

Towards an optimal distribution of b-values for IVIM imaging

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Introduction:

The intra voxel incoherent motion (IVIM) theory is a sophisticated method to separate diffusion and perfusion effects using diffusion weighted imaging [1], which is particularly useful for the investigation of abdominal lesions [2,3]. However the quality of the parametric maps, especially of the perfusion related maps, is usually limited due to large uncertainties of the fitted parameters [4], in particular when clinical feasible scan times are used. The choice of the number and distribution of b-values varies greatly [2,4] and a fundamental basis for this choice is currently not available. Therefore, the aim of this work was to find an optimal distribution of b-values. To this end we used Monte Carlo simulations where the reduction of the error of the fit was used as criterion to find this distribution.

Materials and Methods:

The signal curve was generated according to the IVIM theory (equation 1) using three different parameter sets of perfusion fraction f , diffusion coefficient D and pseudo diffusion coefficient D^* . The values used for the three parameter sets were taken from the literature for brain, liver and kidney (brain: $f=5\%$, $D=1\ \mu\text{m}^2/\text{ms}$, $D^*=10\ \mu\text{m}^2/\text{ms}$ [5]; liver: $f=30\%$, $D=1\ \mu\text{m}^2/\text{ms}$, $D^*=60\ \mu\text{m}^2/\text{ms}$ [2]; kidney: $f=30\%$, $D=1.5\ \mu\text{m}^2/\text{ms}$, $D^*=15\ \mu\text{m}^2/\text{ms}$ [6]). For every b-value, Gaussian noise was added to the complex signal to simulate a Rician distribution of the magnitude signal. Equation 1 was then fitted to the noisy signal curve using the Levenberg-Marquardt algorithm. Initial values were $f=10\%$, $D=0.0001\ \text{s}/\text{mm}^2$ and $D^*=0.001\ \text{s}/\text{mm}^2$. No limiting value was implemented. The complete process was repeated 5000 times for each set of IVIM parameters and b-values and the relative overall error σ (equation 2) were calculated.

Starting from an initial b-value distribution $\{b=0, 40, 1200\ \text{s}/\text{mm}^2\}$, the σ of all distributions with the initial three b-values plus one additional b-value in the range from 0 to 1400 s/mm^2 (in steps of 10 s/mm^2) were tested. The distribution with the minimal σ was then chosen as optimal new b-value distribution. This process was consecutively iterated to obtain distributions with up to 100 b-values. The optimization was run four times: three times for the individual organs and once with the summed σ of all three organs yielding the distribution $\{b_{\text{sum}}\}$. All simulations were implemented in C++.

Results:

Fit quality per unit time defined as $1/(\sigma\sqrt{n})$ is shown in figure 1 of the optimized b-value distributions as a function of the number of b-values n for the three parameter sets at an SNR of 80. Figure 1a shows that the fit quality for the liver increases until 10 b-values are used $\{b=0,40,1200,10,100,170,20,1260,20,0\}$, in order of importance. After this point, additional time can be invested either in repetition of this distribution or in additional b-values without strong differences in the fit quality per unit time. For the kidney, this point is reached even faster, at about eight b-values $\{b=0,40,1200,240,1030,40,240,1070\}$. The fit quality for the brain parameters shows a special feature (figure 1b): using few b-values, the fit quality per unit time decreases until a critical point is reached at about $n=40$. In this n range, the optimization algorithm finds that averaging the three initial b-values is beneficial compared to the addition of novel b-values. Beyond $n>40$ addition of novel b-values improves the fit quality considerably. Figure 2 shows the distribution of b_{sum} ($n=100$). In order to compare this optimized b-value distribution with a previously reported distribution of 16 b-values $\{b_{\text{lit}}=0,10,20,30,40,50,60,70,80,90,100,150,200,400,800,1000\}$ [2] and a distribution that is averaged four times $\{b_{\text{av}}=4*(0,40,100,1200)\}$, we applied the first 16 b-values of b_{sum} $\{0,40,1200,200,20,360,1380,40,280,1300,350,0,70,1020,310,80\}$ and analyzed the relative overall error σ for each distribution (see table 1). The error of b_{sum} is 82 % lower for the brain parameters, 24 % lower for the liver parameters and 24 % lower for the kidney parameters compared to b_{lit} and 74 %, 1% and 9 % lower compared to b_{av} .

