

# South Africa - Glycated haemoglobin threshold for dysglycaemia screening and application to metabolic syndrome diagnosis in HIV-infected Africans

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## Overview

### Identification

#### ID NUMBER

HbA1cfordysglycaemiascreeninginHIV

### Overview

#### ABSTRACT

**Background:** Glycated haemoglobin (HbA1c) test has been increasingly promoted as an alternative to fasting plasma glucose (FPG) or oral glucose tolerance test (OGTT) to diagnose dysglycaemia but its performance in HIV-infected Africans has yet to be established. This study aimed to assess the diagnostic accuracy of HbA1c for dysglycaemia including FPG-defined and OGTT-defined dysglycaemia, and OGTT-defined diabetes in HIV-infected Africans, and the effect of HbA1c-predicted dysglycaemia on Joint Interim Statement (JIS)-based prevalent metabolic syndrome (MS).

**Methods:** A cross-sectional study included HIV-positive patients recruited across public healthcare facilities in the Western Cape. The recommended HbA1c cut-points were tested alongside the optimal cut-points obtained from receiver operating characteristic curve analyses, while the agreement between the MS criteria were assessed using kappa statistic.

**Results:** 748 participants (157 men), median age 38 years, 93% on anti-retroviral drugs were included. The optimal HbA1c cut-points of 5.75% (39.3 mmol/mol) showed 54% sensitivity, 84% specificity for FPG-defined dysglycaemia, and 52% sensitivity, 85% specificity for OGTT-defined dysglycaemia. The HbA1c value of 5.85% (40.4 mmol/mol) (63% sensitivity, 99% specificity) was optimal for diabetes. The internationally advocated cut-point of 6.5% (48 mmol/mol) had 37% sensitivity and 99% specificity for diabetes, while HbA1c =5.7% (=39 mmol/mol) yielded similar performance with the study-specific cut-point for any dysglycaemia. MS prevalence by the JIS criteria (28.2%) increased to 29.7% when using HbA1c =5.75% (=39.3 mmol/mol) and to 32.9% with HbA1c =5.7% (=39 mmol/mol); agreement between the original and modified criteria was generally good.

**Conclusions:** This study agrees with the internationally recommended HbA1c cut-point for detecting dysglycaemia, but not for diabetes in HIV-infected Africans. In line with previous studies in general African populations, our findings suggest that similar factors interfere with HbA1c values regardless of HIV infection status. Replacing FPG-based with HbA1c-predicted dysglycaemia in the JIS criteria to diagnose MS is feasible in HIV-infected Africans.

### Producers and Sponsors

#### PRIMARY INVESTIGATOR(S)

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## Sampling

No content available

## Questionnaires

No content available

## Data Collection

### Data Collection Dates

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<b>Start</b>	<b>End</b>	<b>Cycle</b>
2014-03-01	2015-02-28	N/A

## Data Processing

No content available

## Data Appraisal

No content available

## **File Description**



## **Variable List**

## Data

Content

Cases 748

Variable(s) 34

Structure Type:  
Keys: ()

Version

Producer

Missing Data

## Variables

ID	Name	Label	Type	Format	Question
V1	PTId	PTId	discrete	character	
V2	Gender	Gender	discrete	numeric	
V3	Age	Age	contin	numeric	
V4	Weight1	Weight1	contin	numeric	
V5	Weight2	Weight2	contin	numeric	
V6	Height1	Height1	contin	numeric	
V7	Height2	Height2	contin	numeric	
V8	Waist1	Waist1	contin	numeric	
V9	Waist2	Waist2	contin	numeric	
V10	nd_SBP	nd_SBP	contin	numeric	
V11	nd_DBP	nd_DBP	contin	numeric	
V12	rd_SBP	rd_SBP	contin	numeric	
V13	rd_DBP	rd_DBP	contin	numeric	
V14	HIVDxy	HIVDxy	discrete	character	
V15	HIVDxm	HIVDxm	discrete	character	
V16	HIVDxw	HIVDxw	discrete	character	
V17	HIVDxd	HIVDxd	discrete	character	
V18	Value_CD4	Value_CD4	discrete	character	
V19	ART_use	ART_use	discrete	character	
V20	HTN_Rx	HTN_Rx	discrete	numeric	
V21	DM_Rx	DM_Rx	discrete	numeric	
V22	Chol_Rx	Chol_Rx	discrete	numeric	
V23	ALT	ALT	contin	numeric	
V24	AST	AST	discrete	character	
V25	HDLC	HDLC	contin	numeric	
V26	CHOLESTEROLS	CHOLESTEROLS	discrete	character	
V27	CREATININES	CREATININES	discrete	character	

