

## Evaluation of the Finnish Diabetes Risk Score (FINDRISC) as a Screening Tool for the Metabolic Syndrome

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### ■ Abstract

**OBJECTIVES:** Traditionally, the Finnish Diabetes Risk Score (FINDRISC) questionnaire is a screening tool to estimate risk of type 2 diabetes. In this study, we evaluated the ability of FINDRISC to predict the development of the metabolic syndrome (MetS) in an Iranian population without diabetes and MetS. **METHODS:** A total of 1,010 first-degree relatives of consecutive patients with type 2 diabetes, 30-70 years old (274 men and 736 women), without diabetes and MetS, were examined and followed up over  $8.0 \pm 1.6$  years (mean  $\pm$  SD) for MetS incidence. The incidence of MetS was examined across quartiles of FINDRISC, and a receiver operating characteristic (ROC) curve was plotted to assess the discrimination. At baseline and through follow-ups, participants underwent a standard 75 g 2-hour oral glucose toler-

ance test (OGTT). Data for determining FINDRISC were available from each participant. **RESULTS:** During 8,089 person-years of follow-up, 69 men and 209 women without MetS and diabetes at baseline subsequently developed MetS. The incidence of MetS was 31.4 per 1000 person-years in men and 35.5 in women. The FINDRISC at baseline was significantly associated with MetS evolution. Participants in the top quartile of FINDRISC were 4.4 times more likely to develop MetS than those in the bottom quartile (rate ratio 4.4; 95% CI 2.7-7.0). The area under the ROC curve was 65.0% (95% CI 61.3-68.7). **CONCLUSION:** The results of this study suggest that FINDRISC can be applied to detect MetS in a high-risk population.

**Keywords:** metabolic syndrome · type 2 diabetes · first-degree relatives · risk score · glucose tolerance

### Introduction

Metabolic syndrome (MetS) constitutes a cluster of metabolic risk factors for cardiovascular disease. It is associated with insulin resistance, which is an important determinant of cardiovascular risk [1]. It is estimated that about a quarter of the world's adult population has MetS [2, 3], and that they are twice as likely to die from and three times as likely to have a heart attack or stroke compared with people without MetS [4]. People with MetS have a fivefold higher risk of developing type 2 diabetes [5]. Thus, living with MetS means a significantly reduced quality of life. The causes for the development of the syndrome

remain unclear, but the pathophysiology seems to be largely attributable to insulin resistance, excessive flux of fatty acids, and a chronic proinflammatory state [6].

There is no specific treatment for MetS. Therapy includes lifestyle changes and pharmaceutical agents, but prevention would be preferable. Current evidence suggests that MetS can be prevented through pharmaceutical and lifestyle interventions in high-risk individuals [7], in whom beneficial changes in dietary and exercise behavior have been associated with reductions in several risk factors for cardiovascular disease and diabetes. Thus, given the strong evidence that pharmaceutical and lifestyle interventions prevent MetS, there is much

**Abbreviations:**

BMI - body mass index  
 BP - blood pressure  
 CI - confidence interval  
 FINDRISC - Finnish Diabetes Risk Score  
 FPG - fasting plasma glucose  
 HbA1c - glycosylated hemoglobin  
 HC - hip circumference  
 HDL - high density lipoprotein  
 IDPS - Isfahan Diabetes Prevention Study  
 IFG - impaired fasting glucose  
 IGT - impaired glucose tolerance  
 LDL - low density lipoprotein  
 LR - likelihood ratio  
 MetS - metabolic syndrome  
 NCEP-ATP III - National Cholesterol Education Program  
 Adult Treatment Panel III  
 NGT - normal glucose tolerance  
 NPV - negative predictive value  
 OGTT - oral glucose tolerance test  
 PG - plasma glucose  
 PPV - positive predictive value  
 ROC - receiver operating characteristic  
 RR - rate ratio  
 SD - standard deviation  
 WC - waist circumference  
 WHR - waist-to-hip ratio

interest in the identification of individuals at high risk of developing MetS, so that preventive action aimed at reducing their risk can be offered.

Early detection of MetS is difficult and population screening, using five clinical and biochemical parameters, would not be practicable or cost-effective, especially in low-income countries. We need a simple, non-invasive, effective tool that uses readily available clinical information for rapid identification of individuals at risk of MetS for use by the general public and in primary health care. These individuals could then be referred to further clinical testing to rule out or diagnose MetS.

The validated Finnish Diabetes Risk Score (FINDRISC) has traditionally been used as a predictor of type 2 diabetes. It takes into account the usual clinical characteristics, such as age, body mass index (BMI), waist circumference (WC), physical activity, dietary consumption of fruits, vegetables, and berries, use of antihypertensive medication, history of high blood glucose, and family history of diabetes.

