Effect of PPARα’s over Expression on the Type 2 Diabetic Rat Cardiomyocytes

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Abstract: To investigate the over expression of PPARα and trimetazidine in the apoptosis of rat cardiomyocyte with type 2 diabetes. All rats were divided into 3 groups: Group N (control group), Group D (type 2 diabetes group); Group T (type 2 diabetes which treated by trimetazidine). After 16 weeks breeding, apoptosis of each group were detected by TUNEL, ultrastructure were detected by electronic microscope, immunohistochemistry staining and western blotting were used to detected each group’s cardiomyocyte about PPARα and NF-κB in the expression of each protein level. The number of apoptosis was increased in Group D and the ultra structure was changed, the expression of PPARα in protein level was decreased also the NF-κB (p<0.05). After treated by trimetazidine in type 2 diabetic rat, the apoptosis of cardiomyocytes was decreased distinctly (p<0.05), the change of ultra structure was improved, the expression of PPARα and NF-κB were all decreased in protein level (p<0.05). The over expression of PPARα might be involved in the regulation of the apoptosis in cardiomyocyte and the ultrastructural change of diabetic cardiomyopathy, suggesting it may participate in the development of diabetic cardiomyopathy and treated by trimetazidine may reverse it.

Key words: Apoptosis, diabetic cardiomyopathy, peroxisome proliferator activated receptor α, trimetazidine, protein

INTRODUCTION

Cardiomyocyte apoptosis in diabetes is one of the important events in diabetic heart failure. This process is regulated by a series of gene. PPARα is one of the member of nuclear receptor superfamily, it can interact with its upper reaches of the specificity of certain DNA response element (Peroxisome Proliferators Responsive Element, PPRE), activate nuclear transcription control mechanisms and control of gene expression, play a key role in the regular of the oxidative metabolism of fatty acids and cell proliferation, apoptosis (Lefebvre et al., 2006). Young et al. (2001) showed that the PPARα agonist can activated pressure overload cardiomyopathy and induced cardiomyocyte apoptosis resulting in mice heart failure. Dyntar et al. (2001) find that PPARα expression level increase in Streptozotocin (STZ)-induced diabetic rat myocardium which resulting in the changes of key enzyme in glucose and lipid metabolism and induced cardiac cell apoptosis. Some researchers pointed out the mice which over expression the PPARα (Myosin Heavy Chain (MHC)-PPAR mice) have the similar cardiac energy metabolism disorder and cardiac cell damage in diabetic rat (Dyntar et al., 2001; Finck et al., 2003). These researches have proved that over expression of PPARα may lead to cardiac cell apoptosis and even heart failure in diabetic model but the mechanism is not clearly. Based on these, researchers chose the type 2 diabetic rats which induced by low dose STZ and high-fat diet. And treated with trimetazidine, the 3-Acyl-CoA S-transferase inhibitor which has been widely used in coronary heart disease, myocardial ischemia and other heart disease, it can reverse the cell damage which induced by the myocardial energy metabolism disorder and prevent the occurrence of advanced cardiovascular events effectively (Aussedat et al., 2006; Tabbt-Arnetti et al., 2003). And detected the expression of PPARα, NF-κB, ultra structure

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and apoptosis of each group's cardiomyocyte, aimed to reveal the connection and mechanism between the over-expression of PPARα and the cardiomyocyte apoptosis in diabetic cardiomyopathy.

MATERIALS AND METHODS

Experimental animal groups and modeling: SD male rats 30 (from the Third Military Medical University Animal Center). Weighing about 250 g, they were divided into 3 groups by the random number law: normal group (fed basis food, 10, N group), diabetes (high fat diet feeding and low dose STZ, 10, Group D) and drug intervention group (low dose STZ, high fat diet and treated by Trimetazidine, 10, T group), the basis food of protein, fat, carbohydrates were (w/w) 21.9, 15.5, 62.6%; high-fat diet of protein, fat, carbohydrates were (w/w) 11.3, 43.6, 45.1%. Intraperitoneal injection, high fat diet feeding 4 weeks post-test blood glucose, blood glucose 16.7 mmol L⁻¹ is the judgment for type 2 diabetes and continued high fat diet feeding 8 weeks (Chen et al., 2004), drug treat group in a model of diabetic rats fed high-fat diet and feed trimetazidine 7.5 mg day⁻¹ (add in the food) 8 weeks (Tabbi-Annemi et al., 2003).

Specimens collected and stored: After 16 weeks feeding, Group D and N rats did high insulin-normal glucose clamp (EICT) to detect the IR and then, on the condition of asepsis, researchers took the cardiac muscle, use silver paper parcels the heart tissue and keep it in refrigerators at -80°C, three groups detect the cardiomyocyte apoptosis rate by TUNEL at the same time and electron microscope was used to detect the change of ultra structural situation.

