Features of the Metabolic Syndrome Predict Higher Risk of Diabetes and Impaired Glucose Tolerance

A prospective study in Mauritius

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OBJECTIVE — To assess the independent and joint effects of the components of the metabolic syndrome, including leptin, which is a recently proposed addition to this syndrome, in predicting the cumulative incidence of impaired glucose tolerance (IGT) and diabetes among individuals with normal glucose tolerance.

RESEARCH DESIGN AND METHODS — This prospective study involved 2,605 residents of Mauritius with normal glucose tolerance who were followed for 5 years for IGT or diabetes onset in relation to total and regional adiposity (BMI, waist-to-hip ratio [WHR]), fasting and 2-h 75-g oral glucose load glucose and insulin, total and HDL cholesterol, blood pressure, serum uric acid, triglyceride, and leptin levels.

RESULTS — A multivariate logistic regression model adjusted for age, sex, ethnicity, and diabetes family history showed a significantly higher linear increase in risk of IGT and diabetes in association with the following variables only: fasting glucose (odds ratio 1.89 [95% CI 1.51–2.34]), 2-h glucose (1.68 [1.50–1.88]), WHR (1.30 [1.10–1.52]), BMI (1.04 [1.00–1.08]), and serum uric acid (1.37 [1.20–1.57]). However, a nonlinear increase was seen with serum triglyceride and plasma leptin concentrations. No risk factors resulted in joint effects that were greater than expected from combining individual effects.

CONCLUSIONS — Metabolic syndrome features independently predict a higher risk of diabetes or IGT in normoglycemic subjects but in combination confer no higher-than-expected risk of these outcomes. At higher concentrations of triglycerides and leptin, risk plateaus and even declines slightly.

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T he more frequent occurrence than expected by chance of multiple characteristics associated with poor cardiovascular and other health outcomes such as hyperglycemia, hypertension, and lipid abnormalities has given rise to the concept of the "metabolic syndrome" (1). A recent review of this syndrome identified 6 components that were included in most of the definitions (fasting or stimulated circulating insulin and glucose levels, triglyceride levels, HDL cholesterol levels, blood pressure lev-els, and overall or central obesity) (1). Also included in some definitions were serum uric acid (2–4) and total or LDL cholesterol (5) levels. Recent data from our group (6), the results of a factor analysis by others (7), and the demonstration of associations between leptin and both insulin sensitivity (8) and hyperinsulinemia (9) argue for leptin's involvement in the metabolic syndrome.

The significance of this syndrome regarding the development of cardiovascular disease is fairly well understood because many of its components are well-accepted cardiovascular disease risk factors, but this is less true regarding the prediction of deterioration in glucose tolerance. Although many studies have examined features of the metabolic syndrome in relation to risk of diabetes and impaired glucose tolerance (IGT), few have examined the effects of all 6 components. A 3.5-year follow-up of an elderly Finnish cohort that considered these 6 components revealed higher diabetes risk in association with presence of IGT or hypertension; higher fasting and 2-h plasma insulin levels, total triglyceride level, waist-to-hip ratio (WHR), and BMI; and lower HDL cholesterol levels (10). The independent effects of these correlated factors were not directly assessed in multivariate models. The importance of assessing whether metabolic syndrome components have independent effects on the risk of this outcome was demonstrated by an analysis conducted using San Antonio Heart Study data (11). A multivariate model developed to predict diabetes identified fasting and 2-h glucose levels, BMI, HDL cholesterol levels, and pulse pressure as having independent effects on the risk of diabetes during 8 years of follow-up (11). In this model, fasting insulin levels, triglyceride levels, subscapular-to-triceps skinfold ratio (a measure of central adiposity), and systolic and diastolic blood pressure did not have independent effects on diabetes risk, even though they were significantly related to higher risk in an age-, sex-, and ethnicity-adjusted analysis. Whether the effects of these independent variables in

combination had effects on diabetes risk that were greater than expected from combining individual effects was not assessed.