FINDRISC has been successfully implemented as a practical screening instrument to assess diabetes risk and to detect undiagnosed type 2 diabetes in European populations [8-24]. However, it has also become evident that it is not universally applicable among all ethnic groups and populations [16, 18, 25, 26]. Three cross-sectional studies are available on the relationship between FIN-

DRISC and prevalence of MetS [16, 21, 27], but no cohort study has examined MetS incidence using FINDRISC. Moreover, little is known about the relationship of FINDRISC and MetS in first-degree relatives of patients with type 2 diabetes, who are at higher risk of diabetes and MetS than the general population.

The objective of this longitudinal study was to evaluate the ability of FINDRISC to predict the incidence of MetS in an Iranian population without diabetes and MetS.

## Subjects and methods

### *Data collection*

The recruitment methods and examination procedures of the Isfahan Diabetes Prevention Study (IDPS) have been described previously [28]. Briefly, IDPS is an ongoing cohort in central Iran to assess the various potential risk factors for diabetes in subjects with a family history of type 2 diabetes (one of the main risk factors for diabetes). Our study sample comprised 3,409 (895 male and 2,514 female) first-degree relatives of patients with type 2 diabetes. All patients attended clinics at the Isfahan Endocrine and Metabolism Research Center, which is affiliated to the Isfahan University of Medical Sciences, Iran.

The study was conducted between 2003 and 2005. All participants were from Isfahan city and adjoining areas. They completed laboratory tests including a standard 75 g 2-h oral glucose tolerance test (OGTT), a questionnaire on their health status and on various potential risk factors for diabetes, and the FINDRISC questionnaire. The participants were examined in follow-up tests according to the standard of medical care in diabetes [29] to update information on demographic, anthropometric, and lifestyle factors and to diagnose diabetes onset. Accordingly, if OGTT was normal at baseline, repeat testing was carried out at 3-year intervals at least. Otherwise, repeat testing was usually carried out annually. Tenets of the current version of the Declaration of Helsinki were followed, institutional ethical committee approval was granted, and an informed consent form was signed by each participant.

### *Follow-up and identification of MetS*

MetS cases were identified according to the consensus criteria released in 2009 [30], which is equivalent to the third report of the National Cholesterol Education Program Adult Treatment Pan-

el III (NCEP-ATP III) [31]. MetS was considered to be present when at least three of the following five characteristics were determined:

1. Central obesity, defined using ethnic-specific cut-points of waist (waist circumference  $\geq 102$  cm in men and  $\geq 88$  cm in women)
2. Triglycerides  $\geq 150$  mg/dl
3. High-density lipoprotein (HDL) cholesterol  $<40$  mg/dl in men and  $<50$  mg/dl in women
4. Blood pressure (BP)  $\geq 130/85$  mmHg or on antihypertensive medication
5. Raised plasma glucose, defined as fasting plasma glucose (FPG)  $\geq 100$  mg/dl.

Pregnant women and people with type 1 and type 2 diabetes were excluded. Participants with type 2 diabetes were excluded because the diagnosis of MetS may have additional meaning in individuals with type 2 diabetes who are treated to counteract the high cardiovascular risk. Other than these, individuals who already had MetS or subjects with a history of taking antidiabetic or lipid-lowering compounds were also excluded. Among the 3,409 persons who participated at baseline, 1,340 subjects were excluded because of diagnosis of type 1 or type 2 diabetes, MetS, or a history of taking antidiabetic or lipid-lowering agents at baseline. Another 1,059 had no follow-up, leaving 1,010 participants included in the study.

The 1,010 participants had a mean age of 42.2 (6.5) years (range 30-70). All participants underwent at least one subsequent review during the follow-up period of 8.0 (1.6) years (range 3-10). Attendees at follow-up visits did not differ significantly from non-attendees regarding most baseline characteristics such as: height, weight, BMI, WC, hip circumference (HC), waist-to-hip ratio (WHR), HbA1c, triglycerides, fasting plasma glucose, systolic and diastolic blood pressure, and obesity.

However, non-attendees had slightly lower:

- Plasma glucose (PG) at 30 min (134.6 mg/dl versus 140.7 mg/dl,  $p < 0.001$ )
- PG at 60 min (133.7 mg/dl versus 142.7 mg/dl,  $p < 0.001$ )

**Table 1.** Prevalence of components of the modified version of the Finnish Diabetes Risk Score (FINDRISC) in men and women

