

RSC Advances

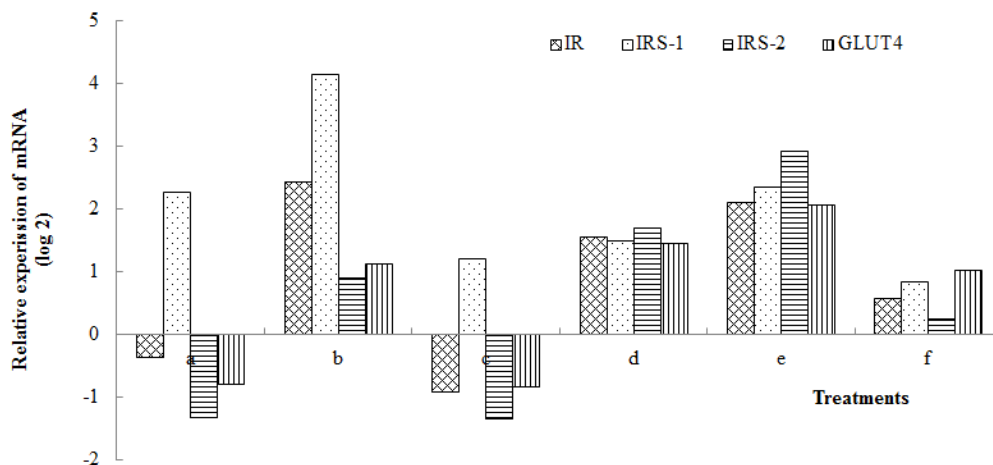


This is an *Accepted Manuscript*, which has been through the Royal Society of Chemistry peer review process and has been accepted for publication.

Accepted Manuscripts are published online shortly after acceptance, before technical editing, formatting and proof reading. Using this free service, authors can make their results available to the community, in citable form, before we publish the edited article. This *Accepted Manuscript* will be replaced by the edited, formatted and paginated article as soon as this is available.

You can find more information about *Accepted Manuscripts* in the [Information for Authors](#).

Please note that technical editing may introduce minor changes to the text and/or graphics, which may alter content. The journal's standard [Terms & Conditions](#) and the [Ethical guidelines](#) still apply. In no event shall the Royal Society of Chemistry be held responsible for any errors or omissions in this *Accepted Manuscript* or any consequences arising from the use of any information it contains.



Relative expression of IR, IRS-1, IRS-2 and GLUT4 mRNA was determined in insulin-resistant 3T3-L1 adipocytes (a-f), which were (a) 1 μ g/mL GTF-treated, (b) 1 μ g/mL GTF- and insulin-treated, (c) 1 μ g/mL Cr(pic)₃-treated, (d) 1 μ g/mL Cr(pic)₃- and insulin-treated, (e) 5 μ mol/L rosiglitazone- and insulin-treated, (f) 5 μ mol/L rosiglitazone-treated.

ARTICLE

Effect of glucose tolerance factor (GTF) from high chromium yeast on glucose metabolism in insulin-resistant 3T3-L1 adipocytes

Cite this: DOI: 10.1039/x0xx00000x

Received 00th January 2012,
Accepted 00th January 2012

DOI: 10.1039/x0xx00000x

www.rsc.org/

L. Liu,^{a,b} W.M. Cui,^{a,b} S.W. Zhang,^{a,b} F.H. Kong,^{b,d} M.A. Pedersen^c, Y. Wen,^e and J.P. Lv,^{a,b*}

The purpose of this paper was to assess and compare the impact of GTF, CrCl₃ and Cr(pic)₃ on glucose metabolism and explore the underlying mechanism of GTF in insulin-resistant 3T3-L1 adipocytes. The insulin-resistant 3T3-L1 adipocytes were induced by incubation with insulin for 48h. Purified GTF from high chromium yeast was used in this study, with m/z of 769 to 712, and glutamic acid, glycine, and cysteine in an approximate ratio of 1:1:1, as well as nicotinic acid and Cr (III). GTF, CrCl₃, Cr(pic)₃ and rosiglitazone (positive control) were applied to the cells. The effective dose of GTF ranged from 0.5ug/mL to 1.5ug/mL. GTF decreased cell viability significantly ($P < 0.01$) at doses of 3ug/mL or higher. Glucose consumption in insulin-resistant 3T3-L1 adipocytes induced by GTF increased significantly ($P < 0.05$) when incubated with GTF after 12h. Among GTF, Cr(pic)₃ and CrCl₃, GTF stimulated glucose consumption the greatest. In the presence of insulin, the relative expression level of insulin receptor (IR), insulin receptor substrate-1 (IRS-1), insulin receptor substrate-2 (IRS-2) and glucose transporter 4 (GLUT4) mRNA were increased by GTF by 2.4, 4.1, 0.9 and 1.1-fold, respectively, however, only IRS-1 was increased by 2.3-fold in the absence of insulin. GTF affected mRNA levels of IR and IRS-1 significantly ($P < 0.01$) as compared to the other two. This study not only further demonstrates that chromium containing complexes show promise in reducing insulin resistance in instances of type 2 diabetes, but also that among the chromium complexes, GTF performs best. Additionally, new mechanistic details of how GTF affects mRNA levels of insulin signalling proteins were revealed.

1 Introduction

Chromium (Cr) is required for normal insulin function, and low Cr levels have been linked with insulin resistance. Severe chromium deficiency is accompanied by symptoms resembling diabetes mellitus. Experiments in rats and squirrel monkeys demonstrated that dietary deficiency of chromium can result in elevated levels of blood glucose, triglycerides, and cholesterol.¹ These metabolic abnormalities are readily reversed by the administration of inorganic chromium. Chromium supplements are available as trivalent chromium in chloride, picolinate and other forms.

The best-selling and most studied chromium nutraceutical is chromium picolinate (Cr(pic)₃). A recent study suggests, however, that when administered intravenously and in the presence of reactive oxygen species, Cr(pic)₃ causes damage as it diffuses into cells.² Compared with Cr(pic)₃, organic chromium yeast seems to be a safer form of chromium supplementation. Many studies have reported that

supplementation with high-chromium yeast can improve diabetes more efficiently.³⁻⁶ When treated with both chromium-enriched yeast and reduced exogenous insulin, type 2 diabetic patients exhibit significantly decreased plasma glucose and glycated haemoglobin when compared to insulin treatment alone.⁷

