

# Hemorheological parameters better classify metabolic syndrome than novel cardiovascular risk factors and peripheral vascular disease marker

Prajwal Gyawali<sup>a,\*</sup>, Ross S Richards<sup>a</sup>, Paul Tinley<sup>b</sup>, Ezekiel Uba Nwose<sup>c</sup> and Phillip T Bwititi<sup>d</sup>

<sup>a</sup>*Biomedical Sciences, School of Community Health, Charles Sturt University*

<sup>b</sup>*Podiatry, Charles Sturt University*

<sup>c</sup>*School of Community Health, Charles Sturt University*

<sup>d</sup>*School of Biomedical Sciences, Charles Sturt University*

## 1. Background

Hemorheological parameters are altered in metabolic syndrome (MetS) and its components [8–13]. Oxidative stress and chronic inflammation present in MetS are shown to be responsible for hemorheological changes to certain extent [7, 8]. In this brief report, we have presented the data that compare the association of MetS with hemorheological parameters (erythrocyte aggregation, erythrocyte deformability and whole blood viscosity (WBV)), oxidative stress (urinary isoprostanes), inflammation (high sensitivity C-reactive protein (hsCRP)), coagulopathy (D-dimer) and peripheral arterial disease (toe brachial pressure index (TBPI)).

## 2. Materials and methods

Erythrocyte deformability and erythrocyte aggregation was measured by RheoScan-AnD 300 system (RheoMeditech Inc., Korea). WBV measurement was carried out using a Brookfield DV-II+ programmable viscometer (MA, USA), using a CP40 spindle at 37°C at a shear rate of 150 s<sup>-1</sup>. Erythrocyte morphology was studied by scanning electron microscopy (JCM 5000, Benchtop SEM, Neoscope). All the rheological measurements were performed within two hours of blood collection after adjusting EDTA anticoagulated whole blood to the hematocrit of 40%. TBPI was measured by using SysToe (ATYS Medical). MetS was defined by National Cholesterol Education Program, Adult Treatment Panel III definition [6]. Inflammatory markers high sensitivity C-reactive protein (hsCRP) and thrombotic marker D-dimer were measured in the day of collection in a commercial clinical pathology laboratory. 15-isoprostanes F2t was measured in urine sample (NWLSS<sup>TM</sup>) and was expressed as ng of isoprostanes per mmol of

\*Corresponding author: Dr. Prajwal Gyawali (PhD), Biomedical sciences, School of Community health, Charles Sturt University, NSW 2640, Australia. E-mail: clbioprajwal@gmail.com.

28 urinary creatinine (Cayman chemical). The details of instrumentation and demographic characteristics of  
29 the participants have been published elsewhere [7–9, 13]. Briefly, 100 participants were recruited from  
30 a rural town of Australia from June–Dec 2013. Pregnant women, non-ambulatory patients, and children  
31 under 18 years of age were excluded from the study. Recruited participants were divided into three groups  
32 on the basis of absence or presence of MetS and its components. Group I consists of the participants with-  
33 out any positive components of MetS (healthy controls); group II consists of the participants with one or  
34 two positive components; and group III consists of participants with three or more positive components.  
35 Participants in groups I and II are non- MetS whereas participants of group III are with MetS.

### 36 3. Results

37 Of the 100 participants, 36 participants had MetS, 33 had one or two positive components and 33 were  
38 healthy controls.

#### 39 3.1. Binomial logistic regression analysis

40 Binomial logistic regression analysis (adjusted for age and sex) was performed to predict the chances of  
41 having MetS by altered hemorheological parameters; urinary isoprostanes, hsCRP, D-Dimer and TBPI.  
42 All of the markers were divided into quartiles and the odds of having MetS after increase or decrease  
43 ( $EI_{max}$ , TBPI) in one quartile of the markers was estimated. The results show that all of the markers  
44 significantly predicted MetS and the Odds ratio was highest for erythrocyte aggregation followed by  
45 erythrocyte deformability.

#### 46 3.2. ROC Curve analysis

47 The values of odds ratio obtained in the regression analysis depend on the range of data and the  
48 scaling. The regression coefficient represents the expected change in y (Mets/non-MetS) for a one unit  
49 change in x (the predictor: markers), hence, the magnitude of that coefficient is partly determined by  
50 the magnitude of the units used. Therefore, to confirm the outputs of logistic regression analysis, ROC  
51 curve was used to compare the association of different markers with MetS. The ROC curve shows the  
52 diagnostic performance of a test, or the accuracy of a test to discriminate two groups (MetS and non-  
53 MetS [14] and the area under the ROC curve (AUC) is a measure of how well a parameter can distinguish  
54 between two groups [14]. ROC curve analysis demonstrated that all the hemorheological components  
55 significantly classified MetS participants ( $P$ -values for all curves were  $< 0.0005$ ). AUC was higher for the  
56 hemorheological parameters (erythrocyte aggregation and erythrocyte deformability) than for the TBPI  
57 or other oxidative stress and inflammatory markers (Table 2 and Fig. 1).

