Detection of C-Peptide in Urine as a Measure of Ongoing Beta Cell Function

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Abstract

C-peptide is a protein secreted by the pancreatic beta cells in equimolar quantities with insulin, following the cleavage of proinsulin into insulin. Measurement of C-peptide is used as a surrogate marker of endogenous insulin secretory capacity. Assessing C-peptide levels can be useful in classifying the subtype of diabetes as well as assessing potential treatment choices in the management of diabetes.

Standard measures of C-peptide involve blood samples collected either fasted or, most often, after a fixed stimulus (such as oral glucose, mixed meal, or IV glucagon). Despite the established clinical utility of blood C-peptide measurement, its widespread use is limited. In many instances this is due to perceived practical restrictions associated with sample collection.

Urine C-peptide measurement is an attractive noninvasive alternative to blood measures of beta-cell function. Urine C-peptide creatinine ratio measured in a single post stimulated sample has been shown to be a robust, reproducible measure of endogenous C-peptide which is stable for three days at room temperature when collected in boric acid. Modern high sensitivity immunoassay technologies have facilitated measurement of C-peptide down to single picomolar concentrations.

Keywords: C-peptide, Urine, UCPCR, Endogenous, Insulin

1 Introduction

Insulin secretory capacity can be assessed by measuring C-peptide, a protein secreted by the pancreas in equimolar quantities with insulin, following the enzymatic cleavage of proinsulin into insulin (Fig. [1](#page-1-0)). C-peptide is an attractive surrogate marker of insulin secretion because it can be used in patients administered exogenous insulin therapy and it is more reflective of insulin secretion than insulin itself, due to the variable clearance of insulin by the liver before it reaches the peripheral circulation $[1]$ $[1]$. C-peptide metabolism largely occurs in the kidney through glomerular filtration and uptake by tubular cells from peritubular capillaries, with 5–10 % normally excreted in the urine. This is in contrast to insulin, 50 % of which is metabolized and extracted in the liver. The total quantity of C-peptide excreted in the urine per day represents 5 % of pancreatic secretion, compared to only 0.1 % of secreted insulin [[2](#page-8-0)].

Fig. 1 Diagram of pro-insulin demonstrating the A chain and B chain of insulin and the connecting peptide (C-peptide), figure adapted from Stoy et al 2007^{[5](http://c/Users/200078/AppData/Local/Microsoft/Windows/Temporary%20Internet%20Files/Content.Outlook/0AYB15KQ/C-peptide_chapter_draft_MPTM060814.docx#_ENREF_5%23Stoy,%202007%20%23528)}. During the biosynthesis of insulin, the C peptide promotes proper protein folding and disulfide bonds between the A and B chains

Consequently, despite equimolar secretion, C-peptide has a longer half-life of approximately 30 min compared with only 6 min for insulin and occurs in the blood in concentrations up to five times higher than insulin $\lceil 3, 4 \rceil$.

The assessment of endogenous C-peptide production in diabetes can be useful in classifying the subtype of diabetes. In a patient with young onset diabetes persistent C-peptide production may reflect the honeymoon period of a patient with Type 1 diabetes, but also enduring C-peptide can be a feature of other types of diabetes including Type 2 diabetes where levels are typically high.

There is extensive evidence that C-peptide can be used to differentiate between the classification of Type 1 diabetes and Type 2 diabetes [[5–9\]](#page-8-0). C-peptide is a good candidate biomarker to differentiate patients with Maturity Onset Diabetes of the Young (MODY) from Type 1 diabetes. Type 1 diabetes is characterized by autoimmune destruction of beta-cells and ultimately results in absolute insulin deficiency, usually within 5 years of diagnosis [[10\]](#page-8-0). In patients with a genetic diagnosis of MODY, beta cell

1.1 C-Peptide and Differentiating Subtypes of Diabetes function is typically maintained with time despite a reduction in insulin secretion and progressive hyperglycemia [[11\]](#page-8-0). The ISPAD Clinical Practise Consensus Guidelines "Compendium on the diagnosis and management of monogenic diabetes in children and adolescents" published in 2009 $\lceil 12 \rceil$ and best practice guidelines for the molecular genetic diagnosis of maturity-onset diabetes of the young, published in 2008 $[13]$ $[13]$ both acknowledge that persistent C-peptide production can be used to identify patients likely to harbor a mutation in a MODY gene. Both guidelines suggest after the initial honeymoon period in patients with suspected Type 1 diabetes (typically >5 years) a blood C-peptide level of >200 pmol/L suggests a possible diagnosis of MODY. These guidelines are supported by recent reports which have shown that a urine C-peptide can be used to discriminate Type 1 diabetes from MODY with a high sensitivity and specificity $[14, 15]$ $[14, 15]$ $[14, 15]$ $[14, 15]$ $[14, 15]$

1.2 Measurement of C-Peptide 1.2.1 Blood C-Peptide Standard measures of C-peptide involve blood samples collected either fasted or, most often, after a fixed stimulus (such as oral glucose, mixed meal or IV glucagon) [[1,](#page-8-0) [16,](#page-8-0) [17](#page-8-0)]. Despite the established clinical utility of blood C-peptide measurement, its widespread use is limited. In many instances this is due to practical restrictions associated with sample collection. Both insulin and C-peptide concentrations are widely considered to be unstable and require specific pre-analytical handling procedures with many laboratory providers stipulating the need for blood samples to be rapidly centrifuged after collection on ice and the serum or plasma immediately frozen following separation from cells [[1,](#page-8-0) [18\]](#page-8-0). However, these strict handling protocols may not be necessary as there is evidence that C-peptide is stable for up to 24 h on whole blood if collected into EDTA preservative [[19](#page-8-0)].