GTF is a low molecular weight substance which possesses an ultraviolet absorbance maximum at 260nm and is comprised of Cr³⁺, glycine, glutamate, a sulfur-containing amino acid, and nicotinic acid. It stems from the report of Mertz and others on the isolation of brewer's yeast GTF in 1977.⁸ GTF is named after its original function, stimulating glucose metabolism in fat pad assays, and typically refers to the yeast-derived material. Therefore, the current term GTF may be different from its initial definition or the original composition proposed. Because of the original harsh isolation conditions and a lack of direct structural data, some researchers have suggested that GTF may be an artifact resulting from hydrolysis of porcine low-molecular-weight chromium-binding substance (LMWCr) and

that the use of the term GTF should be abandoned.^{2, 9-11} Many experiments have shown that high-chromium yeast enhances the action of insulin and improve diabetes in animals.¹¹⁻¹⁴ In diabetic mice supplemented with high-dose, high-chromium yeast (Cr 1000ug/kg.d), the number and size of pancreas islet cells decreased in comparison to those in mice supplemented with low-dose high-chromium yeast (Cr 250ug/kg.d, 500ug/kg.d) and normal yeast (Cr <0.1ug/g dry yeast).¹¹ In this study, our previously established mild purification method was used to purify GTF from high chromium yeast.¹⁵ It was identified, with mass-to-charge ratios (m/z) of 769 and 712, which included glutamic acid, glycine, and cysteine in an approximate ratio of 1:1:1, as well as nicotinic acid and Cr (III). Although previous animal experiments have shown that this high chromium yeast could improve type 2 diabetes, especially for pancreas recovery,¹¹ it is not well understood how purified GTF acts directly on the cells, especially 3T3-L1 insulin-resistant adipocytes. Furthermore, the optimum GTF dosage, cytotoxic effect, and differing effects between GTF and other chromium complexes should be determined. To better understand these questions, we applied the purified GTF and other chromium complexes directly on insulin-resistant 3T3-L1 adipocytes and investigated their effect on glucose metabolism. In this study, rosiglitazone, an insulin sensitizer, was used as a positive control to evaluate the ability of GTF and other chromium sources to adjust glucose consumption.

2 Materials and Methods

2.1 Reagents and cell lines

Mouse 3T3-L1 preadipocytes were obtained from the cell culture center at the Chinese Academy of Medical Sciences. Dulbecco's Modified Eagle's medium (DMEM) and fetal bovine serum (FBS) were purchased from GIBCO BRL (Gaithersburg, MD, USA). Other cell culture solutions and supplements such as trypsin, insulin, dexamethasone, 3-isobutyl-1-methylxanthine (IBMX), Cr(pic)₃, CrCl₃, rosiglitazone were obtained from Sigma (St Louis, MO, USA). The glucose assay kit was purchased from Yihua (Shanghai, China). Organic solvents and other chemicals used for extraction and purification were of the highest analytical grade. Sephadex exclusion media (sizes G-75, G-25, and G-15) were obtained from Pharmacia Ltd., UK.

2.2 Cell culture, differentiation, induction and evaluation

The 3T3-L1 preadipocytes were cultured in DMEM containing 10% fetal bovine serum, 1% penicillin and 1% streptomycin. 3T3-L1 fibroblasts were differentiated into adipocytes following standard protocols with slight modifications. Completely confluent plates were incubated in DMEM containing 10% FBS with 500uM 3-isobutyl-1-methylxanthine (IBMX), 1uM dexamethasone and 172nM insulin. Two days after incubation, the medium was replaced with DMEM containing 172nM insulin. Culture medium was replaced with fresh medium after 2d and then every other day until the cells attained adipocyte morphology. With this protocol, >90% adipocyte differentiation was achieved.

Differentiated cells were then incubated for a period of 48h in DMEM containing 10% FBS and 100nM insulin as insulin-resistant cells or in DMEM containing 10% FBS without insulin as the control (normal 3T3-L1 adipocytes), which was used to evaluate the insulin-resistant model. These cells were maintained at 37°C under a humidified 5% CO₂ atmosphere.

The insulin-resistant adipocytes and the control were incubated for a period of 24h in DMEM with or without 100nM insulin. Supernatants of these treatments were then collected and the levels of glucose concentration were measured and analysed.

2.3 Preparation of high chromium yeast

High chromium yeast was obtained as previously described.^{5, 15} The strain of yeast used in this work was *Saccharomyces cerevisiae*, mutated by spaceflight-induced mutagenesis and deposited in the China General Microbiological Culture Collection Center (WDCM550 CGMCC No. 2687). The strain was obtained from the Institute of Agro-products Processing Science and Technology, Chinese Academy of Agricultural Science, Beijing, China. After incubation with 400ug/mL CrCl₃ in Yeast Extract Peptone Dextrose (YPD) medium, for 44h at 28°C on a rotary shaker (200r/min), cells were harvested by centrifugation, washed three times with de-ionized water, and then freeze-dried. The high-chromium yeast was a tan powder with a total chromium content of 1512ug/g dry yeast, and organic chromium content of 1200ug/g dry yeast, as assayed by atomic absorption spectrometry using flame atomization.

2.4 Extraction and purification of GTF

GTF was extracted and purified from high chromium yeast as previously described.¹⁵ The freeze-dried, high-chromium yeast was extracted with 0.1mol/L ammonia by shaking for 3h at 37°C. The supernatants obtained by centrifugation were freeze-dried. The dried material was suspended in distilled water and centrifuged at 10000r/min for 30min at 4°C. The supernatants were collected, condensed to the desired volume using vacuum evaporation at 45°C and then was applied to three chromatography columns (Sephadex G75, Sephadex G25, Sephadex G15) consecutively. The chromium content of the chromatographic effluents was determined by atomic absorption spectrometry using flame atomization. Chromium-containing fractions from these columns were pooled, concentrated, and lyophilized for further cell experiments. All chromatography procedures were carried out in a cold-room maintained at 4°C.

2.5 Measurement of glucose consumption

3T3-L1 preadipocytes were plated into a 48-well plate. After following the differentiation and induction protocol, the insulin-resistant cells were treated as follows:

A: For the GTF dose response assay, insulin-resistant cells were treated with various concentrations of GTF (calculated as Cr) in the presence or absence of 100nM insulin for 24h.

B: For the GTF time response assay, insulin-resistant cells were treated with GTF (0ug/mL, 1ug/mL) for a scheduled time in the presence or absence of 100nM insulin.

C: For the chromium complex comparison, insulin-resistant cells were treated with 1ug/mL of GTF, Cr(pic)₃, and CrCl₃ (calculated as Cr) in the presence of 100nM insulin for 24h. Rosiglitazone (5umol/L) was set as positive control.

Supernatants of these treatments were collected and glucose concentrations were measured using the glucose assay kit for a quantitative determination. Glucose consumption rates were calculated by subtracting residual glucose in the treated medium from glucose in non-treated medium.

2.6 Cell viability assay - MTT assay

Cell viability assay was performed using the modified MTT methods. Cell viability is assessed by measuring the degree of formazan crystal formation. Light-yellow colored MTT is decomposed to dark-blue coloured formazan by mitochondria, and only live cells are able to generate formazan. Thus, amount of formazan formed in the cells reflects the degree of cell viability.