Few studies have examined in a comprehensive fashion the effect of most metabolic syndrome components and leptin concentration on the risk of diabetes and IGT in a manner that permitted the assessment of the independent effects and testing for interaction between individual components. We therefore investigated this issue among residents of Mauritius with normal glucose tolerance at baseline who were followed for 5 years for the development of IGT or diabetes.

RESEARCH DESIGN AND METHODS

Background and subjects
Mauritius is a subtropical island located in the southwestern Indian Ocean about 800 km east of Madagascar. The multiethnic population consists of 70% individuals of Indian origin (54% Hindu and 16% Muslim), 2% individuals of Chinese origin, and the remaining 28% are Creoles predominantly of African and Malagasy ancestry with some European admixture.

All adults 25–74 years of age living within 11 geographically defined areas were eligible for the 1987 baseline survey as previously described (12,13). Overall response ($n = 5,083$) consisted of 83% of eligible men and 89% of eligible women. All subjects alive and willing to participate were eligible for the 1992 follow-up survey. Longitudinal 1987–1992 data were available for 3,793 subjects (74.6%). This analysis focuses on the development of diabetes and IGT among individuals who were initially normoglycemic (as defined below) in 1987 ($n = 3,467$), of whom 2,605 (75.1%) had longitudinal information available from both surveys. The survey protocol was reviewed and approved by the Alfred Healthcare Group Ethics Committee (Melbourne, Australia).

Survey procedures
The following procedures were carried out on all subjects in 1987 as previously described (12–15). Height was measured without shoes to the nearest centimeter. Weight was measured to the nearest 0.1 kg in light clothes and without shoes. BMI was calculated as weight in kilograms by height in meters squared. Waist and hip circumferences were measured in duplicate to the nearest 0.5 cm with a measuring tape while

the subject was standing relaxed and wearing 1 layer of light clothing. A third measurement was taken if the first 2 were not within 2 cm of each another. The mean of the closest 2 measurements was used to calculate the WHR. The waist measurement was taken at the midpoint between the iliac crest and the lower rib margin, and the hip circumference was taken around the maximum circumference of the buttocks posteriorly and the symphysis pubis anteriorly. Participants were seated and rested for at least 5 min before systolic and fifth-phase diastolic blood pressure measurements were taken in duplicate using a standard mercury sphygmomanometer. Ethnicity was determined by self-report.

All subjects not taking diabetes medication had a 2-h 75-g (glucose monohydrate) oral glucose tolerance test after an overnight fast in 1987 and 1992. Diabetes was diagnosed if subjects reported a history of diabetes and were taking oral hypoglycemic medication or insulin or if the fasting plasma glucose level was ≥ 7.0 mmol/l or the 2-h value was ≥ 11.1 mmol/l (16). Subjects with a fasting plasma glucose level < 7.0 mmol but a 2-h value from ≥ 7.7 to < 11.1 were defined as having IGT. Subjects who did not meet the criteria for either IGT or diabetes were deemed normoglycemic (16).

Fasting and 2-h venous blood samples were centrifuged and separated immediately, and plasma glucose was measured on-site using Yellow Springs Instruments glucose analyzers (Yellow Springs, OH) within 3 h of collection. Serum samples were stored at $-20°C$ and were transported on dry ice from Mauritius to Newcastle upon Tyne, U.K., in 1987 and later to Melbourne in 1996. Serum fasting and 2-h insulin measurements were analyzed in Newcastle upon Tyne using a modified method of Soeldner and Slone (17). The interassay and intraassay coefficients of variation were 6 and 4%, respectively. Serum uric acid, fasting serum triglyceride, and HDL cholesterol levels in plasma were measured using manual enzymatic methods at the central laboratory in Mauritius. For quality assurance, every 10th sample was also analyzed in Newcastle upon Tyne. Triglyceride and total cholesterol values were consistently overestimated across the distribution and were adjusted downward using a calculated regression equation. Leptin was measured in 1996 by radioimmunoassay (Linco, St. Charles, MO). The limit of detection for the assay is 0.5 ng/ml in human serum or plasma with acceptable interassay (8.2%) and intra-assay

(4.1%) coefficients of variation. A measure of insulin sensitivity was estimated using homeostasis model assessment (HOMA %S) (18).

