High Sucrose Low Magnesium Diet Modulates the Expression of PI3K and ERK2 in Different Tissues of Weanling Rats

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Abstract

Activation of the insulin receptor initiates signaling through both the phosphatidylinositol (PI) 3-kinase and the mitogen-activated protein kinase [MAPK, also referred to as extracellular signal-regulated kinases (ERK1/2)] pathways. The present study was undertaken to evaluate the effects of feeding high sucrose low magnesium diet to weanling rats on the insulin signal transduction pathways. To accomplish this, a non-genetic type 2 diabetic rat model was developed in the laboratory by feeding high sucrose+ Low Mg (HSLM) diet to weanling rats and the expression of the downstream effectors of insulin signaling pathway was studied. The Expression of ERK2 and PI3 Kinase was assessed in adipose tissue, skeletal muscle and liver of control and experimental diet fed rats. The results obtained clearly indicated a decrease in the expression of ERK2 and PI3 Kinase in all the three tissues from HSLM diet fed rats when compared to control rats, thereby confirming that Sucrose+ Low Mg feeding ultimately leads to insulin resistance via a decrease in the expression of the downstream effectors of the insulin signaling pathway.

Keywords: PI3K, ERK expression, Sucrose+ Low Mg, adipose tissue, skeletal muscle, liver.

1. Introduction

Diet composition has been proposed as one of the causes of the increasing prevalence of obesity, insulin resistance and the metabolic syndrome [1]. Increased access to diets enriched in fat or foods and beverages containing added simple sugars, e.g. sucrose or high fructose corn syrup lead to positive energy balance and therefore, fat or weight gain [1]. Dietary nutrients can also modulate insulin action. In rats, diets enriched in sucrose or fructose produce hepatic insulin resistance prior to adipose tissue and skeletal muscle, and independently of the changes in body composition [2]. Magnesium deficiency is...
frequently associated with changes in carbohydrate homeostasis [3]. Attempts have been made to correlate hypomagnesaemia and reduced erythrocyte Mg concentration with poor glyemic control and the development of complications in diabetic patients [4]. Because a low Mg concentration may be a consequence or a cause of insulin resistance [4], it may be premature to assign a primary role to Mg in abnormal carbohydrate metabolism.

Insulin action in target tissues is mediated by the heterotetrameric insulin receptor. Insulin signaling is initiated by the activation of insulin receptor tyrosine kinase, leading to the phosphorylation of insulin receptor substrates 1 and 2, thereby creating binding sites for the regulatory subunits of phosphatidylinositol 3-kinase (PI3-kinase) and Grb/Sos complex [5]. PI3-kinase appears to be particularly important for mediating most of the metabolic effects of insulin and its downstream signaling events [6]. PI 3-kinase is activated by the docking of its p85 subunit to phosphorylated IRS-1 and is linked to the activation of glucose transport, p70 S6 kinase activation, and nuclear DNA synthesis and gene expression. PI 3-kinase also activates protein kinase Akt (protein kinase B) and leads to glucose transporter 4 (GLUT4) vesicle translocation and glucose transport activation. On the other hand, Shc and the Grb2/Sos complex link insulin receptor signaling to activation of the mitogen-activated protein kinase [MAPK or extracellular signal-regulated kinase (ERK1/2)] cascade [7]. MAPK activity is increased when Grb2/Sos complex is recruited to the plasma membrane by an increase in the rate of exchange of guanosine triphosphate (GTP) for guanosine diphosphate (GDP) on Ras, which, in turn, activates MAPK kinase kinase (MEKK) leading to activation of MAPK kinase (MEK1), a threonine-tyrosine kinase that phosphorylates and activates ERK1 and ERK2 [8]. ERK2 is the most abundant MAPK isoform in skeletal muscle. Several studies have shown that activation of the PI3-kinase signaling pathway in response to insulin is reduced in muscles of obese and Type 2 diabetic subjects and in animal models of insulin resistance [9, 10]. One such animal model of insulin resistance is the non-diabetogenic HSLM diet fed rat model. The rats in this model are characterized by hyperglycemia, hyperinsulinemia and hypertriglyceridemia [11], conditions similar to those observed in the early stage of human Type 2 diabetes. Thus this HSLM diet fed rat model is ideal for the present study. Therefore, the present study was planned to raise a HSLM diet fed rat model by feeding a synthetic diet to weanling rats on the same basis as the adult rats [11] and to determine whether HSLM feeding results in change in the expression of ERK2 and PI3 Kinase in adipose tissue, skeletal muscle and liver.