3T3-L1 preadipocytes were plated into a 48-well plate. After following the differentiation and induction protocol, the insulin-resistant cells were exposed to one the following media:

A: For the dose response assay, cells were treated with medium containing various concentrations (0 μ g/mL, 0.01 μ g/mL, 0.1 μ g/mL, 0.5 μ g/mL, 1 μ g/mL, 1.5 μ g/mL, 3 μ g/mL and 5 μ g/mL) of GTF (calculated as Cr) in the presence of 100nM insulin for 24h.

B: For the chromium complex comparison, cells were treated with medium containing 1 μ g/mL of GTF, Cr(pic)₃ or CrCl₃ (calculated as Cr) in the presence 100nM insulin for 24h. Rosiglitazone (5 μ mol/L) was set as positive control.

After incubation for the indicated times, the supernatants were removed. The cells were washed with PBS for three times and replaced with DMEM medium. Then 50 μ L of sterile filtered MTT solution (5mg/mL) in phosphate buffered saline (PBS, pH 7.4) was added to each well, reaching a final concentration of 0.5mg/mL MTT. After 4h, the supernatants of cells were removed. Then, formazan crystals formed in the wells were dissolved in 375 μ L dimethylsulfoxide (DMSO) and the plates were shaken for 10min. The absorbance was measured at 490nm using a Model680 Microplate Reader (BIO-RAD). Cell viability was calculated relative to the untreated control cells: viability (%) = (absorbance of treated sample)/(absorbance of control) \times 100.

2.7 RNA preparations and real-time fluorescence quantitative PCR (RT-qPCR)

Total RNA was isolated using the Trizol system (Watson, Shanghai) according to the manufacturer's guidelines. A reverse transcription system commercial kit (Promega) was used for cDNA synthesis. The primers of target genes and reference gene were designed using Primer Premier 5.0. The details of primers are described in Tab.1. RT-qPCR was performed with SYBR Green I using ABI 7500 (Applied Biosystems). β -actin was selected as the reference gene. PCR amplification was performed on a final 20 μ L containing cDNA template (4ng) and primers (400nM each). The following PCR program was used: one cycle of 95 $^{\circ}$ C for 10min and 35 cycles of 95 $^{\circ}$ C for 15s, 50 $^{\circ}$ C for 15s and 72 $^{\circ}$ C for 15s. A melting curve analysis was done at the end of each run for all primer sets. This resulted in single-product-specific melting curves, and no primer dimers were generated during the runs. A "no template control" (distilled H₂O) and a "negative control" (RNA samples which had not undergone the reverse transcription step) were included in each run in order to confirm the absence of DNA contamination. Quantifications were performed in duplicate and the experiments were repeated three times independently.

Tab.1 Nucleotide sequence of real-time PCR primers

Gene	Primes	Tm	Product length	Gene No.
β -actin	GAGACCTTCAACACCCAGC	50 $^{\circ}$ C	446 bp	NM_007393.3
	CCACAGGATCCATACCCAA			
Glucose transporter 4 (GLUT4)	ATTGGACGCTCTCTCCAA	50 $^{\circ}$ C	168 bp	NM_009204.2
	GATTCTGCTGCCCTTCTGTC			
Insulin receptor (IR)	AAATCGTCAACCTGCTC	50 $^{\circ}$ C	137 bp	NM_010568.2
	ATCCAACGGGACATTC			
Insulin receptor substrate 1 (IRS-1)	GGATCGTCAATAGCGTAA	50 $^{\circ}$ C	232 bp	NM_010570.4
	GCTTGGACAATGTAGAA			
Insulin receptor substrate 2 (IRS-2)	CATGTCCCTGACGAGTATG	50 $^{\circ}$ C	225 bp	NM_001081212
	TTCTCAGTCTCTATCCAG			

2.8 Statistical Analysis

Measurement data were presented as means \pm standard error. A pooled Student's test was used in each group to evaluate the difference in mean concentrations between the control and other groups. Multiple comparisons were performed with Duncan's Multiple Range Test. $P < 0.05$ was considered to be a significant difference. $P < 0.01$ was considered extremely significant.

3 Results

Adipocytes, major target cells of insulin, are used as a model system for studying insulin resistance in vitro. Treating hyperglycemia with insulin could induce the development of insulin resistance in 3T3-L1 adipocytes, which closely mimic the pathophysiological abnormality of type 2 diabetes.¹⁹ The ability of chromium complexes, including GTF extract, to enhance glucose uptake and metabolism in normal 3T3-L1 adipocytes and fat cells has been reported previously.^{16, 17, 18, 30} Based on our previous experiment, high chromium yeast has no effect on blood variables and pancreas pathology in normal Balb/c mice, however, it improved blood variables and pancreas pathology in diabetic type 2 mice at high dosages⁵. Therefore, in this study, purified GTF was applied directly to insulin-resistant 3T3-L1 adipocytes to investigate its effects including viability, glucose metabolism and relative expression of mRNA of IR, IRS-1, IRS-2 and GLUT4. Furthermore, the effects of different chromium complexes on insulin-resistant 3T3-L1 adipocytes were also examined and compared.

As shown in Fig.1, glucose consumption in insulin-resistant 3T3-L1 adipocytes treated with and without insulin was 6.49mmol/L and 6.19mmol/L, respectively. This showed that insulin could not stimulate the insulin-resistant model to consume more sugar. Glucose consumption in normal 3T3-L1 adipocytes treated with and without insulin was 13.8mmol/L and 8.00mmol/L, respectively. Glucose consumption in insulin-resistant 3T3-L1 adipocytes and 3T3-L1 adipocytes treated with insulin was significantly different ($P < 0.01$). This demonstrates that the model of insulin-resistant 3T3-L1 adipocytes was successfully established.

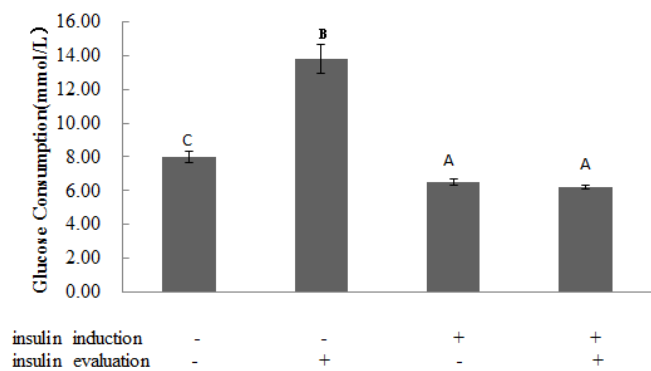


Fig.1 Evaluation of insulin-resistant model. 3T3-L1 adipocytes were incubated with 100 nM insulin as induction or without insulin as control for 24h. Insulin-resistant adipocytes and non insulin induced control cells were then exposed to DMEM in the absence or presence of 100 nM insulin for 24h before evaluation. Supernatants were then collected for the glucose assay. Results are expressed as mean values \pm SD (n=6). Multiple comparisons were carried out and capital letters means different at $P < 0.01$.