Statistical analysis
All statistical analyses were performed using Stata Version 6.0 (College Station, TX). Logistic regression analysis was used to estimate odds ratios (95% CIs) and P values for the independent associations between metabolic syndrome components and the odds of developing diabetes or IGT coded as a dichotomous variable (0 = normoglycemia, 1 = IGT or diabetes) (19). A backwards selection algorithm was used to arrive at the best final model. To assess whether nonlinear effects were present between diabetes or IGT occurrence and metabolic syndrome components, the risk of the dependent variable was plotted against the independent variable divided into 20 quantiles, and visual inspection was used to assess departures from linearity, which were modelled with appropriate transformations. To assess whether effect modification existed when multiple components of the metabolic syndrome were present simultaneously, first-order interaction terms were tested for significance in logistic regression models with the threshold for rejection of the null hypothesis of no interaction set at $P < 0.05$. This analysis permitted formal assessment of whether joint effects of metabolic syndrome components exceeded the expected combination of individual effects on the incidence of the outcomes of interest. Relative risk (RR) of diabetes or IGT was calculated in crude and stratified analyses using standard methods (19).

RESULTS — A total of 2,605 subjects with normal glucose tolerance in 1987 and follow-up data in 1992 were available for analysis. Mean age for study subjects was 40.4 years and ranged from a low of 25 years to a high of 74 years. Subjects included 48.9% men and 51.1% women. Among these individuals, 356 (13.7%) developed IGT and 159 (6.1%) developed diabetes during the 5-year follow-up, which gave annual incidence rates of 2.9 and 1.3%, respectively.

Characteristics of study subjects by whether diabetes or IGT developed during follow-up in univariate analyses are shown in Table 1. Sex, ethnicity, and diabetes family history were not significantly related to diabetes or IGT odds. Regarding features of

Table 1—*Baseline characteristics of study subjects in relation to the occurrence of IGT and diabetes at the end of the follow-up period*

| Characteristic	Normal glucose tolerance	IGT	Diabetes	Diabetes or IGT vs. normal glucose tolerance
n	2,090	356	159	
Age (years)	39.5 ± 11.8	43.4 ± 12.1	45.3 ± 12.3	3.3 (2.4–4.5)
Sex				
Men	48.7	40.7	70.4	1.0 (referent)
Women	51.3	59.3	29.6	1.0 (0.8–1.2)
Ethnicity				
Indian (Hindu)	54.8	55.1	54.7	1.0 (referent)
Indian (Muslim)	14.9	17.4	11.3	1.0 (0.8–1.4)
African Creole	24.9	21.1	26.4	0.9 (0.7–1.2)
Chinese	5.4	6.5	7.6	1.3 (0.8–1.9)
Family history of diabetes	20.1	22.8	21.4	1.1 (0.9–1.4)
BMI (kg/m^2)	22.8 ± 3.8	24.6 ± 4.5	24.2 ± 4.3	1.5 (1.4–1.7)
WHR	0.83 ± 0.07	0.85 ± 0.07	0.88 ± 0.08	1.7 (1.5–1.9)
Systolic blood pressure (mmHg)	122.0 ± 16.8	127.4 ± 19.3	131.2 ± 19.1	1.3 (1.2–1.5)
Diastolic blood pressure (mmHg)	75.3 ± 11.2	78.5 ± 11.2	81.5 ± 12.7	1.4 (1.3–1.5)
Serum triglycerides (mmol/l)	1.35 ± 1.04	1.50 ± 0.93	1.83 ± 1.31	See triglycerides model below
Plasma HDL cholesterol (mmol/l)	1.30 ± 0.34	1.27 ± 0.30	1.22 ± 0.32	0.8 (0.8–0.9)
Serum uric acid (mmol/l)	0.34 ± 0.08	0.36 ± 0.09	0.40 ± 0.09	1.5 (1.4–1.7)
Fasting plasma insulin (μU/ml)	6.48 ± 5.53	8.60 ± 6.50	8.38 ± 7.88	1.4 (1.3–1.5)
2-h plasma insulin (μU/ml)	37.67 ± 34.62	54.05 ± 49.81	48.85 ± 50.23	1.4 (1.3–1.6)
HOMA %S	114.6 ± 108.8	85.6 ± 92.1	98.9 ± 101.3	0.7 (0.6–0.8)
Fasting plasma glucose (mmol/l)	5.10 ± 0.49	5.25 ± 0.54	5.51 ± 0.57	1.5 (1.4–1.7)
2-h plasma glucose (mmol/l)	5.61 ± 1.14	6.39 ± 0.98	6.20 ± 1.07	2.1 (1.9–2.4)
Fasting plasma leptin (ng/ml)	8.52 ± 9.69	10.56 ± 8.80	7.52 ± 6.86	See leptin model below
Leptin model (leptin + log$_e$ plasma leptin, 25th vs. 1st percentile)				1.7 (1.4–2.1)
Leptin quintiles				
1 (lowest)				1.0 (referent)
2				1.0 (0.7–1.4)
3				1.9 (1.3–2.7)
4				2.2 (1.6–3.1)
5				2.6 (1.8–3.6)
Triglycerides model (triglycerides + log$_e$ serum triglycerides, 25th vs. 1st percentile)				2.1 (1.6–2.8)
Triglyceride quintiles				
1 (lowest)				1.0 (referent)
2				1.3 (0.9–1.9)
3				1.8 (1.3–2.5)
4				1.7 (1.2–2.4)
5				2.8 (2.0–3.9)

