



Assessment of myocardial perfusion by dynamic O-15-labeled water PET imaging: validation of a new fast factor analysis

Adachi, I ; Gaemperli, O ; Valenta, I ; Schepis, T ; Siegrist, P T ; Treyer, V ; Burger, C ; Morita, K ; Kaufmann, P A

Abstract: BACKGROUND: Factor analysis (FA) is an established method for separating myocardium from blood pool by use of oxygen 15-labeled water and positron emission tomography for analyzing myocardial blood flow (MBF). Conventional FA methods generating images from sinograms (sinoFA) are time-consuming, whereas FA can be performed on the reconstructed images (reconFA) in a fraction of time. We validated the MBF values obtained by reconFA versus sinoFA. METHODS AND RESULTS: In 23 volunteers (mean age, 26.6 +/- 3.4 years) MBF was calculated from sinoFA and reconFA and blindly reanalyzed 1 month later by the same observer. Intraobserver agreement and reconFA-versus-sinoFA agreement were assessed according to Bland and Altman (BA). Reproducibility proved excellent for global sinoFA ($r = 0.968$; $P < .001$; BA limits, -0.617 to 0.676 mL x min⁻¹ x g⁻¹) and slightly superior for reconFA ($r = 0.979$; $P < .001$; BA limits, -0.538 to 0.558 mL x min⁻¹ x g⁻¹), with wider limits of agreement for segmental MBF from sinoFA ($r = 0.777$; $P < .001$; BA limits, -1.676 to 1.656 mL x min⁻¹ x g⁻¹) and reconFA ($r = 0.844$; $P < .001$; BA limits, -1.999 to 1.992 mL x min⁻¹ x g⁻¹). In addition, sinoFA and reconFA showed excellent correlation ($r = 0.975$, $P < .001$) and agreement (BA limits, -0.528 to 0.648 mL x min⁻¹ x g⁻¹) for global and segmental values ($r = 0.955$; $P < .001$; BA limits, -1.371 to 1.491 mL x min⁻¹ x g⁻¹). CONCLUSIONS: Use of reconFA allows rapid and reliable quantitative MBF assessment with O-15-labeled water.

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Assessment of myocardial perfusion by dynamic O-15-labeled water PET imaging: Validation of a new fast factor analysis

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Background. Factor analysis (FA) is an established method for separating myocardium from blood pool by use of oxygen 15-labeled water and positron emission tomography for analyzing myocardial blood flow (MBF). Conventional FA methods generating images from sinograms (sinoFA) are time-consuming, whereas FA can be performed on the reconstructed images (reconFA) in a fraction of time. We validated the MBF values obtained by reconFA versus sinoFA.

Methods and Results. In 23 volunteers (mean age, 26.6 ± 3.4 years) MBF was calculated from sinoFA and reconFA and blindly reanalyzed 1 month later by the same observer. Intraobserver agreement and reconFA-versus-sinoFA agreement were assessed according to Bland and Altman (BA). Reproducibility proved excellent for global sinoFA ($r = 0.968$; $P < .001$; BA limits, -0.617 to 0.676 $\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) and slightly superior for reconFA ($r = 0.979$; $P < .001$; BA limits, -0.538 to 0.558 $\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$), with wider limits of agreement for segmental MBF from sinoFA ($r = 0.777$; $P < .001$; BA limits, -1.676 to 1.656 $\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) and reconFA ($r = 0.844$; $P < .001$; BA limits, -1.999 to 1.992 $\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$). In addition, sinoFA and reconFA showed excellent correlation ($r = 0.975$, $P < .001$) and agreement (BA limits, -0.528 to 0.648 $\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) for global and segmental values ($r = 0.955$; $P < .001$; BA limits, -1.371 to 1.491 $\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$).

Conclusions. Use of reconFA allows rapid and reliable quantitative MBF assessment with O-15-labeled water. (J Nucl Cardiol 2007;14:698-705.)

