Intestinal alkaline phosphatase deficiency increases the risk of diabetes

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ABSTRACT

Introduction Our previous case–control study demonstrated that a high level of intestinal alkaline phosphatase (IAP), an endotoxin-detoxifying anti-inflammatory enzyme secreted by villus-associated enterocytes and excreted with stool, plays a protective role against type 2 diabetes mellitus (T2DM) irrespective of obesity. In the current study, we investigated the long-term effect of IAP deficiency (IAPD) on the pathogenesis of T2DM.

Research design and methods A healthy cohort of participants without diabetes (30–60 years old), comprising 188 without IAPD (IAP level: ≥65 U/g stool) and 386 with IAPD (IAP level: <65 U/g stool), were followed up for 5 years. We measured stool IAP (STAP) and fasting plasma glucose, and calculated the risk ratio (RR) using log-binomial regression model.

Results T2DM incidence rates were 8.0%, 11.7%, 25.6%, and 33.3% in participants with ‘persistent no IAPD’ (IAP level: always ≥65 U/g stool), ‘remittent IAPD’ (IAP level: increased from <65 U/g stool to ≥65 U/g stool), ‘persistent IAPD’ (IAP level: always <65 U/g stool), and ‘incident IAPD’ (IAP level: decreased from ≥65 U/g stool to <65 U/g stool), respectively. Compared with ‘persistent no IAPD’ the risk of developing T2DM with ‘incident IAPD’ was 270% higher (RR: 3.69 (95% CI 1.76 to 7.71), p<0.001). With ‘incident IAPD’ the risk was 230% higher (RR: 2.24 (95% CI 0.99 to 5.11), p=0.0541). Sensitivity analyses of persistent IAP levels revealed that, compared with participants of the highest persistent IAP pentile (always >115 U/g stool), the rate of increase of fasting glycemia was double and the risk of developing T2DM was 1280% higher (RR: 13.80 (95% CI 8.87 to 101.3), p=0.0099) in participants of the lowest persistent IAP pentile (always <15 U/g stool). A diabetes pathogenesis model is presented.

Conclusions IAPD increases the risk of developing T2DM, and regular STAP tests would predict individual vulnerability to T2DM. Oral IAP supplementation might prevent T2DM.

INTRODUCTION

Type 2 diabetes mellitus (T2DM), a chronic metabolic disease manifested with hyperglycemia and insulin resistance, is a major global health problem with devastating consequences in terms of morbidity, mortality, and healthcare costs. T2DM affected approximately 463 million adults worldwide (9.3% of world population) in 2019 and is projected to affect approximately 723 million (10.3%) in 2030 and 700 million (10.8%) in 2045. The pathogenesis of T2DM is poorly understood. Various factors have been postulated to be involved in the development of T2DM, notably genetic polymorphism, ethnicity, metabolic syndrome, obesity, diets, infection, dysbiosis, autoimmunity, drugs, stress, and pregnancy.1–11

Significance of this study

What is already known about this subject?

► The deficiency of intestinal alkaline phosphatase (IAP), an anti-inflammatory gut enzyme secreted by intestinal enterocytes and excreted with stool, is directly associated with type 2 diabetes mellitus (T2DM), and a high level of IAP plays a protective role against T2DM irrespective of obesity.

► IAP deficiency (IAPD) is also directly associated with type 1 diabetes mellitus.

What are the new findings?

► This 5-year prospective cohort study shows that IAPD increases the rate of fasting plasma glucose.

► Intervention of IAPD by oral IAP supplementation might prevent T2DM.

► Eradication of the worldwide pandemic of T2DM is possible by early diagnosis of IAPD-associated vulnerability to T2DM followed by pre-emptive preventive measures.