3.1 Effect of GTF on glucose metabolism of insulin-resistant 3T3-L1 adipocytes

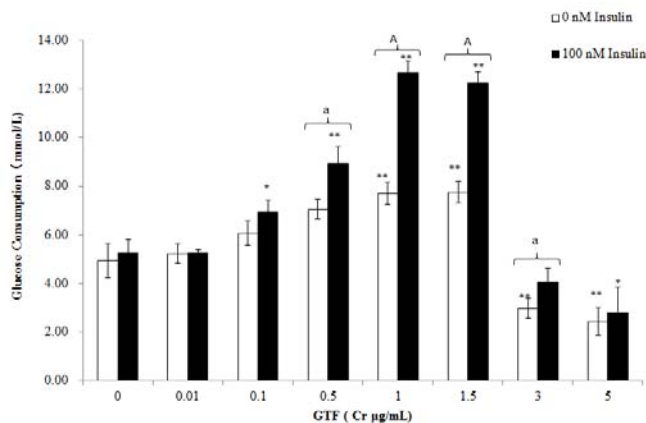


Fig.2 GTF increased glucose consumption of insulin-resistant 3T3-L1 adipocytes in a dose-dependent manner. The insulin-resistant 3T3-L1 adipocytes were exposed to multiple concentrations of GTF (calculated as Cr) in the absence or presence of 100 nM insulin for 24h. Supernatants were then collected for the glucose assay. Results are expressed as mean values \pm SD (n=6). Asterisks indicate a difference compared with the corresponding control (in the presence or absence of insulin, 0 μ g/mL GTF treatment), * $P < 0.05$, ** $P < 0.01$. Letters indicate a difference between 0nM insulin- and 100nM insulin-treated cells for the same GTF dosage. Lower letters means different at $P < 0.05$. Capital letters means different at $P < 0.01$.

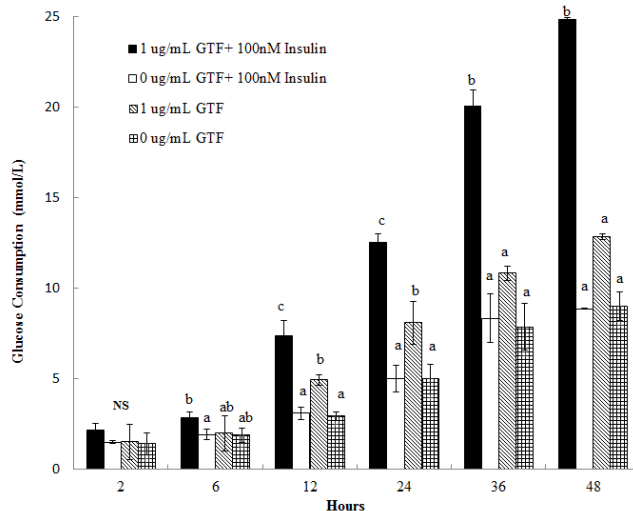


Fig.3 GTF increased glucose consumption of insulin-resistant 3T3-L1 adipocytes in a time-dependent manner. The insulin-resistant 3T3-L1 adipocytes were exposed to 1 μ g/mL or 0 μ g/mL GTF (calculated as Cr) in the absence or presence of 100nM insulin for a scheduled time. Supernatants were then collected for the glucose assay. Results are expressed as mean values \pm SD (n=3). Multiple comparisons were carried out among the four treatments for each incubation time. Bars topped by different lower case letters displayed above the columns of the figure are significantly different at $P < 0.05$. Time periods with no significant differences between treatments are labelled non-significant (NS).

Glucose consumption was examined when insulin-resistant 3T3-L1 adipocytes were exposed to multiple concentrations of GTF in the presence or absence of insulin for a fixed time. As shown in Fig.2, compared with the treatment of 0nM insulin and 0 μ g/mL GTF, glucose consumption in insulin-resistant 3T3-L1 adipocytes treated by 100nM insulin and 0 μ g/mL GTF was only enhanced by 6%. No significant difference in glucose consumption was observed for 0 μ g/mL GTF-treated insulin-resistant 3T3-L1 adipocytes in the presence or absence of insulin. Fig.2 showed that GTF increased glucose consumption significantly in a dose-dependent manner in the presence of insulin, while GTF alone could not increase glucose consumption as high as it did in the presence of insulin. At a dose between 1 μ g/mL to 1.5 μ g/mL, GTF increased glucose consumption by 56% in the absence of insulin ($P < 0.01$). In the presence of insulin, glucose consumption was further augmented by GTF, which led to a 141% increase at 1 μ g/mL ($P < 0.01$). GTF accelerated glucose consumption of insulin-resistant 3T3-L1 adipocytes by potentiating the effect of insulin. The dose-dependent experiment showed that stimulation of glucose consumption was maximal at 1 μ g/mL GTF, which was employed in subsequent experiments to compare effects of different chromium complexes. But at a dose of 3 μ g/mL and 5 μ g/mL, GTF decreased glucose consumption greatly. Glucose consumption decreased by 22%-46% in the presence of insulin and 39%-50% in the absence of insulin, respectively. The cells were inhibited by high concentrations of chromium, which would be proved by the MTT assay.

At low doses of GTF (from 0.01 μ g/mL to 0.1 μ g/mL), there was no significant difference in glucose consumption between 0nM insulin- and 100nM insulin-treated cells. As GTF increased (from 0.5 μ g/mL to 1.5 μ g/mL), the difference in glucose consumption between 0nM insulin- and 100nM insulin-treated

cells became significant for the 0.5ug/mL treatment and greatly significant for the 1ug/mL and 1.5ug/mL treatments. GTF treatment also increased basal glucose consumption (without insulin) as shown in Fig.2. At higher doses of GTF, glucose consumption decreased significantly. The difference in glucose consumption between 0nM insulin- and 100nM insulin-treated cells was significant ($P < 0.05$) for 3ug/mL treatment and insignificant at the higher GTF dose.

To further illuminate the effect of GTF on glucose metabolism, insulin-resistant 3T3-L1 adipocytes were exposed to 0ug/mL and 1ug/mL GTF both in the presence and absence of 100nM insulin, and glucose consumption was evaluated after 2h, 6h, 12h, 24h, 36h and 48h. As shown in Fig.3, without GTF treatment, no significant difference in glucose consumption was observed for the treatment with and without 100nM insulin for 48h. This means that the cell model for insulin resistance in vitro was successfully established. Significant differences in glucose consumption were observed between the treatment with 1ug/mL GTF and without GTF in the presence of 100nM insulin starting after 6h of incubation. Interestingly, without insulin treatment, significant differences in glucose consumption between treatment with and without GTF were observed only after 12h and 24h of incubation.

As a whole, GTF enhanced glucose consumption significantly in the presence of 100nM insulin after 6h and glucose consumption increased to 24.88mmol/L after 48h. GTF caused a 2.5 and 1.5-fold increase in glucose consumption with and without 100nM insulin, respectively, after 24h, and a 2.8-fold and 1.9-fold increase in glucose consumption after 48h. Considering the glucose consumption rate, half of the glucose in the medium was consumed after 24h incubation. Almost all glucose in the medium was consumed after 48h. 24h incubation periods were used for all further experiments.

