Clinical Implications of Non-Steatotic Hepatic Fat Fractions on Quantitative Diffusion-Weighted Imaging of the Liver

Hildebrand Dijkstra1,2*, Astri Handayani1, Peter Kappert2, Matthijs Oudkerk1, Paul E. Sijens2

1 Center for Medical Imaging - North East Netherlands, University of Groningen, University Medical Center Groningen, Groningen, The Netherlands, 2 Department of Radiology, University of Groningen, University Medical Center Groningen, Groningen, The Netherlands

Abstract

Diffusion-weighted imaging (DWI) is an important diagnostic tool in the assessment of focal liver lesions and diffuse liver diseases such as cirrhosis and fibrosis. Quantitative DWI parameters such as molecular diffusion, microperfusion and their fractions, are known to be affected when hepatic fat fractions (HFF) are higher than 5.5% (steatosis). However, less is known about the effect on DWI for HFF in the normal non-steatotic range below 5.5%, which can be found in a large part of the population. The aim of this study was therefore to evaluate the diagnostic implications of non-steatotic HFF on quantitative DWI parameters in eight liver segments. For this purpose, eleven healthy volunteers (2 men, mean-age 31.0) were prospectively examined with DWI and three series of in-/out-of-phase dual-echo spoiled gradient-recalled MRI sequences to obtain the HFF and T2*. DWI data were analyzed using the intravoxel incoherent motion (IVIM) model. Four circular regions (ø22.3 mm) were drawn in each of eight liver segments and averaged. Measurements were divided in group 1 (HFF ≤2.75%), group 2 (2.75< HFF ≤5.5%) and group 3 (HFF>5.5%). DWI parameters and T2* were compared between the three groups and between the segments. It was observed that the molecular diffusion (0.85, 0.72 and 0.49 x10^-3 mm^2/s) and T2* (32.2, 27.2 and 21.0 ms) differed significantly between the three groups of increasing HFF (2.18, 3.50 and 19.91%). Microperfusion and its fraction remained similar for different HFF. Correlations with HFF were observed for the molecular diffusion (r = -0.514, p<0.001) and T2* (−0.714, p<0.001). Similar results were obtained for the majority of individual liver segments. It was concluded that fat significantly decreases molecular diffusion in the liver, also in absence of steatosis (HFF=5.5%). Also, it was confirmed that fat influences T2*. Determination of HFF prior to quantitative DWI is therefore crucial.

Introduction

The effect of fat on the self-diffusion of water has been assessed since the onset of nuclear magnetic resonance. Already in 1983, it was demonstrated that water diffusion drops six-fold inside of Cheddar and Swiss cheeses [1]. Later it was observed in vitro that water diffusion is hindered by lipid-rich cores in susceptible plaque [2]. The clinical assessment of water diffusion in the liver became feasible with the introduction of diffusion weighted imaging (DWI) in the abdomen [3]. DWI reflects the mobility of water molecules (molecular diffusion) in a tissue which can be described by the apparent diffusion coefficient (ADC) or the intravoxel incoherent motion (IVIM) model [4–6]. Since then DWI has been successfully applied in the assessment of focal liver lesions and diffuse liver diseases such as cirrhosis, fibrosis and steatosis [7–11]. However, the effect of fat on hepatic DWI is still subject of debate. In an animal study it was concluded that steatosis may confound determination of hepatic fibrosis with DWI [12]. This was confirmed in two clinical studies where the ADC decreased significantly in patients with hepatic steatosis [13,14]. Similarly, a study which applied the IVIM model demonstrated that steatosis can reduce the molecular diffusion significantly and thus act as a potential confounder when IVIM is used to assess diffuse liver diseases such as cirrhosis [15]. These studies discussed the effect of hepatic fat on DWI for patients with steatosis, which is defined as fat fractions higher than 5.56% [16]. However, a detailed insight in the dependency of IVIM parameters on normal (non-steatotic) fat fractions ranging between 0 and 5.56% has not been reported yet; it has been suggested however that there might be a nonlinear relationship [15]. Also, the relation between the different segmental regions of the liver and the effect of fat on IVIM modelled DWI has not been studied up to now. Considering that fat content has been demonstrated to differ between liver segments, its effect on IVIM modelled DWI may be expected to be location dependent [17]. In addition, it has been reported that next to the effects of fat on diffusion, fat also affects T2* estimation [18–20]. The purpose of this study was therefore to evaluate the diagnostic implications of non-steatotic HFF (< 5.5%) on quantitative DWI by assessing the HFF and T2* of healthy subjects in eight liver segments.
Materials and Methods

