

1 **A systematic review of diagnostic and prognostic models of chronic kidney disease in low-**  
2 **and middle- income countries**

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27 **ABSTRACT**

28 **Objective:** To summarize available chronic kidney disease (CKD) diagnostic and prognostic models  
29 in Low- and Middle-Income countries (LMIC)

30 **Method:** Systematic review (PRISMA guidelines). We searched Medline, EMBASE, Global Health  
31 (these three through OVID), Scopus and Web of Science from inception to April 9th, 2021, April 17th,  
32 2021 and April 18th, 2021, respectively . We first screened titles and abstracts, and then studied in  
33 detail the selected reports; both phases were conducted by two reviewers independently. We followed  
34 the CHARMS recommendations and used the PROBAST for risk of bias assessment.

35 **Results:** The search retrieved 14,845 results, 11 reports were studied in detail and nine (n= 61,134)  
36 were included in the qualitative analysis. The proportion of women in the study population varied  
37 between 24.5%-76.6%, and the mean age ranged between 41.8-57.7 years. Prevalence of  
38 undiagnosed chronic kidney disease ranged between 1.1%-29.7%. Age, diabetes mellitus and sex  
39 were the most common predictors in the diagnostic and prognostic models. Outcome definition varied  
40 greatly, mostly consisting of urinary albumin-to-creatinine ratio and estimated glomerular filtration rate.  
41 The highest performance metric was the negative predictive value. All studies exhibited high risk of  
42 bias, and some had methodological limitations.

43 **Conclusion:** There is no strong evidence to support the use of a CKD diagnostic or prognostic model  
44 throughout LMIC. The development, validation and implementation of risk scores must be a research  
45 and public health priority in LMIC to enhance CKD screening to improve timely diagnosis.

46

47 **Keywords:** population health; prognosis research; non-communicable diseases

48 **Strengths and limitations of this study**

49 **Strengths**

- 50 • An extensive search was conducted, involving five major databases (Medline, Embase,  
51 Global Health, Scopus and Web of Science).
- 52 • A comprehensive list of available CKD diagnostic and prognostic models and their limitations  
53 is provided, which were not previously accounted for in the LMIC population.
- 54 • This study adhered to PRISMA, CHARMS and PROBAST guidelines.

55 **Limitations**

- 56 • Meta-analysis was not possible due to the heterogeneity in the measurement of outcomes.
- 57 • Additional data sources such as grey literature were not retrieved.

58

## 59 **INTRODUCTION**

60 Chronic kidney disease (CKD) is a condition with a large burden globally. Between 1990 and 2017,  
61 the health metrics of CKD showed a bleak profile: mortality, incidence and kidney transplantation  
62 rates increased by 3%, 29% and 34%, respectively.<sup>1</sup> CKD led to 1.2 million deaths in 2017 and in the  
63 best-case scenario, CKD mortality will increase to 2.2 million deaths and become the 5<sup>th</sup> cause of  
64 years of life lost (YLL) by 2040.<sup>2</sup> CKD reveals disparities between low- and middle-income countries  
65 (LMIC) and high-income countries (HIC). In the period 1990-2016, the age-standardised disability-  
66 adjusted life-years (DALY) due to CKD was the highest in LMIC,<sup>3</sup> where they need to optimize CKD  
67 early diagnosis.

68 Risk scores are a cost-effective alternative for CKD screening and early diagnosis.<sup>4</sup> These equations  
69 require less resources and contribute to decision making,<sup>5</sup> and allow screening of large populations.<sup>4</sup>  
70 Many of the available CKD risk scores have been developed in HIC,<sup>6-8</sup> and they may not be used in  
71 LMIC without recalibration to secure accurate predictions. How many CKD risk scores there are for  
72 LMIC, and what their strengths and limitations are, remains largely unknown.<sup>9,10</sup> This limits our  
73 knowledge of what tools there are to enhance CKD screening in LMIC. Similarly, this lack of evidence  
74 prevents planning research to overcome the limitations of available models. To fill these gaps and to  
75 inform CKD screening strategies in LMIC, we summarized available CKD diagnostic and prognostic  
76 models in LMIC.

## 77 **METHODS**

### 78 **Protocol and registration**

79 This systematic review and critical appraisal of the scientific literature was conducted following the  
80 Preferred Reporting Items for Systematic Reviews and Meta-Analysis guidelines (PRISMA)  
81 statement<sup>11</sup> (S1 Table). Protocol is available elsewhere<sup>12</sup> and in the S1 Text. We followed the  
82 CHecklist for critical Appraisal and data extraction for systematic Reviews of prediction Modelling  
83 Studies (CHARMS) guidelines.<sup>13,14</sup>

### 84 **Information sources**

85 We searched Medline, EMBASE, Global Health (these three through OVID), Scopus and Web of  
86 Science from inception to April 9<sup>th</sup>, 2021, April 17<sup>th</sup>, 2021 and April 18<sup>th</sup>, 2021, respectively. The

87 search strategy is available in S2 Table. We also screened the references of relevant systemic  
88 reviews<sup>10</sup> and of the selected studies.

### 89 **Eligibility criteria**

90 We sought models which assessed the current CKD status (i.e., diagnostic) or future CKD risk (i.e.,  
91 prognostic), aiming to inform physicians, researchers, and the general population (Table 1). Reports  
92 could include model derivation, external validation, or both. The target population was adults ( $\geq 18$   
93 years) in LMIC according to The World Bank.<sup>15</sup>

### 94 **Study selection**

95 Reports were selected if the study population included people who were from and currently living in  
96 LMIC. Cross-sectional (diagnostic models) and longitudinal studies (prognostic models) with a  
97 random sample of the general population were included. The outcome was CKD based on a  
98 laboratory or imaging test (isolated or in combination with self-reported diagnosis): urine albumin-  
99 creatinine ratio, urine protein-creatinine ratio, albumin excretion ratio, urine sediment, kidney images,  
100 kidney biopsy or the estimated glomerular filtration rate (eGFR).<sup>12</sup>

101 Reports had to present the development and/or validation of a multivariable model. On the other  
102 hand, reports with LMIC populations outside LMIC, or those including foreigners living in LMIC, were  
103 excluded. Reports that only studied people with underlying conditions (e.g., patients with diabetes),  
104 people with a specific risk factor (e.g., alcohol consumption), or a hospital-based population, were  
105 excluded. We also excluded models that were developed using machine learning techniques due to  
106 their usually poor report of performance metrics, as noted from previous reviews.<sup>16 17</sup> To overcome  
107 this limitation, CHARMS and PROBAST tools are currently being adapted to machine learning  
108 methodology but are yet to be published.<sup>18</sup>

### 109 **Data collation**

110 We used EndNote20 and Rayyan<sup>19</sup> to remove duplicates from the search results. We used Rayyan<sup>19</sup>  
111 to screen titles and abstracts by two reviewers independently (DJA-G and EJA); discrepancies were  
112 solved by consensus. Two reviewers independently (DJA-G and EJA) studied the full length of the  
113 reports selected in the screening phase; discrepancies were solved by consensus. If consensus was  
114 not reached, a third party was consulted (RMC-L). A data extraction form based on the CHARMS  
115 guidelines<sup>14</sup> was developed and not modified during data collation. Data was extracted as presented

116 in the original reports by two reviewers independently (DJA-G and EJA); discrepancies were solved  
117 by consensus.

### 118 **Risk of bias of individual studies**

119 We used the PROBAST (Prediction model Risk of Bias ASsessment Tool) to assess the risk of bias of  
120 diagnostic and prognostic models.<sup>20 21</sup> Two reviewers (EJA and DJA-G) independently ascertained the  
121 risk of bias of individual reports; discrepancies were solved by consensus or a third party (RMC-L).

### 122 **Synthesis of results**

123 A qualitative synthesis was conducted whereby the characteristics of the selected models was  
124 comprehensively described.<sup>12</sup> Quantitative analysis (meta-analysis) was not conducted because the  
125 selected models used different predictors and they had different outcome definitions.

### 126 **Ethics**

127 This review was deemed as a low risk because human subjects were not directly involved. The funder  
128 did not have any role in the conception, conduction, results interpretation, and drafting of this work.  
129 Results and opinions expressed in the article are entirely the authors.

### 130 **Patient and public involvement**

131 No patient involved.

## 132 **RESULTS**

### 133 **Reports selection**

134 The search yielded 14,845 reports. After removing duplicates (1,462 articles), we screened 13,383  
135 titles and abstracts. Then, 11 reports were selected, one of them was not available as full-text,<sup>22</sup> and  
136 the rest (10 articles) were studied in detail. We excluded one report because the study population was  
137 not randomly selected,<sup>23</sup> and another report because it was conducted in a HIC.<sup>24</sup> Additionally, one  
138 report was identified by reference searching.<sup>25</sup> Finally, nine reports (n=61,134) were included in the  
139 qualitative synthesis (Figure 1).

### 140 **General characteristics of the selected reports**

141 Original reports were from Iran,<sup>26</sup> India,<sup>27</sup> Peru,<sup>28</sup> South Africa,<sup>25</sup> two from China<sup>29 30</sup> and three from  
142 Thailand<sup>31-33</sup> (S1 Figure). All studies were developed on community-based populations with random  
143 sampling (S3 Table).

144 Overall, Wu and colleagues studied the largest sample size (n=14,374) which was a population of  
145 workers who underwent health checks;<sup>30</sup> conversely, the smallest sample was studied by Mogueo *et al*  
146 *al* (n=902).<sup>25</sup> The oldest data was collected in 1999<sup>26</sup> whereas the most recent study was published in  
147 2018.<sup>26</sup>

148 The sample size analysed to derive the diagnostic models ranged from 2,368<sup>28</sup> to 14,374 people,<sup>30</sup>  
149 and from 902<sup>25</sup> to 4,940<sup>27</sup> for the validation models. The mean age of participants in the derivation  
150 models varied from 44.9 to 57.7 years, and the proportion of male subjects ranged from 46.8% to  
151 70.5%.<sup>27-30 32 33</sup> The mean age of participants in the validation models varied from 41.8 to 57.1 years,  
152 and the proportion of male subjects ranged from 23.4% to 75.5%<sup>25-28 30-32</sup> (Table 2; S3 Table).

153 The number of CKD cases varied greatly in the derivation models, from 81<sup>28</sup> to 947;<sup>27</sup> the  
154 corresponding numbers in the validation models were 27<sup>32</sup> and 1,359<sup>26</sup>. Of note, number of CKD  
155 cases could not be extracted from the validation work by Bradshaw *et al*<sup>27</sup>. The ratio of outcome  
156 events per number of candidate predictors in the derivation models ranged from 2.3<sup>28</sup> to 135.3<sup>27</sup>. This  
157 ratio could not be calculated for the derivation models by Wen *et al*<sup>29</sup> and Wu *et al*<sup>30</sup>. Across all  
158 reports, missing data were handled by conducting a complete-case analysis;<sup>25-32</sup> this information was  
159 not available in the study by Thakkinstian's *et al*<sup>33</sup> (Table 2; S3 Table).

## 160 **What has been done?**

161 In 2011, Thakkinstian *et al* derived one model using cross-sectional data.<sup>33</sup> In 2015, Mogueo *et al*  
162 used cross-sectional data to validate two models that were previously developed in South Korea and  
163 Thailand using two different outcome definitions for each model, i.e., they provided estimates for four  
164 model validations.<sup>25</sup> In 2016, Wu *et al* used cross-sectional data to derive and validate one model, i.e.,  
165 they provided estimates for two models (one derivation and one validation).<sup>30</sup> In 2017, Carrillo-Larco  
166 *et al* used cross-sectional data to derive and validate two models, i.e., they provided estimates for four  
167 models (two derivations and two validations).<sup>28</sup> In 2017, Saranburut *et al* prospectively validated the  
168 Framingham Heart Study risk score on a cohort using two different outcome definitions, i.e., they  
169 provided estimates for two model validations.<sup>31</sup> In 2017, Saranburut *et al* prospectively developed four  
170 models and validated two of them using cohort data, i.e., they provided estimates for six models (four  
171 derivations and two validations).<sup>32</sup> In 2019, Bradshaw *et al* used cross-sectional data to derive four  
172 models, one of them was validated on two populations (rural and urban), i.e. they provided estimates

173 for six models (four derivations and two validations).<sup>27</sup> In 2020, Asgari and colleagues prospectively  
174 validated a model from the Netherlands for 6- and 9-years CKD prediction, i.e. they provided  
175 estimates for two model validations.<sup>26</sup> In 2020, Wen *et al* prospectively derived two models.<sup>29</sup> Overall,  
176 fourteen models were derived and fifteen underwent validation (hence the 29 rows in Table 4).

### 177 **Outcome ascertainment**

178 Across all reports, CKD was defined as eGFR <60 mL/min/1.73m<sup>2</sup> <sup>25-33</sup> assessed by either the  
179 Modification of Diet Renal Disease (MDRD) formula<sup>25 26 28 29 31 33</sup> or the Chronic Kidney Disease  
180 Epidemiology Collaboration (CKD-EPI) formula.<sup>27 30-32</sup> In addition to the eGFR assessment, Bradshaw  
181 *et al*<sup>27</sup> and Wen *et al*<sup>29</sup> defined CKD as a urinary albumin-to-creatinine ratio (UACR) ≥30 mg/g.  
182 Mogueo *et al* validations also considered CKD as any nephropathy including stages I to V of the  
183 “Kidney Disease: Improving Global Outcomes (KDIGO)” classification.<sup>25</sup> Thakkestian *et al*, also  
184 considered CKD as eGFR ≥60 mL/min/1.73m<sup>2</sup> if it had haematuria or UACR ≥30 mg/g <sup>33</sup> (Table 2).

### 185 **Predictors and modelling**

186 Logistic regression analysis was conducted in all derivation models.<sup>27-30 32 33</sup> Selection of the final  
187 predictors was based on modelling techniques: backward<sup>27 28</sup> and forward selection<sup>29 30 32 33</sup> (S3  
188 Table). All studies categorized numerical variables. The most frequent predictors included in the  
189 models were: age, diabetes mellitus and sex (S2 Figure).

### 190 **Model performance**

191 All studies reported calibration and discrimination metrics, except for the validations by Bradshaw *et*  
192 *al*<sup>27</sup> and Carrillo-Larco *et al*<sup>28</sup> (S3 Table). Regarding discrimination metrics, the area under the  
193 Receiver Operating Characteristic (ROC) curve and C-statistic were over 63%<sup>31</sup> and 70%,<sup>27</sup>  
194 respectively. Among all studies, sensitivity ranged from 56.8%<sup>29</sup> to 84.0%,<sup>25</sup> specificity ranged from  
195 65.1%<sup>29</sup> to 86.3%,<sup>30</sup> positive predictive value (PPV) ranged from 8.8%<sup>28</sup> to 33.8%,<sup>29</sup> and negative  
196 predictive value (NPV) ranged from 89.4%<sup>29</sup> to 99.1%.<sup>28</sup> The NPV was the best metric, consistently  
197 above 89.4% (Table 3).



198 **Risk of bias**

199 All studies showed a high risk of bias due to insufficient or inadequate analytical reporting. The flaw  
200 regarding the analysis criteria can be explained by how original reports handled missing data and  
201 predictors categorization. The participants and predictors criteria had low risk of bias in most of the  
202 reports. Most of the individual reports demonstrated an inappropriate evaluation of performance  
203 metrics.<sup>26 28-33</sup> Low applicability concern was noted (Table 4; S4Table).

204 **DISCUSSION**

205 **Main findings**

206 This systematic review summarized all available risk scores for CKD in LMIC. In so doing, we  
207 provided the most comprehensive list of CKD risk scores to enhance primary prevention and early  
208 diagnosis of CKD in LMIC. Although the available models had acceptable discrimination metrics and,  
209 when available, acceptable calibration metrics, these models had serious methodological limitations  
210 such as a reduced number of outcome events. The best performance metric across risk scores was  
211 the negative predictive value. Overall, CKD risk prediction tools in LMIC need rigorous development  
212 and validation so that they can be incorporated into clinical practice and interventions. The available  
213 evidence would not support using any of the available CKD risk scores across LMIC.

214 **Limitations of the review**

215 We did not search grey literature. We argue that this limitation would not substantially change our  
216 results because these sources are most likely not to have included a random sample of the general  
217 population and are likely to have included a small sample size with few outcome events. That is, we  
218 would not expect to find a report in the grey literature with a much better methodology than that of the  
219 studies herein summarised.

220 **Limitations of the selected reports**

221 Several LMIC do not have a CKD risk score, particularly countries in Central America and Oceania.  
222 This should encourage public health officers and researchers to develop CKD prediction models.  
223 They could conduct new epidemiological studies or leverage on available health surveys with kidney  
224 biomarkers. These models could have pragmatic and direct applications in clinical medicine, by  
225 providing a tool for early identification of CKD cases. Similarly, these models could inform public

226 health interventions and planning, by providing a tool to quantify the size of the population likely to  
227 have or to develop CKD.

228 Clinical guidelines state that CKD is defined as a sustained structural or functional kidney damage for  
229  $\geq 3$  months.<sup>34</sup> In the studies herein summarised, CKD was defined at one point in time. Future work  
230 could expand the definition of CKD to also incorporate the lapse during which the patient had kidney  
231 damage. In addition, different procedures were used to define CKD including eGFR, proteinuria, and  
232 UACR. Even amongst those studies in which CKD was defined with eGFR, they used different  
233 equations to compute the eGFR. Researchers and practitioners in LMIC could agree on the best and  
234 most pragmatic as well as cost-effective definition of CKD, so that future models could use this  
235 definition. This would improve the comparability and extrapolability of the models.

236 All reports in which a new CKD risk score was developed selected the predictors through univariate  
237 analyses,<sup>27-30 32 33</sup> which is not be the best approach to choose predictors.<sup>35-37</sup> Ideally, predictors  
238 should be selected based on expert knowledge, or amongst those with the strongest association  
239 evidence with CKD. In a similar vein, predictors selection should be guided by the target population.  
240 For example, CKD prediction models for populations in LMIC should prioritize simple biomarkers or  
241 inexpensive clinical evaluations (e.g., blood pressure). In this way, the risk score is likely to be used in  
242 clinical practice in resource-limited settings. Another relevant methodological limitation was how the  
243 original reports handled missing data. To the extent possible, multiple imputation should be  
244 implemented to maximize available data and to avoid potential bias by studying only observations  
245 with complete information.

246 Calibration assesses the degree of agreement between actual outcomes and model prediction,  
247 whereas discrimination is the ability of the model to differentiate people with and without the outcome.  
248 Calibration metrics need to be consistently reported and should inform the direction of the  
249 miscalibration. Most of the studies used the Hosmer-Lemeshow  $X^2$  test as the calibration metric.  
250 Unfortunately, this test does not inform on whether the model prediction is overestimating or  
251 underestimating the observed risk; calibration plots are a useful alternative. Therefore, it was not  
252 always possible to reach strong conclusions about the performance of the available models.  
253 Prognostic models should be updated before they can be applied in a new target population. This  
254 process is known as recalibration. Because we found a handful of prognostic models in some

255 countries, it is debatable whether these can be successfully used in other populations. Available  
256 prognostic models for CKD would need to be recalibrated and independently validated in new target  
257 populations.

