

Prophylactic efficacy and possible mechanisms of oligosaccharides based standardized fenugreek seed extract on high-fat diet-induced insulin resistance in C57BL/6 mice

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ABSTRACT

The present work was aimed to study the efficacy and possible mechanism of oligosaccharides based standardized fenugreek seed extract (SFSE-OS) on high-fat diet (HFD)-induced insulin resistance in male C57BL/6 mice. The effects of 12 weeks of oral administration of SFSE-OS (30, 60 and 100 mg/kg, twice daily) were evaluated on HFD fed mice for anthropomorphic, glycemic, gene expression related and histopathological parameters. Separate groups of mice with vehicle co-administered with HFD and low-fat diet (LFD) were maintained as HFD control and LFD control respectively. Twelve weeks of SFSE-OS (60 and 100 mg/kg, p.o.) administration showed significant prophylactic effects on HFD induced insulin resistance in terms of body weight, plasma glucose and insulin levels, glycated hemoglobin, insulin resistance (IR), area under the curve (AUC) of plasma glucose during oral glucose tolerance and intraperitoneal insulin tolerance. Furthermore, HFD-induced mRNA expression changes in adipose tissue, liver and skeletal muscle were prevented by SFSE-OS co-administration. Histology of sections of the pancreas showed the normal architecture in all groups of mice. SFSE-OS showed promising efficacy in prevention of HFD-induced insulin resistance through modulation of Glut-2, Glut-4, IRS-2 and SREBP-1c expression.

INTRODUCTION

Diabetes mellitus (DM) is the most common endocrine disorder of the 21st century. The prevalence of DM is more than 194 million people worldwide (Liao *et al.*, 2010) and estimated to increase to 333 million by 2025 (Zhu *et al.*, 2010). Lifestyle modification measures such as diet control and exercise have been found useful in short term control of body weight and T2D management. However, long-term solution is still elusive. Existing oral anti-diabetic medications including insulin sensitizers and exogenous insulin therapy address hyperglycemia but leads to many side effects such as increased body weight, osteoporosis sodium retention, hypoglycemia and lactic acidosis (Chiang *et al.*, 2007, Hamza *et al.*, 2010, Stades *et al.*, 2004). The type 2 DM (T2DM) is characterized by insulin resistance in

target tissues due to both insulin action and insulin secretion with subsequent pancreatic β -cell dysfunction (Nyenwe *et al.*, 2011). Insulin resistance (IR) is a physiological condition in which cells fail to respond to the normal actions of the hormone insulin.

The body produces insulin, but the cells in the body become resistant to insulin and are unable to use it effectively, leading to hyperglycemia. High plasma levels of insulin and glucose due to IR are a major component of the metabolic syndrome. The association between obesity and IR is well established (Hardy *et al.*, 2012, Masuo *et al.*, 2010).

IR is undoubtedly known as a hallmark of feature of obesity that compromises function of the pancreatic β -cell so as to cause insufficient insulin secretion and leads to T2DM (Das *et al.*, 2013, McGarry, 2002). Strongly association between IR and microvascular (Groop *et al.*, 2005), cardiovascular (Fontbonne, 1996) and renal (Masuo *et al.*, 2010) complications of DM and obesity is also established.

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Therefore, safer medications with effective control on both arms (blood glucose and IR) in DM are in need (Turner *et al.*, 1998). In recent years, many new plant derived natural products have demonstrated the potential for treatment of T2DM and its complications (Matsui *et al.*, 2006). One such promising plant is Fenugreek (*Trigonella foenum-graecum* L., family: Fabaceae). Plethora of evidence confirmed beneficial effects of fenugreek seeds powder and extract on glucose and fat metabolism in animal models and patients of DM (Roberts, 2011). Fenugreek seeds reported to cause improvement in glycemic control by decreasing IR in mild T2DM patients (Gupta *et al.*, 2001). Fenugreek seed extract in various forms demonstrated promising effects on high-fat (Chaturvedi *et al.*, 2013, Hamza *et al.*, 2012a, Handa *et al.*, 2005) and high sucrose (Muraki *et al.*, 2012) diet-induced hyperlipidemia. However, the responsible component of fenugreek seed extract and its mode of action for beneficial effects on IR not been elucidated.

Among all bioactive phytoconstituents of fenugreek seeds, only component that showed efficacy towards effective restoration of both glucose and fat metabolism in DM is its fibre component (Roberts, 2011)). The endosperm of the fenugreek seed is a rich source of fiber (20%) and gum (32.4 %) (Sharma, 1986, Sharma *et al.*, 1990) which include water soluble (low molecular weight) and insoluble (high molecular weight) fibres. Interestingly, only water soluble low molecular weight soluble fibre fraction of fenugreek seeds demonstrated excellent efficacy in glucose and lipid management in animal and human studies (Ali *et al.*, 1995, Evans *et al.*, 1992). Recently, we have reported promising and dose-dependent anti-hyperglycemic effects of water-soluble low molecular weight soluble fibre fraction in diabetic mice (Kamble *et al.*, 2013).

The soluble fibres of fenugreek seeds contain oligosaccharides along with Pinitol (methylated cyclic alcohol derivative of sugar). Oliosugers with structural similarity with a synthetic anti-diabetic oligosaccharide compound, acarbose (an α -glucosidase inhibitor). Oligosaccharides from various natural sources have been shown anti-diabetic potential in animal studies by effective control of glucose and fat metabolism (Costa *et al.*, 2012, Jo *et al.*, 2014, Lee *et al.*, 2003, Zhang *et al.*, 2004). Furthermore, eleutheroside E (an oligosaccharide component of *Eleutherococcus senticosus*) was shown to ameliorates IR in animal model of type 2 diabetic db/db mice (Ahn *et al.*, 2013). On the other hand, Pinitol obtained from other plant sources is reported to have good potential against IR for maintenance of glycemic control (Dang *et al.*, 2010, Hernandez-Mijares *et al.*, 2013, Kim *et al.*, 2012, Kim *et al.*, 2005a) through alternations in glucose (Kang *et al.*, 2006, Kim *et al.*, 2007) and fat metabolism (Choi *et al.*, 2009, Geethan and Prince, 2008). The combination of naturally occurring oligosaccharide with pinitol in fenugreek seeds offers a unique option for management of insulin resistance (IR) and effective glycemic control (Watson and Preedy, 2010).