Ethics statement

The protocol of the study was approved by the Medical Ethics Review Board of the University Medical Center Groningen, and written informed consent was obtained for each volunteer.

Study population

In April 2011, healthy volunteers were randomly selected by local advertisement in the university to ensure a diverse population. Volunteers were required to be without any history of hepatic pathology or any other pathology related to liver function. The minimum age for inclusion was 18 years old. Exclusion criteria included MRI contra-indications such as pacemakers, clips, stents and implants. In total, 11 subjects were included (2 men) with an age between 18 to 56 years old (mean 31.0) and a body mass between 55 and 116 kg. Body mass-index (BMI) ranged between 19.9 and 34.4 kg/m² (mean 25.4 kg/m²). The only preparation before the examination was an 8-h fasting period.

MR protocols

All subjects were prospectively examined on a 1.5 T MRI system (Magnetom Avanto, Siemens Medical Solutions, Erlangen, Germany). The body coil served as transmitter and a 24-element spine matrix coil in combination with a 6-element body matrix as receiver.

After the localiser scans, a series of diffusion weighted images (DWI) were obtained using a spin echo based single shot echoplanar imaging (SS-EPI) sequence in combination with spectral adiabatic inversion recovery (SPAIR) fat suppression. The DWI acquisitions (b = 0, 50, 100, 250, 500, 750 and 1000 s/mm²) were gated using PACE respiratory triggering (TR = 3065–5947 ms) and tuned with the following parameters: TE 90 ms; FA 90°; slice-thickness 5 mm; FOV 300×242 mm²; matrix 144×116; bandwidth 1335 Hz/pixel; 4 averages and parallel acquisition technique GRAPPA with acceleration factor 2. Diffusion gradients (25 mT/m) were applied in the phase-, read-, and z-directions respectively using bipolar diffusion-encoding schemes. For each subject, 16 transverse slices were acquired in interleaved mode to cover the liver in an acquisition time between 7.2 and 13.5 minutes.

After the DWI scans, a dual-echo spoiled gradient recalled (SPGR) sequence was acquired to obtain two series of in-phase (IP) images with echo times of 4.5 and 18 ms, TR = 220 ms and FA = 70° to calculate T₂*. Then, to calculate the hepatic fat fraction (HFF), a second dual-echo SPGR was acquired to obtain two series of both out-phase (OP) and in-phase (IP) images with echo times of 2.38 and 4.76 ms respectively tuned with TR = 206 ms and FA = 70°. Finally, a third series of dual-echo SPGR was acquired with equal TE/TR settings as the second series hence a flip angle of 20°. All three SPGR scans were acquired with slice-thickness 6 mm; FOV 375×196 mm²; matrix 256×134; bandwidth 434 Hz/pixel; 1 averages and parallel acquisition technique GRAPPA with acceleration factor 2 and an acquisition time between 1.5 and 2.0 minutes. Total acquisition time for DWI and HFF measurements was between 12 and 20 minutes.