Key Words: Positron emission tomography • myocardial blood flow • water • factor analysis • reproducibility

Positron emission tomography (PET) with oxygen 15-labeled water is a noninvasive method for accurately quantifying the regional myocardial blood flow (MBF), because O-15-labeled water has the following major advantages: free diffusion, kinetics independent of changes in the myocardial metabolism, and a short half-life (2.1 minutes), allowing sequential measure-

ments with a low radiation burden for patients.¹⁻⁴ The myocardial tissue cannot be directly distinguished from blood pool on the dynamic O-15-labeled water images, because O-15-labeled water is not trapped by the myocardial cells. This limitation hinders the use of this tracer in routine clinical practice. O-15-labeled carbon monoxide, which permits labeling of the vascular volume, has been used in combination with O-15-labeled water for delineation of the myocardial wall from other anatomic structures to obtain tissue O-15-labeled water time-activity curves (TACs) suitable for MBF measurement via compartmental analysis.⁵ This method is standard in several centers for the determination of myocardial perfusion.⁶⁻⁸ Alternatively, processing of O-15-labeled water PET dynamic image sequences with factor analysis (FA) methods has been proposed and validated to generate high-quality myocardial factor images from the sinograms, avoiding the use for the O-15-labeled carbon monoxide scan.^{9,10} This method is relatively time-consuming, because of the involved iterative reconstructions. Some investigators proposed another method that

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was close to FA and based on the linear reduction of signal in the ventricular cavity.¹¹ This method can perform FA on the reconstructed images (reconFA), not the sinograms, and therefore it can create factor images more quickly than the conventional method. However, the MBF values obtained from images by reconFA have not been validated. The aim of this study, therefore, was to validate the MBF values obtained by reconFA versus those obtained from sinograms (sinoFA) and to document their interchangeability.

METHODS

Study population

We consecutively enrolled 23 healthy volunteers (17 men and 6 women; mean age, 26.6 ± 3.4 years) in our study. This group was chosen because the incidence of coronary artery disease (CAD) increases sharply beyond 50 years of age.¹² None of the volunteers had a history of a prior cardiac event or elevated total or low-density lipoprotein cholesterol levels, hypertension, or diabetes mellitus. The inclusion criteria were normal resting electrocardiogram, normal heart rate, normal blood pressure, and unremarkable excursive cardiopulmonary examination. All female study participants had a negative urine pregnancy test. The study protocol was approved by the local institutional review board, and all subjects gave written informed consent before enrollment in the study.

PET

All subjects refrained from ingesting caffeinated beverages or theophylline-containing medications for 24 hours before the PET study. The MBF at rest and during continuous administration of adenosine, $0.140 \text{ mg} \cdot \text{kg}^{-1} \cdot \text{min}^{-1}$, infused intravenously over a period of at least 6 minutes (stress) was quantified noninvasively with O-15-labeled water and serial PET imaging.¹³ Scanning was performed with an Advance PET scanner (GE Healthcare Technologies, Waukesha, Wis). This device records 35 image planes simultaneously in 2-dimensional mode. The axial field of view was 14.5 cm. A 30-minute blank scan was recorded as part of the daily routine practice. A 20-minute transmission scan via an external germanium 68 source was performed for attenuation correction. Starting after the background frame, a dose of 500 to 700 MBq of O-15-labeled water was injected as an intravenous bolus over a period of 20 seconds at an infusion rate of 24 mL/min to assess MBF. Dynamic images were obtained for 4 minutes 40 seconds (14 frames for 5 seconds each, 3 frames for 10

seconds, 3 frames for 20 seconds, and 4 frames for 30 seconds).

Image processing

Transaxial images of the O-15-labeled water distribution over time were reconstructed via filtered back-projection with an 8-mm Hanning filter and a zoom factor of 1.72. The resulting image volumes had a matrix of $128 \times 128 \times 35$ and a pixel size of $2.34 \times 2.34 \times 4.25 \text{ mm}^3$. Images of the transmission data were reconstructed with the same geometry so that matched emission and transmission images were available. In addition, sinograms were calculated from the raw emission data, which were corrected for attenuation and scatter.