3.2 Comparison of glucose metabolism of insulin-resistant 3T3-L1 adipocytes for different chromium complexes

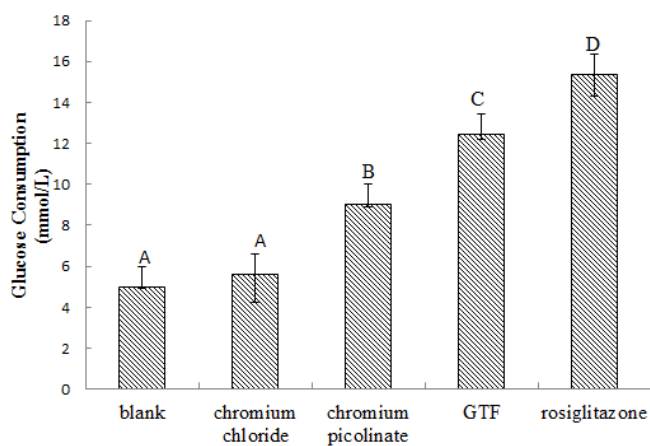


Fig.4 Effects of different chromium complexes on glucose consumption in insulin-resistant 3T3-L1 adipocytes. The insulin-resistant 3T3-L1 adipocytes were incubated with 1ug/mL GTF, Cr(pic)₃, and CrCl₃ (calculated as Cr) respectively in the presence of 100nM insulin for 24h. Rosiglitazone (5umol/L) was set as positive control. Supernatants were then collected for the glucose assay. Results are expressed as mean values \pm SD (n=3). Bars topped by different capital letters are significantly different at $P < 0.01$.

The effects of three chromium complexes on glucose metabolism in insulin-resistant 3T3-L1 adipocytes were compared in this section. Among these chromium complexes, CrCl₃ and Cr(pic)₃ were representative of inorganic chromium and organic chromium, respectively. Rosiglitazone was set as a positive control to evaluate their ability to regulate glucose consumption. As shown in Fig.4, all treatments, except the control and CrCl₃, could cause the stimulation of glucose consumption to different degrees in the presence of 100nm insulin. Rosiglitazone had the highest stimulation on glucose consumption (15.33mmol/L) among the four treatments. The stimulation on glucose consumption for GTF, Cr(pic)₃ and CrCl₃ was 12.43mmol/L, 9.01mmol/L and 6.53mmol/L, respectively. Compared with the untreated control, rosiglitazone, GTF, Cr(pic)₃ and CrCl₃ improved the glucose consumption by a 3.1-, 2.5-, 1.7- and 1.2-fold increase, respectively. There was no significant difference between the control and CrCl₃ treatment. GTF-induced glucose consumption was significantly higher than that induced by either CrCl₃ or Cr(pic)₃. These results suggested that GTF could potentiate the effect of insulin very significantly ($P < 0.01$) enhancing glucose consumption compared with other chromium supplements, which was not comparable to rosiglitazone by itself.

3.3 Cytotoxicity of GTF and two other chromium complexes on insulin-resistant 3T3-L1 adipocytes

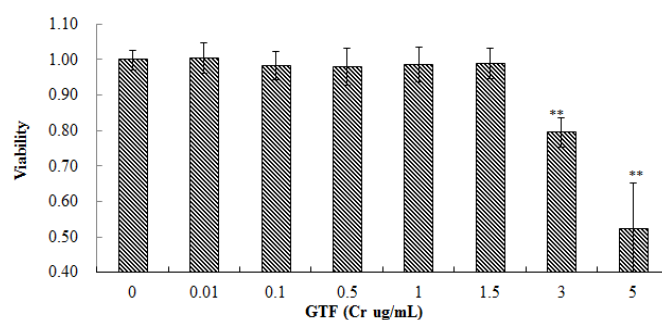


Fig.5 Effect of GTF on the viability of insulin-resistant 3T3-L1 adipocytes in the presence of 100nM insulin. The insulin-resistant 3T3-L1 adipocytes were incubated with multiple concentrations of GTF (calculated as Cr) in the presence of 100nM insulin for 24h. The MTT assay was used to evaluate cell viability after GTF treatment. Results are expressed as mean values \pm SD (n=3). Asterisks indicate a difference compared with the corresponding control (100nM insulin, 0 ug/mL GTF), ** $P < 0.01$.

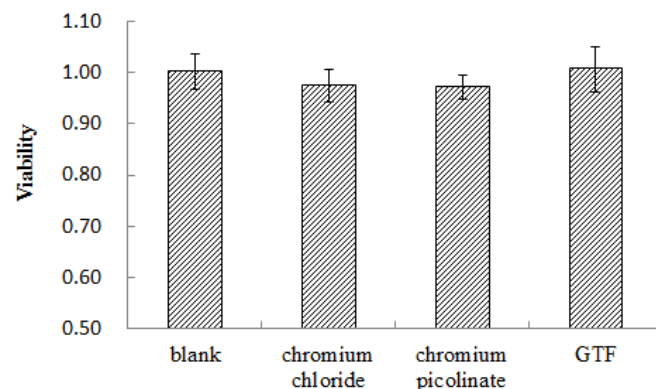


Fig.6 Effects of three chromium complexes on the viability of insulin-resistant 3T3-L1 adipocytes in the presence of 100nM insulin. The insulin-resistant 3T3-L1 adipocytes were incubated with three chromium complexes (Cr 1ug/mL) separately in the presence of 100nM insulin for 24h. The MTT assay was used to evaluate cell viability after each treatment. Results are expressed as mean values \pm SD (n=3). Bars topped by different lower case letters are significantly different at $P < 0.05$.

Considering GTF contains chromium, which is an essential part for its bioactivity, GTF inevitably has dual effects on the cells. Fig.5 shows the cells' survival profile was concentration dependent. A range of concentrations was tested and cell viability did not change significantly up to the concentration of 1.5ug/mL. Upon exposure to 3ug/mL and 5ug/mL GTF for 24h, cell viability was further reduced to 80% and 52%, respectively. This data showed that high concentrations of GTF, beyond 1.5ug/mL, were cytotoxic to the cells, which was correlated with the trends of GTF on glucose consumption. Fig.6 shows the comparison of three chromium complexes on insulin-resistant 3T3-L1 adipocytes. In the case of 1ug/mL Cr, there was no significant difference in the cell viability among these chromium complexes. The three chromium complexes did not affect the cell viability in the concentration of 1ug/mL Cr.

