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# Linearity, Bias, and Precision of Hepatic Proton Density Fat Fraction Measurements by Using MR Imaging: A Meta-Analysis<sup>1</sup>

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## Purpose:

To determine the linearity, bias, and precision of hepatic proton density fat fraction (PDFFF) measurements by using magnetic resonance (MR) imaging across different field strengths, imager manufacturers, and reconstruction methods.

## Materials and Methods:

This meta-analysis was performed in accordance with Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines. A systematic literature search identified studies that evaluated the linearity and/or bias of hepatic PDFFF measurements by using MR imaging (hereafter, MR imaging-PDFFF) against PDFFF measurements by using colocalized MR spectroscopy (hereafter, MR spectroscopy-PDFFF) or the precision of MR imaging-PDFFF. The quality of each study was evaluated by using the Quality Assessment of Studies of Diagnostic Accuracy 2 tool. De-identified original data sets from the selected studies were pooled. Linearity was evaluated by using linear regression between MR imaging-PDFFF and MR spectroscopy-PDFFF measurements. Bias, defined as the mean difference between MR imaging-PDFFF and MR spectroscopy-PDFFF measurements, was evaluated by using Bland-Altman analysis. Precision, defined as the agreement between repeated MR imaging-PDFFF measurements, was evaluated by using a linear mixed-effects model, with field strength, imager manufacturer, reconstruction method, and region of interest as random effects.

## Results:

Twenty-three studies (1679 participants) were selected for linearity and bias analyses and 11 studies (425 participants) were selected for precision analyses. MR imaging-PDFFF was linear with MR spectroscopy-PDFFF ( $R^2 = 0.96$ ). Regression slope (0.97;  $P < .001$ ) and mean Bland-Altman bias ( $-0.13\%$ ; 95% limits of agreement:  $-3.95\%$ ,  $3.40\%$ ) indicated minimal underestimation by using MR imaging-PDFFF. MR imaging-PDFFF was precise at the region-of-interest level, with repeatability and reproducibility coefficients of 2.99% and 4.12%, respectively. Field strength, imager manufacturer, and reconstruction method each had minimal effects on reproducibility.

## Conclusion:

MR imaging-PDFFF has excellent linearity, bias, and precision across different field strengths, imager manufacturers, and reconstruction methods.

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**H**epatic steatosis, or intracellular accumulation of triglycerides in hepatocytes, is a common histologic manifestation of many liver diseases. In particular, obesity-related steatosis, or nonalcoholic fatty liver disease, has become one of the leading causes of liver disease worldwide (1) paralleling the obesity pandemic. Nonalcoholic fatty liver disease can progress to cirrhosis and liver cancer, and it is the most rapidly growing indication for liver transplantation in the United States (2). Moreover, even in those patients with nonprogressive disease, nonalcoholic fatty liver disease is a risk factor for future development of diabetes, cardiovascular death, and other cancers (3–9). Control of nonalcoholic fatty liver disease and its complications has thus become a major public health priority.

Percutaneous liver biopsy has been the clinical reference standard for diagnosis and grading of hepatic steatosis (10). However, biopsy is costly, painful, and invasive with rare but serious complication risks including hemorrhage

and death (11). In addition, inherently small tissue volume of biopsy (approximately 1/50000th of the entire liver) is a concern for sampling variability and misclassification of disease severity (12,13). Because of the limitations of biopsy, especially for longitudinal disease monitoring (14,15), interest in noninvasive methods to diagnose and grade hepatic steatosis has increased.

Recently, proton density fat fraction (PDFFF) has emerged as the leading noninvasive quantitative imaging biomarker (QIB) of hepatic steatosis (16,17). PDFFF is a fundamental tissue property and an objective magnetic resonance (MR) imaging–based measure of tissue triglyceride concentration, calculated as the ratio of MR imaging–visible triglyceride protons to the sum of triglyceride and water protons. Spatially localized MR spectroscopy has been the accepted noninvasive reference standard for quantifying hepatic steatosis, used in epidemiologic studies and randomized controlled clinical trials to derive the highest level of evidence (18–34). Acquired and analyzed by using a standardized approach, MR spectroscopy can measure proton densities of triglyceride and water in a small volume of liver tissue *in vivo*, from which PDFFF is calculated. However, PDFFF measurements by using MR spectroscopy can be technically challenging for several reasons, including potential biases because of the selection of sampling volume in livers with nonuniform distribution of fat, difficulty in colocalizing measurement volumes across longitudinal time points, and a need for offline spectral analysis and data quality

assessment. To address these technical challenges, advanced chemical shift–encoded MR imaging methods have been developed to automatically “map” hepatic PDFFF values pixel by pixel throughout the entire liver. These specialized imaging methods are now commercially available with many 1.5-T and 3.0-T MR imaging systems, and the opportunity for widespread use of hepatic PDFFF measurements as a QIB has become a reality.

According to the Radiological Society of North America Quantitative Imaging Biomarkers Alliance (QIBA), three key technical performance metrics of QIBs are linearity, bias, and precision (35). Linearity and bias together assess the degree to which a QIB (eg, PDFFF measurements by using MR imaging [hereafter, MR imaging–PDFFF]) provides an estimate of the true value (eg, a phantom with known triglyceride concentration) or of an accepted *in vivo* reference value (eg, PDFFF measurements by using MR spectroscopy [hereafter, MR spectroscopy–PDFFF]) over the entire range of expected values (eg,

### Advances in Knowledge

- Hepatic proton density fat fraction (PDFFF) measured by using MR imaging (hereafter, MR imaging–PDFFF) has excellent pooled linearity against the reference values measured by using MR spectroscopy (hereafter, MR spectroscopy–PDFFF), with a linear model coefficient of determination of 0.96.
- The pooled linear regression analysis of data from MR imaging–PDFFF and MR spectroscopy–PDFFF demonstrated intercept of  $-0.07\%$  and slope of 0.97, indicating minimal underestimation by using MR imaging–PDFFF.
- The pooled bias, or the mean difference between MR imaging–PDFFF and MR spectroscopy–PDFFF, was  $-0.13\%$ .
- Hepatic MR imaging–PDFFF is highly precise at the region-of-interest level, with repeatability and reproducibility coefficients of 2.99% and 4.12%, respectively.

