ORIGINAL ARTICLE



Autoantibodies against β_1 -adrenoceptor induce blood glucose enhancement and insulin insufficient via T lymphocytes

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Abstract Diabetes mellitus is a chronic metabolic disorder with a high morbidity and mortality, but its pathogenesis is not fully understood. An increasing amount of evidence indicates that an immune mechanism plays an important role in the pathogenesis of diabetes. We demonstrated previously that the long-term presence of autoantibodies against the second extracellular loop of the β_1 -adrenoceptor (β_1 -AA) could change the ratio of peripheral CD4+T/CD8+T in rats, which was accompanied by lymphocytes infiltration in the rat heart, liver, and kidneys. To investigate whether β_1 -AA is involved in the pathogenesis of diabetes, BALB/c or nude mice were passively immunized with monoclonal antibodies against β₁-AR (β_1 -AR mAb). Compared with vehicle control mice, β_1 -AA-positive *BALB/c* mice exhibited significantly increased blood glucose (P < 0.01) and increased fasting insulin (P < 0.05). However, the same changes did not occur in the nude mice. And altered islet morphology was found at week 28 in β_1 -AA immunization group compared with vehicle control. The basal insulin level of NIT-1 βcells was decreased markedly (P < 0.01), and the lactate dehydrogenase level was increased (P < 0.01) after the

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administration of conditioned media from T lymphocytes that had been treated with $\beta_1\text{-}AA$ alone. However, these effects were reversed by treatment with metoprolol or peptides of the second extracellular loop of $\beta_1\text{-}adrenoceptor$ ($\beta_1\text{-}AR\text{-}EC_{II}$). These results suggest that $\beta_1\text{-}AA$ could induce hyperglycemia in both rats and mice, and also impair insulin secretion and change islet structure. T lymphocytes may play a key role in the pathogenesis of these changes in the islets.

Keywords Autoantibody against β_1 -adrenoceptors \cdot Diabetes \cdot Hyperglycemia \cdot T lymphocytes \cdot Insulin secretion

Abbreviations

Abbreviations	
DM	Diabetes mellitus
β_1 -AR	B ₁ -adrenoceptor
β_1 -AR-EC _{II}	The second extracellular loop of β_1 -AR
β_1 -AR mAb	Monoclonal antibody against β ₁ -AR-EC _{II}
β_1 -AA	Autoantibodies against the second
	extracellular loop of β_1 -AR
T1DM	Type 1 diabetes mellitus
T2DM	Type 2 diabetes mellitus
DCM	Dilated cardiomyopathy
IPGTT	Intraperitoneal glucose tolerance test
AUC	Area under the curve
LDH	Lactate dehydrogenase
IDF	International Diabetes Federation
Treg	Regulator T cells
Teff	Effector T cells
CTfh	Circulating follicular helper T cells
Th	Helper T cells
GK rat	Goto-Kakizaki rat
HPLC	High-performance liquid chromatography
SC	Subcutaneously



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ELISA Enzyme-linked immunosorbent assay

OD Optical density

IgG Immunoglobulin G fractions

ConA Concanavalin A

Introduction

Diabetes mellitus is known as a chronic metabolic disease that is characterized by hyperglycemia and insulin resistance. According to the new data from the International Diabetes Federation (IDF), 387 million people are living with diabetes worldwide, and China has the largest number of people suffering from diabetes [1]. Therefore, diabetes causes a heavy burden in both China and worldwide. Diabetes is a complex syndrome that is the result of the combination of genetic factors and environmental stimuli [2, 3]; however, its pathogenesis is still not fully understood.

