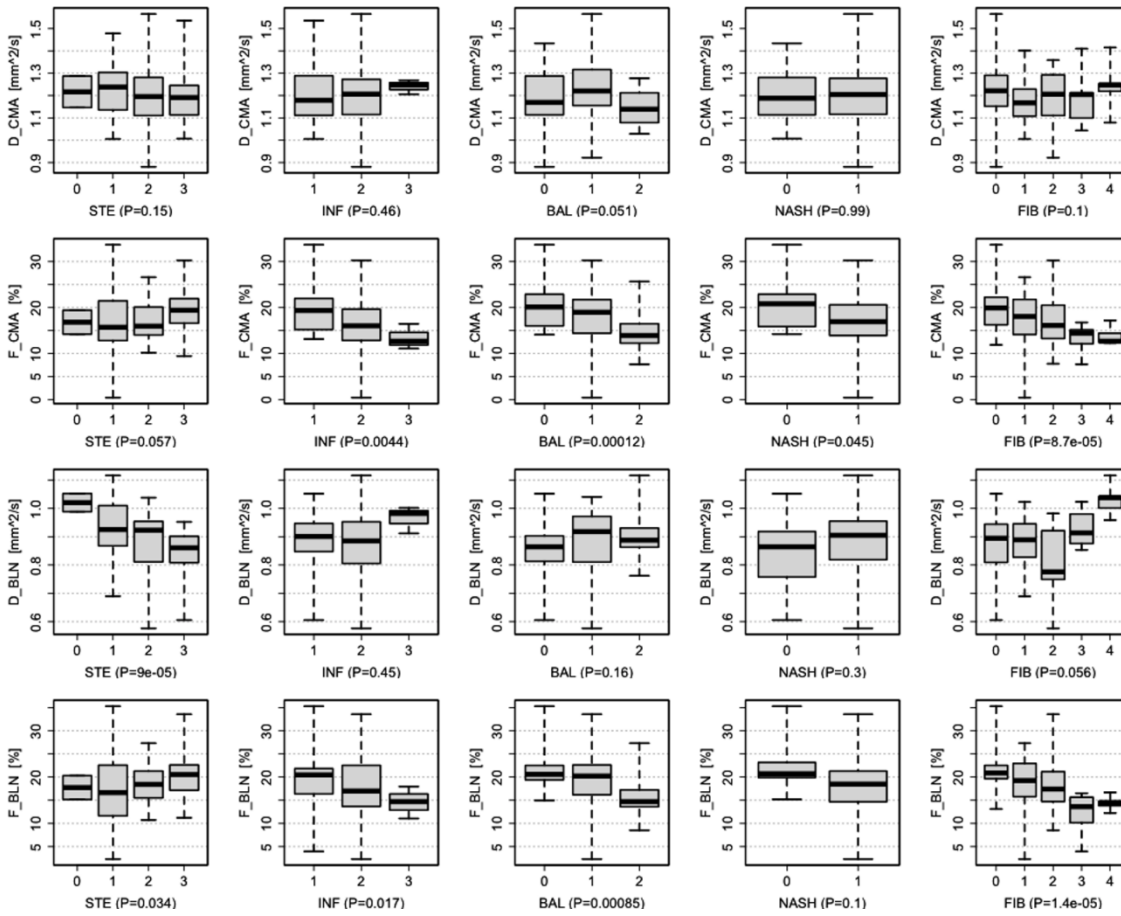


**PDFF** 15 (1-36) %

<b>DWI Parameter</b>	<b>CMA</b>	<b>P</b>	<b>BLN</b>	<b>P</b>
<i>D</i> [ $10^{-3}mm^2/s$ ]	1.2 (0.88-1.6)	<0.001	0.89 (0.58-1.1)	<0.001
<i>F</i> [%]	18 (5 - 34)	<0.001	19 (2.3-35)	<0.001

**Table S1 – Distribution of PDFF and of IVIM parameters after fat fraction adjustment.** Mean and range of each parameter is given. P-value corresponds to paired student's t-test between fat fraction-adjusted parameter versus unadjusted parameter.



**Figure S1 – Trend tests after fat fraction adjustment.** Trends in IVIM parameters (D middle, F bottom) after fat fraction adjustment for each histologic feature (STE: steatosis, INF: inflammation, BAL: ballooning, NASH score, and FIB: fibrosis) are shown above. The barplot indicates the minimum, 1<sup>st</sup>, 2<sup>nd</sup>, 3<sup>rd</sup> quartiles, and maximum value of each DWI parameter for each observed score of histologic feature. Each plot is annotated with the p-value of the Jonckheere-Terpstra test for trends or the p-value of the Wilcoxon signed rank test for NASH. Parameters derived from the LS (top) and BLN (bottom) reconstruction methods are shown.

Regression		FF Adjusted CMA Reconstruction DWI -Derived Parameter	
Model	Coefficient	D	F
1	Intercept	1.285 ± 0.064	21.209 ± 2.397
	STE	-0.015 ± 0.017, P=0.393	<b>1.476 ± 0.659, P=0.028 *</b>
	INF	0.015 ± 0.030, P=0.615	-1.342 ± 1.138, P=0.241
	BAL	-0.023 ± 0.030, P=0.442	-2.245 ± 1.145, P=0.053
	FIB	-0.005 ± 0.016, P=0.750	-0.562 ± 0.590, P=0.344
2	Intercept	1.233 ± 0.088	22.818 ± 3.386
	STE	-0.020 ± 0.017, P=0.244	1.150 ± 0.649, P=0.080
	NASH	0.029 ± 0.048, P=0.541	-3.094 ± 1.833, P=0.095
	FIB	-0.012 ± 0.012, P=0.336	<b>-1.327 ± 0.472, P=0.006 **</b>

Regression		FF Adjusted BLN Reconstruction DWI -Derived Parameter	
Model	Coefficient	D	F
1	Intercept	0.952 ± 0.046	21.026 ± 2.604
	STE	<b>-0.046 ± 0.013, P=0.0004 ***</b>	1.406 ± 0.715, P=0.053 .
	INF	-0.006 ± 0.022, P=0.770	-0.659 ± 1.236, P=0.595
	BAL	0.030 ± 0.022, P=0.178	-1.252 ± 1.244, P=0.317
	FIB	0.009 ± 0.011, P=0.410	<b>-1.410 ± 0.641, P=0.031 *</b>
2	Intercept	0.894 ± 0.063	22.938 ± 3.599
	STE	<b>-0.047 ± 0.012, P=0.0001 ***</b>	1.305 ± 0.690, P=0.062 .
	NASH	0.057 ± 0.034, P=0.101	-2.407 ± 1.948, P=0.220
	FIB	0.012 ± 0.009, P=0.160	<b>-1.753 ± 0.502, P= 0.0007 ***</b>

**Table S2 – Multiple Linear Regression Tables after Fat Fraction Adjustment.** The coefficient ± standard error and significance of the effects of each histologic feature of NAFLD are reported above for both reconstruction methods and both model after fat fraction adjustment. The p-value range is specified by the number of adjacent asterisks, as follows: \*\*\* = 0-0.001, \*\* = 0.001-0.010, \* = 0.010-0.050.