FINDRISC component	Score	Men	Women	Total
<b>Age (yr)</b>				
<45	0	166(61.0)	489(66.9)	655(65.3)
45-54	2	89(32.7)	277(31.1)	316(31.5)
55-64	3	17 (6.3)	14 (1.9)	31 (3.1)
>64	4	0 (0.0)	1 (0.1)	1 (0.1)
<b>BMI (kg/m<sup>2</sup>)</b>				
$\leq 25$	0	69(25.7)	137(18.8)	206(20.6)
25-30	1	160(59.5)	401(55.0)	561(56.2)
>30	3	40(14.9)	191(26.2)	231(23.1)
<b>Waist circumference (cm)</b>				
Men <94; women <80	0	136(51.7)	205(28.4)	341(34.6)
Men 94-101; women 80-87	3	109(41.4)	327(45.2)	436(44.2)
Men >101; women >87	4	18 (6.8)	191(26.4)	209(21.2)
<b>History of drug treatment*</b>				
Yes	0	224(86.2)	680(95.1)	904(92.7)
No	2	36(13.8)	35 (4.9)	71 (7.3)
<b>Previous high BG**</b>				
Yes	0	163(59.5)	448(60.9)	611(60.5)
No	5	111(40.5)	288(39.1)	399(39.5)
<b>Family history of diabetes</b>				
Yes, first-degree	1	274(100)	736(100)	1010(100)

**Legend:** Data are number (%). The total of each variable may vary because of missing values. \* Antihypertensive drug treatment. \*\* Previously measured high blood glucose. *Abbreviations:* BG – blood glucose, BMI – body mass index, FINDRISC – Finnish Diabetes Risk Score.

- PG at 120 min (107.8 mg/dl versus 115.9 mg/dl,  $p < 0.001$ )
- FINDRISC (10.6 vs. 11.2,  $p < 0.001$ )

On the other hand, non-attendees had higher levels of:

- Cholesterol (196.2 mg/dl versus 190.9 mg/dl,  $p < 0.001$ )
- Low-density lipoprotein (LDL) cholesterol (120.3 mg/dl versus 116.7 mg/dl,  $p < 0.05$ )
- HDL cholesterol (49.0 mg/dl versus 47.2 mg/dl,  $p < 0.01$ )

Also, non-attendees were slightly older (42.8 year versus 42.2 year,  $p < 0.05$ ).

#### *Follow-up and test procedures*

Information on age, gender, body size, HbA1c, cholesterol, LDL cholesterol, HDL cholesterol, triglycerides, BP, family, and personal medical his-

**Table 2.** Characteristics of first-degree relatives of patients with type 2 diabetes according to FINDRISC quartiles in the Isfahan Diabetes Prevention Study

Characteristic	Total	1 <sup>st</sup> quartile	2 <sup>nd</sup> quartile	3 <sup>rd</sup> quartile	4 <sup>th</sup> quartile
Number (%)	1010 (100)	238 (23.5)	328 (32.5)	244 (24.1)	200 (19.8)
FINDRISC	11.2 (3.8)	6.3 (1.2)	9.9 (0.9)	13.1 (0.9) <sup>1,2</sup>	16.8 (1.2) <sup>*</sup>
Age (yr)	42.2 (6.5)	41.0 (6.3)	40.3 (5.2)	43.2 (6.3) <sup>1</sup>	45.4 (7.3) <sup>*</sup>
Height (cm)	159.8 (8.3)	161.0 (8.6)	159.8 (8.2)	160.0 (8.2)	157.9 (7.8) <sup>*</sup>
Weight (kg)	70.7 (10.7)	64.0 (9.1)	69.8 (9.7)	74.8 (11.0)	75.3 (9.2)
WC (cm)	86.2 (9.2)	79.7 (6.8)	84.9 (7.2)	89.8 (8.4) <sup>1,2</sup>	91.2 (6.8) <sup>*</sup>
HC (cm)	105.5 (7.8)	100.1 (5.2)	104.6 (5.8)	108.5 (9.4) <sup>1</sup>	109.4 (7.1) <sup>*</sup>
Waist-to-hip ratio	0.82 (0.07)	0.80 (0.07)	0.81 (0.06)	0.83 (0.07) <sup>1,2</sup>	0.84 (0.06) <sup>*</sup>
BMI (kg/m <sup>2</sup> )	27.7 (3.6)	24.7 (2.6)	27.3 (2.4)	29.2 (4.0) <sup>1,2</sup>	30.2 (3.1) <sup>*</sup>
Follow-up duration (yr)	8.1 (1.6)	8.4 (1.5)	8.1 (1.5)	8.0 (1.6) <sup>1</sup>	7.4 (1.8) <sup>*</sup>
FPG (mmol/l)	92.9 (11.3)	86.7 (7.5)	90.8 (10.4)	95.4 (11.8) <sup>1,2</sup>	100.7 (10.7) <sup>*</sup>
PG 30 min (mmol/l)	140.7 (31.0)	128.0 (26.4)	136.4 (28.1) <sup>1</sup>	147.1 (31.4) <sup>1</sup>	154.9 (32.8) <sup>*</sup>
PG 60 min (mmol/l)	142.7 (42.2)	119.4 (32.4)	137.5 (39.2) <sup>1</sup>	151.2 (40.9) <sup>1,2</sup>	168.2 (42.3) <sup>*</sup>
PG 120 min (mmol/l)	115.9 (33.1)	97.8 (21.8)	110.8 (30.9)	119.7 (32.7) <sup>1,2</sup>	141.1 (31.8) <sup>*</sup>
HbA1c (%)	5.0 (0.8)	4.9 (0.7)	5.0 (0.7) <sup>1</sup>	5.1 (0.9)	5.2 (0.9) <sup>*</sup>
TC (mmol/l)	190.9 (38.9)	183.5 (37.9)	186.7 (38.4)	194.7 (39.4) <sup>1</sup>	202.1 (37.2) <sup>*</sup>
LDL (mmol/l)	116.7 (35.9)	111.0 (37.1)	113.1 (32.9)	119.1 (32.2)	126.5 (31.2) <sup>*</sup>
HDL (mmol/l)	47.2 (11.8)	45.8 (12.0)	45.9 (10.7)	48.0 (12.6)	50.0 (11.9) <sup>*</sup>
TG (mmol/l)	140.6 (81.7)	138.0 (76.6)	143.8 (81.2)	144.5 (97.4)	133.7 (65.1)
Systolic BP (mm Hg)	110.4 (13.4)	107.1 (13.7)	109.2 (12.9)	112.4 (12.7) <sup>1</sup>	113.7 (13.6) <sup>*</sup>
Diastolic BP (mm Hg)	71.7 (10.6)	69.3 (11.1)	70.9 (9.9)	73.5 (9.9) <sup>1</sup>	73.6 (11.2) <sup>*</sup>
Women (no., %)	736 (72.9)	149 (62.9)	248 (75.4)	179 (73.4)	160 (80.0) <sup>*</sup>
Overweight (no., %)	794 (79.5)	106 (45.9)	275 (84.6)	214 (88.1)	199 (99.5) <sup>*</sup>