Discussion:

This work shows that an optimal choice of b-values can substantially minimize the overall measurement error and thus is of critical importance. The presented distribution b_{sum} can be seen as a first step towards an optimal b-value set up for IVIM imaging. Further evaluation is warranted to solidify these results. Furthermore, our results confirm findings of Pekar [5], which indicated that an extremely high SNR is needed for IVIM of the brain. On the other hand, IVIM of the abdomen can be performed at moderate SNR with a low overall measurement error.

References:

- [1] Le Bihan D, Radiology 1988;168(2):497-505. [2] Luciani A, Radiology. 2008 Dec;249(3):748-52. [3] Thoeny HC., Radiology 2009;252:721-728.
 [4] Lee SS, J Magn Reson Imaging 2008;28(4):928-936 [5] Pekar J., Magn Reson Med 1992;23:122-129 [6] Zhang JL, ISMRM 2009 Honolulu; 4111

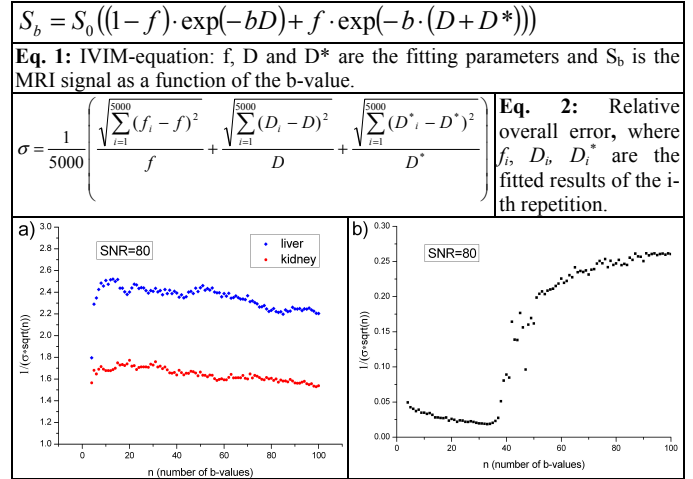


Fig. 1: Fit quality per unit time of the optimized b-value distributions as a function of the number of b-values at SNR=80 for the liver and kidney parameters (a) and the brain parameters (b)

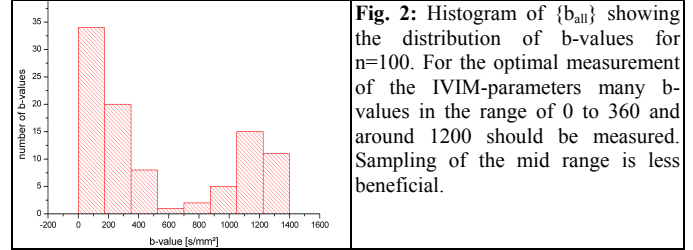


Fig. 2: Histogram of $\{b_{\text{all}}\}$ showing the distribution of b-values for $n=100$. For the optimal measurement of the IVIM-parameters many b-values in the range of 0 to 360 and around 1200 should be measured. Sampling of the mid range is less beneficial.

	b_{sum}	b_{lit}	b_{av}
brain	$\sigma=16.3\%$	$\sigma=92.7\%$	$\sigma=63.6\%$
liver	$\sigma=0.81\%$	$\sigma=1.07\%$	$\sigma=0.82\%$
kidney	$\sigma=1.20\%$	$\sigma=1.58\%$	$\sigma=1.32\%$

Tab.1: Comparison of the relative overall error of the optimized b-value distribution b_{all} to a previously reported distribution b_{lit} and an averaged distribution b_{av} at SNR=200. The error decreases significantly using the optimized b-value distribution b_{sum} compared to b_{lit} for all parameter sets.