1.2.2 Urine C-Peptide There is a body of evidence that suggests 24 h urine C-peptide (UCP) levels provide an accurate means of assessing beta cellsecretory capacity and correlate with both fasting and stimulated serum insulin and C-peptide [\[2](#page-8-0), [7](#page-8-0), [20,](#page-8-0) [21](#page-8-0)].

> However, the cumbersome nature and difficulties in obtaining an accurate and complete 24 h urine collection has limited the utility of this test $\lceil 22-24 \rceil$. Urine samples collected over a period of 4 h and single post stimulated UCP have been shown to be significantly correlated with serum insulin and serum C-peptide in nondiabetic subjects $[25]$ $[25]$ $[25]$, and in insulin treated diabetic patients [[6\]](#page-8-0). Correcting for creatinine adjusts urine C-peptide concentration for variation in urine concentration and enables the use of "spot" urine samples in place of 24-h urine collection. Urinary C-peptide when collected in boric acid is stable for up to 3 days at room temperature making it possible to be collected remotely from the processing laboratory and even posted into the laboratory by

patients [[26\]](#page-9-0). Urine C-peptide creatinine ratio correlates strongly with blood C-peptide and has been validated in nondiabetics, type 1 diabetes, type 2 diabetes, adults and children [[14,](#page-8-0) [15](#page-8-0), [27–34\]](#page-9-0).

1.2.3 C-Peptide Assays Early assays for C–peptide were radioimmunoassays and suffered from poor analytical sensitivity and specificity and are timeconsuming to perform (and therefore expensive) $[1, 35, 36]$ $[1, 35, 36]$ $[1, 35, 36]$ $[1, 35, 36]$ $[1, 35, 36]$ $[1, 35, 36]$. The introduction of non-isotopic assays (chemiluminescence, fluorescence, etc.) that utilize monoclonal antibodies has improved analytical sensitivity, specificity, reproducibility and reduced assay costs (to approximately $\pounds 10$). These technologies are also amenable to automation, allowing these assays to be incorporated into large high throughput clinical analysers.

2 Materials

- Urine collected into boric acid container. For interpretation of beta-cell reserve, the sample should be a stimulated sample, collected 2 h after the biggest meal of the day (see Methods section on patient preparation).
- Roche Modular Analytics E170 immunoassay analyzer (Roche Diagnostics, Mannheim, Germany).
- C-peptide calibration material: Roche C-peptide Calibration Set (Roche Diagnostics, Mannheim, Germany).
- C-peptide quality control material: Elecsys Precicontrol Multianalyte (Roche Diagnostics, Mannheim, Germany).
- Roche C-peptide chemiluminescence assay kit (Roche Diagnostics, Mannheim, Germany).
- Diluent used for C-peptide assay (Equine serum albumin): Roche Diluent MultiAssay (Roche Diagnostics, Mannheim, Germany).
- Roche Creatinine modified Jaffe assay kit (Roche Diagnostics, Mannheim, Germany).
- Creatinine calibration material: Roche Calibrator for automated systems (Roche Diagnostics, Mannheim, Germany).
- Creatinine QC material: Randox assayed urine control level 2 and 3 (Randox UK, Co. Antrim, UK).

3 Methods

Patient preparation: To assess the maximum endogenous C-peptide response, a stimulated urine sample should be collected. Patients should be briefed with the following information before sample collection:

1. Pass urine just before the biggest meal of the day and discard. 2. Eat the meal as usual with a glass or more of water. 3. Do not eat anything else for the next 2 h unless you have a

- hypoglycemic episode, in which case you should do this test on another day. You can drink water freely throughout the duration of the collection.
- 4. 2 h after this meal, collect urine and send for analysis.

Urine C-peptide concentration is unchanged at room temperature for 24 h and at 4 \degree C for 72 h even in the absence of preservative.

Urine C-peptide collected in boric acid is stable at room temperature for 72 h. Urine C-peptide remained stable after seven freeze–thaw cycles but decreased with freezer storage time and dropped to 82–84 % of baseline by 90 days at -20 °C [\[26](#page-9-0)].