Recently, effects of oliosugers based standardized fenugreek seeds extract (SFSE-OS) containing oligosaccharide (with eleutheroside-C, methyl- α -D-galactoside), Pinitol

(methylated cyclic alcohol derivative of sugar) has been reported against experimental neuropathy models in rats (Bhaskaran and Vishwaraman, 2014). However, potential of SFSE-OS to achieve effective control of glycemia and/or IR is not been explored yet. Therefore, we undertook present study with an objective to evaluate efficacy and probable mechanism of action SFSE-OS on high-fat diet (HFD)-induced IR in C57BL/6 mice.

METHODS

Animals

Male C57BL/6 mice (age 4-weeks, 20-22 g) were used for the study. They were housed in cages at a temperature of 19.1-23.6 °C, and relative humidity 56- 66 %, with 12 h fluorescent light and 12 h dark cycle in an accredited animal facility (Bioneed, Bangalore, India). The mice had free access to water ad libitum throughout the study duration except during actual measurements. All experiments were carried out between 09:00 h and 17:00 h. Institutional Animal Ethics Committee (IAEC) of Bioneed (Bangalore, India) approved the experimental protocol (BIO-IAEC-477). The experiments were performed in accordance with the guidelines on animal experimentation recommended by the Committee for Control and Supervision of Experimentation on Animals (CPCSEA), Government of India.

The test compound

The test compound, SFSE-OS supplied by Indus Biotech Private Limited (Pune, India) after preparation and characterization (HPLC and LC-MS) as per reported procedure (Bhaskaran and Vishwaraman, 2014). The sample of SFSE-OS used for the present study was standardized to Eleutheroside-C (30%), Pinitol (48%), Sucrose (6%), Raffinose (6.5%), and Stachyose (9.5%). As SFSE-OS was water soluble, the solution was freshly prepared daily as 1% w/v in distilled water and orally administered twice a day to mice in volume of 10 mg/kg. The doses of SFSE-OS (30, 60 and 100 mg/kg, twice a day, oral) were derived from LD50 dose of 2000 mg/kg found during acute oral toxicity (AOT) study (Bodhankar *et al.*, 2013). The vehicle or SFSE-OS were orally administered twice daily with difference of approximately 8 h between morning and evening administration using intragastric tube for 12 weeks.

Research diets

The Low fat diet, LFD (10 kcal % Fat, #D12450B, 3.85 kcal/g) and HFD (60 kcal %fat, # D12492, 5.24 kcal/g) were purchased from Research Diet Inc., New Brunswick, NJ , USA.

Development of HFD induced IR

Mice were randomly divided into five groups of twelve mice each and administrated with treatments and diets as follows: G1 (LFD control): received vehicle (10 mg/kg of distilled water, twice daily); G2 (HFD control): received vehicle (10 mg/kg of distilled water, twice daily) + HFD; G3 (SFSE-OS (30): received SFSE-OS (30 mg/kg of SFSE-OS, twice daily) + HFD; G4 (SFSE-

OS (60) : received SFSE-OS (60 mg/kg of SFSE-OS, twice daily) + HFD; G5 (SFSE-OS (100): received SFSE-OS (100 mg/kg of SFSE-OS, twice daily) + HFD. The mice were caged individually and respective diet was supplied during entire study period of 12 weeks.

Body weights of individual mouse were recorded before grouping and three times a week thereafter, during the treatment period. At end of treatment period of 12-weeks, 100 μ L blood were withdrawn from the retro-orbital plexus and used for glycated hemoglobin (HbA1c) levels measurement using mouse hemoglobin A1c kit (Crystal Chem, Inc., Downers Grove, IL, USA) as per manufacturer's instructions.

Oral Glucose Tolerance Test (OGTT)

On the next day of HbA1c measurement, mice were fasted for 12 h and OGTT was performed as per reported procedure (Gallou-Kabani *et al.*, 2007). At 0 h (before administration of glucose) fasting plasma glucose (FPG) was measured using Accu-Chek Performa Blood Glucose Meter (Roche Diagnostics India Private Ltd, Mumbai, India) and glucose load (2 g/kg) was administered orally. Blood was withdrawn at 30, 60, 120, and 180 min, plasma was obtained after centrifugation and PG measurements were done. Additional blood was collected at 0 and 120 min, plasma was separated and used for insulin measurement with mouse insulin ELISA kit (ALPCO Diagnostics, Salem, NH, USA).

From the values of FPG and fasting insulin, the HOMA-IR was calculated as per reported formula (Matsuda and DeFronzo, 1999). The graph of PG levels versus time (min) was plotted and area under the OGTT curve (AUC-OGTT) was calculated with the help of Prism v6.0 (GraphPad Software Inc, La Jolla, CA, USA).

Intraperitoneal Insulin Tolerance Test (IPITT)

Three days after OGTT, 6 mice per group were selected for IPITT. Mice were fasted for 4 h and FPG was measured using Accu-Chek Performa Blood Glucose Meter (Roche Diagnostics India Private Ltd). After 30 min, mice were administered intraperitoneally with insulin (0.5 IU/kg, Actrapid[®], Novo Nordisk India Pvt Ltd, Bangalore, India). PG was measured at 30, 60, and 120 min from insulin administration. The graph of PG levels versus time (min) was plotted and area under the IPITT curve (AUC-IPITT) was calculated with the help of Prism v6.0 (GraphPad Software Inc, La Jolla, CA, USA).