High insulin-normal glucose clamp (EICT): The rats were fasted for 12-14 h overnight before the clamp study began the next morning. Rats were anesthetized by the intraperitoneal injection of amobarbital sodium (25 mg kg⁻¹) which individualized to each subject's body weight. Rats were punctured in the right jugular vein for infusion of glucose and insulin (using separate lines) and in the left carotid artery for blood sampling. Two digital syringe pumps jointed by a Y connector to the jugular catheter which stored the glucose and insulin solutions each.

Insulin (Humulin R, Lilly, France SAS) was infused at a rate of 1.67 mU/kg/min through the jugular vein catheter from 0-120 min. Glucose concentrations were clamped at euglycemic levels (5.0±0.5 mmol L⁻¹) by a variable rate infusion of 10% glucose. Blood glucose levels were serially monitored with a Glucometer (Lifescan, USA) and Glucose Infusion Rates (GIR) were adjusted every 5-10 min as needed to maintain the blood glucose at euglycemic levels (5.0±0.5 mmol L⁻¹). Clamping was achieved by 90 min and maintained for 30 min. The mean GIR was calculated based on GIR readings corresponding to the last 8 samplings over 90-120 mg/kg/min (Chen et al., 2004).

Immunohistochemical detection of the expression of PPARα: Myocardial tissue removed, embedded in paraffin sections after conventional serum gamma dew axing closed to water, 3% hydrogen peroxide elimination of exogenous catalase with 0.01 mol L⁻¹ of PBS after washing 3 times by adding to 1:200 dilution of goat anti-rat cardiac-specific PPARα antibody (SANTACRUZE original, rat anti-goat) were incubated overnight at 4°C.

Then, with 1:100 dilution respectively HRP labeled goat anti-rabbit IgG antibody (SANTACRUZE companies, Beijing Zhongshan Biotechnology Limited packed) were incubated for 1 h. At last, DAB staining the sections. The staining of cardiomyocytes were detected at 250x magnification in light microscopy.

Detection of ultra structural situation: The fragments were cut into 1 mm cubic pieces which were soaked in 1% OSO₄ solution for 2 h at 4°C and then dehydrated and embedded in araldite. Silver or gray thin sections (60-90 mm) were selected and cut on a Porter-Blum MT-B ultra microtome, mounted on copper silver grids with 200 patches and stained with uranyl acetate and lead citrate. The samples were selected, collected and deposed on copper wire with 200 patches. The preparations were examined with an electronic microscope (Model EM 90, Carl Zeiss) using a tension of 80 kV.

Detection of apoptosis: PBS will be organized by washing three times after the 4% more than in the PGM fixed 25 min was detected by TUNEL apoptosis, the kit used by the Nanjing Kai Ji Biological Limited. Proteinase K digestion will be used after the biopsy of coverage by TUNEL, 37°C incubated for 90 min; SSC termination reaction, antibody-HPHR incubated 15 min at room temperature, PBS washed DAB colorimetric.

Light microscopy showed apoptotic cells depth varying brown. Semi-quantitative analysis of apoptotic cells: according distribution of apoptotic cells in 400 times the light microscope, each slice shooting seven positive vision, each counting 200 myocardial per-vision to calculate the average percentage of apoptotic cells as a down-death index.

Western blot analysis: Nucleus protein were diluted 1:1.5 in SDS-Laemmli buffer (30) and boiled for 5 min. Equivalent amounts of each treatment group were run on
10% SDS polyacrylamide gels. Proteins were transferred electrically (12 V, 30 min) to PVDF membrane and incubated with a rat anti-goat PPARα, NF-κB (P65) and β-actin polyclonal antibody (1:300, Santa Cruz Biotechnology, Santa Cruz, CA; 2 h at 37°C). After 3 times washed with TBST followed by incubation with goat radish Streptavidin-HRP-linked anti-mouse IgG (1:100, Santa Cruz Biotechnology, Santa Cruz, CA; 2 h at 37°C). After Lumiglo reagent (Photo top-HRP Western blot Detection kit; Biolabs, Beverly, MA) was added, the emitted light was captured on X-ray film. Densitometric analysis was performed with quantity one software.

Statistical analysis: All results were analyzed by Spss13.0 Software. Data were analyzed by 1-factor Analysis of Variance (ANOVA). If a statistically significant effect was found, the Newman-Keuls test was performed to isolate the difference between the groups. The p<0.05 were considered as indicative of significance.