C-peptide analysis is performed on the high throughput Roche Modular Analytics E170 immunoassay analyzer (Roche Diagnostics, Mannheim, Germany). The assay is a heterogeneous sandwich immunoassay. Urine is diluted 10-fold with Multi Assay Diluent. One biotinylated anti-C-peptide specific monoclonal mouse antibody and a second monoclonal antibody to C-peptide labeled with a ruthenium complex, react with C-peptide in 20 μL of diluted urine sample to form an antigen–antibody–antigen sandwich complex. Separation is achieved via interaction of biotin and streptavidin attachment to paramagnetic microparticles (solid phase). The detection system employs electrochemiluminescence with ruthenium trisbipyridyl as the label.

Electrochemiluminescence occurs at 620 nm and readings are taken by the photomultiplier tube. The intensity of light signal is proportional to the concentration of C-peptide in the serum. All urine samples were pre-diluted 1:10 with equine serum albumin (diluent multianalyte, Roche Diagnostics, Mannheim, Germany) (see Note [2](#page-6-0)). The analysis time is 18 min per specimen. The E170 is able to analyze 170 samples per hour.

The assay was calibrated using Roche C-peptide CalSet calibration material (Roche Diagnostics, Mannheim, Germany), traceable to WHO International Reference Reagent (IRR) for C-peptide of human insulin for immunoassay (IRR code 84/510) [[37](#page-9-0)]. Quality Control was performed on each day of analysis using low level (0.67 nmol/L) and high level (3.33 nmol/L) PreciControlMultiAnalyte.

3.2.1 *Measuring Range* The measuring range of the urine C-peptide assay is from 0.03 to 133 nmol/L.

> There is approximately 30 % cross-reactivity with pro-insulin (see Note [3](#page-6-0))

3.1 Sample Storage and Stability

3.2 Roche Assay Used to Measure C-Peptide in Urine

4 Notes

- 1. Agreement of urine C-peptide measurements in different commercially available assays—There is poor comparability between commercial and in-house C-peptide assays making transferability of C-peptides values/cutoffs from research to clinical practice problematic [\[39–41](#page-9-0)]. International working groups (in the USA and Europe) are currently addressing the widely disparate values between analytical methods for both C-peptide and insulin $\lceil 39, 41, 42 \rceil$ $\lceil 39, 41, 42 \rceil$. The groups are combining efforts to improve standardization with the aim of establishing a complete reference measurement system and certified primary reference materials based on pure biosynthetic insulin and C-peptide [[42\]](#page-9-0).
- 2. Using other diluents in the Roche C-peptide assay—Cpeptide is approximately 20-fold more concentrated in urine samples and therefore require dilution before analysis. We have found that the diluent must be proteinaceous (e.g., equine serum albumin) in order to achieve both a concentration and matrix that will generate a comparable and reproducible result. This has been shown for three different platforms measuring C-peptide (Roche, Immulite, and Centaur). (*unpublished data derived at the Blood Sciences Dept., RD&E).
- 3. Cross reactivity of the Roche assay with proinsulin— Proinsulin does cross react with the Roche assay (30 % cross reactivity) despite the fact that the assay utilizes two site monoclonal antibodies. However, proinsulin and split products exist in much lower concentrations than C-peptide in serum \langle <2 % in nondiabetics) and are therefore of little clinical significance except for in rare conditions such as insulinoma $[43]$ $[43]$. In addition, the total quantity of C-peptide excreted in the urine per day represents 5–10 % of pancreatic secretion, compared to only 0.05 % of secreted proinsulin (2–3 times lower than insulin of which only 0.1 % is excreted into the urine $[44]$ $[44]$). With only 30 % cross reactivity in the Roche assay the contribution of proinsulin to urine C-peptide measurement would be negligible.
- 4. Making a diagnosis in insulin treated patients—The principal role of urinary C-peptide is to identify insulin insufficiency, a feature of long term Type 1 diabetes, in insulin treated patients. We have found that >95 % Type 1 diabetes (>5 year duration) have a UCPCR value < 0.2 nmol/mmol.
	- Type 1 v $MOD\rightarrow$ cutoff UCPCR of 0.2 nmol/mmol differentiates HNF1A/4A MODY from Type 1 diabetes with a sensitivity of 97 % and specificity of 96 % (ROC 0.98) $[14]$.
	- Type 1 v Type 2—A cutoff of 0.2 nmol/mmol differentiates Type 2 diabetes from Type 1 diabetes with 94 % sensitivity and 94 % specificity (ROC AUC 0.94).
- 5. Monitoring the honeymoon period in Type 1 diabetes—In patients with Type 1 diabetes it is possible to measure the extent to which they are progressing through the honeymoon phase. UCPCR is highly correlated with the serum C peptide in a formal mixed tolerance test [\[31\]](#page-9-0).
- 6. Assessing in insulin treated Type 2 diabetes the extent of endogenous insulin secretion—Endogenous insulin secretion can be measured in Type 2 diabetes. UCPCR is highly correlated with the serum C-peptide in a formal mixed tolerance test [28]. Patients with a high endogenous secretion >75 th centile are likely be those that benefit most from metformin and other oral agents. Patients with low endogenous insulin secretion <25th centile and particularly if UCPCR <0.02 nmol/mmol will most likely require insulin therapy.

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