The FA approach developed by Hermansen et al⁹ was used to calculate anatomic images of the myocardium and the ventricular blood volume. As a first step, volumes of interest (VOIs) were outlined within the left and right lungs by use of the anatomic information available in the transmission images. These lung VOIs were projected to the dynamic images for calculation of the average lung TAC. Approximations of the activity in the left ventricle (LV) and right ventricle (RV) were derived by time-shifting the lung TAC (RV, -5 seconds; LV, 5 seconds). A third curve representing myocardial uptake was obtained by solving the 1-compartment water model with the shifted lung TAC (8 seconds) as the input curve, a mean MBF value of $1 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$ tissue, and a partition coefficient of 0.96. Two FA procedures were applied to the TACs as described by Hermansen et al. The first used the myocardium TAC as the variate and the LV and RV TACs as the covariate, resulting in the myocardium factors. The second used the lung TAC as the variate and the myocardium TAC as the covariate, resulting in the blood factors.

These factors were then applied to the emission data in two different ways. (1) As described in the original approach (sinoFA),⁹ the corrected sinograms were weighted by the respective factors and added, resulting in a set of myocardial factor sinograms and a set of blood volume sinograms. The factor sinograms were then reconstructed into factor images of the myocardium and the blood volume via an ordered-subsets expectation maximization algorithm with 4 iterations, 14 subsets, and 2 final expectation maximization iterations. (2) As a fast alternative (reconFA), the factors were directly applied to the dynamic emission images.

The resulting factor images of either approach were used for the quantification process as follows. First, the factor images were resliced into the standard short-axis orientation (Figure 1). Next, the myocardium centerline was traced in all slices from the apex to the base (at least 12 consecutive slices). From this

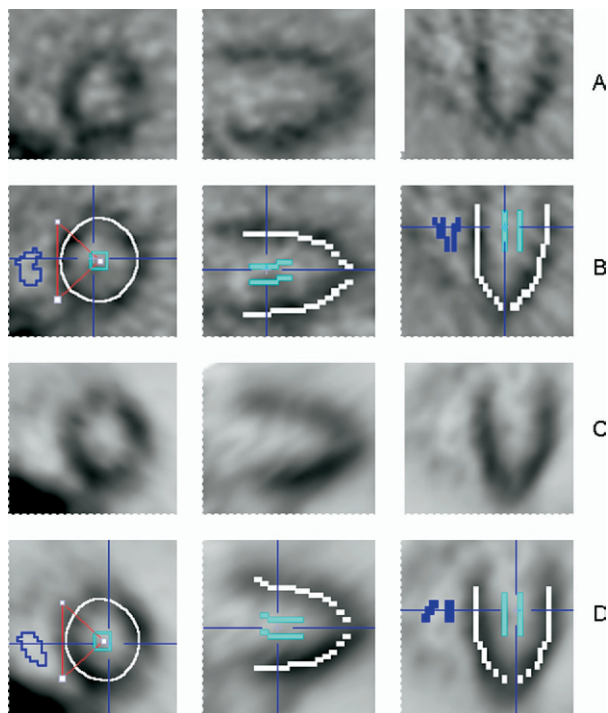


Figure 1. After FA, cardiac images were reoriented to short axis (left column), vertical long axis (middle column), and horizontal long axis (right column) for both sinoFA (A and B) and reconFA (C and D) on which the regions of interest were drawn. Both methods allow clear discrimination of myocardium from blood pool.

centerline definition, 16 segmental VOIs were generated, representing the segments as recommended by the American Heart Association.¹⁴ All image pixels within a range of ± 1 pixel from the centerline were included in the VOIs. By use of these VOIs, the myocardial TACs in the 16 segments were calculated from the dynamic O-15-labeled water images. Two additional VOIs were outlined in the cavities of the LV and RV on 6 consecutive slices to calculate the blood LV and RV TACs.

The MBF in the segments was quantified by use of a 1-tissue compartment model with the LV TAC as the input curve and 2 spillover terms. Spillover from the LV was considered in all segments, whereas spillover from the RV was set to 0 for all non-septal segments. The analysis resulted in segmental values of MBF in milliliters per minute per gram, and a global MBF value was obtained by volume-weighted averaging of the segmental results. All post-processing was performed with commercially available software (PMOD Technologies, Adliswil, Switzerland) that includes the FA and the modeling as a part of the myocardial quantification module.