Gross Necropsy

At end of the experimental period, all mice were sacrificed using CO₂ asphyxiation and subjected to gross necropsy. Four samples of adipose tissues, liver, pancreas and skeletal muscle from each group were preserved in liquid nitrogen for gene expression analysis by reverse transcriptase-polymerase chain reaction (RT-PCR). Two samples of pancreas from each group were preserved in 10% neutral buffered formalin for histopathology.

Gene expression analysis by RT-PCR

Four tissue samples of adipose tissue, liver, pancreas and skeletal muscle from each group was used for RT-PCR analysis using kits (Biotools B & M Labs, Spain) according to manufacturer's instructions. Total RNA was extracted from each tissue. The PCR mixture was amplified in a DNA thermal cycler (Eppendorf India Ltd, Chennai, India) by using gene specific primers (Table 1). PCR products were run on 1 % agarose gels, stained with ethidium bromide. The expression of all the genes was assessed by generating densitometry data for band intensities in different sets of experiments and was generated by analyzing the gel images on the Image J program (Version 1.33, Wayne Rasband, National Institutes of Health Bethesda, MD, USA). The band intensities were compared with constitutively expressed β -actin which served as a control for sample loading and integrity. The intensity of mRNAs was standardized against that of the β -actin mRNA from each sample, and the results were expressed in a quantitative manner as PCR-product/ β -actin mRNA ratio.

Histopathology

The samples from pancreas were embedded in paraffin wax, sectioned at 4 μ m and stained with haematoxylin and eosin (H&E) and observed under compound microscope using a Zeiss intravital microscopy setup (Zeiss Axioscope A1, Carl Zeiss MicroImaging, Jena, Germany) with a 40X magnification for histological examinations.

Statistical analysis

All statistical analysis was performed using GraphPad Prism 6.0 (GraphPad Software, Inc, La Jolla, CA, USA). Data of body weight and biochemical measurements was analyzed by separate One-way ANOVA followed by Dunnett's test separately for each parameter. A value of $P < 0.05$ was considered to be statistically significant.

RESULTS

Effect on body weight and Glycemic parameters

The data of effects of treatments on body weight and glycemic parameters (fasting glucose level, OGTT and IPITT, plasma insulin, HOMA-IR and HbA1c levels) are presented in Table 2. At the end of study (12-weeks), significant gain in the body weight ($P < 0.05$) and HbA1c ($P < 0.001$) was observed in HFD control mice as compared to LFD control mice (Table 2). The mean body weights in LFD fed mice was 20.66 g whereas that of HFD control mice were 33.40 g (gain of increase 4.74 g., 17.7%).

The body weights of HFD mice co-administered with SFSE-OS (60) group showed significant ($P < 0.01$) reduction by 3.73 g (15.17%) as compared to HFD control group. At the end of 12-week of study period, HFD control group showed significant ($P < 0.001$) increase in mean HbA1c values of 5.43% as compared to HbA1c values of LFD control mice (2.94%). Mean HbA1c levels of HFD mice co-administered with SFSE-OS (30, 60 and

100 mg/kg, p.o.) were 4.11%, 2.96% and 3.13% respectively. These values were significantly ($P < 0.001$) less as compared to HFD control mice (5.43%).

Effect of SFSE-OS during OGTT

At the end of study (12-weeks), during OGTT, the LFD fed mice showed values of FPG (114.44 mg/dl), fasting plasma insulin (0.36 $\mu\text{g/L}$), 2 h post-glucose plasma insulin (0.52 $\mu\text{g/L}$), IR (0.11) and AUC-OGTT (31998.30) (Table 2). In the same period, HFD control mice showed significantly ($P < 0.001$) higher values of FPG (175.44 mg/dl, 53.30% increase), fasting plasma insulin (1.12, 211.11% increase), IR (0.51, 363.63% increase), AUC-OGTT (42966.70, 34.28% increase) respectively. However, the values of 2 h post-glucose plasma insulin (0.73 $\mu\text{g/L}$) was not significantly different as compared with corresponding value of LFD control mice during OGTT.

At the end of study (12-weeks), during OGTT, HFD fed mice co-administrated with SFSE-OS (30, 60 and 100 mg/kg, p.o.) showed significantly ($P < 0.001$) less values of FPG levels of 129.33, 119.83 and 125.67 mg/dl respectively. These values were 26.28%, 31.69 % and 28.36 % less as compared with plasma glucose values of HFD control mice.

Fasting insulin levels of SFSE-OS (60 and 100 mg/kg, p.o.) co-administrated HFD mice showed significantly less values of 0.70 and 0.57 $\mu\text{g/L}$ (37.5% and 49.10% less) as compared to that of HFD control mice. However, fasting plasma insulin levels of SFSE-OS (30 mg/kg) co-administered mice show values of 0.89, (20.53% less) which was not significant as compared HFD control mice. The values of 2 h post-glucose plasma insulin was not significantly different in any of SFSE-OS co-administrated group as compared to HFD control mice.

During OGTT, the mean value of IR of SFSE-OS (60 and 100 mg/kg, p.o.) co-administrated mice was 0.23 (54.90 % less) and 0.19 (62.74 % less) and were significantly lower ($P < 0.05$ and 0.01 respectively) as compared to mean values of IR of HFD control mice (0.51). However, IR of SFSE-OS (30 mg/kg) co-administrated mice (0.33, 35.29% less) showed no significant difference as compared HFD control mice (0.51).

The values of AUC-OGTT in SFSE-OS (60 and 100 mg/kg) co-administrated mice were 33290 and 36336.30 respectively. These values were significantly ($P < 0.001$ and $P < 0.01$) less by 22.52% and 15.43% as compared to that of HFD control group. However, SFSE-OS (30 mg/kg) co-administrated mice showed reduction of 6.74% in AUC-OGTT which was not significant as compared with that of HFD control group (42967).

Effect of SFSE-OS during IPITT

The effects of treatments on AUC-IPITT are presented in Table 2. On insulin (0.5 U/kg, i.p.) load, AUC-IPITT of control mice was 17227.50, which was significant ($P < 0.001$) rise (103.0 %) in AUC-IPITT over corresponding value of LFD control group (8482.50). SFSE-OS (30, 60 and 100 mg/kg, p.o.) co-administrated mice showed AUC-IPITT values of 11350.00, 9822.50 and 12795.00 respectively which are 34.12 %, 42.98 %

and 25.73 % less as compared to corresponding values observed in HFD control mice (17227.50).