Fitting of DWI signal

Bi-exponential fitting procedures and exact positioning of ROIs were performed using a programmable graphical and calculus environment (Matlab, The Mathworks, Natick, MA, USA) according to the instructions of a radiologist (M.O.) with more than 35 years of experience. For all analyses, the diffusion weighted signal intensities S were fitted bi-exponentially using the parameters prescribed by the IVIM model [4,21]:

\[ \frac{S}{S_0} = f_{fast} \exp(-bD_{fast}) + f_{slow} \exp(-bD_{slow}) \]  

where \( S_0 \) is the maximum signal intensity, \( D_{fast} \) is the fast pseudodiffusion component, \( f_{fast} \) is the fraction of the fast component, \( D_{slow} \) is the slow diffusion component and \( f_{slow} \) is the fraction of the slow component (\( b_{slow} = 1 - f_{fast} \)) as defined previously by Le Bihan et al. [21]. In this study, \( D_{fast} \) is referred to as microperfusion, and \( f_{fast} \) as the fraction of microperfusion in accordance with the study of Lemke et al. who suggested that the IVIM-model separates DWI measurements into a “contribution of microperfusion and diffusion” [22]. \( D_{slow} \) is referred to as the molecular diffusion in accordance with the study of Luciani et al. [23].

Equation 1 was fitted by the Nelder-Mead simplex direct search method with bound constraints, which performs a constrained non-linear minimisation of the sum of the squared residuals [3,24]. The initial guess \( D_{slow} \) was estimated by calculating the slope of the asymptote of the slow signal component between \( b = 500 \) and \( 1000 \) s/mm², and \( D_{slow} \) was bound between 0.5 and \( 5 \times 10^{-3} \) mm²/s. The intercept of the asymptote with the \( y \)-axis at \( S_0 \) resulted in an initial guess \( f_{fast} \) and \( f_{slow} \) was bound between \( f_{fast} - 0.02 \) and \( f_{fast} + 0.02 \). The slope of the signal between \( b = 0 \) and \( b = 100 \) s/mm² was used to guess the initial value of the fast signal component (\( D_{fast} \)), and \( D_{slow} \) was bound between \( D_{slow} \) and \( 100 \times 10^{-3} \) mm²/s.

Hepatic fat fraction

The hepatic fat fraction (HFF) was calculated by Dixon’s in- and out-of-phase SPGR imaging modified with dual flip angles (70°, 20°) as proposed by Hussain et al.: HFF = HFF20 if HFF20 ≤ HFF70 and otherwise HFF = 100 – HFF20 [25,26]. The second SPGR series with a flip angle of 70 degrees were used to calculate HFF70:

\[ HFF70(\%) = \frac{S_{70,IP,corr} - S_{70,OP}}{2 + S_{70,IP,corr}} \times 100 \]  

where

\[ S_{70,IP,corr} = S_{70,IP} \times e^{\frac{T_2^*}{2}} \]  

and \( S_{70,IP} \) and \( S_{70,OP} \) are the signal intensities of the IP and OP images of the second SPGR series using \( \tau = 2.38 \) ms (\( T_{2P} - T_{2OP} \)). Similarly, the calculation of HFF20 was done using the third SPGR series with a flip angle of 20°. \( T_2^* \) was estimated using the first dual echo SPGR series:

\[ \frac{1}{T_2^*} = \frac{1}{ATE} \times \ln \left( \frac{S_{IP,1}}{S_{IP,2}} \right) \]  

where ATE = 13.5 ms (\( T_{2P} - T_{2OP} \)) and \( S_{IP,1} \) and \( S_{IP,2} \) are the respective signal intensities of both echoes.
Image analysis

First the DWI data were loaded. For each of the 11 subjects, four circular regions-of-interest (ROI) with a diameter of 22.3 mm were drawn in each of the eight segmental regions (II – VIII) according to the Couinaud-Bismuth classification [27,28]. The four ROIs were drawn on four different slices when possible; hence when no additional slices were available a second ROI was drawn on the same slice (yet in another location of the segment). For each ROI the average signal intensity S was obtained and the IVIM-DWI parameters ($D_{slow}$, $D_{fast}$, and the respective fractions) were fitted. The exact locations of the ROIs were stored as xy-coordinates, and for each ROI the HFF and T2* were recorded. Finally, the four ROIs measured in each segment were averaged, resulting in 88 measurements totally (11 subjects, 8 segments). During the assessments, any visible vascular and biliary structures nearby were avoided.