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Low-grade systemic inflammation has been implicated to play a pivotal role in the pathogenesis of T2DM.\(^8\)\(^-\)\(^11\) Death of Gram-negative bacteria results in fragmentation of the cell wall, releasing its lipopolysaccharide (LPS), an endotoxin that causes inflammation. Intestinal luminal LPS is usually excreted with stool; however, under certain circumstances, such as increased gut permeability and increased intake of a high-fat diet, alcohol, and fructose, LPS may translocate to blood circulation, causing endotoxemia.\(^\)\(^12\) Recently, ‘metabolic endotoxemia’, defined as persistently increased levels of LPS in blood, has been shown to induce low-grade systemic inflammation, leading to insulin resistance, hyperglycemia (T2DM), dyslipidemia, and fatty liver in mice.\(^7\)\(^12\)\(^13\) We have previously shown that LPS is detoxified by intestinal alkaline phosphatase (IAP), a gut enzyme secreted by villus-associated enterocytes and partially excreted with stool.\(^7\)\(^12\) IAP functions as an anti-inflammatory enzyme by detoxifying LPS and other bacterial toxins through dephosphorylation (see online supplemental appendix B for IAP functions).\(^7\)\(^12\)\(^14\)\(^15\) We have previously demonstrated that mice deficient in IAP (IAP knockout, Akp\(^-/-\)) develop metabolic syndrome (hyperglycemia, dyslipidemia, and fatty liver) that can be prevented by oral IAP supplementation.\(^\)\(^12\) Our subsequent case–control human study identified that the average level of IAP was approximately 65 U/g stool in the healthy population,\(^7\) and accordingly an IAP level of ≥65 U/g stool is considered high (normal) and protective against T2DM. In contrast, the average IAP level detected in patients with T2DM was only 35 U/g stool.\(^7\) Obese people with normal IAP level did not develop T2DM. The study established that IAP deficiency (IAPD), defined as <65 U of IAP per gram of stool, is directly associated with T2DM.\(^7\)\(^12\) Based on the above-mentioned coherent animal and human studies,\(^7\)\(^12\) we hypothesized that IAPD might be an independent risk factor of T2DM in humans. Thus, we conducted a 5-year prospective cohort study and here we report that IAPD plays a pivotal role in the development of T2DM. We also provide a model defining the pathway of IAPD-mediated pathogenesis of diabetes.

**METHODS**

**Study design and participants**

We followed up the health status of a healthy cohort of 671 participants without diabetes (30–60 years old) of our previous case–control study\(^7\) for 5 years. IAPD was considered as the major exposure variable and incidence of T2DM as the outcome variable, where baseline physical and biochemical profiles, including body mass index (BMI), fasting plasma glucose (FPG), and IAP values of each participant, were taken into account for outcome measurement (figure 1; see online supplemental appendix C for details).

Each participant consented to participate in the study and signed an informed consent form approved by the National Research Ethics Committee (see Ethics approval).

**RESULTS**

Both baseline and follow-up data were available for 574 participants and were thus included in the study, and 97 participants were excluded because of unavailability of follow-up data due to migration, refusal, pregnancy, and death (figure 1). There was no significant difference between the included and excluded groups in terms of physical and biochemical parameters (online supplemental table S1). The characteristics of all included participants are shown in online supplemental table S2.

Based on the follow-up data, we observed that 20% of the participants developed T2DM in 5 years.

**Laboratory procedures**

We measured blood pressure and body mass index (BMI), and serum cholesterol, high-density lipoproteins (HDL), low-density lipoproteins (LDL), triglycerides, creatinine, and alanine aminotransferase (ALT). The T2DM status of each participant was confirmed by measuring fasting plasma glucose (FPG) (at least 10 hours) concentration and/or hemoglobin A1c (HbA1c) concentration. An FPG level of ≥7.0 mmol/L (126 mg/dL) or HbA1c level ≥6.5% was considered diagnostic for T2DM (see online supplemental appendix C for details).\(^1\) Stool alkaline phosphatase (STAP) assay was performed following the procedures as previously described (see online supplemental appendix C for details).\(^7\) In brief, a specific amount of stool was dissolved in a specific amount of stool dilution buffer (eg, 100 mg stool to 5 mL stool dilution buffer), homogenized and centrifuged, and supernatant containing STAP was collected and followed by measuring STAP concentration by an automatic chemistry analyzer. STAP assays were performed by laboratory technologists who were blinded to the diagnoses of participants. Because 80% of STAP is IAP and 20% of STAP is of bacterial origin,\(^7\) IAP is synonymously used for STAP in this article.