3.4 Effects of GTF and Cr(pic)₃ on mRNA levels of insulin receptor (IR), insulin receptor substrate-1 (IRS-1), insulin receptor substrate-2 (IRS-2), and glucose transporter 4 (GLUT4)

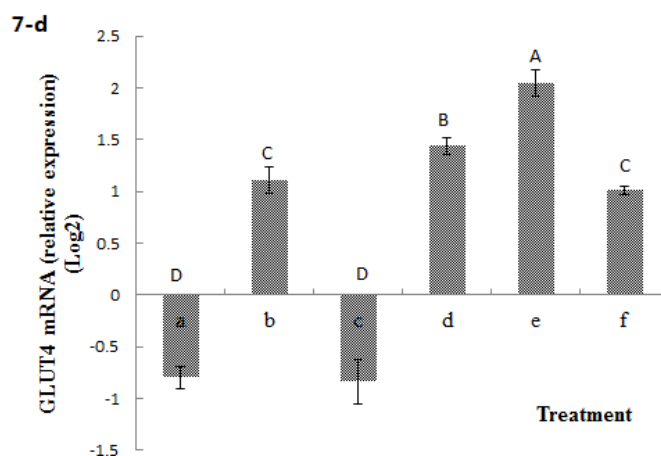
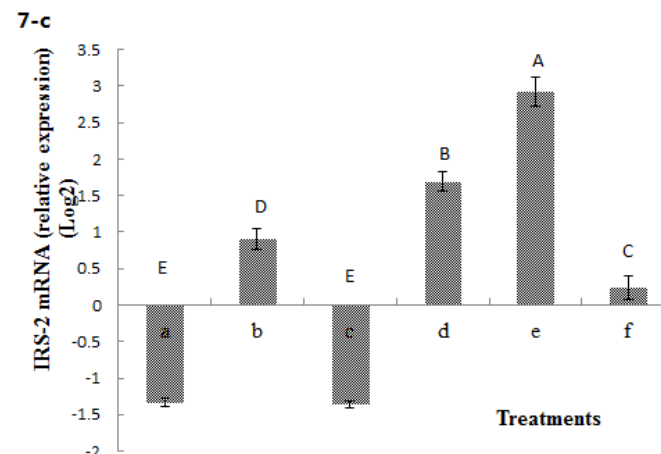
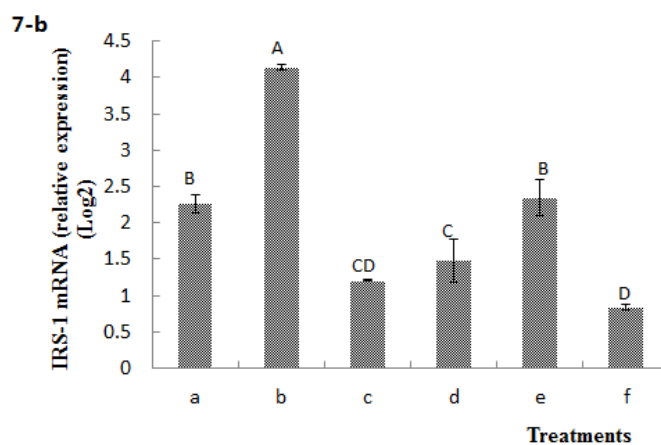
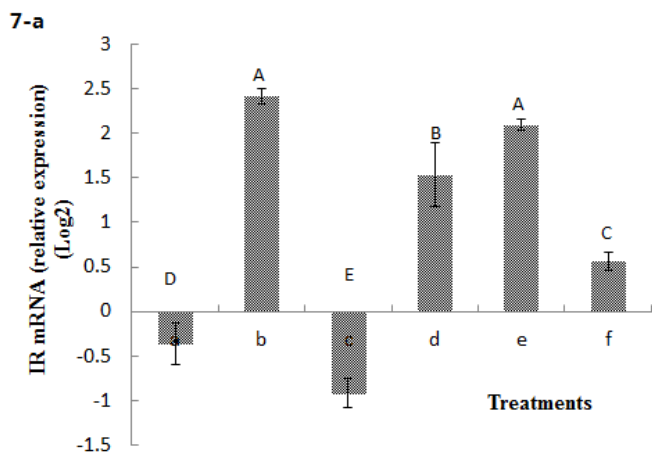


Fig.7 Relative expression of IR, IRS-1, IRS-2 and GLUT4 mRNA in insulin-resistant 3T3-L1 adipocytes (7-a, 7-b, 7-c, 7-d). Total RNA was prepared for RT-PCR from insulin-resistant 3T3-L1 adipocytes treated with GTF (Cr 1ug/mL), Cr(pic)₃ (Cr 1ug/mL) or rosiglitazone (5umol/L) in the presence or absence of 100nM insulin for 24h. PCR of β -actin was performed to verify that the initial cDNA contents of samples were similar. Relative expression of IR(7-a), IRS-1(7-b), IRS-2(7-c) and GLUT4(7-d) mRNA was determined in insulin-resistant 3T3-L1 adipocytes (a-f), which were (a)1ug/mL GTF-treated, (b)1ug/mL GTF- and insulin-treated, (c)1ug/mL Cr(pic)₃-treated, (d) 1ug/mL Cr(pic)₃- and insulin-treated, (e) 5umol/L rosiglitazone- and insulin-treated, (f) 5umol/L rosiglitazone-

treated. Multiple comparisons were carried out among the six treatments. Bars topped by different upper case letters displayed above the columns of the figure are significantly different at $P < 0.01$.

To understand the underlying mechanism for GTF, the effect of different chromium complexes on expression of IR, IRS-1, IRS-2 and GLUT4 were investigated and compared. The results showed that GTF significantly ($P < 0.01$) enhanced the level of IR, IRS-1, IRS-2 and GLUT4 mRNA expression by 2.4, 4.1, 0.9 and 1.1-fold, respectively, in the presence of insulin. GTF alone just enhanced the level of IRS-1 mRNA by 2.3-fold. While in the presence of insulin, $\text{Cr}(\text{pic})_3$ significantly ($P < 0.01$) enhanced the level of IR, IRS-2 and GLUT4 mRNA by 1.5, 1.7 and 1.4-fold, respectively. $\text{Cr}(\text{pic})_3$ alone just increased the level of IRS-1 mRNA by 1.2-fold. These results demonstrated that there were synergistic effects between chromium complexes and insulin, especially for GTF. Overall, GTF treatment had much more effect on the level of IR and IRS-1 mRNA as compared to $\text{Cr}(\text{pic})_3$. Fig 7-a and 7-b showed that GTF enhanced the IR and IRS-1 mRNA expression significantly ($P < 0.01$) in the presence of insulin.

In the presence of insulin, rosiglitazone treatment significantly ($P < 0.01$) increased the level of IR, IRS-1, IRS-2 and GLUT4 mRNA by 2.09, 2.3, 2.9 and 2.04-fold, respectively, whereas, without insulin 0.56, 0.84, 0.23 and 1-fold changes were observed. Compared with other treatments, rosiglitazone had a greater effect ($P < 0.01$) on IRS-2 and GLUT4 in the presence of insulin. GTF alone enhanced the level of IRS-1 mRNA much more than rosiglitazone alone ($P < 0.01$).