**Legend:** Data are mean (SD) or number (%). FINDRISC values at baseline. 1<sup>st</sup> quartile:  $\leq 8.0$ ; 2<sup>nd</sup> quartile: 8.1-11.0; 3<sup>rd</sup> quartile: 11.1-14.0; 4<sup>th</sup> quartile: 14.1-23.0. \*  $p < 0.001$ , comparison across all four groups. Difference in the mean value of variables compared to the 1<sup>st</sup> quartile and 2<sup>nd</sup> quartile. **Abbreviations:** BMI – body mass index, BP – blood pressure, CI – confidence interval, FG – fasting glucose, PG – plasma glucose, FINDRISC – Finnish Diabetes Risk Score, HDL – high-density lipoprotein, LDL – low-density lipoprotein, IFG – impaired fasting glucose, IGT – impaired glucose tolerance, NGT – normal glucose tolerance, TC – total cholesterol.

tory was collected at baseline and through follow-ups. The same methodology was applied for baseline and follow-up studies.

The participants included siblings and children of patients with type 2 diabetes. They reported to the clinics in the morning after an overnight fast. They were asked to abstain from vigorous exercise in the evening and in the morning of their visit. Smokers were encouraged to abstain from smoking in the morning of the investigations.

Firstly, on arrival at the clinic, the information provided by the participants in the questionnaire on family history was verified. Then, with the subjects in light clothing and without shoes, height, weight, WC, and HC were measured using standard apparatus. Weight was measured to the nearest 0.1 kg on a calibrated beam scale. Height, WC,

and HC were measured to the nearest 0.5 cm with a measuring tape. The waist was measured midway between the lower rib margin and the iliac crest at the end of gentle expiration. Hip circumference was measured over the greater trochanters directly over the underwear. Resting BP was measured after the participants had been seated for 10 min with a mercury sphygmomanometer and appropriately sized cuffs, using standard techniques.

FPG was measured with the glucose oxidase method. Participants with FPG  $\geq 200$  mg/dl or pharmacological treatment were considered to be persons with diabetes. If FPG was  $\geq 126$  mg/dl and  $< 200$  mg/dl, a second FPG was measured on another day. If the second FPG was also  $\geq 126$  mg/dl, participants were considered to be diabetic. Those with FPG  $< 126$  mg/dl underwent a standard OGTT (75 g glucose 2-h) at baseline and follow-up visits. Venous blood was

sampled 0, 30, 60, and 120 min after oral glucose administration. Plasma samples were centrifuged and analyzed the same day.

HbA1c (measured by ion-exchange chromatography), total cholesterol, triglycerides, HDL, and LDL (calculated with the Friedewald equation [32] provided total triglycerides did not exceed 400 mg/dl) were recorded. All blood sampling procedures were performed in the central laboratory of the Isfahan Endocrine and Metabolism Research Center using the enzyme-linked method.