### Implications for Patient Care

- As measured in published studies, hepatic MR imaging–PDFFF has excellent technical performance as a quantitative imaging biomarker for widespread use in clinical trials and patient care.
- Hepatic MR imaging–PDFFF can be measured with excellent linearity, negligible bias, and high precision by using different field strengths, imager manufacturers, and reconstruction methods.

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#### Abbreviations:

CI = confidence interval  
 PDFFF = proton density fat fraction  
 QIB = quantitative imaging biomarker  
 QIBA = Quantitative Imaging Biomarkers Alliance  
 ROI = region of interest

#### Author contributions:

Guarantors of integrity of entire study, T.Y., S.D.S., M.S.M., S.B.R.; study concepts/study design or data acquisition or data analysis/interpretation, all authors; manuscript drafting or manuscript revision for important intellectual content, all authors; approval of final version of submitted manuscript, all authors; agrees to ensure any questions related to the work are appropriately resolved, all authors; literature research, T.Y., S.D.S., A.P., M.R.B., G.H., D.H., H.H.H., H.H., J.P.K., G.M.K., R.L., M.S.M., X.W., S.B.R., C.B.S.; clinical studies, T.Y., S.D.S., H.H., J.P.K., G.M.K., A.T., X.W., S.B.R.; experimental studies, S.D.S., G.M.K., R.L., X.W.; statistical analysis, T.Y., S.D.S., M.R.B., J.P.K., N.A.O., S.B.R.; and manuscript editing, T.Y., S.D.S., A.P., M.R.B., G.H., D.H., H.H.H., H.H., J.P.K., G.M.K., R.L., M.S.M., N.A.O., J.S.S., A.T., S.B.R., C.B.S.

Conflicts of interest are listed at the end of this article.

See also the editorial by Jackson in this issue.

PDFF measurements of approximately 0%–55%, or the range from normal lean liver to the most severe steatosis observed in patients). Precision assesses the agreement between repeated measurements of a QIB (eg, MR imaging–PDFF) and can be reported in two different ways: repeatability, or the agreement between repeated QIB measurements under identical or near-identical conditions (eg, scan-rescan repeatability of MR imaging–PDFF after interscan recalibration), and reproducibility, or the agreement between repeated measurements under different conditions (eg, MR imaging–PDFF by using equipment with different field strengths, imager manufacturers, and/or reconstruction methods).

Multiple previous studies, almost all single center, have shown MR imaging–PDFF to have high agreement with MR spectroscopy–PDFF in terms of the in vivo reference value (36–58), scan-rescan repeatability (53,56,59–61), cross-imager reproducibility (55,62,63), and cross-field strength reproducibility (48,55,62,63). Despite the excellent performance reported in these studies, there are limited comprehensive data available on the performance of MR imaging–PDFF in multicenter research or clinical settings in which participants may undergo MR imaging–PDFF by using equipment that varies in field strength, imager manufacturer, and/or reconstruction method. An understanding of the technical performance of MR imaging–PDFF in such settings is needed to inform, qualify, and support its use as a QIB for clinical trials and patient care.

Therefore, the purpose of this meta-analysis was to determine the linearity, bias, and precision of hepatic MR imaging–PDFF across different field strengths, imager manufacturers, and reconstruction methods.

## Materials and Methods

No industry funding was used to support this meta-analysis. None of the authors were industry employees. The lead author (T.Y.) had full control of the data and the information submitted for publication.

## Definition and Criteria of PDFF

PDFF is defined as follows:

$$\text{PDFF} = \frac{\sum_{\text{all}} \text{PD}_{\text{FP}}}{(\text{PD}_{\text{WP}} + \sum_{\text{all}} \text{PD}_{\text{FP}})},$$

where PD is MR imaging–visible proton density, or equivalently the spectral peak area, of the water molecules having a single resonance frequency of 4.7 ppm, or the triglyceride molecules having multiple frequencies as described elsewhere; FP is fat peak; and WP is water peak (64). Various MR imaging–based methods of PDFF measurement have been proposed, including chemical shift two- and three-dimensional spoiled gradient-recalled echo sequences at 1.5 T and 3.0 T by using different reconstruction methods (45,49,58,65), as detailed in Appendix E1 (online).

The QIBA PDFF Biomarker Committee currently adopts the following protocol design criteria for MR imaging– and MR spectroscopy–based methods to measure PDFF (detailed in Table E1 [online]). Briefly, three major confounders of PDFF must be either minimized or corrected (66,67): the T1 relaxation effect, the T2 or T2\* relaxation effects, and multiple proton resonance frequencies of triglycerides (so-called spectral complexity). Various pulse sequence–specific confounders (eg, phase errors [68–72] and J coupling [73]) also need to be addressed either at acquisition or during reconstruction steps.

