Diagnostic Test Accuracy of Urine C-peptide Creatinine Ratio for the Correct Identification of the Type of Diabetes: A Systematic Review

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Objective: To examine the accuracy of urine c-peptide creatinine ratio (UCPCR) for identifying the type of diabetes in appropriate clinical settings. Design: Systematic review of test accuracy studies on patients with different forms of diabetes. Data sources: Medline, Embase and Cochrane library databases from 1 January 2000 to 15 November 2020. Eligibility criteria: Studies reporting the use of UCPCR for diagnosing patients with type 1 diabetes mellitus (T1DM), type 2 diabetes mellitus (T2DM) and monogenic forms of diabetes (categorized as maturity-onset diabetes of the young [MODY]). Study selection and data synthesis: Two reviewers independently assessed articles for inclusion and assessed the methodological quality of the studies using the Quality Assessment of Diagnostic Accuracy Studies-2 tool, with input from a third reviewer to reach consensus when there was a dispute. Meta-analysis was performed with the studies reporting complete data to derive the pooled sensitivity, specificity and diagnostic odds ratio (DOR), and narrative synthesis only for those with incomplete data. Results: Nine studies with 4,488 patients were included in the qualitative synthesis, while only four of these (915 patients) had complete data and were included in the quantitative synthesis. All the studies had moderate risk of bias and applicability concerns. Meta-analysis of three studies (n=130) revealed sensitivity, specificity and DOR of 84.4% (95% confidence interval [CI] 68.1–93.2%), 91.6% (82.8–96.1%) and 59.9 (32.8–106.0), respectively, for diagnosing T1DM using a UCPCR cut-off of <0.2 nmol/mmol. For participants with T2DM (three studies; n=739), UCPCR >0.2 nmol/mmol was associated with sensitivity, specificity and DOR of 92.8% (84.2–96.9%), 81.6% (61.3–92.5%) and 56.9 (31.3–103.5), respectively. For patients with MODY in the appropriate clinical setting, a UCPCR cut-off of >0.2 nmol/mmol showed sensitivity, specificity and DOR of 85.2% (73.1–92.4%), 98.0% (92.4–99.5%) and 281.8 (57.5–1,379.7), respectively. Conclusions: Based on studies with moderate risk of bias and applicability concerns, UCPCR confers moderate to high sensitivity, specificity, and DOR for correctly identifying T1DM, T2DM and monogenic diabetes in appropriate clinical settings. Large multinational studies with multi-ethnic participation among different age groups are necessary before this test can be routinely used in clinical practice.

Study registration: Protocol was registered as PROSPERO CRD42017060633.

Keywords
- C-peptide, urine C-peptide creatinine ratio (UCPCR), endogenous insulin reserve, type 1 diabetes mellitus (T1DM), type 2 diabetes mellitus (T2DM), monogenic diabetes, maturity-onset diabetes of the young (MODY)

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However, there are practical difficulties in measuring plasma C-peptide levels, mainly the requirement that specimens be kept in ice because of the short biological half-life (30 minutes). As such, collection and transport of samples between outpatient clinics, home settings and testing laboratories is challenging.

C-peptide is largely metabolized by the kidney through glomerular filtration and tubular uptake, with only 5% of the total amount produced excreted in the urine. Urinary C-peptide remains stable at room temperature for up to 72 hours when preserved in boric acid, and can be easily transported to testing laboratories, even from remote settings. Therefore, urine C-peptide becomes an attractive and non-invasive alternative for testing β-cell function and insulin production capacity. To avoid having to collect urine for 24 hours to estimate β-cell reserve, spot urine C-peptide creatinine ratio (UCPCR) is an easy alternative option. By using modern, high-sensitivity immunoassay technologies, an Exeter research group later popularized UCPCR to measure C-peptide down to picomolar concentrations. However, the sensitivity, specificity and predictive value of UCPCR for identifying the type of diabetes is highly variable in these published reports, reducing its clinical application, especially when disease duration is considered. Therefore, the aim of this diagnostic test accuracy (DTA) systematic review was to critically appraise the use of UCPCR in correctly identifying the type of diabetes.

**Methods**

We performed the study as per the guidelines specified in the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA). We searched the Medline, EMBASE and Cochrane Library databases to obtain articles from 1 January 2000 to 15 November 2020 (detailed search strategy is shown in Appendix 1). Search was limited to articles published in English, and the references of relevant studies were also searched for inclusion if relevant. Overlapping or potential duplicate data were carefully excluded using EndNote Version X8 (Clarivate™, London, UK).

