Annals of Internal Medicine

Estimating Equations for Glomerular Filtration Rate in the Era of Creatinine Standardization

A Systematic Review

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Background: Clinical laboratories are increasingly reporting estimated glomerular filtration rate (GFR) by using serum creatinine assays traceable to a standard reference material.

Purpose: To review the performance of GFR estimating equations to inform the selection of a single equation by laboratories and the interpretation of estimated GFR by clinicians.

Data Sources: A systematic search of MEDLINE, without language restriction, between 1999 and 21 October 2011.

Study Selection: Cross-sectional studies in adults that compared the performance of 2 or more creatinine-based GFR estimating equations with a reference GFR measurement. Eligible equations were derived or reexpressed and validated by using creatinine measurements traceable to the standard reference material.

Data Extraction: Reviewers extracted data on study population characteristics, measured GFR, creatinine assay, and equation performance.

Data Synthesis: Eligible studies compared the MDRD (Modification of Diet in Renal Disease) Study and CKD-EPI (Chronic Kidney Disease Epidemiology Collaboration) equations or modifications

Estimates of glomerular filtration rate (GFR) from serum creatinine levels are now reported by more than 80% of clinical laboratories in the United States (1). Accurate estimation of GFR is important for detecting and staging chronic kidney disease (CKD), determining drug dosages, and stratifying risk (2). The equation from the MDRD (Modification of Diet in Renal Disease) Study is most frequently used but is known to be less accurate at higher GFRs and in racial and ethnic groups outside of North America, Europe, and Australia. Thus, researchers have developed and validated other GFR estimating equations to overcome these limitations.

The increasing use of GFR estimating equations has led to an appreciation of the effect of differences in creatinine assays on the accuracy of GFR estimates (3, 4). In 2006, a serum matrix standard reference material (SRM) was prepared by the National Institute of Standards and Technology and submitted to the Joint Committee for Traceability in Laboratory Medicine (5). Use of this material, in combination with the isotope-dilution mass spectrometry reference method, was intended to assist reagent manufacturers in achieving better consensus among methods (6). By the end of 2009, the calibration of most clinical laboratory methods was traceable to the SRM and isotopedilution mass spectrometry (1).

Our goal was to systematically review the performance of creatinine-based GFR estimating equations for use with

thereof. In 12 studies in North America, Europe, and Australia, the CKD-EPI equation performed better at higher GFRs (approximately >60 mL/min per 1.73 m²) and the MDRD Study equation performed better at lower GFRs. In 5 of 8 studies in Asia and Africa, the equations were modified to improve their performance by adding a coefficient derived in the local population or removing a coefficient.

Limitation: Methods of GFR measurement and study populations were heterogeneous.

Conclusion: Neither the CKD-EPI nor the MDRD Study equation is optimal for all populations and GFR ranges. Using a single equation for reporting requires a tradeoff to optimize performance at either higher or lower GFR ranges. A general practice and public health perspective favors the CKD-EPI equation.

Primary Funding Source: Kidney Disease: Improving Global Outcomes.

 Ann Intern Med. 2012;156:785-795.
 www.annals.org

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 This article was published at www.annals.org on 7 February 2012.

standardized serum creatinine measurements to inform selection of a single equation that laboratories could use to estimate GFR and to help clinicians interpret estimated GFR.

METHODS

Data Sources and Searches

We conducted a systematic search in MEDLINE from 1999 (the year before the 4-variable MDRD Study equation was published [7]) to 21 October 2011, with no language restrictions. Because the MDRD Study equation was the first equation to be reexpressed for use with standardized creatinine measurements (8), our search allowed us to identify other equations based on the MDRD Study equation that could also be reexpressed for use with standardized creatinine measurements. We also supplemented our

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Appendix Tables CME quiz (preview on page I-29) Conversion of graphics into slides

REVIEW | Estimating Equations for GFR in the Era of Creatinine Standardization

Context

Multiple methods are used to estimate glomerular filtration rate (GFR) from serum creatinine level.

Contribution

This review summarized data from cross-sectional studies that compared 2 or more creatinine-based GFR estimating equations to a reference GFR measurement. Studies from North America, Europe, and Australia showed that the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) equation performed better at higher GFRs and the Modification of Diet in Renal Disease (MDRD) Study equation performed better at lower GFRs. Neither equation performed as well in Asian or African populations as it did in North American or European populations.

Implication

The performance of the CKD-EPI and MDRD Study equations varies across populations and GFR ranges.

—The Editors

search with references from 2 previous reviews (9, 10). Appendix Table 1 (available at www.annals.org) lists our keywords and search strategy.

Study Selection

We included cross-sectional studies in any adult clinical or research population comparing GFR estimates from at least 2 creatinine-based estimating equations with a reference method of GFR measurement. Acceptable reference methods for measuring GFR in the development and validation populations were the urinary or plasma clearance of an exogenous filtration marker. We included equations that were originally developed by using standardized serum creatinine measurements or those for which the coefficients were subsequently recalculated for use with standardized creatinine measurements, which we refer to as reexpressed. We also included 1 study for which a conversion factor could be applied post hoc (11). We required evaluation of the equation in a data set that was external to the one in which it was developed, with independent sampling of the development and validation populations. Creatinine assays had to be traceable to the SRM (5). Acceptable assay methods included those standardized against isotope-dilution mass spectrometry of the reference material of the National Institute of Standards of Technology. Studies were also acceptable if they used a conversion factor derived from calibration of samples across the range of creatinine concentrations represented in the study population, with methods traceable to isotope-dilution mass spectrometry. We excluded studies that used an assay that was not traceable to the SRM and those with unclear traceability. In some cases, these studies were developed with assays and analytic material that are no longer available; the performance of the equations described in such studies was considered to be irrelevant to current practice. The minimum size was arbitrarily set at 100. We also looked for results of performance in subgroups by GFR, age, and race that had at least 50 members.

Data Extraction and Quality Assessment

We extracted data on study population characteristics, reference standards, measured GFRs, methods of creatinine calibration, and equation performance (bias, precision, and accuracy). We noted whether participants had conditions that could affect serum creatinine level through non-GFR determinants, such as conditions that alter creatinine generation (chronic illness or steroids), or medications that inhibit creatinine secretion (trimethoprim). Data from each article were systematically extracted by 1 of the authors. Other authors reviewed the methods section of studies describing development and evaluation of an equation to identify the creatinine assays and to confirm the study design and results. Our stringent selection criteria ensured that all reviewed studies were of good methodological quality.

Statistical Analysis

Summary measures of the differences between measured and estimated GFR were compared for each equation. No single metric captures all of the important information for evaluating performance of GFR estimating equations (10). In general, measures on the raw scale tend to emphasize errors at higher GFRs, whereas measures on the percentage or log scale tend to emphasize errors at lower GFRs (10). Bias, an expression of systemic error in estimated GFR, is defined as the median or mean of the differences between estimated and measured GFR. We used bias on the raw scale because it is easier to interpret. Precision is an expression of the random variation or "spread" of estimated GFR values around the measured GFR. The root mean square error of the regression of estimated GFR versus measured GFR, or log estimated GFR versus log measured GFR, is considered to be a direct measure of precision. Indirect measures, such as interquartile range or SD for the differences between estimated GFR and measured GFR, were included if the root mean square error was not provided. Accuracy is affected by both bias and imprecision and was expressed as the percentage of estimated GFR values within 30% of measured GFR (P_{30}). This measure of P₃₀ was first reported as a measure of the accuracy of the MDRD Study equation (12) and was subsequently recommended by the National Kidney Foundation Disease Outcomes Quality Initiative in its CKD guidelines (13). An error of 30% is considered large; for example, a 30% error in a patient with measured GFR of 50 mL/min per 1.73 m^2 could lead to an estimated GFR as low as 35 or as high as 65 mL/min per 1.73 m². Some have suggested that 20% would be a more appropriate threshold for a large error. We have included alternative values for accuracy (such as P10, P15, or P20) if they were reported. For our conclusions, we focused primarily on P₃₀ because it is a measure of large errors that would be important to

Figure 1. Summary of evidence search and selection.

clinicians, is less influenced by small bias due to differences in measurement methods or regression to the mean, and was most consistently reported across studies.

We tabulated results separately for adult populations of largely Northern European ancestry (North American, European, and Australian populations) and those from other locations and separated the results for subgroups by GFR range, race or ethnicity, and age. Within each table, we organized the studies first by similar reference measurement method and then by size.

Role of the Funding Source

The authors are members of the evidence review team and 2 workgroup experts of the ongoing Kidney Disease: Improving Global Outcomes (KDIGO) guideline on evaluation and management of CKD. The evidence review team was supported by KDIGO to conduct systematic reviews and provide methods support. The judgments and interpretations in the article are those of the authors. The funding source did not participate in the design, conduct, or reporting of the study.

RESULTS

Our search yielded 3250 abstracts; of these, 100 articles were reviewed in full text and 23 met our inclusion criteria (Figure 1). The main reason for exclusion was that the equations had not been developed or reexpressed for use with creatinine assays traceable to the SRM. Appendix Table 2 lists equations that met our criteria, and Appendix Table 3 (both available at www.annals.org) lists the equations that were not traceable to the SRM.

Estimating Equations Developed in Adult Populations in North America, Europe, or Australia

Twelve studies, comprising 12 898 patients, met our criteria (Table 1) (14–26). Study populations were the general population in 3 studies (14–17), kidney transplant recipients in 3 studies (18–20), individuals before kidney donation in 1 study (21) and before and after kidney donation in another study (22), patients with cancer in 1 study (23), and a heterogeneous population in 3 studies (24–26). All studies compared the CKD-EPI (Chronic Kidney Disease Epidemiology Collaboration) equation with the MDRD Study equation. Methods of measuring GFR included clearance of iothalamate in 5 studies, technetium–diethylenetriamine pentacetic acid (Tc-DTPA) in 3 studies, inulin in 1 study, iohexol in 1 study, chromium–ethylenediamine tetraacetic acid (Cr-EDTA) in 1 study, and various markers in 1 study.