## Literature Search

This study was conducted according to Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines (74,75). Separate literature searches were performed for two pooled analyses for performance of MR imaging–PDFF: (a) linearity and bias and (b) precision. A systematic search of PubMed/MEDLINE, the Cochrane Library, and the Web of Science databases was performed on February 22, 2017, by a senior radiology resident (A.P.) to identify primary research studies satisfying inclusion criteria (or equivalent, depending on the syntax requirement of the specific search engine,

listed in Tables E2 and E3 (online) as follows: (a) linearity and bias analysis: liver “fat fraction” (imaging AND spectroscopy) “magnetic resonance” NOT review [publication type]; (b) precision analysis: liver “fat fraction” imaging (repeatability OR reproducibility OR precision OR agreement) “magnetic resonance” NOT review [publication type].

Only in vivo studies on humans that were either published in or translated to English were included.

## Study Selection

Titles and abstracts, followed by the full text of these eligible studies, were screened by the same author performing the systematic search (A.P.) and then independently verified by another author (T.Y., with 10 years of experience in MR imaging fat quantification methods) using the following exclusion criteria: (a) For either analysis: secondary analysis of previously published data, unable to verify or did not meet the above criteria for PDFF; (b) linearity and bias analysis: PDFF measurements were not performed by using both MR imaging and MR spectroscopy; (c) precision analysis: multiple PDFF measurements were not performed per participant.

## Data Collection and Quality Assessment

After all articles satisfying the selection criteria were identified, the corresponding authors of these articles were invited to submit anonymized individual participant data from MR imaging–PDFF and MR spectroscopy–PDFF for a meta-analysis. The local investigational review board of each participating institution either approved or waived formal review for the transfer and central analysis of anonymized study data. For each participant’s MR imaging–PDFF and MR spectroscopy–PDFF measurements, data listed in Table 1 were recorded in a pooled database. Data quality of included studies (ie, the bias and applicability of each study) was assessed by one author (S.D.S.) using the Quality Assessment of Studies of Diagnostic Accuracy 2 tool (76,77). The possibility of publication bias across studies was assessed by using funnel plots and Egger test (78).

**Table 1****Data Collected from Included Studies**

Parameter	Details
Field strength	1.5 T or 3.0 T
Manufacturer	GE, Siemens, or Philips
Reconstruction method	Magnitude, complex, or hybrid
No. of examinations	Repeated image acquisitions (if any) in two or more examinations by using independent instrument setup and/or calibration per examination
No. of acquisitions	Repeated image acquisitions (if any) during a single examination by using identical instrument setup and/or calibration
ROI	Repeated ROI placement in different locations in the liver per methods described in each study
MR imaging–PDFF (%)	Average PDFF within an ROI
MR spectroscopy–PDFF (%)	PDFF measurement colocalized to MR imaging ROI, if available

Note.—ROI = region of interest.

**Statistical Analysis**

The pooled data were analyzed by using R version 3.1.3 (R Foundation for Statistical Computing, Vienna, Austria) by one author (T.Y., with >10 years of experience in statistical computing) under supervision of an expert biostatistician (N.A.B., with >20 years of experience).

Linearity was initially evaluated by using a second-degree polynomial regression model of MR imaging–PDFF against MR spectroscopy–PDFF. A quadratic (second-order) term was first evaluated in the model; if the quadratic term did not reach statistical significance at the  $\alpha$  level of .05 or if its relative effect size was two orders of magnitude smaller than the linear (first-order) term, then the model was reduced to linear regression and the first- and zeroth-order terms (ie, slope and intercept) were estimated. A linear mixed-effects model was used for linear regression to account for clustered measurements within the same participants. The coefficient of determination ( $R^2$ ) was calculated as the strength metric of linearity (35).

Bias, defined as the average difference between MR imaging–PDFF and MR spectroscopy–PDFF measurements per participant (by using MR spectroscopy as the reference technique) was evaluated by using Bland-Altman analysis (79). The 95% limits

of agreement were calculated. To determine the contributing factors to bias, a linear mixed-effects model was used with field strength, imager manufacturer, and reconstruction method as fixed effects and with participant, ROI (nested within participant), examination, and acquisition as random effects.

Precision, defined as the closeness of agreement between repeated MR imaging–PDFF measurements (35), was evaluated by using a linear mixed-effects model, with field strength, manufacturer, reconstruction, participant, ROI (nested within participant), examination, and acquisition as random effects. Precision was measured as the standard deviation of PDFF measurement associated with each random effect. Repeatability was assessed as the precision of using identical equipment (same field strength, imager manufacturer, and reconstruction method), such as in a scan-rescan setting. Reproducibility was assessed as precision under varying circumstances (ie, different field strength, imager manufacturer, and/or reconstruction method), such as would be encountered in a multicenter setting. Both participant-level and ROI-level repeatability and reproducibility were assessed, because differences in ROI placement within the liver may contribute to PDFF variability due to biologic factors (rather than technical factors), especially in participants with nonuniform distribution of hepatic fat.

Heterogeneity of bias or precision across sites was not explicitly evaluated because differences in sites are modeled by (and expected to be strongly correlated with) the differences in the MR imaging equipment (ie, field strength, imager manufacturer) or reconstruction method. The 95% confidence intervals (CIs) and/or  $P$  values were computed for each statistic when appropriate.  $P$  values < .05 were considered to indicate statistical significance.

