

Postprandial Insulin and Glucagon Secretions during a Mixed-Meal Tolerance Test in Dietary Induced Metabolic Syndrome Cynomolgus Monkeys

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Abstract

Objective: The objective of this investigation was to assess the postprandial glucose, insulin and glucagon responses to a mixed-meal tolerance test (MMTT) in Cynomolgus monkeys with dietary-induced obese, pre-diabetes or overt type-2 diabetes mellitus (T2DM) and compare them to chow-fed controls.

Method: Mixed meal was administered via nasogastric gavage in eighteen adult male Cynomolgus monkeys stratified into four groups - either T2DM (n=3), pre-diabetes (n=5), metabolically healthy obese (n=5) or chow-fed control (n=5). The plasma was sampled for fasting and postprandial glucose, insulin and glucagon concentrations. And based on these, we calculated areas under curves (AUCs) of postprandial glucose, insulin and glucagon. Homeostasis model assessment-estimated insulin resistance (HOMA-IR) and insulinogenic Index (IGI) were used to estimate pancreatic secretion and insulin resistance.

Results: Compared to control group, the T2DM, pre-diabetes and obese groups had higher plasma fasting glucose and insulin concentrations, and a greater postprandial glucose and insulin AUC_{0-240 min}. The HOMA-IR and IGI indicated that the obese, pre-diabetes and T2DM animals are more insulin resistant and have impaired beta cell function compared to animals in control group. The highest and lowest fasting plasma glucagon concentrations were found in the pre-diabetes and T2DM groups, respectively and both groups showed moderate postprandial glucagon AUC_{0-240 min}. Mild fasting plasma glucagon but a greatest postprandial glucagon AUC_{0-240 min} was found in obese group. Moderate fasting plasma glucagon and a smallest postprandial glucagon AUC_{0-240 min} were found in the control group.

Conclusion: The results of postprandial insulin and glucagon secretions in responses to the postprandial glucose challenge enclosed with anthropometric and lipidemic parameters well demonstrated the insulin sensitivity with beta and alpha cells function in each classified group and the very similar patterns to humans. It revealed that dietary induced metabolic disorders in Cynomolgus monkeys are useful models to study the therapeutically agents for human metabolic syndrome and T2DM.

Keywords: NHP; Type 2 diabetes; Insulin resistance; Obesity; Glucagon; Insulin

lipoprotein-cholesterol; LDL-C: Low density lipoprotein-cholesterol; TG: Total triglyceride; FFA: Free fat acid.

Abbreviations

MMTT: Mixed-meal tolerance test; MS: Metabolic syndrome; HbA1C: Hemoglobin A1C; IVGTT: Intravenous glucose tolerance test; OGTT: Oral glucose tolerance test; IACUC: Institutional animal care and use committee; T2DM: Type 2 diabetes; AUC: Areas under curve; IGI: Insulinogenic index; TC: Total cholesterol; HDL-C: High density

Introduction

Insulin and glucagon are the primary gluco-regulatory hormones; regulate the levels of blood glucose in our body. The two hormones work in balance to maintaining the euglycemia level in normal physiological conditions.

Insulin is produced from beta islet cells of pancreas. As an anabolic hormone, in response to postprandial glucose removal from circulating

by signaling the highly insulin – responsive peripheral tissues such as skeletal muscle, adipose, and liver to increasing glucose uptake, and triglyceride storage and promoting of glycogenesis in liver while also suppresses the glucose production and inhibiting the suppress of glucagon secretion [1].

With an opposite function of insulin, glucagon is produced from alpha islet cells and its secretion is complex. Involving the combination of paracrine, autocrine hormonal as well as autonomic neural mechanisms [2]. However, in addition to glucose and insulin, glucagon secretion is regulated by gut incretin hormones of glucagon-like peptide 1 (GLP-1) and glucose dependent insulinotropic polypeptide (GIP) [3]. As a catabolic hormone, glucagon acts in response to raising and maintaining circulating glucose level to meet physiological needs during the fasting state, or the hypoglycemia condition mainly through the hepatic glucose production [1]. Both hormones play a vital role in regulating blood glucose levels, and if the level of one hormone is outside the ideal range, blood sugar levels may elevate or drop.

Imbalanced blood glucose levels related to genetic insulin hormone defect and the insulin resistance with further overt type 2 diabetes. It has also been well recognized that overtime of excessive intake and physical inactivity causes obesity that consequent hyperglycemia, metabolic syndrome, insulin resistance (pre-diabetes) and then type 2 diabetes.