### 258 **Clinical and public health relevance**

259 The Latin American Society of Nephrology and Hypertension (Sociedad Latinoamericana de  
260 Nefrología e Hipertensión - SLANH) recommends to annually screen for CKD with several markers:  
261 blood pressure, serum creatinine, proteinuria and urinalysis.<sup>38</sup> The South African Renal Society  
262 (SARS) guidelines also recommend CKD screening annually, yet they focus on high-risk populations:  
263 people with diabetes, hypertension, or HIV.<sup>39</sup> This recommendation is endorsed by the Asian Forum  
264 for Chronic Kidney Disease Initiatives (AFCKDI), extending it to individuals  $\geq 65$  years, people  
265 consuming nephrotoxic substances, and those with family history of CKD and past history of acute  
266 kidney injury.<sup>40</sup> Although it seems reasonable to screen people with risk factors such as hypertension  
267 and diabetes, this approach may miss a large proportion of the high-risk population because they  
268 could be unaware of their condition.<sup>41 42</sup> In this case, risk scores could be useful because they can be  
269 applied to large populations regardless of whether they are aware of their hypertension or diabetes  
270 status. Unfortunately, our work would not support nor encourage the inclusion of available risk scores  
271 for CKD in clinical guidelines in LMIC. Instead, our results urgently call to improve risk prediction  
272 research in LMIC. Therefore, CKD risk scores could be included into clinical practice to identify high-  
273 risk individuals and to inform the patient's management plan as is the case in other fields such as  
274 cardiovascular primary prevention.

### 275 **Conclusions**

276 This systematic review of diagnostic and prognostic models of CKD did not find conclusive evidence  
277 to recommend the use of a single CKD score across LMIC. Nonetheless, we identified relevant efforts  
278 in Iran, India, Peru, South Africa, China and Thailand; these models would require further external  
279 validation before they can be applied in other LMIC. We encourage researchers and practitioners to  
280 develop and validate CKD risk scores, which are cost-efficient tools to early identify CKD prevalent  
281 and incident cases so that they can receive timely treatment.

282

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284 search. DJA-G and EJA wrote the manuscript. All authors approved the submitted version.

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288 **Patient consent for publication:** Not required.

289 **Data availability statement:** Data sharing not applicable as no data sets generated for this study.  
290 Given the nature of systematic reviews, the data set generated and analysed for the current study is  
291 already available. All studies analysed for the present review are referenced for readers.

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424

425 **TABLES:**

426 **Table 1. CHARMS criteria to define research question and strategy.**

Concept	Criteria
Prognostic or diagnostic?	Both - this review focused on diagnostic and prognostic risk scores for chronic kidney disease (CKD)
Scope	Diagnostic/prognostic models to inform physicians, researchers and the general population whether they are likely to have CKD (i.e., diagnostic) or will be likely to have CKD (i.e., prognostic)
Type of prediction modelling studies	<ul style="list-style-type: none"> <li>• Diagnostic/prognostic models with external validation</li> <li>• Diagnostic/prognostic models without external validation</li> <li>• Diagnostic/prognostic models validation</li> </ul>
Target population to whom the prediction model applies	General adult population in Low- and Middle-Income Countries (LMIC). No age or gender restrictions
Outcome to be predicted	CKD (diagnostic or prognostic)
Time span of prediction	Any, prognostic models will not be included/excluded based on the prediction time span
Intended moment of using the model	Diagnostic/prognostic models to be used in asymptomatic adults of LMIC to ascertain current CKD status or future risk of developing CKD. These models could be used for screening, treatment allocation in primary prevention, or research purposes

427 Based on the CHARMS checklist.<sup>14</sup>



428 Table 2. General characteristics.

Nº of report	Study	Country	Outcome prevalence (%)	Mean age (years)	Men (%)	Outcome details	Baseline sample size	Number of outcome events	Outcome events per candidate predictors
1	Asgari <i>et al</i> , 2020	Iran	6-years validation: 22.08 9-years validation: 41.94	6-years validation: 46.02 9-years validation: NI	6-years validation: 40.1 9-years validation: 40.6	CKD was defined as eGFR <60 mL/min/1.73 m <sup>2</sup> , provided by the MDRD formula	6-years validation: 3,270 9-years validation: 3,240	6-years validation: 722 9-years validation: 1,359	For every model validation: n/a
2	Bradshaw <i>et al</i> , 2019	India	For every model derivation: 10.89 For every model validation: NI	For every model derivation: 44.9 For every model validation: NI	For every model derivation: 46.8 For every model validation: NI	CKD was defined as an eGFR rate <60 mL/min/1.73 m <sup>2</sup> (estimated with the CKD-EPI equation) or UACR ≥30 mg/g	For every model derivation: 8,698 Urban model validation: 4,065 Rural model validation: 4,940	For every model derivation: 947 For every model validation: NI	Model 1 derivation: 31.6 Model 2 derivation: 41.2 Model 3a derivation: 135.3 Model 3b derivation: 118.4 For every model validation: n/a
3	Carrillo-Larco <i>et al</i> , 2017	Peru	For every model derivation: 3.42 For every model validation: NI	For every model derivation: 57.7 For every model validation: NI	For every model derivation: 49.4 For every model validation: NI	CKD was defined as eGFR <60 mL/min/1.73 m <sup>2</sup> , provided by the MDRD formula	For every model derivation: 2,368 For every model validation: NI	For every model derivation: 81 For every model validation: NI	Complete model derivation: 2.25 Lab-free model derivation: 3.1

			validation: 5.41	validation: 57.1	validation: 47.7		validation: 1,459	validation: 79	For every model validation: n/a
4	Mogueo <i>et al</i> , 2015	South Africa	For every eGFR model validation: 28.71  For every eGFR or proteinuria model validation: 29.71	For every model validation: 55	For every model validation: 23.4	CKD was defined as eGFR <60 mL/min/1.73 m <sup>2</sup> , provided by the 4- variable MDRD formula	For every model validation: 902	For every eGFR model validation: 259  For every eGFR or proteinuria model validation: 268	For every model validation: n/a
5	Saranburut <i>et al</i> , 2017 - Framingham Heart Study	Thailand	MDRD model validation: 10.37  CKD-EPI model validation: 10.01	MDRD model validation: 54.6  CKD-EPI model validation: 54.7	MDRD model validation: 70.8  CKD-EPI model validation: 71.5	MDRD model validation: CKD was defined as eGFR <60 mL/min/1.73 m <sup>2</sup> , provided by the MDRD formula  CKD-EPI model validation: CKD was defined as eGFR <60 mL/min/1.73 m <sup>2</sup> , provided by the CKD-EPI equation	MDRD model validation: 2,141  CKD-EPI model validation: 2,328	MDRD model validation: 222  CKD-EPI model validation: 233	For every model validation: n/a
6	Saranburut <i>et al</i> , 2017	Thailand	For every model derivation: 8.51  For every model validation: 1.94	For every model derivation: 51.3  For every model validation: 45.6	For every model derivation: 70.5  For every model validation: 70.5	CKD was defined as a preserved GFR (eGFR ≥60 mL/min/1.73m <sup>2</sup> ) at baseline and subsequently developed decreased GFR (eGFR <60 mL/min/1.73m <sup>2</sup> ) at the 10-year follow-up, provided by the Two- level Race Variable CKD-EPI equation (using the non-black coefficient)	For every model derivation: 3,186  For every model validation: 1,395	For every model derivation: 271  For every model validation: 27	Model 1 derivation: 18.1  Model 1 BMI derivation: 18.1  Model 2 derivation: 16.9

									<b>Model 3 derivation: 12.3</b>  <b>For every validation model: n/a</b>
7	Thakkinstian <i>et al</i> , 2011	Thailand	18.10	45.2	45.5	CKD was defined as a combination of stages I to V. CKD stage I & II was defined as eGFR $\geq 90$ and eGFR 60-89 ml/min/1.73 m <sup>2</sup> , respectively; with haematuria or UACR $\geq 30$ mg/g. CKD stage III, IV, and V was defined as eGFR 30-59, 15-29, and $< 15$ ml/min/1.73 m <sup>2</sup> , respectively; regardless of kidney damage (eGFR was calculated using the MDRD formula)	3,459	626	16.9
8	Wen <i>et al</i> , 2020	China	<b>For every derivation model: 18.06</b>	<b>For every derivation model: 50</b>	<b>For every derivation model: 44.7</b>	CKD was defined as an eGFR rate $< 60$ mL/min/1.73 m <sup>2</sup> (assessed with the modified Chinese MDRD equation) or UACR $\geq 30$ mg/g	<b>For every derivation model: 3,266</b>	<b>For every derivation model: 590</b>	<b>For every derivation model: NI</b>
9	Wu <i>et al</i> , 2016	China	<b>Model derivation: 2.05</b>  <b>Model validation: 1.10</b>	<b>Model derivation: 45.3</b>  <b>Model validation: 41.8</b>	<b>Model derivation: 56.7</b>  <b>Model validation: 63.7</b>	CKD was defined as eGFR $< 60$ mL/min/1.73 m <sup>2</sup> , provided by the CKD-EPI equation	<b>Model derivation: 14,374</b>  <b>Model validation: 4,371</b>	<b>Model derivation: 294</b>  <b>Model validation: 48</b>	<b>Model derivation: NI</b>  <b>Model validation: n/a</b>

429 CKD, chronic kidney disease; CKD-EPI, chronic kidney disease epidemiology collaboration; eGFR, estimated glomerular filtration rate; GFR, glomerular

430 filtration rate; KDIGO, MDRD, modification of diet renal disease; n/a, not applicable; NI, no information; UACR, urinary albumin-to-creatinine ratio.

431 Table 3. Performance metrics.

N <sup>o</sup>	Study	Discrimination (%)	Classification measures
1	Asgari <i>et al</i> , 2020	<p><b>6-years validation:</b> AUC (95% CI) for final intercept adjusted model = Male: 76 (72-79) and Female: 71 (69-73)</p> <p><b>9-years validation:</b> AUC (95% CI) for final intercept adjusted model = Male: 71 (67-74) and Female: 70 (68-73)</p>	<p><b>6-years validation:</b> For men at a cut-off of 25: sensitivity=72.7%; specificity=67.6%. For women at a cut-off of 19: sensitivity=66.8%; specificity=65.6%</p> <p><b>9-years validation:</b> For men at a cut-off of 25: sensitivity=64.5%; specificity=69.5%. For women at a cut-off of 23: sensitivity=56.9%; specificity=76.6%</p>
2	Bradshaw <i>et al</i> , 2019	<p><b>Model 1 derivation:</b> C-statistic (95% CI) = 79 (78-81)</p> <p><b>Model 2 derivation:</b> C-statistic (95% CI) = 73 (72-75)</p> <p><b>Model 3a derivation:</b> C-statistic (95% CI) = 77 (75-79)</p> <p><b>Model 3b derivation:</b> C-statistic (95% CI) = 77 (76-79)</p> <p><b>Urban validation:</b> C-statistic (95% CI) = 74 (73-74)</p> <p><b>Rural validation:</b> C-statistic (95% CI) = 70 (69-71)</p>	<p><b>Model 1 derivation:</b> At a cut-off of 0.09: sensitivity=72%; specificity=72%; positive predictive value=24%; negative predictive value=96%</p> <p><b>Model 2 derivation:</b> At a cut-off of 0.09: sensitivity=68%; specificity=67%; positive predictive value=20%; negative predictive value=95%</p> <p><b>Model 3a derivation:</b> At a cut-off of 0.09: sensitivity=71%; specificity=70%; positive predictive value=22%; negative predictive value=95%</p> <p><b>Model 3b derivation:</b> At a cut-off of 0.09: sensitivity=71%; specificity=70%; positive predictive value=22%; negative predictive value=95%</p> <p><b>Urban model validation:</b> NI</p> <p><b>Rural model validation:</b> NI</p>
3	Carrillo-Larco <i>et al</i> , 2017	<p><b>Complete model derivation:</b> AUC = 76.2</p> <p><b>Lab-free model derivation:</b> AUC = 76</p> <p><b>Complete model validation:</b> AUC = 70</p> <p><b>Lab-free model validation:</b> AUC = 70</p>	<p><b>Complete model derivation:</b> At a cut-off of 2: sensitivity=82.5%; specificity=70.0%; positive predictive value=8.8%; negative predictive value=99.1%; likelihood ratio positive=2.8; likelihood ratio negative=0.3</p> <p><b>Lab-free model derivation:</b> At a cut-off of 2: sensitivity=80%; specificity=72%; positive predictive value=9.1%; negative predictive value=99%; likelihood ratio positive=2.9; likelihood ratio negative=0.3</p>

			<p><b>Complete model validation:</b> At a cut-off of 2: sensitivity=70.5%; specificity=69.1%; positive predictive value=11.4%; negative predictive value=97.6%; likelihood ratio positive=2.3; likelihood ratio negative=0.4</p> <p><b>Lab-free model validation:</b> At a cut-off of 2: sensitivity=70.5%; specificity=69.7%; positive predictive value=11.6%; negative predictive value=97.7%; likelihood ratio positive=2.3; likelihood ratio negative=0.4</p>
4	Mogueo <i>et al</i> , 2015	<p><b>South Korean eGFR model validation:</b> C-statistic (95% CI) = 79.7 (76.5-82.9)</p> <p><b>Thai eGFR model validation:</b> C-statistic (95% CI) = 76 (72.6-79.3)</p> <p><b>South Korean eGFR or proteinuria model validation:</b> C-statistic (95% CI) = 81.1 (78.0-84.2)</p> <p><b>Thai eGFR or proteinuria model validation:</b> C-statistic (95% CI) = 77.2 (73.9-80.5)</p>	<p><b>South Korean eGFR model validation:</b> At a cut-off of 0.30: sensitivity=82%; specificity=67%</p> <p><b>Thai eGFR model validation:</b> At a cut-off of 0.31: sensitivity=73%; specificity=72%</p> <p><b>South Korean eGFR or proteinuria model validation:</b> At a cut-off of 0.31: sensitivity=84%; specificity=68%</p> <p><b>Thai eGFR or proteinuria model validation:</b> At a cut-off of 0.32: sensitivity=74%; specificity=73%</p>
5	Saranburut <i>et al</i> , 2017 - Framingham Heart Study	<p><b>MDRD model validation:</b> AUC (95% CI) = 69 (66-73)</p> <p><b>CKD-EPI model validation:</b> AUC (95% CI) = 63 (57-65)</p>	<p><b>MDRD model validation:</b> NI</p> <p><b>CKD-EPI model validation:</b> NI</p>
6	Saranburut <i>et al</i> , 2017 - Model 1 (derivation Clinical only)	<p><b>Model 1 derivation:</b> AUC (95% CI) = 72 (69-75)</p> <p><b>Model 1 BMI derivation:</b> AUC (95% CI) = 72 (69-75)</p> <p><b>Model 2 derivation:</b> AUC (95% CI) = 79 (76-82)</p> <p><b>Model 3 derivation:</b> AUC (95% CI) = 80 (77-82)</p> <p><b>Model 1 validation:</b> AUC (95% CI) = 66 (55-78)</p> <p><b>Model 2 validation:</b> AUC (95% CI) = 88 (80-95)</p>	<p><b>Model 1 derivation:</b> NI</p> <p><b>Model 1 BMI derivation:</b> NI</p> <p><b>Model 2 derivation:</b> NI</p> <p><b>Model 3 derivation:</b> NI</p> <p><b>Model 1 validation:</b> NI</p> <p><b>Model 2 validation:</b> NI</p>

7	Thakkinstian <i>et al</i> , 2011 (derivation)	C-statistic of internal validation = 74.1	At a cut-off of 5: sensitivity=76%; specificity=69%
8	Wen <i>et al</i> , 2020 - Simple Risk Score (derivation)	<b>Simple model derivation:</b> AUC (95% CI) = 71.7 (68.9-74.4) <b>Best-fit model derivation:</b> AUC (95% CI) = 72.1 (69.3-74.8)	<b>Simple model derivation:</b> At a cut-off of 14: sensitivity=70.5%; specificity=65.1%; positive predictive value=29.8%; negative predictive value=91.3%; likelihood ratio positive=2.0; likelihood ratio negative=0.5 <b>Best-fit model derivation:</b> At a cut-off of 24: sensitivity=56.8%; specificity=76.6%; positive predictive value=33.8%; negative predictive value=89.4%; likelihood ratio positive=2.4 likelihood ratio negative=0.6
9	Wu <i>et al</i> , 2016 (derivation)	<b>Model derivation:</b> AUC (95% CI) = 89.4 (86.1-92.6) <b>Model validation:</b> AUC (95% CI) = 88.0 (82.9-93.1)	<b>Model derivation:</b> At a cut-off of 36: sensitivity=82%; specificity=86.3% <b>Model validation:</b> NI

432 AUC, area under the curve; CI, confident interval; NI, no information.

433 **Table 4: Risk of bias assessment of individual diagnostic/prediction models**

Study	Objective	Risk of Bias (RoB)				Applicability			Overall	
		Participants	Predictors	Outcome	Analys s	Participants	Predictors	Outcome	RoB	Applicability
Asgari <i>et al</i> , 2020 European Risk Assessment tool (6-years)	Validation	+	+	?	-	+	+	+	-	+
Asgari <i>et al</i> , 2020 European Risk Assessment tool (9-years)	Validation	+	+	?	-	+	+	+	-	+
Bradshaw <i>et al</i> , 2019 - Model 1	Derivation	+	+	?	-	+	+	+	-	+
Bradshaw <i>et al</i> , 2019 - Model 2	Derivation	+	+	?	-	+	+	+	-	+
Bradshaw <i>et al</i> , 2019 - Model 3a	Derivation	+	+	?	-	+	+	+	-	+
Bradshaw <i>et al</i> , 2019 - Model 3b	Derivation	+	+	?	-	+	+	+	-	+
Bradshaw <i>et al</i> , 2019 - Model 3a (CARRS-I urban)	Validation	+	+	?	-	+	+	+	-	+
Bradshaw <i>et al</i> , 2019 - Model 3a (UDAY rural)	Validation	+	+	?	-	+	+	+	-	+
Carrillo-Larco <i>et al</i> , 2017 - CRONICAS-CKD (complete)	Derivation	+	+	+	-	+	+	+	-	+
Carrillo-Larco <i>et al</i> , 2017 - CRONICAS-CKD (lab-free)	Derivation	+	+	+	-	+	+	+	-	+
Carrillo-Larco <i>et al</i> , 2017 - CRONICAS-CKD (complete)	Validation	+	+	+	-	+	+	+	-	+

Carrillo-Larco <i>et al</i> , 2017 - CRONICAS-CKD (lab-free)	Validation	+	+	+	-	+	+	+	-	+
Mogueo <i>et al</i> , 2015 – South Korean model (eGFR)	Validation	+	+	?	-	+	+	+	-	+
Mogueo <i>et al</i> , 2015 - Thai model (eGFR)	Validation	+	+	?	-	+	+	+	-	+
Mogueo <i>et al</i> , 2015 – South Korean model (eGFR or proteinuria)	Validation	+	+	?	-	+	+	+	-	+
Mogueo <i>et al</i> , 2015 - Thai model (eGFR or proteinuria)	Validation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Framingham Heart Study (MDRD)	Validation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Framingham Heart Study (CKD-EPI)	Validation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Model 1 (Clinical only)	Derivation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Model 1 BMI (Clinical only)	Derivation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Model 2 (Clinical + Limited laboratory tests)	Derivation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Model 3 (Clinical + Full laboratory tests)	Derivation	+	+	?	-	+	+	+	-	+
Saranburut <i>et al</i> , 2017 - Model 1 (Clinical only)	Validation	+	+	?	-	+	+	+	-	+



Saranburut <i>et al</i> , 2017 - Model 2 (Clinical + Limited laboratory tests)	Validation	+	+	?	-	+	+	+	-	+
Thakkestian <i>et al</i> , 2011	Derivation	+	+	?	-	+	+	+	-	+
Wen <i>et al</i> , 2020 - Simple Risk Score	Derivation	+	+	?	-	+	+	+	-	+
Wen <i>et al</i> , 2020 - Best-fit Risk Score	Derivation	+	+	?	-	+	+	+	-	+
Wu <i>et al</i> , 2016	Derivation	+	+	?	-	+	+	+	-	+
Wu <i>et al</i> , 2016	Validation	+	+	?	-	+	+	+	-	+

434 PROBAST = Prediction model Risk of Bias ASsessment Tool;<sup>20 21</sup> RoB = risk of bias. + indicates low RoB/low concern regarding applicability; - indicates high  
435 RoB/high concern regarding applicability; and ? indicates unclear RoB/unclear concern regarding applicability.