Statistical analysis

The MBF values are expressed as mean \pm SD at rest, at stress, and as the average of both (all pooled). Intraobserver reproducibility and comparison of MBF values obtained by sinoFA versus reconFA was performed by linear regression analysis and by the assessment of the limits of agreement as proposed by Bland and Altman¹⁵ for global and segmental MBF for rest and stress values separately and for all values pooled. Coronary flow reserve was calculated as the ratio of hyperemic over resting MBF (relative values).

RESULTS

All rest and adenosine stress procedures were well tolerated, and no subjects were withdrawn from the analysis. None of the subjects had any significant electrocardiographic changes or intolerable alterations of blood pressure.

Intraobserver reproducibility study

There were no significant differences between baseline and repeat analysis for the resting (Table 1) and stress MBF (Table 2) by use of sinoFA and reconFA on global MBF. The percentage of reproducibility coefficient at stress was slightly lower than that at rest. Intraobserver reproducibility of baseline and repeat global MBF (rest, stress, and pooled) was excellent, as evidenced by a high correlation coefficient and narrow Bland-Altman (BA) limits of agreement for sinoFA ($n = 46$; $r = 0.968$; standard error of the estimate [SEE], $0.335 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$; $P < .001$; BA limits, -0.617 to $0.676 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) and even slightly superior for reconFA ($n = 46$; $r = 0.979$; SEE, $0.279 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$; $P < .001$; BA limits, -0.538 to $0.558 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) (Figure 2). No significant differences between baseline and repeat analysis for the resting and stress MBF were observed by use of sinoFA and reconFA on segmental MBF. The reproducibility coefficient for segmental MBF was higher than that for global MBF. However, good reproducibility was maintained for segmental analysis with sinoFA ($r = 0.777$; SEE, $0.992 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$; $P < .001$; BA limits, -1.676 to $1.656 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) and with reconFA ($r = 0.844$; SEE, $0.824 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$; $P < .001$; BA limits, -1.999 to $1.992 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) (Figure 3).

Comparison of MBF from sinoFA versus reconFA

MBF by sinoFA and reconFA compared well at rest ($1.08 \pm 0.16 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$ vs $0.97 \pm 0.14 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) and at stress ($3.56 \pm 0.44 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$ vs 3.54 ± 0.47

Table 1. Intraobserver reproducibility at rest for global and segmental MBF ($\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$)

Segment	Rest				Rest			
	sinoFA baseline	sinoFA repeated	abs	%	reconFA baseline	reconFA repeated	abs	%
1 APEX_Septum	1.17 ± 0.42	1.20 ± 0.49	0.10	8	1.16 ± 0.34	1.05 ± 0.23	0.18	16
2 MID_Septum-Inferior	1.03 ± 0.26	0.97 ± 0.26	0.14	14	0.88 ± 0.21	0.87 ± 0.31	0.04	5
3 MID_Septum-Anterior	1.06 ± 0.35	1.00 ± 0.29	0.08	8	0.98 ± 0.23	1.11 ± 0.37	0.29	27
4 BASAL_Septum-Inferior	1.16 ± 0.56	1.18 ± 0.36	0.18	16	1.00 ± 0.36	0.98 ± 0.38	0.08	8
5 BASAL_Septum-Anterior	0.94 ± 0.34	0.92 ± 0.18	0.02	2	1.05 ± 0.41	0.99 ± 0.16	0.12	12
6 APEX_Anterior	1.02 ± 0.29	1.01 ± 0.42	0.02	2	1.03 ± 0.28	1.09 ± 0.38	0.16	15
7 MID_Anterior	1.17 ± 0.34	1.07 ± 0.24	0.22	20	1.11 ± 0.36	0.99 ± 0.22	0.25	24
8 BASAL_Anterior	1.25 ± 0.54	1.1 ± 0.23	0.45	39	1.04 ± 0.27	0.94 ± 0.21	0.18	18
9 APEX_Lateral	1.10 ± 0.25	1.12 ± 0.23	0.08	7	0.99 ± 0.23	0.96 ± 0.22	0.06	6
10 MID_Lateral	1.23 ± 0.38	1.16 ± 0.24	0.20	17	1.09 ± 0.29	0.98 ± 0.13	0.22	21
11 BASAL_Lateral	1.07 ± 0.32	0.98 ± 0.20	0.18	17	0.93 ± 0.25	0.92 ± 0.21	0.02	2
12 APEX_Inferior	1.05 ± 0.37	1.13 ± 0.39	0.16	15	0.97 ± 0.35	0.84 ± 0.35	0.25	27
13 MID_Inferior-Posterior	0.98 ± 0.35	0.90 ± 0.28	0.08	8	0.90 ± 0.37	0.75 ± 0.24	0.25	30
14 MID_Inferior-Inferior	1.12 ± 0.73	1.13 ± 0.53	0.02	2	0.93 ± 0.29	0.8 ± 0.27	0.25	29
15 BASAL_Inferior-Posterior	1.10 ± 0.60	1.06 ± 0.28	0.31	30	0.91 ± 0.34	0.85 ± 0.22	0.12	14
16 BASAL_Inferior-Inferior	1.16 ± 0.66	1.10 ± 0.34	0.37	35	0.87 ± 0.29	0.91 ± 0.29	0.02	2
Global	1.12 ± 0.20	1.04 ± 0.16	0.16	15	1.00 ± 0.22	0.95 ± 0.12	0.10	11