<b>ID</b>	<b>Name</b>	<b>Label</b>	<b>Type</b>	<b>Format</b>	<b>Question</b>
V28	hsCRP	hsCRP	discrete	character	
V29	GAMMAGT	GAMMAGT	discrete	character	
V30	GLUCOSE2h	GLUCOSE2h	discrete	character	
V31	GLUCOSEf	GLUCOSEf	contin	numeric	
V32	HbA1c	HbA1c	contin	numeric	
V33	LDLC	LDLC	contin	numeric	
V34	TGS	TGS	contin	numeric	



## PTId (PTId)

### File: Data

#### Overview

Type: Discrete	Valid cases: 748
Format: character	Invalid: 0
Width: 3	

## Gender (Gender)

### File: Data

#### Overview

Type: Discrete	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 0	
Range: 1-2	

## Age (Age)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 0	Minimum: 18
Range: 18-72	Maximum: 72
	Mean: 38.5
	Standard deviation: 9

## Weight1 (Weight1)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 2	Minimum: 37.8
Range: 37.75-140.75	Maximum: 140.8
	Mean: 71.3
	Standard deviation: 17.6

## Weight2 (Weight2)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 2	Minimum: 37.7
Range: 37.7-140.75	Maximum: 140.8
	Mean: 71.3
	Standard deviation: 17.6

## Height1 (Height1)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 2	Minimum: 137.2
Range: 137.2-185.6	Maximum: 185.6
	Mean: 161.1
	Standard deviation: 7.6

## Height2 (Height2)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 2	Minimum: 136.8
Range: 136.8-185.4	Maximum: 185.4
	Mean: 161.1
	Standard deviation: 7.6

## Waist1 (Waist1)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 2	Minimum: 62
Range: 62-149.1	Maximum: 149.1
	Mean: 88.9
	Standard deviation: 14.4

## Waist2 (Waist2)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 2	Minimum: 61.8
Range: 61.8-149.5	Maximum: 149.5
	Mean: 89.1
	Standard deviation: 14.4

## nd\_SBP (nd\_SBP)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 0	Minimum: 77
Range: 77-218	Maximum: 218
	Mean: 121.3
	Standard deviation: 20.1

## nd\_DBP (nd\_DBP)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 0	Minimum: 48
Range: 48-134	Maximum: 134
	Mean: 83.7
	Standard deviation: 12.8

## rd\_SBP (rd\_SBP)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 0	Minimum: 76
Range: 76-223	Maximum: 223
	Mean: 120
	Standard deviation: 19.7

## rd\_DBP (rd\_DBP)

### File: Data

#### Overview

Type: Continuous	Valid cases: 748
Format: numeric	Invalid: 0
Decimals: 0	Minimum: 49
Range: 49-135	Maximum: 135
	Mean: 83
	Standard deviation: 12.8