It is widely accepted that type 1 diabetes mellitus (T1DM) is an autoimmune disease that is characterized by the destruction of pancreatic β-cells, which causes an absolute deficiency of insulin secretion. An increasing number of notable discoveries in recent years have suggested that the immune system also plays an important role in the pathogenesis of type 2 diabetes mellitus (T2DM) [4– 7]. A recent study reported that both the percentage of peripheral CD4+T cells and the ratio of CD4+T to CD8+T cells in T2DM patients were increased significantly, as well as the imbalance between regulatory T cells (Treg) and effector T cells (Teff). [8] Another study reported that proportion of circulating follicular helper T cells (CTfh) in peripheral CD4+T cells was increased significantly in T2DM patients than in controls [9]. A link between obesity, T2DM, and inflammation was established in the 1990s [10]. Subsequently, an increasing number of studies have demonstrated that as the resources of many different kinds of proinflammatory cytokines, T cells and immune system, both play a role in T2DM [11, 12]. Consistent with this, data that analyzed patients with prediabetes and T2DM showed that immunity-related biomarkers were altered dynamically during the progression of T2DM [13]. Similar immune changes also appear in animal models of diabetes; non-obese type 2 diabetes rats (Goto-Kakizaki rat) show higher levels of natural IgM and T cell ratios with elevated helper T(Th)2 cells compared with Wister rats. T cell ratios with elevated Th2 cells were observed in diabetic GK rats [14]. Taken together, these data suggest that T lymphocytes and the immune system contribute to the pathogenesis of both T1DM and T2DM.

Catecholamines can interfere with the function of T lymphocytes by activating β -adrenoceptors (β -ARs) [15, 16]. Furthermore, Yu et al. [17] demonstrated that β_1 -AR was expressed on the surface of T lymphocytes. Autoantibodies against the second extracellular loop of β₁adrenoceptors (β₁-AAs; 100 % sequence identity between humans and rats) were first reported in 1987, and β_1 -AA acts similarly with agonist of β_1 -AR [18, 19]. We demonstrated previously that rats that had undergone longterm active immunization with the second extracellular loop of β_1 -AR (β_1 -AR-EC_{II}) produced β_1 -AA and exhibited an elevated CD4+/CD8+T cells ratio, as well as lymphocyte infiltration into the heart, kidneys, and liver. [20] Taken together with observations that β_1 -AA enhanced the proliferation of T lymphocytes, this suggests that β_1 -AAs are an autoimmune product that participates in changes in multiple organs, thereby playing a key role in immunity in diabetes [21]. The aim of the current study was to investigate whether β_1 -AA is associated with diabetes, and whether T lymphocytes are implicated in this process.

Materials and methods

Animals

All animal experiments were performed in accordance with the Guide for the Care and Use of Laboratory Animals protocol, published by the Ministry of the People's Republic of China (issued June 3, 2004) and were approved by the Institutional Committee on Animal Care of Capital Medical University. All *BALB/c* mice, *BALB/c* nude mice, and *Wistar* rats (180–220 g) used in the present study were obtained from the Animal Center of Capital Medical University. The mice were housed in pathogen-free conditions at 20–26 °C and were exposed to the conditions with 12/12-h light–dark periods daily. The mice and rats were fed rat chow and water ad libitum. The food was free from insulin and other oral glucose-lowering drugs.

Peptide synthesis and active immunization

The peptide β_1 -AR-EC_{II} was synthesized as described previously using the sequence from the second extracellular loop of β_1 -AR (197–223 amino acid; H-W-W-R-A-E-S-D-E-A-R-R-C-Y-N-D-P-K-C-C-D-F-V-T-N-R-A), which shares 100 % homology between humans and mice [21]. Analytical high-performance liquid chromatography (HPLC) determined that the peptide preparations were 98 % pure. This work was performed by China Peptides Co. Ltd (Shanghai, China).



Eight-week-old healthy Wistar rats (weighing 180–200 g) that were sera-negative for β_1 -AA were divided randomly into two groups. Rats in the vehicle group (n = 15) were immunized with a mixture of normal saline and immunoadjuvant, whereas those in the β_1 -AA group (n = 15) were immunized with a mixture of artificial synthetic β_1 -AR-EC_{II} and immunoadjuvant. β_1 -AR-EC_{II} was dissolved in Na₂CO₃ solution (pH 11.0), mixed with the same volume of complete Freund's adjuvant (Sigma-Aldrich, St. Louis, MO, USA), and then injected subcutaneously (SC) at a concentration of 0.4 µg/g posteriorly along the back at multiple sites (initial immunization). One week after the initial immunization, a booster SC injection of a mixture of an equal volume of peptide solution and incomplete Freund's adjuvant (Sigma-Aldrich) administered to one posterior site. A booster immunization was then given every 2 weeks subsequently. Specifically, rats in the vehicle group were injected SC with 1 mL saline mixed with 1 mL incomplete Freund's adjuvant as described above. Blood samples were collected 1 day before each booster injection.