abs, Absolute value of reproducibility coefficient; %, percent of reproducibility coefficient.

$\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$), resulting in similar coronary flow reserve values (3.34 ± 0.5 vs 3.69 ± 0.66) without a significant difference ($P =$ not significant for all comparisons). The comparison of sinoFA and reconFA revealed excellent correlation ($n = 46$, $r = 0.978$, $P < .001$) for global MBF values with narrow BA limits of agreement (-0.528 to $0.648 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$). The excellent correlation was preserved also for segmental values ($n = 736$, $r = 0.955$, $P < .001$) with narrow BA limits of agreement (-1.371 to $1.491 \text{ mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$) (Table 3 and Figure 4).

DISCUSSION

This study confirms an excellent agreement of MBF assessed with O-15-labeled water and PET via both methods for FA (ie, reconFA and sinoFA). The validity and reliability of reconFA are further substantiated by its high intraobserver reproducibility, which was slightly superior than that for sinoFA. This equally applies to global and segmental MBF estimates.

The use of factor images has been introduced into nuclear cardiology by Hermansen et al,⁹ and its reproducibility for assessing MBF has been extensively validated by use of various stimuli such as adenosine, bicycle exercise stress, dobutamine, and cold pressor test.¹⁶⁻¹⁹

The novelty of the present method lies in the fact that the FA no longer needs to be performed on the sinograms. With sinograms, there is usually a very large data set, and therefore the processing is rather cumbersome and time-consuming. With the new method, the analysis can be performed on the reconstructed images. As these have a much smaller size, the FA on such data sets can now be done considerably more quickly, shortening the processing time by about 1 order of magnitude. This, however, is not achieved at the cost of any loss in the signal-to-noise ratio or image quality, resulting in highly comparable MBF values. A strength of our method, representing an advantage over previously reported approaches by other groups using regularized factor images,¹¹ is that we use lung regions of interest to determine lung kinetics, which improves the signal-to-noise ratio. Therefore, with our method, there was no loss in image quality at stress compared with rest, as opposed to the latter method reported by Frouin et al.¹¹ A direct comparison between the sinoFA and reconFA methods by BA analysis revealed an excellent agreement of the 2 methods for global and segmental MBF over a wide range of flow. In addition, the MBF values obtained in our study with reconFA and sinoFA compare well with the numbers in healthy volunteers reported in the literature.^{17,19-21}

Table 2. Intraobserver reproducibility at stress for global and segmental MBF ($\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$)