Statistical analysis

Statistical analyses were performed using SPSS (SPSS 20, Chicago, IL, USA). All data were tested for normality using Shapiro–Wilk tests. Non-steatotic measurements (HFF ≤ 5.5%) were divided into two groups: group 1 (HFF ≤ 2.75%) and group 2 (2.75 < HFF ≤ 5.5%). Steatotic measurements (HFF > 5.5%) were assigned to group 3. For normally distributed data ($D_{slow}$, $D_{fast}$, $f_{fast}$ and T2*) one-way ANOVA tests were used to compare intergroup differences. Familywise type I error protection when the number of comparisons equals three, while providing increased power compared to Bonferroni correction [29]. For non-normally distributed data (HFF) the differences between the three groups were determined using non-parametric Kruskal-Wallis tests, where post-hoc comparisons after ANOVA were implemented using Fisher’s LSD tests which provides familywise type I error protection when the number of comparisons equals three, while providing increased power compared to Bonferroni correction [29]. For non-normally distributed data (HFF), the differences between the three groups were determined using non-parametric Kruskal-Wallis tests, where post-hoc Mann-Whitney U multiple comparisons were performed with Bonferroni type I error correction.

Guiu et al. reported a potential nonlinear effect between $D_{slow}$ and HFF, especially for HFF below 3% [15]. To investigate whether the relationship between HFF and $D_{slow}$, $D_{fast}$, $f_{fast}$ and T2*, was linear or potentially nonlinear, Pearson’s correlations for all data points were calculated using a linear ($Y = aHFF + b$) and a log-linear ($Y = a\log(HFF) + b$) model. The best-fit model, with the highest correlations for all data points was used to investigate the individual Pearson’s correlations in each of the eight individual liver segments. The significance of the multiple correlation analysis was adjusted by Bonferroni correction to avoid type I errors. Correlations (in absolute values) were classified as weak ($r < 0.36$), moderate ($r = 0.36$ to 0.67) and strong ($r > 0.67$) [30]. For all statistical tests $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Effect of fat on IVIM-DWI parameters

The HFF were non-normally distributed ($p < 0.001$) and ranged between 1.5 and 29.9% for the eleven subjects. Five subjects yielded non-steatotic HFF (≤5.5%) measurements only, two subjects had steatotic HFF (>5.5%) measurements only, and four subjects had both steatotic and non-steatotic measurements (Fig. 1).

IVIM-DWI parameters were normally distributed ($p \geq 0.319$). Molecular diffusion ($D_{slow}$) differed significantly between the three groups of different HFF, and also the intergroup comparisons were significantly different ($p < 0.001$, Table 1). $D_{slow}$ was 0.85 x 10^-3 mm^2/s for the first group (HFF ≤ 2.75%) and decreased steadily to 0.72 x 10^-3 mm^2/s in group 2 (2.75 < HFF ≤ 5.5%) and 0.49 x 10^-3 mm^2/s in group 3 (HFF > 5.5%). $D_{fast}$ and $f_{fast}$ did not show differences between the three HFF groups ($p \geq 0.194$). Pearson’s correlation analysis (Table 2) showed a significant negative linear relationship with moderate correlation between HFF and $D_{slow}$ using both the linear model ($r = -0.446$, $p < 0.001$) and the log-linear model ($r = -0.514$, $p < 0.001$, Fig. 2). The log-linear model showed overall higher correlations compared to the linear model and was therefore used for the individual segment analysis. The average HFF varied from 3.00±9.17% to 7.32±9.36% between the individual segments (Table 3). In segment VII a significant negative linear relationship with strong correlation ($r = -0.840$; $p \leq 0.008$) was observed between HFF and $D_{slow}$ using the log-linear model (Table 4). No significant correlations between HFF and $D_{fast}$ or $f_{fast}$ were observed.