Across the 12 studies, P_{30} ranged from 59% to 95%. The CKD-EPI equation was more accurate than the MDRD Study equation in 10 studies and less accurate in 2 studies. Bias ranged from 14.6 to -22 mL/min per 1.73 m². The CKD-EPI equation was less biased than the MDRD Study equation in 7 studies and more biased in 5 studies. Two studies reported the root mean square error as a mea-



SCr = serum creatinine; SRM = standard reference material.

sure of precision. In 6 of the 10 studies that reported a measure of precision, the CKD-EPI equation was more precise than the MDRD Study equation; precision for the MDRD Study equation was better or the same in the other 4 studies. In 5 studies (14–17, 25, 26), performance measures were consistently better for the CKD-EPI equation than for the MDRD Study equation. In 2 studies (18, 19), both of which were conducted in kidney transplant recipients, performance measures were better for the MDRD Study equation. In 1 study (18), all patients received trimethoprim.

Figure 2 shows differences in accuracy and bias according to measured GFR and GFR measurement method. The CKD-EPI equation seems to be more accurate and less biased in studies with higher mean measured GFRs (approximately >60 mL/min per 1.73 m^2), whereas the MDRD Study equation has greater accuracy and less bias at lower GFRs. No clear pattern by GFR measurement method was observed. Appendix Table 4 (available at www .annals.org) shows subgroups stratified by GFR or clinical characteristics in studies in which these subgroup data were reported (19, 20, 25, 26). Within each study, the differences in accuracy and bias were larger at higher GFRs and smaller at lower GFRs.

One study (27) compared a modification of the CKD-EPI equation that used a 4-level race or ethnicity coeffi-

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Table 1. Performance Comparison of Creatinine-Based GFR Estimating Equations in North America, Europe, and Australia

Study, Year (Reference)	Country	Population	Patients, n	Me	asured GFR
(Reference)				Reference Standard	Value (SD), <i>mL/min per</i> 1.73 m ²
Murata et al, 2011 (26)	United States	Adults who underwent an outpatient iothalamate clearance at the Mayo Clinic in Rochester, Minnesota (women, 45%; African American, 2%; mean age, 56 y; donor or potential donor, 13%; KTRs, 26%)	5238	¹²⁵ I-Iothalamate (urine)	55.9 (29.7)
Levey et al, 2009 (25)	United States	External validation set comprising 16 studies (women, 45%; white or other, 87%; black, 10%; Hispanic, 2%; Asian, 2%; mean age, 50 y; diabetes, 28%; donor, 16%; KTRs, 29%)	3896	¹²⁵ I-Iothalamate (urine) and others	68 (36)
Lane et al, 2010 (21)	United States	Patients before and after nephrectomy for causes other than donation at the Cleveland Clinic (women, 93%; white, 91%; black, 7%; median age, 58 y)	425	¹²⁵ I-Iothalamate (urine)	50 (IQR, 29 to 69)
Michels et al, 2010 (17)	The Netherlands	Potential kidney donors and adults who underwent GFR measurement for clinical reasons at the Academic Medical Center in Amsterdam (women, 56%; black, 12%; mean age, 44 y)	271	¹²⁵ I-Iothalamate (urine)	78.2 mL/min (33.4)
Tent et al, 2010 (22)	The Netherlands	Living kidney donors who donated from 1996–2007 (women, 57%; white, 100%; mean age, 50 y)	253 before donation 253 after donation	¹²⁵ I-Iothalamate (urine)	115 mL/min (20) 73 mL/min (13)
Kukla et al, 2010 (18)	United States	KTRs receiving immunosuppression (women, 40%; white, 86%; mean age, 49 y; receiving trimethoprim, 100%)	107 on steroid-free immunosuppression early posttransplantation 81 on steroid-free immunosuppression at 1 y	¹²⁵ I-Iothalamate (urine)	55.5 (17.0) 56.8 (17.7)
White et al, 2010 (20)	Canada	Stable KTRs (women, 36%; white, 92%; receiving trimethoprim, 19%)	207	^{99m} Tc-DTPA (plasma)	58 (22)
Pöge et al, 2011 (19)	Germany	Patients with stable renal function after transplantation (women, 40%; white, 99%; mean age, 49 y)	170	^{99m} Tc-DTPA (plasma)	39.6 (IQR, 11.8 to 82.9)
Jones and Imam, 2009 (15), and Jones, 2010 (16)	Australia	Australian patients referred for routine GFR measurements (women, 43%; mean age, 61 y)	169	^{99m} Tc-DTPA (plasma)	75 (IQR, 5 to 150)
Cirillo et al, 2010 (14)	Italy	White adults with and without kidney disease (with kidney disease, 49%; women, 41%; mean age, 47 y; diabetic nephropathy, 26 cases; glomerulonephritis, 40 cases; PKD, 15 cases)	356	Inulin (plasma)	71.5 (36.3)
Eriksen et al, 2010 (24)	Norway	Patients participating in the 6th Tromsø population survey who had no previous myocardial infarction, angina, stroke, diabetes, or renal disease (women, 51%; mean age, 57 y)	1621	Iohexol (plasma)	91.7 (14.4)
Redal-Baigorri et al, 2011 (23)	Denmark	Patients with cancer who were referred for determination of GFR before chemo- therapy (women, 57%; mean age, 62 y)	185	⁵¹ Cr-EDTA (plasma)	85.1 (20.3)

CKD-EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; IDMS = isotope-dilution mass spectrometry; IQR = interquartile range; KTR = kidney transplant recipient; MDRD = Modification of Diet in Renal Disease; ND = not documented; NIST = National Institute of Standards andTechnology; P₃₀ = percentage of estimated GFR values within 30% of measured GFR; PKD = polycystic kidney disease; SCr = serum creatinine.

* Computed as estimated GFR minus measured GFR. Positive numbers indicate overestimation and negative numbers indicate underestimation of measured GFR. Smaller absolute values indicate lesser bias.

† Lower values indicate greater precision.

¹ Higher values indicate greater precision. ² Higher values indicate greater accuracy. Among the 3 studies (14, 18, 19) that reported alternative measures of accuracy, results were consistent with P_{30} in all. In addition to P_{30} , references 14, 18, and 19 reported P_{10} ; reference 14 also reported P_{20} . § Evaluated as the root mean square error for the regression of estimated GFR on measured GFR.

|| Evaluated as the IQR for the differences between estimated and measured GFR.

¶ Evaluated as the SD of the differences between estimated and measured GFR. ** Converted to raw scale by multiplying percentage of bias by measured GFR.

Table 1—Continued

GFR Estimation			Results	
SCr Calibration and Assay	Equation	Bias (95% Cl), <i>mL/min</i> per 1.73 m ² *	Precision (95% CI)†	P ₃₀ (95% CI), %‡
Roche Jaffe assay (Roche P- or D-Modular or Roche Cobas C501 with Roche Creatininase Plus assay; Roche Diagnostics, Indianapolis, Indiana) with demonstrated IDMS alignment	MDRD CKD-EPI	-4.1 -0.7	ND	77.6 78.4
Roche enzymatic assay (Roche–Hitachi P-Module instrument with Roche Creatininase Plus assay; Hoffman-La Roche, Basel, Switzerland) recalibrated to standardized SCr at the Cleveland Clinic	MDRD CKD-EPI	-5.5 (-5.0 to -5.9) -2.5 (-2.1 to -2.9)	0.274 (0.265 to 0.283)§ 0.250 (0.241 to 0.259)§	80.6 (79.5 to 82.0) 84.1 (83.0 to 85.3)
Measured at the Cleveland Clinic; assay standardized against NIST	MDRD CKD-EPI	-1.0 -1.7	15.0 13.8	75 80
Hitachi enzymatic assay (Hitachi H911; Boehringer Mannheim, Mannheim, Germany) validated against IDMS	MDRD CKD-EPI	14.6 mL/min 12.3 mL/min	19.9¶ 12.1¶	81.2 84.5
Roche enzymatic assay or Jaffe assay on MEGA analyzer (Merck, Darmstadt, Germany); both methods calibrated to reference standard at the Cleveland Clinic	MDRD CKD-EPI MDRD CKD-EPI	-22 mL/min (20 to 25) -14 mL/min (11 to 16) -15 mL/min (14 to 16) -11 mL/min (9 to 11)	20 (14 to 26) 18 (14 to 22) 12 (9 to 15) 12 (10 to 16)	73 (68 to 79) 89 (85 to 93) 71 (65 to 76) 89 (85 to 93)
Measured by using a Jaffe CXR Synchron method and, later, an IDMS-traceable assay; Jaffe assay-based SCr converted to IDMS-traceable values	MDRD CKD-EPI MDRD CKD-EPI	8.23 13.30 2.40 6.91	17.9§ 21.1§ 15.8§ 17.3§	71.7 58.5 75.0 66.7
For the reexpressed MDRD Study and CKD-EPI equations, SCr adjusted to IDMS standard	MDRD CKD-EPI	-7.4 -5.2	14.4 15.7	79 (73 to 84) 84 (78 to 88)
Jaffe assay on Dimension RxLTM analyzer (Dade Behring, Marburg, Germany); assay adjusted for calibration with the IDMS method	MDRD CKD-EPI	4.49 8.07	10.0¶ 10.9¶	71.8 64.1
Roche Jaffe assay (Roche, Australia) with demonstrated IDMS alignment	MDRD CKD-EPI	-3** -1.5**	ND	81 86
Kinetic Jaffe assay (Bayer Express Plus; Siemens, Munich, Germany) standardized to NIST	MDRD CKD-EPI	-5.2 -0.9	14.9¶ 13.2¶	87.4 88.2
Hitachi enzymatic method (CREA Plus; Roche Diagnostics, Mannheim, Germany) standardized against IDMS	MDRD CKD-EPI	1.3 (0.4 to 2.1) 2.9 (2.2 to 3.5)	18.2 (17.2 to 19.5)∥ 15.4 (14.5 to 16.3)∥	93 (91 to 94) 95 (94 to 96)
Jaffe (Abbott Architect C systems 8000, reagent 7D64; Abbott Laboratories, Abbott Park, Illinois) method standardized to IDMS	MDRD CKD-EPI	0.81 (IQR, −1.56 to 3.19) 1.16 (IQR, −0.76 to 3.09)	16.49¶ 13.37¶	88.6 89.7