Enzyme-linked immunosorbent assay (ELISA)

ELISAs were used to detect autoantibodies against β_1 -AR-EC_{II} in the sera of animals before and after immunization with β_1 -AR-EC_{II} or β_1 -AR mAb or vehicle, and the results were expressed as optical density (OD) units according to the methods published previously [22]. The antibody titer was also calculated according to the ratio of the OD values of positive and negative (P and N) controls as follows: ([Specimen OD – blank control OD]/[negative control OD – blank control OD]) [23].

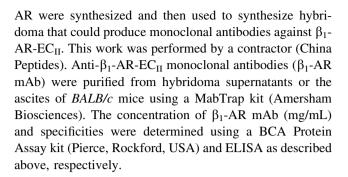
Sera positive or negative for β_1 -AA were defined as samples with P/N \geq 2.1 and P/N \leq 1.5, respectively. The supernatants were then divided into small aliquots and stored for future use.

Preparation of immunoglobulin G

Immunoglobulin G fractions (IgG) were prepared from the β_1 -AA-positive sera of active immunized *Wistar* rats or the ascites of *BALB/c* mice using MabTrap Kit (Amersham Bioscience, Uppsala, Sweden). The concentrations (mg/mL) and specificities of the purified IgGs were determined using a bicinchoninic acid (BCA) protein assay kit (Pierce, Rockford, USA) and ELISA as described above, respectively.

Preparation of β₁-AR monoclonal antibodies

Long and short peptides corresponding to amino acids 197-223 of the second extracellular loop of the human β_1 -



Passive immunization

Passive immunization rat model. The IgG was purified from the β_1 -AA-positive sera of active immunized rats. Wistar rats in the β_1 -AA group (n=32) were injected with the purified IgG (0.7 µg/g) via the vena caudalis of rats every 2 weeks for 40 weeks, and four rats were died in the process of immunization. Rats in the vehicle group (n=28) were injected with β_1 -AA-negative IgG via the same method. Blood samples were collected via the vena caudalis at weeks 0, 12, 24, and 36 for blood glucose measurements.

Passive immunization of BALB/c mice and BALB/c nude mice. BALB/c mice (n = 18) were injected intraperitoneally (IP) with β_1 -AR mAb (5 μ g/g) every 2 weeks for 12 weeks; mice in the vehicle group (n = 18) were injected IP with the same volume of saline. BALB/c nude mice were administered as the same method of BALB/c mice, n = 18 each group.

Intraperitoneal glucose tolerance test

Mice were fasted for 6 h before an IP glucose tolerance test (IGTT) was performed. An IP glucose solution (20 % glucose; 1 g glucose/kg bodyweight) was administered, and the blood glucose was monitored after 30, 60, and 120 min. The area under the curve (AUC) was determined by calculating the sum of rectangular area between each time point.

Analysis of glucose, insulin, and glucagon

Mice and rats were fasted for at least 12 h but were allowed free access to water. Blood glucose levels were measured using an ACCU-CHEK active glucometer (Roche, Mannheim, Germany). Plasma insulin and glucagon were measured at 4-week intervals using an iodine [125I] insulin radioimmunoassay kit (F01PZB; Beijing North Institute of Biological Technology, China) or an iodine [125I] glucagon radioimmunoassay kit (F03PJA: Beijing North Institute of Biological Technology, China). Supernatant insulin of



NIT-1 cells was measured using the same iodine [125I] insulin radioimmunoassay kit.

Histopathological examination

Rats were killed after 24, 28, 32, and 36 weeks of passive immunization, and the pancreas was harvested, fixed in 10% formaldehyde, and embedded in paraffin. The specimens were processed into 4- μ m-thick paraffin sections, and the relative islet area was determined based on the method reported previously [24] after staining with hematoxylin and eosin (H&E).