Effect of fat on T2*

T2* was normally distributed ($p = 0.116$) and differed significantly between the three groups of different HFF, and also the intergroup comparisons were significantly different ($p < 0.001$, Table 1). T2* was 32.2 ms for the first group (HFF ≤ 2.75%) and decreased steadily to 27.2 ms in group 2 (2.75 < HFF ≤ 5.5%) and 21.0 in group 3 (HFF > 5.5%). Pearson’s correlation analysis (Table 2) showed a significant negative linear relationship with moderate to strong correlation between HFF and T2* using both the linear model ($r = -0.607$, $p < 0.001$) and the log-linear model ($r = -0.714$, $p < 0.001$, Fig. 3). The log-linear model showed overall higher correlations compared to the linear model and was therefore used for the individual segment analysis. In 5 of 8 segments a significant negative linear relationship with strong
Correlation (r = 0.767 to 0.804; p < 0.048) was observed between HFF and T2* using the log-linear model (Table 4).

**Discussion**

**Effect of fat on IVIM-DWI parameters**

In this study it was demonstrated that molecular diffusion (Dslow) in the liver is affected by hepatic fat, also in the absence of steatosis (HFF below 5.5%). Dslow differed significantly between three groups of different HFF, and a steady significant decrease of Dslow with moderate correlation was found for increasing HFF. These results complement existing knowledge of the reduction of molecular diffusion by steatotic HFF (>5.5%). Previous IVIM studies showed comparable negative correlations (r = 0.59 and r = 0.18) between HFF and molecular diffusion using a linear model.

Figure 2. Regression plot between HFF and molecular diffusion. The correlation between HFF (%) and molecular diffusion (×10⁻³ mm²/s) was assessed by using a log-linear model. Pearson’s product-moment correlation and its significance were calculated. The log-linear regression line is displayed together with its 95% confidence interval.

doii:10.1371/journal.pone.0087926.g002

correlation (r = -0.767 to -0.804; p < 0.048) was observed between HFF and T₂* using the log-linear model (Table 4).

**Table 2. Correlations of IVIM-DWI parameters and T₂* with hepatic fat fraction (HFF) using two models.**

<table>
<thead>
<tr>
<th></th>
<th>Linear model</th>
<th>Log-linear model</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Pearson’s r</td>
<td>F</td>
</tr>
<tr>
<td>Dslow</td>
<td>-0.446*</td>
<td>21.317</td>
</tr>
<tr>
<td>Dfast</td>
<td>-0.040</td>
<td>0.138</td>
</tr>
<tr>
<td>ffast</td>
<td>0.041</td>
<td>0.148</td>
</tr>
<tr>
<td>T₂*</td>
<td>-0.607*</td>
<td>50.277</td>
</tr>
</tbody>
</table>

Two models were used to assess the effect of hepatic fat fraction (HFF) on the IVIM-DWI parameters (Dslow, Dfast, and ffast) or T₂* measured in 11 patients and 8 segments (n = 88). The linear model assumed a linear relationship between HFF and the IVIM-DWI parameters or T₂* (Y = aHFF + b). The log-linear model assumed a linear relationship between the logarithmic of HFF and the IVIM-DWI parameters or T₂* (Y = a+log(HFF) + b). Pearson’s correlations increased when the log-linear model was applied. *Indicates significant correlations.

doii:10.1371/journal.pone.0087926.t002
model [15,31]. In addition, Guin et al. noticed a potential nonlinear effect between $D_{\text{slow}}$ and HFF, especially for HFF below 3% [15]. This was confirmed in our study: the relationship ($r = \pm 0.39$) [13]. In an animal study similar results were found, increasing degrees of hepatic steatosis correlated fairly well ($r = \pm 0.56$) with decreasing liver ADCs [12].

With regard to the effect of fat on IVIM-DWI parameters in the different segmental regions, we used the log-linear model which had the highest correlations in the overall analysis compared to the linear model. Others compared non-steatotic with steatotic livers and nonlinear effect between $D_{\text{slow}}$ and HFF, especially for HFF below 5%, with moderate correlations [15]. This was confirmed in our study: the relationship between HFF (%) and $T_2^*$ (ms) was assessed by using a log-linear model. Pearson’s product-moment correlation and its significance were calculated. The log-linear regression line is displayed together with its 95% confidence interval.