436 **FIGURES**

437 **Figure 1. PRISMA 2020 flow diagram.**

438 **SUPPLEMENTARY MATERIAL**

439 **S1 Text. Protocol.**

440 **S1 Table. PRISMA 2020 checklist.**

441 **S2 Table. Search terms.**

442 **S3 Table. Data extraction form.**

443 **S4 Table. Risk of bias and applicability**

444 **S1 Figure. Countries where studies were conducted.** LMIC that developed and/or validated models  
445 included in this review (Green). Moreover, Asgari *et al*,<sup>26</sup> Mogueo *et al*<sup>25</sup> and Saranburut *et al*<sup>31</sup> validated  
446 risk models that were originally derived in the Netherlands, South Korea and the United States,  
447 respectively (Blue).

448 **S2 Figure. Predictors included in the final models.** The colours of the bars identify the underlying  
449 characteristic of predictors inherent to: the subject (purple), anthropometrics (blue), clinical assessment  
450 and history (green), and laboratory measures (yellow).

451

### Identification of studies via databases and registers

### Identification of studies via other methods

Identification

Records identified from Databases (n = 14,845):  
OVID (Medline, Embase, Global Health) (n = 11,026)  
Scopus (n = 1,654)  
Web of Science (n = 2,165)

Records removed before screening:  
Duplicate records removed (n = 1,462)  
Records marked as ineligible by automation tools (n = 0)

Records identified from:  
Citation searching (n = 1)

Screening

Records screened (n = 13,383)

Records excluded by manual screening (n = 13,372)

Reports sought for retrieval (n = 11)

Reports not retrieved (n = 1)

Reports sought for retrieval (n = 1)

Reports not retrieved (n = 0)

Reports assessed for eligibility (n = 10)

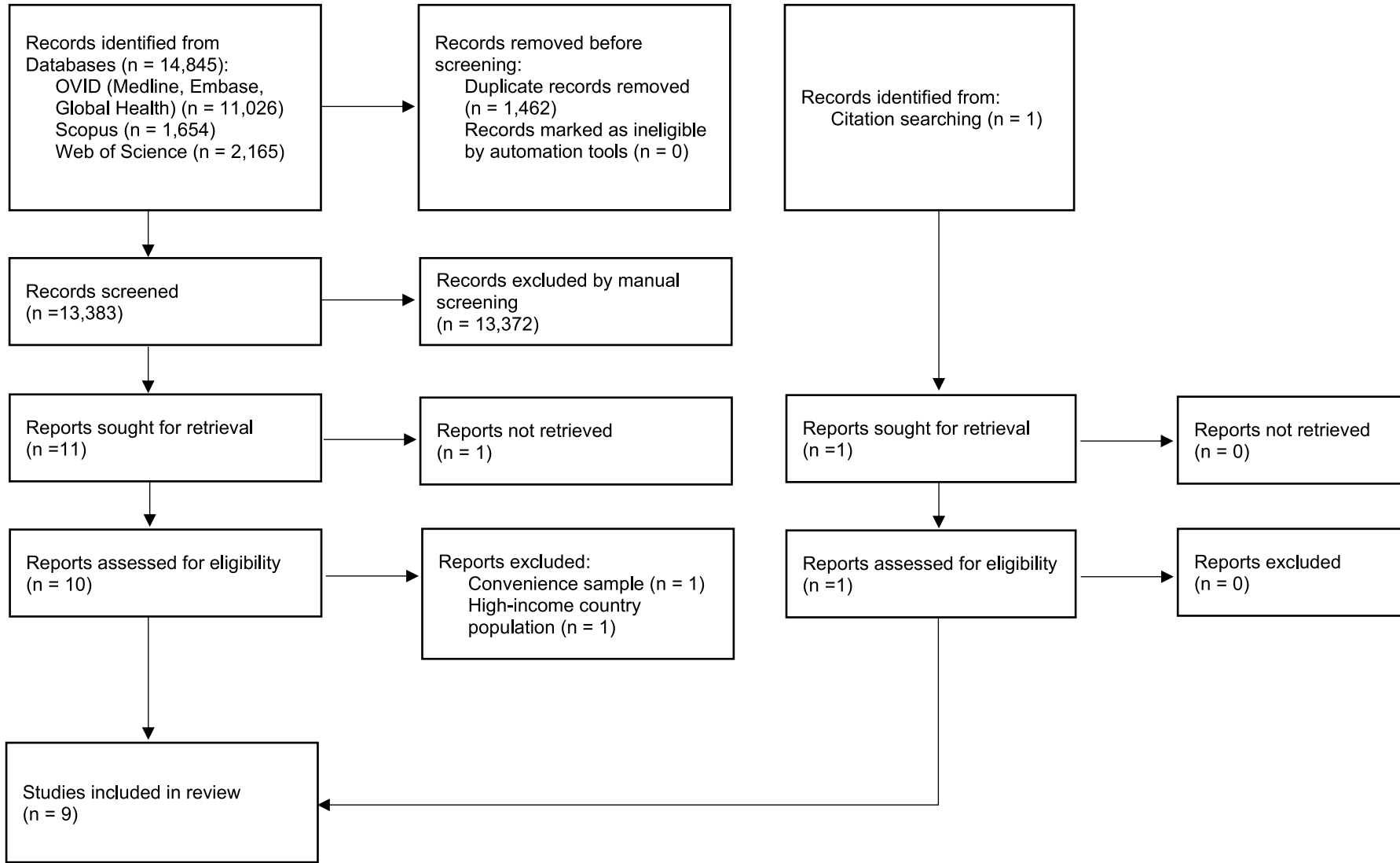
Reports excluded:  
Convenience sample (n = 1)  
High-income country population (n = 1)

Reports assessed for eligibility (n = 1)

Reports excluded (n = 0)

Included

Studies included in review (n = 9)



## **Supplementary Material**

- . A systematic review of diagnostic and prognostic models of Chronic kidney disease in Low- and Middle- Income Countries**

<b>S1 Text: Protocol (also available at <a href="https://doi.org/10.1101/2021.04.24.21256041">https://doi.org/10.1101/2021.04.24.21256041</a>)</b>	1
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**S1 Text: Protocol (also available at <https://doi.org/10.1101/2021.04.24.21256041>)**

**Chronic Kidney Disease in Low- and Middle- Income Countries: Protocol for a  
systematic review of diagnostic and prognostic models**

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**ABSTRACT**

**Background:** Chronic Kidney Disease (CKD) is a highly prevalent condition with a large disease burden globally. In low- and middle-income countries (LMIC) the CKD screening challenges the health system. This systematic and comprehensive search of all CKD diagnostic and prognostic models in LMIC will inform screening strategies in LMIC following a risk-based approach.

**Objective:** To summarize all multivariate diagnostic and prognostic models for CKD in adults in LMIC.

**Methods:** Systematic review. Without date or language restrictions we will search Embase, Medline, Global Health (these three through Ovid), SCOPUS and Web of Science. We seek multivariable diagnostic or prognostic models which included a random sample of the general population. We will screen titles and abstracts; we will then study the selected reports. Both phases will be done by two reviewers independently. Data extraction will be performed by two researchers independently using a pre-specified Excel form (CHARMS model). We will evaluate the risk of bias with the PROBAST tool.

**Conclusion:** This systematic review will provide the most comprehensive list and critical appraisal of diagnostic and prognostic models for CKD available for the general population in LMIC. This evidence could

inform policies and interventions to improve CKD screening in LMIC following a risk-based approach, maximizing limited resources and reaching populations with limited access to CKD screening tests. This systematic review will also reveal methodological limitations and research needs to improve CKD diagnostic and prognostic models in LMIC.

**Keywords:** Chronic Kidney Disease; Diagnostic Models; Prognostic Models; Low- and Middle-income countries.

## INTRODUCTION

Chronic kidney disease (CKD) is a highly prevalent condition that contributes to a large part of disease burden globally. Between 1990 and 2017, the health metrics of CKD showed a bleak profile: mortality rate, incidence and kidney transplantation rate increased by 2.8%, 29.3% and 34.4%, respectively.<sup>1</sup> CKD led to 1.2 million deaths in 2017 and in the best-case scenario, mortality is projected to increase to 2.2 million deaths<sup>2</sup> and become the 5th cause of years of life lost (YLL) by 2040.<sup>3</sup> Currently, 2.5 million of patients receive kidney transplantation therapy and it is projected to increase to 5.4 million by 2030.<sup>1</sup> CKD also reveals disparities between low- and middle-income countries (LMIC) and high income countries (HIC); for example, the age-standardised disability-adjusted life-year (DALY) rate due to CKD was the highest in LMIC between 1990-2017.<sup>4</sup> In LMIC, that remain as resource-constrained settings, there is a need for optimization of the CKD screening strategies which usually challenge the health system.<sup>5</sup>

Risk equations or risk scores are a cost-effective alternative for CKD screening.<sup>6</sup> These equations are less invasive and accepted by the general population;<sup>7</sup> also, they require less resources like laboratory tests.<sup>8</sup> Many scores were developed in high-income countries,<sup>9-11</sup> and they may not be used in LMIC because their accuracy is better where they have been developed.<sup>12</sup> Current strategies for CKD screening suggest studying people with risk factors (e.g. diabetes, hypertension).<sup>13-15</sup> These recommendations rely on studies where albuminuria and proteinuria were used as screening tools for identifying CKD patients.<sup>16</sup> Nevertheless, a systematic review found that using risk scores allows screening of a larger population and therefore can be useful for detecting more CKD cases.<sup>6</sup>

To date, there are no systematic reviews of diagnostic or prognostic models for CKD with a focus on LMIC.<sup>17, 18</sup> This limits our knowledge of what tools we have to enhance CKD screening in LMIC; similarly, this dearth of evidence prevents from planning future research to overcome the limitations of available models. This will be the first systematic review to fill these knowledge gaps in LMIC to improve and complement the CKD screening programmes in LMIC.

## METHODS

### Objective

To synthesise CKD diagnostic and prognostic models for the adult population of LMIC.

### Study design

This systematic review and meta-analysis will be conducted following the preferred reporting items for systematic reviews and meta-analyses (PRISMA) 2020 guidelines.<sup>19</sup> We will also adhere to the recommendations for systematic reviews of diagnostic and prognostic models following the CHARMS guidelines<sup>20</sup> and the PROBAST tool to assess risk of bias.<sup>21</sup>

### Eligibility criteria

*Participants/population:* We will include the general adult population (18 years and above) of LMIC with no gender restrictions. Studies following a population-based random sampling approach will be included. We will only include populations from LMIC according to The World Bank.<sup>22</sup> Conversely, studies with a study population of only patients (e.g., people with hypertension) or high-risk individuals (e.g., smokers) will be excluded. We will exclude studies with LMIC populations outside a LMIC.

*Intervention, exposure:* None (this review is looking at CKD diagnostic and prognostic models in LMIC).

*Comparator, control:* None (this review is looking at CKD diagnostic and prognostic models in LMIC).

*Outcome:* Diagnostic and prognostic models for CKD. The CKD diagnosis should have been based on a laboratory or imaging test including: urine albumin-creatinine ratio, urine protein-creatinine ratio, albumin excretion ratio, urine sediment, kidney images, kidney biopsy or the estimated glomerular filtration rate (eGFR). In other words, research in which CKD diagnosis was based on self-reported information only will not be considered. However, if a study combined both self-reported information and a laboratory or imaging tests, this will be included.

*Types of studies:* Studies with an observational design will be included, which encompasses cross-sectional (for diagnostic models) and prospective longitudinal studies (for prognostic models). If we retrieve any systematic review on this subject, we will revise its reference list to identify relevant original sources.

### **Literature Search and Data collation**

The search will be conducted in five search engines: Embase, Medline, Global Health (these three through Ovid), SCOPUS and Web of Science. No date or language restrictions will be set. The complete search strategy can be found in Supplementary Material.

Titles and abstracts will be screened by two researchers independently (DJA-G and EJA), looking for studies that meet the selection criteria above detailed. Full-text reports of the selected publications will be studied by two researchers independently (DJA-G and EJA). Discrepancies at any stage will be solved by consensus or by a third party (RMC-L).

During the full-text phase, if there are any original reports in which the population, methodology or results are not clear enough to assess the inclusion/exclusion criteria, we will contact the corresponding author by email. We will wait for two weeks, if we receive no answer and cannot solve our doubts through other means, this report will be excluded based on the lack of clarity to assess inclusion/exclusion criteria.

We will record the reasons for exclusion in the full-text phase and summarize the number of included/excluded reports following the PRISMA flow diagram.

### **Data extraction**

We will develop a data extraction form following the CHARMS recommendations.<sup>20</sup> Data extraction will be conducted by two researchers independently; discrepancies will be solved by consensus or by a third party (RMC-L).

### **Risk of bias of individual studies**

The risk of bias assessment of individual reports will be conducted using the Prediction model Risk Of Bias ASsessment Tool (PROBAST) tool.<sup>21</sup>

### **Statistical Analysis**

A qualitative synthesis is planned, whereby we will narratively synthesise the findings from the selected studies. We will summarize the key elements from each report such as study design, study population and characteristics of the study population. Also, we will summarize the key features of the risk scores as provided by each report, including discrimination, calibration, sensitivity, specificity, and predictive values. A quantitative synthesis will be carried out if the included studies are found to be sufficiently homogenous and we have at least four original reports.

### **Ethics**

This review did not directly include human subjects. We considered this work as 'low risk' and did not request approval by an Ethics Committee. Results and opinions included in this protocol, and those included in the final report, are the author's alone and do not represent those of the institutions to which they belong.



## CONCLUSIONS

This systematic review will provide a comprehensive list of diagnostic and prognostic models for CKD for people in LMIC, along with their accuracy metrics. Currently, information lacks in LMIC where diagnostic and prognostic models could inform CKD screening strategies. Similarly, this work will elucidate the limitations of available diagnostic and prognostic models for CKD in LMIC, so that future research can be planned accordingly to overcome these caveats and deliver robust models to advance

CKD screening strategies in LMIC.

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**S1 Table: PRISMA Checklist**

Section and Topic	Item #	Checklist item	Location where item is reported
<b>TITLE</b>			
Title	1	Identify the report as a systematic review.	page 01
<b>ABSTRACT</b>			
Abstract	2	See the PRISMA 2020 for Abstracts checklist.	page 02
<b>INTRODUCTION</b>			
Rationale	3	Describe the rationale for the review in the context of existing knowledge.	page 03
Objectives	4	Provide an explicit statement of the objective(s) or question(s) the review addresses.	page 04
<b>METHODS</b>			
Eligibility criteria	5	Specify the inclusion and exclusion criteria for the review and how studies were grouped for the syntheses.	page 04-05
Information sources	6	Specify all databases, registers, websites, organisations, reference lists and other sources searched or consulted to identify studies. Specify the date when each source was last searched or consulted.	page 04

Search strategy	7	Present the full search strategies for all databases, registers and websites, including any filters and limits used.	supplementary page 03-07
Selection process	8	Specify the methods used to decide whether a study met the inclusion criteria of the review, including how many reviewers screened each record and each report retrieved, whether they worked independently, and if applicable, details of automation tools used in the process.	page 05
Data collection process	9	Specify the methods used to collect data from reports, including how many reviewers collected data from each report, whether they worked independently, any processes for obtaining or confirming data from study investigators, and if applicable, details of automation tools used in the process.	page 05-06
Data items	10a	List and define all outcomes for which data were sought. Specify whether all results that were compatible with each outcome domain in each study were sought (e.g. for all measures, time points, analyses), and if not, the methods used to decide which results to collect.	page 04-05, table 1
	10b	List and define all other variables for which data were sought (e.g. participant and intervention characteristics, funding sources). Describe any assumptions made about any missing or unclear information.	page 04-05, table 1
Study risk of bias assessment	11	Specify the methods used to assess risk of bias in the included studies, including details of the tool(s) used, how many reviewers assessed each study and whether they worked independently, and if applicable, details of automation tools used in the process.	page 06
Effect measures	12	Specify for each outcome the effect measure(s) (e.g. risk ratio, mean difference) used in the synthesis or presentation of results.	NA
Synthesis methods	13a	Describe the processes used to decide which studies were eligible for each synthesis (e.g. tabulating the study intervention characteristics and comparing against the planned groups for each synthesis (item #5)).	page 06
	13b	Describe any methods required to prepare the data for presentation or synthesis, such as handling of missing summary statistics, or data conversions.	NA
	13c	Describe any methods used to tabulate or visually display results of individual studies and syntheses.	page 06

	13d	Describe any methods used to synthesize results and provide a rationale for the choice(s). If meta-analysis was performed, describe the model(s), method(s) to identify the presence and extent of statistical heterogeneity, and software package(s) used.	page 06
	13e	Describe any methods used to explore possible causes of heterogeneity among study results (e.g. subgroup analysis, meta-regression).	NA
	13f	Describe any sensitivity analyses conducted to assess robustness of the synthesized results.	NA
Reporting bias assessment	14	Describe any methods used to assess risk of bias due to missing results in a synthesis (arising from reporting biases).	NA
Certainty assessment	15	Describe any methods used to assess certainty (or confidence) in the body of evidence for an outcome.	page 11
<b>RESULTS</b>			
Study selection	16a	Describe the results of the search and selection process, from the number of records identified in the search to the number of studies included in the review, ideally using a flow diagram.	page 06-07
	16b	Cite studies that might appear to meet the inclusion criteria, but which were excluded, and explain why they were excluded.	page 06-07
Study characteristics	17	Cite each included study and present its characteristics.	page 08-09
Risk of bias in studies	18	Present assessments of risk of bias for each included study.	page 11, supplementary page 39-45

Results of individual studies	19	For all outcomes, present, for each study: (a) summary statistics for each group (where appropriate) and (b) an effect estimate and its precision (e.g. confidence/credible interval), ideally using structured tables or plots.	page 9-11
Results of syntheses	20a	For each synthesis, briefly summarise the characteristics and risk of bias among contributing studies.	page 9-11
	20b	Present results of all statistical syntheses conducted. If meta-analysis was done, present for each the summary estimate and its precision (e.g. confidence/credible interval) and measures of statistical heterogeneity. If comparing groups, describe the direction of the effect.	table 3
	20c	Present results of all investigations of possible causes of heterogeneity among study results.	NA
	20d	Present results of all sensitivity analyses conducted to assess the robustness of the synthesized results.	NA
Reporting biases	21	Present assessments of risk of bias due to missing results (arising from reporting biases) for each synthesis assessed.	NA
Certainty of evidence	22	Present assessments of certainty (or confidence) in the body of evidence for each outcome assessed.	page 11
<b>DISCUSSION</b>			
Discussion	23a	Provide a general interpretation of the results in the context of other evidence.	page 11
	23b	Discuss any limitations of the evidence included in the review.	page 11-13
	23c	Discuss any limitations of the review processes used.	page 11-13

	23d	Discuss implications of the results for practice, policy, and future research.	page 14-15
<b>OTHER INFORMATION</b>			
Registration and protocol	24a	Provide registration information for the review, including register name and registration number, or state that the review was not registered.	page 04
	24b	Indicate where the review protocol can be accessed, or state that a protocol was not prepared.	page 04
	24c	Describe and explain any amendments to information provided at registration or in the protocol.	NA
Support	25	Describe sources of financial or non-financial support for the review, and the role of the funders or sponsors in the review.	page 01
Competing interests	26	Declare any competing interests of review authors.	page 01
Availability of data, code and other materials	27	Report which of the following are publicly available and where they can be found: template data collection forms; data extracted from included studies; data used for all analyses; analytic code; any other materials used in the review.	page 15