cient (black, Asian, and Native American or Hispanic vs. white and other) with a modification that used a 2-level coefficient (black vs. white and other) in racial or ethnic groups in North America and Europe (**Appendix Table 5**, available at www.annals.org). The development data set had few Asian or Native American or Hispanic patients, and the improvements in bias and precision in these groups were not clinically meaningful. Only 2 studies (26, 28) provided results by age subgroups. In both studies, the

differences in bias between equations were generally smaller in subgroups of older patients (data not shown).

Estimating Equations Developed in Adult Populations Outside North America, Europe, or Australia

Eight studies met our criteria (**Table 2**) (29–34). The study populations were from Asia and Africa, including Japanese, Chinese, Thai, Korean, South African black, or multiethnic Singapore populations and included general

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Study, Year (Reference)	Country	Population	Patients, n	Me	asured GFR
(Reference)				Reference Standard	Value (SD), <i>mL/min</i> per 1.73 m ²
Stevens et al, 2011 (27)	China, Japan, and South Africa	Populations from 3 studies in China, Japan, and South Africa (women, 48%; Asian, 90%; black, 10%; mean age, 49 y; diabetes, 6%;	99 black patients (South Africa)	¹²⁵ I-Iothalamate (urine) and other filtration	61 (32)
		KTRs, 0%)	248 Asian patients (Japan)	markers	55 (35)
			675 Asian patients (China)		53 (31)
Matsuo et al, 2009 (34)	Japan	Hospitalized Japanese patients; external validation set (women, 42%; mean age, 54 y; diabetes, 44 cases; donors, 10; KTRs, 2; glomerulo- nephritis, 176 cases; PKD, 0 cases; lupus, 3 cases)	350	Inulin (urine)	57.2 (34.7) (GFR ≥60, 41%; GFR <60, 59%)
Horio et al, 2010 (29)	Japan	Japanese patients (women, 42%; mean age, 54 y; diabetes, 22%; donors, 3%; KTRs, 1%; hypertension, 58%)	350	Inulin (urine)	45 (25)
Yeo et al, 2010 (33)	Korea	Korean KTRs in the early postoperative period (women, 43%; mean age, 42 y; diabetes, 16%)	102	⁵¹ Cr-EDTA (plasma)	76.8 (17.0)
van Deventer et al, 2008 (30)	South Africa	Black South African patients (women, 49%; median age, 47 y; diabetes, 25%; donors, 7%; HIV, 20%; hypertension, 36%)	100	⁵¹ Cr-EDTA (plasma)	61.5 (49.6)
Teo et al, 2011 (32)	Singapore	Outpatients with CKD in the nephrology clinic at National University Hospital, Singapore (women, 48%; Chinese, 41%; Malay, 31%; Indian or other, 28%; median age, 58 y; diabetes, 23%; hypertension, 50%)	232	⁹⁹ Tc-DTPA (plasma)	51.7 (27.5) (GFR ≥60, 69%; GFR <60, 31%)
Ma et al, 2006 (11)	China	Patients with CKD from 9 renal institutes at university hospitals located in 9 geographic regions of China	230	⁹⁹ Tc-DTPA (plasma)	ND
Praditpornsilpa et al, 2011 (31)	Thailand	Thai patients with CKD who were in stable condition	100	⁹⁹ Tc-DTPA (plasma)	51.1 (28.4)

Table 2. Performance Comparison of Creatinine-Based GFR Estimating Equations Outside of North America, Europe, and Australia

CKD = chronic kidney disease; CKD-EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; IDMS = isotope-dilution mass spectrometry; JSN-CKDI = Japanese Society of Nephrology-Chronic Kidney Disease Initiatives; KTR = kidney transplant recipient; MDRD = Modification of Diet in Renal Disease; ND = not documented; NIST = National Institute of Standards and Technology; P₃₀ = percentage of estimated GFR values within 30% of measured GFR; PKD = polycystic kidney disease; SCr = serum creatinine; SRM = standard reference material.

+ Lower values indicate greater precision.

 \ddagger Higher values indicate greater accuracy. Among the 5 studies (11, 31–34) that reported alternative measures of accuracy, results were consistent with P₃₀ in 3. Of the studies reporting results consistent with P₃₀, reference 31 reported P₁₀ and P₁₅, reference 33 reported P₁₀, and reference 34 reported P₁₅. References 11 and 32 reported inconsistent results between P₁₅ and P₃₀.

§ Evaluated as the interquartile range for the differences between estimated and measured GFR.

|| Evaluated as the root mean square error for the regression of estimated GFR on measured GFR.

¶ Evaluated as the SD of the differences between estimated and measured GFR.

** African American coefficient of the MDRD equation.

++ Derived by using the original MDRD equation (with the 186 coefficient). For use with an SCr traceable to the SRM, the 186 coefficient should be replaced with the 175 coefficient from the reexpressed MDRD equation.

 \pm Derived by using the original MDRD equation (with the 186 coefficient). For use with an SCr traceable to the SRM, the SCr should be replaced by SCr \times 0.95, which represents the calibration factor relating the SCr assay in the Cleveland Clinic laboratory to the standardized SCr assay.

^{*} Computed as estimated GFR minus measured GFR. Positive numbers indicate overestimation and negative numbers indicate underestimation of measured GFR. Smaller absolute values indicate lesser bias.

Table 2—Continued

GFR Es	timation		Results	
SCr Calibration and Assay	Equation	Bias (95% Cl), <i>mL/min</i> per 1.73 m ² *	Precision (95% CI)†	P ₃₀ (95% CI), %‡
Calibrated to standardized SCr measurements by using Roche	CKD-EPI equation with 2-level race or ethnicity coefficient	12.4 (7.6 to 18.3)	0.326 (0.292 to 0.361)	55.6 (46.5 to 64.6)
(Roche–Hitachi P-Module instrument with Roche Creatininase Plus assay;	CKD-EPI equation with 4-level race or ethnicity coefficient	12.5 (7.6 to 18.4)	0.327 (0.292 to 0.362)	55.6 (46.5 to 64.6)
Hoffmann-La Roche, Basel, Switzerland) enzymatic assay at the	CKD-EPI equation with 2-level race or ethnicity coefficient	17.8 (14.7 to 20.1)	0.469 (0.424 to 0.515)	29.4 (23.8 to 35.1)
Cleveland Clinic	CKD-EPI equation with 4-level race or ethnicity coefficient	21.4 (18.2 to 23.3)	0.507 (0.463 to 0.553)	36.3 (30.6 to 42.3)
	CKD-EPI equation with 2-level race or ethnicity coefficient	-2.7 (-3.7 to -1.9)	0.325 (0.302 to 0.348)	73.2 (69.9 to 76.6)
	CKD-EPI equation with 4-level race or ethnicity coefficient	-1.3 (-2.2 to -0.6)	0.318 (0.295 to 0.343)	72.1 (68.7 to 75.7)
Hitachi enzymatic assay (Hitachi, Tokyo,	MDRD	12.0	25.2	59 (54 to 64)
Japan) with excellent agreement	Japanese-modified MDRD (equation 1)	-5.9	19.9	72 (67 to 76)
against the Cleveland Clinic methods	JSN-CKDI equation (equation 2)	-7.9	20.3	73 (69 to 78)
-	MDRD with Japanese coefficient (equation 3)	-1.3	19.4	73 (59 to 78)
	3-variable Japanese equation (equation 4)	-2.1	19.1	75 (70 to 79)
	5 variable supariese equation (equation 4)	2.1	13.1	/ 5 (/ 6 (6 / 5))
Enzymatic assay validated by using the	MDRD with Japanese coefficient	-1.3	19.4	73 (69 to 78)
calibration panel of the Cleveland Clinic	CKD-EPI with Japanese coefficient	-0.4	17.8	75 (70 to 79)
Rate-blanked, Toshiba-compensated,	MDRD	-0.33	12.57¶	94.1
kinetic Jaffe assay (Toshiba Medical Systems, Tokyo, Japan) using a Roche calibrator (Roche Diagnostics, Indianapolis, Indiana) traceable to the IDMS reference method	Japanese-modified MDRD	17.95	11.06¶	68.6
Rate-blanked, Roche-compensated, kinetic Jaffe assay (Roche Modular analyzer; Roche Diagnostics, Indianapolis, Indiana) with calibration traceable to IDMS, compared with Roche enzymatic assay (Creatinine Plus; Roche Diagnostics, Indianapolis, Indiana) at the Cleveland Clinic with values adjusted by regression	MDRD with race or ethnicity factor** MDRD without race or ethnicity factor**	13.1 (5.5 to 18.3) 1.9 (−0.8 to 4.5)	28.5 16.6	52 74
Siemens enzymatic assay (Siemens Advia	MDRD	-3.0 (-4.2 to -1.7)	12.2 (10.0 to 14.4)§	79.7 (74.6 to 84.9)
2400; Siemens, Munich, Germany) calibrated with manufacturer-provided materials traceable to standardized creatinine (NIST SRM 967) measured by using IDMS	CKD-EPI	-1.2 (-2.7 to 0.3)	12.1 (9.0 to 15.1)§	82.8 (77.9 to 87.6)
Hitachi kinetic Jaffe assay with values adjusted by regression analysis to	Original MDRD equation, calibrated to the Cleveland Clinic creatinine measure-	-7.8 (-21.5 to -1.8)	ND	66.1
Cleveland Clinic Beckman CX3 assay (Beckman Coulter, Fullerton, California)	ments (study equation 2)++ Original MDRD equation, calibrated to Cleveland Clinic creatinine measure- ments, with Chinese coefficient (study equation 4)++	-0.9 (-9.6 to 7.4)		77.8
	Chinese equation (study equation 6)‡‡	-0.8 (-9.7 to 7.4)		79.6
Roche enzymatic assay (Roche	MDRD	-11.9	8.8¶	62.7
Diagnostics, Indianapolis, Indiana) with	CKD-EPI	-10.9	7.8¶	68.0
values adjusted to IDMS reference serum SRM 967	MDRD equation with Thai racial factor correction	-10.3	8.5¶	73.3