**Statistical analysis**

Statistical analysis was performed using the SAS V.9.4 software. \(^2\)\(^*\) test was used to examine the association between IAPD group and T2DM status. The correlation between IAP and different risk factors of T2DM was evaluated by Pearson’s correlation coefficient. Adjusted generalized linear model of regression analysis was used to determine adjusted mean difference in IAP between T2DM groups (T2DM vs non-T2DM). Log-binomial regression analysis was used to calculate the unadjusted and adjusted risk ratios (RR) (see online supplemental appendix C for details). We also performed repeated measure analysis of variance (ANOVA) to examine the association of the IAP level with T2DM. A p value of <0.05 was considered statistically significant.
Confirming our previous results,\textsuperscript{7} we found that a high IAP level plays a protective role against T2DM (figure 2A). The percentile distribution shows that IAP levels are low in patients with T2DM at all percentile points compared with healthy participants without diabetes (controls) (figure 2B). We also found that obese people with high IAP do not develop T2DM (figure 2C).

The baseline characteristics of participants grouped by IAPD status are shown in online supplemental table S3.
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and the follow-up characteristics of the IAPD groups are shown in table 1.

Validating our previous data,7 Pearson’s correlation coefficient analysis showed no correlation between IAP level and age, gender, FPG, BMI, ALT, serum creatinine, blood pressure, or lipid level (online supplemental table S4). A generalized linear regression model predicted a strong association of IAP with T2DM, confirming our previous observation (table 2).

A repeated measure ANOVA also showed a very strong association of IAP with T2DM (online supplemental table S5).

We assessed unadjusted relative risks of developing T2DM in different IAPD groups (figure 3). Compared
with the ‘persistent no IAPD’ group, we observed that the ‘incident IAPD’ and ‘persistent IAPD’ groups had significantly higher risk of developing T2DM, especially approximately 7-fold increased risk in younger men (30–39 years old) with ‘incident IAPD’. In contrast, like ‘persistent no IAPD’, ‘remittent IAPD’ was associated with less risk of developing T2DM.

Table 3 shows the incidence rates of T2DM, which were 8.0%, 11.7%, 25.6%, and 33.3% in participants with ‘persistent no IAPD’, ‘remittent IAPD’, ‘persistent IAPD’, and ‘incident IAPD’, respectively (also see figure 2D).

The data indicated that compared with the ‘persistent no IAPD’ group, the incidence rate was more than 4-fold higher in the ‘incident IAPD’ group. Table 3 also shows the relative risks of developing T2DM in different IAPD groups after adjusting for various confounding factors, such as age, gender, BMI, systolic blood pressure, diastolic blood pressure, cholesterol, HDL, LDL, triglycerides, creatinine, ALT, and FPG. Compared with ‘persistent no IAPD’, ‘incident IAPD’ was associated with 3.7-fold increased risk of developing T2DM (RR: 3.69 (95% CI 1.76 to 7.71), p<0.001). Similarly, ‘persistent IAPD’ was associated with 3.3-fold increased risk (RR: 3.27 (95% CI 1.64 to 6.50), p<0.001). In contrast, the relative risk of developing T2DM in the ‘remittent IAPD’ group was insignificant (RR: 2.24 (95% CI 0.99 to 5.11), p=0.0541).

To delineate the sensitivity of different persistent levels of IAP on developing T2DM, we calculated T2DM incidence rates, relative risks, and rates of increase of FPG for five persistent levels of IAP (table 4).

Compared with participants with the highest levels of persistent IAP (always >150 U/g stool, pentile 5), participants with the lowest level of persistent IAP (always ≤15.0 U/g stool, pentile 1) had approximately 16.0-fold higher incidence rate of T2DM (2.9% vs 46.7%, respectively). As expected, the age-adjusted and gender-adjusted relative risk also increased by 13.80-fold in pentile 1 (RR: 13.80 (95% CI 1.87 to 101.3), p=0.0099).