NA: Not applicable

## S2 Table: Search terms

### S2.1 Table: Embase, Medline and Global Health (OVID)

01	chronic renal insufficiency.mp.
02	chronic kidney disease.mp.
03	chronic kidney failure.mp.
04	CKD.mp.
05	exp Renal Insufficiency, Chronic/
06	(chronic adj2 kidney adj2 disease).mp.
07	(chronic adj2 kidney adj2 failure).mp.
08	chronic renal failure.mp.
09	chronic renal disease.mp.
10	chronic kidney insufficiency.mp.
11	end stage renal disease.mp.
12	ESRD.mp.
13	kidney function.mp.
14	renal function.mp.
15	kidney dysfunction.mp.
16	renal dysfunction.mp.
17	01 or 02 or 03 or 04 or 05 or 06 or 07 or 08 or 09 or 10 or 11 or 12 or 13 or 14 or 15 or 16
18	(("Afghanistan") or ("Benin") or ("Burkina Faso") or ("Burundi") or ("Central African Republic") or ("Chad") or ("Comoros") or ("Democratic Republic of the Congo") or ("Eritrea") or ("Ethiopia") or ("Gambia") or ("Guinea") or ("Guinea-Bissau") or ("Haiti") or ("Democratic People's Republic of Korea") or ("Liberia") or ("Madagascar") or ("Malawi") or ("Mali") or ("Mozambique") or ("Nepal") or ("Niger") or ("Rwanda") or ("Senegal") or ("Sierra Leone") or ("Somalia") or ("South Sudan") or ("Tanzania") or ("Togo") or ("Uganda") or ("Zimbabwe") or ("Armenia") or ("Bangladesh") or ("Bhutan") or ("Bolivia") or ("Cape Verde") or ("Cambodia") or ("Cameroon") or ("Congo") or ("Cote d'Ivoire") or ("Djibouti") or ("Egypt") or ("El Salvador") or ("Ghana") or ("Guatemala") or ("Honduras") or ("India") or ("Indonesia") or ("Kenya") or ("Micronesia") or ("Kosovo") or ("Kyrgyzstan") or ("Laos") or ("Lesotho") or ("Mauritania") or ("Moldova") or ("Mongolia") or ("Morocco") or ("Myanmar") or ("Nicaragua") or ("Nigeria") or ("Pakistan") or ("Papua New Guinea") or ("Philippines") or ("Samoa") or ("Atlantic Islands") or ("Melanesia") or ("Sri Lanka") or ("Sudan") or ("Swaziland") or ("Syria") or ("Tajikistan") or ("Timor-Leste") or ("Tonga") or ("Tunisia") or ("Ukraine") or ("Uzbekistan") or ("Vanuatu") or ("Vietnam") or ("Middle East") or ("Yemen") or ("Zambia") or ("Albania") or ("Algeria") or ("American Samoa") or ("Angola") or ("Argentina") or ("Azerbaijan") or ("Republic of Belarus") or ("Belize") or ("Bosnia and Herzegovina") or ("Botswana") or ("Brazil") or ("Bulgaria") or ("China") or ("Colombia") or ("Costa Rica") or ("Cuba") or ("Dominica") or ("Dominican Republic") or ("Equatorial Guinea") or ("Ecuador") or ("Fiji") or ("Gabon") or ("Georgia") or ("Grenada") or ("Guyana") or ("Iran") or ("Iraq") or ("Jamaica") or ("Jordan") or ("Kazakhstan") or ("Lebanon") or ("Libya") or ("Macedonia") or ("Malaysia") or ("Indian Ocean Islands") or ("Mexico") or ("Montenegro") or ("Namibia") or ("Palau") or ("Panama") or ("Paraguay") or ("Peru") or ("Russia") or ("Serbia") or ("South Africa") or ("Saint Lucia") or ("Saint Vincent and the Grenadines") or ("Suriname") or ("Thailand") or ("Turkey") or ("Turkmenistan") or ("Venezuela") or (developing countr*) or (lowincome countr*) or (middle-income countr*) or (low-middle income countr*) or (upper-middle income countr*))
19	risk assessment.mp.
20	risk functions.mp.
21	Risk Assessment/mt
22	risk equation\$.mp.
23	risk chart?.mp.
24	(risk adj3 tool\$.mp.
25	risk assessment function?.mp.
26	risk assessor.mp.
27	risk appraisal\$.mp.
28	risk calculation\$.mp.
29	risk calculator\$.mp.



30	risk factor\$ calculator\$.mp.
31	risk factor\$ calculation\$.mp.
32	risk engine\$.mp.
33	risk equation\$.mp.
34	risk table\$.mp.
35	risk threshold\$.mp.
36	risk disc?.mp.
37	risk disk?.mp.
38	risk scoring method?.mp.
39	scoring scheme?.mp.
40	risk scoring system?.mp.
41	risk scal\$.mp.
42	risk prediction?.mp.
43	risk algorith\$.mp.
44	prediction model\$.mp.
45	predictive instrument?.mp.
46	project\$ risk?.mp.
47	predictive model?.mp.
48	scoring method\$.mp.
49	(prediction\$ adj3 method\$.mp.
50	exp Risk Assessment/
51	(risk? adj1 assess\$.mp.
52	screening.mp.
53	diagnostic test.mp.
54	19 or 20 or 21 or 22 or 23 or 24 or 25 or 26 or 27 or 28 or 29 or 30 or 31 or 32 or 33 or 34 or 35 or 36 or 37 or 38 or 39 or 40 or 41 or 42 or 43 or 44 or 45 or 46 or 47 or 48 or 49 or 50 or 51 or 52 or 53
55	17 and 18 and 54
56	exp animals/ not humans.sh.
57	55 not 56
58	Remove duplicates from 57

## S2.2 Table: SCOPUS

((TITLE-ABS-KEY("Afghanistan") OR TITLE-ABS-KEY("Benin") OR TITLE-ABS-KEY("Burkina Faso") OR TITLE-ABS-KEY("Burundi") OR TITLE-ABS-KEY("Central African Republic") OR TITLE-ABS-KEY("Chad") OR TITLE-ABS-KEY("Comoros") OR TITLE-ABS-KEY("Democratic Republic of the Congo") OR TITLE-ABS-KEY("Eritrea") OR TITLE-ABS-KEY("Ethiopia") OR TITLE-ABS-KEY("Gambia") OR TITLE-ABS-KEY("Guinea") OR TITLE-ABS-KEY("Guinea-Bissau") OR TITLE-ABS-KEY("Haiti") OR TITLE-ABS-KEY("Democratic People's Republic of Korea") OR TITLE-ABS-KEY("Liberia") OR TITLE-ABS-KEY("Madagascar") OR TITLE-ABS-KEY("Malawi") OR TITLE-ABS-KEY("Mali") OR TITLE-ABS-KEY("Mozambique") OR TITLE-ABS-KEY("Nepal") OR TITLE-ABS-KEY("Niger") OR TITLE-ABS-KEY("Rwanda") OR TITLE-ABS-KEY("Senegal") OR TITLE-ABS-KEY("Sierra Leone") OR TITLE-ABS-KEY("Somalia") OR TITLE-ABS-KEY("South Sudan") OR TITLE-ABS-KEY("Tanzania") OR TITLE-ABS-KEY("Togo") OR TITLE-ABS-KEY("Uganda") OR TITLE-ABS-KEY("Zimbabwe") OR TITLE-ABS-KEY("Armenia") OR TITLE-ABS-KEY("Bangladesh") OR TITLE-ABS-KEY("Bhutan") OR TITLE-ABS-KEY("Bolivia") OR TITLE-ABS-KEY("Cape Verde") OR TITLE-ABS-KEY("Cambodia") OR TITLE-ABS-KEY("Cameroon") OR TITLE-ABS-KEY("Congo") OR TITLE-ABS-KEY("Cote d'Ivoire") OR TITLE-ABS-KEY("Djibouti") OR TITLE-ABS-KEY("Bolivia") OR TITLE-ABS-KEY("Cape Verde") OR TITLE-ABS-KEY("Cambodia") OR TITLE-ABS-KEY("Cameroon") OR TITLE-ABS-KEY("Congo") OR TITLE-ABS-KEY("Cote d'Ivoire") OR TITLE-ABS-KEY("Djibouti") OR TITLE-ABS-KEY("Egypt") OR TITLE-ABS-KEY("El Salvador") OR TITLE-ABS-KEY("Ghana") OR TITLE-ABS-KEY("Guatemala") OR TITLE-ABS-KEY("Honduras") OR TITLE-ABS-KEY("India") OR TITLE-ABS-KEY("Indonesia") OR TITLE-ABS-KEY("Kenya") OR TITLE-ABS-KEY("Micronesia") OR TITLE-ABS-KEY("Kosovo") OR TITLE-ABS-KEY("Kyrgyzstan") OR TITLE-ABS-KEY("Laos") OR TITLE-ABS-KEY("Lesotho") OR TITLE-ABS-KEY("Mauritania") OR TITLE-ABS-KEY("Moldova") OR TITLE-ABS-KEY("Mongolia") OR TITLE-ABS-KEY("Morocco") OR TITLE-

ABS-KEY("Myanmar") OR TITLE-ABS-KEY("Nicaragua") OR TITLE-ABS-KEY("Nigeria") OR TITLE-ABS-KEY("Pakistan") OR TITLE-ABS-KEY("Papua New Guinea") OR TITLE-ABSKEY("Philippines") OR TITLE-ABS-KEY("Samoa") OR TITLE-ABS-KEY("Atlantic Islands") OR TITLE-ABSKEY("Melanesia") OR TITLE-ABS-KEY("Sri Lanka") OR TITLE-ABS-KEY("Sudan") OR TITLE-ABSKEY("Swaziland") OR TITLE-ABS-KEY("Syria") OR TITLE-ABS-KEY("Tajikistan") OR TITLE-ABSKEY("Timor-Leste") OR TITLE-ABS-KEY("Tonga") OR TITLE-ABS-KEY("Tunisia") OR TITLE-ABSKEY("Ukraine") OR TITLE-ABS-KEY("Uzbekistan") OR TITLE-ABS-KEY("Vanuatu") OR TITLE-ABSKEY("Vietnam") OR TITLE-ABS-KEY("Middle East") OR TITLE-ABS-KEY("Yemen") OR TITLE-ABS-KEY("Zambia") OR TITLE-ABS-KEY("Albania") OR TITLE-ABS-KEY("Algeria") OR TITLE-ABSKEY("American Samoa") OR TITLE-ABS-KEY("Angola") OR TITLE-ABS-KEY("Argentina") OR TITLE-ABSKEY("Azerbaijan") OR TITLE-ABS-KEY("Republic of Belarus") OR TITLE-ABS-KEY("Belize") OR TITLE-ABSKEY("Bosnia and Herzegovina") OR TITLE-ABS-KEY("Botswana") OR TITLE-ABS-KEY("Brazil") OR TITLEABS-KEY("Bulgaria") OR TITLE-ABS-KEY("China") OR TITLE-ABS-KEY("Colombia") OR TITLE-ABS-KEY("Costa Rica") OR TITLE-ABS-KEY("Cuba") OR TITLE-ABS-KEY("Dominica") OR TITLE-ABSKEY("Dominican Republic") OR TITLE-ABS-KEY("Equatorial Guinea") OR TITLE-ABS-KEY("Ecuador") OR TITLE-ABS-KEY("Fiji") OR TITLE-ABS-KEY("Gabon") OR TITLE-ABS-KEY("Georgia") OR TITLE-ABSKEY("Grenada") OR TITLE-ABS-KEY("Guyana") OR TITLE-ABS-KEY("Iran") OR TITLE-ABS-KEY("Iraq") OR TITLE-ABS-KEY("Jamaica") OR TITLE-ABS-KEY("Jordan") OR TITLE-ABS-KEY("Kazakhstan") OR TITLEABS-KEY("Lebanon") OR TITLE-ABS-KEY("Libya") OR TITLE-ABS-KEY("Macedonia (Republic)") OR TITLEABS-KEY("Malaysia") OR TITLE-ABS-KEY("Indian Ocean Islands") OR TITLE-ABS-KEY("Mexico") OR TITLE-ABS-KEY("Montenegro") OR TITLE-ABS-KEY("Namibia") OR TITLE-ABS-KEY("Palau") OR TITLEABS-KEY("Panama") OR TITLE-ABS-KEY("Paraguay") OR TITLE-ABS-KEY("Peru") OR TITLE-ABSKEY("Russia") OR TITLE-ABS-KEY("Serbia") OR TITLE-ABS-KEY("South Africa") OR TITLE-ABS-KEY("Saint Lucia") OR TITLE-ABS-KEY("Saint Vincent and the Grenadines") OR TITLE-ABS-KEY("Suriname") OR TITLE-ABS-KEY("Thailand") OR TITLE-ABS-KEY("Turkey") OR TITLE-ABS-KEY("Turkmenistan") OR TITLEABS-KEY("Venezuela") OR TITLE-ABS-KEY(developing countr\*) OR TITLE-ABS-KEY(lowincome countr\*) OR TITLE-ABS-KEY(middle-income countr\*) OR TITLE-ABS-KEY(low-middle income countr\*) OR TITLEABS-KEY(upper-middle income countr\*) OR TITLE-ABS-KEY("low resource") OR TITLE-ABS-KEY ("underresourced") OR TITLE-ABS-KEY("resource poor") OR TITLE-ABS-KEY("under-developed") OR TITLE-ABSKEY("underdeveloped") OR TITLE-ABS-KEY("developing world") OR TITLE-ABS-KEY("third world") OR TITLE-ABS-KEY(lmic) OR TITLE-ABS-KEY(low AND middle AND income) AND (TITLE-ABS-KEY(Risk Assessment) OR TITLE-ABS-KEY(risk? adj1 assess\*) OR TITLE-ABS-KEY(risk function) OR TITLE-ABS-KEY(Risk Assessment) OR TITLE-ABS-KEY(risk functions) OR TITLE-ABS-KEY(risk equation\*) OR TITLEABS-KEY(risk chart?) OR TITLE-ABS-KEY(risk adj3 tool\*) OR TITLE-ABS-KEY(risk assessment function?) OR TITLE-ABS-KEY(risk assessor) OR TITLE-ABS-KEY(risk appraisal\*) OR TITLE-ABS-KEY(risk calculation\*) OR TITLE-ABS-KEY(risk calculator\*) OR TITLE-ABS-KEY(risk factor\* calculator\*) OR TITLEABS-KEY(risk factor\* calculation\*) OR TITLE-ABS-KEY(risk engine\*) OR TITLE-ABS-KEY(risk equation\*) OR TITLE-ABS-KEY(risk table\*) OR TITLE-ABS-KEY(risk threshold\*) OR TITLE-ABS-KEY(risk disc?) OR TITLE-ABS-KEY(risk disk?) OR TITLE-ABS-KEY(risk scoring method?) OR TITLE-ABS-KEY(scoring scheme?) OR TITLE-ABS-KEY(risk scoring system?) OR TITLE-ABS-KEY(risk prediction?) OR TITLE-ABSKEY(risk algorith\*) OR TITLE-ABS-KEY(prediction model\*) OR TITLE-ABS-KEY(predictive instrument?) OR TITLE-ABS-KEY(project\* risk?) OR TITLE-ABS-KEY(predictive model?) OR TITLE-ABS-KEY(scoring method\*) OR TITLE-ABS-KEY(prediction\* adj3 method\*) OR TITLE-ABS-KEY(screening) OR TITLE-ABSKEY(risk scal\*) OR TITLE-ABS-KEY(diagnostic test)) AND (TITLE-ABS-KEY(chronic renal insufficiency) OR TITLE-ABS-KEY(chronic kidney disease) OR TITLE-ABS-KEY(chronic kidney failure) OR TITLE-ABS-KEY(CKD) OR TITLE-ABS-KEY(chronic renal failure) OR TITLE-ABS-KEY(chronic renal disease) OR TITLE-ABS-KEY(chronic kidney insufficiency) OR TITLE-ABS-KEY(end stage renal disease) OR TITLE-ABSKEY(ESRD) OR TITLE-ABS-KEY(kidney function) OR TITLE-ABS-KEY(renal function) OR TITLE-ABSKEY(kidney dysfunction) OR TITLE-ABS-KEY(renal dysfunction) OR TITLE-ABS-KEY(chronic W/2 kidney W/2 disease) OR TITLE- ABS-KEY(chronic W/2 kidney W/2 failure) AND NOT DBCOLL(medl))

### S2.3 Table: WEB OF SCIENCE

((chronic renal insufficiency) OR (chronic kidney disease) OR (chronic kidney failure) OR (CKD) OR (Renal Insufficiency, Chronic) OR (chronic NEAR/2 kidney NEAR/2 disease) OR (chronic NEAR/2 kidney NEAR/2 failure) OR (chronic renal failure) OR (chronic renal disease) OR (chronic kidney insufficiency) OR (end stage renal disease) OR (ESRD) OR (kidney function) OR (renal function) OR (kidney dysfunction) OR (renal dysfunction)) AND (("Afghanistan") OR ("Benin") OR ("Burkina Faso") OR ("Burundi") OR ("Central African Republic") OR ("Chad") OR ("Comoros") OR ("Democratic Republic of the Congo") OR ("Eritrea") OR ("Ethiopia") OR ("Gambia") OR ("Guinea") OR ("Guinea-Bissau") OR ("Haiti") OR ("Democratic People's Republic of Korea") OR ("Liberia") OR ("Madagascar") OR ("Malawi") OR ("Mali") OR ("Mozambique") OR ("Nepal") OR ("Niger") OR ("Rwanda") OR ("Senegal") OR ("Sierra Leone") OR ("Somalia") OR ("South Sudan") OR ("Tanzania") OR ("Togo") OR ("Uganda") OR ("Zimbabwe") OR ("Armenia") OR ("Bangladesh") OR ("Bhutan") OR ("Bolivia") OR ("Cape Verde") OR ("Cambodia") OR ("Cameroon") OR ("Congo") OR ("Cote d'Ivoire") OR ("Djibouti") OR ("Egypt") OR ("El Salvador") OR ("Ghana") OR ("Guatemala") OR ("Honduras") OR ("India") OR ("Indonesia") OR ("Kenya") OR ("Micronesia") OR ("Kosovo") OR ("Kyrgyzstan") OR ("Laos") OR ("Lesotho") OR ("Mauritania") OR ("Moldova") OR ("Mongolia") OR ("Morocco") OR ("Myanmar") OR ("Nicaragua") OR ("Nigeria") OR ("Pakistan") OR ("Papua New Guinea") OR ("Philippines") OR ("Samoa") OR ("Atlantic Islands") OR ("Melanesia") OR ("Sri Lanka") OR ("Sudan") OR ("Swaziland") OR ("Syria") OR ("Tajikistan") OR ("Timor-Leste") OR ("Tonga") OR ("Tunisia") OR ("Ukraine") OR ("Uzbekistan") OR ("Vanuatu") OR ("Vietnam") OR ("Middle East") OR ("Yemen") OR ("Zambia") OR ("Albania") OR ("Algeria") OR ("American Samoa") OR ("Angola") OR ("Argentina") OR ("Azerbaijan") OR ("Republic of Belarus") OR ("Belize") OR ("Bosnia and Herzegovina") OR ("Botswana") OR ("Brazil") OR ("Bulgaria") OR ("China") OR ("Colombia") OR ("Costa Rica") OR ("Cuba") OR ("Dominica") OR ("Dominican Republic") OR ("Equatorial Guinea") OR ("Ecuador") OR ("Fiji") OR ("Gabon") OR ("Georgia") OR ("Grenada") OR ("Guyana") OR ("Iran") OR ("Iraq") OR ("Jamaica") OR ("Jordan") OR ("Kazakhstan") OR ("Lebanon") OR ("Libya") OR ("Macedonia (Republic) ") OR ("Malaysia") OR ("Indian Ocean Islands") OR ("Mexico") OR ("Montenegro") OR ("Namibia") OR ("Palau") OR ("Panama") OR ("Paraguay") OR ("Peru") OR ("Russia") OR ("Serbia") OR ("South Africa") OR ("Saint Lucia") OR ("Saint Vincent and the Grenadines") OR ("Suriname") OR ("Thailand") OR ("Turkey") OR ("Turkmenistan") OR ("Venezuela") OR (developing countr) OR (lowincome countr\*) OR (middle-income countr\*) OR (low-middle income countr\*) OR (upper-middle income countr\*)) AND ((risk assessment) OR (risk equation\$) OR (risk chart?) OR (risk NEAR/3 tool\$) OR (risk assessment function?) OR (risk assessor) OR (risk appraisal\$) OR (risk calculation\$) OR (risk calculator\$) OR (risk factor\$ calculation\$) OR (risk engine\$) OR (risk equation\$) OR (risk table\$) OR (risk threshold\$) OR (risk disc?) OR (risk disk?) OR (risk scoring method?) OR (scoring scheme?) OR (risk scoring system?) OR (risk scal\$) OR (risk prediction?) OR (risk algorithm\$) OR (prediction model\$) OR (predictive instrument?) OR (project\$ risk?) OR (predictive model?) OR (scoring method\$) OR (prediction\$ NEAR/3 method\$) OR (risk? NEAR/1 assess\$) OR (screening) OR (diagnostic test))) NOT ((animal\*) OR ("not humans"))