2. Materials and Methods

Chemicals: TRIzol® reagent for total RNA isolation was procured from Invitrogen Corp, California, USA. Primers for ERK2 and PI3K were obtained from Sigma Chemicals Co., St. Louis, Mo., USA. M-MuLV RT-PCR Kit was purchased from Bangalore GeNei™, India. Kit for estimation of plasma glucose was obtained from Reckon Diagnostics Ltd., India. All other chemicals used were of analytical grade.

Animals and Diet: 24 Weanling male Wistar rats (bred in our facility), 21 day old and weighing about 30 g at the beginning of the study were randomly divided into four groups containing 6 rats each and were housed in polypropylene cages in a temperature-controlled room on a 12-h-light 12-h-dark cycle. The rats in the four groups were fed either a Starch+Normal Mg (Control) or Sucrose+Normal Mg (HS) diet or Starch+Normal Mg (LM) or Sucrose+Low Mg (HSLM) diet for a total period of 16 weeks. The diets were made up according to Table 1, and were fed to the rats in the form of pellets. Diets were freshly made up on a weekly basis and stored at 4°C. Rats in the Control and HS group had free access to tap water whereas rats in the LM and HSLM group had free access to double distilled water. All the institutional guidelines for the care of animals were followed strictly.

2.1 Biochemical Analyses

Estimation of plasma glucose: Blood was drawn from the supraorbital sinus of ether anaesthized rats once the animals had reached an appropriate age and weight. To attain this aim, rats
were fed different diets for a period of 6 weeks and blood was drawn every 2 weeks hence. Plasma was separated and the levels of plasma glucose were measured by the method of Trinder [12] using glucose oxidase. Briefly, 10µl of plasma was added to 1 ml of working reagent.

Equal volumes of glucose standard and distilled water were added respectively to tubes marked standard and blank, containing 1ml of the working reagent. The contents of the tubes were then mixed well and incubated for 15 min at 37°C in a water bath. Colour developed was read at 510nm.

Table 1: Experimental diet composition in g/kg diet &

<table>
<thead>
<tr>
<th>Ingredient</th>
<th>Starch +Normal Mg diet</th>
<th>Starch+Low Mg diet</th>
<th>Sucrose+Normal Mg diet</th>
<th>Sucrose +Low Mg diet</th>
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</thead>
<tbody>
<tr>
<td>Sucrose</td>
<td>-</td>
<td>-</td>
<td>658</td>
<td>658</td>
</tr>
<tr>
<td>Starch</td>
<td>658</td>
<td>658</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Casein</td>
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<td>188</td>
<td>188</td>
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<tr>
<td>Methionine</td>
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<td>1.9</td>
<td>1.9</td>
<td>1.9</td>
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<tr>
<td>Gelatin</td>
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<td>14.1</td>
<td>14.1</td>
<td>14.1</td>
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<tr>
<td>Safflower oil</td>
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<td>45.4</td>
<td>45.4</td>
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<tr>
<td>Bran</td>
<td>37.6</td>
<td>37.6</td>
<td>37.6</td>
<td>37.6</td>
</tr>
<tr>
<td>Mineral Mix#*</td>
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<tr>
<td>Vitamin Mix#*</td>
<td>9.4</td>
<td>9.4</td>
<td>9.4</td>
<td>9.4</td>
</tr>
</tbody>
</table>

*By calories, the diets were 69% carbohydrate, 21% protein, and 10% fat.

#Supplied per kilogram of vitamin mix: 3 g thiamine mononitrile, 3 g riboflavin, 3.5 g pyridoxine HCl, 15 g nicotinamide, 8 g d-calcium pantothenate, 1 g folic acid, 0.1 g d-biotin, 5 mg cyanocobalamin, 500 000 IU (12.5 mg) cholecalciferol, 25 mg acetonenaphthone, 2 000 000 IU (600 mg) vitamin A acetate, 25 000 IU (22 g) d-a-tocophenyl acetate and 10 g choline chloride.