### Definitions

Based on the OGTT results, participants were assigned to the following categories:

**Table 3.** Incidence rates and rate ratios (RR) of the metabolic syndrome according to FINDRISC quartiles at baseline in the Isfahan Diabetes Prevention Study

Characteristic	1 <sup>st</sup> quartile	2 <sup>nd</sup> quartile	3 <sup>rd</sup> quartile	4 <sup>th</sup> quartile
Number of cases (%)	33 (11.9)	76 (27.3)	82(29.5)	87(31.3)
Person-years	1997	2664	1954	1483
Incidence/1000 p-y (95% CI)	16.5 (11.4-23.1)	28.5 (22.5-35.6)	42.0(33.5-51.8)	58.7(47.3-71.8)
Unadjusted RR (95% CI)	1.00	1.9 (1.2- 2.9)	3.1 (2.0- 5.0)	4.8 (3.0- 7.6)
Gender-adjusted RR (95% CI)	1.00	1.9 (1.2- 2.9)	3.1 (2.0- 4.9)	4.7 (3.0- 7.5)
Age- and gender-adjusted RR (95% CI)	1.00	1.9 (1.2- 2.9)	3.0 (1.9- 4.7)	4.4 (2.7- 7.0)

**Legend:** Data are FINDRISC values at baseline. 1<sup>st</sup> quartile: ≤8.0; 2<sup>nd</sup> quartile: 8.1-11.0; 3<sup>rd</sup> quartile: 11.1-14.0; 4<sup>th</sup> quartile: 14.1-23.0. Rate ratio (95% CI) calculated by binary logistic regression. *Abbreviations:* CI – confidence interval, FINDRISC – Finnish Diabetes Risk Score, p-y – person-years, RR – rate ratio.

1. Normal glucose tolerance (NGT, FPG below 100 mg/dl and the 2-h plasma glucose (2hPG) <140 mg/dl)
2. Impaired fasting glucose (IFG, FPG in the range of 100 to 126 mg/dl and the 2hPG was <140 mg/dl)
3. Impaired glucose tolerance (IGT, FPG <126 mg/dl, but with 2hPG concentration ≥140 and <200 mg/dl)
4. Diabetes (FPG ≥200 mg/dl or pharmacological treatment, FPG ≥126 and/or 2hPG of ≥200 mg/dl) [33].

#### Finnish Diabetes Risk Score (FINDRISC)

FINDRISC was computed for each participant using clinical and questionnaire data collected at baseline. The FINDRISC usually comprises the following eight items [6, 9]:

- Age
- Body mass index (BMI, weight (kg) / height squared (m<sup>2</sup>))
- Waist circumference (WC)
- Physical activity
- Dietary consumption of fruits, vegetables, and berries
- Use of antihypertensive medication
- History of high blood glucose
- Family history of diabetes.

The maximum achievable score is 26. In the current study, a modified version of the FINDRISC was used. In this shortened version, the following variables were omitted because these items did not add much power for the prediction of diabetes risk in previous studies, as suggested in the original publication [9] and in subsequent studies [18, 25, 34]:

- Dietary consumption of fruits, vegetables, and berries
- Physical activity

Thus, the maximum achievable score on the modified FINDRISC was 23. High blood glucose was defined as IFG and/or IGT at

baseline. The modified version of the FINDRISC and the prevalence of FINDRISC questionnaire components in men and women are presented in **Table 1**.

#### Determination of MetS incidence

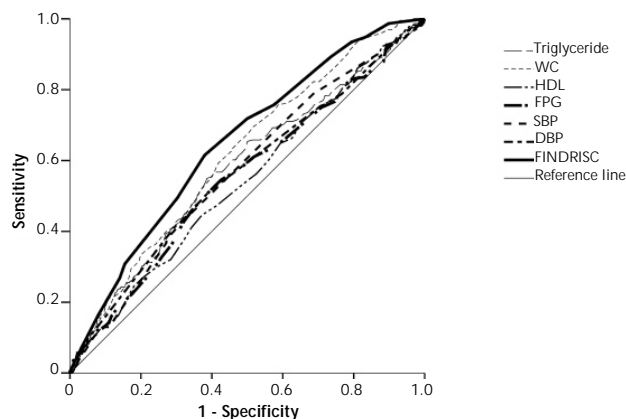
The incidence of MetS was expressed as the number of MetS cases per 1000 person-years of follow-up, beginning with the date of completion of the baseline examination in 2003 to 2005 and continuing until the occurrence of MetS, the date of the last completed follow-up, death, or end of follow-up on September 31, 2011, whichever came first.

#### Statistical analysis

Statistical methods included Student's *t*-test, chi-squared test, and binary logistic regression. Univariate and multivariate binary logistic regression equations were fitted to identify predictors of new-onset MetS using SPSS version 18 for Windows (SPSS Inc., Chicago, IL, USA).

We calculated the FINDRISC for each participant using baseline age, BMI, WC, use of antihypertensive medication, history of high blood glucose (IFG and/or IGT), and family history of diabetes. We re-coded the FINDRISC into quartile and compared the risk of developing MetS in each quartile with the lowest category of risk (reference group). The ability of FINDRISC, and each component of MetS, to predict the incidence of MetS was examined with receiver operating characteristic (ROC) curves and their respective areas under the curve. Sensitivity was plotted as a function of 1-specificity.