Effect on mRNA expressions in adipose tissues

The effects of SFSE-OS on mRNA expressions of Glut-2, Glut-4, IRS-2 and SREBP-1c in adipose tissue are presented as Fig.1. There was significant down-regulation in the mRNA expressions of Glut-2, Glut-4 and IRS-2 in adipose tissues of HFD control mice ($P < 0.01$, $P < 0.05$ and $P < 0.001$ respectively) as compared with LFD control mice. Treatment with SFSE-OS (100 mg/kg, p.o.) significantly up-regulated the mRNA expression of Glut-2 ($P < 0.01$) in adipose tissue as compared to HFD control mice. The mRNA expression of Glut-4 in adipose tissue was significantly up-regulated ($P < 0.01$ and $P < 0.01$, resp.) in SFSE-OS (60 and 100 mg/kg, p.o.) treated mice as compared to HFD control mice. As compared with HFD control mice, the mRNA expression of IRS-2 in adipose tissue was significantly up-regulated ($P < 0.01$) by SFSE-OS (100 mg/kg, p.o.) treatment.

On the other hand, SFSE-OS (30 mg/kg, p.o.) failed to produce any significant up-regulation in mRNA expression of Glut-2, Glut-4 and IRS-2 in adipose tissue when compared with HFD control mice (Fig. 1A, 1B, 1C and 1D). The level of SREBP-1c mRNA expression in adipose tissue of HFD control mice was significantly up-regulated ($P < 0.001$) as compared to LFD control mice. Up-regulation in SREBP-1c mRNA expression was significantly reduced ($P < 0.001$) by the SFSE-OS (100 mg/kg, p.o.) treatment when compared with HFD control mice. But when compared with HFD control mice, the SREBP-1c mRNA expression in adipose tissue of SFSE-OS (30 mg/kg, p.o.) treated mice was not significantly down-regulated (Fig. 1E).

Effect of SFSE-OS on mRNA expressions in liver

The effects of SFSE-OS on gene expressions of Glut-2, Glut-4, IRS-2 and SREBP-1c in liver are presented as Fig. 2. The mRNA expression of Glut-2 and IRS-2 in liver was significantly down-regulated ($P < 0.001$) in HFD control mice as compared to LFD control mice. This down-regulated mRNA expression of Glut-2 and IRS-2 in liver was significantly up-regulated ($P < 0.001$) by the SFSE-OS (100 mg/kg, p.o.) treatment as compared to HFD control mice. However, treatment with SFSE-OS (30 and 60 mg/kg, p.o.) did not produce any significant up-regulation in the mRNA expression level of Glut-2 and IRS-2 in liver as compared to HFD control mice. Moreover, the Glut-4 mRNA expression was non-significantly down-regulated in HFD control mice as compared to LFD control mice and SFSE-OS (30, 60 and 100 mg/kg, p.o.) treated mice also did not show any significant change in Glut-4 mRNA expression as compared to HFD control mice (Fig. 2A, 2B, 2C and 2D). The liver SREBP-1c mRNA expression of HFD control mice was significantly up-regulated ($P < 0.001$) as compared to LFD control mice. When compared with HFD control mice, the liver SREBP-1c mRNA expression was significantly and dose dependently down-regulated ($P < 0.01$, $P < 0.001$ and $P < 0.001$, resp.) by SFSE-OS (30, 60 and 100 mg/kg, p.o.) treatment (Fig. 2A and 2E).

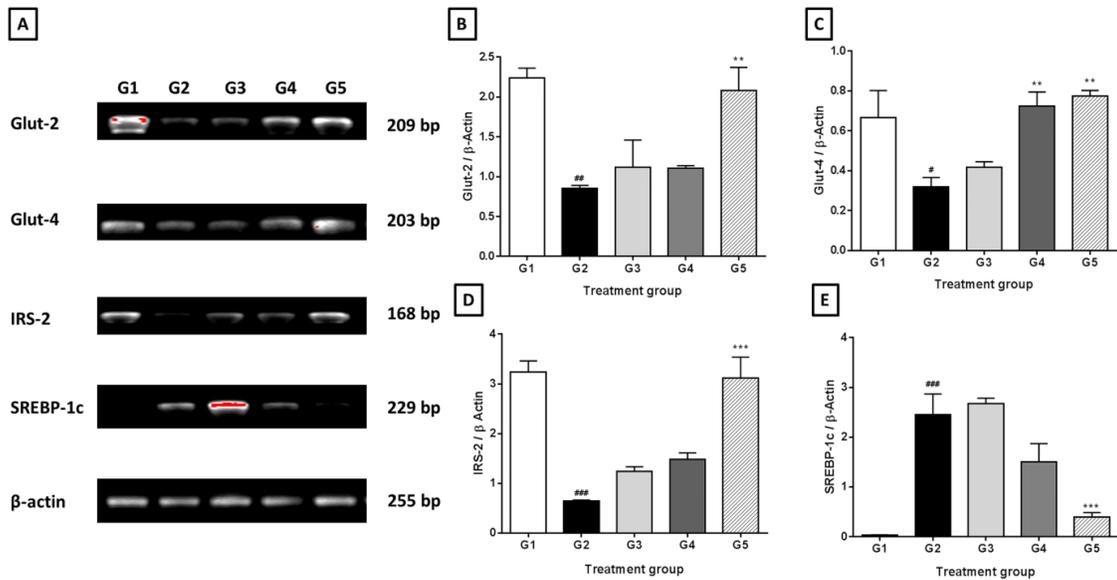


Fig. 1: Effect of SFSE-OS on HFD induced alteration in adipose tissue. Qualitative (A) and quantitative representation mRNA expression of Glut-2 (B), Glut-4 (C), IRS-2 (D) and SREBP-1c (E). Data are expressed as mean \pm S.E.M. (n=4) and analyzed by one-way ANOVA followed by Dunnett's test for each parameter separately. G1 - LFD control; G2 - HFD control; G3 - SFSE-OS (30) + HFD; G4 - SFSE-OS (60) + HFD; G5 - SFSE-OS (100) + HFD. **P < 0.01, ***P < 0.001 as compared to HFD control and ###P < 0.01, ###P < 0.001 as compared to LFD control.