**S3 Table: Data extraction form (by chapters)**

**S3.1 Table: Source of data and participants**

		Source of data	Participants									
N°	Study	Source of data	Participant location	Baseline year	End year (cohorts)	Sampling	Inclusion criteria	Exclusion criteria	Outcome prevalence (%)	Outcome incidence (for cohorts)	Baseline mean age	Baseline % men
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	Cohort	Community	1999-2005	2011	Random	Tehran lipids and glucose study (TLGS) cohort participants.	Persons with prevalent Cardiovascular Disease (CVD), Type 2 Diabetes Mellitus or End-stage Renal Disease with (eGFR) <15 mL/min/1.73 m <sup>2</sup> . Also excluded those with missing data at baseline for creatinine (Cr), fasting plasma glucose (FPG), 2-hour postchallenge plasma	46.02 (11.95)	40.1%	58.34	29.53

								glucose (2 h-PCG), body mass index (BMI), waist circumference (WC) and smoking status as well as participants with missing data during follow-up on Cr, FPG, 2 h-PCG and CVD status				
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	Cohort	Community	1999-2005	2009-2018	Random	Tehran lipids and glucose study (TLGS) cohort participants.	Persons with prevalent Cardiovascular Disease (CVD), Type 2 Diabetes Mellitus or End-stage Renal Disease with (eGFR) <15 mL/min/1.73 m <sup>2</sup> . Also excluded those with missing data at baseline for creatinine (Cr), fasting plasma glucose (FPG), 2-hour postchallenge plasma glucose (2 h-PCG), body mass index (BMI), waist circumference (WC) and smoking status as well as participants with missing data during follow-up on Cr, FPG, 2 h-PCG and CVD status	NI	40.6%	48.20	49.70
2	Bradshaw, 2019 - Model 1 (derivation)	Cross-sectional	Community	2015	n/a	Random	Any individual aged ≥20 years and permanently residing in at Delhi and Chennai (CARRS-II). A permanent resident was defined as a person living in the selected household, was related to the household head and ate at least 3 meals in a week with the family.	Bedridden individuals, pregnant women, participants with missing both or either serum creatinine or urine albumin-to-creatinine ratio data and participants on dialysis.	44.9 (13.5)	46.8%	48.20	49.70

							Households were defined as “a group of people who live together, usually pool their income and eat at least one meal together a day when they are at home. This does not include people who have migrated permanently or are considered visitors”					
2	Bradshaw, 2019 - Model 2 (derivation)	Cross-sectional	Community	2015	n/a	Random	Any individual aged $\geq 20$ years and permanently residing in at Delhi and Chennai (CARRS-II). A permanent resident was defined as a person living in the selected household, was related to the household head and ate at least 3 meals in a week with the family. Households were defined as “a group of people who live together, usually pool their income and eat at least one meal together a day when they are at home. This does not include people who have migrated permanently or are considered visitors”	Bedridden individuals, pregnant women, participants with missing both or either serum creatinine or urine albumin-to-creatinine ratio data and participants on dialysis.	44.9 (13.5)	46.8%	48.20	49.70
2	Bradshaw, 2019 - Model 3a (derivation)	Cross-sectional	Community	2015	n/a	Random	Any individual aged $\geq 20$ years and permanently residing in at Delhi and Chennai (CARRS-II). A permanent resident was defined as a person living in the selected household,	Bedridden individuals, pregnant women, participants with missing both or either serum creatinine or urine albumin-to-creatinine ratio data and participants on dialysis.	44.9 (13.5)	46.8%	39.90	46.97

							was related to the household head and ate at least 3 meals in a week with the family. Households were defined as “a group of people who live together, usually pool their income and eat at least one meal together a day when they are at home. This does not include people who have migrated permanently or are considered visitors”					
2	Bradshaw, 2019 - Model 3b (derivation)	Cross-sectional	Community	2015	n/a	Random	Any individual aged ≥20 years and permanently residing in at Delhi and Chennai (CARRS-II). A permanent resident was defined as a person living in the selected household, was related to the household head and ate at least 3 meals in a week with the family. Households were defined as “a group of people who live together, usually pool their income and eat at least one meal together a day when they are at home. This does not include people who have migrated permanently or are considered visitors”	Bedridden individuals, pregnant women, participants with missing both or either serum creatinine or urine albumin-to-creatinine ratio data and participants on dialysis.	44.9 (13.5)	46.8%	39.90	46.97
2	Bradshaw, 2019 - Model	Cross-sectional	Community	2010-2012	n/a	Random	Any individual aged ≥20 years and permanently residing in at Delhi	Bedridden individuals, pregnant women, participants with missing	NI	NI	47.20	38.00

	3a (CARRS-I urban validation)						(CARRS-I). A permanent resident was defined as a person living in the selected household, was related to the household head and ate at least 3 meals in a week with the family. Households were defined as “a group of people who live together, usually pool their income and eat at least one meal together a day when they are at home. This does not include people who have migrated permanently or are considered visitors”	both or either serum creatinine or urine albumin-to-creatinine ratio data and participants on dialysis.				
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	Cross-sectional	Community	2014	n/a	Random	UDAY cohort participants ((a) adults aged ≥30 years residing in the sampled urban and rural areas of Sonipat and Vizag, respectively; and (b) willing to participate and provide informed consent).	Participants with missing both or either serum creatinine or urine albumin-to-creatinine ratio data, unwilling to provide informed consent, with serious chronic illnesses [such as that of the liver (cirrhosis), kidneys (renal failure) or malignancies], and pregnant women.	NI	NI	47.20	38.00
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation)	Cross-sectional	Community	2013-2014	n/a	Random	Full time resident, capable of giving informed consent, one subject per household.	Being pregnant, having active pulmonary tuberculosis, and having any disability preventing from undergoing anthropometric assessments, having CKD, missing values in the prediction variables, missing	57.7 (12.4)	49.4%		



	complete)							values in key variables to calculate eGFR, subjects with BMI >40 kg/m <sup>2</sup> or BMI <18.5 kg/m <sup>2</sup> .				
3	Carrillo-Larco, 2017 - CRONI CAS-CKD (derivation lab-free)	Cross-sectional	Community	2013-2014	n/a	Random	Full time resident, capable of giving informed consent, one subject per household.	Being pregnant, having active pulmonary tuberculosis, and having any disability preventing from undergoing anthropometric assessments, having CKD, missing values in the prediction variables, missing values in key variables to calculate eGFR, subjects with BMI >40 kg/m <sup>2</sup> or BMI <18.5 kg/m <sup>2</sup> .	57.7 (12.4)	49.4%		
3	Carrillo-Larco, 2017 - CRONI CAS-CKD (validation complete)	Cross-sectional	Community	2004-2006	n/a	Random	PREVENCION cohort participants.	Report having CKD, missing values in key variables to calculate eGFR, subjects with BMI >40 kg/m <sup>2</sup> or BMI <18.5 kg/m <sup>2</sup> , age < 35 years, missing values in prediction variables.	57.1 (12.6)	47.7%		
3	Carrillo-Larco, 2017 - CRONI CAS-CKD (validation lab-free)	Cross-sectional	Community	2004-2006	n/a	Random	PREVENCION cohort participants.	Report having CKD, missing values in key variables to calculate eGFR, subjects with BMI >40 kg/m <sup>2</sup> or BMI <18.5 kg/m <sup>2</sup> , age < 35 years, missing values in prediction variables.	57.1 (12.6)	47.7%		
4	Moguelo, 2015 -	Cross-sectional	Community	2008-2011	n/a	Random	Cape Town Bellville-South study cohort participants.	Participants with missing data on all variables, except	55 (15)	23.4%		

	Korean model (eGFR validation)							anaemia				
4	Mogueo, 2015 - Thai model (eGFR validation)	Cross-sectional	Community	2008-2011	n/a	Random	Cape Town Bellville-South study cohort participants.	Participants with missing data on all variables, except kidney stones	55 (15)	23.4%		
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Cross-sectional	Community	2008-2011	n/a	Random	Cape Town Bellville-South study cohort participants.	Participants with missing data on all variables, except anaemia	55 (15)	23.4%		
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Cross-sectional	Community	2008-2011	n/a	Random	Cape Town Bellville-South study cohort participants.	Participants with missing data on all variables, except kidney stones	55 (15)	23.4%		
5	Saranbut, 2017 - Framingham Heart	Cohort	Community	2002	2012	Random	Employees of the Electric Generating Authority of Thailand (EGAT) who participated in a health survey in 2002	Subjects who had CKD at baseline or did not have serum creatinine at baseline or at follow-up.	54.6 (5.6)	70.8%		

	Study (MDRD validation)												
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Cohort	Community	2002	2012	Random	Employees of the Electric Generating Authority of Thailand (EGAT) who participated in a health survey in 2003	Subjects who had CKD at baseline or did not have serum creatinine at baseline or at follow-up.	54.7 (5.7)	71.5%			
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	Cohort	Community	2002-2003	2012-2013	Random	EGAT 1-2 cohort participants with preserved GFR (estimate glomerular filtration rate (eGFR) $\geq$ 60 mL/min/1.73m <sup>2</sup> ) at baseline who attended both the examinations (EGAT 1 5rd examination and EGAT 2 4nd examination).	Patients who died, retired, moved, did not want to participate or had with missing baseline serum creatinine data. Also, patients with eGFR<60 at baseline in 2002-2003 were excluded	51.3 (7.4)	70.5%			
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Cohort	Community	2002-2003	2012-2013	Random	EGAT 1-2 cohort participants with preserved GFR (estimate glomerular filtration rate (eGFR) $\geq$ 60 mL/min/1.73m <sup>2</sup> ) at baseline who attended both the examinations (EGAT 1 5rd examination and EGAT 2 4nd examination).	Patients who died, retired, moved, did not want to participate or had with missing baseline serum creatinine data. Also, patients with eGFR<60 at baseline in 2002-2003 were excluded	51.3 (7.4)	70.5%			

6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Cohort	Community	2002-2003	2012-2013	Random	EGAT 1-2 cohort participants with preserved GFR (estimate glomerular filtration rate (eGFR) $\geq$ 60 mL/min/1.73m <sup>2</sup> ) at baseline who attended both the examinations (EGAT 1 5rd examination and EGAT 2 4nd examination).	Patients who died, retired, moved, did not want to participate or had with missing baseline serum creatinine data. Also, patients with eGFR<60 at baseline in 2002-2003 were excluded	51.3 (7.4)	70.5%		
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Cohort	Community	2002-2003	2012-2013	Random	EGAT 1-2 cohort participants with preserved GFR (estimate glomerular filtration rate (eGFR) $\geq$ 60 mL/min/1.73m <sup>2</sup> ) at baseline who attended both the examinations (EGAT 1 5rd examination and EGAT 2 4nd examination).	Patients who died, retired, moved, did not want to participate or had with missing baseline serum creatinine data. Also, patients with eGFR<60 at baseline in 2002-2003 were excluded	51.3 (7.4)	70.5%		
6	Saranburut, 2017 - Model 1 (validation Clinical only)	Cohort	Community	2009	2014	Random	EGAT 3 cohort participants with preserved GFR (eGFR $\geq$ 60) at baseline in 2009 (EGAT 3 1st examination) who were followed up 5 years later in 2014 (EGAT 3 2nd examination).	Participants younger than 40 years old at baseline, with missing serum creatinine values, participants who died, retired and moved, unwilling to participate and with an eGFR <60 at baseline.	45.6 (4.2)	75.5%		
6	Saranburut, 2017 - Model 2 (validation Clinical +	Cohort	Community	2009	2014	Random	EGAT 3 cohort participants with preserved GFR (eGFR $\geq$ 60) at baseline in 2009 (EGAT 3 1st examination) who were followed up 5 years later in 2014 (EGAT 3 2nd examination).	Participants younger than 40 years old at baseline, with missing serum creatinine values, participants who died, retired and moved, unwilling to participate and with an eGFR <60 at baseline.	45.6 (4.2)	75.5%		

	Limited laboratory tests)												
7	Thakkinstian, 2011 (derivation)	Cross-sectional	Community	2007-2008	n/a	Random	Global Screening and Early Evaluation of Kidney Disease (SEEK) study subjects: being 18 years or older, had no menstruation period for at least a week prior to the examination date if women, and whom were willing participants of the study and provided signed consent forms.	Subjects without blood or urine specimens.	45.2 (0.79)	45.5%			
8	Wen, 2020 - Simple Risk Score (derivation)	Cohort	Community	2006-2007	2012-2013	Random	Handan Eye Study (HES) participants (rural residents aged ≥30 years old living in Yongnian County).	Subjects who were diagnosed with CKD, unwilling to participate, missing follow up data (eGFR or UACR).	50 (10)	44.7%			
8	Wen, 2020 - Best-fit Risk Score (derivation)	Cohort	Community	2006-2007	2012-2013	Random	Handan Eye Study (HES) participants (rural residents aged ≥30 years old living in Yongnian County).	Subjects who were diagnosed with CKD, unwilling to participate, missing follow up data (eGFR or UACR).	50 (10)	44.7%			
9	Wu, 2016 (derivation)	Cross-sectional	Community	2012	n/a	Random	Adults older than 18 years and having given consent to this study.	Participants without: age information; body mass index (BMI) information; blood pressure (BP) measurement; serum creatinine test.	45.3 (14.3)	56.7%			
9	Wu, 2016	Cross-sectional	Community	2012	n/a	Random	Adults older than 18 years and having given consent to this study.	Participants without: age information; body mass index (BMI) information;	41.8 (11.7)	63.7%			

	(validation)							blood pressure (BP) measurement; serum creatinine test.				
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### S3.2 Table:: Outcome

		Outcome					
N°	Study	Outcome	Outcome details	Same outcome definition for all patients?	Blinded outcome	Predictors part of the outcome	Mean follow-up (years) (cohorts)
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	CKD composite	CKD was defined as eGFR < 60 mL/min/1.73 m <sup>2</sup> , provided by the Modification of Diet in Renal Disease (MDRD).	Yes	NI	No	6.2
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	CKD composite	CKD was defined as eGFR < 60 mL/min/1.73 m <sup>2</sup> , provided by the Modification of Diet in Renal Disease (MDRD).	Yes	NI	No	9.2
2	Bradshaw, 2019 - Model 1 (derivation)	CKD composite	CKD was defined as an eGFR rate <60 mL/min/1.73 m <sup>2</sup> (estimated by the CKD-EPI equation) or UACR ≥30 mg/g	Yes	NI	No	0
2	Bradshaw, 2019 - Model 2 (derivation)	CKD composite	CKD was defined as an eGFR rate <60 mL/min/1.73 m <sup>2</sup> (estimated by the CKD-EPI equation) or UACR ≥30 mg/g	Yes	NI	No	0
2	Bradshaw, 2019 - Model 3a (derivation)	CKD composite	CKD was defined as an eGFR rate <60 mL/min/1.73 m <sup>2</sup> (estimated by the CKD-EPI equation) or UACR ≥30 mg/g	Yes	NI	No	0
2	Bradshaw, 2019 - Model 3b (derivation)	CKD composite	CKD was defined as an eGFR rate <60 mL/min/1.73 m <sup>2</sup> (estimated by the CKD-EPI equation) or UACR ≥30 mg/g	Yes	NI	No	0
2	Bradshaw, 2019 - Model 3a	CKD composite	CKD was defined as an eGFR rate <60 mL/min/1.73 m <sup>2</sup> (estimated by the CKD-EPI equation) or UACR ≥30 mg/g	Yes	NI	No	0