RESULTS

Rats GIR: The determination of N rats average GIR was 12.3±4.14 mg kg⁻¹ min, D rats GIR average for 6.37±1.90 mg kg⁻¹ min, there are significant differences between the two groups, p<0.01, the high note a small amount of fat diet feeding+STZ can successfully established type 2 diabetes rat model (Table 1).

Apoptosis rate, ultra structure and PPAR, NF-κB’s protein expression

Apoptosis: Compared with Group N and D, N group rarely have the apoptosis of cells, organization arranged neatly (Fig. 1 and 2); Group D have more apoptotic cells than group N, organizations arranged disorder (Fig. 2 and 3, p<0.05, Table 2); compared with Group D and T, cell apoptosis of T group was less than Group D, organization arranged neatly (Fig. 1-3, p<0.05, Table 2).

Table 1: GIR and glucose in D group compared with contrast. (x ± s.n=8)

<table>
<thead>
<tr>
<th>Groups</th>
<th>GIR (mg kg⁻¹ min)</th>
<th>Glucose (mmol L⁻¹)</th>
</tr>
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<tbody>
<tr>
<td>N</td>
<td>12.3±4.14</td>
<td>5.21±1.74</td>
</tr>
<tr>
<td>D</td>
<td>6.37±1.90</td>
<td>19.7±3.28</td>
</tr>
</tbody>
</table>

Group N: *p<0.01

Table 2: The expression of PPARα, NF-κB in protein level and apoptosis rate in N and D group. (x ± S)

<table>
<thead>
<tr>
<th>Groups</th>
<th>Apoptosis rate n = 7</th>
<th>The expression of nucleus PPARα in protein level of each group n = 3</th>
<th>The expression of nucleus NF-κB in protein level of each groups n = 3</th>
</tr>
</thead>
<tbody>
<tr>
<td>N</td>
<td>6.14±0.02</td>
<td>0.341±0.2502</td>
<td>0.62±0.9 ± 0.0561</td>
</tr>
<tr>
<td>D</td>
<td>45.71±0.04</td>
<td>1.975±0.5682</td>
<td>2.70±0.3860</td>
</tr>
<tr>
<td>T</td>
<td>27.43±0.05</td>
<td>1.174±0.2418</td>
<td>1.23±0.21920</td>
</tr>
</tbody>
</table>

Group N: *p<0.05; group D: *p<0.05

PPARα’s protein expression: Group N muscle fibers arranged neatly, PPARα positive staining and expressing (Fig. 4, Table 2), Group D muscle fiber disarranged, PPARα positive staining were deeper than N group also the nuclear protein expression (Fig. 5-7, p<0.05, Table 2), Group T myocardial PPARα positive staining shallow than Group D also the nuclear protein expression (p<0.05, Table 2).

NF-κB’s protein expression: The NF-κB expression in the nuclear protein of Group D is higher than group N (Fig. 8, p<0.05, Table 2). The NF-κB expression in the nuclear protein of Group D is higher than Group T (Fig. 8, p<0.05, Table 2). The NF-κB expression in the nuclear protein of Group T is higher than Group N (Fig. 8, p<0.05, Table 2).

Ultra structure: In Group N, cardiomyocytes arranged regularly, intensive; in Group D, nuclear swelling, muscle

Fig. 1: The cardiomyocytes of N group TUNEL staining (x400)

Fig. 2: The cardiomyocytes of group D TUNEL staining (x400)
Fig. 3: The cardiomyocytes of T group; TUNEL staining (x400)

Fig. 4: Expression of PPARα in cardiomyocytes of N group-immunohistochemistry staining (x250)

Fig. 5: Expression of PPARα in cardiomyocytes of D group-immunohistochemistry staining (x250)

Fig. 6: Expression of PPARα in cardiomyocytes of T group-immunohistochemistry staining (x250)

Fig. 7: The expression of nucleus PPARα in protein level of each group-western blotting; 55 kD: PPARα 42 kD: β-actin; Line 1: Group D; Line 2: Group T; Line 3: Group T

Fig. 8: The expression of nucleus NF-κB in protein level of each group-western blotting; 65 kD: NF-κB 42 kD: β-actin; Line 1: Group D; Line 2: Group T; Line 3: Group T

wire dissolved, fracture, the intermittent of disks are significantly expanded. In Group T, cardiomyocytes were
Diabetic cardiomyopathy is one of the chronic cardiovascular complications of diabetes, it has been recognized since 1974 but its mechanism has not yet been clarified. In recent years, many study found that apoptosis of cardiac cell and oxidative stress may play a key role in its process (Fang et al., 2004).