The decrease of molecular diffusion by fat can be due to several mechanisms. MR relaxation is determined predominantly by water-macromolecular interactions [33]. The MR signal from protons bound to macromolecules such as fat will interfere with the MR signal from freely diffusing water molecules. This can partly explain the observed decreased molecular diffusion in the presence of fat. To prevent interference, fat suppression techniques serve to suppress the signal originating from protons bound to fat in order to reduce the chemical shift artifacts and eliminate signals arising from adipose tissue [34]. However, fat suppression techniques are not perfect and come with disadvantages and pitfalls such as the dependency on the homogeneity of the main static magnetic field. Hence there will always be some interference of MR signal between free protons and macromolecular protons that cannot be neglected. However, as indicated previously by the breast DWI study of Baron et al., low molecular diffusion in the presence of fat may reflect either direct contributions from the protons of the relatively immobile fat molecules or low water content, thereby restricting the diffusion of water (trapped water) [35]. Similarly, we hypothesize that the reduction of molecular diffusion in the liver is caused by physical hindrance of the movement of water molecules by the presence of macrovesicular fat droplets in hepatocytes. The fat present in the liver, is stored as triglycerides in sphere shaped vacuoles, which usually appear as large droplets with diameters larger than 15 μm [36]. These vacuoles reside in the hepatocytes, which are polygonal cells with six or more faces and a mean diameter ranging between 20 and 40 μm [37]. A considerable fraction of the volume of the hepatocyte can therefore be occupied by the macrovesicular fat droplet (Fig. 4). Considering a diffusion length [38] of about 17 μm ($l_D = \sqrt{6}D_{t}t$, $D_t = 1.0 \times 10^{-5}$ mm$^2$/s, $t = 50$ ms), which is in the same order of magnitude of the hepatocyte’s diameter, we suspect that the movement of water molecules can be physically hindered by the presence of macrovesicular fat droplets in hepatocytes. This would be a mechanical process rather than signal interference between protons bound to fat molecules and free water protons.

**Table 3.** Hepatic fat fraction (HFF), IVIM-DWI parameters and $T_2^*$ per segment.

<table>
<thead>
<tr>
<th>Seg</th>
<th>HFF (%)</th>
<th>$D_{\text{slow}}$ (10$^{-3}$ mm$^2$/s)</th>
<th>$D_{\text{fast}}$ (10$^{-3}$ mm$^2$/s)</th>
<th>$f_{\text{fast}}$ (%)</th>
<th>$T_2^*$ (ms)</th>
</tr>
</thead>
<tbody>
<tr>
<td>II</td>
<td>4.99 ± 1.80</td>
<td>0.48 ± 0.28</td>
<td>48.4 ± 8.5</td>
<td>50 ± 5.1</td>
<td>25.5 ± 5.8</td>
</tr>
<tr>
<td>III</td>
<td>7.32 ± 9.36</td>
<td>0.65 ± 0.16</td>
<td>40.9 ± 8.8</td>
<td>44 ± 6.7</td>
<td>28.1 ± 5.9</td>
</tr>
<tr>
<td>IVa</td>
<td>3.29 ± 8.64</td>
<td>0.85 ± 0.28</td>
<td>40.0 ± 7.0</td>
<td>38 ± 5.4</td>
<td>26.8 ± 8.4</td>
</tr>
<tr>
<td>IVb</td>
<td>3.00 ± 9.17</td>
<td>0.85 ± 0.19</td>
<td>48.4 ± 8.6</td>
<td>39 ± 6.1</td>
<td>29.0 ± 7.1</td>
</tr>
<tr>
<td>V</td>
<td>3.46 ± 7.38</td>
<td>0.60 ± 0.25</td>
<td>38.0 ± 8.1</td>
<td>33 ± 5.9</td>
<td>27.6 ± 4.8</td>
</tr>
<tr>
<td>VI</td>
<td>3.41 ± 8.73</td>
<td>0.74 ± 0.21</td>
<td>37.5 ± 7.2</td>
<td>32 ± 5.3</td>
<td>27.7 ± 5.8</td>
</tr>
<tr>
<td>VII</td>
<td>3.98 ± 9.18</td>
<td>0.75 ± 0.23</td>
<td>34.3 ± 6.6</td>
<td>31 ± 5.6</td>
<td>26.2 ± 6.0</td>
</tr>
<tr>
<td>VIII</td>
<td>3.27 ± 8.54</td>
<td>0.59 ± 0.26</td>
<td>31.4 ± 5.8</td>
<td>29 ± 5.3</td>
<td>25.6 ± 5.6</td>
</tr>
</tbody>
</table>

