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應用 HRM 分析方法快速偵測糖尿病患 Hb Iraq-Halabja 基因突變

劉素卿<sup>1,2</sup>, 林建佑<sup>1</sup>, 林彩秀<sup>1</sup>, 王秀娟<sup>1</sup>, 張建國<sup>1</sup>, 彭慶添<sup>2</sup>  
中國醫藥大學附設醫院<sup>1</sup>檢驗醫學部, <sup>2</sup>兒童醫學中心**Application of the rapid HRM analysis to detected Hb Iraq-Halabja in Diabetes patient**Su-Ching Liu<sup>1,2</sup>, Chien-Yu Lin<sup>1</sup>, Tsai-Hsiu Lin<sup>1</sup>, Shioh-Jain Wang<sup>1</sup>, Jan-Gowth Chang<sup>1</sup> and Ching-Tien Peng<sup>2</sup><sup>1</sup>Department of Laboratory Medicine and <sup>2</sup>Children's Medical Center, China Medical University Hospital, Taichung, Taiwan**BACKGROUND:**

Hemoglobin A1c (HbA1c) is used routinely to monitor long-term blood sugar control in people with diabetes, as HbA1c is related directly to risks for diabetic complications. Hb gene disorders are common hereditary diseases: e.g., alpha- or beta-thalassemia and Hb variant-related disease. More than 1000 abnormal Hb variants have been identified. This study was application of the rapid HRM analysis to detected Hb Iraq-Halabja in diabetes.

**METHODS and MATERIALS:**

In the present case study was a 50-year-old Taiwanese man and following a long time history of diabetes. We modify Shih et al.(2009) establish rapid identification of HBB gene mutations by HRM analysis.

**RESULTS:**

In this study, CBC data showed RBC  $4.73 \times 10^6 / \mu\text{L}$ , Hb 16.4 g/dL, mean corpuscular value (MCV) 100.5 fL, mean corpuscular hemoglobin (MCH) 34.6 pg, mean corpuscular hemoglobin concentration (MCHC) 34.4 g/dL, Ferritin 290.1 ng/ml and hemoglobin electrophoretic pattern (HbA 97.5%, HbA2 2.5%, HbA1C 2.5%). Herein, we application rapid HRM method to detected and find new Hb variant. In addition, we confirmed Hb  $\beta 10$  (A7) Ala  $\rightarrow$  Val (GCC  $\rightarrow$  GTC) (Hb Iraq-Halabja) with directly sequencing and PCR-RFLP

**CONCLUSIONS:**

This study demonstrates that HbA1C values determined by ion-exchange HPLC did not reflect the glycemic state in our patient. The ion-exchange HPLC to measurement HbA1C necessity combine with repeated using additional method e.g., application HRM method or hemoglobin gene whole gene sequence.

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利用成人健檢報告資訊評估腎絲球過濾速率之變異因子

林湘雲, 翁詩玉, 郭怡君, 李文琮  
郭綜合醫院檢驗科, 台南, 台灣**Utilization of adult health screening information to evaluate the variation factors of estimated glomerular filtration rate**Lin, Siang-Yun, Weng, Shih-Yu, Kuo, Yi-Chun, Lee, Wen-Tsung.  
Kuo General Hospital, Tainan, Taiwan

血清肌酸酐(Serum creatinine, Scr)是最常被用來評估腎臟功能的指標,但因肌酸酐的產生會受個體肌肉質量、飲食、年齡、性別..等多種因素影響,因此並非評估腎臟功能的最好指標。根據美國腎臟基金會建議,檢測腎絲球過濾速率(estimated glomerular filtration rate, eGFR)才能更精準的來判定慢性腎臟病的嚴重度,並定義只要 eGFR 持續低於 60 ml/min/1.73 m<sup>2</sup> 達 3 個月以上,就可以認定為慢性腎臟病。

本研究利用 2012 年 1~6 月 1190 位接受成人健檢者之報告資訊,初步依據美國腎臟基金會標準將 eGFR 結果分成腎功能正常、輕度、中度、重度慢性腎衰竭及末期腎臟疾病等 5 種類別( $\geq 90$ , 60-90, 30-59, 15-29,  $<15$ ),分別與 Scr ( $>1.3$  mg/dl)之臨床意義比較,除第五類無個案外,其餘 1~4 類別 eGFR 結果為  $102.4 \pm 11.0$ ,  $76.8 \pm 7.6$ ,  $51.8 \pm 6.6$ ,  $22.5 \pm 5.5$ ; Scr 結果為  $0.7 \pm 0.1$ ,  $0.9 \pm 1.5$ ,  $1.3 \pm 0.2$ ,  $2.6 \pm 0.9$ 。我們發現在 eGFR 分類上已能呈現輕微功能異常之報告(第 2, 3 類),但 Scr 數據可能無法提早呈現相同之結果。進一步實際評估影響 eGFR 之可能變異因子(包含性別、年齡, Scr, Glucose, GOT, GPT, TG, Cholesterol (Chol), HDL-C, LDL-C, Urine protein (UP) 十一項),利用逐步多元迴歸分析方法,保留有統計意義之影響項目,發現性別,年齡, Scr, UP, Chol 等 5 個項目與 eGFR 之多元相關性達 0.886,影響 eGFR 之變異量達 78.5%。而影響變異量依序為 Scr (63%), 性別 (10.5%), 年齡 (4.7%), UP (2%), Chol (1%), 但有 21.5% 尚無法由上述項目解釋其變異來源。