*Supplied in per kilogram of mineral mix: 30.5 g MgSO₄·7H₂O, 65.2 g NaCl, 105.7 g KCl, 200.2 g KH₂PO₄, 38.8 g MgCO₃, 3H₂O, 4.0 g FeCH₂O₂·5H₂O, 512.4 g CaCO₃, 0.8 g KI, 0.9 g NaF, 1.4 g CuSO₄·5H₂O, 0.4 g MnSO₄ and 0.05 g CoNH₃₃·6H₂O.

* A similar composition of mineral mixture was used in all the experimental groups, except for the addition of MgO and MgSO₄·2H₂O to provide (per kg) 507.0 mg of Mg in the control and high sucrose diets and 40.0 mg of Mg in the low magnesium and the high sucrose low magnesium diets.

Table 2: PCR primers and Programs for quantitation of RNA*.

<table>
<thead>
<tr>
<th>cDNA</th>
<th>Forward Primer</th>
<th>Reverse Primer</th>
<th>PCR Program</th>
</tr>
</thead>
<tbody>
<tr>
<td>β-actin</td>
<td>5‘CTGACCGAGCTGGCTAC 3’</td>
<td>5’ CCTGCTTGCTGATCCACA 3’</td>
<td>94/30/55/30/72/45/22</td>
</tr>
<tr>
<td>ERK2</td>
<td>5’TGAAGACACAGCCTCAGC 3’</td>
<td>5’ GTCACGGTGACAGAAGTTAG 3’</td>
<td>94/30/60/30/72/60/40</td>
</tr>
<tr>
<td>PI3K</td>
<td>5’ TTAACCGCGAAGGCAACGA 3’</td>
<td>5’ CAGTCTCCTCCTGCTGATCAG 3’</td>
<td>94/15/60/60/72/60/40</td>
</tr>
</tbody>
</table>

*Primers were designed using the known sequences for the respective genes. Programs are given as denaturation temperature (°C)/denaturation time(s)/annealing temperature (°C)/annealing time(s)/elongation temperature (°C)/elongation time(s)/number of cycles. Additionally, all programs started with a period of 94°C for 10 min and finished with 5 extra min at 72°C. For example, the program for β-actin was 94°C for 10 min followed by 22 cycles of 94°C for 30 sec, 55°C for 30 sec, 72°C for 45 sec, and ending with 5 min at 72°C.
Estimation of magnesium: At the end of 6 weeks of diet feeding, magnesium levels were determined every 2 weeks [13]. Suitable volumes of plasma were mixed well with 2 ml of the working reagent in small tubes and read immediately at 570 nm against a blank containing equal volumes of double deionized water and working reagent. A range of appropriate standards were run similarly.

2.2 Tissue Preparation and RNA isolation

At the end of the 4 months of diet feeding the overnight fasted rats were sacrificed by cervical dislocation and the liver, skeletal muscle from the thigh and the adipose tissue from the epididymal fat pad was separated, washed in ice cold normal saline and then used for Total RNA isolation. Total RNA was extracted using a commercially available reagent TRizol® (Invitrogen Corp, California, USA). Briefly, 50–100 mg of tissue was homogenized in 1 ml of TRizol® reagent and processed according to Kit manufacturer’s instructions. After that the concentration of total RNA was measured by spectrophotometry (Beckman Instruments, Fullerton, CA). The samples were stored in -80°C freezer for carrying out RT-PCR study.