Segment	Stress				Stress			
	sinoFA baseline	sinoFA repeated	abs	%	reconFA baseline	reconFA repeated	abs	%
1 APEX_Septum	3.49 ± 0.71	4.04 ± 1.35	1.23	32	4.24 ± 0.90	3.99 ± 0.96	0.51	12
2 MID_Septum-Inferior	3.65 ± 1.08	3.88 ± 1.21	0.37	10	3.56 ± 1.06	3.77 ± 0.99	0.04	1
3 MID_Septum-Anterior	3.57 ± 1.18	3.76 ± 1.93	0.29	8	3.93 ± 1.25	3.99 ± 1.16	0.49	13
4 BASAL_Septum-Inferior	3.24 ± 1.28	3.03 ± 0.95	0.25	8	2.74 ± 0.63	3.04 ± 1.01	0.76	26
5 BASAL_Septum-Anterior	3.59 ± 1.51	3.75 ± 1.33	0.1	3	3.48 ± 1.26	3.91 ± 1.92	0.45	13
6 APEX_Anterior	3.57 ± 1.17	3.61 ± 1.06	0.1	3	3.55 ± 0.79	3.86 ± 0.91	0.59	16
7 MID_Anterior	3.51 ± 0.99	3.66 ± 1.05	0.12	3	3.55 ± 0.92	3.64 ± 0.61	0.06	2
8 BASAL_Anterior	3.70 ± 0.99	3.43 ± 0.72	0.98	28	3.41 ± 1.04	3.33 ± 1.18	0.29	9
9 APEX_Lateral	3.60 ± 0.83	3.49 ± 0.78	0.22	6	3.62 ± 0.96	3.64 ± 0.76	0.16	4
10 MID_Lateral	3.72 ± 1.08	3.82 ± 0.91	0.33	9	3.59 ± 0.61	3.81 ± 0.88	0.51	14
11 BASAL_Lateral	3.91 ± 1.29	3.36 ± 0.93	1.27	35	3.23 ± 0.97	3.44 ± 0.95	0.18	6
12 APEX_Inferior	3.47 ± 0.91	3.59 ± 1.26	0.43	12	3.23 ± 0.96	3.25 ± 0.86	0.25	8
13 MID_Inferior-Posterior	3.56 ± 0.76	3.97 ± 2.00	0.78	21	3.87 ± 1.25	3.64 ± 1.10	0.31	8
14 MID_Inferior-Inferior	3.44 ± 0.89	3.47 ± 1.05	0.16	5	3.73 ± 1.01	3.76 ± 1.18	0.22	6
15 BASAL_Inferior-Posterior	3.34 ± 1.12	3.42 ± 1.34	0.02	1	3.15 ± 1.33	3.26 ± 1.05	0.35	11
16 BASAL_Inferior-Inferior	3.20 ± 1.35	3.75 ± 1.42	1.22	35	3.70 ± 1.54	3.38 ± 1.12	0.69	20
Global	3.55 ± 0.49	3.58 ± 0.49	0.50	14	3.51 ± 0.51	3.58 ± 0.50	0.37	11

abs, absolute value of reproducibility coefficient; %, percent of reproducibility coefficient.

Reproducibility

As mentioned previously, reconFA tended to be more reproducible as compared with sinoFA. This most probably reflects that the improved signal-to-noise ratio increased the validity of the analysis despite being faster.

The repeatability of PET O-15-labeled water resting MBF measurement has been reported in 5 publications to date. Four studies were conducted on human subjects,¹⁶⁻¹⁹ whereas the last one was an experimental animal study.²² Our method is different in several aspects. All of the previous studies investigated the agreement of repeat MBF, whereas our study has assessed purely intraobserver reproducibility of the same data sets. By examining the resting and stress MBF, 1 observer calculated MBF using different methods (reconFA and sinoFA). One month later, the same data sets were reanalyzed by the same observer. The reproducibility of the global MBF was excellent for both sinoFA and reconFA methods in this study. In addition, we performed segmental comparisons, as the LV was further subdivided into a total of 16 segments. In most previous studies the subsegments were grouped to obtain the mean value of global wall perfusion. By contrast, we provide segmental data, revealing a marked variation of reproducibility among different segments. This dispersion, however, was within the spatial heterogeneity of MBF as previously reported both within the same individual and

between different individuals.²³ Another study demonstrated a good reproducibility of 16-segmental resting MBF using an animal model.²² However, it did not include hyperemic flow conditions.