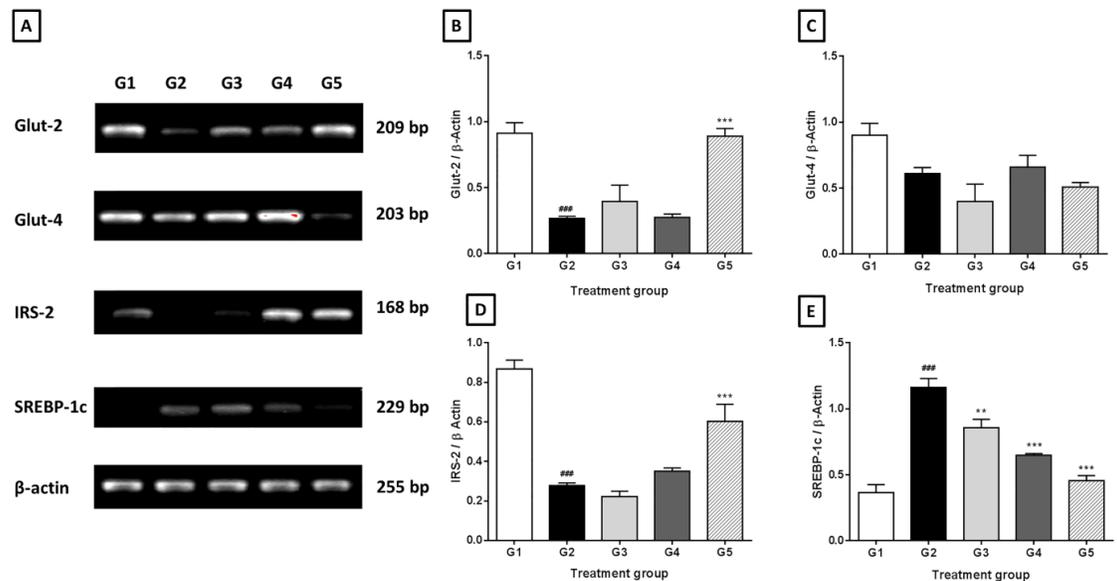


Fig. 2: Effect of SFSE-OS on HFD induced alterations in mRNA expression (reverse transcriptase PCR method) of Liver. Qualitative (A) and quantitative representation mRNA expression of Glut-2 (B), Glut-4 (C), IRS-2 (D) and SREBP-1c (E). Data are expressed as mean \pm S.E.M. (n=4) and analyzed by one-way ANOVA followed by Dunnett's test for each parameter separately. G1 - LFD control; G2 - HFD control; G3 - SFSE-OS (30) + HFD; G4 - SFSE-OS (60) + HFD; G5 - SFSE-OS (100) + HFD. **P < 0.01, ***P < 0.001 as compared to HFD control and ###P < 0.001 as compared to LFD control.

Table 1: Gene specific Primer sequences for Glut-2, Glut-4, IRS-2, IL-6, SREBP-1c and β -actin used for RT-PCR analysis.

Sr. No.	Gene	Primer sequences (5'-3')	Base Pair
1	Glut-2	5'-CTGCACCATCTTCATGTCGG-3' 5'-AATTGCAGACCCAGTTGCTG-3'	209
2	Glut-4	5'-GCCATCGTCATTGGCATTCT-3' 5'-CGCTTTAGACTCTTTCGGGC-3'	203
3	IRS-2	5'-TGGTCCAATAGCTGCAAGA-3' 5'-CCTCCGTTTACTGGGAAGGT-3'	168
4	IL-6	5'-GCCAGAGTCCTTCAGAGAGA-3' 5'-GGTCTTGGTCCTTAGCCACT-3'	153
5	SREBP-1c	5'-CCCACCTCAAACCTGGATCT-3' 5'-AAGCAGCAAGATGTCCTCT-3'	229
6	β -actin	5'-CCTCTATGCCAACACAGTGC-3' 5'-CCTGCTTGCTGATCCACATC-3'	255

Table 2: Effect of SFSE-OS on body weight and glyceimic parameters (fasting glucose level, OGTT and IPITT, plasma insulin, HOMA-IR and HbA1c levels) in mice.

Parameters	G1	G2	G3	G4	G5
	LFD control	HFD control	SFSE-OS (30) + HFD	SFSE-OS (60) + HFD	SFSE-OS (100) + HFD
Body weight (g)	28.66 ± 0.78	33.40 ± 0.65 [#]	29.67 ± 1.10 ^{ns1}	28.33 ± 1.01 ^{**}	29.62 ± 0.80 ^{ns1}
HbA1c (%)	2.94 ± 0.23	5.43 ± 0.22 ^{###}	4.11 ± 0.22 ^{***}	2.96 ± 0.09 ^{***}	3.13 ± 0.12 ^{***}
OGTT					
FPG (mg/dL)	114.44 ± 5.53	175.44 ± 3.73 ^{###}	129.33 ± 2.01 ^{***}	119.83 ± 1.38 ^{***}	125.67 ± 2.73 ^{***}
Fasting Plasma insulin (µg/L)	00.36 ± 0.12	01.12 ± 0.13 ^{###}	00.89 ± 0.13 ^{ns1}	00.70 ± 0.11 [*]	00.57 ± 0.07 ^{**}
2 h post-glucose plasma insulin (µg/L)	00.52 ± 0.08	00.73 ± 0.09 ^{ns}	00.70 ± 0.11 ^{ns1}	00.73 ± 0.11 ^{ns1}	00.89 ± 0.10 ^{ns1}
HOMA-IR	00.11 ± 0.04	00.51 ± 0.06 ^{###}	00.33 ± 0.05 ^{ns1}	00.23 ± 0.04 [*]	00.19 ± 0.03 ^{**}
AUC-OGTT	31998.30 ± 1777.04	42966.70 ± 861.36 ^{###}	40068.80 ± 1885.73 ^{ns1}	33290.00 ± 834.66 ^{***}	36336.30 ± 944.46 ^{**}
AUC-IPITT	8482.50 ± 191.01	17227.50 ± 1132.05 ^{###}	11350.00 ± 192.04 ^{***}	9822.50 ± 291.80 ^{***}	12795.00 ± 673.00 ^{***}