population in 3 studies (27, 29, 30), patients with CKD in 3 studies (11, 31, 32), kidney transplant recipients in 1 study (33), and a heterogeneous population in 1 study (34). All studies examined the MDRD Study and CKD-EPI equations or modifications thereof. In 6 studies (11, 28–31, 34), the MDRD Study or CKD-EPI equations were modified by adding or removing a coefficient to improve the performance of the equation in the development data set or a new equation was developed by using the same variables (**Appendix Table 2**). Methods for measuring GFR included Tc-DTPA in 3 studies, inulin in 2 studies, Cr-EDTA in 2 studies, and iothalamate in 1 study.

Comparing the performance of equations across studies is limited because locally derived equations were generally not tested in other studies or populations. In these studies, the unmodified MDRD Study and CKD-EPI equations were less accurate (P_{30} ranging from 29% to 94%) than in the studies of populations in North America, Europe, and Australia. In 5 studies, adding (11, 29, 31, 34) or removing (30) a coefficient improved the accuracy of

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Difference in accuracy (*top*), as measured by P_{30} (P_{30} for CKD-EPI minus P_{30} for MDRD), is plotted against mean measured GFR in the study population. Difference in bias (*bottom*) (absolute value for bias for estimated GFR by MDRD Study equation minus absolute value for bias for estimated GFR by CKD-EPI equation) is plotted against mean measured GFR in the study population. CKD-EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; MDRD = Modification of Diet in Renal Disease; P_{30} = percentage of estimated GFR values within 30% of measured GFR.

* Denotes study in which all patients received trimethoprim.

+ Could be reported in mL/min.

GFR estimation (11, 29–31, 34). However, in 1 study (27), modifying the CKD-EPI equation by substituting a 4-level race or ethnicity coefficient derived in a North American and European population for the 2-level coefficient did not significantly improve accuracy in Chinese, Japanese, or South African black populations. Coefficients developed in 1 study of an ethnic or racial population did not improve equation accuracy in a study of another ethnic or racial population (30, 33). The Japanese and Chinese coefficients also vary widely (Appendix Table 2) (29, 34). In 3 studies that compared the CKD-EPI and MDRD Study equations (29, 31, 32) (with or without modification), the CKD-EPI equation was more accurate.

Three studies examined the performance of estimating equations by GFR strata (Appendix Table 6, available at www.annals.org) (29, 30, 32). In 2 studies, the differences

in accuracy and bias were larger at higher GFRs and smaller at lower GFRs. In 1 study (29), the differences in bias between equations were smaller in older subgroups than in younger subgroups (data not shown).

DISCUSSION

Only the MDRD Study and CKD-EPI equations and modifications thereof have been expressed for use with creatinine assays traceable to the SRM and concurrently compared with measured GFR in adults. In studies of North American, European, or Australian populations, the CKD-EPI equation was more accurate (had a higher P_{30}) than the MDRD Study equation in 10 of 12 studies. The comparison of bias was more variable. For both accuracy and bias, the CKD-EPI equation performed better at higher GFRs (approximately >60 mL/min per 1.73 m²) and the MDRD Study equation performed better at lower GFRs. Within studies, the differences in bias were greater at higher GFRs than at lower GFRs.

Data on adults outside of North America, Europe, and Australia are more limited. Neither the MDRD Study nor the CKD-EPI equation performs as well in these locations as it does in North America and Europe. Equation performance can be improved by deriving local "race/ethnicity" coefficients; however, the modified or new equations generally do not exhibit the same level of accuracy as the CKD-EPI or MDRD Study equations do in North American, European, or Australian populations. The coefficients also do not seem to be generalizable beyond the local population, possibly because of differences in GFR measurement methods or differences in populations in addition to race or ethnicity (35).

Despite estimating GFR from the same variables (age, sex, race, and serum creatinine level), the CKD-EPI equation generally yields higher values for estimated GFR than does the MDRD Study equation. The difference in equation performance can be explained by differences in the development populations. The MDRD Study equation was developed in a study population with CKD and a mean GFR of 40 mL/min per 1.73 m², whereas the CKD-EPI equation was developed in a more diverse study population, including participants with and without CKD, with a mean GFR of 68 mL/min per 1.73 m². Our observation of differences in equation performance in various GFR ranges probably reflects differences in non-GFR determinants of creatinine (in particular, creatinine generation due to muscle mass and diet) and regression to the mean. Our observation of larger differences in estimated GFR between the equations at higher GFRs probably reflects equation development on the logarithmic scale.

Other differences in equation performance can be explained by differences between the development and the validation populations. The CKD-EPI and MDRD Study equations were developed in North American and European populations that mainly consisted of black and white

persons. Our observation that both equations perform less well in other racial and ethnic groups is consistent with known racial and ethnic differences in muscle mass and diet (35). No participants were receiving trimethoprim in either development population. Our observation of marked underestimation of GFR with both equations in persons receiving trimethoprim is consistent with the known inhibition of creatinine secretion by this drug (36).

Both the MDRD Study and CKD-EPI equations were developed by using iothalamate clearance as the reference standard for GFR. The validation studies used various filtration markers, with small differences in clearance compared with iothalamate (37). Use of filtration markers other than iothalamate in validation studies of these equations would introduce a systematic bias. The observation of variation in differences in equation performance among studies with similar mean GFRs may be due in part to differences in GFR measurement method.

The observed and expected differences in performance by range of GFR suggest that we cannot optimize the performance of any equation for all clinical populations across a wide range of GFRs. Because the goal is to select a single estimating equation for routine use by clinical laboratories, the tradeoff of optimizing performance at either higher or lower GFR ranges must be accepted. Because the difference in bias (on the raw scale) between the equations is greater at higher GFRs, using the CKD-EPI equation would lead to smaller average bias in clinical populations with a wide range of GFRs.

As in diagnostic test evaluation, the magnitude of differences in performance between the CKD-EPI and MDRD Study equations are best appreciated by considering the implications for clinical practice. Reporting estimated GFR by using the MDRD Study equation is widespread, so a change in the estimating equation used by clinical laboratories would have profound implications. Because the differences between the equations are greater at higher GFRs, the implications of introducing the CKD-EPI equation would be larger for public health and general clinical practice than for nephrology practices. Applying the CKD-EPI rather than the MDRD Study equation to the U.S. adult population would lead to a higher average estimated GFR, less sensitivity but more specificity for detecting a GFR less than 60 mL/min per 1.73 m², and a lower prevalence estimate but a higher risk profile for persons with an estimated GFR in this range (25, 38-40). This would potentially enable more efficient use of resources in caring for patients with decreased estimated GFR. Another consequence of using the CKD-EPI equation would be to allow reporting of estimated GFR as a numerical value throughout the full range, rather than limit it to lower values (for example, <60 mL/min per 1.73 m², as currently recommended for the MDRD Study equation in the United States) (1). However, using the CKD-EPI equation would slightly increase bias (overestimation) at lower GFRs. Nephrologists and others caring for patients with low GFR would need to be aware of this limitation.

Estimation methods need to be further improved. Even in North America, Europe, and Australia, the CKD-EPI equation does not meet the 2002 KDOQI benchmark of P_{30} greater than 90% (13). This level of performance may be beyond expectation for creatinine-based equations. Standardization has reduced but not eliminated bias due to differences in creatinine assays, and differences among GFR measurement methods remain another source of bias. Imprecision in estimating equations is quantitatively more important, probably because of variation in non-GFR determinants of serum creatinine level and imprecision in measured GFR. The former reflects the large number of factors affecting creatinine generation and reducing imprecision may require including additional variables, such as measures of muscle mass or diet; the latter reflects the fallibility of the reference standard rather than errors in the estimating equations and can be improved only by more precise measures of GFR. Although urinary clearance of inulin is the gold standard, it is difficult to use, and studies are needed to compare the bias and precision of alternative methods to enable calibration of GFR measurement methods. In addition, statistical adjustment for measurement error may allow more accurate evaluation of estimated GFR (41).

Outside North America, Europe, and Australia, bias remains an issue because of systematic variation in differences in creatinine generation. Incorporating locally derived coefficients can minimize this bias. However, the lack of generalizability of these coefficients probably reflects the contributions of multiple factors that differ across populations in various locations. Developing or reexpressing existing equations for local populations requires significant resources and may not be feasible in all locations.