For each subject four ROIs were drawn in each segment and then averaged, resulting in 11 measurements per segment. * Data are medians ± median. ‡ Data are means ± standard deviations.

doi:10.1371/journal.pone.0087926.t003
The microperfusion parameters (D\text{fast} and f\text{fast}) were overall not affected by the HFF, neither in the individual segments. The average fraction of microperfusion f\text{fast} was comparable to previously published numbers (29–35%) on healthy livers [15,23]. In agreement with earlier findings, f\text{fast} was highest (≥ 44%) in the left lobe (segments II and III) [39]. Microperfusion (D\text{fast}) has been found to be lower in patients with steatosis compared to patients without steatosis [15]. In contrast, in this study we did not find a relation between D\text{fast} and HFF. This can be partially due to the limited accuracy of D\text{fast} in this study. Because of software limitations on the MR system, it was not possible to acquire any data of D\text{fast} between b = 0 and 50 s/mm². The lack of b-values below 50 s/mm² can also explain the relatively low standard deviation of D\text{fast} in our study. Previously published numbers. In a large population-based project conducted in northeast Germany, HFF ranged between 4.6% and 34.9% for the majority of a group of 88 healthy volunteers [40]. Also in a comparative methodological study, HFF of healthy volunteers ranged up to 21.1% showing high correlations between MR spectroscopy and two-point Dixon-based MRI fat quantification [41].

**Effect of fat on T2**

In this study, T2\text{*} differed significantly between three groups of different HFF, also for non-steatotic HFF below 5.5%, and a steady significant decrease of T2\text{*} with strong correlation was found for increasing HFF. Also in the different segmental regions, we found that T2\text{*} correlated significantly with HFF in 5 of the 8 segments. Correlations were higher for the log-linear model compared to the linear model, suggesting a nonlinear relationship between HFF and T2\text{*} as well. Hepatic T2\text{*} variations among different segments have been shown to be low in healthy subjects, ranging between 19.5 and 29.9 ms [42], which is in accordance with our observations. Similar results were obtained in an animal study where the T2\text{*} of liver parenchyma of rats decreased from 31.4 ms for the control group (0.9% HFF) to 19.1 ms for rats fed by a four week choline-deficient diet (26.0% HFF) [43]. A clinical study on patients with non-alcoholic fatty liver disease (NAFLD) reported significant decreases of T2 relaxation times of water with increasing fat fractions [19]. In addition, T2\text{*} shorting by fat has been confirmed using various phantoms with different fat-water mixtures [44].

However, for a number of studies no correlation was found between T2\text{*} relaxation and HFF [20,45]. For example, Hernandez and Kuhli et al. demonstrated that T2\text{*} estimations are inaccurate in tissues with high fat content due to the complex fat spectrum, and concluded that these issues can be solved when multipoint spectral modeling of fat is applied: this way they showed that T2\text{*} is independent of the fat fraction [18,40]. In our study, T2\text{*} was not corrected for the spectral complexity of the fat signal, which can explain the dependency of T2\text{*} on the HFF in our study.
The dependency of $T_2^*$ on HFF can also be understood from the perspective of Bottomley et al. who suggested a fast exchange two-state (FETS) model to describe proton $T_1$ and $T_2$ relaxation in normal tissue [46]. They identified three chemically different proton species: macromolecular protons (excluding fatty acids), free water protons, and mobile fatty acid protons, relaxing with $T_2$ times of $\sim 10-100 \mu s$, $\sim 50 \text{ ms}$ and $\sim 0.2 \text{ s}$ respectively. If the amount of fat in the liver changes, the interference pattern of the different proton signals, causing dephasing, also changes, and the overall effect can be shortening of the $T_2^*$ relaxation time. This is in concordance with Yu et al. who suggested that when fat coexists with water in a voxel, $T_2^*$ relaxometry may be disturbed by the chemical shift of fat, due to constructive and destructive interference of fat and water signals [47].