In early of pre-diabetes pathological process, insulin secretion has initially increased in beta cell function compensatory way to facilitate entry of excess blood glucose to peripheral tissues and liver for utilization of energy and storage via glycogenesis. But over time of insulin compensatory secretion causes the exhausting of beta cell function, the beta cell postprandial action becomes abnormal, as evidenced by the loss of immediate insulin response to a meal [1,4]. The elevated levels of fasting glucose and insulin, along with an impaired glucose tolerance test are well defined as pre-diabetes, and also defined as the insulin resistance syndrome [1,3].

If any of food intake restriction and physical activity are still missing in the pre-diabetes stage, the T2DM is essentially developed from worsening hyperglycemia comes with the significantly increased fasting glucose and A1C value with peripheral insulin resistance coupled with progressive beta cell dysfunction.

Besides of insulin, people with T2DM have defects in both insulin and glucagon secretion [5-7]. It is also known that type-2 diabetes is associated with elevated levels of glucagon [4,8-10] and defective suppression of glucagon secretion. Moreover, it also had a numerous study described that glucagon suppression is impaired after an oral or intravenous glucose load in pre-diabetes and T2DM [11-18] the increased alpha-cell function in pre-diabetes and T2DM consequent hyper glucagonemia by stimulating hepatic glucose production [4,19] As the regulators to glucagon secretion, the glucagon-like peptides 1 (GLP-1) and glucose-dependent insulinotropic polypeptide (GIP) are most important ones to consider when studying glucagon secretion *in vivo* [4,16,17].

It has been reported that glucose-induced insulin response is greater after an oral load as compared with an IV glucose injection because of the effect of incretins [20]. So a standard mixed meal tolerance test (MMTT) corresponding to a standardized oral glucose tolerance test (OGTT) may yield a more physiological and suitable model to test the impact of insulin and glucagon on postprandial glucose regulation [21]. But a major limitation to the study of the MS in humans, and especially the pathogenesis of these disorders and their related

metabolic abnormalities, is the lack of appropriate animal models with which to rigorously investigate the progression and biological mechanisms of the disease processes. Currently, more studies related to obesity/diabetes research are moving toward using nonhuman primates (NHPs) which have better pathophysiological metabolism similarities to humans [22,23]. Many characteristics of the Cynomolgus monkey (*Macaca fascicularis*) make the species ideal for this purpose [24,25]. But the report of postprandial relationship between glucagon, insulin and glucose in Cynomolgus monkeys is scarce.

In this investigation, we aimed to characterize the glucose, insulin and glucagon responses and indices of insulin resistance during a MMTT in a high fat and high fructose diet-induced Cynomolgus metabolic model of obese pre-diabetes, and overt T2DM in comparison to control.

Materials and Methods

Thirteen monkeys were fed a high-calorie (high fat diet & high fructose) diet to induce the metabolic syndrome diseases models. Five monkeys were fed normal monkey diet and with normal intravenous glucose tolerance test (IVGTT) profile served as age-matched chow-fed controls. The high-calorie fed animals developed varying metabolic alterations and all animals could be stratified into groups of metabolically healthy obese, pre-diabetes, T2DM, and controls according to their hemoglobin A1C% (HbA1C%) values and glycemic status defined from plasma glucose level at 60 minutes in an IVGTT. The detail criteria for stratify animals were report previously [26].

All animals were obtained from an approved vendor (Hainan Jingang Biotechnology, Hainan, China), and housed in a controlled animal facility with a 12:12-h light:dark cycle; reverse-osmosis-purified water was available without restriction to all animals. Enrichment toys and treats were provided on a daily basis.

All eighteen Cynomolgus macaques underwent a MMTT via nasogastric gavage administration. The MMTT procedure in Cynomolgus macaques was conducted as approved by the IACUC of WuXi Apptec (Suzhou, Jiangsu Province, People's Republic of China), which is in compliance with the Animal Welfare Act and guidelines in the Guide for the Care and Use of Laboratory Animals.