Limitations

MBF measurements in this study were not assessed independently from PET by another reference method because no other noninvasive method was available by which to quantitatively assess the myocardial perfusion. The radioactive microsphere method, as the gold standard for measurement of MBF, requires postmortem tissue samples and can only be used in animals, whereas for humans, PET represents the established gold standard for MBF measurement.²⁴ A further limitation of the study is that no CAD patients were included. However, as the maximal MBF is typically higher in healthy volunteers than in CAD patients, our study seems to appropriately cover the whole range of values.

Finally, correlation between sinoFA and reconFA revealed a markedly decreased correlation coefficient if rest and stress were analyzed separately as compared with pooled analysis. This, however, had little impact on the agreement of the 2 methods, as reflected by the limits of agreement (Table 3), which are very similar for resting (30%), hyperemic (21%), and pooled data (26%).

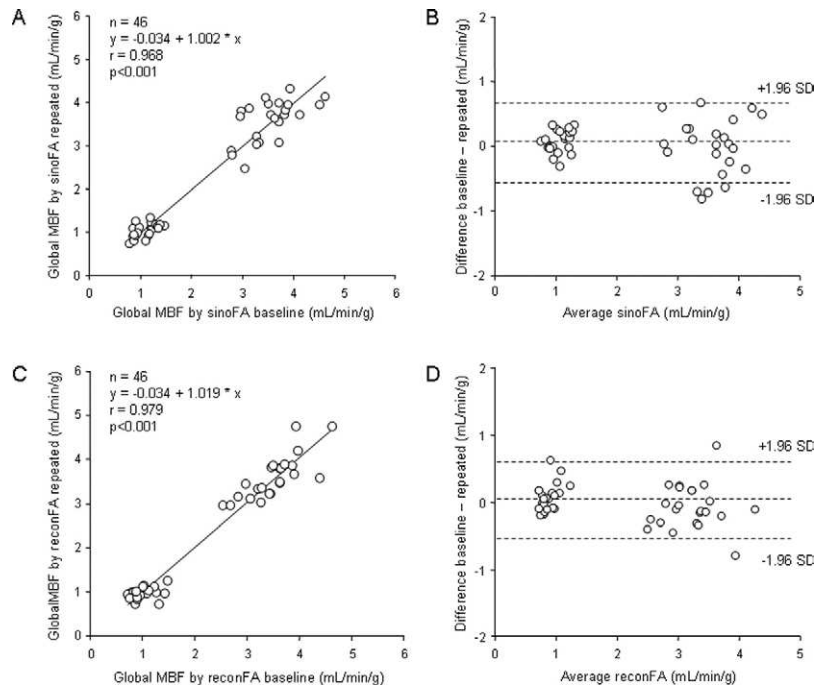


Figure 2. Linear regression analysis and Bland-Altman plot comparing baseline and repeat global MBF values documenting intraobserver reproducibility for sinoFA (A and B) and reconFA (C and D), where 1.96 SD denotes reproducibility coefficient.