Data are *n*, means ± SD, %, or odds ratios (95% CIs). Odds ratios for continuous variables reflect a 1-SD magnitude increase.

the metabolic syndrome, all measures were significantly related to diabetes or IGT odds in the expected directions. Higher plasma leptin was significantly related to higher odds of diabetes or IGT. By visual examination, plots of the risk of diabetes or IGT in relation to triglyceride quantiles appeared nonlinear with a plateau followed by a slight decline at higher levels (Fig. 1*A*). A similar association was seen for plasma leptin (Fig. 1*B*). Because leptin levels differ markedly by sex in this and other populations, this rela-

tionship was reexamined by sex, but similar associations were seen (Fig. 1*C* and *D*). Therefore, multivariate models were fit with log$_e$ transformations of these variables in addition to the linear term (Table 1). Because both the linear and log$_e$ transformations of these variables were significant in these models at $P < 0.05$, this finding rejects the hypothesis that a linear association exists between diabetes or IGT odds and leptin or triglyceride levels. The relative odds of diabetes or IGT in relation to quin-

tiles of leptin and triglycerides inserted into each model shown in Table 1 as a set of dummy variables demonstrated the leveling off of risk with increasing level, more so for leptin than for triglycerides.

Analysis of leptin effects on diabetes or IGT risk was further examined by estimating the RR of diabetes or IGT in crude and stratified analyses. In crude analysis, the RR (95% CIs) of diabetes or IGT for leptin concentration above the median value was 1.7 (1.4–2.0). A higher RR was seen in men (2.0
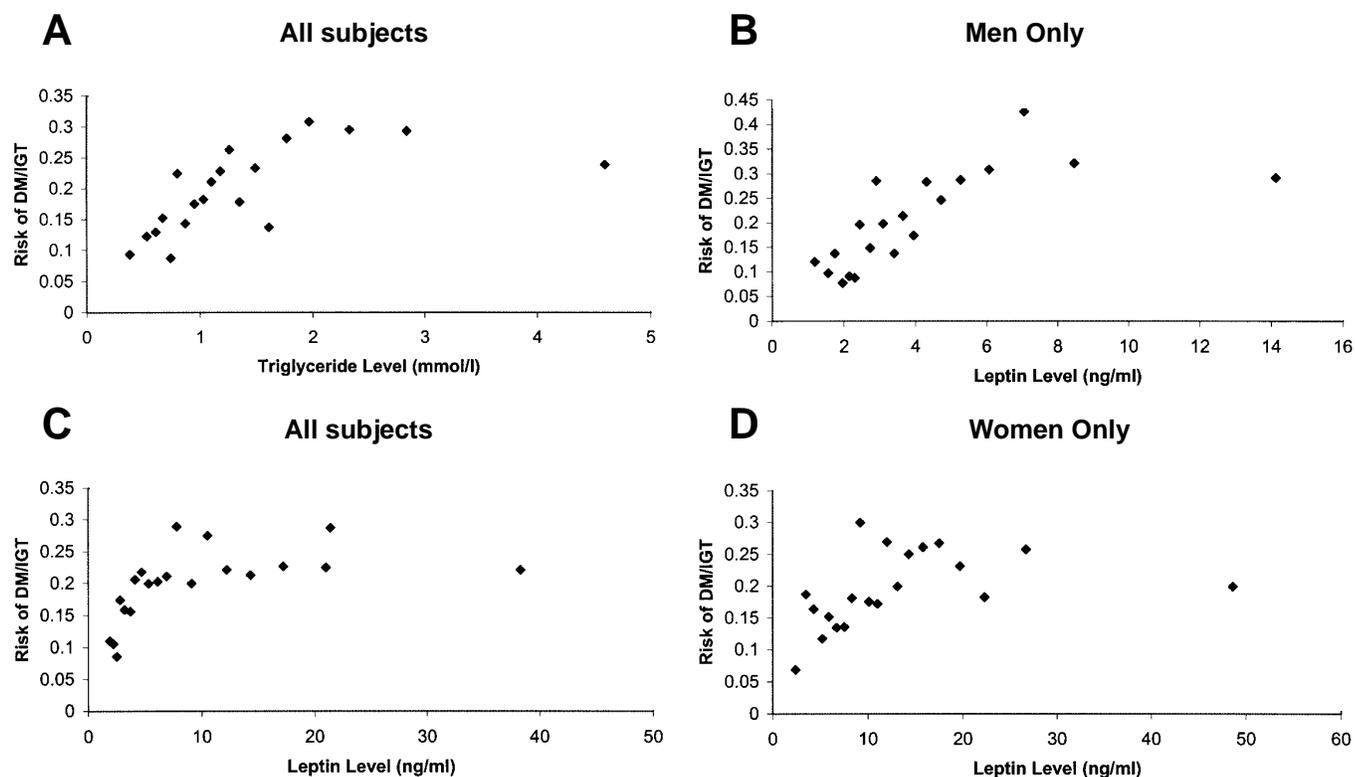
Figure 1—*Risk of the combined outcome in relation to quantiles (n = 20) of serum triglycerides (A) and plasma leptin levels (B–D). Mean triglyceride and leptin levels for each quantile are shown on the x-axis, and risk of diabetes or IGT (DM/IGT) is shown on the* y-*axis.*