Animals were trained for oral gavage administration prior to the test. The MMTT was performed in the morning of the test day under slight sedation with ketamine, after about 14 hours overnight fast. And following the baseline sampling (-30, -5 minutes) the animals have received a test meal via the gavage administration. Ensure plus® was purchased from Abbott company, and used as the test meal in this MMTT. A single dose of Ensure plus® (350 Kcal, 17% carbohydrate, 17% fat, and 26% protein) to each monkey is calculated by 6 mL per kilogram of body weight multiple the individual body weight. Postprandial sampling for glucose, insulin and glucagon assessments collected at 15, 30, 45, 60, 90, 120, 180, 240 minutes after gavage administration.

Blood chemistries of total cholesterol, triglyceride and liver enzymes were determined by using Biochemical analyzer (HITACHI 7180). Glucose was measured by glucose oxidase method, using a Glucose analyzer (EKF Biosen C-line). Insulin ELISA kits (Catalog no.: 10-1113-01 & 10-1113-10) and glucagon ELISA kits (Catalog no.: 10-1271-01) were purchased from Mercodia Developing Diagnostics.

Hemoglobin A1c (HbA1c) was determined by using DCA Vantage analyzer (Siemens). Total body fat composition was determined by using Dual Energy X-ray bone density machine (HOLOGIC Discovery A).

Statistical analysis and calculations

All results are expressed as mean ± SEM (except HOMA-IR & IGI were expressed as mean). AUC value was calculated by using Phoenix WinNonlin software (version 6.2.1, Pharsight, Mountain View, CA), the non-compartmental model (Linear/log trapezoidal, best fit) was used.

Differences between groups and within groups were evaluated using one-way ANOVA followed by Tukey's multiple comparison post tests using Graph Pad statistic software. A p value of less than 0.05 is considered statistically significant.

The insulinogenic index (IGI), as a surrogate marker of beta cell function has largely used in human clinical trials to estimate the level of insulin secretion with a more physiological route of glucose administration and evaluate subjects in different metabolic status. It measured of first-phase insulin response to glucose challenge during the first 30 minutes of the OGTT or MMTT and is computed by following the equation as $IGI = \frac{\delta \text{ insulin (0-30 min)}}{\delta \text{ glucose (0-30 min)}}$. Insulin is measured in micro units per milliliter, whereas glucose is measured in milligrams per deciliter [27].

The HOMA-IR is also a useful method have been used with numbers of human epidemiological studies in quantifying the degree of insulin resistance and beta cell function in human subjects among ethnic groups, as well in comparison of HOMA-IR and beta cell function between or within groups. The equation known as $HOMA-IR = \frac{\text{fasting insulin concentration} \times \text{fasting glucose concentration}}{22.5}$, calculations are based on mmol/L for glucose and mU/mL for insulin [28].

Results

The anthropometries and measured lipid parameters

Anthropometric characteristics and lipids parameters of 18 monkeys in T2DM, pre-diabetes, obese and control groups are presented as Mean ± SEM in Table 1.

| Group | T2DM | Pre diabetes | Obese | Control |
|---------------------------------|-----------------|--------------|--------------|--------------|
| Age (Year) | 14.3 ± 0.33 | 14.4 ± 0.40 | 14.0 ± 0.00 | 12.8 ± 1.28 |
| Body Weight (kg) | 10.5 ± 2.01 | 9.47 ± 0.76 | 10.1 ± 0.63 | 9.02 ± 0.76 |
| Total Body Fat Composition (%) | 46.0 ± 3.68** | 41.1 ± 2.70 | 40.5 ± 1.43* | 27.7 ± 3.56 |
| HbA 1c% | 6.20 ± 0.40**** | 4.80 ± 0.15 | 4.18 ± 0.09 | 4.20 ± 0.17 |
| Fasting Plasma Glucose (mmol/L) | 7.98 ± 1.27** | 5.84 ± 0.328 | 5.26 ± 0.414 | 4.66 ± 0.380 |
| Fasting Plasma Insulin (mU/L) | 47.3 ± 22.7 | 183 ± 108 | 44.3 ± 7.95 | 40.7 ± 16.2 |

| Fasting Plasma Glucose (pmol/L) | 21.2 ± 7.33 | 30.5 ± 5.84 | 23.9 ± 4.55 | 28.6 ± 6.48 |
|---------------------------------|---------------|--------------|-------------|-------------|
| IVGTT Glucose at Baseline | 4.86 ± 0.86* | 3.94 ± 0.22 | 3.47 ± 0.40 | 3.14 ± 0.11 |
| IVGTT Glucose at 60 minutes | 8.29 ± 0.40** | 5.97 ± 0.80 | 4.29 ± 0.58 | 3.71 ± 0.50 |
| TC (mmol/L) | 8.26 ± 4.32 | 9.15 ± 1.93 | 8.50 ± 3.02 | 2.91 ± 0.17 |
| TG (mmol/L) | 0.72 ± 0.25 | 1.89 ± 1.34 | 0.98 ± 0.47 | 0.67 ± 0.19 |
| FFA (mmol/L) | 0.67 ± 0.09 | 0.91 ± 0.15* | 0.53 ± 0.08 | 0.42 ± 0.11 |