2.3 RT-PCR

The following primer sets were used for RT-PCR. Rat ERK2 forward primer: 5’tgaagacacagcactcag 3’, reverse primer: 5’ gtcaggtgagaacccctag 3’, rat PI3K forward primer: 5’ taaaeggaagggcaacga 3’, reverse primer: 5’ cagtcctccctgctgtag 3’ and beta actin forward primer: 5’ctgacgcgagctggctacc 3’and reverse primer: 5’ cctgcttgctgatccacca3’. We used M-MuLV RT-PCR Kit (GeNei™, Bangalore, India) according to the manufacturer’s instructions. Oligo d(T) primers were used with this kit to reverse transcribe the mRNA from total RNA. cDNA was produced with 1 µg of RNA as template. The gene-specific primers (Table 2) were designed by using available gene sequences. The oligomers were purchased from Sigma Chemicals Co. (St. Louis, Mo., USA). PCRs were run on a MiniCycler (Eppendorf, Germany). The programs (Table 2) were optimized for each primer pair to get a linear proportion between the amount of template used and the amount of product after the PCR reaction. All programs started with a 10-min period at 94°C and ended with a 5-min extra elongation period at 72°C. The PCR products were run on 2% agarose gels (Sigma). Gels were stained with ethidium bromide, visualized by ultraviolet light and photographed using Gel Doc XR System (BioRad). The bands were identified and quantified using the Multi-Analysis software of Gel Doc XR system. The mRNA level of each enzyme was expressed as a ratio of each enzyme/ β-actin RNA. All the reactions were performed thrice.

Table 3: Percentage downregulation of MAPK as calculated by the normalization of the test with that of the control gene.

<table>
<thead>
<tr>
<th>Enzyme</th>
<th>% Downregulation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Adipose MAPK (LM)</td>
<td>12.9</td>
</tr>
<tr>
<td>Adipose MAPK (HS)</td>
<td>10.12</td>
</tr>
<tr>
<td>Adipose MAPK (HSLM)</td>
<td>16.3</td>
</tr>
<tr>
<td>Muscle MAPK (LM)</td>
<td>9.8</td>
</tr>
<tr>
<td>Muscle MAPK (HS)</td>
<td>6.14</td>
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<td>Muscle MAPK (HSLM)</td>
<td>12.8</td>
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<tr>
<td>Liver MAPK (LM)</td>
<td>26.6</td>
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<tr>
<td>Liver MAPK (HS)</td>
<td>21.2</td>
</tr>
<tr>
<td>Liver MAPK (HSLM)</td>
<td>32.1</td>
</tr>
</tbody>
</table>

Table 4: Percentage downregulation of PI3K as calculated by the normalization of the test with that of the control gene.

<table>
<thead>
<tr>
<th>Enzyme</th>
<th>% Downregulation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Adipose PI3K (LM)</td>
<td>25.1</td>
</tr>
<tr>
<td>Adipose PI3K (HS)</td>
<td>15.9</td>
</tr>
<tr>
<td>Adipose PI3K (HSLM)</td>
<td>31.4</td>
</tr>
<tr>
<td>Muscle PI3K (LM)</td>
<td>39.2</td>
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<tr>
<td>Muscle PI3K (HS)</td>
<td>30.4</td>
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<tr>
<td>Muscle PI3K (HSLM)</td>
<td>40.5</td>
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<tr>
<td>Liver PI3K (LM)</td>
<td>28.4</td>
</tr>
<tr>
<td>Liver PI3K (HS)</td>
<td>5.7</td>
</tr>
<tr>
<td>Liver PI3K (HSLM)</td>
<td>38.4</td>
</tr>
</tbody>
</table>

2.4 Statistical analysis

Statistical analysis was performed using GraphPad InStat (GraphPad Inc., San Diego, CA, USA) software package. Results were expressed as mean and SD of six observations in each group. Further, the statistical significance of the
differences among the various dietary groups was determined by subjecting the data to two way ANOVA with diet as the main effect, followed by inspection of all differences between pairs of means by Post Hoc test. Differences were considered statistically significant at $p < 0.05$.

In all cases $p<0.001$
* Significantly different from the initial value, $^\$$ Significantly different from the control group, $^@$Significantly different from the LM group, $^@$Significantly different from the HS group.

**Fig. 1:** Levels of plasma glucose as measured at 6th to 16th week of diet feeding. The figure clearly shows that rats fed HSLM diet had increased plasma glucose levels as compared to the control rats.

In all cases $p<0.001$
* Significantly different from the initial value, $^\$$ Significantly different from the control group, $^@$Significantly different from the HS group.