### 58 4. Conclusions

59 Age and sex adjusted odds ratio for predicting MetS was higher for hemorheological parameters  
60 when compared to TBPI. The ROC curve analysis also showed that two of the three haemorheologi-  
61 cal parameters (critical stress and  $EI_{max}$ ) better classified MetS than TBPI. The finding suggests that  
62 hemorheology better identifies with MetS than macrovascular circulation abnormalities. Microvascular  
63 dysfunction (lower functional capillary density) has been shown in MetS participants [5]. Superiority of

Table 1

Age and sex adjusted odds ratio for predicting MetS by hemorheological parameters, Oxidative stress and inflammatory markers and TBPI

Parameters	Odds Ratio	95% CI	P-Value
Critical stress (quartile)	3.896	2.174 to 6.985	<0.0005
EI <sub>max</sub> (quartile)	2.840	1.666 to 4.830	<0.0005
WBV (quartile)	1.823	1.030 to 1.114	0.009
TBPI (quartile)	1.828	1.059 to 3.154	0.030
Urinary isoprostanes (quartile)	1.715	1.096 to 2.683	0.018
hsCRP (quartile)	2.090	1.297 to 3.370	0.002
D-dimer (quartile)	1.639	1.035 to 2.595	0.035

Table 2

AUC and 95% CI obtained from ROC curve analysis for differentiating MetS from non-MetS

Parameters	AUC	95% CI	P-value
Critical stress	0.818	0.715 to 0.922	<0.0005
EI <sub>max</sub>	0.782	0.688 to 0.876	<0.0005
TBPI	0.774	0.679 to 0.869	<0.0005
WBV	0.719	0.616 to 0.821	<0.0005
Urinary isoprostanes	0.706	0.603 to 0.809	0.001
D-dimer	0.695	0.583 to 0.807	0.001
hsCRP	0.661	0.549 to 0.774	0.008

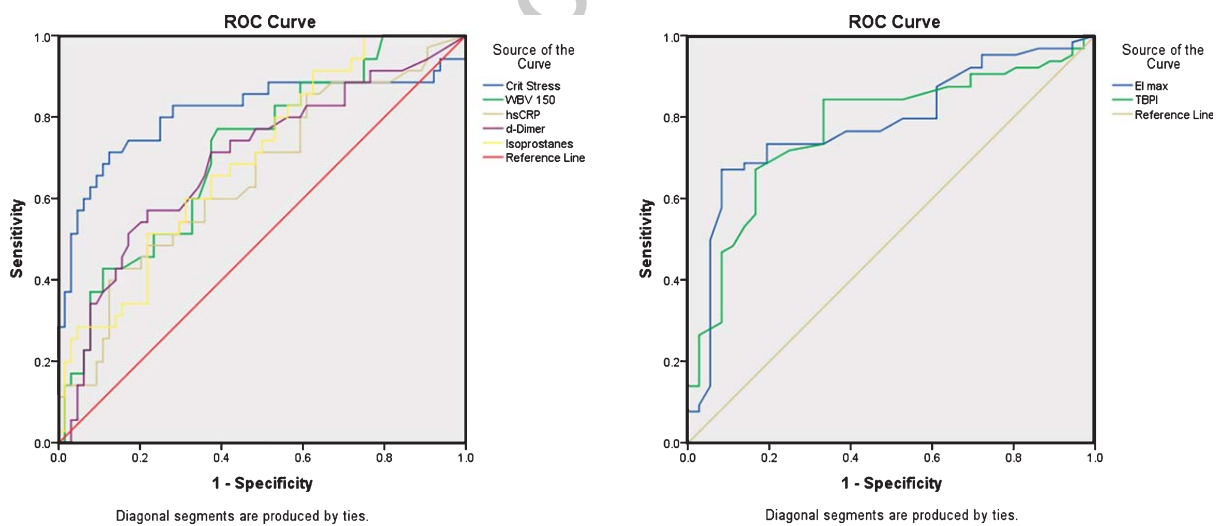


Fig. 1. ROC curve for haemorheological parameters, novel cardiovascular risk factors and peripheral vascular diseases marker for correctly classifying MetS.

64 the hemorheological parameters in predicting MetS than that of peripheral arterial disease marker further  
65 emphasises the importance that should be given to rheological changes occurring in the MetS along with  
66 macrovascular assessment. The present findings also suggests that rheological changes may occur earlier  
67 or more frequently than the peripheral vasculopathy in MetS and its early identification may provide  
68 clinical benefits to the MetS patients.

69 Insulin resistance is generally considered as a major factor for the pathogenesis of MetS [15]. Insulin  
70 resistance is associated with increased erythrocyte aggregation [4]. Brun JF et al. suggested that increased  
71 erythrocyte aggregation is an early phenomenon that characterises insulin resistance at an initial stage  
72 where it is compensated by an increase in insulin secretion [4] and the increased erythrocyte aggregation  
73 could be considered as a major hemorheological alteration of insulin resistance [3]. Moreover, increased  
74 erythrocyte aggregation has been reported among the obese subjects who are not under the state of MetS  
75 [2] signifying that role of adipocytokines and adiposity in hemorheological alterations. Similarly, in the  
76 present study, the AUC for erythrocyte aggregation (critical stress) was found to be higher than that  
77 of hsCRP and urinary isoprostanes. Also, since erythrocyte aggregation is significantly associated with  
78 oxidative stress and chronic inflammation generated in MetS, it could be included as a component of MetS.  
79 No studies have reported the ROC curve analysis of hemorheological parameters for the correct prediction  
80 of MetS making it difficult to make comparisons. However, it has been shown that increased erythrocyte  
81 aggregation correctly classified patients with vascular disease [1]. Furthermore, from the ROC curve  
82 analysis, AUC of erythrocyte aggregation for the correct classification of vascular disease was shown  
83 to be higher than that of ESR, fibrinogen and hsCRP [1]. Similarly, it has been shown that although  
84 conventional cardiovascular risk parameters such as triglyceride, HDL-C, LDL-C, total cholesterol, BMI  
85 and fibrinogen did not significantly predicted cardiac death, haematocrit/WBV significantly predicted the  
86 same (AUC = 0.716;  $P = 0.028$ ) [16]. The possibilities of the hemorheological components to be identified  
87 as better cardiovascular risk markers due to their strong association with MetS cannot be precluded from  
88 present findings.

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