The accuracy and worldwide generalizability of GFR estimating equations might be improved by using such alternative filtration markers as cystatin C, which is less dependent on muscle mass. However, all filtration markers have non-GFR determinants, so it is unlikely that imprecision will be eliminated by any single marker. Using multiple markers can improve precision by minimizing the contribution of any 1 non-GFR determinant (42). In June 2010, the Institute for Reference Materials and Measurements released a reference material for cystatin C measurement (43–45). Reagent manufacturers are in the process of recalibrating their assays against this standard, and estimating equations for use with standardized cystatin C measurements are being evaluated (46).

A strength of our review is that the stringent selection criteria eliminated older and smaller studies. This generated an evidence base of good-quality studies and makes the conclusions directly relevant to the current era of creatinine assays. Our review also highlights the shortcomings of the existing literature. We propose criteria for studies that are developing and validating GFR estimating equa-

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Table 3. Suggested Criteria for Developing and Validating GFR Estimating Equations

- The equation should be developed and validated in separate populations. Both populations should be representative of the population of intended use. A study should have sufficient sample size to detect a substantial improvement in equation performance, which we suggest is a relative reduction in bias of 50% or RMSE of 20% in relevant age and GFR strata. These values correspond to the improvements found in performance of the CKD-EPI over the MDRD Study equation.
- The measured GFR should be based on urine or plasma clearance of an exogenous filtration marker. The clearance method and filtration marker assay, including typical performance characteristics, should be stated. The accuracy of the GFR measurement procedure compared with urinary inulin clearance should be documented if not already known.
- The serum creatinine assay used in both the development and validation data sets should be standardized against SRM. The method, reagent manufacturer, and analytic platform should be stated together with typical performance characteristics of the assay (such as imprecision). When available, traceability data should be provided from robust crossover studies against the reference method. These same recommendations apply to other filtration markers, including cystatin C.
- Measures of equation performance should include bias, precision, and accuracy. Accuracy is probably the best single measure for comparing equations because it incorporates bias and precision. Studies should report P_{30} and P_{20} . For precision, RMSE should be reported. However, the best measure may vary depending on the particular population and study hypotheses.
- Validation should include comparison of a newly developed estimating equation with ≥1 previously developed equation in the population of interest to identify which equation is superior. To facilitate comparison of equations across studies in heterogeneous populations, we suggest including a common comparator. We currently suggest comparing the performance of any new or adapted equations with that of the unmodified CKD-EPI equation.

CKD-EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; MDRD = Modification of Diet in Renal Disease; RMSE = root mean square error; SRM = standard reference material.

tions (Table 3) to enhance the quality and comparability of future studies.

Our review has limitations. It was restricted to studies that compared at least 2 equations. Although this omits studies that developed and validated a single GFR estimating equation, it probably did not alter our conclusions because such equations are unlikely to have been developed in large populations and tested as widely as the equations that we included. We may also have inadvertently excluded studies that did use creatinine assays standardized against SRM. This emphasizes the importance of our recommendation that investigators provide full information about their methods.

In summary, neither the CKD-EPI nor the MDRD Study equation is optimal across all populations and GFR ranges. Using a single equation for reporting estimated GFR requires a tradeoff to optimize performance at either higher or lower GFR ranges. A general practice and public health perspective favors adopting the CKD-EPI equation in North America, Europe, and Australia and using it as a comparator for new equations in all locations. Whether the precision of creatinine-based equations can be substantially improved without adding other variables is uncertain. Equations using other filtration markers instead of or in addition to creatinine hold promise in this regard. Going forward, new equations should be rigorously developed, validated, and reported.

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Grant Support: By KDIGO.

Potential Conflicts of Interest: Ms. Earley and Drs. Miskulin and Uhlig report the following: Grant (money to institution): National Kidney Foundation; Support for travel to meetings for the study or other purposes: National Kidney Foundation. Ms. Earley and Dr. Uhlig further report: Fees for participation in review activities such as data monitoring boards, statistical analysis, end point committees, and the like (money to institution): National Kidney Foundation. Dr. Miskulin further reports: Grant: National Kidney Foundation; Employment: Dialysis Clinic. Dr. Lamb: Support for travel to meetings for the study or other purposes: Kidney Disease: Improving Global Outcomes; Grants/grants pending (money to institution): National Institutes of Health. Dr. Levey: Support for travel to meetings for the study or other purposes (money to institution): National Kidney Foundation; Board membership (money to institution): National Kidney Foundation; Grants/grants pending (money to institution): National Kidney Foundation. Disclosures can also be viewed at www.acponline.org/authors /icmje/ConflictOfInterestForms.do?msNum=M11-2267.

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Obtaining of funding: K. Uhlig.

Administrative, technical, or logistic support: A. Earley, E.J. Lamb, K. Uhlig.

Collection and assembly of data: A. Earley, D. Miskulin, E.J. Lamb, A.S. Levey, K. Uhlig.

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Appendix Table 1. Search Strategy

- 1. exp kidney glomerulus/
- 2. exp kidney disease/
- 3. exp kidney function tests/
- 4. exp renal replacement therapy/
- 5. exp kidney transplantation/
- exp kidney, artificial/
 renal.af. or renal.tw.
- 8. kidney.af. or kidney.tw.
- 9. or/1-8
- 10. limit 9 to humans
- 11. limit 9 to (guideline or meta analysis or practice guideline or "review")
- 12. 10 not 11
- 13. glomerular filtration rate.af. or glomerular filtration rate.tw.
- 14. gfr.af.
- 15. exp kidney function tests/
- 16. serum creatin\$.af. or serum creatin\$.tw.
- 17. creatin\$.af. or creatin.tw.
- 18. cystat\$.af. or cystat\$.tw.
- 19. or/13-18
- 20. predict\$.af.
 21. formula.af.
- 22. equation.af.
- 23. exp regression analysis/ or regression analysis.mp.
- 24. 20 or 21 or 22 or 23
- 25. 12 and 19 and 24
- 26. limit 25 to yr="1999-2011"