**Results****Study Selection and Subjects**

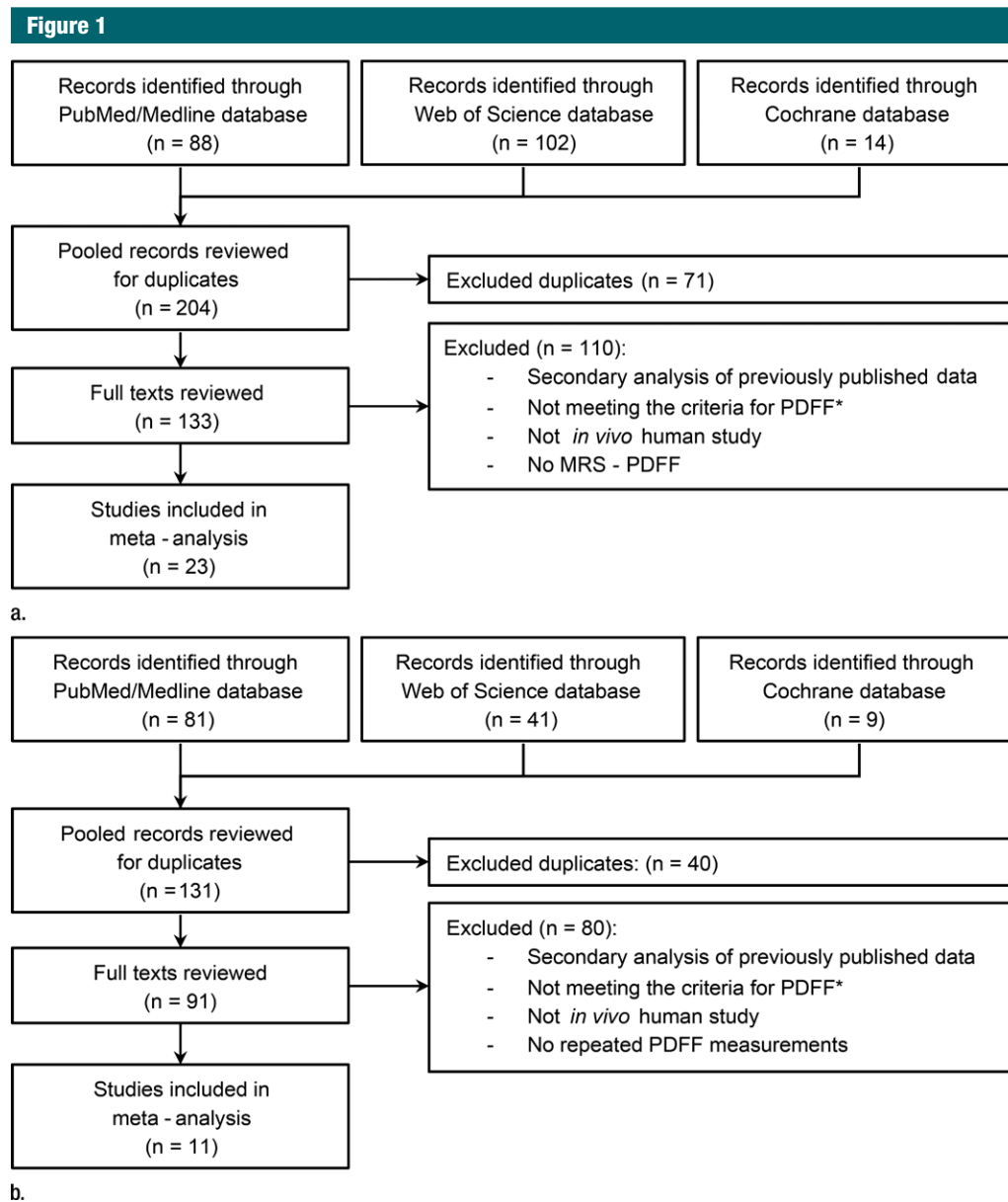
A flow diagram summarizing study selection for inclusion and exclusion criteria according to Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines is shown in Figure 1. Twenty-eight studies fulfilled the selection criteria. Anonymized data sets from all 28 selected studies were submitted for quantitative synthesis.

The pooled data included 16624 MR imaging–PDFF measurements in 1960 unique participants. For three studies (36,38,63) data were unavailable on the age and/or sex of 48, 50, and 10 participants, respectively. Mean participant age was 43 years (range, 8–89 years), with a man-to-woman ratio of 52%:48%. Mean MR imaging–PDFF value was 9.6% (range, –2.8% to 55.4%). A total of 1679 participants from 23 different studies were included in the linearity and bias analysis (36–58), 425 participants from 11 studies were included in the precision analysis (36,41,48,53,55,56,59–63), and 195 participants from six studies were included in both analyses.

Table 2 summarizes the characteristics of the included studies.

**Quality Assessment and Publication Bias**

Figure 2 summarizes quality assessment of the included studies by using the Quality Assessment of Studies of Diagnostic Accuracy 2 tool. No study met criteria for high risk of bias or applicability concern. Categories 2, 3, and 4 had greater than 90% compliance, suggesting that the studies were adequately blinded to



**Figure 1:** Flow diagrams show study selection according to Preferred Reporting Items for Systematic Reviews and Meta-Analyses guidelines for **(a)** linearity and bias analysis and **(b)** precision analysis. \* = Based on RSNA-QIBA PDFF Biomarker Committee for PDFF criteria as described in Materials and Methods section and Table E1 (online).

the reference standard and variations in reference standard were minimal. Categories 1 and 5 had approximately 60% compliance because some studies used case-control selection based on age, sex, and/or status of the following risk factors: healthy, obesity, diabetes, and/or nonalcoholic steatohepatitis. Because these risk factors were not considered to be high risk for bias or applicability

for this technical validation study, no specific subanalyses were performed. Funnel plots and Egger test *P* values are shown in Figure E2 (online). No statistically significant asymmetry was found to indicate publication bias (all *P* values > .5).