4 Discussion

Type 2 diabetes is associated with insulin resistance and relative insulin deficiency. In this disorder, tissues such as muscle, fat, and liver become less responsive or resistant to insulin. Some strategies to improve insulin sensitivity using pharmacological and nutritional supplements could alleviate insulin resistance in many individuals and improve glycemic control in some instances of diabetics. In this study, rosiglitazone was used as a positive control to evaluate glucose consumption. Rosiglitazone acts as an insulin sensitizer and modulates nuclear receptor proteins (peroxisome proliferator-activated receptors, PPARs), and it is an anti-diabetic drug in the thiazolidinedione class of drugs, which was once the best-selling diabetes therapy in the world.²⁰ This study was undertaken to explore how GTF affects glucose metabolism as compared to other chromium complexes and its potential mechanism in treating type 2 diabetes in insulin-resistant 3T3-L1 adipocytes.

The effects of GTF on insulin-resistant 3T3-L1 adipocytes were examined, including its optimal dose and incubation time [Fig. 2 and Fig. 3]. Fig. 2 shows that GTF could increase both basal (0 nM insulin) and insulin-stimulated glucose consumption in the insulin-resistant cells to different degrees. However, only when GTF reached 1 $\mu\text{g}/\text{mL}$, glucose consumption in basal cells was significantly higher than its corresponding control in the absence of insulin. In the presence of insulin, a minimum GTF concentration of 0.1 $\mu\text{g}/\text{mL}$ increased glucose consumption significantly for insulin-resistant 3T3-L1 adipocytes. It was apparent that there is a potentiation of GTF effect in the presence of insulin. Glucose consumption was significantly higher after GTF treatment in the presence of insulin when compared to GTF treatment without insulin at GTF

concentrations between 0.5 and 3 $\mu\text{g}/\text{mL}$. Yi-qun and Ming-hui reported that a significant difference in glucose uptake was observed for insulin-resistant cells treated with and without $\text{Cr}(\text{pic})_3$ in the absence of insulin. They suggested that effects of $\text{Cr}(\text{pic})_3$ were independent of insulin, which could be regarded as an insulin analogue²⁸. It is our finding, however, that the activities of insulin and GTF are not independent.

As shown in Fig. 3, significant differences could be observed between 1 $\mu\text{g}/\text{mL}$ and 0.1 $\mu\text{g}/\text{mL}$ GTF treatments from 6h to 48h incubation time. GTF significantly increases glucose consumption in a dose-dependent manner in the presence of insulin, especially at concentrations between 0.1 $\mu\text{g}/\text{mL}$ ($P < 0.05$) to 1.5 $\mu\text{g}/\text{mL}$ ($P < 0.01$), suggesting GTF could partially reduce hyperglycemia and insulin-induced insulin resistance.

Previous studies show that Cr bioavailability of different chromium complexes varies greatly, and Cr absorption ranges between 0.4% and 2.0% for inorganic complexes while the availability of organic Cr is more than 10 times higher.^{21, 22} Chromium complexes will break down in the digestive system when administered orally. If they are administered intravenously or applied to the cells directly, chromium complexes briefly diffuse into the cells and affect the cells' metabolism.²³ Three chromium complexes were applied to the cells separately in this study. CrCl_3 , as inorganic chromium, is authorized to be used as food fortifier in China. At a dose of 1 $\mu\text{g}/\text{mL}$ Cr, no cytotoxic effect was observed with any of the three chromium complexes. While in the case of glucose consumption, these chromium complexes exhibited different effects. GTF could increase glucose consumption most significantly ($P < 0.01$), followed by $\text{Cr}(\text{pic})_3$ and CrCl_3 .

The pathophysiology of insulin resistance involves a complex network of signalling pathways, activated by the insulin receptor, which regulates intermediary metabolism and its organization in cells. Insulin receptor substrate (IRS) proteins are a family of molecules that are directly phosphorylated by the insulin receptor, which leads to the recruitment and activation of additional signalling proteins.²⁴ Insufficient expression of IRS-1, IRS-2 or abnormal phosphorylation will affect glucose metabolism and lipid metabolism during signalling pathways. Weksler-Zangen et al applied GTF on the 3T3-L1 adipocytes and suggested that GTF does not exert its effects via the insulin receptor but in a different pathway²⁵. In their study, however, the GTF did not include chromium. The requirement of chromium in GTF preparation is still controversial.

Understanding the molecular basis of the mechanism of action of GTF and insulin is of crucial importance. Based on the cytotoxicity and cell metabolism study, the effects of GTF, $\text{Cr}(\text{pic})_3$ and rosiglitazone on the levels of IR, IRS-1, IRS-2 and GLUT4 mRNA in the insulin resistant adipocytes were examined. Earlier reports showed that GTF's natural function was to bind insulin, enhancing its interaction with insulin receptor.²⁶ Vincent thought that GTF did not have an intrinsic activity in the cells but was simply acting as a source of chromium, restoring the chromium pool of the cells from the Cr-deficient rats.^{25, 27} Our studies showed that different chromium complexes did exert various effects on the expression of mRNA. CrCl_3 and $\text{Cr}(\text{pic})_3$ have been reported to improve the glucose uptake and to upregulate mRNA levels of the insulin receptor, GLUT4, glycogen synthase and uncoupling protein-3 in skeletal muscle cells.²⁸ Yin reported that $\text{Cr}(\text{pic})_3$ had no effect on insulin receptor in insulin-resistant 3T3-L1 adipocytes.²⁹ Our study showed that $\text{Cr}(\text{pic})_3$ increased the level of IR mRNA significantly ($P < 0.01$) in the

presence of insulin as compared to Cr(pic)₃ treatment alone. GTF increased the level of IR mRNA higher and more significantly ($P<0.01$) than the Cr(pic)₃ treatment did. In the case of GLUT4 and IRS-2, however, Cr(pic)₃ increased mRNA levels higher and more significantly ($P<0.01$) than GTF did. Laurène Martinez et al., reported that rosiglitazone increased cell surface GLUT4 levels by increasing its endosomal recycling and restoring insulin-induced GLUT4 translocation in insulin resistance.³⁰ Our studies showed that rosiglitazone alone influenced the expression of GLUT4 mRNA in the insulin-resistant 3T3-L1 adipocytes. Rosiglitazone exhibited a significant effect ($P<0.01$) on GLUT4 and IRS-2 in the presence of insulin.