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MORD equation without ethnicity 175 × SC ⁻¹¹⁵⁴ × age ^{-0.200} × 0.742 Tete MURD study frema age, 306 5) Tete MURD study of the advisor Tete MURD study (if ternale) Tete MURD study (if ternal	North America, Europe, and					
CCD-EPI equation 141 × min(SC/A, 1)* × max(SC/A, main(SC/A, 1)* × max(SC/A, so 2) for backs and 0 for males, and 4 clinical so 0 for finates and 0 for males, are and 4 clinical so 0 for finates and 0 for males, are and 4 clinical so 0 for finates and - 0.411 so 0 for males, minimum of SC/A or 1, and max indicates the maximum of SC/A or 1 2354 participants from resured in areas and - 0.411 areas areas are -0.303 × 0.742 (if female) 13 Japanese patients are -0.303 × 0.742 (if female) 13 Japanese patients area -0.303 × 0.742 (if female) 10 areas and - 0.411 areas areas area -0.303 × 0.742 (if female) 10 areas areas area -0.303 areas areas area -0.303 × 0.742 (if female) 10 areas areas areas -0.303 × 0.742 (if female) 10 areas areas -0.713 × 0.742 (if female) 10 areas -0.71	Australia Levey et al, 2006 (8)	MDRD equation MDRD equation without ethnicity factor*	$\begin{array}{l} 175 \times SCr^{-1.154} \times age^{-0.203} \times 0.742 \\ (ff \ female) \times 1.212 \ (ff \ black) \\ 175 \times SCr^{-1.154} \times age^{-0.203} \times 0.742 \\ (ff \ female) \end{array}$	1628 patients enrolled in the MDRD study (mean age, 50.6 y)	 ¹²⁵I-lothalamate (urine); GFR measured in mL/min per 1.73 m² (39.8 mL/min per 1.73 m² SD, 21.21) 	Samples from MDRD Study were assayed from 1988 to 1994 with the Beckman Synchron CX3 kinetic Jaffe assay (Global Medical Instrumentation, Ramsey, Minnesotia). Samples were reassayed in 2004 with the same instrument. The Beckman assay was calibrated to the Roche enzymatic assay (Roche Diagnostics, Basel, Switzerland), which is
MDRD equation with Japanese 0.308 × 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 (if female) 413 Japanese patients in 80 medical measured in mL/min development or MDRD equation Inulin (urine); GFR development or internal validation set) Le Japanese-modified 0.741 × 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 (if female) 248 inpatients with nullin (urine); GFR and determining the equation for Japanese patients 10-101 (urine); GFR development or internal validation set) Le MDRD equation 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 (if female) 248 inpatients with nullin (urine); GFR and determining the equation for Japanese patients Paraforese patients Le MDRD equation 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 250 cases of Thai Parafor set) Paraforese patients MDRD equation 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 250 cases of Thai Paraforese patients Paraforese patients MDRD equation 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 250 cases of Thai Paraforese patients Paraforese patients MDRD equation 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 250 cases of Thai Paraforese patients Paraforese patients With Thai (If female) × 1.1.129 (If Thai) Patients with CKD Paraforese patients Parafore Wote 175 × 5Cr ^{-1,154} × age ^{-0,203} × 0.742 250 cases of Thai Parafore Parafore Patients Patients Patient	Levey et al, 2009 (25)	CKD-EPI equation	141 × min(SCr/ κ , 1) ^{α} × max(SCr/ κ , 1) ^{-1.209} × 0.993 ^{age} × 1.018 (if female) × 1.159 (if black), where κ is 0.7 for females and 0.9 for males, α is -0.329 for females and -0.411 for males, min indicates the minimum of SCr/ κ or 1, and max indicates the maximum of SCr/ κ or 1	8254 participants from 6 research studies and 4 clinical populations (mean age, 47 y)	 ¹²⁵I-lothalamate (urine); GFR measured in mL/min per 1.73 m² (68 mL/min per 1.73 m² [SD, 40]) 	traceable to an IUMS assay at NIS . Values of SCr were recalibrated to standardized SCr measurements at the Cleveland Clinic by using a Roche enzymatic assay (Roche-Hitachi P-Module instrument with Roche Creatininase Plus assay; Hoffman-La Roche, Basel, Switzerland).
MDRD equation 0.808 × 175 × SCr ^{-11,154} × 13 Apanese patients 413 Apanese patients Inilin (urine); GR Le with Japanese age ^{-0.203} × 0.742 (if female) in 80 medical measured in mL/min coefficient 0.041 × 175 × SCr ^{-11,154} × age ^{-0.203} 248 inpatients with Inilin (urine); GR Le Japanese-modified 0.741 × 175 × SCr ^{-11,154} × age ^{-0.203} 248 inpatients with Inulin (urine); GR Le MDRD equation × 0.742 (if female) × 0.742 (if female) 248 inpatients with Inulin (urine); GR Le MDRD equation × 0.742 (if female) × 0.742 (if female) 248 inpatients with Inulin (urine); GR Le MDRD equation × 0.742 (if female) × 0.742 (if male) 248 inpatients with Inulin (urine); GR Le MDRD equation × 0.742 (if female) × 0.742 (if male) internal validation set) eerologneent or MDRD equation × 0.742 (if female) × 0.742 (if male) internal validation set) erologneent or MDRD equation 175 × SCr ⁻¹¹⁴⁴ × age ^{-0.203} × 0.742 250 cases of Thai internal validation set) erologneent or MDRD equation 175 × SCr ⁻¹¹⁴⁴ × age ^{-0.203} × 0.742 250 cases of Thai PortGeDTPA; GR fm with Thai (if female) × 1.129 (if Thai) with	Outside of North America, Europe, and Australia					
Japanese-modified 0.741 × 175 × SCr ^{-1,154} × age ^{-0.203} 248 inpatients with Inulin (urine), GFR Le MDRD equation × 0.742 (if female) × 0.742 (if female) × 0.742 (if female) RCND for estimating per 1.73 m ² (ND for carginating the per 1.73 m ² (ND for carginating the per 1.73 m ² (ND for cargination for cargination for cargination set) per 1.73 m ² (ND for cargination set) MDRD equation 1.75 × Scr ^{-1,154} × age ^{-0.203} × 0.742 250 cases of Thai Pointer of the measured in mL/min with Thai MDRD equation 1.75 × Scr ^{-1,154} × age ^{-0.203} × 0.742 250 cases of Thai Pointer CBTPA; GFR Fa with Thai (if female) × 1.129 (if Thai) who were in stable Pointer CDTPA; GFR Fa coefficient (if female) × 1.129 (if Thai) who were in stable Pointer development or internal or internal vith Or development	Horio ét al, 2010 (29)	MDRD equation with Japanese coefficient	0.808 × 175 × SCr ^{-1,154} × age ^{-0.203} × 0.742 (if female)	413 Japanese patients in 80 medical centers (mean age, 51.4 y)	Inulin (urine); GFR measured in mL/min per 1.73 m ² (ND for development or internal validation set)	Levels of SCr were measured by using a Hitachi enzymatic creatinine assay and the values obtained were compared with those of the Cleveland Clinic.
MDRD equation 175 × SCr ^{-1.154} × age ^{-0.203} × 0.742 250 cases of Thai ^{99m} Tc-DTPA; GFR Fa with Thai (if female) × 1.129 (if Thai) patients with CKD measured in mL/min coefficient (if female) × 1.129 (if Thai) who were in stable per 1.73 m ² (plasma) coefficient (ND for development age, 59.5 y) or internal validation set)	Imai et al, 2007 (47)	Japanese-modified MDRD equation	0.741 × 175 × SCr ^{-1.154} × age ^{-0.203} × 0.742 (if female)	248 inpatients with CKD for estimating and determining the coefficient of the MDRD Study equation for Japanese patients (mean age, 50.1 y)	Inulin (urine); GFR measured in mL/min per 1.73 m ² (ND for development or internal validation set)	Levels of SCr were assayed by both enzymatic and noncompen- sated kinetic Jaffe methods for the 116 patients in the inulin clinical study and by an enzymatic method for the 132 inpatients at Tokyo Women's Medical University between 2003 and 2004 and the 168 patients with CKD at the University of Tsukuba Hospital from 1988 to 1994. The SCr levels of the samples from the 101 patients from Tokyo Women's Medical University between 2001 and 2002 were measured by a noncompensated kinetic Jaffe method. The SCr levels used in the inulin clinical trial were simultaneously measured by both the noncompensated kinetic Jaffe method and the enzymatic method at the international Organization for Istandardization-approved central laboratory. The SCr values from the noncompen- sated Jaffe method were 0.207 higher than those from the enzymatic method throughout the measured range, based on calibrated standards that were indirectly calibrated with
	Praditpornsipa et al, 2011 (31)	MDRD equation with Thai coefficient	175 × SCr ^{-1.154} × age ^{-0.203} × 0.742 (if female) × 1.129 (if Thai)	250 cases of Thai patients with CKD who were in stable condition (mean age, 59.5 y)	^{99m} Tc-DTPA; GFR measured in mL/min per 1.73 m ² (plasma) (ND for development or internal validation set)	Fasting SC: levels measured by using a Roche enzymatic assay (Roche Diagnostics, Indianapolis, Indiana), and values adjusted by using IDMS reference SCr (SRM 967) from NIST. Levels of SCr also measured by a Roche kinetic Jaffe assay (Roche Diagnostics) without adjustments to IDMS reference. The SCr levels obtained from enzymatic and Jaffe assays were used in each estimated GFR equation accordingly.

Appendix Table 2—Continued	inued				
Study, Year (Reference)	Equation Name	Equation	Development and Internal Validation Population	GFR Measurement Method (Mean GFR)	SCr Assay Method
Horio et al, 2010 (29)	CKD-EPI equation with Japanese coefficient	0.813 \times 141 \times min(SCr/ κ , 1) ^a \times max(SCr/ κ , 1) ^{-1.209} \times 0.993 ^{age} \times 1.018 (if female) \times 1.159 (if black), where κ is 0.7 for females and 0.9 for males, α is -0.329 for females and -0.411 for males, min indicates the minimum of SCr/ κ or 1, and max indicates the maximum of SCr/ κ or 1.	413 Japanese patientsin 80 medicalcenters (mean age,51.4 y)	Inulin (urine); GFR measured in mL/min per 1.73 m ² (ND for development or internal validation set)	Levels of SCr measured by Hitachi enzymatic assay and values obtained compared with those of the Cleveland Clinic.
Matsuo et al, 2009 (34)	JSN-CKDI equation (equation 2) 3-variable Japanese equation (equation 4)	171 × SCr ^{-1 004} × age ^{-0.287} × 0.782 (if female) 194 × SCr ^{-1 094} × age ^{-0.287} × 0.739 (if female)	 413 Japanese patients in 80 medical centers; data collected from 1 December 2006 to 20 April 2007 (mean age, 51.4 y) 	Inulin (urine) (59.1 mL/min per 1.73 m² [SD, 35.4])	Levels of SCr measured by using a Hitachi enzymatic assay (Hitachi, Tokyo, Japan) and values obtained compared with those of the Cleveland Clinic.
Levey et al, 2000 (7)	Original MDRD equation, calibrated to the Cleveland Clinic creationic measurements (study equation 2)†	186 × SCr ^{-1,154} × age ^{-0.203} × 0.742 (if female)	1628 patients whose GFR was measured as part of the MDRD Study (mean age, 50.6 y)	 ¹²⁵I-lothalamate (urine) GFR measured in mL/min per 1.73 m² [SD, 21.2]) 	Serum and urine creatinine levels measured by using kinetic Jaffe assay.
Ma et al, 2006 (11)	Original MDRD equation, calibrated to Cleveland Clinic creatinine measure- ments, with Chinese coefficient (study equation 4)† Chinese equation (study equation 6)‡	 186 × SCr^{-1,154} × age^{-0.203} × 0.742 (if female) × 1.227 (if Chinese), with SCr calibrated to the Cleveland Clinic 206 × SCr^{-1.234} × age^{-0.227} × 0.803 (if female), with SCr calibrated to the Cleveland Clinic 	454 patients from 9 renal institutes of university hospitals located in 9 geo- graphic regions of China (mean age, 49.9 y)	^{99m} Tc-DTPA (plasma); GFR measured in mL/min per 1.73 m ² (ND for development or internal validation set)	Levels of SCr measured by using a Hitachi kinetic Jaffe assay and values calibrated to the Cleveland Clinic Laboratory.
Praditpornsilpa et al, 2011 (31)	Thai estimated GFR equation	375.5 × SCr ^{-0.848} × age ^{-0.364} × 0.712 (if female); r ² = 0.869	250 Thai patients with CKD who were in stable condition (mean age, 59.5 y)	^{99m} Tc-DTPA (plasma); GFR measured in mL/min per 1.73 m ² (ND for development (ND for development or internal validation set)	Fasting SCr levels measured by using a Roche enzymatic assay (Roche Diagnostics) and values adjusted by using IDMS reference SCr (SRM 967) from NIST. Levels of SCr were also measured by using a Roche kinetic Jaffe assay (Roche Diagnostics) without adjustments to IDMS reference. The SCr levels obtained from enzymatic and Jaffe assays were used in each estimated GFR equation accordingly.