[1.6–2.5]) than in women (1.5 [1.2–1.8]), but this difference was not significant by the heterogeneity χ^2 test ($P = 0.07$). In analyses stratified by BMI and WHR dichotomized at the median value (BMI: men 22.5 kg/m^2, women 23.0 kg/m^2; WHR: men 0.88, women 0.79), RRs associated with leptin concentration above the median value did not differ at higher versus lower levels of overall adiposity (BMI less than or equal to the median 1.3 [0.9–1.7]; BMI greater than the median 1.3 [1.0–1.7]) or at higher versus lower levels of WHR (WHR less than or equal to the median 1.5 [1.1–2.0]; WHR greater than the median 1.4 [1.1–1.7]).

All metabolic syndrome features were inserted into 2 logistic regression models (1 containing HOMA %S and the other containing insulin levels to avoid collinearity) for the prediction of diabetes or IGT that also contained age, sex, ethnicity, and diabetes family history (Table 2, full model). In these full models, the following variables were no longer significantly related to the outcome: BMI, HDL cholesterol level, HOMA %S, systolic and diastolic blood pressure, and fasting and 2-h insulin levels. Using backward stepwise logistical regression on these full models yielded 1 reduced

model (Table 2) that contained variables that independently predict diabetes or IGT occurrence. Nonlinear associations between these continuous variables (except for leptin and triglycerides) and diabetes incidence were tested using exponential and logarithmic transformations, but none represented an improvement in fit compared with the model shown that used simple linear terms. This analysis showed that WHR, fasting and 2-h plasma glucose levels, serum uric acid levels, leptin concentrations, and triglyceride levels significantly predicted the occurrence of diabetes or IGT, whereas BMI was nearly significantly related to this outcome ($P = 0.053$).