Cell lines and cell culture

NIT-1 cells, a pancreatic beta-cell line isolated from *NOD/* Lt mice, were purchased from Tongji Medical School, Huazhong University of Science and Technology, Wuhan, China [25]. They were cultured in RPMI1640 (Hyclone, Utah, USA) medium supplemented with 10 % FBS, 100 μg/mL streptomycin, and 100 U/mL penicillin [26]. T lymphocyte suspensions were prepared from the spleen of BALB/c mice as described previously [21]. T lymphocytes were incubated at 37 °C and 5 % CO2 for 24 h, and were then activated with 5 mg/L Concanavalin A (ConA) (Sigma-Aldrich). After activation, the T cells were seeded in 96-well plates and treated with saline (vehicle group), 0.1 μ mol/L isoproterenol (positive control), 0.1 μ mol/L β_1 -AA (β_1 -AA group), 0.1 μ mol/L β_1 -AA + 3 μ mol/L β_1 -AR-EC_{II}, or 0.1 μ mol/L β_1 -AA + 1 μ mol/L metoprolol for 48 h. The supernatant was collected and added to NIT-1 cells for 6 or 24 or 48 h for measurement of insulin and LDH.

Analysis of lactate dehydrogenase (LDH) levels

NIT-1 cells (1×10^4 cells/well) were seeded in 96-well plate and cultured for 24 h. The media were then discarded and replaced with 100 μ L of conditioned media collected as described above for another 6 or 12 or 24 or 48 h. LDH levels were then measured using commercially available kits (Beyotime, Shanghai, China).

Statistical analysis

The results are presented as mean \pm standard deviation (S.D), and Student's t test was used to compare two independent sample means; one-way ANOVA was used to compare the means of more than two samples. All statistical analyses were performed using SPSS 13.0, and $P \le 0.05$ was considered to indicate significance (*). Nonsignificant differences are noted in the figures.

Results

β_1 -AA induced hyperglycemia in rats

The β_1 -AA-positive rat model was generated using passive immunization, which increased the blood glucose of the immunized rats gradually. After week 24 of immunization, the blood glucose of rats in the β_1 -AA group was significantly higher than that of rats in the vehicle group (24.6 \pm 7.41 vs. 6.7 \pm 0.30 mmol/L; P < 0.01) and reached a peak level (Fig. 1). The blood glucose level remained higher than the vehicle group at week 36 after immunization, suggesting that the long-term presence of β_1 -AA increased blood glucose.

β₁-AA increased blood glucose levels and impaired pancreatic function in vivo

The results obtained from the β_1 -AA passive immunization rat model suggested that β_1 -AA increased blood glucose. To investigate the mechanism behind this change, a β_1 -AR mAb passive immunization BALB/c mouse model was generated (supplementary information). An IP GTT was performed on mice in both the β₁-AR mAb and vehicle groups every 2 weeks from the beginning of passive immunization. There was no significant difference in the blood glucose levels of the two groups before the start of the experiment or after 4 weeks of immunization (Fig. 2a). However, GTTs performed after 8 and 12 weeks demonstrated that the blood glucose was increased in the β_1 -AR mAb group at 30 and 60 min compared with the vehicle group (week 30 min: 13.47 ± 3.85 8, $8.3 \pm 1.39 \text{ mmol/L}$, respectively [*P* < 0.01]; 60 min: 10.43 ± 1.15 vs. 8 ± 2.03 mmol/L [P < 0.05]; week 12, 30 min: 9.83 ± 0.63 vs. 8.16 ± 1.32 mmol/L, respectively