A graphical presentation of percentage of risk of T2DM associated with each pentile is shown in online supplemental figure S1. Table 4 also shows that, compared with pentile 5, the rate of increase in glycemia was double in pentile 1 (17.4% vs 34.1%, respectively, p<0.001). The rate of increase in FPG was not dependent on age (online supplemental table S6).

**DISCUSSION**

This observational prospective cohort study shows that IAPD is an independent risk factor of T2DM. In the context of confounding factors affecting IAP levels, confirming our previous observations,7 Pearson’s correlation coefficient analysis showed that age, gender, and
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various physical and biochemical risk factors do not have any significant effect on IAP levels (online supplemental table S4). The participants were on an unrestricted diet, and it is expected that if the sample sizes of related groups are high the diet should not affect the outcome of the study because both groups would be similarly affected by diet. The sample sizes of related groups in this study were large enough to achieve more than 90% post-hoc statistical power (figure 3) and hence we believe it is unlikely that diet had any effect on the outcome of study. The participants were from diverse socio-economic background and therefore we think that socio-economic conditions did not impact the outcome of the study. Obesity has been implicated in the pathogenesis of T2DM;1–3 however, we found that obese people with a high level of IAP do not develop T2DM (figure 2C), indicating a critical role of IAP in preventing T2DM even in the presence of obesity.

This study established IAPD as an independent risk factor of T2DM, and adjusted relative risk analyses demonstrated that various confounding factors did not have any effect on the outcome of the study (table 3). Regarding the cause and consequence relationship between IAPD and T2DM, at the beginning of study all participants were without diabetes and 386 participants had IAPD and therefore IAPD cannot be a consequence of T2DM; instead T2DM is the consequence of IAPD. In the context of IAPD being a causal factor of T2DM, we anticipate that a case–control interventional study with oral supplementation of IAP will be required. Based on our previous study, we proposed that ‘temporal IAP profiling’, defined as regular monitoring of stool IAP, should be able to diagnose if a person has ‘incipient (latent) metabolic syndrome’, including ‘incipient T2DM’.7 The current study also established that ‘temporal IAP profiling (regular STAP tests)’ will identify an individual vulnerable to develop T2DM.

We have previously shown that oral IAP supplementation prevents T2DM in mice,12 and the current human study shows that ‘remittent IAPD’ prevents T2DM (table 3). Based on these studies, we anticipate that oral IAP supplementation might prevent T2DM in humans. We strongly believe that IAPD as an independent risk factor plays a pivotal role in the pathogenesis of T2DM in humans based on the following findings of our previous and current studies: (1) IAPD (IAP gene knockout) causes diabetes (hyperglycemia) in mice;13 (2) chronic inhibition of intestinal luminal IAP activity in mice by oral supplementation of phenylalanine, an inhibitor of IAP, in drinking water causes diabetes; (3)
IAPD status | Total | T2DM (n, %) | RR (95% CI)
--- | --- | --- | ---
Males
Persisten no IAPD | 31 | 2 (6.5) | 1.00
Incident IAPD | 16 | 7 (43.8) | 6.78 (1.59-28.94)
Persistent IAPD | 80 | 19 (23.8) | 3.68 (0.91-14.88)
Remittent IAPD | 41 | 3 (7.3) | 1.13 (0.20-6.38)
Females
Persisten no IAPD | 94 | 8 (8.5) | 1.00
Incident IAPD | 47 | 14 (29.8) | 3.50 (1.08-7.75)
Persistent IAPD | 186 | 49 (26.3) | 3.10 (1.53-6.26)
Remittent IAPD | 79 | 11 (13.9) | 1.64 (0.89-3.67)
Age 30 – 39 years
Persisten no IAPD | 60 | 3 (5.0) | 1.00
Incident IAPD | 30 | 11 (36.7) | 7.33 (2.21-24.32)
Persistent IAPD | 87 | 16 (18.4) | 3.68 (1.12-12.07)
Remittent IAPD | 68 | 5 (7.4) | 1.47 (0.37-5.90)
Age 40 – 49 years
Persisten no IAPD | 40 | 5 (12.8) | 1.00
Incident IAPD | 15 | 6 (40.0) | 3.20 (1.14-8.94)
Persistent IAPD | 96 | 28 (29.2) | 2.33 (0.97-5.61)
Remittent IAPD | 27 | 4 (14.8) | 1.19 (0.35-4.02)
Age 50 – 60 years
Persisten no IAPD | 25 | 2 (8.0) | 1.00
Incident IAPD | 18 | 4 (22.2) | 2.78 (0.57-13.56)
Persistent IAPD | 83 | 24 (28.9) | 3.61 (0.92-14.24)
Remittent IAPD | 25 | 5 (20.0) | 2.50 (0.53-11.70)