	(CARRS-I urban validation)						
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	CKD composite	CKD was defined as an eGFR rate <60 mL/min/1.73 m2 (estimated by the CKD-EPI equation) or UACR ≥30 mg/g	Yes	NI	No	0
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	CKD composite	CKD defined as an eGFR <60 mL/min/1.73m2, using the MDRD (Modification of Diet in Renal Disease) formula, also known as CKD stage III	Yes	Yes	No	0
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	CKD composite	CKD defined as an eGFR <60 mL/min/1.73m2, using the MDRD (Modification of Diet in Renal Disease) formula, also known as CKD stage III	Yes	Yes	No	0
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	CKD composite	CKD defined as an eGFR <60 mL/min/1.73m2, using the MDRD (Modification of Diet in Renal Disease) formula, also known as CKD stage III	Yes	Yes	No	0
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	CKD composite	CKD defined as an eGFR <60 mL/min/1.73m2, using the MDRD (Modification of Diet in Renal Disease) formula, also known as CKD stage III	Yes	Yes	No	0
4	Mogueo, 2015 - Korean model (eGFR validation)	CKD composite	eGFR <60 ml/min/1.73 m2 based on the 4-variable Modification of Diet in Renal Disease (MDRD) formula	Yes	NI	No	0
4	Mogueo, 2015 - Thai model (eGFR validation)	CKD composite	eGFR <60 ml/min/1.73 m2 based on the 4-variable Modification of Diet in Renal Disease (MDRD) formula	Yes	NI	No	0
4	Mogueo, 2015 - Korean model (eGFR or	CKD composite	eGFR <60 ml/min/1.73 m2 based on the 4-variable Modification of Diet in Renal Disease (MDRD) formula and 'any nephropathy' including any of the stages I to V of the Kidney Disease: Improving Global Outcomes Chronic	Yes	NI	No	0



	proteinuria validation)		Kidney Disease (KDIGO) classification				
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	CKD composite	eGFR <60 ml/min/1.73 m <sup>2</sup> based on the 4-variable Modification of Diet in Renal Disease (MDRD) formula and 'any nephropathy' including any of the stages I to V of the Kidney Disease: Improving Global Outcomes Chronic Kidney Disease (KDIGO) classification	Yes	NI	No	0
5	Saranburut, 2017 - Framingham Heart Study (MDRD validation)	CKD composite	CKD was defined as estimate glomerular filtration rate (eGFR) <60 mL/min/1.73 m <sup>2</sup> using the Modification of Diet in Renal Disease (MDRD)	Yes	NI	No	10
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	CKD composite	CKD defined as (eGFR) <60 mL/min/1.73 m <sup>2</sup> using the CKD-EPI equation.	Yes	NI	No	10
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	CKD composite	Preserved GFR (eGFR ≥60) at baseline and subsequently developed decreased GFR (eGFR < 60 mL/min/1.73m <sup>2</sup> ) at the 10 year follow-up calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation using the non-black coefficient. The outcome is a modification from the KDIGO definition of CKD stage 3-5	Yes	NI	No	10
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	CKD composite	Preserved GFR (eGFR ≥60) at baseline and subsequently developed decreased GFR (eGFR < 60 mL/min/1.73m <sup>2</sup> ) at the 10 year follow-up calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation using the non-black coefficient. The outcome is a modification from the KDIGO definition of CKD stage 3-5	Yes	NI	No	10
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	CKD composite	Preserved GFR (eGFR ≥60) at baseline and subsequently developed decreased GFR (eGFR < 60 mL/min/1.73m <sup>2</sup> ) at the 10 year follow-up calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation using the non-black	Yes	NI	No	10

			coefficient. The outcome is a modification from the KDIGO definition of CKD stage 3-5				
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	CKD composite	Preserved GFR (eGFR $\geq 60$ ) at baseline and subsequently developed decreased GFR (eGFR $< 60$ mL/min/1.73m <sup>2</sup> ) at the 10 year follow-up calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation using the non-black coefficient. The outcome is a modification from the KDIGO definition of CKD stage 3-5	Yes	NI	No	10
6	Saranburut, 2017 - Model 1 (validation Clinical only)	CKD composite	Preserved GFR (eGFR $\geq 60$ ) at baseline and subsequently developed decreased GFR (eGFR $< 60$ mL/min/1.73m <sup>2</sup> ) at the 10 year follow-up calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation using the non-black coefficient. The outcome is a modification from the KDIGO definition of CKD stage 3-5	Yes	NI	No	5
6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	CKD composite	Preserved GFR (eGFR $\geq 60$ ) at baseline and subsequently developed decreased GFR (eGFR $< 60$ mL/min/1.73m <sup>2</sup> ) at the 10 year follow-up calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation using the non-black coefficient. The outcome is a modification from the KDIGO definition of CKD stage 3-5	Yes	NI	No	5
7	Thakkinstian, 2011 (derivation)	CKD composite	CKD was defined as stage I & II if GFR $\geq 90$ and GFR 60-89 ml/min/1.73 m <sup>2</sup> with haematuria and/or albumin-creatinine ratio 30 mg/g or greater, stage III, IV, and V if the GFR of 30-59, 15-29, and $< 15$ ml/min/1.73 m <sup>2</sup> respectively, regardless of kidney damage. eGFR was calculated using the MDRD equation for IDMS traceable serum creatinine values.	Yes	NI	No	0
8	Wen, 2020 - Simple Risk Score (derivation)	CKD composite	CKD was defined as an eGFR rate $< 60$ mL/min/1.73 m <sup>2</sup> ((assessed by the modified Chinese MDRD equation) or UACR $\geq 30$ mg/g	Yes	NI	No	5.6
8	Wen, 2020 - Best-fit Risk Score (derivation)	CKD composite	CKD was defined as an eGFR rate $< 60$ mL/min/1.73 m <sup>2</sup> ((assessed by the modified Chinese MDRD equation) or UACR $\geq 30$ mg/g	Yes	NI	No	5.6

9	Wu, 2016 (derivation)	CKD composite	Reduced eGFR was defined as eGFR<60 mL/min/1.73 m2 using the CKD-EPI equation.	Yes	NI	No	0
9	Wu, 2016 (validation)	CKD composite	Reduced eGFR was defined as eGFR<60 mL/min/1.73 m2 using the CKD-EPI equation.	Yes	NI	No	0

CKD, chronic kidney disease; CKD-EPI, chronic kidney disease epidemiology collaboration; eGFR, estimated glomerular filtration rate; GFR, glomerular filtration rate; KDIGO, MDRD, modification of diet renal disease; n/a, not applicable; NI, no information; UACR, urinary albumin-to-creatinine ratio.

**S3.3 Table: Candidate predictors**

		Candidate Predictors						
N°	Study	Number of candidate predictors	Number of predictors in the final model	Predictors timing	List of predictors in the final model	Predictors definition	Predictors ascertainment	Predictors modelling
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	n/a	18	NI	Age; BMI (body mass index); waist circumference; use of antihypertensives; current smoking, parent and/or sibling with myocardial infarction or stroke; parent and/or sibling with diabetes.	Age (<45, ≥45 to <50, ≥50 to <55, ≥55 to <60, ≥60 to <65, ≥65 to <70, ≥70 to <75, ≥75 to <85); Body mass index (<25, ≥25 to <30, ≥30); Waist circumference [<94, ≥94 to <102, ≥102 (for men) and <80, ≥80 to <88, ≥88 (for women)]; use of antihypertensive medications; current smoking ('who smokes cigarettes daily or occasionally'); family history of cardiovascular disease (CVD) and/or diabetes (previously diagnosed CVD in first-degree male and female relatives aged < 55 and < 65 years, respectively)	BMI was calculated as weight (kg) divided by height (m <sup>2</sup> ). Data collected by trained interviewer using a standard questionnaire	n/a
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	n/a	18	NI	Age; BMI (body mass index); waist circumference; use of antihypertensives; current smoking, parent and/or sibling with myocardial infarction or stroke; parent and/or sibling with diabetes.	Age (<45, ≥45 to <50, ≥50 to <55, ≥55 to <60, ≥60 to <65, ≥65 to <70, ≥70 to <75, ≥75 to <85); Body mass index (<25, ≥25 to <30, ≥30); Waist circumference [<94, ≥94 to <102, ≥102 (for men) and <80, ≥80 to <88, ≥88 (for women)]; use of antihypertensive medications; current smoking ('who smokes cigarettes daily	BMI was calculated as weight (kg) divided by height (m <sup>2</sup> ). Data collected by trained interviewer using a standard questionnaire	n/a

						or occasionally'); family history of cardiovascular disease (CVD) and/or diabetes (previously diagnosed CVD in first-degree male and female relatives aged < 55 and < 65 years, respectively)		
2	Bradshaw, 2019 - Model 1 (derivation )	30	NI	NI	NI	NI	NI	All continuous variables used cubic spline terms with knots placed at fixed quantiles of the predictor's marginal distribution, categorical variables were summarized using percentages and counts.
2	Bradshaw, 2019 - Model 2 (derivation )	23	NI	NI	NI	NI	NI	All continuous variables used cubic spline terms with knots placed at fixed quantiles of the predictor's marginal

									distribution, categorical variables were summarized using percentages and counts.
2	Bradshaw, 2019 - Model 3a (derivation )	NI	NI	NI	NI	NI	NI	NI	All continuous variables used cubic spline terms with knots placed at fixed quantiles of the predictor's marginal distribution, categorical variables were summarized using percentages and counts.
2	Bradshaw, 2019 - Model 3b (derivation )	8	NI	NI	NI	NI	NI	NI	All continuous variables used cubic spline terms with knots placed at fixed quantiles of the

									predictor's marginal distribution, categorical variables were summarized using percentages and counts.
2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	n/a	NI	NI	NI	NI	NI	NI	n/a
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	n/a	NI	NI	NI	NI	NI	NI	n/a
3	Carrillo-Larco, 2017 - CRONICA S-CKD (derivation complete)	36	7	NI	Age; hypertension; anemia.	Age (< 50, 50-69, ≥ 70 years), hypertension (blood pressure ≥ 140/90 mmHg OR previous diagnosis of hypertension and currently under treatment) and anemia (haemoglobin < 13 g/dL if male and < 12 g/dL if female).	Age (information was collected by trained fieldworkers through face-to-face interviews), hypertension (blood pressure measurements were conducted according to the recommendations of the 7th Joint National Committee on the diagnosis and management of High Blood Pressure in adults (JNC-7), NI on anemia.	NI	
3	Carrillo-Larco, 2017 -	26	5	NI	Age; hypertension.	Age (< 50, 50-69, ≥ 70 years), hypertension (blood pressure ≥ 140/90 mmHg OR previous diagnosis of	Age (information was collected by trained fieldworkers through face-to-	NI	

	CRONICA S-CKD (derivation lab-free)					hypertension and currently under treatment).	face interviews), hypertension (blood pressure measurements were conducted according to the recommendations of the 7th Joint National Committee on the diagnosis and management of High Blood Pressure in adults (JNC-7), NI on anemia.	
3	Carrillo-Larco, 2017 - CRONICA S-CKD (validation complete)	n/a	7	NI	Age; hypertension; anemia.	Age (< 50, 50-69, ≥ 70 years), hypertension (blood pressure ≥ 140/90 mmHg OR previous diagnosis of hypertension and currently under treatment) and anemia (haemoglobin < 13 g/dL if male and < 12 g/dL if female).	Age (information was collected by trained fieldworkers through face-to-face interviews), hypertension (blood pressure measurements were conducted according to the recommendations of the 7th Joint National Committee on the diagnosis and management of High Blood Pressure in adults (JNC-7), NI on anemia.	n/a
3	Carrillo-Larco, 2017 - CRONICA S-CKD (validation lab-free)	n/a	5	NI	Age; hypertension.	Age (< 50, 50-69, ≥ 70 years), hypertension (blood pressure ≥ 140/90 mmHg OR previous diagnosis of hypertension and currently under treatment).	Age (information was collected by trained fieldworkers through face-to-face interviews), hypertension (blood pressure measurements were conducted according to the recommendations of the 7th Joint National Committee on the diagnosis and management of High Blood Pressure in adults (JNC-7), NI on anemia.	n/a
4	Mogueo, 2015 -	n/a	8	NI	Age; sex; diabetes mellitus; hypertension; use	Age (50-59, 60-69, ≥70); Female gender; Hypertension (history of	Participants received a standardized interview (Age	NI



	Korean model (eGFR validation)				of statins; proteinuria	illness, taking antihyper-tensive drug(s) or had systolic blood pressure $\geq 140$ mmHg or diastolic blood pressure $\geq 90$ mmHg); Diabetes (history of illness, taking oral hypoglycaemic agents or fasting plasma glucose levels $\geq 126$ mg/dL); Use of statins; Proteinuria	and sex) and physical examination during which blood pressure was measured according to the World Health Organisation (WHO) guidelines using a semi-automated digital blood pressure monitor (Rossmax PA, USA) on the right arm in the sitting position. Participants with no history of doctor diagnosed diabetes mellitus underwent a 75 g oral glucose tolerance test (OGTT) as recommended by the WHO	
4	Mogueo, 2015 - Thai model (eGFR validation)	n/a	8	NI	Age; diabetes mellitus; hypertension	Age (<40, 40-59, 60-69, >70); Hypertension (history of illness, taking antihyper-tensive drug(s) or had systolic blood pressure $\geq 140$ mmHg or diastolic blood pressure $\geq 90$ mmHg); Diabetes (history of illness, taking oral hypoglycaemic agents or fasting plasma glucose levels $\geq 126$ mg/dL)	Participants received a standardized interview (Age) and physical examination during which blood pressure was measured according to the World Health Organisation (WHO) guidelines using a semi-automated digital blood pressure monitor (Rossmax PA, USA) on the right arm in the sitting position. Participants with no history of doctor diagnosed diabetes mellitus underwent a 75 g oral glucose tolerance test (OGTT) as recommended by the WHO	NI
4	Mogueo, 2015 - Korean model (eGFR or	n/a	8	NI	Age; sex; diabetes mellitus; hypertension; use of statins; proteinuria	Age (50-59, 60-69, $\geq 70$ ); Female gender; Hypertension (history of illness, taking antihyper-tensive drug(s) or had systolic blood pressure $\geq 140$ mmHg or diastolic blood pressure	Participants received a standardized interview (Age and sex) and physical examination during which blood pressure was measured	NI

	proteinuria validation)					≥90 mmHg); Diabetes (history of illness, taking oral hypoglycaemic agents or fasting plasma glucose levels ≥126 mg/dL); Use of statins; Proteinuria	according to the World Health Organisation (WHO) guidelines using a semi-automated digital blood pressure monitor (Rossmax PA, USA) on the right arm in the sitting position. Participants with no history of doctor diagnosed diabetes mellitus underwent a 75 g oral glucose tolerance test (OGTT) as recommended by the WHO	
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	n/a	8	NI	Age; diabetes mellitus; hypertension	Age (<40, 40-59, 60-69, >70); Hypertension (history of illness, taking antihypertensive drug(s) or had systolic blood pressure ≥140 mmHg or diastolic blood pressure ≥90 mmHg); Diabetes (history of illness, taking oral hypoglycaemic agents or fasting plasma glucose levels ≥126 mg/dL)	Participants received a standardized interview (Age) and physical examination during which blood pressure was measured according to the World Health Organisation (WHO) guidelines using a semi-automated digital blood pressure monitor (Rossmax PA, USA) on the right arm in the sitting position. Participants with no history of doctor diagnosed diabetes mellitus underwent a 75 g oral glucose tolerance test (OGTT) as recommended by the WHO	NI
5	Saranburu t, 2017 - Framingham Heart Study (MDRD validation)	n/a	5	NI	Diabetes mellitus; hypertension; eGFR category	Diabetes mellitus (yes); hypertension (yes); eGFR category (60-74, 75-89, 90-119)	Hypertension was defined as systolic blood pressure ≥ 140 mmHg or diastolic blood pressure ≥ 90 mmHg or use of oral antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of ≥126 mg/dl or use	n/a

							of medications. eGFR was estimated using the Modification of Diet in Renal Disease (MDRD) equation.	
5	Saranburu t, 2017 - Framingham Heart Study (CKD-EPI validation)	n/a	16	NI	Age; diabetes mellitus; hypertension; eGFR category	Age (30-34, 35-39, 40-44, 45-49, 50-54, 55-59, 60-64, 65-69, 70-74, 75-79, 80-85); diabetes mellitus (yes); hypertension (yes); eGFR category (60-74, 75-89, 90-119)	Age was obtained by a survey. Hypertension was defined as systolic blood pressure $\geq$ 140 mmHg or diastolic blood pressure $\geq$ 90 mmHg or use of oral antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of $\geq$ 126 mg/dl or use of medications. eGFR was estimated using the chronic kidney disease–epidemiology collaboration (CKD-EPI) equation	n/a
6	Saranburu t, 2017 - Model 1 (derivation Clinical only)	15	15	NI	Age; sex; systolic blood pressure; waist circumference; diabetes mellitus	Age (<45, 45-54, 55-59, $\geq$ 55); Sex (male, female); Waist circumference ( $\leq$ 80 for male or $\leq$ 90 for female or $>$ 80 for male); Diabetes (yes, no); Systolic blood pressure (<120, 120-129, 130-139, 140-149, 150-159, $\geq$ 160)	Age (health survey), sex (health survey). Hypertension was defined as systolic blood pressure $\geq$ 140 mmHg or diastolic blood pressure $\geq$ 90 mmHg or use of oral antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of $\geq$ 126 mg/dl or a positive history of diabetes. Waist circumference was measured midway between the lowest ribs and the iliac crest.	NI
6	Saranburu t, 2017 - Model 1 BMI (derivation)	15	15	NI	Age; sex; systolic blood pressure; body mass index (BMI); diabetes mellitus	Age (<45, 45-54, 55-59, $\geq$ 55); Sex (male, female); BMI (<25, $\geq$ 25); Diabetes (yes, no); Systolic blood pressure (<120, 120-129, 130-139, 140-149, 150-159, $\geq$ 160)	Age (health survey), sex (health survey). Hypertension was defined as systolic blood pressure $\geq$ 140 mmHg or diastolic blood pressure $\geq$ 90 mmHg or use of oral	NI

	Clinical only)						antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of $\geq 126$ mg/dl or a positive history of diabetes. Body mass index was defined as weight in kilograms divided by the square of height in meters	
6	Saranburu t, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	16	16	NI	Age; sex; systolic blood pressure; diabetes mellitus; glomerular filtration rate at baseline	Age (<45, 45-54, 55-59, $\geq 55$ ); Sex (male, female); Diabetes (yes, no); Systolic blood pressure (<120, 120-129, 130-139, 140-149, 150-159, $\geq 160$ ); eGFR ( $\geq 90$ , 75-89, 60-74)	Age (health survey), sex (health survey). Hypertension was defined as systolic blood pressure $\geq 140$ mmHg or diastolic blood pressure $\geq 90$ mmHg or use of oral antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of $\geq 126$ mg/dl or a positive history of diabetes. Serum creatinine (sCr) was measured by the enzymatic assay on the Vitros 350 analyzer (Ortho-Clinical Diagnostics, USA) using IDMS-Standard Reference Material (SRM) 967 as the standard. Estimate glomerular filtration rate (eGFR) was calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation	NI
6	Saranburu t, 2017 - Model 3 (derivation Clinical + Full)	22	20	NI	Age; sex; systolic blood pressure; diabetes mellitus; glomerular filtration rate at baseline; uric acid; hemoglobin	Age (<45, 45-54, 55-59, $\geq 55$ ); Sex (male, female); Diabetes (yes, no); Systolic blood pressure (<120, 120-129, 130-139, 140-149, 150-159, $\geq 160$ ); eGFR ( $\geq 90$ , 75-89, 60-74); Uric acid (>6 for female or >7 for male, $\leq 6$	Age (health survey), sex (health survey). Hypertension was defined as systolic blood pressure $\geq 140$ mmHg or diastolic blood pressure $\geq 90$ mmHg or use of oral	NI

	laboratory tests)					for female or $\leq 7$ for male); Hemoglobin ( $<12$ for female or $<13$ for male, $\geq 12$ for female or $\geq 13$ for male)	antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of $\geq 126$ mg/dl or a positive history of diabetes. Serum creatinine (sCr) was measured by the enzymatic assay on the Vitros 350 analyzer (Ortho-Clinical Diagnostics, USA) using IDMS-Standard Reference Material (SRM) 967 as the standard. Estimate glomerular filtration rate (eGFR) was calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation. There is no information about uric acid and hemoglobin	
6	Saranburu t, 2017 - Model 1 (validation Clinical only)	n/a	15	NI	Age; sex; systolic blood pressure; waist circumference; diabetes mellitus	Age ( $<45$ , 45-54, 55-59, $\geq 55$ ); Sex (male, female); Waist circumference ( $\leq 80$ for male or $\leq 90$ for female or $>80$ for male); Diabetes (yes, no); Systolic blood pressure ( $<120$ , 120-129, 130-139, 140-149, 150-159, $\geq 160$ )	Age (health survey), sex (health survey). Hypertension was defined as systolic blood pressure $\geq 140$ mmHg or diastolic blood pressure $\geq 90$ mmHg or use of oral antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of $\geq 126$ mg/dl or a positive history of diabetes. Waist circumference was measured midway between the lowest ribs and the iliac crest.	n/a
6	Saranburu t, 2017 - Model 2 (validation	n/a	16	NI	Age; sex; systolic blood pressure; diabetes mellitus; glomerular filtration rate at baseline	Age ( $<45$ , 45-54, 55-59, $\geq 55$ ); Sex (male, female); Diabetes (yes, no); Systolic blood pressure ( $<120$ , 120-129, 130-139, 140-149, 150-159,	Age (health survey), sex (health survey). Hypertension was defined as systolic blood pressure $\geq 140$ mmHg or	n/a