Table 1: Anthropometric Characteristics and Lipids Parameters. Data were presented as mean ± SEM, T2DM, pre diabetes and obese groups were compared with control group. *p<0.05, **p<0.01, ****P<0.0001.

The anthropometric data showed animals in T2DM, pre-diabetes and obese groups were slight older, and heavier than the animals in control group, also had a greater percentages of body fat composition than control.

The lipids parameters including total cholesterol, triglyceride and free fatty acid showed the higher levels were found in the T2DM, pre-diabetes and obese groups than control group animals.

HbA1C% and fasting glucose

As glycemic parameters are also shown in Table 1, the higher HbA1c% value in most groups but obese associates with higher fasting glucose and defined the impaired glucose tolerance status based on the plasma glucose level at 60 minutes from previous IVGTT.

T2DM group had a highest HbA1C value than others, which is statistically significant higher than control group (P<0.0001). T2DM group also had a highest fast plasma glucose level than others, which is statistically significant higher than control (p<0.01). Also, the glucose levels at baseline and 60 minutes from previously IVGTT showed the T2DM had the worsened impaired IVGTT profile.

Contrast to T2DM, the control group had the lowest HbA1C value and fast plasma glucose level, also associated with a normal IVGTT profile. The pre-diabetes group showed having a moderate elevated HbA1c value and fast plasma glucose level, and that is as well associated with an impaired IVGTT profile. Unlike the positively association between HbA1C and fast plasma glucose and IVGTT profile have found in other groups, obese group had a mild elevated fast plasma glucose level and an impaired IVGTT profile but associated with the lowest HbA1C% value similar to the control group.

Postprandial plasma glucose appearances in MMTT

The postprandial plasma glucose profile of each groups and calculated AUC_{0-240 min} as mean ± SEM are given in Table 2 and Figure 1.

The glucose profiles during MMTT are different between control and groups of T2DM, pre-diabetes and obese.

Postprandial glucose levels in all groups started rising at 15 min after meal administration, but the time for achieving peak value are differed as earliest at 30 minutes in control compared to 45 minutes in

obese and pre-diabetes. Moreover, there were two peak values with the postprandial glucose appearance have found in T2DM, as the first peak occurred at 30 min, then glucose gradually rose again after a small nadir to the second peak value at 210 min.