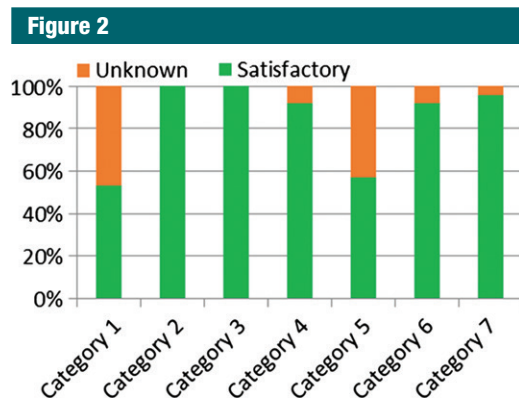
#### Assessment of Linearity

Figure 3a illustrates the relationship between the 3191 paired MR

imaging-PDFF and MR spectroscopy-PDFF measurements. In the second-degree polynomial model, the quadratic term was smaller than two orders of magnitude than was the linear term (ratio of quadratic to linear terms, 0.002), so it was removed and the model was reduced to a first-degree linear model. Linearity was strong ( $R^2 = 0.96$ ). The

**Table 2**  
**Characteristics of 28 Selected Studies**

First Author	Year	Study Region	Data Collection	No. of Participants	Mean Age (y)	Manufacturer	Field Strength (T)	Reconstruction Method	PDFF		Analysis
									Minimum (%)	Maximum (%)	
Hines (56)	2011	North America	Prospective	42	51	GE	1.5	Hybrid	-2.8	5.0	Both
Kang (55)	2011	North America	Prospective	21	55	GE, Siemens	1.5, 3.0	Magnitude	-0.1	15.1	Both
Johnson (53)	2014	North America	Prospective	30	38	GE	3.0	Hybrid	0.5	12.1	Both
Artz (48)	2015	North America	Prospective	25	48	GE	1.5, 3.0	Hybrid, magnitude	0.2	10.8	Both
Tyagi (41)	2015	North America	Prospective	29	24	GE	3.0	Hybrid, magnitude	0.1	13.0	Both
Motosugi (36)	2017	North America	Prospective	48	45	GE	1.5	Hybrid, magnitude	-0.2	6.2	Both
Maisamy (57)	2011	North America	Prospective	53	70	GE	1.5	Hybrid	0.5	6.6	Linearity and bias
Yokoo (58)	2011	North America	Prospective	163	39	GE	3.0	Magnitude	0.0	10.6	Linearity and bias
Motosugi (47)	2015	North America	Prospective	50	56	GE	3.0	Hybrid	0.0	6.6	Linearity and bias
Le (54)	2012	North America	Prospective	48	48	GE	3.0	Magnitude	1.9	18.4	Linearity and bias
Levin (52)	2014	North America	Prospective	84	36	GE	3.0	Magnitude	-0.8	10.8	Linearity and bias
Kühn (51)	2014	Europe	Prospective	50	57	Siemens	3.0	Hybrid	-0.2	13.1	Linearity and bias
Kühn (50)	2014	Europe	Prospective	33	57	Siemens	3.0	Hybrid	3.1	14.0	Linearity and bias
Bashir (49)	2015	North America	Prospective	42	53	Siemens	3.0	Hybrid	0.3	8.6	Linearity and bias
Rehm (44)	2015	North America	Prospective	132	13	GE	3.0	Hybrid	0.8	4.4	Linearity and bias
Zand (46)	2015	North America	Retrospective	286	14	GE	3.0	Magnitude	-0.1	12.2	Linearity and bias
Kukuk (45)	2015	Europe	Prospective	59	52	Philips	3.0	Complex	0.0	12.8	Linearity and bias
Kramer (38)	2016	North America	Prospective	50	56	GE	1.5	Hybrid	1.0	7.1	Linearity and bias
Kim (43)	2015	Asia	Retrospective	156	58	Siemens	3.0	Hybrid	0.3	5.2	Linearity and bias
Hetterich (42)	2016	Europe	Prospective	215	57	Siemens	3.0	Hybrid	0.6	9.2	Linearity and bias
Yu (40)	2016	North America	Prospective	35	60	Philips	3.0	Magnitude	1.0	14.3	Linearity and bias
Cui (39)	2016	North America	Prospective	50	54	GE	3.0	Magnitude	5.1	16.8	Linearity and bias
Park (37)	2016	North America	Prospective	31	53	GE	3.0	Magnitude	0.5	7.6	Linearity and bias
Wu (59)	2016	Asia	Prospective	15	33	GE	3.0	Hybrid	-1.8	3.5	Precision
Mashhood (63)	2013	North America	Prospective	10	50	GE, Siemens	1.5, 3.0	Magnitude	-0.1	8.1	Precision
Negrete (61)	2014	North America	Prospective	29	24	GE	3.0	Magnitude	-0.1	11.8	Precision
Sofue (60)	2015	North America	Prospective	150	57	Siemens	3.0	Hybrid	0.5	6.9	Precision
Serai (62)	2017	North America	Prospective	24	39	GE, Philips	1.5, 3.0	Complex, hybrid	0.3	5.8	Precision



**Figure 2:** Bar chart shows estimates of percentage compliance by using Quality Assessment of Studies of Diagnostic Accuracy 2 tool. All 28 studies had moderate to high scores and low risk of bias in all seven categories; all fulfilled five or more of the seven quality categories. Summary of scores for each category were as follows: Category 1, Risk of bias—Patient selection, 57% (16 of 28); Category 2, Risk of bias—Index test, 100% (28 of 28); Category 3, Risk of bias—Reference standard, 100% (28 of 28); Category 4, Risk of bias—Flow and timing, 93% (26 of 28); Category 5, Applicability concerns—Patient selection, 61% (17 of 28); Category 6, Applicability concerns—Index test, 93% (26 of 28); and Category 7, Applicability concerns—Reference standard, 96% (27 of 28).

estimated intercept of  $-0.07$  was not significantly different from zero ( $P = .70$ ; 95% CI:  $-0.50, 0.32$ ). The estimated slope of the regression line was significantly below unity at  $0.97$  ( $P < .001$ ; 95% CI:  $0.96, 0.98$ ), indicating underestimation by using MR imaging–PDFF compared with MR spectroscopy–PDFF (corresponds to about 1.5% underestimation at 50% MR spectroscopy–PDFF).