The reduced model was repeated with diabetes only as the outcome (Table 2). Odds ratios of a similar magnitude were seen when compared with the diabetes or IGT reduced model except in several instances. The odds ratios for serum triglycerides and the log of this value were closer to 1, whereas the fasting glucose odds ratio was greater. Statistical significance of these associations differed as would be expected because of the smaller number of diabetes ($n = 159$) versus diabetes or IGT ($n = 515$) outcomes.

Interaction was tested by the insertion of first-order interaction terms between all possible pairs ($n = 35$) of independent variables in the reduced model predicting diabetes or IGT. None of these interactions was significant. The interaction between BMI and leptin came closest with a P value of 0.107, but P values for the remaining terms were generally much larger. Also, no significant difference in the effects of the variables in the reduced model predicting diabetes or IGT were seen according to age, sex, ethnicity, or diabetes family history. Relative odds were calculated using the reduced logistic model coefficients for different comparisons of leptin and triglyceride levels adjusted for all covariates (Table 2). These comparisons demonstrate an increase in the odds of diabetes or IGT when these levels increase from near the minimum value, with leveling off and eventually a small decline in the odds at higher values (Table 2).

CONCLUSIONS — This analysis demonstrates that all features of the metabolic syndrome that we considered were related in the univariate analysis to a higher risk of worsening of glucose tolerance among members of this multiethnic study

population with normal glucose tolerance at baseline. In multivariate models, however, several of these features did not contribute independently to this risk, including blood pressure, insulin levels, HOMA %S, and HDL cholesterol levels.

Insulin resistance and/or hyperinsulinemia have been proposed as the common underlying pathophysiological mechanism(s) for the features of the metabolic syndrome, which include hyperglycemia. That hyperinsulinemia and HOMA %S, which is an indirect measure of insulin sensitivity, are not independently related to higher diabetes or IGT risk was surprising. This result should not be interpreted as refuting a role for insulin resistance in the pathogenesis of diabetes because glucose level in the fasting and stimulated states is in part a function of insulin sensitivity, possibly more so for the stimulated value (20,21). Plasma glucose levels also depend on several other factors that may not be adequately captured by insulin level or HOMA %S, including the ability of glucose to promote its own disposal (glucose effectiveness) and counterregulatory hormones (20). These other phenomena may influence future risk of worsening of glucose tolerance. Our finding is supported by an analysis of San Antonio Heart Study data, which showed no independent effect of fasting insulin on diabetes risk in a stepwise logistic model that contained fasting and 2-h glucose levels (11). Further research would help to clarify these associations.

Leptin, a hormone secreted by adipose tissue and implicated in body weight regulation, has been reported to increase the risk of diabetes in a cohort of nondiabetic Japanese Americans followed prospectively (22). In that analysis, plasma leptin concentration was positively correlated with a higher odds of diabetes or IGT, although a nonlinear association best described this relationship. As can be seen in Fig. 1 and in the analysis of leptin quintiles (Table 1), diabetes or IGT risk appears to plateau with increasing leptin concentration. After adjustment for covariates, the odds ratio for diabetes or IGT initially increases and then diminishes with increasing leptin concentration. Leptin has been shown to inhibit insulin binding in isolated rat adipocytes (23), thereby possibly decreasing peripheral insulin sensitivity, although other investigators have shown enhanced glucose uptake in rat skeletal muscle because of leptin (24). Also, leptin may inhibit release of insulin by pancreatic islet cells, which has been demonstrated in

Table 2—*Multivariate logistic regression models of the odds of conversion to diabetes or IGT from normal glucose tolerance in relation to features of the metabolic syndrome adjusted for age, sex, ethnicity, and diabetes family history*