Data are expressed as mean ± S.E.M. and analyzed by one-way ANOVA followed by Dunnett's test for each parameter separately. ns - not significant, ###P < 0.001 as compared to LFD control and ns1 - not significant, *P < 0.05, **P < 0.01, ***P < 0.001 as compared to HFD control.

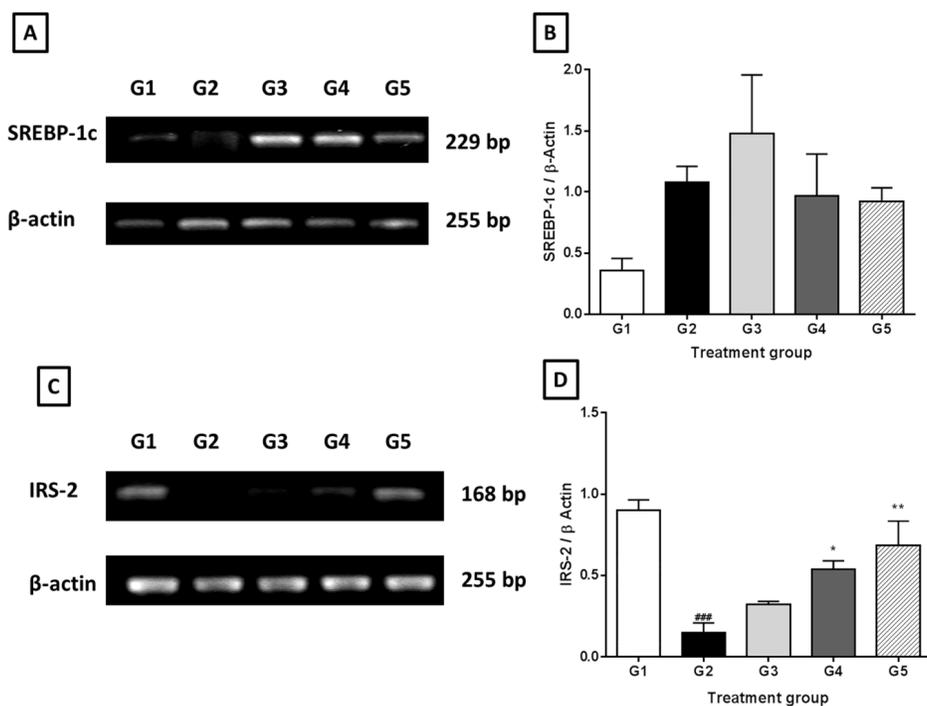


Fig. 3: Effect of SFSE-OS on HFD induced alteration in mRNA expression (reverse transcriptase PCR method) of pancreas (A, B) and skeletal muscle (C and D). Data are expressed as mean ± S.E.M. (n=4) and analyzed by one-way ANOVA followed by Dennett's test for each parameter separately. G1 - LFD control; G2 - HFD control; G3 - SFSE-OS (30) + HFD; G4 - SFSE-OS (60) + HFD; G5 - SFSE-OS (100) + HFD. *P < 0.05, **P < 0.01, ***P < 0.001 as compared to HFD control and ###P < 0.001 as compared to LFD control.

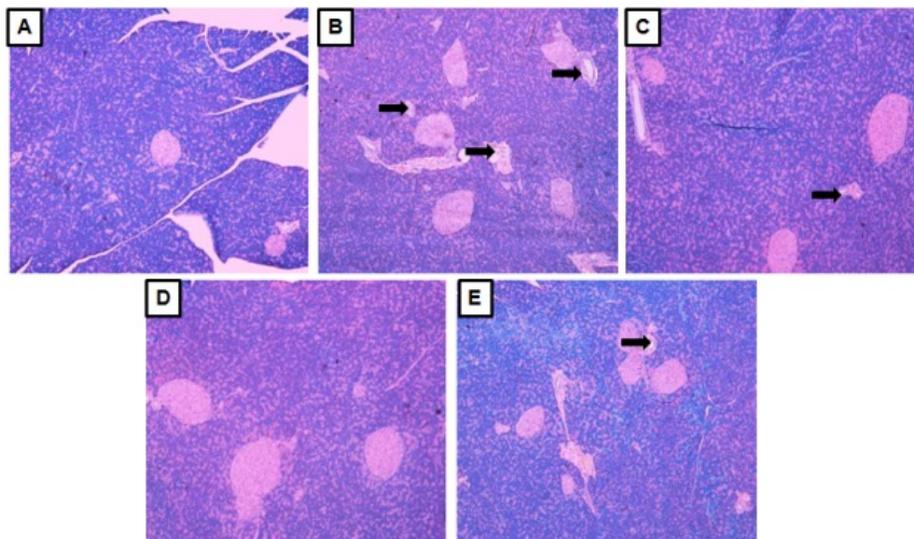


Fig. 4: Photomicrographs of sections of representative sample of pancreas of mice from LFD control (A); HFD control; (B), SFSE-OS (30 mg/kg) treated (C); SFSE-OS (60 mg/kg) treated (D) and SFSE-OS (100 mg/kg) treated (E). H&E staining at 40X. Arrow indicates necrosis in pancreatic beta cell of Langerhans.