### Assessment of Bias

Figure 3b illustrates the MR imaging–PDFF measurement bias. The mean bias was small ( $-0.13\%$ ; 95% limits of agreement:  $-3.95\%, 3.70\%$ ). In the analysis of individual bias components (Table 3), all effects except manufacturer (between GE and Siemens) had statistically significant effects on bias, but most bias components had effects smaller than 1.5% in absolute PDFF. Bias because of manufacturer for the Philips system was the exception, with approximately 2% higher PDFF values compared with those obtained with either GE or Siemens systems.

### Assessment of Precision

Figure 4 illustrates the precision profiles, represented as the difference between the repeated measurements as a function of the per-participant (Fig 4a) and per-ROI (Fig 4b) mean MR imaging–PDFF values based on 9103 measurements in 425 participants.

Under repeatability conditions (same field strength, imager manufacturer, and reconstruction method), within-participant standard deviation was 1.69% and within-ROI standard deviation was 1.08% in absolute PDFF value. Under reproducibility conditions (different field strength, imager manufacturer, or reconstruction method), within-participant standard deviation was 1.97% and within-ROI standard deviation was 1.48% in absolute PDFF value.

The estimated variance components from linear mixed-effects models (Fig 4) indicated that the main contributors to MR imaging–PDFF measurement variability were ROI locations (standard deviation, 1.30%) and random measurement error because of

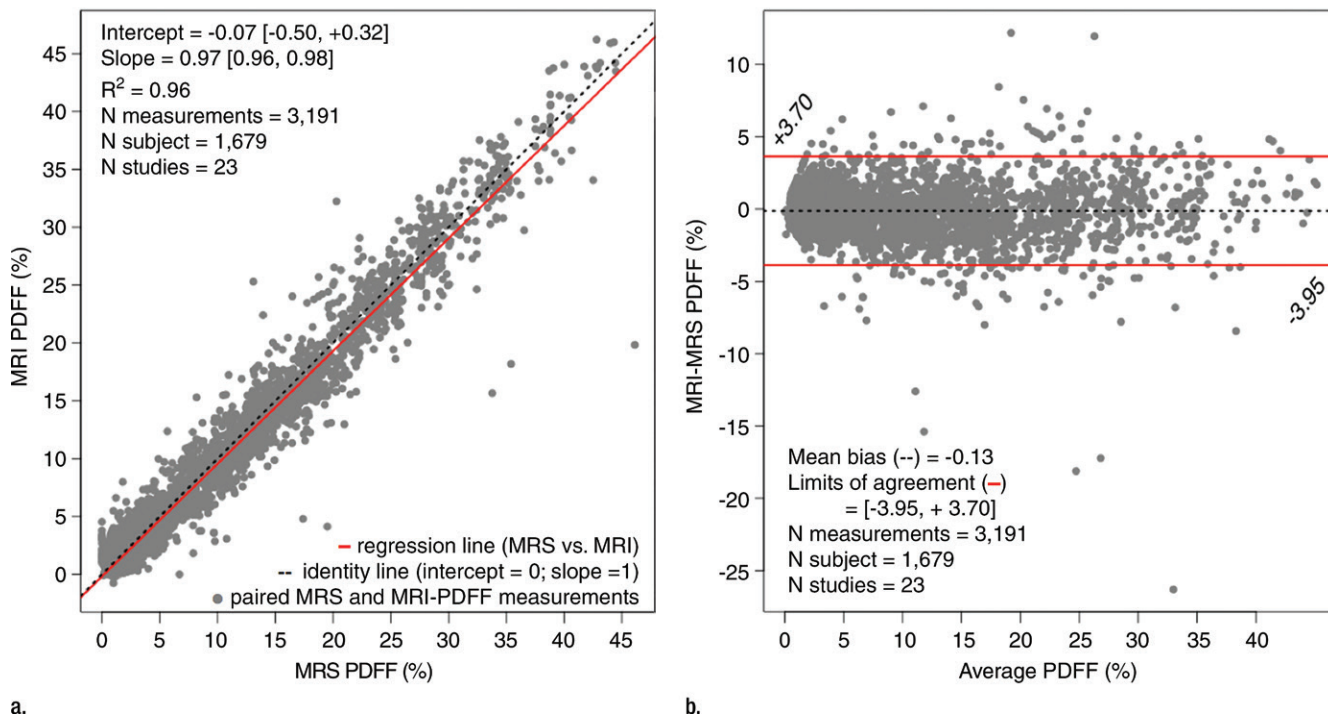
repeated acquisitions (standard deviation, 1.07%). Other technical factors (field strength, imager manufacturer, and reconstruction method) invariably had smaller standard deviations of less than 0.8% in absolute PDFF value.