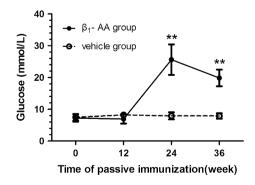
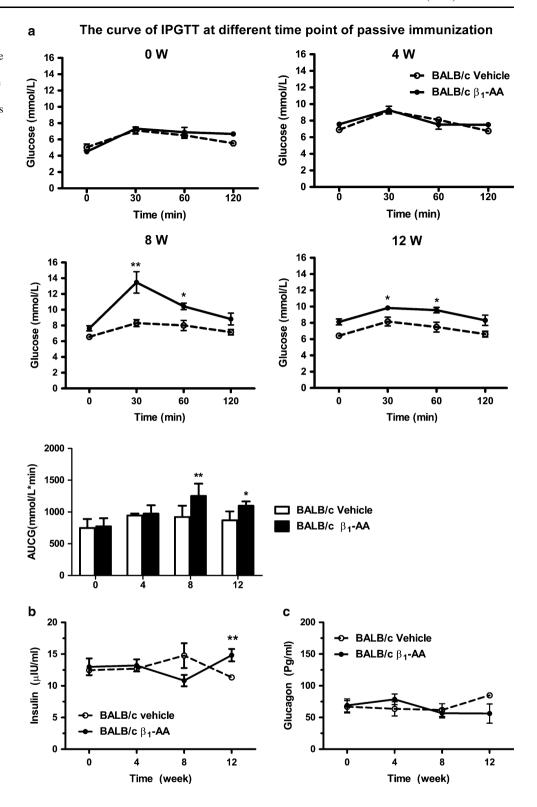


Fig. 1 Blood glucose levels in the sera of rats after $β_1$ -AA passive immunization. Vehicle group (n=32): rats were immunized with IgG from $β_1$ -AA-negative sera; $β_1$ -AA group (n=28): rats passively immunized with $β_1$ -AA. Sera sample was collected and measured at weeks 0, 12, 24, 36 during the process of immunization. Data are presented as mean ± SD of three independent experiments. **P < 0.01 $β_1$ -AA group versus vehicle group at the same time point



Fig. 2 Changes in glucose, insulin, and glucagon in the process of BALB/c mice passive immunization. a IP GTT at different time points of passive immunization. *P < 0.05, ** $P < 0.01 \beta_1$ -AA group versus vehicle group at the same time point. Fasting insulin (b) and glucagon (c) levels in the passive immunization. $^*P < 0.01 \; \beta_1$ -AA *BALB/c* group versus vehicle BALB/c group at the same time point. Data are presented as mean \pm SD of three independent experiments. n = 18/group



[P < 0.05]; 60 min, 9.57 \pm 0.81 vs. 7.48 \pm 1.50 mmol/L [P < 0.05]). The fasting insulin level was also measured in both groups. No significant differences were observed between during the first 8 weeks. However, fasting insulin was increased significantly in the β_1 -AR mAb group

compared with the vehicle group on week 12 after immunization (14.83 \pm 2.38 vs. 11.33 \pm 0.75 μ U/mL, respectively; P < 0.05; Fig. 2b). In contrast, there was no difference in glucagon between the β_1 -AR mAb and vehicle groups (Fig. 2c).



β_1 -AA induced an irregular structure in pancreatic islets

The pancreases were harvested from passively immunized *Wistar* rats and analyzed using H&E staining. An irregular structure was observed in the β_1 -AR mAb group on week 28 (Fig. 3). Specifically, the islets exhibited an abnormal morphology, including unclear edges, a decreased area, and pancreatic acinar invasive into the islet. T lymphocyte infiltration was also observed, suggesting that β_1 -AA could disrupt the structure of the pancreatic islet and that T cells might be involved in this process.

T lymphocytes mediated the β_1 -AA-induced hyperglycemia

We next investigated the role of T cells in β_1 -AA-induced hyperglycemia. To test this, in vitro T lymphocytes were treated with β_1 -AA alone or β_1 -AA+ metoprolol for 48 h as described in method above. The supernatants were then harvested and used to treat NIT-1 cells for 6, 24, or 48 h. Insulin levels were decreased markedly in NIT-1 cells treated with β₁-AA-stimulated T cell supernatant compared with those treated with vehicle-treated supernatant (6 h, 55.82 ± 3.23 vs. 79.23 ± 8.82 µU/mL, respectively [P < 0.01]; 24 h, 111.45 \pm 3.91 vs. 180.58 \pm 19.05 μ U/mL [P < 0.01]; 48 h, 132.26 ± 1.64 vs. 175.52 ± 15.71 μ U/ mL [P < 0.01]; Fig. 4a-c). Metoprolol partially counteracted the effects of β_1 -AA on reducing insulin at 24 and 48 h, but not 6 h (P < 0.01). Furthermore, LDH assays demonstrated that incubation with T cell supernatant increased LDH levels of β₁-AA group in 6, 24, and 48 h compared with vehicle group. (6 h, 808.37 ± 138.46 vs.