**Fig. 2:** Plasma magnesium levels at 6th to 16th week of diet feeding. The figure indicates a marked decrease in the plasma magnesium content in rats fed HSLM diet in comparison to control rats.
**Fig. 3:** Agarose gel electrophoresis of RT-PCR products of ERK2 and β-actin from adipose tissue, skeletal muscle and liver of Control, LM, HS and HSLM diet fed rats.
Fig. 4: Agarose gel electrophoresis of RT-PCR products of PI3K and β-actin from adipose tissue, skeletal muscle and liver of Control, LM, HS and HSLM diet fed rats.
3. Results

3.1 Animal model preparation by feeding High Sucrose Low Magnesium diet to rats

Levels of Plasma glucose and magnesium were monitored on a biweekly basis after feeding rats the experimental diets for a period of 4 weeks. Results obtained clearly show marked hyperglycemia and hypomagnesemia in rats fed HSLM diet when compared to rats fed a control diet (Table 3, 4; Fig 1, 2).

3.2 RNA Isolation and RT-PCR:

At the end of 4 months of diet feeding the overnight fasted rats were sacrificed and RNA was isolated from adipose tissue, skeletal muscle and liver. Total RNA was then subjected to RT-PCR analysis and the products thus obtained were run on a 2% agarose gel. Results obtained clearly show a marked decrease in the density of the cDNA products from the tissues of rats fed a high sucrose low magnesium diet when compared to that of control rats indicating a downregulated expression of both ERK2 and PI3K in all the three tissues. (Fig. 3-6).

Fig. 5: MAPK/ β-actin ratio in adipose, skeletal muscle and liver tissue of rats fed different diets.

3.3 Change in expression

Percentage change in the expression of both the enzymes was calculated. Our results clearly show a significant percentage downregulation of both of the kinases in all the three tissues in rats fed HSLM diet when compared to those fed a control diet.

Fig. 6: PI3K/ β-actin ratio in adipose, skeletal muscle and liver tissue of rats fed different diets.

4. Discussion

Feeding rats a high sucrose low magnesium diet leads to hyperglycemia, hypertriglyceridemia, hyperinsulinemia [11], insulin resistance, and if continued long enough, impaired glucose tolerance or diabetes[14, 15]. The present study was carried out to develop an insulin resistant rat model by feeding HSLM diet to weanling rats and to study the expression of the downstream effectors of the insulin signaling pathway, viz. ERK2 and PI3 Kinase in this rat model. To achieve this, rats in different groups were fed respective diets and the plasma concentration of glucose and magnesium were monitored on a biweekly basis for 12 weeks to observe any difference as compared to control rats. Our results clearly show that sucrose overfeeding and inadequate magnesium intake in experimental animals resulted in hyperglycemia [p< 0.05] in both low magnesium (LM) as well as high sucrose (HS) groups. The animals in the group fed HSLM diet also had elevated plasma glucose levels, with the likelihood that both carbohydrate (sucrose)
overfeeding and hypomagnesaemia contribute to the abnormal glucose levels in the plasma of experimental rats. Studies have illustrated the detrimental effects of sucrose rich diets on glucose homeostasis [16] and it is possible that the hyperglycemia associated with sucrose-rich diets might be due partially to several reasons including perturbations in the activities of crucial enzymes leading to a change in flux through the pathways. Magnesium deficient diets are implicated in causing hypomagnesaemia (in approximately 3 months of diet feeding [11]). It has been observed that omission of extracellular magnesium inhibited glucose stimulation of insulin homeostasis in isolated islet cells [17]. The mechanisms of this inhibition could be linked to the ability of glucose and several other stimulators of insulin biosynthesis to increase magnesium uptake by islet cells [18, 19]. Our results on the concentration of magnesium in the plasma of experimental animals are presented in Fig 2. Plasma magnesium levels in HS diet group were comparable to control in our study indicating that feeding a low magnesium diet, irrespective of the carbohydrate component of the diet, resulted in a marked lowering [p<0.05] of plasma magnesium levels. Our results are consistent with the previous studies reporting significantly decreased plasma magnesium levels in rats fed a low magnesium diet [20-22].