**Eligibility criteria**

We included a study if: (1) UCPCR was used as a diagnostic test for subtyping diabetes, and (2) positive antibody testing was used to confirm the diagnosis in patients with T1DM, and candidate genes were genetically confirmed for monogenic diabetes (MODY). A UCPCR value <0.2 nmol/mol was considered evidence of severe insulin deficiency. The gold-standard criteria used for diagnosing T1DM were continuous insulin treatment (exogenous insulin dependency) within 3 years of diagnosis, and absolute insulin deficiency (UCPCR <0.2 nmol/mmol ≥5 years post-diagnosis). Diagnosis of MODY was based on the International Society for Paediatric and Adolescent Diabetes (ISPAD) criteria: a family history of diabetes in one parent and a first-degree relative of that parent; lack of characteristics of T1DM (lack of islet-cell antibodies, low or no insulin requirement 5 years after diagnosis; a serum C-peptide levels >200 pmol/L), and lack of classical features of T2DM (marked obesity or acanthosis nigricans). The remainder were considered to be diagnosed with T2DM.

**Data extraction and quality assessment**

Two independent investigators (JM and BS) independently assessed all the titles and abstracts to identify studies eligible for inclusion in the review. The same authors then reviewed the full texts of these studies for eligibility to be included in the DTA study, and extracted data from the eligible studies using a standardized data-extraction form. The form included the following characteristics of each trial: first author's name; year of publication; study population characteristics, including sample size, geographical location, mean age and sex; and diagnostic criteria, including screening and confirmatory tests for the type of diabetes. Differences between reviewers were resolved by discussion with the third reviewer (APA).

The risk of bias and applicability of the identified studies were assessed by two independent reviewers (JM and CJF) using the modified Quality Assessment of Diagnostic Accuracy Studies-2 (QUADAS-2) criteria, for patient selection, performance of the index test, performance of the reference test, and flow and timing. Conflicts were resolved by consensus between the two reviewers and, when necessary, with additional input from a third reviewer (APA).

**Statistical analysis and data synthesis**

In our quantitative analysis, we included only studies reporting full data of true positives, true negatives, false positives and false negatives, or data obtained from eligible study publications or from authors directly. Other studies were used for qualitative synthesis only. Meta analysis was performed only if the absolute numbers of true-positive, true-negative, false-positive and false-negative results were provided or could be derived from at least two of the studies included.

The DTA measures are reported as point estimates with 95% confidence intervals (CIs). We derived the sensitivity, specificity and likelihood ratios (positive likelihood ratio +LR and negative likelihood ratio −LR) based on the true-positive, true-negative, false-positive and false-negative rates from each trial. Summary sensitivity, specificity, +LRs, −LRs and diagnostic odds ratios (DORs) also were derived using a bivariate random-effects model.

To visualize heterogeneity of diagnostic accuracy among the included studies, we plotted sensitivity and specificity of our index tests on forest plots using RevMan version 5.4 (Cochrane Training, London, UK). When meta analysis was appropriate (given the number of studies and extent of clinical heterogeneity), we pooled results from the included studies. Because our random-effects meta analysis was performed for a single threshold, we chose a bivariate model for binary results to determine summary estimates of sensitivity and specificity with 95% confidence and prediction regions. All meta-analyses were performed using MetaDTA version 2.01, an online application that uses statistical software R and the existing packages Shiny and lm4.

**Results**

**Study characteristics**

Our search on 15 November 2020 identified 1,389 citations in all three databases and, after removing duplicates, 994 titles were screened for eligibility for inclusion. The study flow diagram is shown in Figure 1. Of these, four studies reported the role of UCPCR among 836 patients with T2DM; seven studies studied that in 3,395 patients with T1DM and six studies studied the role of UCPCR among 257 patients with MODY (Table 1).

The assessment of risk of bias and applicability via QUADAS-2 is presented in Figure 2 and Figure 3. Overall, the risk of bias was scored as ‘low’ or ‘concern’ in 100% of studies for the index test, ‘unclear’ in 42% for the reference standard test, and ‘high’ or ‘concern’ in 55% for flow and timing and 80% in the patient selection domain. The applicability of studies was scored as ‘low concern’ in 100% for patient selection and index test, and ‘unclear’ in 35% for reference standard test.