 $468.09 \pm 32.82 \text{ U/L}$, respectively [P < 0.01]: 24 h. 1528.67 ± 33.82 vs. 1426.67 ± 39.41 U/L, [P < 0.01]; 1641.04 ± 54.91 $1511.38 \pm 85.56 \text{ U/L}$ 48 h, vs. [P < 0.05]; Fig. 5a–c) And the LDH level of β_1 -AA group in 24 and 48 h was apparently higher than 6 h. However, the presence of the β_1 -AR blocker metoprolol or β_1 -AR-EC_{II} suppressed the increasing LDH levels compared with β_1 -AA alone at 6 and 48 h, but not 24 h (6 h, 808.37 \pm 138.46 vs. β_1 -AA + β_1 -AR-EC_{II} group 914.55 \pm 32.25 U/ L, [P < 0.05]; 48 h, 1641.04 ± 54.91 vs. β_1 -AA + β_1 -AR-EC_{II} group 1527.04 \pm 47.18 U/L, [P < 0.01], 1641.04 \pm 54.91 vs. β_1 -AA + metoprolol group 1432.99 \pm 27.91 U/L, [P < 0.01]; Fig. 5a, c).

β_1 -AA did not cause glucose imbalance and pancreatic dysfunction in nude mice

The in vitro results demonstrated that T cells might contribute to the β_1 -AA-induced increased glucose levels. Nude mice lack the thymus and so are often used in immune-related research. Therefore, we generated a β_1 -AR mAb passive immunization model in BALB/c nude mice and measured IP GTT, and fasting insulin and glucagon levels. There were no significant differences in blood glucose, fasting insulin, or glucagon levels between the β_1 -AR mAb and vehicle groups at any time after immunization (Fig. 6a-c). These results demonstrated that the increased blood glucose levels and pancreatic islet dysfunction observed in β₁-AR mAb passively immunized BALB/c mice did not occur in nude mice. This suggests that T lymphocytes might play a role in the glucose imbalance and pancreatic islet injury induced by β_1 -AA.

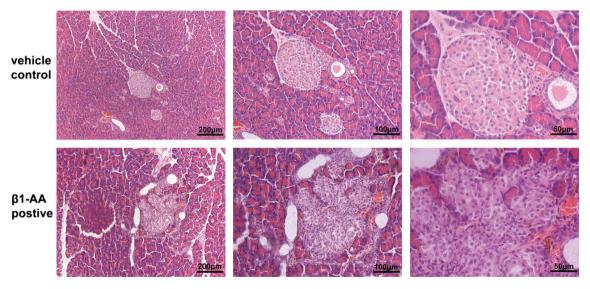


Fig. 3 Morphology change in pancreatic islet. H&E staining of pancreatic islet on week 28 was measured by optical microscope. An irregular islet structure and decreased islet area were observed in the

 β_1 -AA group compared with vehicle. *Scale bar* of 100 times, 200 times, 400 times magnification = 20, 10, 5 μ m, respectively



Fig. 4 Insulin level of NIT-1 cells treated by supernatant of T cells stimulated by $β_1$ -AA or $β_1$ -AA+ metoprolol. Insulin level of NIT-1 cells after the treatment of T cells supernatant for 6 h (**a**), 24 h (**b**), 48 h (**c**).**P < 0.01 versus vehicle group; **#P < 0.01 versus $β_1$ -AA group. *MET* metoprolol. n = 6 in the 6 h group, and n = 4 in both the 24 and 48 h groups. Data are presented as mean ± SD of three independent experiments