Effect of SFSE-OS on mRNA expression in skeletal muscle

The effects of SFSE-OS on gene expression of IRS-2 in skeletal muscle are presented as Fig. 3. There was significant down-regulation ($P < 0.001$) in skeletal muscle IRS-2 mRNA expression in HFD control mice as compared to LFD control mice. Treatment with SFSE-OS (60 and 100 mg/kg, p.o.) significantly and dose dependently up-regulated ($P < 0.001$) the mRNA expression of IRS-2 in skeletal muscle as compared to HFD control mice. However, SFSE-OS (30 mg/kg, p.o.) treated mice failed to produce any significant up-regulation in IRS-2 mRNA expression in skeletal muscle when compared with HFD control mice (Fig. 3A and 3B).

Effect of SFSE-OS on m-RNA expressions in pancreas

The effects of SFSE-OS on gene expression of SREBP-1c in pancreas are presented as Fig. 3. The pancreatic SREBP-1c mRNA expression of HFD control mice did not differ significantly as compared to LFD control mice and SFSE-OS (30, 60 and 100 mg/kg, p.o.) treatment also did not produce any significant change in the pancreatic SREBP-1c mRNA expression as compared to HFD control mice (Fig. 3C and 3D).

Effect on histopathological alteration in mice pancreas

The effects of treatments on histopathological alternations found during histology of pancreatic tissues are presented as Fig. 4. The sections of pancreas from LFD control, HFD control mice and SFSE-OS (30, 60 and 100 mg/kg, p.o.) treated mice showed no histopathological alterations in pancreatic tissue and showed normal size and shape of acini and islet of Langerhans.

DISCUSSION

In the etiology of IR, diet played a important role. Features like weight and fat mass gain, polydipsia and hyperglycemia are hallmark of progressive worsening of IR.

Several reports of postprandial blood glucose reduction and improved insulin sensitivity by fenugreek seed extracts in non-diabetic (Fairchild *et al.*, 1996, Jenkins *et al.*, 1980, O'Connor *et al.*, 1981) and diabetic (Gupta *et al.*, 2001, Losso *et al.*, 2009, Mohan and Balasubramanyam, 2001, Roberts, 2011) subjects exist. The combination of naturally occurring oligosaccharide with pinitol from fenugreek seeds unique and offers excellent potential for management of IR. Therefore, the present study was aimed at evaluation of efficacy and mechanism of SFSE-OS, a unique composition with oligosaccharide and pinitol, in well-validated animal model of diet induced IR (HFD in C57/BL6 mice). Recent advance in preclinical pharmacology allow us to mimic human obese condition in laboratory animals with ease of weight gain as well as IR. HFD feeding for 12 weeks is known to mimic modern fat rich dietary conditions (Surwit *et al.*, 1988). HFD induced IR model is well validated and known to have excellent predictive validity (Buettner *et al.*, 2007, Pagliassotti *et al.*, 1994). This model exhibits an array of features of IR including hyperglycemia

and hyperinsulinemia (Ahren and Pacini, 2002). In the present study, 12-weeks of HFD feeding resulted in increase in body weight, and FPG and insulin levels in HFD control mice as compared to LFD control mice. Our observations of weight gain, hyperglycemia, and hyperinsulinemia in mice confirmed induction of IR and in line with earlier reports (Azman *et al.*, 2012, Veerapur *et al.*, 2012, Zhang *et al.*, 2010).

Treatment with SFSE-OS showed significant antihyperglycemic effects with prevention of weight gain despite of HFD feeding. Prevention of body weight gain shown by SFSE-OS can be attributed to balance between fat burning catabolic (reduction of weight) effect and anabolic (increased muscle mass) effect. In our study during OGTT, HFD control mice showed elevated plasma insulin levels, which represent important characteristics of IR. The elevated insulin levels during the OGTT represents reduced glucose elimination and impaired insulin utilization (Fonseca, 2003). Furthermore, HFD control mice exhibited higher AUC than LFD control mice similar to earlier reports (Abdin *et al.*, 2010, Zhang *et al.*, 2010). In the present study, SFSE-OS co-administration with HFD prevented hyperglycemia during OGTT and IPITT and better utilization of exogenous insulin during IPITT. Administration of SFSE-OS showed time-dependent suppression of increased blood glucose and insulin levels as shown by reduced AUC-OGTT and AUC-IPITT. The results of present study are also consistent with the earlier reports of beneficial effects of chronic administration of fenugreek seeds significantly decreased plasma insulin, AUC of OGTT and HOMA-IR (Hamza *et al.*, 2012b).

These observation can be attributed oligosaccharide content of SFSE-OS similar to reports of slowing glucose intestinal glucose absorption either by carbohydrate hydrolyzing enzyme inhibition (Jo *et al.*, 2014) or induction of intestinal gluconeogenesis (Mithieux and Gautier-Stein, 2014). Another probable mechanism behind observed effects during present study can be increased utilization of insulin-regulated glucose into peripheral tissues. This notion is supported by past reports of activation of an insulin-signaling pathway by fenugreek seed extract in general (Vijayakumar *et al.*, 2005). The reports of insulinotropic activity of oligosaccharides (Kim *et al.*, 2009, Kim *et al.*, 2005b, Lee *et al.*, 2003) and pinitol (Bates *et al.*, 2000, Dang *et al.*, 2010, Kim *et al.*, 2012) suggested their major role in reduction of blood glucose and IR observed in the present study.