Only four studies reported (or authors shared) the full data for appropriate quantitative synthesis; therefore, complete meta-analysis could be performed. The study by Yılmaz Ağacıoğlu et al. included...
Total number of articles identified through database search: 1,389 (Medline 449, Embase 701, Cochrane Library 239)

395 duplicates removed

Titles and abstracts screened for eligibility: 994

969 records excluded

3 articles with full DTA data for T1DM, 2 with full DTA for MODY and 3 with full DTA data on T2DM

7 articles with DTA data for T1DM, 6 with DTA for MODY and 4 with DTA data on T2DM

13 full text papers were excluded (diagnostic accuracy test not done: 13, review articles: 2)

Full text articles assessed for eligibility: 25

13 full text papers were excluded (diagnostic accuracy test not done: 13, review articles: 2)

Studies included in the qualitative synthesis: 10

Studies included in the quantitative synthesis: 4

DTA = Data Transfer Agreement; MODY = maturity-onset diabetes of the young; T1DM = type 1 diabetes mellitus; T2DM = type 2 diabetes mellitus.

Table 1: Characteristics of individual studies included in the review

| Study, year, country | N | Index test (UCPCR cut-off in nmol/mol) | Sensitivity, % | Specificity, % | PPV, % | NPV, % |
|----------------------|---|---------------------------------------|---------------|---------------|--------|--------|
| **Genetic testing for MODY and exclusion of T1DM by antibody testing** | | | | | | |
| Shields et al, 2017, UK | 51 | >0.2 | 98.0 | 85.0 | 20.0 | 99.9 |
| Shepherd et al, 2016, UK | 20 | >0.2 | 99.0 | N/A | N/A | N/A |
| Yılmaz Ağabagli et al, 2015, Turkey | 27 | >0.2 | 96.3 | 85.7 | 81.3 | 97.9 |
| Besser et al, 2013, UK | 39 | >0.7 | 100.0 | 97.0 | N/A | N/A |
| Besser et al, 2011, UK | 97 | >0.2 | 97.0 | 96.0 | N/A | N/A |
| Wang et al, 2019, China | 23 | >0.2 | 87.0 | 92.7 | 83.3 | 94.6 |
| **Positive T1DM by antibody tests and exclusion of MODY by genetic testing** | | | | | | |
| Shields et al, 2017, UK | 1,362 | <0.2 | N/A | N/A | N/A | N/A |
| Shepherd et al, 2016, UK | 817 | <0.2 | N/A | N/A | N/A | N/A |
| Yılmaz Ağabagli et al, 2015, Turkey | 42 | <0.2 | 85.7 | 96.3 | 97.3 | 81.3 |
| Oram et al, 2015, UK | 924 | <0.2 | N/A | N/A | N/A | N/A |
| Besser et al, 2013, UK | 120 | <0.7 | 100.0 | 97.0 | N/A | N/A |
| Liu et al, 2019, China | 61 | <0.2 | 68.9 | 95.8 | 67.7 | 96.0 |
| Besser et al, 2011, UK | 69 | <0.2 | N/A | N/A | N/A | N/A |
| **90 minutes post-mixed-meal serum C-peptide (if UCPCR <0.2 nmol/mmol) and exclusion of T1DM by antibody testing** | | | | | | |
| Hope et al, 2013, UK | 188 | >0.2 | 96.0 | 100.0 | 100.0 | 27.0 |
| Hope et al, 2016, UK | 41 | <0.2 | 83.0 | 93.0 | 90.0 | N/A |
| **WHO criteria for T2DM diagnosis and exclusion of T1DM by antibody testing** | | | | | | |
| Liu et al, 2019, China | 471 | >0.2 | 95.8 | 68.9 | 96.0 | 67.7 |
| **WHO diagnostic criteria for T2DM, exclusion of T1DM by antibody testing and exclusion of MODY by genetic testing** | | | | | | |
| Wang et al, 2019, China | 136 | >0.2 | 81.6 | 92.9 | 96.5 | 67.5 |

MODY = maturity-onset diabetes of the young; N/A = not available; N = number; NPV = negative predictive value; PPV = positive predictive value; T1DM = type 1 diabetes mellitus; T2DM = type 2 diabetes mellitus; UCPCR = urine C-peptide creatinine ratio; WHO = World Health Organization.
adolescents and children aged <20 years with diabetes duration ≥2 years. In the study by Hope et al., patients with T2DM had a median duration of disease of 13 years (interquartile range [IQR] 9–17 years) and a median age of 58 years (IQR 50–66 years). In the study by Liu et al., the mean age was 42.9 ± 18.5 years for patients with T1DM (age at diagnosis: 31.8 ± 17.2 years) and 56.2 ± 13.4 years for patients with T2DM (age at diagnosis: 44.1 ± 11.5 years). In the Wang et al. study, median duration of diabetes in patients with T1DM, monogenic diabetes and T2DM was 6.5 years (IQR 2.3–13.0 years), respectively. In the Wang et al. study, the median age of patients was 46.0 years (IQR 26.5–59.5 years), 35.0 years (IQR 30.0–47.0 years) and 53.0 years (IQR 42.0–60.0 years), respectively.