To evaluate the effect of adding components of MetS together, we put each component into a model going from most to least significant as follows:



	Area under the curve (95% CI)
Finnish Diabetes Risk Score	0.650 (0.613, 0.687)
Waist circumference	0.620 (0.580, 0.659)
Triglyceride	0.588 (0.546, 0.630)
Systolic blood pressure	0.572 (0.531, 0.613)
Diastolic blood pressure	0.565 (0.522, 0.608)
Fasting plasma glucose	0.550 (0.507, 0.593)
HDL	0.542 (0.499, 0.585)

**Figure 1.** Receiver operating characteristic (ROC) curves for FINDRISC, triglyceride, waist circumference (WC), high-density lipoprotein cholesterol (HDL), fasting plasma glucose (FPG), systolic (SBP) and diastolic blood pressure (DBP) to predict metabolic syndrome in first-degree relatives of patients with type 2 diabetes without diabetes or metabolic syndrome at baseline. The estimates of the area under the ROC curves and their 95% CI are shown.

**Model 1:** WC and triglyceride

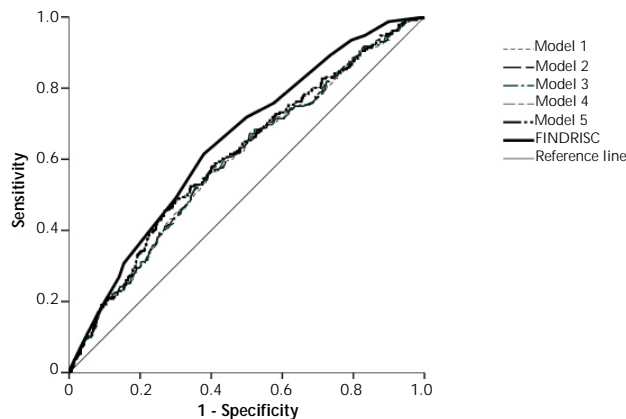
**Model 2:** WC, triglyceride, and systolic blood pressure

**Model 3:** WC, triglyceride, systolic and diastolic blood pressure

**Model 4:** WC, triglyceride, systolic and diastolic blood pressure, and FPG

**Model 5:** WC, triglyceride, systolic and diastolic blood pressure, FPG, and HDL

The area under the ROC curve is a global summary statistic of the discriminative value of a model, describing the probability that the score is higher in an individual developing MetS than in an individual not developing MetS. Areas under the ROC curves were compared by the algorithm developed by DeLong *et al.* [35]. All tests for statistical significance were two-tailed, and all tests were performed assuming a type I error probability of <0.05.



	Area under the curve (95% CI)
Finnish Diabetes Risk Score	0.650 (0.609, 0.688)
Model 1	0.596 (0.554, 0.639)
Model 2	0.598 (0.555, 0.640)
Model 3	0.599 (0.556, 0.641)
Model 4	0.609 (0.567, 0.651)
Model 5	0.609 (0.567, 0.651)

**Figure 2.** Receiver operating characteristic (ROC) curves for FINDRISC, model 1 (WC, triglyceride), model 2 (WC, triglyceride, SBP), model 3 (WC, triglyceride, SBP, and DBP), model 4 (WC, triglyceride, SBP, DBP, and FPG) and model 5 (WC, triglyceride, SBP, DBP, FPG, and HDL) to predict metabolic syndrome in first-degree relatives of patients with type 2 diabetes without diabetes or metabolic syndrome at baseline. The estimates of the area under the ROC curves and their 95% confidence intervals are shown.

## Results

Baseline characteristics of the 732 (72.5%) participants without and 278 (27.5%) with MetS were as expected. Those participants who developed MetS were older and had higher mean weight, BMI, WC, HC, WHR, FPG, and PG at 30, 60, and 120 min, higher HbA1c, triglyceride, cholesterol, systolic and diastolic BP, and FINDRISC at baseline (table not shown, can be requested from the authors). A higher proportion of them also had overweight, used antihypertensive medication, and had higher IFG and IGT. The mean (SD) age was 43.3 (6.8) years for those with MetS and 41.8 (6.3) years for those without MetS. The mean (SD) FINDRISC was 12.7 (3.7) for those with MetS and 10.7 (3.7) for those without MetS. The total score ranged from 5 to 21.

The baseline characteristics of the study participants according to FINDRISC quartiles are shown in **Table 2**. In comparison with variables at

baseline, all variables except triglyceride were more likely to increase, height and follow-up duration was more likely to decrease across all four subject groups.