W-272 5 June 2012 Annals of Internal Medicine Volume 156 • Number 11

CKD = chronic kidney disease; CKD-EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; IDMS = isotope-dilution mass spectrometry; JPN-CKDI = Japanese Society of Nephrology-Chronic Kidney Disease Initiatives; MDRD = Modification of Diet in Renal Disease; ND = not documented; NIST = National Institute of Standards and Technology; SCr = serum creatinine; SRM = standard * African American coefficient of the MDRD equation. (with the 186 coefficient, For use with an SCr traceable to the SRM, the 186 coefficient should be replaced with the 175 coefficient from the reexpressed MDRD equation (with the 186 coefficient). For use with an SCr traceable to the SRM, the SCr should be replaced with the 175 coefficient from the reexpressed MDRD equation (with the 186 coefficient). For use with an SCr traceable to the SRM, the SCr should be replaced with the 175 coefficient from the reexpressed MDRD equation (with the 186 coefficient). For use with an SCr traceable to the SRM, the SCr should be replaced by SCr × 0.95, which represents the calibration factor relating the SCr assay. In the Cleveland Clinic laboratory to the standardized SCr assay.

Appendix Table 3. Overview Table of Equations Developed to Predict GFR Based on Serum Creatinine Assays Not Traceable to the Standard Reference Material

Study, Year (Reference)	Equation Name	Expression	Formula
Cockcroft and Gault, 1976 (48)	Cockcroft–Gault	CrCl in mL/min	(140 – age) \times weight/(72 \times SCr) \times 0.85 (if female) SCr in mg/dL
Levey et al, 2006 (8)	4-variable MDRD	GFR in mL/min per 1.73 m ²	186 \times SCr $^{-1.154}$ \times age $^{-0.203}$ \times 0.742 (if female) SCr in mg/dL
Levey et al, 1999 (12)	6-variable MDRD	GFR in mL/min per 1.73 m ²	$170 \times Scr^{-0.999} \times age^{-0.176} \times 1.180$ (if black) \times 0.762 (if female) \times BUN ^{-0.170} \times albumin ^{0.318} SCr in mg/dL, BUN in mg/dL, and albumin in g/dL
Ma et al, 2006 (11)	Chinese-modified MDRD	GFR in mL/min per 1.73 m ²	1.233 × 186 × SCr ^{-1.154} × age ^{-0.203} × (0.742 if female) SCr in mg/dL
Rule et al, 2004 (49)	Quadratic equation by Rule	GFR in mL/min per 1.73 m ²	exp[1.911 + (5.249/SCr) - (2.114/SCr ²) - 0.00686 × age - 0.205 (if female)] SCr in mg/dL
Jelliffe, 1971 (50)	Jelliffe, 1971	CrCl in mL/min	Men: (100/SCr) – 12 Women: (80/SCr) – 7 SCr in mg/dL
Jelliffe, 1973 (51)	Jelliffe, 1973	CrCl in mL/min	[98 – 0.8 × (age – 20)]/SCr × (0.9 if female) SCr in mg/dL
Mawer et al, 1972 (52)	Mawer	CrCl in mL/min	Men: weight × [29.3 - (0.203 × age)] × $[1 - (0.03 × SCr)]/$ (14.4 × SCr) × weight/70 Women: weight × [25.3 - (0.175 × age)] × $[1 - (0.03 × SCr)]/(14.4 × SCr) × weight/70$ SCr in mg/dL
Hull et al, 1981 (53)	Hull	CrCl in mL/min	[(145 – age)/SCr – 3] \times (weight/70) \times 0.85 (if female) SCr in mg/dL
Gates, 1985 (54)	Gates	CrCl in mL/min	Men: $(89.4 \times SCr^{-1.2}) + [(55 - age) \times (0.447 \times SCr^{-1.1})]$ Women: $(60 \times SCr^{-1.1}) + [(56 - age) \times (0.3 \times SCr^{-1.1})]$ SCr in mg/dL
Bjornsson et al, 1983 (55)	Bjornsson	CrCl in mL/min	Men: $[27 - (0.173 \times age)] \times weight \times 0.007/SCr$ Women: $[25 - (0.175 \times age)] \times weight \times 0.007/SCr]$ SCr in mg/dL
Walser et al, 1993 (56)	Walser	GFR in mL/min	Men: 7.57/SCr - 0.103 × age) + (0.096 × weight) - 6.66 Women: 6.05/SCr - (0.08 × age) + (0.08 × weight) - 4.81 SCr in μ mol/L
Nankivell et al, 1995 (57)	Nankivell	GFR in mL/min	 (6.7/SCr) + (weight/4) – (BUN/2) – (100/height²) + (35 if male or 25 if female) SCr in μmol/L and BUN in mmol/L
Imai et al, 2007 (47)	JSN-CKDI	GFR in mL/min per 1.73 m ²	$1.223 \times 186 \times {\rm SCr}^{-1.154} \times {\rm age}^{-0.203} \times 0.742$ (if female) SCr in mg/dL

BUN = blood urea nitrogen; CrCl = creatinine clearance; GFR = glomerular filtration rate; JSN-CKDI = Japanese Society of Nephrology-Chronic Kidney Disease Initiatives; MDRD = Modification of Diet in Renal Disease; SCr = serum creatinine.

Appendix Table 4. Performance of Creatinine-Based GFR Estimating Equations in North America, Europe, and Australia in Subgroups by GFR

Study, Year (Reference)	Country	Population	Patients, n	Measure	ed GFR
				Reference Standard	Value (SD), <i>mL/min</i> per 1.73 m ²
Murata et al, 2011 (26)	United States	5238 adults who underwent an outpatient iothalamate	583	¹²⁵ I-Iothalamate (urine)	98.9 (20.1)
		clearance at the Mayo Clinic in Rochester,	97		65.8 (17.4)
		Minnesota (women, 45%; African American, 2%;	2324		57.3 (24.1)
		mean age, 56 y; donor or potential donor, 13%; KTRs, 26%)	1375		52.3 (19.5)
		KTKS, 20%)	859		46.2 (29.5)
Levey et al, 2009 (25)	United States	External validation set comprising 16 studies (women, 45%; white or other, 87%; black, 10%; Hispanic, 2%; Asian, 2%; mean age, 50 y; diabetes, 28%; donor, 16%; KTRs, 29%)	3896	¹²⁵ I-Iothalamate (urine) and others	68 (36)
White et al, 2010 (20)	Canada	Stable KTRs (women, 36%; white, 92%; receiving trimethoprim, 19%)	109 98	^{99m} Tc-DTPA (plasma)	58 (22)
Pöge et al, 2011 (19)	Germany	Patients with stable renal function after transplan- tation (women, 40%; white, 99%; mean age,	98 150	^{99m} Tc-DTPA (plasma)	35.9
		49 y)	20		67.9

CKD- $EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; IDMS = isotope-dilution mass spectrometry; KTR = kidney transplant recipient; MDRD = Modification of Diet in Renal Disease; <math>P_{30}$ = percentage of estimated GFR values within 30% of measured GFR; SCr = serum creatinine. * Computed as estimated GFR minus measured GFR. Positive numbers indicate overestimation and negative numbers indicate underestimation of measured GFR. Smaller absolute values indicate lesser bias.

+ The difference between the absolute values for bias.

I he difference between the absolute values for blas.
‡ Lower values indicate greater precision.
§ Higher values indicate greater accuracy.
|| The difference between the P₃₀ values for accuracy (CKD-EPI minus MDRD).
¶ Evaluated as the root mean square error for the regression of estimated GFR on measured GFR.
** Evaluated as the interquartile range for the differences between estimated and measured GFR.

++ Evaluated as the SD of the differences between estimated and measured GFR.

Appendix Table 4—Continued

GFR	Estimation				Results		
SCr Calibration and Assay	Subgroup	Equation	Bias (95% Cl), mL/min per 1.73 m ² *	Difference in Bias, <i>mL/min per 1.73</i> <i>m</i> ² †	Precision (95% CI)‡	P ₃₀ (95% CI), %§	Difference in Accuracy, percentage points
Roche Jaffe assay (Roche	Potential kidney	MDRD	-19.2	9	ND	75.8	12.9
P- or D-Modular or Roche Cobas C501	donors	CKD-EPI	-10.2			88.7	
with Roche Creat-	Postnephrectomy kidney donors	MDRD	-11	5.3		82.5	11.3
ininase Plus assay;	Other organ	CKD-EPI MDRD	-5.7 -1.5	-0.5		93.8 80.6	-2.1
Roche Diagnostics,	recipients	CKD-EPI	2.0	-0.5		78.5	-2.1
Indianapolis, Indiana)	Kidney recipient	MDRD	-1.9	0.3		80.1	-1.7
with demonstrated IDMS alignment	ritario) recipiente	CKD-EPI	1.6	0.0		78.4	
ibinis alignment	Native patients	MDRD	-2.3	1.9		75.2	-0.2
	with CKD	CKD-EPI	-0.4			75.0	
Roche enzymatic assay (Roche–Hitachi P-Module instrument with Roche	$\begin{array}{l} \text{GFR} \geq 60 \\ \text{mL/min per} \\ 1.73 \text{ m}^2 \end{array}$	MDRD	-10.6 (-9.8 to -11.3)	7.1	0.248 (0.238 to 0.258)¶	84.7 (83.0 to 86.3)	3.6
Creatininase Plus assay; Hoffman-La Roche, Basel,		CKD-EPI	-3.5 (-2.6 to -4.5)		0.213 (0.203 to 0.223)¶	88.3 (86.9 to 89.7)	
Switzerland) recalibrated to standardized SCr at the Cleveland Clinic	GFR <60 mL/min per 1.73 m ²	MDRD	-3.4 (-2.9 to -4.0)	1.3	0.294 (0.280 to 0.308)¶	77.2 (75.5 to 79.0)	2.7
		CKD-EPI	-2.1 (-1.7 to -2.4)		0.284 (0.270 to 0.298)¶	79.9 (78.1 to 81.7)	
For the reexpressed	GFR >60	MDRD	-14.0	5.8	15.8**	79 (69 to 86)	10
MDRD Study and CKD-EPI equations,	mL/min per 1.73 m ²	CKD-EPI	-8.2		17.9**	89 (81 to 94)	
SCr adjusted to IDMS	GFR <60	MDRD	-4.0	1.9	11.6**	80 (71 to 86)	-1
standard	mL/min per 1.73 m ²	CKD-EPI	-2.1		12.3**	79 (70 to 85)	
Jaffe assay on Dimension	GFR >60	MDRD	1.23	-5.77	16.3††	85.0	-15
RxLTM analyzer (Dade Behring, Marburg,	mL/min per 1.73 m ²	CKD-EPI	7.00		15.2††	70.0	
Germany); assay adjusted for calibration	GFR <60	MDRD	4.93	-3.29	8.75††	70.0	-6.7
with the IDMS method	mL/min per 1.73 m ²	CKD-EPI	8.22		10.1††	63.3	