Figure 3 Association between intestinal alkaline phosphatase deficiency (IAPD) and incidence of type 2 diabetes mellitus (T2DM) stratified by gender and age. A prospective cohort of healthy participants without diabetes (n=574, 30–60 years old) were followed up for 5 years. Based on baseline and follow-up IAP values, the participants were classified as having ‘persistent no IAPD’ (intestinal alkaline phosphatase (IAP) level: always ≥65 U/g stool), ‘persistent IAPD’ (IAP level: always <65 U/g stool), ‘incident IAPD’ (IAP level: decreased from ≥65 U/g stool to <65 U/g stool), or ‘remittent IAPD’ (IAP level: increased from <65 U/g stool to ≥65 U/g stool). Post-hoc statistical power analyses revealed adequate power (conventionally, >80%) for different comparative groups (‘persistent no IAPD’ vs ‘incident IAPD’: 98.6%; ‘persistent no IAPD’ vs ‘remittent IAPD’: 99.4%; ‘persistent no IAPD’ vs ‘remittent no IAPD’: 90.0%), validating the adequacy of sample size for each group as calculated by an online program (http://clinicalcalc.com/Stats/Power.aspx). Data are unadjusted relative risk (RR) and 95% CI.

Chronic oral supplementation of IAP prevents diabetes in IAP knockout mice; (4) IAPD is directly associated with T2DM in humans; (5) overall incidence rate of T2DM is approximately 4.2-fold higher in humans with IAPD; (6) overall relative risk of developing T2DM is increased by approximately 3.7-fold in humans with IAPD; (7) the lower the persistent IAP level the higher (approximately 16.1-fold) the incidence rate of T2DM in humans; (8) the lower the persistent IAP level the higher (approximately 13.8-fold) the relative risk of developing T2DM in humans; (9) the lower the persistent IAP level the higher (approximately twofold) the rate of increase of fasting glycemia in humans; and (10) remission of IAPD prevents T2DM in humans, as shown in this study and in Malo’ and Kaliannan et al. It is to note that an association between high serum alkaline phosphatase (liver/bone/kidney alkaline phosphatase) and new-onset T2DM has been reported in a population with hypertension. Serum alkaline phosphatase is a proinflammatory marker and we think increased serum alkaline phosphatase might indicate IAPD-induced systemic inflammation in this patient population. Based on this study and other published data, we developed a model depicting the molecular mechanisms of IAPD-mediated diabetes pathogenesis (figure 4). Brief description and related references on the steps of the diabetes pathogenesis pathway are provided in the following sections.

Steps 1 and 2: induction of dysbiosis
Dysbiosis is defined as an imbalance in type and number of normal microbes, and numerous diseases have been linked to dysbiosis including T2DM, type 1 diabetes mellitus (T1DM), cancer, coronary artery disease, and AIDS. The homeostasis of normal intestinal microbiota could be disrupted by various factors, leading to dysbiosis such as diets, infections, antibiotics, chemotherapy, and radiation. It is anticipated that some dysbiotic factors might reduce the number of bacteria that produce sodium butyrate (NaBu) (see steps 3 and 4).