278 incident cases of MetS (27.5%, 69 men and 209 women) occurred during 8,089 (2,197 men and 5,892 women) person-years of follow-up. The overall incidence of subsequent MetS development was 34.3 (95% CI: 30.4-38.3) per 1000 person-years. The incidence rates were slightly higher in women (35.5, 95% CI: 30.7-40.2 per 1000 person-years) than in men (31.4, 95% CI: 24.5-39.6), but the difference was not statistically significant.

The FINDRISC was associated with MetS incidence. The incidence of MetS was 58.7 per 1000 person-years (95% CI 47.3-71.8) for participants in the highest quartile of FINDRISC, and 16.5 per 1000 person-years (95% CI 11.4-23.1) for the lowest quartile. The risk of MetS increased with increasing quartiles of FINDRISC. Compared with participants in the lowest quartile, the risk of MetS was:

1. 4.8 times higher for those in the highest quartile at baseline (rate ratio (RR) 4.8, 95% CI: 3.01-7.59)
2. 3.1 times higher for those in the 3rd quartile (RR 3.1, 95% CI: 2.00-4.95)
3. 1.9 times higher in those in the 2nd quartile (RR 1.9, 95% CI: 1.20-2.93) in unadjusted models

Controlling for age and gender did not appreciably alter the RR compared with the unadjusted model (Table 3).

The ROC curves for the incidence of MetS by FINDRISC, and each component of MetS, are shown in Figure 1. The areas under the ROC curves from the greatest to the least area were:

- 0.650 (95% CI: 0.611-0.689) for FINDRISC
- 0.620 (95% CI: 0.580-0.659) for WC

**Table 4.** The predictive performance of different Finnish Diabetes Risk Score (FINDRISC) cut-off values for predicting the metabolic syndrome

Score	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)	Correctly classified (%)	Youden index	LR+	LR-
≥5	100.0	0.0	26.4	-	27.5	0.00	1.00	0.25
≥6	97.5	10.3	28.1	92.1	34.3	0.08	1.09	0.38
≥7	93.5	16.9	28.7	87.7	38.0	0.10	1.13	0.36
≥8	92.5	20.8	29.5	88.2	40.5	0.13	1.17	0.42
≥9	88.1	28.0	30.5	86.8	44.6	0.16	1.22	0.57
≥10	75.5	43.2	32.3	83.1	52.1	0.19	1.33	0.56
≥11	71.6	50.7	34.2	83.2	56.4	0.22	1.45	0.63
<b>≥12</b>	<b>60.8</b>	<b>62.4</b>	<b>37.0</b>	<b>70.8</b>	<b>62.0</b>	<b>0.23</b>	<b>1.62</b>	<b>0.71</b>
≥13	50.0	70.2	37.4	79.7	64.7	0.20	1.68	0.76
≥14	43.2	74.9	37.2	78.5	66.1	0.18	1.72	0.81
≥15	31.2	84.6	42.1	77.4	69.9	0.16	2.03	0.84
≥16	27.7	86.1	41.7	76.9	70.0	0.14	1.99	0.89
≥17	17.6	92.1	44.3	75.7	71.6	0.10	2.22	0.94
≥18	9.0	96.7	48.9	74.7	72.6	0.06	2.74	0.96
≥19	5.4	98.4	52.0	74.3	72.8	0.04	3.29	0.99
≥20	1.1	99.9	50.0	75.0	73.8	0.01	7.90	0.99
≥21	0.4	100.0	0.0	100.0	73.7	0.00	-	1.00

**Legend:** Values in bold represent best cut-points. Youden index was defined as the maximum of (sensitivity + specificity - 1). *Abbreviations:* LR+: positive likelihood ratio; LR-: negative likelihood ratio.

- 0.588 (95% CI: 0.546-0.630) for triglyceride
- 0.572 (95% CI: 0.531-0.613) for systolic and
- 0.565 (95% CI: 0.522-0.608) for diastolic BP
- 0.550 (95% CI: 0.507-0.593) for FPG
- 0.542 (0.499-0.585) for HDL

All parameters, except HDL, were significant predictors for future risk of MetS ( $p < 0.001$ ). Although WC, FPG, triglyceride, HDL, and BP had areas smaller than that of FINDRISC, there were no statistical significances between the areas under the ROC curves for FINDRISC, WC, and triglyceride. The area for FPG, HDL, and systolic and diastolic BP was significantly smaller than that for FINDRISC ( $p < 0.05$ ).

Test characteristics for various FINDRISC cut-off values are presented in Table 4. At a score of 12 or higher, the sum of sensitivity and specificity was maximal. Accordingly, the optimal cut-point for detecting MetS was a FINDRISC greater than or equal to 12. At a FINDRISC greater than or equal to 12, sensitivity was 60.8% and specificity was 62.4%. The corresponding positive and negative predictive values were 37.0% and 70.8%, respectively.

The ROC curves for the incidence of MetS by FINDRISC and the models are shown in Figure 2. Adding more components of MetS to the model did not appreciably alter the area under the curve. However there was no significant difference between each model and FINDRISC.

