$) and R_1 ($=1/T_1$) of a representative normal rat. **b)** Multiple linear regression analysis between mean kurtosis and R_1 , R_2 of the same animal.

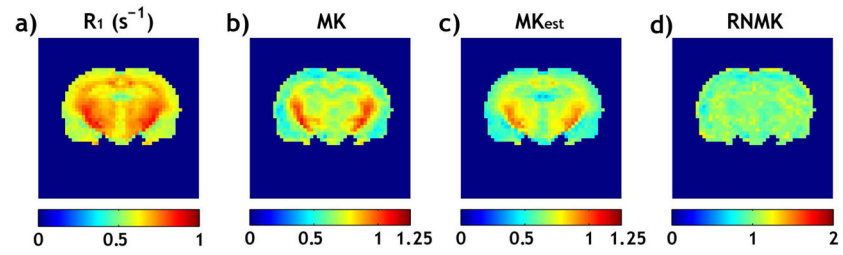


Fig 3.

Fast DKI with Inherent CORrelation-based Normalization (ICON) analysis in a representative normal rat. a) R_1 map. b) MK map with substantial regional variation. c) MK map estimated from R_1 map. d) Relaxation-normalized MK (RNMK) map ($RNMK = MK/MK_{est}$), which was substantially more homogeneous.

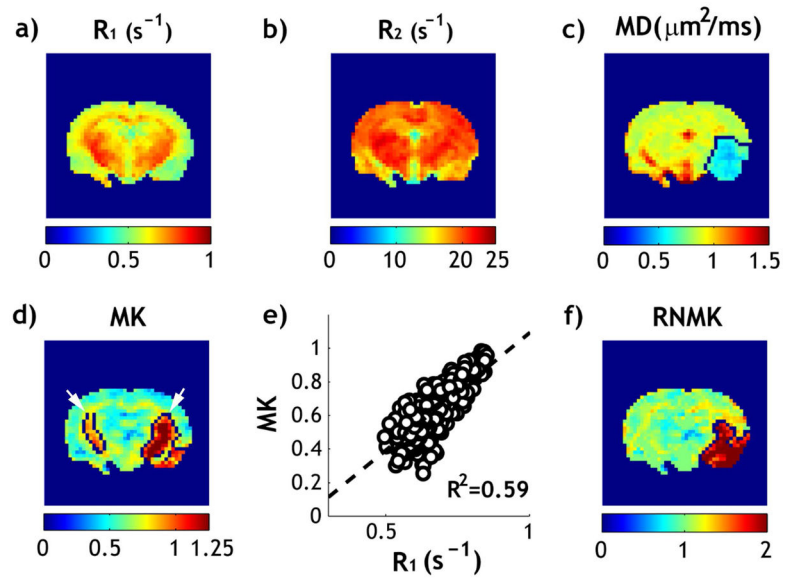


Fig 4. Relaxation-normalized fast DKI of a representative rat after acute ischemic stroke. a) R_1 map. b) R_2 map. c) MD map with outlined ischemic lesion. d) MK map with outlined ischemic lesion. e) Linear regression analysis of R_1 and MK in the contralateral normal brain. f) RNMK map with outlined ischemic lesion.

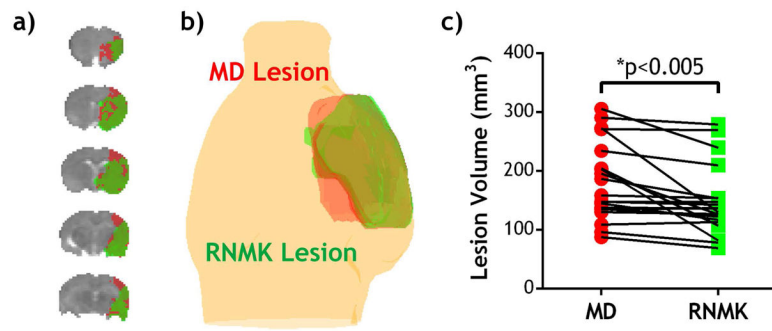


Fig 5. Ischemic tissue segmentation using the proposed relaxation-normalized kurtosis MRI. a) MD lesion (red) vs. RNMK lesion (green). b) 3-D overlay of diffusion and kurtosis lesions. c) Comparison of ischemic diffusivity and kurtosis lesions.

Table 1

Comparisons of cerebral blood flow (CBF), MD and RNMK values in different brain regions.

N=20	Contralateral Normal Region	Diffusivity Lesion	Kurtosis Lesion	Kurtosis/Diffusivity Lesion Mismatch
CBF (ml/g/min)	1.27 ± 0.42	0.64 ± 0.40 ^{***}	0.59 ± 0.37 ^{***}	0.79 ± 0.51 ^{**}
MD (μm ² /ms)	0.86 ± 0.03	0.63 ± 0.06 ^{***}	0.65 ± 0.06 ^{***}	0.68 ± 0.06 ^{***†}
RNMK	1.00 ± 0.01	1.60 ± 0.27 ^{***}	1.77 ± 0.21 ^{***§}	1.17 ± 0.15 ^{*§§§†††}

One-way ANOVA with Bonferroni's multiple comparisons test was performed.

^{***}
p<0.001,

^{**}
p<0.01,

^{*}
p<0.05 significantly different from contralateral normal region;

^{§§§}
p<0.001,

[§]
p<0.05 significantly different from diffusivity lesion;

^{†††}
p<0.001,

[†]
p<0.05 significantly different from kurtosis lesion. Note all the measurements passed D'Agostino-Pearson normality test, indicating Gaussian distribution.