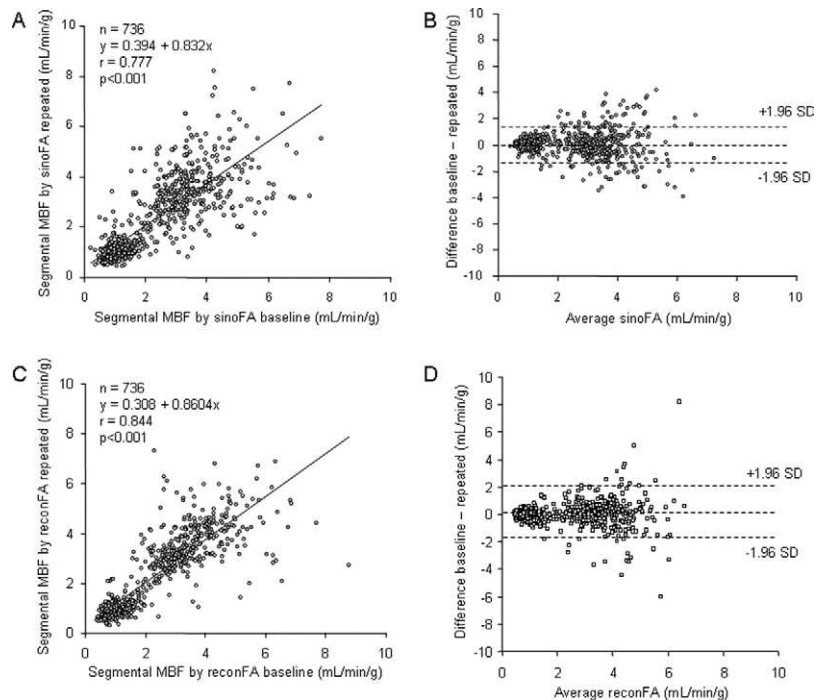


Figure 3. Linear regression analysis and Bland-Altman plot comparing baseline and repeat segmental MBF values documenting intraobserver reproducibility for sinoFA (A and B) and reconFA (C and D), where 1.96 SD denotes reproducibility coefficient.

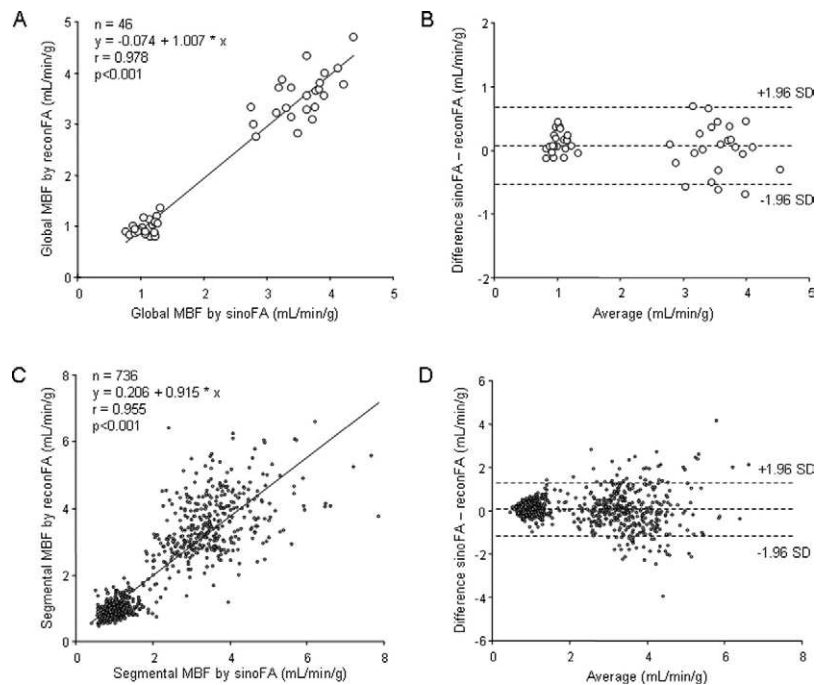


Figure 4. Linear regression analysis and Bland-Altman plot of global (A and B) and segmental (C and D) MBF for comparison between sinoFA and reconFA, where 1.96 SD denotes reproducibility coefficient.

Table 3. Comparison of global MBF: rest and stress separate and pooled

	Value ($\text{mL} \cdot \text{min}^{-1} \cdot \text{g}^{-1}$)		Correlation			BA agreement	
	sinoFA	reconFA	n	r	P	abs	%
Rest MBF	1.08 ± 0.16	0.97 ± 0.144	23	0.61	<.001	0.31	30
Stress MBF	3.56 ± 0.44	3.54 ± 0.47	23	0.56	<.001	0.76	21
Pooled MBF	2.32 ± 1.29	2.26 ± 1.34	46	0.97	<.001	0.58	26

abs, absolute value of reproducibility coefficient; %, percent of reproducibility coefficient.

CONCLUSIONS

Our results suggest that reconFA should be used for quantitative MBF measurement with O-15-labeled water and PET, as it considerably shortens analysis time and provides accurate values as compared with the validated sinoFA method. Its reliability is further substantiated by the excellent reproducibility, which is slightly superior to that for sinoFA.

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