## Discussion

In this study, the FINDRISC showed a reasonably good ability to predict MetS in a cohort of first-degree relatives of patients with type 2 diabetes, with an area under the ROC of 65%. This was similar to its ability to predict MetS in other populations tested to date [16, 21, 27]. The FINDRISC was originally developed in a prospective setting to identify persons at high risk of developing type 2 diabetes. In this study, we analyzed the score's performance in a prospective setting as screening tool for the detection of MetS, which is independently associated with future risk of type 2 diabetes [4, 36, 37]. It may therefore be regarded as a useful screening instrument for the identification of high-risk individuals who could benefit from early life-style and pharmacological interventions [7, 38].

Although the cause and role of MetS are still controversial, many authors [31, 39] believe that insulin resistance is the 'core aspect' of the syndrome. Under this assumption, it is reasonable to postulate that the factors that contribute most to the development of the MetS should relate most closely to insulin resistance.

The efficiency of risk scores may vary between populations with different ethnic backgrounds. Therefore, risk scores should be validated in each population before use [40, 41]. Although the FINDRISC tool was developed and validated in European populations [8-10, 14-21], it is also valid for Middle-Eastern populations, despite different life-styles.

Participants in the top quartile of FINDRISC were 4.8 times more likely to progress to MetS than those in the bottom quartile. Those with higher WC at baseline had lower risk of progression to MetS than those with high FINDRISC, further emphasizing the usefulness of FINDRISC in predicting MetS. Thus, the study confirms the reliability of FINDRISC for MetS prediction. FINDRISC may serve as an initial assessment tool to identify persons with a risk of developing type 2 diabetes. The FINDRISC, which uses information routinely available in primary care records, is a simple, inexpensive, and non-invasive tool.

Two cross-sectional studies in Europe and one in Taiwan have assessed the risk of developing MetS by using FINDRISC. Saaristo *et al.* examined the prevalence of MetS in a Finnish population; the area under the ROC curve for the prevalence of MetS (NCEP-ATP III definition) was 72.4% in men and 75.3% in women [21]. A Greek cross-sectional study found an area under the ROC curve of 70.7% in men and 75.7% in women using

a modified version of FINDRISC [16]. A Taiwanese cross-sectional study evaluated a FINDRISC questionnaire, and compared it with 9 other available risk scores for the detection of unknown MetS, demonstrating an area under the ROC of 77.0% using FINDRISC [27]. The area under the ROC curve for the prevalence of MetS in all these studies was higher than that in the present study. However, the other studies could only provide information on the present condition of the tested individuals as they all had a cross-sectional design.

Our study has several strengths and limitations. The strengths include the use of a sample consisting of both men and women, performance of standard OGTT, and information on potential determinants of MetS. There was unlikely to be selection or information bias because of the prospective design. At follow-up, non-attendees did not differ from attendees in the entire population according to major risk factors for the development of MetS, although a difference too small to explain the high progression rate to MetS in our study was seen in the mean levels of lipid profiles, PG, and FINDRISC.

Our database is one of the few that followed first-degree relatives of patients with type 2 diabetes, thereby enabling us simultaneously to control the genetic factors that may predict insulin resistance status. Our study was limited to a cohort of individuals who are at increased risk of developing type 2 diabetes, because they were first-degree relatives of patients with type 2 diabetes. Therefore, a possible selection bias is more likely to underestimate associations than to exaggerate them.

The length of follow-up in the IDPS cohort was relatively short. Thus, while the study proved effective in identifying first-degree relatives of patients with type 2 diabetes who rapidly progress to MetS, those with slower disease onset may not have been identified. Alternatively, longer follow-up periods may increase the RR for the association between FINDRISC and the incidence of MetS if more people in the highest risk score category go on to develop MetS. Therefore, it is necessary to validate the association of FINDRISC and MetS in other populations.

This study evaluated the modified FINDRISC instead of the full risk model, which also includes daily consumption of vegetables, fruits, and berries, and physical activity [10]. As our information on these two risk factors was insufficient we excluded them. Both factors are considered to be important and were primarily targeted during recent diabetes prevention trials. However, in the Lind-



strom and Tuomilehto model, both daily intake of vegetables and fruits or berries and physical inactivity were not statistically significant. They were included in the model mainly because prevention studies have demonstrated their importance [10, 42, 43]. It is difficult to apply simple variables for such complex behavioral patterns such as physical activity and diet. In terms of the definition of MetS incidence used, a selection bias may be present as participants who attend for screening may have been more likely to be tested. Consequently, they have been diagnosed with MetS. On the other hand, participants with MetS who had a low risk score may not have been identified because they had not been tested.

## Conclusion

In conclusion, our data provide further evidence that FINDRISC can be a suitable tool to predict

MetS in a high-risk population and to identify undetected MetS in clinical practice.

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