PPARα is mainly distributed in cardiac, skeletal muscle, liver and kidney tissues and play a key role in the activation of oxidative metabolism of fatty acids, cell proliferation and apoptosis (Lefebvre et al., 2006). Recent studies found that overexpression of PPARα can change the myocardial energy supply mode and lead to cardiac myocyte apoptosis (Dyntas et al., 2001; Finck et al., 2003; Boudina and Abel, 2007; Indu et al., 2006). NF-κB can be activated by PPARα directly (Philippe Lefebvre et al., 2006) in the mediating of chronic inflammation, oxidative stress and apoptosis of cells directly or indirectly in many paths (Fang et al., 2004; Farhangkhoei et al., 2006; Matsuzawa, 2006). Many scholars believe that Type 2 Diabetes is a chronic inflammatory diseases, the activation of NF-κB and the increasing of oxidative stress level may stand an important position in the development of diabetes and its cardiovascular complications (Indu et al., 2006; Farhangkhoei et al., 2006; Matsuzawa, 2006).

Trimetazidine is widely used in the treatment of coronary heart disease and myocardial ischemia, its can block 3-Acyl-CoA S-transferase and inhibit the fatty acid oxidation, correct myocardial cell energy metabolism disorders, decrease the oxidative stress level and reduce the apoptosis of cardiac cell (Aussedat et al., 2006;
Tabbi-Anneni et al., 2003; Rosano et al., 2003). In this experiment, compared to the Group N, the Gf of Group D is decreased and the blood glucose is increased, it has proved that researchers established the type 2 diabetic rat model successfully. In Group D, PPARα is more expression than Group N (p<0.05), the nuclear is swelling, muscle wire dissolved, fracture, the intermittent of disks are significantly expanded and TUNEL pictures show that Group D has more cardiomyocyte apoptosis than the Group N (p<0.05), so it has proved that the existence of diabetic cardiomyopathy in the diabetic rat model. From the Fig. 7 and 8 and Table 2, PPARα’s expression of Group D are significantly higher than Group N also the NF-κB and apoptotic cells in Group D is increasing compare with Group N (p<0.05).

It can prove that long period diabetes may induce the over expression of PPARα and NF-κB, increase the apoptosis of cardiac cell and then change the cardiac ultra structure of diabetes rat. In the Group T, we found that after the therapy of trimetazidine, compared to Group D, the apoptosis of cardiac cells decreased, cardiomyocytes and capillaries are normal, Z lines are less fuzzy, the expression of PPARα is lower than Group D also the NF-κB (p<0.05), it can prove that trimetazidine may have improvement effect on the change of cardiac ultra structure of diabetic cardiomyopathy, decrease the apoptosis of cardiac cell in rat’s diabetes rat and these effect may due to the reduction of over expressed PPARα and the NF-κB downstream.

Although, the previous researches believed that the activation of PPARα in cardiac cell has protective effect, this is also the theoretical basis of application of fibrates (Lefebvre et al., 2006). Recent studies suggested that over expression of PPARα activate the downstream path of the fatty acid metabolism through the key enzyme MCPT-1 (carnitine Acyl-CoA transacylase -1), FAT/CD36 (Acyl-transposition of enzyme) and promote the oxidation of free fatty acids. At the early stage of diabetes, it can compensate for the shortage of energy caused by the lack of glucose intake but the long-term over expression of PPARα may cause the providing of energy is depended on β-oxidation of fatty acid almost. This changing pattern of metabolism can cause myocardial cells and myocardial oxygen consumption increased, induce ROS accumulation and also the intracellular concentration ceramide increasing, endothelial Nitric Oxide (eNO) reduction inducible Nitric Oxide (iNO) increasing, all of these may induce the apoptosis of cardiac cell (Dyntar et al., 2001; Finek et al., 2003; How et al., 2006; Finek et al., 2003; Carley and Severson, 2005).

CONCLUSION

Through the experiment, researchers conclude that long-term diabetes metabolic disorder may cause the over-expression of PPARα and the NF-κB downstream expression, it may be involved in the cardiac cells apoptosis and diabetic cardiomyopathy. The intervention of trimetazidine stand the anti-apoptotic role, it can reduce the excessive expression of PPARα and NF-κB, the mechanism may lies in the blockage of PPARα can reverse fatty acid β-oxidation, corrected myocardial energy metabolism disorder, decrease the NF-κB expression, reducing myocardial oxygen consumption and superoxide clusters (ROS) aggregation (Lefebvre et al., 2006; Boudina and Abel, 2007; Indu et al., 2006; Farhangkhoee et al., 2006; Matsuzawa, 2006; Rosano et al., 2003).

ACKNOWLEDGEMENTS

This study was supported by research grants from the National Natural Science Foundation of China (No.: 30670838) and Health grants Council of Chongqing (07-2-78). Liang ZiWen and Wu QiNan are contributed equal to this research.

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