Model	Odds ratios (95% CIs)*	P
Full model		
BMI	1.1 (1.0–1.3)	0.119
WHR	1.3 (1.1–1.5)	0.004
Systolic blood pressure	1.1 (0.9–1.3)	0.398
Diastolic blood pressure	1.0 (0.8–1.2)	0.842
Fasting insulin level	1.1 (0.9–1.2)	0.441
2-h insulin level	1.0 (0.9–1.1)	0.820
Fasting plasma glucose level	1.4 (1.2–1.6)	<0.001
2-h plasma glucose level	1.8 (1.6–2.1)	<0.001
Serum uric acid	1.4 (1.2–1.6)	<0.001
Plasma HDL cholesterol	1.0 (0.9–1.1)	0.959
Serum triglycerides + $\log_e$ serum triglycerides (25th vs. 1st percentile)	1.5 (1.1–2.1)	0.030
Plasma leptin + $\log_e$ plasma leptin (25th vs. 1st percentile)	1.3 (1.0–1.7)	0.002
Reduced model (outcome = diabetes or IGT)		
BMI	1.2 (1.0–1.4)	0.053
WHR	1.3 (1.1–1.5)	0.002
Fasting glucose level	1.4 (1.2–1.6)	<0.001
2-h glucose level	1.8 (1.6–2.1)	<0.001
Serum uric acid	1.4 (1.2–1.6)	<0.001
Serum triglycerides + $\log_e$ serum triglycerides (25th vs. 1st percentile)	1.6 (1.2–2.1)	0.0159
Plasma leptin + $\log_e$ plasma leptin (25th vs. 1st percentile)	1.3 (1.0–1.7)	0.003
Reduced model (outcome = diabetes)		
BMI	1.1 (0.8–1.4)	0.710
WHR	1.2 (0.9–1.6)	0.160
Fasting glucose level	1.8 (1.5–2.1)	<0.001
2-h glucose level	1.4 (1.1–1.7)	0.002
Serum uric acid	1.4 (1.1–1.7)	0.001
Serum triglycerides + $\log_e$ serum triglycerides (25th vs. 1st percentile)	1.4 (0.9–2.1)	0.255
Plasma leptin + $\log_e$ plasma leptin (25th vs. 1st percentile)	1.3 (0.9–2.0)	0.171
Diabetes or IGT reduced model (adjusted for age, sex, ethnicity, family history of diabetes, BMI, WHR, fasting and 2-h glucose, and uric acid level)		
Leptin comparisons (ng/ml)		
5.4 vs. 0.4 ng/ml	2.06 (0.97–4.40)	
10.4 vs. 5.4	1.02 (0.87–1.18)	
15.4 vs. 10.4	0.92 (0.83–1.01)	
20.4 vs. 15.4	0.88 (0.81–0.97)	
25.4 vs. 20.4	0.86 (0.79–0.95)	
Triglyceride comparisons (mmol/l)		
1.1 vs. 0.1	3.61 (1.51–8.67)	
2.1 vs. 1.1	1.14 (1.00–1.31)	
3.1 vs. 2.1	0.96 (0.85–1.09)	
4.1 vs. 3.1	0.90 (0.77–1.04)	
5.1 vs. 4.1	0.86 (0.73–1.02)	

The full model contains all independent variables shown in Table 1, whereas the reduced model results from backwards removal of nonsignificant independent variables from the full model. *Odds ratios are shown for a 1-SD increase in the value of the independent variable except where indicated otherwise.

vitro in human cells (25), thereby contributing to inadequate insulin secretion. Although serum leptin and insulin concentrations have been shown to be positively correlated independent of obesity (26), differences have been noted in the associations between these hormones and insulin sensitivity. The association between insulin sensitivity and leptin depends on BMI, whereas the association between insulin concentration and insulin sensitivity does not (9,27). The conflicting effects of leptin on glucose metabolism may explain the nonlinear association between leptin and diabetes or IGT odds if these effects occur differentially depending on the serum leptin concentration. Also, leptin concentration may reflect other physiological processes not directly measured in this study that alter diabetes or IGT odds independent of BMI, insulin concentration, or insulin sensitivity.