### Discussion

Our meta-analysis included nearly 2000 participants in 28 independent primary research studies from geographically diverse sites that varied in field strength, imager manufacturer, and reconstruction method. These studies were generally considered satisfactory in quality per Quality Assessment of Studies of Diagnostic Accuracy 2 criteria with minimal bias in index test (MR imaging–PDFF) and reference standard (MR spectroscopy–PDFF for linearity and bias analyses and repeated MR imaging–PDFF for precision analyses). Although patient selection criteria in some studies were not uniform because of emphasis in different subpopulations of various age range, sex, and steatosis risk factors (eg, obesity, diabetes), the potential impact of the variability is thought to be small in this meta-analysis of technical validation.

Results of our meta-analysis demonstrated that MR imaging–PDFF has excellent linearity and negligible bias with respect to the reference standard of MR spectroscopy–PDFF measurements over the entire range of observed steatosis severity. A slight deviation of the MR imaging versus MR spectroscopy regression from perfect agreement was statistically significant, but the effect size was small (up to 1.5% absolute PDFF value in very high PDFF range) and is unlikely to be meaningful either clinically or in research studies. In addition, our results demonstrated that the largest contributors to the variability in MR imaging–PDFF measurements (repeatability and reproducibility) were inherent heterogeneity of steatosis across the liver (ie, because of different ROI locations; standard deviation, 1.3% in PDFF) and the random measurement error (1%), with smaller contributions ( $<0.8\%$ ) from technical factors such

Figure 3



**Figure 3:** Scatterplots show linearity and bias of MR imaging-PDF. **(a)** Linearity of MR imaging-PDF against MR spectroscopy-PDF, both measured in colocalized ROIs in the liver. Red line represents linear regression fit (after correcting for within-participant correlations between replicated measurements, if any) with coefficient of determination indicating very strong linear fit ( $R^2 = 0.96$ ). Estimated intercept ( $-0.07\%$ ) was not significantly different from zero; estimated slope was close to but less than unity (0.97), indicating very slight underestimation of MR imaging-PDF values at larger MR spectroscopy-PDF values. **(b)** Bland-Altman plot of MR imaging-PDF relative to MR spectroscopy-PDF as reference technique demonstrates very small mean bias ( $-0.13\%$ ) and limits of agreement within  $\pm 4\%$ . Estimated size of various bias components are presented in Table 3.

as field strength, imager manufacturer, and reconstruction method. Based on these observations, we conclude that MR imaging-PDF has excellent technical performance characteristics as a QIB for widespread use in clinical trials and patient care.

Our meta-analysis findings are consistent with previous single-center studies demonstrating high technical accuracy and precision of MR imaging-PDF as a QIB of hepatic steatosis. These earlier data suggested that PDFF should be well suited for multicenter trials and longitudinal assessment of hepatic steatosis. Our meta-analysis provides higher-level evidence supporting this claim. Several observational studies and therapeutic trials of nonalcoholic fatty liver disease demonstrated the feasibility of using PDFF as a study end point (44,46,80), as well as a patient-level diagnostic and

Table 3

## Bias Components of MR Imaging-PDF

Bias Components	Estimated Change PDFF (%)	P Value
Intercept	0.42 (0.21, 0.63)	<.0001
Manufacturer		
GE	Reference	...
Siemens	-0.01 (-0.19, 0.17)	.933
Philips	2.24 (1.51, 2.96)	<.0001
Reconstruction method		
Magnitude	Reference	...
Complex	-1.36 (-2.22, -0.50)	.002
Hybrid	0.92 (0.80, 1.05)	<.0001
Field strength		
1.5 T	Reference	...
3.0 T	-1.20 (-1.40, -1.00)	<.0001

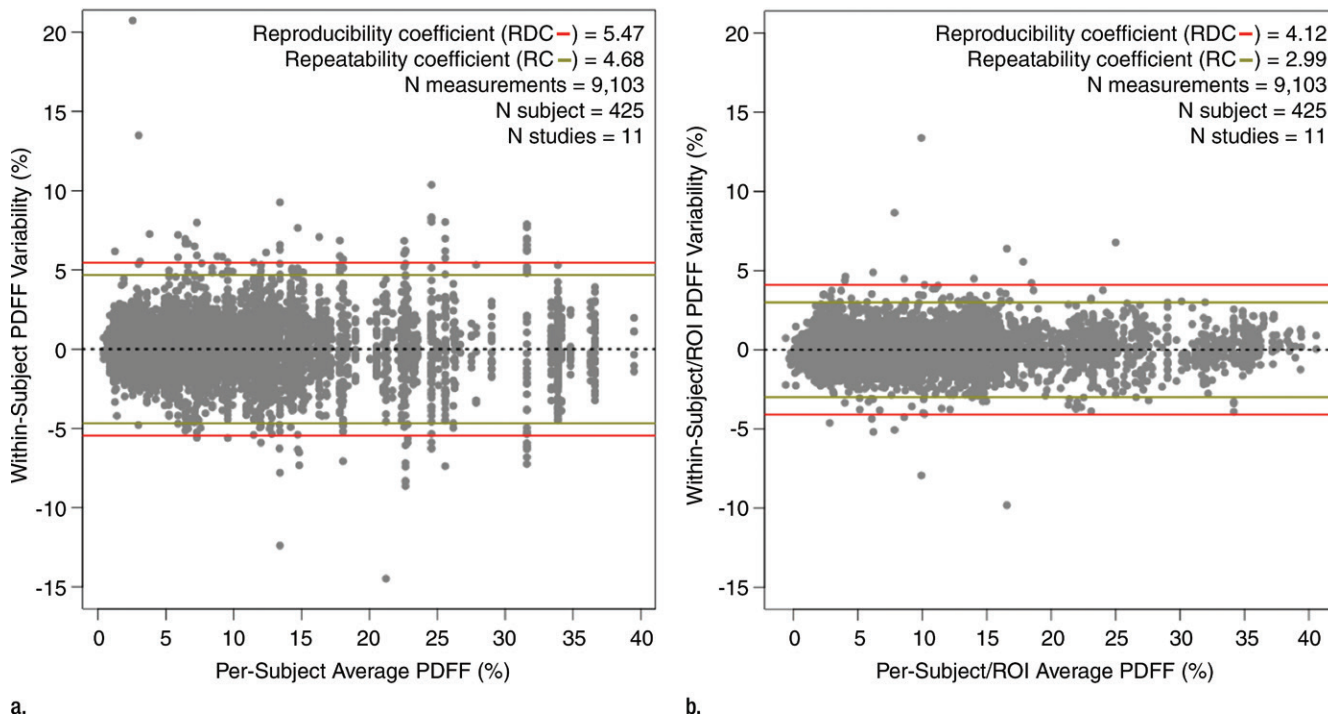
Note.—Data in parentheses are 95% CIs.