	Clinical + Limited laboratory tests)					≥160); eGFR (≥90, 75-89, 60-74)	diastolic blood pressure ≥ 90 mmHg or use of oral antihypertensive medication. Diabetes mellitus was defined as a fasting glucose of ≥126 mg/dl or a positive history of diabetes. Serum creatinine (sCr) was measured by the enzymatic assay on the Vitros 350 analyzer (Ortho-Clinical Diagnostics, USA) using IDMS-Standard Reference Material (SRM) 967 as the standard. Estimate glomerular filtration rate (eGFR) was calculated according to two-level race variable Chronic Kidney Disease–Epidemiology Collaboration (CKDEPI) equation	
7	Thakkinstian, 2011 (derivation)	37	10	NI	Age; history of kidney stones; diabetes mellitus; hypertension	Age (<40, 40-59, 60-69, ≥70); Hypertension (taking antihypertensive drug(s) or had systolic blood pressure ≥140 mmHg or diastolic blood pressure ≥90 mmHg); Diabetes (taking oral hypoglycaemic agents or fasting plasma glucose levels ≥126 mg/dL); History of kidney stone was measured by self-reporting kidney stone	Age (survey), diabetes (history of illness, relevant medicines used or laboratory tests/physical examinations), hypertension (history of illness, relevant medicines used or laboratory tests/physical examinations), and history of kidney stones (self-reported in survey).	NI
8	Wen, 2020 - Simple Risk Score (derivation)	NI	15	Time-varying	Waist circumference; systolic blood pressure; sex; education; diabetes	Waist circumference [<80/<75, 80-84.9/75-79.9, 85-89.9/80-84.9, 90-94.9/85-89.9, ≥95/≥90 (for male/female)]; systolic blood pressure (<120, 120-139, 140-159, >160); sex (male, female); education (illiterate, primary school and above); diabetes (no or yes)	During medical examinations, participants took two blood pressure measurements using a non-invasive automatic HEM-907 blood pressure monitor after 5 minutes of rest. Systolic blood pressure was identified as the	NI

							<p>average values of two independent measurements; Diabetes was defined as: (1) FPG <math>\geq 7.0</math> mmol/L, or (2) self-reported diagnosis of diabetes, or (3) the use of antidiabetic medications; According to the number of years of education, they were divided into four groups (illiterate for 0 years, primary school for 1–6 years, junior high school for 7–9 years, and senior high school for <math>\geq 10</math> years); Sex was self-reported; Information about waist circumference was no available</p>	
8	Wen, 2020 - Best-fit Risk Score (derivation )	NI	19	Time-varying	Urinary Albumin-to-creatinine ratio; systolic blood pressure; C-reactive protein; triglycerides; sex; education; diabetes	Urinary Albumin-to-creatinine ratio (<5.0, 5.0-10.0, >10.0); systolic blood pressure (<120, 120-139, 140-159, >160); C-reactive protein (<1.0, 1-3, >3.0); triglycerides (<1.0, 1.0-1.7, >1.7); sex (male, female); education (illiterate, primary school and above); diabetes (no or yes)	<p>Urinary albumin and creatinine were measured from fresh morning spot urine samples; During medical examinations, participants took two blood pressure measurements using a non-invasive automatic HEM-907 blood pressure monitor after 5 minutes of rest. Systolic blood pressure was identified as the average values of two independent measurements; Diabetes was defined as: (1) FPG <math>\geq 7.0</math> mmol/L, or (2) self-reported diagnosis of diabetes, or (3) the use of antidiabetic medications; According to the number of years of education, they were divided into four groups</p>	NI

							(illiterate for 0 years, primary school for 1–6 years, junior high school for 7–9 years, and senior high school for ≥10 years); Sex was self-reported; Information about waist circumference, C-reactive protein and triglycerides were no available	
9	Wu, 2016 (derivation)	NI	10	Baseline	Age, gender and body mass index (BMI) status.	Age (≤ 40, 41–50, 51–60, 61–70, ≥71), gender (male, female) and body mass index (BMI) status (normal, overweight: 23–24.9 kg/m <sup>2</sup> , obesity: ≥25 kg/m <sup>2</sup> ).	Age (self-reported), gender (self-reported) and body mass index (BMI) status (calculated from participant's measured body weight and height).	NI
9	Wu, 2016 (validation)	n/a	10	Baseline	Age, gender and body mass index (BMI) status.	Age (≤ 40, 41 – 50, 51 – 60, 61 – 70, 71+), gender (male, female) and body mass index (BMI) status (normal, overweight: 23–24.9 kg/m <sup>2</sup> , obesity: ≥25 kg/m <sup>2</sup> ).	Age (self-reported), gender (self-reported) and body mass index (BMI) status (calculated from participant's measured body weight and height).	n/a



**S3.4 Table: Sample size and missing data**

N°	Study	Sample Size			Missing Data		
		Baseline sample size	Number of outcome events	Total outcome events per candidate predictors	Missing data	Number of participants with missing data	Missing data per candidate predictors
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	3270	722	n/a	Complete-case	2817	n/a
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	3240	1359	n/a	Complete-case	2847	n/a
2	Bradshaw, 2019 - Model 1 (derivation)	8698	947	31,57	Complete-case	896	29,87
2	Bradshaw, 2019 - Model 2 (derivation)	8698	947	41,17	Complete-case	896	38,96
2	Bradshaw, 2019 - Model 3a (derivation)	8698	947	NI	Complete-case	896	NI
2	Bradshaw, 2019 - Model 3b (derivation)	8698	947	118,38	Complete-case	896	112,00
2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	4065	NI	n/a	Complete-case	1300	n/a
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	4940	NI	n/a	Complete-case	1233	n/a
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	2368	81	2,25	Complete-case	235	6,53
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	2368	81	3,12	Complete-case	235	9,04
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	1459	79	n/a	Complete-case	79	n/a
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	1459	79	n/a	Complete-case	79	n/a
4	Mogueo, 2015 - Korean model (eGFR validation)	902	259	n/a	Complete-case	383	n/a
4	Mogueo, 2015 - Thai model (eGFR validation)	902	259	n/a	Complete-case	383	n/a
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	902	268	n/a	Complete-case	383	n/a
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	902	268	n/a	Complete-case	383	n/a
5	Saranburut, 2017 - Framingham Heart Study (MDRD validation)	2141	222	n/a	Complete-case	NI	n/a
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	2328	233	n/a	Complete-case	NI	n/a
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	3186	271	18,07	Complete-case	NI	NI
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	3186	271	18,07	Complete-case	NI	NI
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	3186	271	16,94	Complete-case	NI	NI
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	3186	271	12,32	Complete-case	NI	NI
6	Saranburut, 2017 - Model 1 (validation Clinical only)	1395	27	n/a	Complete-case	NI	NI

6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	1395	27	n/a	Complete-case	NI	NI
7	Thakkinstian, 2011 (derivation)	3459	626	16,92	NI	NI	NI
8	Wen, 2020 - Simple Risk Score (derivation)	3266	590	NI	Complete-case	992	NI
8	Wen, 2020 - Best-fit Risk Score (derivation)	3266	590	NI	Complete-case	992	NI
9	Wu, 2016 (derivation)	14374	294	NI	Complete-case	3135	NI
9	Wu, 2016 (validation)	4371	48	n/a	Complete-case	911	n/a

**S3.5 Table: Model development**

		Model Development					
N°	Study	Regression method	Were the model assumptions verified?	Predictors selection	If the prediction model was a replication, which was the original model?	If there were pre-selection, describe the method	Was a shrinkage method used?
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	n/a	n/a	n/a	n/a	n/a	n/a
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	n/a	n/a	n/a	n/a	n/a	n/a
2	Bradshaw, 2019 - Model 1 (derivation)	Logistic	NI	Pre-selection	n/a	Step-down selection procedure based on the Akaike information criterion to select the final predictors	No
2	Bradshaw, 2019 - Model 2 (derivation)	Logistic	NI	Pre-selection	n/a	Step-down selection procedure based on the Akaike information criterion to select the final predictors	No
2	Bradshaw, 2019 - Model 3a (derivation)	Logistic	NI	Pre-selection	n/a	Step-down selection procedure based on the Akaike information criterion to select the final predictors	No
2	Bradshaw, 2019 - Model 3b (derivation)	Logistic	NI	Pre-selection	n/a	Step-down selection procedure based on the Akaike information criterion to select the final predictors	No
2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	n/a	n/a	n/a	n/a	n/a	n/a
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	n/a	n/a	n/a	n/a	n/a	n/a
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Logistic	NI	Pre-selection	n/a	Stepwise backward elimination method	No

3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	Logistic	NI	Pre-selection	n/a	Stepwise backward elimination method	No
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	n/a	n/a	n/a	n/a	n/a	n/a
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	n/a	n/a	n/a	n/a	n/a	n/a
4	Mogueo, 2015 - Korean model (eGFR validation)	n/a	n/a	n/a	n/a	n/a	n/a
4	Mogueo, 2015 - Thai model (eGFR validation)	n/a	n/a	n/a	n/a	n/a	n/a
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	n/a	n/a	n/a	n/a	n/a	n/a
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	n/a	n/a	n/a	n/a	n/a	n/a
5	Saranburut, 2017 - Framingham Heart Study (MDRD validation)	n/a	n/a	n/a	n/a	n/a	n/a
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	n/a	n/a	n/a	n/a	n/a	n/a
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	Logistic	NI	Pre-selection	n/a	Variables were sequentially added in a pre-specified order and incorporated using a p< 0.05 threshold for entry and retention in the final model	No
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Logistic	NI	Pre-selection	n/a	Variables were sequentially added in a pre-specified order and incorporated using a p< 0.05 threshold for entry and retention in the final model	No
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Logistic	NI	Pre-selection	n/a	Variables were sequentially added in a pre-specified order and incorporated using a p< 0.05 threshold for entry and retention in the final model	No
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Logistic	NI	Pre-selection	n/a	Variables were sequentially added in a pre-specified order and incorporated using a p< 0.05 threshold for entry and retention in the final model	No

6	Saranburut, 2017 - Model 1 (validation Clinical only)	n/a	n/a	n/a	n/a	n/a	n/a
6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	n/a	n/a	n/a	n/a	n/a	n/a
7	Thakkinstian, 2011 (derivation)	Logistic	NI	Pre-selection	n/a	Factors with p values < 0.15 in a univariate analysis were considered to be simultaneously included in the multivariate logistic equation. Model selection was performed using F-tests, and thus only significant variables were kept in the final model. C statistic of models with and without a particular variable were then compared; if dropping that variable did not significantly reduce the explanation of the CKD, that variable was omitted in the final parsimonious model.	No
8	Wen, 2020 - Simple Risk Score (derivation)	Logistic	NI	Pre-selection	n/a	Risk factors were investigated by forward stepwise logistic regression and only statistically significant (a two-sided P value <0.05) risk factors were retained.	No
8	Wen, 2020 - Best-fit Risk Score (derivation)	Logistic	NI	Pre-selection	n/a	Risk factors were investigated by forward stepwise logistic regression and only statistically significant (a two-sided P value <0.05) risk factors were retained.	No
9	Wu, 2016 (derivation)	Logistic	NI	Pre-selection	n/a	Stepwise logistic regression model. Variables with a p value less than 0.1 were kept in the final model.	No
9	Wu, 2016 (validation)	n/a	n/a	n/a	n/a	n/a	n/a

n/a: not applicable; NI: no information

**S3.6 Table: Model performance**

		Model Performance				
N°	Study	Calibration	Discrimination (%)	Classification measures	Cut-off point	For replication studies, was the cut-off the same?
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	Hosmer-Lemeshow X2 test (for intercept adjusted model): 13.53 with a p-value 0.09 (for male) and 10.1 with a p-value 0.26 (for women)	AUC (95% CI) for final intercept adjusted model = Male: 0.76 (0.72-0.79) and Female: 0.71 (0.69-0.73)	Men: Sensitivity = 72.7%, Specificity = 67.6%. Women: Sensitivity = 66.8%, Specificity = 65.6%.	Men: 25. Women: 19	No
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	Hosmer-Lemeshow X2 test (for intercept adjusted model): 12.54 with a p-value 0.13 (for male) and 8.19 with a p-value 0.41 (for women)	AUC (95% CI) for final intercept adjusted model = Male: 0.71 (0.67-0.74) and Female: 0.70 (0.68-0.73)	Men: Sensitivity = 64.5%, Specificity = 69.5%. Women: Sensitivity = 56.9%, Specificity = 76.6%	Men: 25. Women: 23	No
2	Bradshaw, 2019 - Model 1 (derivation)	Calibration slope: 0.96	C-statistic (95% CI) = 0.79 (0.78-0.81)	Sensitivity = 72%, Specificity = 72%, PPV = 24%, NPV = 96%	0.09	n/a
2	Bradshaw, 2019 - Model 2 (derivation)	Calibration slope: 0.98	C-statistic (95% CI) = 0.73 (0.72-0.75)	Sensitivity = 68%, Specificity = 67%, PPV = 20%, NPV = 95%	0.09	n/a
2	Bradshaw, 2019 - Model 3a (derivation)	Calibration slope: 0.98	C-statistic (95% CI) = 0.77 (0.75-0.79)	Sensitivity = 71%, Specificity = 70%, PPV = 22%, NPV = 95%	0.09	n/a
2	Bradshaw, 2019 - Model 3b (derivation)	Calibration slope: 0.99	C-statistic (95% CI) = 0.77 (0.76-0.79)	Sensitivity = 71%, Specificity = 70%, PPV = 22%, NPV = 95%	0.09	n/a

2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	NI	C-statistic (95% CI) = 0.74 (0.73-0.74)	NI	0.09	Yes
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	NI	C-statistic (95% CI) = 0.70 (0.69-0.71)	NI	0.09	Yes
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Hosmer-Lemeshow X2 test: 4.13 with a p-value of 0.53 (for final multivariable model).	AUC = 76.2%	Sensibility = 82.5%, Specificity = 70.0%, PPV = 8.8%, NPV = 99.1%, LHR+ = 2.8, LHR- = 0.3	2	n/a
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	Hosmer-Lemeshow X2 test: 4.13 with a p-value of 0.53 (for final multivariable model).	AUC = 76%	Sensibility = 80.0%, Specificity = 72.0%, PPV = 9.1%, NPV = 99.0%, LHR+ = 2.9, LHR- = 0.3	2	n/a
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	NI	AUC = 70.0%.	Sensitivity = 70.5%, Specificity = 69.1%, PPV = 11.4%, NPV = 97.6%, LHR+ = 2.3, LHR- = 0.4	2	Yes
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	NI	AUC = 70.0%.	Sensitivity = 70.5%, Specificity = 69.7%, PPV = 11.6%, NPV = 97.7%, LHR+ = 2.3, LHR- = 0.4	2	Yes
4	Mogueo, 2015 - Korean model (eGFR validation)	Expected/Observed rate (95%) = 0.76 (0.67-0.86); Brier score = 0.164; Yates slope = 0.208	C-statistic (95% CI) = 0.797 (0.765-0.829)	Sensitivity = 82%, Specificity = 67%	0.30	NI
4	Mogueo, 2015 - Thai model (eGFR validation)	Expected/Observed rate (95%) = 0.98 (0.87-1.10); Brier score = 0.165; Yates slope = 0.200	C-statistic (95% CI) = 0.760 (0.726-0.793)	Sensitivity = 73%, Specificity = 72%	0.31	NI
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Expected/Observed rate (95%) = 0.76 (0.67-0.85); Brier score = 0.161; Yates slope = 0.225	C-statistic (95% CI) = 0.811 (0.780-0.842)	Sensitivity = 84%, Specificity = 68%	0.31	NI

4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Expected/Observed rate (95%) = 0.97 (0.86-1.09); Brier score = 0.164; Yates slope = 0.211	C-statistic (95% CI) = 0.772 (0.739-0.805)	Sensitivity = 74%, Specificity = 73%	0.32	NI
5	Saranburut, 2017 - Framingham Heart Study (MDRD validation)	Hosmer-Lemeshow X2 test: 30.2 (p<0.001)	AUC (95% CI) = 0.69 (0.66-0.73)	NI	NI	NI
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Hosmer-Lemeshow X2 test: 256.5 (p<0.001)	AUC (95% CI) = 0.63 (0.57-0.65)	NI	NI	NI
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	Hosmer-Lemeshow X2 test: 9.02 (p=0.34)	AUC (95% CI) = 0.72 (0.69-0.75)	NI	NI	n/a
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Hosmer-Lemeshow X2 test: 8.87 (p=0.35)	AUC (95% CI) = 0.72 (0.69-0.75)	NI	NI	n/a
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Hosmer-Lemeshow X2 test: 10.87 (p=0.21)	AUC (95% CI) = 0.79 (0.76-0.82)	NI	NI	n/a
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Hosmer-Lemeshow X2 test: 8.28 (p=0.41)	AUC (95% CI) = 0.80 (0.77-0.82)	NI	NI	n/a
6	Saranburut, 2017 - Model 1 (validation Clinical only)	Hosmer-Lemeshow X2 test: 4.31 (p=0.229)	AUC (95% CI) = 0.66 (0.55-0.78)	NI	NI	NI
6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	Hosmer-Lemeshow X2 test: 2.29 (p=0.514)	AUC (95% CI) = 0.88 (0.80-0.95)	NI	NI	NI
7	Thakkinstian, 2011 (derivation)	Calibration was assessed by subtracting the two Somer's D correlation coefficients: 0.045 (95% CI: 0.034-0.057)	C-statistic of internal validation = 0.741	Sensitivity = 76%, Specificity = 69%	5	n/a



8	Wen, 2020 - Simple Risk Score (derivation)	Hosmer-Lemeshow X2 test: 4.89 (p=0.769)	AUC (95% CI) = 0.717 (0.689-0.744)	Sensitivity = 70.49%, Specificity = 65.14%, PPV = 29.8%, NPV = 91.3%, LHR+ = 2.02, LHR- = 0.45	14	n/a
8	Wen, 2020 - Best-fit Risk Score (derivation)	Hosmer-Lemeshow X2 test: 2.52 (p=0.961)	AUC (95% CI) = 0.721 (0.693-0.748)	Sensitivity = 56.83%, Specificity = 76.61%, PPV = 33.8%, NPV = 89.4%, LHR+ = 2.43, LHR- = 0.56	24	n/a
9	Wu, 2016 (derivation)	Internal validation dataset: Hosmer-Lemeshow X2 test P=0.798	AUC (95% CI) of internal validation = 0.894 (0.861-0.926)	Sensitivity = 0.820, Specificity = 0.863	36	n/a
9	Wu, 2016 (validation)	Hosmer-Lemeshow X2 test P=397	AUC = 0.880 (95%CI: 0.829-0.931)	NI	NI	NI

AUC, area under the curve; CI, confident interval; NI, no information.