Steps 3 and 4: sodium butyrate deficiency
Butyric acid, a short-chain fatty acid, is produced by fermentation of dietary fibers by the gut microbiota, and the deficiency of sodium butyrate (NaBu, sodium salt of butyric acid) is associated with specific types of dysbioses. Lassenius et al. showed that NaBu deficiency as well as IAPD are associated with T1DM. In a metagenome-wide association study of gut microbiota in T2DM, Qin et al. demonstrated a decrease of some species of universal butyrate-producing bacteria.

Steps 5–7: IAP enzyme deficiency
Histone deacetylases (HDACs) inhibit the transcription of a gene, and an HDAC inhibitor, such as NaBu, increases transcription. We have previously shown that, in human colon cancer HT-29 cells, NaBu transcriptionally activates the IAP gene by H3 histone hyperacetylation. It has been shown that NaBu upregulates the IAP gene in pig intestine explants. Based on the fact that specific types of dysbioses can decrease butyrate production (steps 3 and 4), it is expected that under such a dysbiotic condition the transcription of IAP gene is reduced or silenced, resulting in IAP enzyme deficiency. It is anticipated that factors described in step 1 can also directly affect intestinal physiology, leading to inhibition of IAP activity and/or gene expression.

Steps 8 and 9: development of metabolic endotoxemia
Metabolic endotoxemia is defined as a twofold to threefold persistent increase of bacterial endotoxin LPS in blood circulation compared with normal levels. Previously, we have shown that IAP dephosphorylates and thus...
detoxifies various proinflammatory factors, including LPS, lipoteichoic acids, cytokine-phosphate-guanosine DNA, ATP, uridine diphosphate, and flagellin.7 12 We have shown that IAP leads to endotoxemia.12 It has also been demonstrated that endotoxemia is precipitated by alcohol, fructose, and stress.31

Steps 10–12: chronic systemic inflammation and irreversible IAP gene silencing

We and other laboratories have shown that metabolic endotoxemia in mice precipitates low-grade systemic inflammation as evidenced by increased serum levels of proinflammatory cytokines tumor necrosis factor-alpha (TNF-α) and interleukin 1beta (IL-1β).12 13 Also, human volunteers receiving endotoxin as intravenous bolus injection or infusion showed increased levels of TNF-α, IL-6, and IL-1β.32 33

In the presence of NaBu, we have previously shown that proinflammatory cytokines TNF-α and IL-1β cannot inhibit the expression of IAP in HT-29 cells. However, in the absence of NaBu, these cytokines inhibit the expression of IAP gene, and post-treatment with NaBu cannot reverse this IAP silencing, which indicates that proinflammatory cytokines in the absence of NaBu probably permanently silence the IAP gene.34 It is to point out that inflammation can also cause dysbosis, further promoting IAPD.17

Steps 13 and 14: insulin resistance and hypoinsulinemia

Insulin resistance, defined as the impaired ability of circulating insulin to regulate normal cellular glucose uptake, occurs due to defective insulin-mediated signal transduction.35 Insulin binds to its receptor and activates its kinase domain, which is followed by activation of the insulin receptor substrate family (IRS 1–6), the scaffolding proteins

### Table 3

Intestinal alkaline phosphatase deficiency (IAPD) is associated with increased risk of developing type 2 diabetes mellitus (T2DM)

<table>
<thead>
<tr>
<th>IAPD status</th>
<th>Total participants</th>
<th>T2DM n (%)</th>
<th>Relative risk (95% CI)</th>
<th>Model 1*</th>
<th>Model 2†</th>
<th>Model 3‡</th>
<th>Model 4§</th>
<th>Model 5¶</th>
</tr>
</thead>
<tbody>
<tr>
<td>Persistent no IAPD</td>
<td>125</td>
<td>10 (8.0)</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
<td>1.00</td>
</tr>
<tr>
<td>Incident IAPD</td>
<td>63</td>
<td>21 (33.3)</td>
<td>(2.09 to 8.30)***</td>
<td>3.78</td>
<td>4.31</td>
<td>4.17</td>
<td>3.69</td>
<td></td>
</tr>
<tr>
<td>Persistent IAPD</td>
<td>266</td>
<td>68 (25.6)</td>
<td>(1.89 to 7.54)***</td>
<td>3.01</td>
<td>3.33</td>
<td>3.27</td>
<td>3.27</td>
<td></td>
</tr>
<tr>
<td>Remittent IAPD</td>
<td>120</td>
<td>14 (11.7)</td>
<td>(1.60 to 6.71)***</td>
<td>3.16</td>
<td>3.27</td>
<td>3.27</td>
<td>3.27</td>
<td></td>
</tr>
</tbody>
</table>