To explore molecular mechanism of action in further details we have analyzed mRNA expression important targets in organs related to IR namely skeletal muscle, liver and white adipose tissue (WAT) (Fischer-Posovszky *et al.*, 2007, Korenblat *et al.*, 2008). Skeletal muscle is primary site for IR because insulin-stimulated glucose uptake takes place in skeletal muscle (DeFronzo and Tripathy, 2009). Impairment of insulin signaling transduction pathways leads to IR (Leturque *et al.*, 2009, Previs *et al.*, 2000). Insulin receptor substrate-2 (IRS-2) is known to regulate insulin signaling in liver. In liver, IRS-2 play central role in hepatic nutrient homeostasis via anabolic effects of insulin through the activation of phosphatidylinositol 3-kinase/Akt

pathway cascade (Andersson *et al.*, 2004, Hirashima *et al.*, 2003), inhibit gluconeogenesis, apoptosis (Valverde *et al.*, 2003, Valverde *et al.*, 2004) and GLUT-2 translocation to the plasma membrane, resulting in hepatic IR (Pessin and Saltiel, 2000). Fasted state increases and feeding (and weight increase) reduces expression of IRS-2 (Kubota *et al.*, 2000). In the present study, adipose tissue, liver and skeletal muscles of HFD control mice showed reduced levels of IRS-2 expression whereas pancreatic IRS-2 expression was unaltered. The findings from histology of pancreas are also in line with mRNA expression findings. SFSE-OS co-administration with HFD prevented decrease in the expression of IRS-2 in target organs (adipose tissue, liver and skeletal muscles) and expectedly prevented weight gain and fasting insulin elevation (hyperinsulinemia) in the present study. IRS2 and downstream PI-3K/Akt-->Foxo1 signaling cascade play key roles in many functions of insulin/Insulin like growth factor -1 (IGF-1) that plays important role in control of DM and its complications (Guo, 2013). In the present study, SFSE-OS co-administration with HFD, down regulated the mRNA expression of two important integral isoforms of glucose transporter proteins namely of Glut-2 and Glut-4 in liver and adipose tissues. Glut-2 plays critical role in glucose transport and metabolism to regulate body glucose homeostasis whereas Glut-4 is a key gene that mainly responsible for insulin-stimulated glucose transport in tissues (Charron *et al.*, 1999). Insulin-dependent translocation of Glut-2 resulted in regulation of glucose homeostasis via activation of insulin PI3K/phosphorylated protein kinase B (p-Akt) pathway (Watson *et al.*, 2004). HFD fed mice are known to decrease Glut-2 mRNA expression in liver and adipose tissue to cause hyperglycemia (Azman *et al.*, 2012, Yang *et al.*, 2012). Glut-4 is mainly expressed in liver as well as adipocytes. Clinical evidence of decreased Glut-4 expression in obese and DM patients confirmed its role in glucose homeostasis (Berger *et al.*, 1989, Cushman and Wardzala, 1980, Sivitz *et al.*, 1989). Glut-4 increases insulin sensitivity through glucose uptake via a pathway where insulin binds to its cell surface receptors to facilitate glucose transport in tissue (Kwon *et al.*, 2007) whereas increased expression of Glut-4 in adipose tissue improves glucose disposal and insulin sensitivity *in vivo* (Shepherd *et al.*, 1993, Yang *et al.*, 2012). Moreover, up-regulation in the Glut-4 gene expression in transgenic db/db mice was found to be correlated with amelioration of insulin resistance (Brozinick *et al.*, 2001). Therefore, SFSE-OS is suggested to increase insulin sensitivity and glucose uptake in skeletal muscles through Glut 2 up-regulation. The molecular mechanism of insulin resistance in diabetes involves impaired Glut-4 gene expression via sterol regulatory element binding proteins 1c (SREBP-1c) pathway (Horton *et al.*, 2002, Kang *et al.*, 2010). SREBP-1c gene is one of the responsible factors for development of IR in many organs (Letexier *et al.*, 2003). Up-regulated SREBP-1c expression in HFD control mice in the liver and adipose tissues are in line with earlier reports of up-regulation in SREBP-1c mRNA in HFD-induced insulin resistance mice and subsequent induction of insulin resistance (Horton *et al.*, 2002, Shimomura *et al.*, 2000).

Over-expression of SREBPs, the membrane-bound transcription factors, are implicated in biosynthesis of fatty acids and triglycerides (Foufelle and Ferré, 2002, Osborne, 2000). SREBP-1c play vital role in insulin signaling by inhibiting IRS-2 expression in the liver and adipose tissue (Takahashi *et al.*, 2005) and modulation of glucose-stimulated Glut-2 gene expression (Andersson *et al.*, 2004). In the present study, SFSE-OS administration caused down-regulation in expression of SREBP-1c in liver and adipose tissue that may enhance expression of Glut-2 as well as Glut-4 to facilitate insulin-stimulated glucose transporter in target organ to improve insulin sensitivity. Our study results are in support with earlier reports on mechanism of fenugreek seeds against IR (Vijayakumar *et al.*, 2010, Vijayakumar *et al.*, 2005) and suggested SFSE-OS is a responsible component for action against IR. In the past, dietary oligosaccharides were reported to improve insulin sensitivity in obese horses (Respondek *et al.*, 2011). In addition, dietary pinitol is reported to have antihyperglycemic effects be mediated via the insulin signaling pathway (Do *et al.*, 2008) and reported as an insulin sensitizer or insulin mediator in 3T3-L1 preadipocytes *in vitro* (Do *et al.*, 2008). Therefore, oligosaccharides and pinitol can be envisaged as responsible for efficacy of SFSE-OS against hyperglycemia and IR in the present study. The stimulation of glucose uptake by SFSE-OS against HFD induced IR can be contributed by possible prevention of fat mass accumulation by SFSE-OS. In the past, oligosaccharide (Smith *et al.*, 2010, Wang *et al.*, 2011) and pinitol (Choi *et al.*, 2009, Geethan and Prince, 2008) from other sources demonstrated beneficial effects on lipid and cholesterol metabolism. However, more investigations directed towards effects of SFSE-OS on obesity-induced IR are required to confirm such possibilities.

CONCLUSIONS

SFSE-OS administration showed promising prophylactic efficacy against HFD-induced IR development suggested good potential for management of obese and prediabetes population. This effect, at least in part, mediated through the improved insulin sensitivity and improvement in glucose transport to target organs. The up-regulation of IRS2, Glut2 and Glut4 and downregulation of SREBP-1c in liver and adipose tissues, perhaps plays crucial role in the mechanism of action of SFSE-OS.

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