therapeutic response biomarker for clinical practice (42,81-84). In addition, PDFF methods are used in large

population-based cohort trials (including studies with more than 1000 participants) (85-87). U.S. Food and Drug



Figure 4



**Figure 4:** (a, b) Scatterplots illustrate participant-level and ROI-level precision of MR imaging–PDFF measurements, respectively, because of differences in field strength, imager manufacturer, reconstruction method, equipment setup, and random noise (effect sizes are detailed in Table 4). Repeatability refers to precision under an identical experimental condition (ie, scan-rescan repeatability by using fixed hardware and software). Reproducibility refers to precision under variable experimental condition (ie, by using different hardware or software). At ROI level (b), MR imaging–PDFF is highly precise with repeatability coefficients (RCs) and reproducibility coefficients (RDCs) indicating the 95th percentile of precision to be approximately  $\pm 3\%$  and  $\pm 4\%$ , respectively. At participant level (a), repeatability and reproducibility are approximately  $\pm 5\%$  and  $\pm 5.5\%$ . Subject-level precision is lower than ROI-level precision because of differences in ROI placement, that is, heterogeneity of underlying steatosis. In both cases, RCs and RDCs are similar to each other, indicating small impact of technical factors (field strength, imager manufacturer, reconstruction method) on the precision of the measurement.

Administration–approved MR imaging pulse sequences for hepatic PDFF measurement are now available from major imager manufacturers worldwide including GE, Siemens, and Philips.

Our work was motivated by the activities of QIBA, whose mission is to improve the practical value of QIBs by promoting standardization to reducing variability between hardware and software, thereby facilitating the use of QIBs in clinical trials and patient care. These goals are achieved by preparing a QIBA profile document that is intended to span various commonly encountered potential confounders, such as field strength, imager manufacturer, and reconstruction methods, so that expected variability is included to the maximum extent possible. This profile document is intended for a broad audience, including

imager and third-party device manufacturers, pharmaceutical companies, diagnostic agent manufacturers, medical imaging sites, imaging contract research organizations, physicians, technologists, researchers, professional organizations, educational institutions, and various accreditation and regulatory authorities. To this purpose, it is important for the linearity, bias, and precision claimed in the profile to be realistic and reasonably achievable across imaging centers and readers spanning a relevant range of technical variations. Our meta-analysis aims to provide generalizable technical performance standards to inform future QIBA profile statements.

This study had several strengths and limitations. Major strengths of this meta-analysis included a large and geographically diverse cohort, multiple

studies conducted by independent research groups, and variability in hardware and software as expected in real-world applications. An important and unique strength of this work was the close collaboration among the authors of the selected articles that enabled statistically powerful pooled analyses directly on the collective MR imaging–PDFF and MR spectroscopy–PDFF individual participant data, rather than the summary statistics extracted from the published articles as is done in a traditional meta-analysis. The main limitation of this study was that we used MR spectroscopy as the reference standard for linearity and bias analyses, rather than a tissue-based reference standard. However, MR spectroscopy has been widely accepted as the reference standard for hepatic fat quantification

Table 4

## Within-Subject Variance Components of MR Imaging–PDFF

Component	Variable ROI and Variable Technique (%)	Variable ROI and Fixed Technique (%)	Fixed ROI and Variable Technique (%)	Fixed ROI and Fixed Technique (%)
Nonuniform steatosis (different ROI location)	1.30	1.30	...	...
Manufacturer (GE, Siemens, or Philips)	0.81	...	0.81	...
Field strength (1.5 T or 3.0 T)	0.51	...	0.51	...
Reconstruction method (magnitude, complex, or hybrid)	0.35	...	0.35	...
Interexamination (equipment setup)	0.16	0.16	0.16	0.16
Interacquisition (random error)	1.07	1.07	1.07	1.07
Sum of components	1.98	1.69	1.49	1.08

Note.—Data are standard deviations. Data are from 11 precision studies including 425 participants, with two or three examinations per participant. See Figure 4 for repeatability and reproducibility coefficients.

in research studies to derive highest levels of evidence, including randomized control trials and epidemiologic studies (18–34). Considering the large number of studies available comparing MR imaging and MR spectroscopy, MR spectroscopy is probably the most widely available reference, particularly in the light of high variability of biopsy to quantify hepatic steatosis (12,88).

In summary, our meta-analysis of pooled data collected from 28 published studies demonstrated excellent linearity, negligible bias, and high precision of MR imaging–PDFF across different field strengths, imager manufacturers, and reconstruction methods. We conclude that MR imaging–PDFF has excellent technical performance characteristics for widespread use in clinical trials and patient care.

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