Diagnostic value of urine C-peptide creatinine ratio for diagnosing type 1 diabetes mellitus

The sensitivity of UCPCR for diagnosing T1DM varied from 82% (95% CI 74–88%) to 96% (95% CI 92–98%) using the cut-off value of <0.2 nmol/mmol, while the specificity was 69% (95% CI 56–80%) to 100% (95% CI 29–100%). The pooled estimate of sensitivity was 84.4% (95% CI 68.1–93.2%) and the pooled estimate of specificity was 91.6% (95% CI 82.8–96.1%, Figure 4). The pooled estimate of false-positive rate was 8.4% (95% CI 3.9–17.2%). The DOR was estimated as 59.9 (95% CI 32.8–106.0). The +LR was 10.1 (95% CI 5.3–19.2) and the -LR was 0.17 (95% CI 0.08–0.36).

Full data on the sensitivity, specificity, positive predictive value (PPV) and negative predictive value (NPV) were not available from four studies, and the study by Besser et al. in 2013 used a UCPCR cut-off of <0.7 nmol/mol; therefore, we could not include these studies in the meta-analysis. The available data on these studies are depicted in Table 1.

Diagnostic value of urine C-peptide creatinine ratio for diagnosing type 2 diabetes mellitus

The reported sensitivity of UCPCR for diagnosing T2DM varied from 81.6% to 96.0% keeping the cut-off value of >0.2 nmol/mol, while the
Figure 4: (A) Forest plot for sensitivity and specificity of urine C-peptide creatinine ratio in type 1 diabetes mellitus. (B) Summary receiver operating characteristic curve for sensitivity and specificity of urine C-peptide creatinine ratio in type 1 diabetes mellitus.\(^{21,22,26}\)

| Study                  | TP   | FP   | FN   | TN   | Sensitivity (95% CI) | Specificity (95% CI) | Sensitivity (95% CI) | Specificity (95% CI) |
|------------------------|------|------|------|------|----------------------|----------------------|----------------------|----------------------|
| Yılmaz Ağadioglu et al. 2015\(^{26}\) | 36   | 1    | 6    | 26   | 0.86 (0.71, 0.95)    | 0.96 (0.81, 1.00)    |                      |                      |
| Liu et al., 2019\(^{21}\)   | 42   | 20   | 19   | 451  | 0.69 (0.56, 0.80)    | 0.96 (0.94, 0.97)    |                      |                      |
| Wang et al., 2019\(^{22}\)   | 52   | 24   | 4    | 111  | 0.93 (0.83, 0.98)    | 0.82 (0.75, 0.88)    |                      |                      |

CI = confidence interval; FN = false negative; FP = false positive; TN = true negative; TP = true positive.

Figure 5: (A) Forest plot for sensitivity and specificity of urine C-peptide creatinine ratio in type 2 diabetes mellitus. (B) Summary receiver operating characteristic curve for sensitivity and specificity of urine C-peptide creatinine ratio in type 2 diabetes mellitus.\(^{21,22,24}\)

| Study                  | TP   | FP   | FN   | TN   | Sensitivity (95% CI) | Specificity (95% CI) | Sensitivity (95% CI) | Specificity (95% CI) |
|------------------------|------|------|------|------|----------------------|----------------------|----------------------|----------------------|
| Hope et al., 2013\(^{24}\) | 177  | 0    | 8    | 3    | 0.96 (0.92, 0.98)    | 1.00 (0.29, 1.00)    |                      |                      |
| Liu et al., 2019\(^{21}\)   | 451  | 19   | 20   | 42   | 0.96 (0.94, 0.97)    | 0.69 (0.56, 0.80)    |                      |                      |
| Wang et al., 2019\(^{22}\)   | 111  | 4    | 25   | 52   | 0.82 (0.74, 0.88)    | 0.93 (0.83, 0.98)    |                      |                      |

CI = confidence interval; FN = false negative; FP = false positive; TN = true negative; TP = true positive.
Diagnostic value of urine C-peptide creatinine ratio for diagnosing maturity-onset diabetes of the young

The reported sensitivity of UCPCR for diagnosing MODY varied from 95.0% to 96.0%, keeping the cut-off value of >0.2 nmol/mol, while the specificity was 88–96%. The pooled estimate of sensitivity was 85.2% (95% CI 73.1–92.4%) and the pooled estimate of specificity was 98.0% (95% CI 92.4–99.5%; Figure 6).21,22,26

The pooled estimate of false-positive rate was 2.0% (95% CI 0.5–7.6%). The +LR was 5.04 (95% CI 2.3–11.1) and the -LR was 0.09 (95% CI 0.05–0.17).