Appendix Table 5. Performance of Creatinine-Based GFR Estimating Equations in North America, Europe, and Australia in Subgroups by Race

Study, Year (Reference)	Country	Population	Patients, <i>n</i>	Measured GFR	¥	CFK ESTI	GFR Estimation		Results	
				Reference Standard	Value (SD), mL/min per 1.73 m ²	SCr Calibration and Assay	Equation	Bias (95 % Cl), mL/min per 1.73 m ² *	Precision (95% CI)†	P ₃₀ (95% CI), %‡
Stevens et al, 2011 (27)	United States and Europe	External validation set comprising 16	384 black patients	¹²⁵ I-Iothalamate (urine) and other filtration	62 (34)	Calibrated to standardized SCr measurements by	CKD-EPI equation with 2-level race or	0.8 (-0.6 to 2.0)	0.242 (0.221 to 0.265)†	82 (78 to 85)
		45%; white or other, 87%; black, 10%: Hispanic, 2%;				Hitachi P-Module instrument with Roche Creatininase Plus assay:	CKD-EPI equation with 4-level race or ethnicity coefficient	0.9 (-0.6 to 2.0)	0.243 (0.221 to 0.266)†	82 (80 to 85)
		Asian, 2%; mean age, 50 y; diabetes, 28%; donor, 16%;	185 Native American and Hispanic		105 (47)	Hoffmann-La Roche, Basel, Switzerland) enzymatic assay at the	CKD-EPI equation with 2-level race or ethnicity coefficient	-2.3 (-5.1 to 2.1)	0.265 (0.223 to 0.310)†	80 (74 to 85)
		KTRs, 29%)	patients			Cleveland Clinic	CKD-EPI equation with 4-level race or ethnicitv coefficient	-1.6 (-4.2 to 3.0)	0.264 (0.222 to 0.310)†	81 (76 to 87)
			67 Asian patients		53 (31)		CKD-EPI equation with 2-level race or ethnicity coefficient	-2.1 (-1.1 to -0.3)	0.302 (0.188 to 0.436)†	85 (76 to 93)
							CKD-EPI equation with 4-level race or ethnicity coefficient	-0.8 (-2.6 to 2.2)	0.293 (0.178 to 0.424)†	85 (76 to 93)
			3378 white or other patients		69 (36)		CKD-EPI equation with 2-level race or ethnicity coefficient	-2.8 (-3.2 to -2.4)	0.250 (0.240 to 0.258)†	84 (83 to 86)
							CKD-EPI equation with 4-level race or ethnicity coefficient	-2.9 (-3.4 to -2.5)	0.250 (0.240 to 0.259)†	84 (83 to 85)

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creating. * Computed as estimated GFR minus measured GFR. Positive numbers indicate overestimation and negative numbers indicate underestimation of measured GFR. Smaller absolute values indicate lesser bias. † Lower values indicate greater precision. Evaluated as the root mean square error for the regression of estimated GFR on measured GFR.

Study, Year (Reference)	Country	Population	Patients, n	Measured (GFR
(Kelelelite)				Reference Standard	Value (SD), mL/min per 1.73 m ²
Horio et al, 2010 (29)	Japan	Japanese patients (women, 42%; mean age, 54 y; diabetes, 22%; donors, 3%; KTRs, 1%; hypertension, 58%)	206	Inulin (urine)	45 (25)
			144		
Teo et al, 2011 (32)	Singapore	Outpatients with CKD in the nephrology clinic at National	160	⁹⁹ Tc-DTPA (plasma)	36 (14)
		University Hospital, Singapore (women, 48%; Chinese, 41%; Malay, 31%; Indian or other, 28%; median age, 58 y; diabetes, 23%; hypertension, 50%)	72		86 (19)
van Deventer et al, 2008 (30)	South Africa	Black South African patients (women, 49%; median age, 47 y; diabetes, 25%; donors, 7%; HIV, 20%; hypertension, 36%)	41	⁵¹ Cr-EDTA (plasma)	61.5 (49.6
			39		
			120		

Appendix Table 6. Performance of Creatinine-Based GFR Estimating Equations Outside North America, Europe, and Australia in Subgroups by GFR

CKD-chronic kidney disease; CKD-EPI = Chronic Kidney Disease Epidemiology Collaboration; GFR = glomerular filtration rate; IDMS = isotope-dilution mass spectrometry; KTR = kidney transplant recipient; MDRD = Modification of Diet in Renal Disease; NIST = National Institute of Standards and Technology; P_{30} = percentage of estimated GFR values within 30% of measured GFR; SCr = serum creatinine; SRM = standard reference material.

Computed as estimated GFR minus measured GFR. Positive numbers indicate overestimation and negative numbers indicate underestimation of measured GFR. Smaller absolute values indicate lesser bias.

† The difference between the absolute values for bias (MDRD minus CKD-EPI).

‡ Lower values indicate greater precision.

§ Higher values indicate greater accuracy.

|| The difference between the P₃₀ values for accuracy (CKD-EPI minus MDRD).

¹ Evaluated as the root mean square error for the regression of estimated GFR on measured GFR. ** Evaluated as the interquartile range for the differences between estimated and measured GFR.

++ African American coefficient of the MDRD equation.

Appendix Table 6—Continued

GI	FR Estimation				Results		
SCr Calibration and Assay	Subgroup	Equation	Bias (95% CI), mL/min per 1.73 m ² *	Difference in Bias, mL/min per 1.73 m ² t	Precision (95% CI)‡	P ₃₀ (95% CI), %§	Difference in Accuracy, percentage points
Enzymatic assay validated by using the calibration panel of the Cleveland	GFR >60 mL/min per 1.73 m ²	MDRD with Japanese coefficient	-7.8	0.5	23.5¶	82 (75 to 87)	6
Clinic		CKD-EPI with Japanese coefficient	-7.3		21.8¶	88 (82 to 92)	
	$\begin{array}{l} {\rm GFR} < \!\! 60 \mbox{ mL/min} \\ {\rm per} \mbox{ 1.73 } \mbox{ m}^2 \end{array}$	MDRD with Japanese coefficient	3.3	-1.1	15.9¶	67 (61 to 74)	-2
		CKD-EPI with Japanese coefficient	4.4		14.4¶	65 (58 to 71)	
Siemens enzymatic assay (Siemens Advia 2400;	GFR >60 mL/min per 1.73 m ²	MDRD	-5.3 (-9.5 to -1.2)	4.4	18.3 (10.3 to 26.4)**	81.9 (73.1 to 90.8)	9.8
Siemens, Munich,	·	CKD-EPI	0.9 (-4.1 to 5.9)		22.0 (16.7 to 27.2)**	91.7 (85.3 to 98.1)	
Germany) calibrated with manufacturer-	GFR <60 mL/min per 1.73 m ²	MDRD	-2.4 (-3.7 to -1.1)	0.9	9.2 (7.0 to 11.4)**	78.8 (72.4 to 85.1)	0
provided materials traceable to standardized creatinine (NIST SRM 967) measured by using IDMS		CKD-EPI	−1.5 (−2.8 to −0.1)		9.3 (7.0 to 11.6)**	78.8 (72.4 to 85.1)	
Rate-blanked, Roche- compensated, kinetic Jaffe assay (Roche	Estimated GFR, >60 mL/min per 1.73 m ²	MDRD with ethnicity factort†	20.4 (17.6 to 28)	15.3	35.1¶	51	25
Modular analyzer; Roche Diagnostics, Indianapolis, Indiana)		MDRD without ethnicity factort†	5.1 (-0.3 to 17.0)		26.8¶	76	
with calibration traceable to IDMS; compared with Roche enzymatic assay	Estimated GFR, 30 to 60 mL/min per 1.73 m ²	MDRD with ethnicity factor++	8.8 (-2.2 to 14.8)	8.4	18.0¶	53	22
(Creatinine Plus; Roche Diagnostics, Indiana- polis, Indiana) at the		MDRD without ethnicity factort†	0.4 (-6.4 to 5.1)		11.8¶	75	
Cleveland Clinic with values adjusted by regression	Estimated GFR <30 mL/min per 1.73 m ²	MDRD with ethnicity factort†	1.7 (-1.7 to 4.4)	0.3	7.2¶	55	12
		MDRD without ethnicity factor††	-1.4 (-4.0 to 2.2)		7.0¶	67	