Although significantly associated with the development of diabetes or IGT in univariate analyses, the odds ratios for systolic and diastolic blood pressure approached 1 with adjustment for other important covariates. The association between blood pressure and insulin resistance and its role in the insulin resistance syndrome is inconsistent and difficult to characterize (28). Given that the effects of blood pressure on diabetes or IGT odds diminished with adjustment for covariates, blood pressure likely acts as a marker for other factors associated with a higher risk of glucose intolerance. Central body fat distribution (as reflected by higher WHR) and BMI were related to higher risk of diabetes or IGT, which was expected (29), although the latter was at borderline statistical significance. Serum triglyceride level was related to diabetes or IGT odds, whereas HDL cholesterol level was not related after adjustment for covariates. The association between triglyceride level and the outcome appeared similar to that of leptin, with leveling off of risk at higher levels and even a diminished effect as seen in the multivariate model. This nonlinear association between serum triglycerides and diabetes or IGT was unexpected and will require further investigation. A possible explanation may be that the mechanism driving very high serum triglyceride levels is not as strongly related to glucose metabolism as it is at lower levels and therefore would not be expected to confer the same elevation in risk of progression to diabetes or IGT. Other prospective research has yielded mixed results on the association between concentration of triglyceride or HDL cholesterol levels and diabetes risk in multivariate models (10,11,30–33). Serum uric acid was strongly associated with diabetes or IGT occurrence (Table 2). Uric acid has been proposed as a feature of the metabolic syndrome by several authors (3,4) but has not gained universal acceptance (1). Several other prospective or cross-sectional studies have revealed a high or borderline-high risk of diabetes and IGT in relation to increasing uric acid levels (31,32,34,35).

Features of the metabolic syndrome in combination had effects that can easily be calculated from the individual effects. Because logistic regression analysis was used to model the odds of these outcomes, the independent variables are expected to have multiplicative effects on the odds ratio. For example, the odds ratio for diabetes in relation to both a 1-SD increase in uric acid and a 1-SD increase in 2-h glucose would be $1.4 \times 1.8 = 2.5$. One definition of the presence of interaction, alternately referred to as "effect modification" or "synergism," is deviation from the multiplicative combination of individual effects and is in fact the test of interaction used in logistic regression models (19). We did not find any pairs of metabolic syndrome variables that resulted in a significantly greater- or lesser-than-expected odds ratio for diabetes or IGT. Therefore, this analysis did not support the theory that metabolic features act synergistically in combination to result in a higher (or lower) odds ratio than would be expected from combining the odds ratios for effects estimated individually.

The independent variables from the reduced model for the occurrence of diabetes or IGT were used to predict the development of diabetes only (Table 2). In general, the odds ratios for this outcome were of a similar magnitude to those in the diabetes or IGT model. *P* values were in general larger and in some cases became nonsignificant, although this may reflect the reduced power because of the smaller number of diabetes compared with diabetes or IGT outcomes. The nonlinear association between leptin and diabetes was still apparent in this analysis.

This study has several limitations. No direct measure of insulin sensitivity or body fat mass was available. Surrogate measures that imperfectly classify these variables were used instead. Bias associated with this probable nondifferential misclassification is likely to have resulted in underestimates of odds ratios. In accepting these assumptions, significant differences are therefore likely to represent underestimates of unknown magnitude. Follow-up glucose tolerance could not be assessed for ~25% of subjects. Whether these results can be generalized to ethnic groups not included in this study sample is not clear.

In conclusion, features of the metabolic syndrome were related in a nonsynyergistic manner to deterioration in glucose tolerance of Mauritian subjects during a 5-year follow-up period. Explaining a common pathway or mechanism through which all of these factors contribute to the development of glucose intolerance is difficult, which raises the possibility that multiple mechanisms may lead to this outcome. The unexpected nonlinear associations between worsening glucose tolerance and serum triglyceride or plasma leptin concentrations will require further investigation.

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