A sensitivity analysis for studies with patients with T1DM or T2DM is shown in Table 2.21,22,24

The study by Hope et al. reported the use of UCPCR for identifying insulin deficiency among patients with T2DM, but no other studies reported comparable data in this review and therefore could not be meta-analysed.24

Discussion

Main findings

This systematic review assessed the use of UCPCR for correctly identifying the type of diabetes in ethnically diverse populations from the UK, China and Turkey; however, the ethnic representation of the entire study does not correctly reflect global diversity. Nonetheless, UCPCR is a relatively easy laboratory test to perform, compared with serum C-peptide assay, for identifying absolute insulin deficiency, and can be performed in diverse clinical and experimental settings to replicate the results from this study.

We found considerable heterogeneity among studies, with concerns about moderate risk of bias and applicability of the index test in the studies included in this review. Moreover, several studies did not report full data for the meta-analysis and, therefore, pooled estimates on the sensitivity, specificity and DOR could not be derived for all the studies.

The pooled estimates on the value of a UCPCR cut-off of >0.2 nmol/mol in the studies in patients with T1DM (n=130) indicate moderate sensitivity (84.4%), high specificity (91.6%) and moderate DOR (59.9) for the test in the appropriate clinical settings (as mentioned in the Methods section). Similarly, a UCPCR cut-off of >0.2 nmol/mol among participants with T2DM (n=739) indicates high sensitivity (92.8%), moderate specificity (81.6%) and moderate DOR (58.9) for correctly diagnosing T2DM. However, we have to be mindful about using the test in patients with longstanding and poorly controlled T2DM, or in those with some forms of monogenic diabetes, as insulin deficiency in these patients is not uncommon and the biological behaviour of the disease often simulates that of T1DM.29–31

The pooled estimates on the value
of a UCPCR cut-off of >0.2 nmol/mol in participants with monogenic diabetes (n=46) show the test has moderate sensitivity (85.2%) and high specificity (98.0%) and DOR (281.8) when performed in appropriate clinical settings.

Limitations and the strengths of this review

Although we could procure nine different studies (seven each reporting the value of UCPCR for identifying T1DM and MODY) for the qualitative synthesis, we could obtain full data only from four studies for the meta-analysis. The total number of patients from these four studies was only 915, which is small for a common disease such as diabetes mellitus, and hampers the generalizability of the results. Moreover, lack of representation of major ethnic groups and nationalities other than Caucasian (British), Chinese and Turkish, and the under-representation of different age groups (such as the elderly or children), might limit the applicability of the results across demographics. Despite these limitations, ours is the first review examining the use of UCPCR – a simple and easily performed laboratory test – that shows moderate-to-high sensitivity, specificity and DOR for correctly identifying the main types of diabetes in an appropriate clinical setting. From a clinician’s perspective, classifying the three main subtypes of diabetes can facilitate individualized treatment and thereby optimize glycaemic control, minimize risk and reduce healthcare costs.

Applicability of findings to the review question and conclusions

The studies evaluated in this DTA systematic review revealed moderate concerns of risk of bias and applicability among three ethnically diverse populations from the UK, China and Turkey. A UCPCR cut-off of >0.2 nmol/mol has a sensitivity of 84.4%, a specificity of 91.6% and a DOR of 59.9 for correctly identifying patients with T1DM. A UCPCR cut-off of >0.2 nmol/mol appears to confer sensitivity, specificity and DOR of 92.8%, 81.6% and 56.9, respectively, among patients with T2DM in the appropriate clinical setting. Among patients with monogenic diabetes, a UCPCR cut-off of >0.2 nmol/mol achieves a diagnostic sensitivity of 85.2%, specificity of 98.0% and DOR of 281.8 when T1DM and T2DM are excluded by appropriate clinical and biochemical profiling. Large multinational studies with demographically diverse populations are required to optimally analyse the utility of this laboratory test for routine clinical practice.

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