***p<0.001.

*Model 1: unadjusted model.
†Model 2: adjusted for age and sex at baseline (visit 1).
‡Model 3: adjusted for variables in model 2 plus body mass index, systolic blood pressure, and diastolic blood pressure at baseline (visit 1).
§Model 4: adjusted for variables in model 3 plus creatinine, cholesterol, high-density lipoproteins (HDL), low-density lipoproteins (LDL), triglycerides, and alanine aminotransferase at baseline (visit 1).
¶Model 5: adjusted for variables in model 4 plus fasting plasma glucose at baseline (visit 1).

### Table 4

The lower the persistent IAP level the higher the rate of increase of fasting glycaemia and relative risk of developing T2DM

<table>
<thead>
<tr>
<th>Pentile</th>
<th>Persistent IAP level (U/g stool)</th>
<th>Participants (T2DM, %)</th>
<th>Relative risk (95% CI)</th>
<th>First visit FPG level, mmol/L (mg/dl)</th>
<th>Second visit FPG level, mmol/L (mg/dl)</th>
<th>Percentage increase of FPG level</th>
</tr>
</thead>
<tbody>
<tr>
<td>Pentile 1</td>
<td>0–15.0</td>
<td>30 (14, 46.7)</td>
<td>13.8 (1.87 to 101.3)**</td>
<td>4.4±0.7 (79.2±12.6)</td>
<td>6.9±2.5 (124.2±45.0)</td>
<td>34.1***</td>
</tr>
<tr>
<td>Pentile 2</td>
<td>15.1–33.0</td>
<td>38 (8, 21.1)</td>
<td>6.9 (0.91 to 52.60)</td>
<td>4.5±0.8 (81.0±14.4)</td>
<td>6.1±1.0 (109.8±18.0)</td>
<td>25.9*</td>
</tr>
<tr>
<td>Pentile 3</td>
<td>33.1–55.0</td>
<td>39 (9, 23.1)</td>
<td>7.0 (0.93 to 52.57)</td>
<td>4.3±0.8 (77.4±14.4)</td>
<td>6.0±1.1 (108.0±19.8)</td>
<td>25.2**</td>
</tr>
<tr>
<td>Pentile 4</td>
<td>55.1–115.0</td>
<td>58 (8, 13.8)</td>
<td>4.8 (0.63 to 36.71)</td>
<td>4.3±0.8 (77.4±14.4)</td>
<td>5.6±1.0 (100.8±18.0)</td>
<td>23.9*</td>
</tr>
<tr>
<td>Pentile 5</td>
<td>&gt;115.0</td>
<td>34 (1, 2.9)</td>
<td>1.0</td>
<td>4.5±0.7 (81.0±12.6)</td>
<td>5.5±0.7 (99.0±12.6)</td>
<td>17.3</td>
</tr>
</tbody>
</table>

All participants, 30–60 years old, were stratified into pentiles based on baseline IAP levels, and then participants with persistent IAP level (during follow-up visit) were identified.

Data are summarized as mean (average)±SD for each variable.

Pentile 5 was the reference pentile for calculating the statistical significance of difference in the percentage of increase of FPG levels (Student’s t-test) as well as evaluating age-adjusted and gender-adjusted relative risk between pentiles.

*p<0.05, **p<0.01, ***p<0.001.

FPG, fasting plasma glucose; IAP, intestinal alkaline phosphatase; T2DM, type 2 diabetes mellitus.
required to mediate signal transduction. Gene knockout studies revealed that mice deficient in IRS-1 develop insulin resistance, whereas mice deficient in IRS-2 develop insulin resistance and also beta cell apoptosis, leading to hypoinsulinemia. The proinflammatory cytokines TNF-α and IL-1β inhibit IRS proteins, leading to insulin resistance, hypoinsulinemia, hyperglycemia, and T2DM. Further, it is well known that chronic inflammation inflicts damages to pancreatic beta cells, vascular endothelial cells, and hepatocytes, and dysfunction of these cells precipitates metabolic syndrome, characterized by hyperglycemia, hypertension, and dyslipidemia. Metabolic syndrome ultimately leads to the development of T2DM, heart disease, hypertension, and other metabolic diseases. The model depicted here exemplifies the IAP-mediated pathogenesis pathway of diabetes.

CONCLUSIONS

IAPD increases the risk of diabetes. Regular monitoring of stool alkaline phosphatase (STAP tests) would identify individual vulnerability to diabetes. Oral IAP supplementation might prevent diabetes.

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Acknowledgements We are grateful to Professors Syed S Islam, MBBS, PhD (Epidemiology, West Virginia University), Abu Abdullah, MBBS, MPH, PhD (Epidemiology, Boston University), and Abdullah Baqui, MBBS, MPH, DrPH (Johns Hopkins University), for their critical reviews of the manuscript and statistical analyses. We dedicate the article to Professor Syed S Islam, who recently died.

Contributors Study concept and theory: MSM; research design: MSM, AKAK, MJA, JM, TS, SI, KK, MAH, MFP; participant recruitment: MSM, JM, MMHR, GB, SAT, SKB, TS; data acquisition: MSM, JM, MMHR, GB, SAT, SKB, TS; data analyses and interpretation: MSM, AKAK, MJA, JM, SKB, TS, SI, KK, MAH, SR, MFP; drafting of the manuscript: MSM, MJA, AKAK, JM, SKB, TS, SI, SR, KK, MAH, SR, MFP; critical review of the manuscript for important intellectual content: all authors; obtained funding: MSM; approval of the manuscript: all authors; study supervision: MSM. MSM is the guarantor of this work.

Funding The study was funded by grants (to MSM) from the Ministry of Education, the People’s Republic of Bangladesh (Dhaka, Bangladesh; grant number: LS2019973) and the Bangladesh Medical Research Council (Dhaka, Bangladesh; grant number: BMRC/HN/SP-Grant/2019-2020/569(1-60)). The study was further supported by generous personal donations from Dr Mohammed Salehuddin, MD (Tuscaloosa, Alabama, USA) and Dr ABM Haroon, MBBS (Samorita Hospital, Dhaka, Bangladesh). The funding sources (Ministry of Education, the People’s Republic of Bangladesh, Dhaka, Bangladesh, and the Bangladesh Medical Research Council, Dhaka, Bangladesh) did not play any role in the development of theory and study design. The funding sources were also not involved in the collection, analyses, and interpretation of data, or in writing and submission of the manuscript. The

Figure 4 The pathway of diabetes pathogenesis. IAP, intestinal alkaline phosphatase; IL-1β, interleukin-1β; IRS, insulin receptor substrate; JNK, c-Jun N-terminal kinases; LPS, lipopolysaccharides; LTA, lipoteichoic acids; NF-κB, nuclear factor-κB; NSAID, non-steroidal anti-inflammatory drugs; TNF-α, tumor necrosis factor-α.
corresponding author had full access to all the data in the study and took the
decision for publication of the data.

Competing interests None declared.

Patient consent for publication Not required.

Ethics approval This study involves human participants and was reviewed and
approved by the Bangladesh Medical Research Council, Government of the People’s
Republic of Bangladesh, Dhaka, Bangladesh (BMRC registration number: 150 02
10 10; ref: BMRC/NREC/2016-2018/28). Participants gave informed consent to
participate in the study before taking part.

Provenance and peer review Not commissioned; externally peer reviewed.

Data availability statement All data relevant to the study are included in the
article or uploaded as supplementary information. All data generated or analyzed
during this study are included in the published article. No applicable resources
were generated or analyzed during the current study.

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