## HIVDxy (HIVDxy)

### File: Data

#### Overview

Type: Discrete	Valid cases: 748
Format: character	Invalid: 0
Width: 2	

## HIVDxm (HIVDxm)

### File: Data

#### Overview

Type: Discrete	Valid cases: 748
Format: character	Invalid: 0
Width: 2	

## HIVDxw (HIVDxw)

### File: Data

## HIVD<sub>xw</sub> (HIVD<sub>xw</sub>)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 2

Valid cases: 748  
Invalid: 0

## HIVD<sub>xd</sub> (HIVD<sub>xd</sub>)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 3

Valid cases: 748  
Invalid: 0

## Value\_CD4 (Value\_CD4)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 4

Valid cases: 748  
Invalid: 0

## ART\_use (ART\_use)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 2

Valid cases: 748  
Invalid: 0

## HTN\_Rx (HTN\_Rx)

### File: Data

#### Overview

Type: Discrete  
Format: numeric  
Decimals: 0  
Range: 0-2

Valid cases: 748  
Invalid: 0

## DM\_Rx (DM\_Rx)

### File: Data

#### Overview

Type: Discrete  
Format: numeric  
Decimals: 0  
Range: 0-2

Valid cases: 748  
Invalid: 0



## Chol\_Rx (Chol\_Rx)

File: Data

### Overview

Type: Discrete  
 Format: numeric  
 Decimals: 0  
 Range: 0-1

Valid cases: 748  
 Invalid: 0

## ALT (ALT)

File: Data

### Overview

Type: Continuous  
 Format: numeric  
 Decimals: 0  
 Range: 6-380

Valid cases: 748  
 Invalid: 0  
 Minimum: 6  
 Maximum: 380  
 Mean: 29.3  
 Standard deviation: 26.2

## AST (AST)

File: Data

### Overview

Type: Discrete  
 Format: character  
 Width: 3

Valid cases: 748  
 Invalid: 0

## HDLC (HDLC)

File: Data

### Overview

Type: Continuous  
 Format: numeric  
 Decimals: 2  
 Range: 0.5-3.27

Valid cases: 748  
 Invalid: 0  
 Minimum: 0.5  
 Maximum: 3.3  
 Mean: 1.3  
 Standard deviation: 0.4

## CHOLESTEROLS (CHOLESTEROLS)

File: Data

### Overview

Type: Discrete  
 Format: character  
 Width: 17

Valid cases: 748  
 Invalid: 0

## CREATININES (CREATININES)

File: Data

### Overview

## CREATININES (CREATININES)

### File: Data

Type: Discrete  
Format: character  
Width: 3

Valid cases: 748  
Invalid: 0

## hsCRP (hsCRP)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 17

Valid cases: 748  
Invalid: 0

## GAMMAGT (GAMMAGT)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 3

Valid cases: 748  
Invalid: 0

## GLUCOSE2h (GLUCOSE2h)

### File: Data

#### Overview

Type: Discrete  
Format: character  
Width: 17

Valid cases: 748  
Invalid: 0

## GLUCOSEf (GLUCOSEf)

### File: Data

#### Overview

Type: Continuous  
Format: numeric  
Decimals: 2  
Range: 3.2-21.2

Valid cases: 748  
Invalid: 0  
Minimum: 3.2  
Maximum: 21.2  
Mean: 5.4  
Standard deviation: 1.7

## HbA1c (HbA1c)

### File: Data

#### Overview

## HbA1c (HbA1c)

### File: Data

Type: Continuous  
Format: numeric  
Decimals: 2  
Range: 4.2-13.6

Valid cases: 748  
Invalid: 0  
Minimum: 4.2  
Maximum: 13.6  
Mean: 5.6  
Standard deviation: 0.8

## LDLC (LDLC)

### File: Data

#### Overview

Type: Continuous  
Format: numeric  
Decimals: 2  
Range: 0.4-5.6

Valid cases: 748  
Invalid: 0  
Minimum: 0.4  
Maximum: 5.6  
Mean: 2.6  
Standard deviation: 0.9

## TGS (TGS)

### File: Data

#### Overview

Type: Continuous  
Format: numeric  
Decimals: 2  
Range: 0.35-6.99

Valid cases: 748  
Invalid: 0  
Minimum: 0.4  
Maximum: 7  
Mean: 1.1  
Standard deviation: 0.6