Once the model was established, the expression of PI3 kinase and MAP Kinase (ERK2) was studied in adipose, skeletal muscle and liver tissues of rats. Our results clearly show a significant decrease in the expression of both kinases in all the three tissues in HSLM diet fed rats when compared to control rats. Our findings are consistent with the studies showing that both basal and insulin stimulated ERK2 activity are markedly reduced in obese rats [23]. Reports clearly demonstrate that insulin receptor signaling through the PI3-Kinase pathway is reduced in the Zucker rat model of insulin resistance and the same has been shown in our model of HSLM diet fed rats.

Insulin is known to be an important modulator of the cellular content of Mg [24,25]. Since Mg is a necessary cofactor in all ATP transfer reactions, this implies that [Mg2+] concentration is critical in the phosphorylation of the insulin receptor [26]. Insulin binding to specific cell surface receptors leads to the activation of protein kinase which is an important step in transmembrane signaling for insulin action [27]. There are several examples where alteration in receptor kinase activity could explain an impairment of insulin action [28]. Insulin receptors isolated from various tissues of type 2 diabetics or obese subjects have an impaired capacity to autophosphorylate or express the tyrosine kinase activity when exposed to insulin [29]. Suáres et al. suggested that the insulin resistance observed in the skeletal muscles of magnesium-deficient rats might be attributed to the defective tyrosine kinase activity of the insulin receptor [30]. Studies in multiple insulin resistant cell models have demonstrated that an impaired response of the tyrosine kinase to insulin stimulation is one potential mechanism causing insulin resistant-state in type 2 diabetes [31]. Similarly, a depletion of [Mg2+] may cause a defective tyrosine kinase function at the insulin receptor level.

A decreased concentration of [Mg2+] is associated with a diminution in the ability of insulin to stimulate glucose uptake in insulin-sensitive tissues, such as adipose cells and skeletal muscle tissues [32]. Given in vitro evidence that low Mg concentrations can reduce tissue glucose uptake [33, 34], it seems that reduced [Mg2+] interferes with the insulin signaling mechanism involved in glucose transport [35]. Altered [Mg2+] may also lead to decreased cellular glucose utilization, thus promoting peripheral insulin resistance with a postreceptor mechanism.

High sucrose feeding has long been implicated in leading to insulin resistance at some point of time and the most damage has been attributed to the fructose component of sucrose. Studies of fructose fed models of insulin resistance show that despite the normal number of insulin receptors in the liver and muscle of fructose-fed rats, there was a 29% reduction in insulin receptor autophosphorylation in the liver after stimulation with insulin “in vivo.” The decrease observed here may be of biological importance because a reduction in receptor phosphorylation has been correlated with insulin resistance in different animal models [34-36].
There was a significant reduction in the level of hepatic and muscle IRS-1 tyrosine phosphorylation, followed by a reduction in IRS-1/PI3-K 3-kinase association in liver and muscle. In previous studies of fructose-fed rats, a reduced ability of insulin to suppress hepatic glucose production has been implicated as one of the elements of insulin resistance [37, 38]. The IRS-1/PI3-K 3-kinase association induced by insulin is necessary, and in some cases sufficient to elicit many of the insulin effects on glucose and lipid metabolism. The products of PI3-kinase act as both membrane anchors and allosteric regulators, serving to locate and activate downstream enzymes and their protein substrates [39]. Thus, the reduction in IRS-1/PI3-K 3-kinase association in the liver of fructose-fed rats may have a role in insulin resistance.

On the basis of results of this study, we conclude that feeding high sucrose low magnesium diet to weanling rats resulted in marked hyperglycemia as well as hypomagnesaemia which are hallmarks of type 2 diabetes. This suggests that we have been able to develop a type 2 diabetic rat model by feeding HSLM diet for a period of 4 months. Our results also show a decreased expression of PI3 Kinase and ERK2 in the adipose, skeletal muscle and liver tissues of HSLM diet fed rats when compared to control, although the mechanism leading to this is not yet clear and paves way for future studies.

Abbreviations

MAPK Mitogen-Activated Protein Kinase
ERK Extracellular Signal-Regulated Kinase
PI3 K Phosphatidylinositol- 3-Kinase
IRS Insulin Receptor Substrate
GLUT Glucose Transporter
RT-PCR Reverse Transcriptase- Polymerase Chain Reaction

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