| Items | Group | Time (min) | | | | | | | | |
|----------------|--------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|-------------|
| | | 0 | 15 | 30 | 45 | 60 | 90 | 120 | 180 | 240 |
| Insulin (mU/L) | T2DM | 47.3 ± 22.7 | 42.7 ± 19.5 | 91.0 ± 47.6 | 102 ± 61.8 | 108 ± 74.5 | 137 ± 80.7 | 181 ± 112 | 204 ± 125 | 153 ± 119 |
| | Pre diabetes | 183 ± 108 | 165 ± 105 | 232 ± 110 | 238 ± 119 | 299 ± 169 | 330 ± 158 | 254 ± 99.0 | 141 ± 26.8 | 69.7 ± 21.6 |
| | Obese | 44.3 ± 7.95 | 54.4 ± 8.28 | 83.3 ± 13.3 | 90.3 ± 26.1 | 108 ± 29.4 | 88.2 ± 21.5 | 70.7 ± 15.7 | 51.4 ± 11.8 | 61.2 ± 16.4 |
| | Control | 40.7 ± 16.2 | 65.4 ± 14.9 | 77.9 ± 24.3 | 76.7 ± 27.9 | 58.1 ± 14.1 | 35.8 ± 10.0 | 38.6 ± 9.74 | 30.3 ± 4.25 | 25.5 ± 8.01 |
| Glucagon (pM) | T2DM | 21.2 ± 7.33 | 47.6 ± 18.2 | 34.2 ± 6.55 | 39.5 ± 4.47 | 27.6 ± 4.33 | 39.6 ± 8.96 | 33.9 ± 11.4 | 28.5 ± 2.89 | 20.4 ± 1.08 |
| | Pre diabetes | 30.5 ± 5.84 | 59.8 ± 23.8 | 44.9 ± 8.51 | 45.8 ± 11.3 | 45.5 ± 2.87 | 42.6 ± 7.25 | 31.0 ± 3.05 | 32.7 ± 11.0 | 22.5 ± 6.36 |
| | Obese | 23.9 ± 4.55 | 32.1 ± 5.26 | 30.4 ± 7.42 | 50.1 ± 7.46 | 37.3 ± 5.19 | 37.7 ± 9.81 | 26.1 ± 7.65 | 48.8 ± 27.3 | 43.6 ± 18.7 |
| | Control | 28.6 ± 6.48 | 21.1 ± 7.23 | 19.7 ± 6.90 | 26.9 ± 10.2 | 19.0 ± 4.23 | 20.0 ± 3.33 | 19.8 ± 2.57 | 19.2 ± 1.30 | 33.4 ± 10.3 |
| Glucose (mM) | T2DM | 7.98 ± 1.27 | 9.83 ± 1.93 | 11.1 ± 2.54 | 10.8 ± 2.41 | 10.8 ± 1.94 | 11.6 ± 1.74 | 11.9 ± 2.40 | 11.0 ± 2.27 | 7.94 ± 1.47 |
| | Pre diabetes | 5.84 ± 0.33 | 9.01 ± 1.07 | 9.13 ± 1.06 | 9.64 ± 0.87 | 9.49 ± 1.01 | 9.34 ± 1.38 | 8.64 ± 1.37 | 6.60 ± 1.44 | 4.98 ± 1.08 |
| | Obese | 5.26 ± 0.41 | 6.72 ± 0.45 | 6.91 ± 0.45 | 7.17 ± 0.75 | 7.01 ± 0.64 | 6.25 ± 0.51 | 6.17 ± 0.41 | 5.99 ± 0.35 | 6.21 ± 0.62 |
| | Control | 4.66 ± 0.38 | 4.76 ± 0.25 | 5.02 ± 0.17 | 4.72 ± 0.25 | 4.73 ± 0.35 | 4.48 ± 0.33 | 4.37 ± 0.24 | 4.52 ± 0.10 | 4.33 ± 0.22 |

Table 2: Postprandial Plasma Concentration of Insulin, Glucagon and Glucose. Data were presented as mean ± SEM.

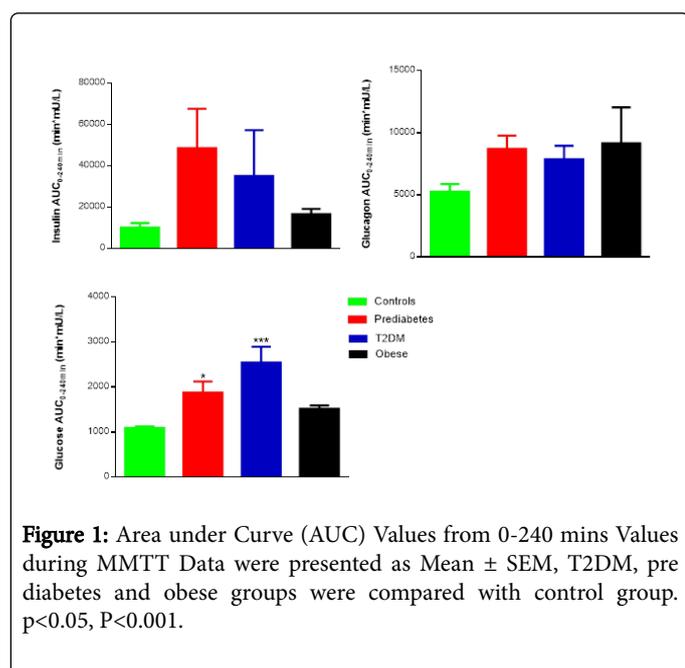


Figure 1: Area under Curve (AUC) Values from 0-240 mins Values during MMTT Data were presented as Mean ± SEM, T2DM, pre diabetes and obese groups were compared with control group. $p < 0.05$, $P < 0.001$.