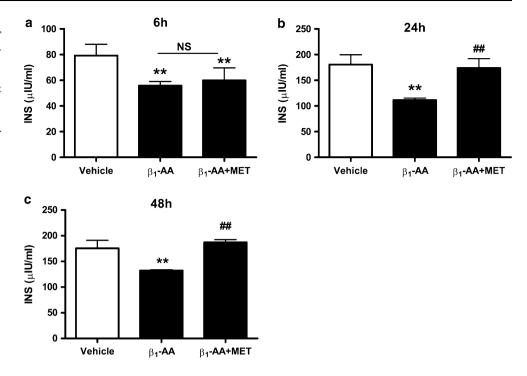


Fig. 5 LDH level of NIT-1 cells treated by supernatant of T cells stimulated by $β_1$ -AA or $β_1$ -AA+ metoprolol or $β_1$ -AA+ $β_1$ -AR-EC_{II}. LDH level of NIT-1 cells after the treatment of T cells supernatant for 6 h (**a**), 24 h (**b**), 48 h (**c**) **P < 0.01 versus vehicle group; *#P < 0.05 versus $β_1$ -AA group; *#P < 0.01 versus $β_1$ -AA group (n = 6/group). Data are presented as mean ± SD of three independent experiments

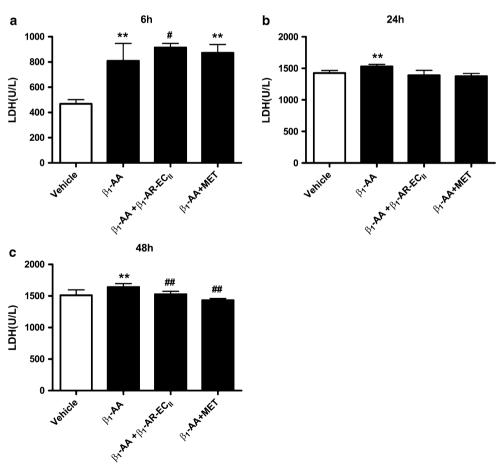
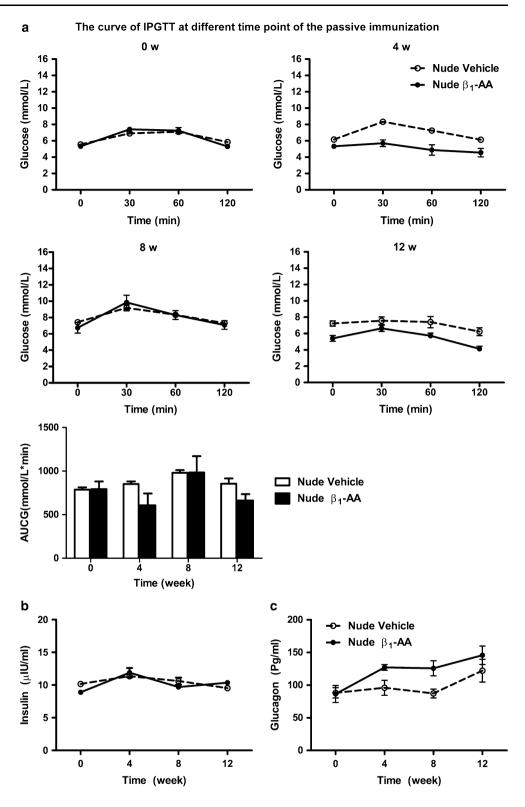




Fig. 6 Changes in glucose, insulin, and glucagon levels during the passive immunization of nude BALB/c mice. a IP GTT curve after different weeks of passive immunization in nude mice. There were no significant difference between the β_1 -AA group and vehicle groups at any time points (n = 18/group). b No significant differences in insulin levels in BALB/c nude mice between the vehicle and immunized groups (n = 18/group). c No significant differences in glucagon levels in BALB/c nude mice between the vehicle and immunized groups (n = 18/group). Data are presented as mean \pm SD of three independent experiments



Discussion

Diabetes is a serious chronic metabolic disease. In addition to the disease itself, the complications caused by diabetes further increase the mortality and morbidity rates. Over time, diabetes can damage the heart, blood vessels, eyes, kidneys, and nerves. However, the mechanisms underlying diabetes have not yet been elucidated fully. The current study introduced a novel mechanism behind diabetes that is caused by autoantibodies against β_1 -AA. Most studies into



 $\beta_1\text{-}AA$ have focused on cardiovascular diseases, including heart failure, idiopathic cardiomyopathy, and arrhythmia [18, 27–29]. In addition, recent studies reported that $\beta_1\text{-}AA$ could induce injury in the kidneys and heart [20, 30]. These results suggest that the impairments caused by $\beta_1\text{-}AA$ might not being limited to the cardiovascular system, but are instead more likely affect multiple systems. The current findings of increased blood glucose levels and pancreatic injury are consistent with this viewpoint.