There were synergistic effects between insulin and the three chromium complexes. The synergistic effect was higher for GTF than the other two complexes. In the presence of insulin, the levels of IR, IRS-1, IRS-2 and GLUT4 mRNA could be enhanced by GTF to different degrees, while only IRS-1 was enhanced by GTF in the absence of insulin. Both GTF and Cr(pic)₃ affected the levels of these four mRNA in the presence of insulin. Furthermore, GTF exerted its effect more on the levels of IR and IRS-1 mRNA than Cr(pic)₃ did in the presence of insulin. It is widely recognized that insulin stimulates glucose uptake via the translocation of GLUT4 to the plasma membrane in the muscle and adipose tissue.²⁹ IR and IRS-1 was significantly increased by GTF treatment in the presence of insulin. Insulin receptor, as the binding spot to insulin, could affect the glucose metabolism by initiating a cascade of phosphorylation of several substances. IRS-1 is responsible for transmitting the signal from insulin receptor to biological endpoints. To further illuminate how GTF affects signalling pathways, phosphorylation of IRS-1, IR- β , AKt and c-Cbl must be compared and analysed.

In summary, the effective dose of GTF ranged from 0.5 μ g/mL to 1.5 μ g/mL. The cells' viability decreased to 80% or lower when the dose of GTF exceeded 1.5 μ g/mL. The glucose consumption of insulin-resistant 3T3-L1 adipocytes increased significantly when treated with GTF for 12h. Among GTF, Cr(pic)₃ and CrCl₃, GTF stimulated glucose consumption most dramatically. In the presence of insulin, the levels of IR, IRS-1, IRS-2 and GLUT4 mRNA could be increased by GTF by 2.4, 4.1, 0.9 and 1.1-fold, respectively, while only IRS-1 was enhanced by 2.3-fold in the absence of insulin. GTF affected mRNA level of IR and IRS-1 significantly ($P<0.01$). Cr(pic)₃ affected mRNA levels of IRS-2 and GLUT4 significantly ($P<0.01$). The investigation showed that GTF could serve as adjuvant therapy for diabetes, lowering doses of current medications that have potential side effects, or delaying the continued onset of the disease. To our knowledge, this is the first report to show that purified GTF increases glucose consumption in insulin-resistant 3T3-L1 adipocytes better than Cr(pic)₃ and CrCl₃ and this is the first description of how GTF affects mRNA levels of proteins involved in insulin signalling.

Acknowledgements

Funding is provided by the National Natural Science Foundation of China (31371763) and the National Space Breeding Project (2006HT00013). L. Liu designed the experiment, interpreted the data and contributed to the writing. F.H. Kong and W.M. Cui contributed to the acquisition of data. S.W. Zhang and Y. Wen contributed with scientific advice. M.A. Pedersen contributed to revise the manuscript. J.P. Lv contributed to the supervision of the manuscript. The authors have declared no conflict of interest.

Notes

^a Institute of Agro-products Processing Science and Technology, Chinese Academy of Agricultural Science, Beijing, PR China, 100193

^b Key Laboratory of Agro-Food Processing and Quality Control, Ministry of Agriculture, Beijing, PR China, 100193

^c Department of Agronomy and Horticulture, University of Nebraska Lincoln, Lincoln, NE 68583-0915, USA

^d School of Chemical Engineering and Technology, Tianjin University, Tianjin, PR China, 300072

^e Chinese Academy for Environmental Planning, Beijing, PR China, 100193

* Corresponding Author: Jia-ping Lv, phone 861062819421, fax: 861062815542, e-mail: lvjp586@vip.sina.com

References

- 1 A.D. Moordian, J.E. Morley, *Am. J. Clin. Nutr.* 1987, 45, 877–895.
- 2 J.B. Vincent, S.T. Love, *Chem Biodivers.* 2012, 9, 1923–41.
- 3 J. Racek, L. Trefil, D. Rajdl, et al., *Biol. Trace Elem. Res.* 2006, 109, 215–230.
- 4 M.H. Lai, Y.Y. Chen, H.H. Cheng, *Int. J. Vitam. Nutr. Res.* 2006, 76, 391–397.
- 5 L. Liu, W. Jin, J.P. Lv, *Biol. Trace Elem. Res.* 2010, 138, 250–264.
- 6 S.M. Bahijiri, S.A. Mira, A.M. Mufti, et al., *Saudi Med. J.* 2000, 21, 831–837.
- 7 J. Racek, C.D. Sindberg, S. Moesgaard, et al., *Biol. Trace Elem. Res.* 2013, 155, 1–4.
- 8 R.A. Anderson, M.M. Polansky, E.E. Roginski, et al. *Agric. Food Chem.* 1978, 26, 858–861.
- 9 K.H. Sumrall, J.B. Vincent, *Polyhedron* 1997, 16, 4171–4177.
- 10 J.B. Vincent, *J. Am. Coll. Nutr.* 1999, 18, 6–12.
- 11 T. Urumow, O.H. Wieland, *Horm. Metab. Res. Suppl.* 1984, 1, 51–4.
- 12 J. A. Vinson, P. Bose, *Nutrition Reports International.* 1984, 30, 911–918.
- 13 J. Racek, L. Trefil, D. Rajdl, et al., *Biol. Trace Elem. Res.* 2006, 109, 215–230.
- 14 M.H. Lai, Y.Y. Chen, H.H. Cheng, *Int. J. Vitam. Nutr. Res.* 2006, 76, 391–397.
- 15 L. Liu, J.P. Lv, H. Uluko, *J. Agric. Food Chem.* 2013, 61, 1279–1287.
- 16 C. L. Broadhurst, M.M. Polansky, R.A. Anderson, *J. Agric. Food Chem.* 2000, 48, 849–852.
- 17 R.A. Anderson, J.H. Brantner, M.M. Polansky, *J. Agric. Food Chem.* 1978, 26, 1219–122.
- 18 J.B. Vincent, *Acc. Chem. Res.* 2000, 33, 503–510.
- 19 S.A. Ross, X. Chen, H.R. Hope, et al., *Biochem. Bioph. Res. Co.* 2000, 273, 1033–1041.
- 20 S.E. Nissen, *Eur. Heart J.* 2010, 31, 773–776.
- 21 W.T. Cefalu, F.B. Hu, *Diabetes Care.* 2004, 27, 2741–2751.
- 22 A. Pechova, L. Pavlata, *Vet. Med-Czeh. Veterinarni Medicina.* 2007, 52, 1–18.
- 23 J.B. Vincent, S.T. Love, *Chem. Biodivers.* 2012, 9, 1923–1941.
- 24 M.J. Brady, *J. Clin. Invest.* 2004, 114, 886–888.
- 25 S. Weksler-Zangen, T. Mizrahi, I. Raz, et al., *Brit. J. Nutr.* 2012, 108, 875–882.
- 26 G.D. Christian, E.C. Knoblock, W.C. Purdy, et al., *Biochim. Biophys. Acta.* 1963, 66, 420–423.
- 27 J.B. Vincen, *J. Nutr.* 1994, 124, 117–118.

- 28 W. Qiao, Z. Peng, Z. Wang, et al., *Biol. Trace Elem. Res.* 2009, 131-133
- 29 Y.Q. Wang, M.H. Yao, *J. Nutr. Biochem.* 2009, 20, 982-991.
- 30 L. Martinez, M. Berenguer, M.C. Bruce, et al. *Biochem. Pharmacol.* 2010, 79, 1300-1309.