**S3.7 Table: Results**

		Results			
N°	Study	Was a simplified model presented?	Were the coefficients of the regression model presented?	Was the baseline risk presented?	Were there alternative results presentation?
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	No	No	Yes	No
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	No	No	Yes	No
2	Bradshaw, 2019 - Model 1 (derivation)	Yes	No	No	No
2	Bradshaw, 2019 - Model 2 (derivation)	Yes	No	No	No
2	Bradshaw, 2019 - Model 3a (derivation)	No	No	No	No
2	Bradshaw, 2019 - Model 3b (derivation)	Yes	No	No	No
2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	No	No	No	No
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	No	No	No	No
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Yes	Yes	No	No
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	No	Yes	No	No
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	Yes	No	No	No
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	No	No	No	No
4	Mogueo, 2015 - Korean model (eGFR validation)	No	No	No	No
4	Mogueo, 2015 - Thai model (eGFR validation)	No	No	No	No
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	No	No	No	No
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	No	No	No	No
5	Saranburut, 2017 - Framingham Heart	No	Yes	No	No

	Study (MDRD validation)				
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	No	Yes	No	No
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	No	Yes	No	Yes
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	No	No	No	Yes
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Yes	Yes	No	Yes
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Yes	Yes	No	No
6	Saranburut, 2017 - Model 1 (validation Clinical only)	No	No	No	Yes
6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	Yes	No	No	Yes
7	Thakkinstian, 2011 (derivation)	No	Yes	No	Yes
8	Wen, 2020 - Simple Risk Score (derivation)	No	Yes	Yes	Yes
8	Wen, 2020 - Best-fit Risk Score (derivation)	No	Yes	Yes	Yes
9	Wu, 2016 (derivation)	No	Yes	No	Yes
9	Wu, 2016 (validation)	No	Yes	No	Yes

**S3.8 Table: Discussion**

		Discussion		
N°	Study	Interpretation of the results	Comparison with other studies in LAC	Generalizability
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	Exploratory	No	Non-generalizability
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	Exploratory	No	Non-generalizability
2	Bradshaw, 2019 - Model 1 (derivation)	NI	No	NI
2	Bradshaw, 2019 - Model 2 (derivation)	NI	No	NI
2	Bradshaw, 2019 - Model 3a (derivation)	Confirmatory	Yes	Non-generalizability
2	Bradshaw, 2019 - Model 3b (derivation)	NI	No	NI
2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	Confirmatory	Yes	Non-generalizability
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	Confirmatory	Yes	Non-generalizability
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Exploratory	Yes	Generalizable
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	Exploratory	Yes	Generalizable
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	Exploratory	Yes	Generalizable
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	Exploratory	Yes	Generalizable
4	Mogueo, 2015 - Korean model (eGFR validation)	Exploratory	Yes	Non-generalizability
4	Mogueo, 2015 - Thai model (eGFR validation)	Exploratory	Yes	Non-generalizability
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Exploratory	Yes	Non-generalizability
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Exploratory	Yes	Non-generalizability
5	Saranburut, 2017 - Framingham Heart Study (MDRD validation)	Exploratory	No	Non-generalizability
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Exploratory	No	Non-generalizability
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	Exploratory	No	Non-generalizability
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Exploratory	No	Non-generalizability
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Exploratory	No	Non-generalizability
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Exploratory	No	Non-generalizability
6	Saranburut, 2017 - Model 1 (validation Clinical only)	Exploratory	No	Non-generalizability
6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	Exploratory	No	Non-generalizability
7	Thakkestian, 2011 (derivation)	Confirmatory	No	Non-generalizability

8	Wen, 2020 - Simple Risk Score (derivation)	Confirmatory	Yes	Non-generalizability
8	Wen, 2020 - Best-fit Risk Score (derivation)	Exploratory	Yes	Non-generalizability
9	Wu, 2016 (derivation)	Exploratory	No	Non-generalizability
9	Wu, 2016 (validation)	Exploratory	No	Non-generalizability

**S4 Table: PROBAST**

**S4.1 Table: Risk of Bias (RoB)**

Study	Participants		Predictors		
	Were appropriate data sources used, e.g., cohort, RCT, or nested case-control study data?	Were all inclusions and exclusions of participants appropriate?	Were predictors defined and assessed in a similar way for all participants?	Were predictor assessments made without knowledge of outcome data?	Are all predictors available at the time the model is intended to be used?
Asgari, 2020 European Risk Assessment tool (6-years validation)	Y	Y	Y	Y	Y
Asgari, 2020 European Risk Assessment tool (9-years validation)	Y	Y	Y	Y	Y
Bradshaw, 2019 - Model 1 (derivation)	Y	Y	Y	Y	PY
Bradshaw, 2019 - Model 2 (derivation)	Y	Y	Y	Y	Y
Bradshaw, 2019 - Model 3a (derivation)	Y	Y	Y	Y	PY
Bradshaw, 2019 - Model 3b (derivation)	Y	Y	Y	Y	PY
Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	Y	Y	Y	Y	PY
Bradshaw, 2019 - Model 3a (UDAY rural validation)	Y	Y	Y	Y	PY
Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Y	Y	Y	Y	PY
Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	Y	Y	Y	Y	Y
Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	Y	Y	Y	Y	PY
Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	Y	Y	Y	Y	Y
Mogueo, 2015 - Korean model (eGFR validation)	Y	Y	Y	Y	PY
Mogueo, 2015 - Thai model (eGFR validation)	Y	Y	Y	Y	Y
Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Y	Y	Y	Y	PY
Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Y	Y	Y	Y	Y

Saranburut, 2017 - Framingham Heart Study (MDRD validation)	Y	Y	Y	Y	PY
Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Y	Y	Y	Y	PY
Saranburut, 2017 - Model 1 (derivation Clinical only)	Y	Y	Y	Y	Y
Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Y	Y	Y	Y	Y
Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Y	Y	Y	Y	PY
Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Y	Y	Y	Y	PY
Saranburut, 2017 - Model 1 (validation Clinical only)	Y	Y	Y	Y	Y
Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	Y	Y	Y	Y	PY
Thakkinstian, 2011 (derivation)	Y	Y	Y	Y	Y
Wen, 2020 - Simple Risk Score (derivation)	Y	Y	Y	Y	Y
Wen, 2020 - Best-fit Risk Score (derivation)	Y	Y	Y	Y	Y
Wu, 2016 (derivation)	Y	Y	Y	Y	Y
Wu, 2016 (validation)	Y	Y	Y	Y	Y

Answer options: Y (yes), PY (probably yes), N (no), PN (probably no), NI (no information), n/a (not applicable).

Study	Outcome					
	Was the outcome determined appropriately?	Was a prespecified or standard outcome definition used?	Were predictors excluded from the outcome definition?	Was the outcome defined and determined in a similar way for all participants?	Was the outcome determined without knowledge of predictor information?	Was the time interval between predictor assessment and outcome determination appropriate?
Asgari, 2020 European Risk Assessment tool (6-years validation)	Y	Y	Y	Y	NI	Y
Asgari, 2020 European Risk Assessment tool (9-years validation)	Y	Y	Y	Y	NI	Y
Bradshaw, 2019 - Model 1 (derivation)	Y	Y	Y	Y	NI	PY
Bradshaw, 2019 - Model 2 (derivation)	Y	Y	Y	Y	NI	Y
Bradshaw, 2019 - Model 3a (derivation)	NI	Y	Y	Y	NI	PY
Bradshaw, 2019 - Model 3b (derivation)	Y	Y	Y	Y	NI	PY
Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	Y	Y	Y	Y	NI	PY
Bradshaw, 2019 - Model 3a (UDAY rural validation)	Y	Y	Y	Y	NI	PY
Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Y	Y	Y	Y	PY	PY
Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	Y	Y	Y	Y	PY	Y
Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	Y	Y	Y	Y	PY	PY



Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	Y	Y	Y	Y	PY	Y
Mogueo, 2015 - Korean model (eGFR validation)	Y	Y	Y	Y	NI	PY
Mogueo, 2015 - Thai model (eGFR validation)	Y	Y	Y	Y	NI	Y
Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Y	Y	Y	Y	NI	PY
Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Y	Y	Y	Y	NI	Y
Saranburut, 2017 - Framingham Heart Study (MDRD validation)	Y	Y	Y	Y	NI	PY
Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Y	Y	Y	Y	NI	PY
Saranburut, 2017 - Model 1 (derivation Clinical only)	Y	Y	Y	Y	NI	Y
Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Y	Y	Y	Y	NI	Y
Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Y	Y	Y	Y	NI	PY
Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Y	Y	Y	Y	NI	PY
Saranburut, 2017 - Model 1 (validation Clinical only)	Y	Y	Y	Y	NI	Y
Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	Y	Y	Y	Y	NI	PY
Thakkinstian, 2011 (derivation)	Y	Y	Y	Y	NI	Y
Wen, 2020 - Simple Risk Score (derivation)	Y	Y	Y	Y	NI	Y

Wen, 2020 - Best-fit Risk Score (derivation)	Y	Y	Y	Y	NI	Y
Wu, 2016 (derivation)	Y	Y	Y	Y	NI	Y
Wu, 2016 (validation)	Y	Y	Y	Y	NI	Y

Answer options: Y (yes), PY (probably yes), N (no), PN (probably no), NI (no information), n/a (not applicable).

Study	Analysis								
	Were there a reasonable number of participants with the outcome?	Were continuous and categorical predictors handled appropriately?	Were all enrolled participants included in the analysis?	Were participants with missing data handled appropriately?	Was selection of predictors based on univariable analysis avoided? [development studies only]	Were complexities in the data (e.g., censoring, competing risks, sampling of control participants) accounted for appropriately?	Were relevant model performance measures evaluated appropriately?	Were model overfitting and optimism in model performance accounted for? [development studies only]	Do predictors and their assigned weights in the final model correspond to the results from the reported multivariable analysis? [development studies only]
Asgari, 2020 European Risk Assessment tool (6-years validation)	Y	Y	N	N	n/a	NI	N	n/a	n/a
Asgari, 2020 European Risk Assessment tool (9-years validation)	Y	Y	N	N	n/a	NI	N	n/a	n/a
Bradshaw, 2019 - Model 1 (derivation)	Y	N	N	N	N	NI	Y	Y	NI
Bradshaw, 2019 - Model 2 (derivation)	Y	N	N	N	N	NI	Y	Y	NI
Bradshaw, 2019 - Model 3a (derivation)	NI	NI	N	N	N	NI	Y	Y	NI
Bradshaw, 2019 - Model 3b (derivation)	Y	N	N	N	N	NI	Y	Y	NI
Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	NI	Y	N	N	n/a	NI	NI	n/a	n/a

Bradshaw, 2019 - Model 3a (UDAY rural validation)	NI	Y	N	N	n/a	NI	NI	n/a	n/a
Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	N	N	N	N	N	NI	N	Y	Y
Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	N	N	N	N	N	NI	N	Y	Y
Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	N	Y	N	N	n/a	NI	N	n/a	n/a
Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	N	Y	N	N	n/a	NI	N	n/a	n/a
Mogueo, 2015 - Korean model (eGFR validation)	Y	Y	N	N	n/a	NI	PY	n/a	n/a
Mogueo, 2015 - Thai model (eGFR validation)	Y	Y	N	N	n/a	NI	PY	n/a	n/a
Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Y	Y	N	N	n/a	NI	PY	n/a	n/a
Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Y	Y	N	N	n/a	NI	PY	n/a	n/a
Saranburut, 2017 - Framingham Heart Study (MDRD validation)	Y	Y	N	N	n/a	NI	N	n/a	n/a
Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Y	Y	N	N	n/a	NI	N	n/a	n/a
Saranburut, 2017 - Model 1 (derivation Clinical only)	PY	N	N	N	N	NI	N	Y	Y
Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	PY	N	N	N	N	NI	N	Y	NI

Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	PY	N	N	N	N	NI	N	Y	Y
Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	PN	N	N	N	N	NI	N	Y	Y
Saranburut, 2017 - Model 1 (validation Clinical only)	N	Y	N	N	n/a	NI	N	n/a	n/a
Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	N	Y	N	N	n/a	NI	N	n/a	n/a
Thakkestian, 2011 (derivation)	PY	N	NI	NI	N	NI	N	Y	Y
Wen, 2020 - Simple Risk Score (derivation)	NI	N	N	N	N	NI	N	N	Y
Wen, 2020 - Best-fit Risk Score (derivation)	NI	N	N	N	N	NI	N	N	Y
Wu, 2016 (derivation)	NI	N	N	N	N	NI	N	N	Y
Wu, 2016 (validation)	N	Y	N	N	n/a	NI	N	n/a	Y

Answer options: Y (yes), PY (probably yes), N (no), PN (probably no), NI (no information), n/a (not applicable).

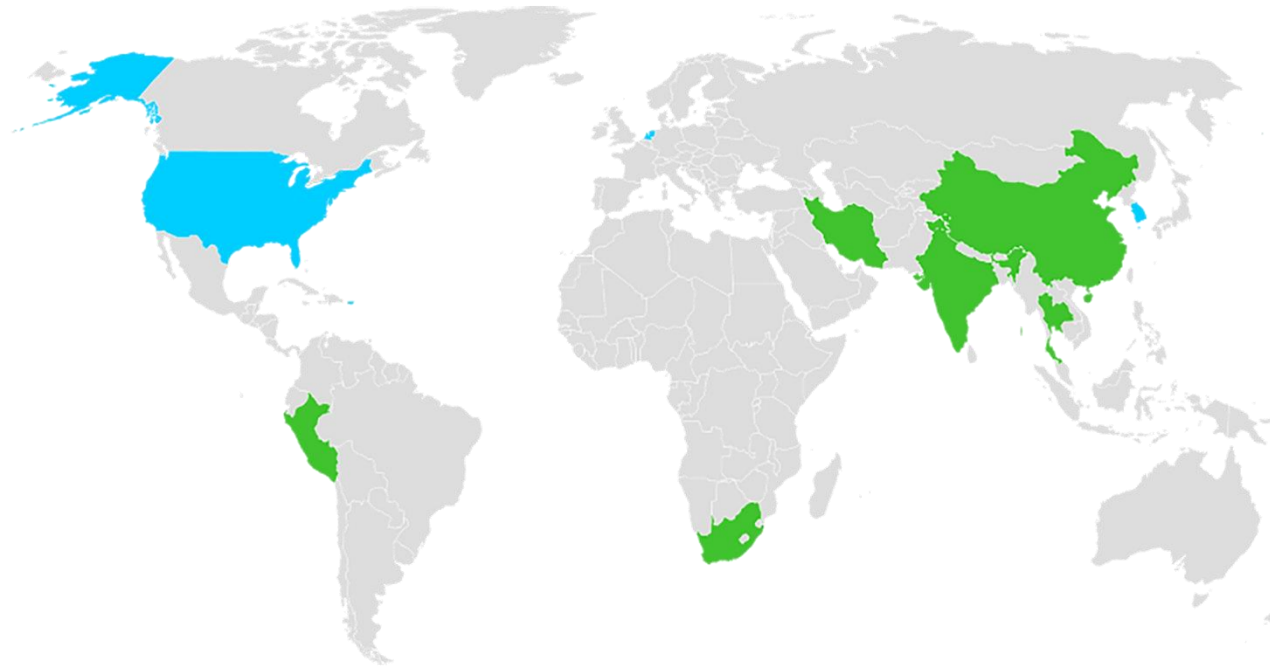
**S4.2 Table: Applicability**

N°	Study	Participants	Predictors	Outcome
1	Asgari, 2020 European Risk Assessment tool (6-years validation)	Low	Low	Low
1	Asgari, 2020 European Risk Assessment tool (9-years validation)	Low	Low	Low
2	Bradshaw, 2019 - Model 1 (derivation)	Low	Low	Low
2	Bradshaw, 2019 - Model 2 (derivation)	Low	Low	Low
2	Bradshaw, 2019 - Model 3a (derivation)	Low	Low	Low
2	Bradshaw, 2019 - Model 3b (derivation)	Low	Low	Low
2	Bradshaw, 2019 - Model 3a (CARRS-I urban validation)	Low	Low	Low
2	Bradshaw, 2019 - Model 3a (UDAY rural validation)	Low	Low	Low
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation complete)	Low	Low	Low
3	Carrillo-Larco, 2017 - CRONICAS-CKD (derivation lab-free)	Low	Low	Low
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation complete)	Low	Low	Low
3	Carrillo-Larco, 2017 - CRONICAS-CKD (validation lab-free)	Low	Low	Low
4	Mogueo, 2015 - Korean model (eGFR validation)	Low	Low	Low
4	Mogueo, 2015 - Thai model (eGFR validation)	Low	Low	Low
4	Mogueo, 2015 - Korean model (eGFR or proteinuria validation)	Low	Low	Low
4	Mogueo, 2015 - Thai model (eGFR or proteinuria validation)	Low	Low	Low
5	Saranburut, 2017 - Framingham Heart Study (MDRD validation)	Low	Low	Low
5	Saranburut, 2017 - Framingham Heart Study (CKD-EPI validation)	Low	Low	Low
6	Saranburut, 2017 - Model 1 (derivation Clinical only)	Low	Low	Low
6	Saranburut, 2017 - Model 1 BMI (derivation Clinical only)	Low	Low	Low
6	Saranburut, 2017 - Model 2 (derivation Clinical + Limited laboratory tests)	Low	Low	Low
6	Saranburut, 2017 - Model 3 (derivation Clinical + Full laboratory tests)	Low	Low	Low
6	Saranburut, 2017 - Model 1 (validation Clinical only)	Low	Low	Low
6	Saranburut, 2017 - Model 2 (validation Clinical + Limited laboratory tests)	Low	Low	Low
7	Thakkinstian, 2011 (derivation)	Low	Low	Low
8	Wen, 2020 - Simple Risk Score (derivation)	Low	Low	Low
8	Wen, 2020 - Best-fit Risk Score (derivation)	Low	Low	Low
9	Wu, 2016 (derivation)	Low	Low	Low

9	Wu, 2016 (validation)	Low	Low	Low
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Answer options: Low (low concern for applicability), Hig (High concern for applicability) and Unclear (Unclear concern for applicability)

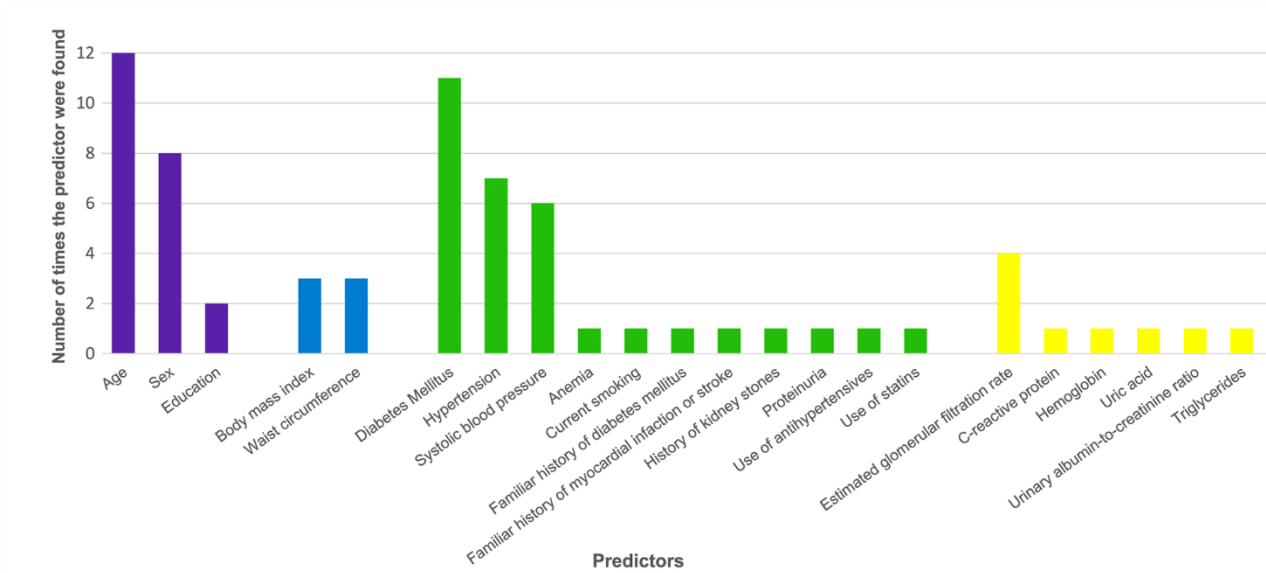
**S1 Figure: Countries where studies were conducted.**



LMIC that developed and/or validated models included in this review (Green). Moreover, Asgari et al, Mogueo et al ] and Saranburut et al validated risk models that were originally derived in the Netherlands, South Korea and the United States, respectively (Blue).



**S2 Figure: Predictors included in the final models.**



The colours of the bars identify the underlying characteristic of predictors inherent to: the subject (purple), anthropometrics (blue), clinical assessment and history (green), and laboratory measures (yellow).