The times as postprandial glucose has returned to or never fall to the basal level are listed in an order from earliest to latest to never as control at 90 minutes, T2DM and pre-diabetes at 240 minutes, but obese has failed to return.

The shapes of postprandial glucose appearance in responses to MMTT resulted the different $AUC_{0-240 \text{ min}}$ and data showed the T2DM group had the greatest postprandial glucose $AUC_{0-240 \text{ min}}$ $2546 \pm 351 \text{ min} \cdot \text{mM}$ than other groups, followed by the pre-diabetes and obese and the control had the smallest.

Postprandial insulin secretions in MMTT

Mean ± SEM of insulin postprandial concentrations and the calculated $AUC_{0-240 \text{ min}}$ during the MMTT are presented in Table 2 and Figure 1.

Overall, the postprandial insulin secretion of every group has two phases, and the patterns of secretion are different between control and groups of T2DM, pre-diabetes and obese.

The postprandial insulin enhancement in T2DM group was started at 30 min to a value of 91.0 ± 47.6 from baseline value of 47.3 ± 22.7 mU/L as a small initial phase rapidly release of performed insulin, then continued up to the peak value of 204 ± 125 mU/L at 180 minutes as second phase release of synthesized insulin to remove the remained postprandial glucose. The insulin level finally dropped to 153 ± 119 mU/L at 240 minutes which resulted a moderate high insulin $AUC_{0-240 \text{ min}}$ as $35121 \pm 22193 \text{ min} \cdot \text{mU/L}$ than obese and control but pre-diabetes groups.

The increasing of postprandial insulin secretion in pre-diabetes group is similar to T2DM, it increased to 232 ± 110 mU/L at post-meal 30 minutes from basal level of 183 ± 108 mU/L as a small initial phase rapidly release of performed insulin, then went up to the peak value of 330 ± 158 mU/L at 90 minutes as second phase release of more synthesized insulin. But insulin level after which has dramatically declined to 141 ± 26.8 mU/L at 180 minutes and ended with a significantly lower value of 69.7 ± 21.6 mU/L to baseline level at 240 min. Moreover, postprandial insulin $AUC_{0-240 \text{ min}}$ was calculated as $48532 \pm 19143 \text{ min} \cdot \text{mU/L}$, a greatest insulin AUC significantly larger ($P < 0.05$) compared to rest groups of T2DM, obese and control.

In obese group, the insulin secretion has increased at 15 minutes at fed state, and faster than T2DM and pre-diabetes. It is rapidly increased to 83.3 ± 13.3 mU/L at 30 minutes from basal 44.3 ± 7.95 mU/L as a small initial phase release of performed insulin and then reached the peak value of 108 ± 29.4 mU/L at 60 minutes as second phase for more synthesized insulin releasing. At 240 minutes, the insulin level is still at a higher value of 61.2 ± 16.4 mU/L more than

baseline. The $AUC_{0-240 \text{ min}}$ for postprandial insulin was calculated as $16720 \pm 2458 \text{ min} \cdot \text{mU/L}$ lower than groups of T2DM and pre-diabetes but control.

Unlike to other groups, the pattern of postprandial insulin secretion acted in age- matched control group increased immediately at post-meal 15 minutes, also rapidly up to the peak value of $77.9 \pm 24.3 \text{ mU/L}$ at 30 minutes, then gradually declined to $25.5 \pm 8.01 \text{ mU/L}$ lower than baseline at 240 minutes. It produced a smallest $AUC_{0-240 \text{ min}}$ as $10141 \pm 2192 \text{ min} \cdot \text{mU/L}$ for postprandial insulin secretion than rest groups.

Postprandial glucagon secretions in MMTT

Mean \pm SEM of glucagon postprandial concentrations and the calculated $AUC_{0-240 \text{ min}}$ during the MMTT are presented in Table 2 and Figure 1.

The postprandial glucagon secretion increased at 15 minutes in T2DM, it was a 15 minutes earlier than the increasing of insulin secretion and achieved a peak value of $47.6 \pm 18.2 \text{ pM}$ at same time, then slowly declined to $20.4 \pm 1.08 \text{ pM}$, a value similar to baseline at 240 minute. The postprandial glucagon $AUC_{0-240 \text{ min}}$ was calculated as 7863 ± 1089 higher than control group, but lower than pre-diabetes and obese groups.

The postprandial glucagon secretion in pre-diabetes group showed increasing from baseline and reaching to the peak value of $59.8 \pm 23.8 \text{ pM}$ were all occurred at 15 minutes after meal administration. The glucagon level has declined slightly to $42.6 \pm 7.25 \text{ pM}$ thought 90 minutes, then fallen to a value of $22.5 \pm 6.36 \text{ pM}$ lower than baseline at 240 min. The postprandial glucagon $AUC_{0-240 \text{ min}}$ calculation resulted a greater glucagon $AUC_{0-240 \text{ min}}$ as $8699 \pm 1072 \text{ min} \cdot \text{pM}$ higher than groups of T2DM and control but obese.

There were two peak values of postprandial glucagon secretion appeared at 45 and 180 min with $50.1 \pm 7.49 \text{ pM}$ and $48.8 \pm 27.3 \text{ pM}$ respectively have found in obese group during a 240 minutes MMTT. Meanwhile, obese group had a greatest $AUC_{0-240 \text{ min}}$ as $9155 \pm 2883 \text{ min} \cdot \text{pM}$ for postprandial glucagon than control, T2DM and pre-diabetes groups.

The control group had a different pattern as postprandial glucagon secretion compared with other groups. While insulin secretion increasing after meal administration, the glucagon has dropped immediately from baseline value of $28.6 \pm 6.48 \text{ pM}$ with a flat level lower than baseline through to 180 minutes, and then increased to $33.4 \pm 10.3 \text{ pM}$ at 240 min as insulin level has dropped significantly. The smallest $AUC_{0-240 \text{ min}}$ as $5270 \pm 601 \text{ min} \cdot \text{pM}$ for postprandial glucagon was found in control group.

Assessments of IGI and HOMA -IR to monkeys in the groups of T2DM, pre-diabetes, obese and control

IGI and HOMA-IR for the monkeys in each group using data from the MMTT and the results are shown in Table 3.

Overall the IGI numbers are shown in a rank from smallest to largest as 0.78 for T2DM, 0.82 for pre-diabetes, 1.31 for obese and 5.67 for control group. The results indicated that the control group had a largest IGI, significantly higher than other three groups and best beta cell function that also proven with the smallest postprandial glucose $AUC_{0-240 \text{ min}}$.

The HOMA-IR indexes shown in the opposite of order to IGI, as largest the number of 47.5 has found in pre-diabetes group, followed

by a 16.8 in T2DM, 10.3 in obese. The control group had smallest number of 8.43. The results of HOMA-IR well associated with the postprandial glucose disappearance and beta cell function in each group.

| Group | Parameters | |
|--------------|------------|---------|
| | IGI | HOMA-IR |
| T2DM | 0.78 | 16.8 |
| Pre diabetes | 0.82 | 47.5 |
| Obese | 1.31 | 10.3 |
| Control | 5.67 | 8.43 |

Table 3: Values of IGI and HOMA-IR Data were presented as Mean.

Discussion

MMTT, OGTT, IVGTT, Arginine Stimulation Tests and hyperglycemic clamp are common methods have been used to assess insulin sensitivity and pancreatic beta cell function. The MMTT is physiological highly relevant mimicking oral challenges routinely encountered daily; as well as the full incretin hormone effect is able to be tested following the oral ingestion of nutrients [29]. We therefore conducted a MMTT in our animal models to assess glucose, insulin and glucagon although we have not assess other incretin hormones in current investigation.

The current study was undertaken to characterize the response to a MMTT in Cynomolgus monkeys with diet-induced metabolic derangements with focus on glucose, insulin and glucagon responses. The monkeys exhibited various degrees of insulin resistance as observed in human subjects according to the HbA 1C% and glucose characteristics (i.e., either T2DM, pre-diabetes, obese or controls).

Control group obtained the smallest HOMA-IR and biggest IGI which suggests control monkeys are more insulin sensitive than other groups, and have the normal beta cell function which shown with rapidly secreted the adequately efficient insulin in response to postprandial glucose appearance and promotes glucose uptake in more insulin sensitive peripheral tissues to accelerate postprandial glucose disappearance. The glucagon secretion in control during the MMTT also showed as expected as normal physiology which the moderate elevated fasting glucagon concentration stimulated the gluconeogenesis to increase the hepatic glucose production to maintaining the euglycemic circulating range. And glucagon secretion has then suppressed when insulin and incretins hormone secretions dominated in postprandial phase until the glucose back to basal level. The insulin and glucagon actions in control group during MMTT are more resemble to normal physiological condition in healthy human [1]

In contrast, a bigger HOMA-IR and a smallest IGI indexes obtained in T2DM group suggesting of the monkeys of T2DM are more insulin resistance and had dysfunctional pancreatic beta cell. Fasting hyperglycemia and the greatest $AUC_{0-240 \text{ min}}$ of postprandial glucose reflected a very slow rate of postprandial glucose disappearance caused by inadequately insulin secretion and a reduction in glucose uptake by insulin resistance peripheral tissue. The insulin secretion had delayed in response to postprandial glucose appearance and shaped with a small portion of preformed insulin in first phase and big portion of synthesized insulin in second phase appeared the dysfunctional beta

cells are unable to provide adequately insulin to help remove the postprandial glucose flux either mediate validate insulin signaling pathway for glucose utilization. This is very like T2DM human associated with insulin resistance and defective insulin secretion [30-32]. It is also known that T2DM human is associated with elevated levels of glucagon [6,33,34] and defective suppression of glucagon secretion [33-36]. In our MMTT, the failure of glucagon inhibition by insulin in T2DM monkeys has been observed which glucagon secretion has increased an average 60% of fasting concentration over the postprandial time period. Our data is resembled to human.

The pre-diabetes group exhibited a similarity of insulin resistance to T2DM. It had a moderate elevated fast plasma glucose and small IGI, but the remarkable fasting hyperinsulinemia resulted in biggest HOMA-IR index than all other groups. The hyperinsulinemia is an adaptive mechanism that enables the maintenance of normoglycemia in the presence of insulin resistance [3]. The postprandial insulin secretion exerts similar as T2DM, a delayed secretion increasing occurred, but had an early started second phase for synthesized insulin that helped glucose flux and back to baseline. It has been documented by many clinical studies that association between hyperinsulinemia and T2DM, or insulin resistance indicated facilitated beta cell compensational efforts takes place to promoting insulin secretion and biosynthesis and beta cell growth to meet insulin demanding from excess glucose. The characterization of compensational hyperinsulinemia with pre-diabetes monkeys indicates the progression to overt T2DM; it resembled insulin resistance and pre-diabetes in human. We also observed that glucagon level in pre-diabetes accompanied with hyperinsulinemia was highest fasting level than others yet has elevated through most postprandial period of MMTT. Obesity is one of the fundamental factors underlying the current epidemic of T2DM [37-39].

From our data, the obese group had a mild high HOMA-IR than control group and moderate high IGI than T2DM and pre-diabetes groups. The slow postprandial glucose disappears with a slight increasing at 240 minutes, reflect the insulin resistance in peripheral tissue reduced the glucose uptake and utilize. Insulin secretion in response to glucose increase in post-meal period exerted as similar to the pre-diabetes. The glucagon suppression in response to MMTT is also diminished in obese monkeys. The increased postprandial level of glucagon, together with moderately elevated fast plasma insulin and postprandial insulin level are tightly coupled to a reduction of insulin sensitivity in the early stages of glucose deregulation [40]. All the data indicated that obesity is the beginning of pre-diabetes and T2DM.

The MMTT is a physiological relevant stimulus that mimics oral daily meal challenges. We conducted this MMTT in Cynomolgus monkeys to assess glucose, insulin and glucagon responses, and generally our findings are in alignment with human data. But our investigations still have some potential limitations. The first is the number of animals, which was limited due to time (development of the diet-induced changes) and costs. Eighteen monkeys were used and for analyses with high variance such as the glucagon measurements, the power to detect clinically relevant minor differences may not have been sufficient. Another potential limitation is lack of incretion data, which may underlay some of the changes in insulin, glucagon and glucose that were detected. Changes in hormones such as glucagon-like peptides 1 (GLP-1) and glucose-dependent insulin-tropic polypeptide (GIP) are clearly needed to be investigated in a NHP model.

Conclusion

In summary, although the number of subjects studied in this investigation was small, and it lacked with data of more incretins though, we believe it confirmed that the postprandial insulin, glucagon and glucose levels in different metabolic syndrome group respond differently to MMTT and the data from this investigation truly reflect the insulin sensitivity and beta cell function for the monkeys in each classified group and different groups response characteristics are very similar to human [9,12,41,42]. It is well demonstrated that cynomolgus monkeys are a good model to study metabolic syndrome and have translational value to human disease investigation.

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