The current study demonstrated that passive immunization with β₁-AA increased blood glucose in both rats and mice. In the passive immunization rat model, a time-dependent increase in blood glucose levels was observed during the immunization period, and levels peaked at week 24. But in the β_1 -AR mAb passive immunization *BALB/c* mice model, increased blood glucose was observed on week 8, which is earlier than the week-24 increase observed in the β_1 -AA-positive IgG immunization rats model. It is possible that β₁-AA might affect glucose homeostasis before diabetes diagnosis. Importantly, impaired glucose tolerance (IGT) is the stage before diabetes, and individuals with IGT or IFG are at a high risk of progressing to T2DM [31]. Taken together, these observations suggest that IGT might occur before β₁-AA-induced increased blood glucose levels. Therefore, we used IP GTTs to examine glucose tolerance and glucose levels. Furthermore, fasting insulin levels were similar between groups on week 4, but a trend toward a reduced insulin level was observed in the β_1 -AA group at week 8. On week 12, insulin levels were increased significantly in the β_1 -AA group compared with the vehicle group. This is consistent with the variations in insulin levels observed in diabetic patients [32], which suggests that β_1 -AA could induce insulin compensation in mice.

IgG purified from the sera of β_1 -AA-positive rats was used to generate a passive immunization rat model. The IgGs used included specific antibodies against the β_1 -AR-EC $_{II}$ as well as other IgGs; therefore, other kinds of antibodies might have interfered with the effects of β_1 -AA. Therefore, we synthesized β_1 -AR mAb, a monoclonal antibody from mice that eliminates nonspecific IgGs and other effects.

In a β_1 -AR-EC_{II} long-term active immunization model, Zuo reported that the CD4+/CD8+T cells' ratio was elevated compared with vehicle control [20]. Another previstudy revealed that β_1 -AA from cardiomyopathy (DCM) patients enhanced the proliferation of T lymphocytes [21]. This suggests that changes in the T lymphocyte population might accompany the presence of β_1 -AA. The current observations of pancreatic morphology also support this hypothesis, since lymphocyte infiltration in the pancreas was also observed during β_1 -AA immunization. In addition to the important role of T lymphocytes in the pathogenesis of diabetes, we also assessed whether T lymphocytes play a role in β_1 -AA-induced high blood glucose levels using a β_1 -AA passive immunization model in nude mice. There were no differences in blood glucose, insulin, or glucagon levels between the β_1 -AA and vehicle groups, suggesting that T lymphocytes play an important role in β₁-AA-induced pancreatic dysfunction. However, T lymphocytes are not the only reason for β_1 -AA-induced elevated blood glucose levels. According to our observations in NIT-1 cells in vitro, increased LDH and decreased insulin levels could be partially counteracted by metoprolol or β_1 -AR-EC_{II}, suggesting that the effects of β_1 -AA on β cell dysfunction were exerted either via β_1 -ARs on T cells or directly actions on β -cells. The decrease in insulin level in 24 and 48 h was severe than 6 h, and the decrease could be counteracted by β -blocker metoprolol, the probable reason of this phenomenon is the long-term effect of β_1 -AA. As far as we know, the injury caused by β_1 -AA in various kinds of cardiac disease is a long time consequence [20, 33–35]. With the stimulation prolonged, the injury becomes more severe; thus, the protective effect of metoprolol seems more obviously. The increase in LDH in 24-h group did not counteract by metoprolol or β_1 -AR-EC_{II}. As far as we know, there was very rare research about $\beta_1\text{-}AA$ and metabolic disease including diabetes. The only two articles in Chinese were focusing on the presence of β_1 -AA in type 2 diabetes with hypertension [36, 37]. Taken together, the current study, for the first time, revealed that long-term exposure to β_1 -AA increased blood glucose levels and impaired insulin secretion. Furthermore, T lymphocytes were implicated in this process, but T cells may not be the only element participant in it.

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