Clinical implications

It is known that the decreases of the molecular diffusion between normal liver tissue and cirrhosis or the different stages of fibrosis are relatively small and technically challenging to detect [23,48,49]. It is therefore important to know what methodological factors can reduce molecular diffusion, regardless of the pathology itself. One of these factors is that diffusion measurements can be heavily dependent on the MR-equipment used, which requires use of the same scanner to ensure comparable measurements [50].

Also, user-dependent factors such as the choice of measurement location within the liver may affect the diffusion measurements. This was demonstrated in a recent study where the apparent diffusion coefficient significantly depended on the segmental region in the liver [39]. In the current study, we added another factor: molecular diffusion is negatively related to the hepatic fat fraction, also at non-steatotic fat levels. This is especially important when in pursuit for quantitative cut-off values for molecular diffusion in order to discriminate healthy liver tissue from pathology. Molecular diffusion is dependent on the hepatic fat fraction, also below 5.5%. This implicates that any derived cut-off value of the molecular diffusion for cirrhosis, or stages of fibrosis, is dependent on the hepatic fat fraction as well, especially because it is known that hepatic fat fractions vary between subjects [51]. Therefore, we recommend that to correctly interpret quantitative hepatic DWI, acquisition of the hepatic fat fraction prior to the hepatic DWI protocol is necessary. In that way, diffusion measurements can be judged along with the fat measurement, which ensures a more reliable assessment of the diffusion properties of pathology.

Similar issues apply to quantitative evaluation of hemochromatosis using $T_2^*$ [ms] estimation. The histopathologic iron grade can be classified using $T_2^*$ measurement of the liver [52]. For that purpose cut-off values have been defined. However, if $T_2^*$ values are dependent on the hepatic fat fraction, then cut-off values used in the classification of hemochromatosis ought to be corrected for the hepatic fat fraction.

Next to quantitative difficulties, also non-quantitative issues may arise due to the effect of fat on hepatic DWI. Hypointensity in DWI can be a side effect of $T_2^*$ shortening. DW images are $T_2$ weighted and changes in $T_2$ relaxation times can therefore affect the signal intensities independently of the tissue diffusion in three ways [53]: shine-through (prolongation of $T_2$), washout (balance between $T_2$ prolongation and increased diffusion) and blackout or hypointensity (shortening of $T_2$). As fat shortens $T_2^*$, locally elevated levels of fat in the liver can reveal as hypointense areas, which might be incorrectly interpreted as increased liver diffusion.

In conclusion, we have demonstrated that hepatic fat fractions significantly decrease the molecular diffusion in the liver, also for non-steatotic fat levels ($\leq 5.5\%$). In addition, it was confirmed that hepatic fat fractions significantly decrease $T_2^*$ measurements when multiprobe spectral modeling is not applied. It is known that the decreases of the molecular diffusion between normal liver tissue and cirrhosis or the different stages of fibrosis are relatively small and technically challenging to be detected [23,48,49]. The knowledge of the effect of low levels of hepatic fat on the molecular diffusion is therefore expected to be of importance in the diagnosis and staging of fibrosis and cirrhosis using quantitative DWI. Therefore, we conclude that to correctly interpret quantitative hepatic DWI, acquisition of the hepatic fat fraction prior to the hepatic DWI protocol is necessary to ensure a reliable assessment of the diffusion properties of pathology.

Acknowledgments

We thank Evelin Hans, Heike Held and Inge Vuijt for their support with the recruitment and data collection.

Author Contributions

Conceived and designed the experiments: HD MO PES. Performed the experiments: HD PK. Analyzed the data: HD AH MO PES. Contributed reagents/materials/analysis tools: HD. Wrote the paper: HD AH PK MO PES.

References

15. Szczepaniak JS, Nurenberg P, Leonard D, Browning JD, Reingold JS, et al. (2005) Magnetic resonance spectroscopy to measure hepatic triglyceride content:


