Ayurvedic dietary formulations and postprandial glycemia in rats

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Abstract: Two dietary formulations were prepared according to the prescriptions made for diabetic people in Ayurvedic classics. One formulation was prepared with barley, brown rice and Bengal gram and parched grains of these food grains were used to prepare second formulation. These formulations were evaluated for their postprandial glycemic effect in rats and were compared with a dietary formula containing mixture of commonly used modern dietary food grains (wheat, polished white rice and pigeon pea). Methanolic extract of these formulations were evaluated for antioxidant activities applying multiple in vitro test methods. It was observed that time dependent increase in the postprandial blood glucose levels of rats up to two hours as well as over all postprandial glycemic loads induced by Ayurvedic dietary formulations were significantly less than the mixture of modern dietary food grains and starch. Total polyphenolic content in the extract of Ayurvedic dietary mixtures and protein content in constituent food grains were higher than the comparative modern dietary food grains. Free radicals scavenging activities, reducing powers as well as antioxidant activity in two Ayurvedic dietary formulations were superior to modern dietary mixture's extract. A SDS-PAGE based protein fingerprint of these formulations was also prepared to identify genuine food grains and standardize dietary preparations. This is the first report of its kind that evaluated and compared postprandial glycemic effect, antioxidant activities of Avurvedic dietary formulations with mixture of modern dietary food grains and provides protein fingerprint as a quality control tool for identification of genuine food ingredients and standardize the finished product.

Keywords: Ayurvedic diet, postprandial glycemia, antioxidant activities, diabetes

Introduction

Modern epidemic of lifestyle related diseases like type-2 diabetes mellitus (T2DM), cardiovascular disorders (CVD) and obesity are thought to be the product of industrialization, progressive modernization and globalization (Zimmet and Alberti, 2006) that has made calorie rich, processed, cheap, and convenient marketed foods the main menu for common man (Sivasankaran, 2010). Researchers argue that human genome which evolved and adjusted to Palaeolithic diet and hunter-gatherer lifestyle some 2.5 million years ago, is finding difficult to adjust to these advancements made during and after Neolithic period (Richards, 2002). The transition and transformation in human dietary patterns and lifestyle conditions started with the dawn of agriculture and animal husbandry practices in Neolithic period merely 10,000 years ago on evolutionary scale. Apart from cultivation of plants and domestication of animals, the phenomenon like increase in population, sedentism and eventually urbanization started with "Neolithic revolution" itself (Richards, 2002). Industrialization, progressive modernization and late 20th century phenomenon of globalization added fire to these practices. Therefore, the inharmonious link between our ancient and genetically determined biology with

the progress made in nutritional, cultural and activity pattern during the course of modernization, may be held responsible for emergence of many of the socalled diseases of civilization (Cordain *et al.*, 2005).

Improvements in public facilities, lifestyle conditions and availability of ample nutrition, paradoxically led to a remarkable increase in the prevalence of risk factors for diseases like T2DM, CVD, obesity, hypertension and Strokes (Zimmet and Alberti, 2006; Zimmet, 2000). These explanations find support with the observations made during animal studies. The Israeli sand rat, which is adapted to a desert environment with frequent scarcities of food, develops obesity and diabetes when maintained in laboratory on a 'westernized rat diet' with abundant food (Haines et al., 1965). Ossabaw Island pigs habituated to live in an environment of uncertain food supply and high physical activity develop obesity and hyperglycemia when raised with food-producing pigs consuming high-calorie diets and living in low activity environment (Whitfield, 2003). Diabetes epidemic among captive population of many primate species in Los Angeles Zoo has also been observed whose zoo lifestyle approximates the high-calorie diet and low-exercise lifestyle of urban humans (Diamond, 2003).

It is important to mention here that the prevalence

of diabetes mellitus has not been a new incidence in India. Ayurvedic texts that originated in India some 4000-2000 year BC (Tiwari, 2006) provide detail descriptions of diabetes mellitus (Madhumeha). Furthermore, the descriptions about the cause and course of diabetes mellitus development made in these texts do not differ with the understanding of modern medical sciences (Tiwari, 2005). Current researches highlight beneficial effects of Paleolithic (Lindeberg et al., 2007; Klonoff, 2009) and Mediterranean diets (Kastorini and Panagiotakos, 2010) on T2DM and other risk factors of CVD. Paleolithic diet has become now a popular diet in Sweden (Lindeberg, 2005). Interestingly, several dietary formulations in the traditional Indian medical texts of Ayurveda have been advocated beneficial for diabetic persons (Shastri, 1949; Shastri et al., 1962). These original texts were scripted in ancient language Sanskrit. In the course of cultural and linguistic transformations in the country, the then known Bharatvarsha through Hindustan to the present day India, the apathy towards this language could not bring this knowledge on the international scientific platform. Therefore, little attention could be paid to evaluate and authenticate the scientific basis of Ayurvedic dietary prescriptions based on modern scientific understandings.

Ayurvedic classics prescribe barley based dietary food preparations and snacks for diabetic persons (Tiwari, 2008). In this research, we compared postprandial glycemic effect of two dietary formulations prepared according to Ayurvedic prescriptions with a dietary mixture containing food grains commonly used in Indian diet today, and soluble potato starch on Wistar rats. Furthermore, this report also evaluated antioxidant properties of these formulations applying various *in vitro* methods and provides an electrophoretic protein-fingerprint of the formulations and individual food grains for standardization and quality control of the mixtures.

Material and Methods

Dietary materials

Barley (*Hordeum vulgare* Linn.), brown rice (dehusked brown grains of *Oryza sativa* Linn.), Bengal gram (*Cicer arietinum* Linn.), wheat (*Triticum aestivam* Linn.), polished rice (dehusked polished-white grains of *Oryza sativa* Linn.), and pigeon pea (*Cajanus indicus* Spreng.), was procured from local food grain stores of Hyderabad city (India). Soluble potato starch was purchased from LOBA CHEMIE, Bombay (India).

Preparation of dietary mixtures

Whole grains of barley, brown rice and Bengal gram were taken in the ratio of 50:25:25 (w/w) and coded as formula 1 (F1). Similarly, whole grains of wheat, polished rice and pigeon pea were taken and coded as formula 2 (F2). Whole grains of barley, brown rice and Bengal gram were parched separately and taken in the same ratio as above and coded as formula 3 (F3). Individual formulations were grinded in flourmill to obtain fine quality raw flour and kept in airtight containers.

Extraction of formulations for determination of antioxidant activities

Extraction of formulations was carried out with slight modification in the method described earlier (Tiwari *et al.*, 2011). In 85% methanol (acidified with 1.0N HCl) each formulation in the ratio of 1:5 (w/v) was soaked for 7 days at room temperature. The formulations in solvent were shaken occasional daily. Supernatant was vacuum filtered on 8th day, concentrated to 1/3 volume under reduced pressure in rotary evaporator ($50\pm1^{\circ}$ C) and lyophilized to dryness. Extracts were stored refrigerated until analysis.

Determination of chemical components in methanol extract

Percentage yield, total polyphenols content (TPC), and total anthocyanins content (TA) were determined as described earlier (Tiwari *et al.*, 2011). The total flavonoids content (TF) in the extracts was determined following method described by Hsieh and Chang (2010).

Evaluation of antioxidant activities

The extracts were evaluated for scavenging of free radicals DPPH, ABTS⁺⁺ cation, and H₂O₂, prevention of ABTS oxidation as reported methods (Tiwari et al., 2011). Reducing activity for FeCl, was assayed as described by Arumugam et al. (2010). Nitroblue tetrazolium (NBT) reducing activity as a measure of presence of ascorbic acid (Concklin et al., 2000) in formulations was determined as follows. In a 96- well plate containing 100 µL phosphate buffer (50 mM, pH10) and equal volume of NBT (1mM), prepared in same buffer), 50 µL (5mg/ml DMSO solution) samples were mixed and incubated for 15 minutes. A blank with each extract in the absence of NBT was run to correct background absorbance. Reduction of NBT was measured at 560 nm using BioTek Synergy 4 multimode microplate reader (BioTek Instruments Inc, USA). The percentage of NBT reduction was calculated applying following formula % reduction=

[(A_t - A_c)/ A_t x100] where, (A_t) represents absorbance of reduced NBT by extracts and (A_c) the absorbance of NBT solution in reaction mixture in the absence of extract. At least four to five dilutions in triplicate of each extract was run in every experiment to find out either free radical scavenging concentration 50% or the reducing power 50% (RC₅₀) of extract. Suitable regression analysis was applied to obtain SC₅₀ and RC₅₀ values.

SDS-PAGE protein fingerprinting

100 mg powder of each formulation and individual food grains was taken into test tubes containing 4 ml of ice-cold protein extraction buffer [62.5 mM Tris-HCl (pH 6.8). 2% SDS, 5% glycerol, 3% β -mercaptoethanol, 5M Urea] according to the method of Sadia *et al.* (2009) with slight modifications. Mixture was vortexed intermittently for five minutes and centrifuged at 15000 rpm for 10 min at 4°C. Supernatant was collected and used for protein estimation (Bradford, 1976) and SDS-PAGE protein separation.

20 µL equal concentration protein samples were mixed with appropriate volume of 2X SDS loading buffer containing 0.5M Tris-HCl (pH 6.8), 10% SDS, Glycerol, 2-mercaptoethanol and bromophenol blue and heated for 5 minutes in boiling water. 20 µl of samples were separated on 12% SDS-PAGE (Bio-Rad mini Protein gel apparatus) along with molecular weight marker. The gel was allowed to run in 1X Tris-Glycine buffer (10X buffer- 250mM Tris base, 1.92M Glycine, 1% SDS), at constant voltage of 100V. The gel was stained with 0.5% Coomassie brilliant blue (250 R) solution for an hour and de-stained several time with fresh methanol: acetic acid: water (50:10:40) solution until appropriate staining of the gel reached. Destained gels were photographed using BioDoc-ItTM Imaging System, UV Transilluminator UVP.

Postprandial glycemic test

Postprandial glycemic activity of dietary formulations was determined by feeding individual formulation to group of rats. Male Wistar rats (176±11 gram body weight) were obtained from National Institute of Nutrition (CPCSEA Reg. No.154, Government of India), Hyderabad. Rats were housed in standard polyvinyl cages in the institute's (IICT) animal house. Room temperature was maintained at $22\pm1^{\circ}$ C with an alternate 12 h light dark cycle. Food and water were provided *ad libitum*. Approval of experimental protocol was obtained from Institutional Animal Ethical Committee (IAEC-IICT). All the animals were kept

for overnight fasting. Next day forenoon, blood was collected from the retro orbital plexus in EDTA containing tubes. Basal plasma glucose level ('0' h) was estimated by glucose oxidase method with auto blood analyzer (Bayer EXPRESS PLUS). Rats were divided into various groups. Each group contained six rats. Animals were grouped such that the mean basal glucose levels do not differ significantly among the groups. Light slurry of formulations F1, F2 and F3 was prepared in normal saline and administered orally (2-gm/kg body weight) to respective group of animals. Control groups of animals were given soluble potato starch in the same dose. A positive control group with standard drug acarbose was also taken. Acarbose (10 mg/kg body weight) was given to animals fifteen minutes before starch feeding. Blood was collected at the intervals of 30, 60, 90 and 120th min post-starch or the formulations feeding. Plasma glucose levels were measured as described above. Time dependent increase in plasma glucose level and two-hour postprandial glycemic load as a measure of area under the curve $(AUC_{0-120minutes})$ was calculated following trapezoidal rules (Raju et al., 2010).

Statistical analysis

One-way ANOVA followed by Dunnett's multiple comparison tests was applied to compare difference between animal study groups. To determine degree of significance between the groups p < 0.05 was considered.

Results and Discussion

Chemical components

Table 1 presents percentage yield and different antioxidant compositions in methanolic extract of three formulations.

 Table 1. Antioxidant components in methanolic extract of dietary formulations

Yield of extract	TPC	TF	TA
7.6	29.8±0.6	16±2	0.61±0.19
7.4	32.8±0.4	35±2	0.25 ± 0.001
6.4	37.8±0.1	26±4	ND
	Yield of extract 7.6 7.4 6.4	Yield of extract TPC 7.6 29.8±0.6 7.4 32.8±0.4 6.4 37.8±0.1	Yield of extract TPC TF 7.6 29.8±0.6 16±2 7.4 32.8±0.4 35±2 6.4 37.8±0.1 26±4

The % yield denotes yield %w/w; gm/100gm. Total polyphenols content (TPC) represents mg GAE/gm, total flavonoids (TF) as mg RE/gm and total anthocyanins (TA) as %w/w. GAE; gallic acid equivalent, RE; rutin equivalent, ND; not detected. Values are mean \pm SD, n=3.

The percentage methanolic yield did not differ among the formulations. Total polyphenols content (TPC) in formula F3 was 21% more than in F1. In formulation F2, total flavonoids content (TF) was found double than F1 and 25% more than F3. Total anthocyanins (TA) could not be detected in formula F3. However, in formula F1 it was 69% more than formula F2. It requires mention here that with increasing time of extraction percentage yield of the extract increases as in our previous study (Tiwari *et al.*, 2011) with only

	Table 2. Pro	otein conten	t in foo	d grains a	and formu	ilations
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Barley	Brown Rice	Bengal gram	Wheat	White Rice	Pigeon Pea	F 1	F 2	F 3
20.9±0.2	18.4±1.7	44.1±4.7	17.9±0.5	17.8±4.7	24.2±3.8	22.1±0.3	24.6±0.65	16.6±0.16
alues (mean±SD) represent mg/gm of grain or formulations powder.								

24-hour extraction time fewer yields was obtained.

In barley, protein content was 14% higher than wheat (Table 2). These two food grains are used mainly to prepare chapatti and breads. Pulse grain Bengal gram contained more protein (45%) than pigeon pea (Table2). Equal protein content in white rice and brown rice was observed. Similarly, there was no difference in protein content of formula F1 and F2. However, protein content in formula F3 was 29% less than formula F1 and 47% less than formula F2.

Antioxidant activities

Highly processed, calorie-dense, nutrient depleted diet leads to exaggerated postprandial spikes in blood glucose and lipids that induces immediate oxidative stress. Induction of oxidative stress has been observed to increase in direct proportion to the increase in postprandial blood glucose level (O'Keefe et al., 2008). Hyperglycemic spikes even in non-diabetic individuals have been shown to markedly increase free radicals generation (Brownlee and Hirsch 2006), Furthermore, oxidative stress has been recognized recently as a major physiological link between CVD and diabetes (Giugliano et al., 1996) and also in the development of diabetic complications (Brownlee, 2001). Therefore, presence of free radicals scavenging antioxidant properties in a diet becomes an important consideration.

Figure 1 presents concentration dependent multiple free radical scavenging properties of the methanol extract of formulations. It was observed that scavenging activities for free radicals like DPPH (Figure 1a), ABTS⁺ cation (Figure 1b), H₂O₂ (Figure 1f), and reduction of FeCl₂ (Figure 1d) and NBT (Figure 1e), or ABTS oxidation (Figure 1c) prevention activities, all the three extract displayed varying degrees of activity potentials. With respect to the IC₅₀ values (Table 3), it was found that extract from the formulation F1 displayed far better antioxidant activity than extract from formulation F2 in scavenging free radicals DPPH, ABTS⁺⁺, reducing FeCl, and NBT. However, methanolic extract of formulation F3 displayed more than twice ABTS oxidation prevention and FeCl₃ reducing activity than F1 extract. Formula F1 was more potent than formula 3 in scavenging DPPH radical however, ABTS⁺⁺ scavenging potency of these two formulas remain same. The antioxidant potency of formula F2 was close to double than formula F1. These results show that whether free radicals scavenging, reducing power or the antioxidant activities, extract of Ayurvedic formulations displayed far better activity than extract of formulation F2.

Table 3. IC₅₀ for methanolic extract of dietary formulations

Formulation	DPPH (µg/mL)	ABTS ⁺⁺ (µg/mL)	ABTS-ox (µg/mL)	NBT (µg/mL)	FeCl (µg/mL)
F1	134.2	31.5	19.0	41.1	226.2
F2	278.4	49.3	11.0	89.0	606.9
F3	211.7	31.8	7.9	48.2	126.9



Figure 1, Concentration dependent antioxidant. activities in methanolic extract of dietary formulations. [a] DPPH scavenging activity. [b] ABTS⁺, scavenging activity. [c] Prevention of ABTS oxidation (ABTS_{ox}). Figures [d] and [e] represent FeCl₃ and NBT reducing activity respectively. [f] Kinetics of hydrogen peroxide (H₂O₂) scavenging at 100µg/ml concentration of samples. Data represent mean ± SD, n=3.



Figure 2. SDS-PAGE protein fingerprint of individual food grains and Ayurvedic dietary formulations. Formulation 3 (F3), formulation 2 (F2), formulation 1 (F1), wheat (W), pigeon pea (PP), white rice (WR), Bengal gram (BG), brown rice (BR), barley (B), Molecular weight marker (MM). Differences in protein band patterns of different food grains are clearly visible. It is clear therefore that each food grain has its unique protein band pattern. Formulation F1 was prepared by mixing barley, brown rice and Bengal gram. It is evident from the band pattern of F1 that it has a different fingerprint than the individual food grains despite the fact that some of the proteins of brown rice, Bengal gram and barley appear in its fingerprint. Similarly, F2 contains mixture of wheat, white rice and pigeon pea. Major proteins of pigeon pea are evident in F2 protein band pattern, altogether however, the protein fingerprint of F2 displays different protein profiles than individual food grains. It is interesting to note that the protein fingerprint of formula F3 is absolutely different than its parent mixture F1. Only few protein bands between 35-45 kDa ranges could be visualized in this formulation. This difference may be due to the reason that F3 was prepared from the parched grains. In the process of parching many of the proteins of individuals food grains might have been lost. The unique electrophoretic protein fingerprint of dietary formulations F1, F2, and F3 from their compositional food grains therefore, may serve as quality control tool and standardization of individual formulation.

In a biological system, multiple sources of free radical generation, oxidants and antioxidants have been identified that possess different chemical and physical characteristics (Prior et al., 2005). Therefore, to balance the multiplicity of oxidants, multiple characteristics in antioxidants are required. It has been observed that in some cases, antioxidants present multiple mechanism of action in a single system (Ishige et al., 2001), however, may display different mechanism of action depending on the reaction system and can respond in a different manner to different radical or oxidant sources (Prior et al., 2005). Because of these multiplicities involved in the characteristics as well as mechanism of antioxidants action, no single assay can optimally reflect true characteristics of antioxidants in a mixture (Wootton-Beard et al., 2011). Variations in the activity level and response to different free radicals in this study presented by dietary Ayurvedic formulations may be due to the presence of mixtures of different type of antioxidants, their synergistic mode of actions, and differences in their physical and chemical characteristics.

Despite the fact that antioxidants present in human diet play important beneficial role in reducing oxidative stress and preventing free radicals induced biomolecules damage, it has recently been observed that the timing (Diano, 2009) and order of antioxidants consumption with meal (Imai and Kajiyama, 2010) affect other metabolic parameters differently. It has been observed that antioxidant when taken on an empty stomach affects appetite and when consumed along with diet affects satiety (Diano, 2009). Furthermore, some antioxidant rich fruits (Alvarez-Parrilla et al., 2010) and antioxidant rich fractions from food grains (Tiwari et al., 2011) have been alleged to induce hyperlipidemia and hyperglycemia respectively. Similarly, antihyperglycemic compounds isolated from fruits have also been reported to increase postprandial blood glucose level when carbohydrate diet is fortified with such compounds (Tiwari et al., 2010). Therefore, selection of antioxidant rich food items meant for diabetic patients that do not adversely affect blood glucose level requires special attention.

SDS-PAGE protein fingerprint

One of the major problems associated with natural food products is the availability of appropriate standardization and quality control tools. Lack of such tools increases the malpractice of adulterations and also erroneous identification of the natural material. Based on molecular characteristics of electrophoretic protein fingerprints, it has become possible to gather information about genetic variations, taxonomic relationship, phylogenetic diversity and even identification of sub-species of a plant material around the world (Emri et al., 2007; El-Hady et al., 2010). The unique band patterns of protein electrophoregram of food grains or the mixture therefore, could serve as an important supplemental tool that can provide passport data for its identification and standardization. Figure 2 presents distinct spectrum of electrophoretic protein band patterns of individual food grains and the formulations prepared by mixing these food grains. Distinct protein band patterns for each food grains were observed. However, protein fingerprint of the mixture of dietary formulations present a clear distinction and differences in protein band pattern than the respective food grains used to prepare formulations (Figure 2). The presence or the absence of a protein band of constituent food grain protein in the respective formula may help identify genuine product. This tool may serve the purpose of correct identification of food grain as well as proper standardization of Ayurvedic dietary formulations.

Postprandial glycemia

Postprandial hyperglycemia (PPHG) has emerged as a prominent and early defect in ensuing T2DM (Carroll et al., 2003). The deterioration of glucose homeostasis in individuals with T2DM progresses from postprandial to fasting hyperglycemia in several steps (Monnier et al., 2007). PPHG has also been identified as an independent predictor of the development of future cardiovascular events even in non-diabetic individuals (O'Keefe et al., 2008). The postprandial state is characterized by several metabolic alterations that may play role in the pathogenesis of CVD. PPHG has been reported as an important independent risk factor contributing to the development of atherosclerosis and amplify generation of oxidative stress (Ceriello, 2000). Indian population has been observed to present increased prevalence of PPHG than other ethnic groups around the world on a daily seven-point scale (Milicevic et al., 2008). Higher intake of carbohydrate in the diet of Asians and South Asians in particular, has been observed in comparison to White Caucasians (Misra and Khurana, 2011). Evolutionarily, this may be due to the reason that adoption of agriculture in Neolithic period occurred independently in different parts of the world that influenced selection of dietary materials and hence dietary patterns. The starch rich food plants such as rice became main domesticated plant in Asia (Imamura, 1996; Crawford, 1992). Now, for most Asian population, white rice has become staple food. Rice is readily available, considered more palatable (Sugiama et al., 2003) and constitute major portion in the menu of this region. The high level of carbohydrate content in the diet of Asian population along with use of white rice as a staple food may be the reason of increased incidence of postprandial hyperglycemia and increasing prevalence of T2DM in this part of the world. Therefore, a diet that can control precipitous rise in postprandial glycemia may become helpful in bringing down the risk factors associated with PPHG.

Figure 3 presents time dependent increase in postprandial blood glucose level in animals fed on different dietary formulations. It was observed that rats fed on F2 diet (which includes mixture of present day common dietary food grains like white rice, wheat flour made breads and Chapatti, and pulse made of pigeon pea), there was a sharp and significant rise in blood glucose level even at the first 30th minute after oral feeding (Fig. 3A) and later blood glucose level paralleled close to the blood glucose level of animal group who received control starch diet. The F2 diet induced increase in blood glucose level was significantly (p<0.05) higher at this time point (first 30th minute), than that induce even by pure starch diet (Control group) fed animals. This may explain the reason Asian population display higher PPHG levels that consume this type of food today. It requires mention here that the development and adoption of this type of fast-glucose releasing diet might have been the requirement of fast paced lifestyle of modernizing society, which looked-for quick energy giving diet.



Figure 3. The glycemic response curve in rats following feeding of F1, F2, F3 and soluble potato starch (Control). [A] Glucose levels at different time points with dietary formulations F1 and F2. Control group received soluble potato starch. [B] Glucose levels with dietary formulation F3. Control group was treated sham as in [A]. Dosage was 2-gm/kg body weights in each group of animals. One group in the experiment [B] with standard antihyperglycemic drug acarbose was also kept wherein acarbose (10 mg/kg body weight) was orally administered 15 minutes before soluble potato starch feeding. ANOVA followed by Dunnet's multiple comparison tests was applied to compare the differences between the means. Degree of significance was chosen "p < 0.05 when compared with control. Values represent mean \pm SD, n = 6.

A verse (Ch.Chi. 6.21) from Ayurvedic text *Charaka Samhita* (Shastri *et al.*, 1962) advocates that the diet of diabetic patients should consist predominantly of barley, wherein various food items may be prepared along with wild variety of rice and barley (Tiwari, 2008). This formed the basis for preparation of dietary formula F1. It was observed that the increase in blood glucose level in animals fed on this formulation was significantly less at 60th and

90th minute (Fig 1A). Although the increase in blood glucose level at 30th and 120th minute was also less in this group, it could not reach degree of significance when compared with either control group or the F2 diet fed group of animals.

Similarly, another verse (Ch. Chi. 6.48) from the same text (Shastri *et al.*, 1962) mentions that use of parched barley flour prevents development of diabetes (Tiwari, 2008). In olden time's parched barley, parched brown rice and parched Bengal gram grains and their flour were used as munching snacks and drinks in India, particularly in Northern part of the country. This practice is still breathing in remote parts of Northern India. These facts laid down the basis for preparation of formula F3. It can be seen in figure 3(B) that by feeding formula F3, the rise in blood glucose level in rats was significantly less (p<0.05) when compared with starch fed control group of animals at 30th and 60th minute and remained lesser than starch fed group up to 120 minute.

Intestinal α -glucosidase inhibitor drugs like acarbose and voglibose slow down the digestion of carbohydrate rich diet. These drugs have been observed better therapeutic in controlling PPHG in Asian people (Scheen, 2009) presumably because of their specific food habits (Chan et al., 1998). Such drugs are now getting special prescription in the population particularly Indian whose diet is starch rich and demonstrate higher glycemic response to all foods (Henry et al., 2008) and present increased prevalence of PPHG (Milicevic et al., 2008) than other ethnic groups around the world. It was interesting to note that postprandial glucose level of formulation F3 treated group of animals was comparable to the postprandial glucose level of acarbose treated group of starch fed animals (Figure 3B). It is possible therefore that if these dietary formulations (F1 and F3) become substitute as dietary therapies, the use of drugs like acarbose can be avoided.

The Paleolithic diet (based on lean meat, fish, fruits, vegetables, root vegetables, eggs and nuts) has been reported significantly potent in improving glucose tolerance test in ischemic heart disease (IHD) patients with impaired glucose tolerance or T2DM than the Mediterranean-like diet that is based on whole grains, low-fat dairy products, fish, fruit and vegetables (Lindeberg *et al.*, 2007.) In another study, Paleolithic diet was found more satiating per calorie than a Mediterranean-like diet in individuals with IHD (Jonsson *et al.*, 2010). However, Paleolithic diet was less liked by the study participants and the drop out rate was observed more in this study group (Lindeberg *et al.*, 2007) despite the fact that it significantly improved over all two hour postprandial

glucose load in terms of area under the curve (AUC $_{0-120 \text{ minutes}}$) than Mediterranean-like diet.

It appears therefore that even though the Stone-Age Paleolithic diet may have better control over PPHG, it may not become diet of preference by the people in modern world. However, a diet or dietary formulation based on whole grains can easily become popular if helps prevent excessive postprandial glycemic excursion. Figure 4 presents postprandial incremental plasma glucose AUC_{0-120 min} with different dietary formulations. It is evident from figure 4 that substitution of Ayurveda based dietary formulations like F1 and F3 if prescribed for impaired glucose tolerant people and diabetic patients, may have significant potential in improving glucose intolerance, and also prevalence and incidence of PPHG caused by modern diet in general population. These formulations may become a suitable dietary substitute to control modern epidemic of PPHG because restoring to drug therapy for an epidemic caused by maladaptive hyperglycemic diet (which represent conventional diet of today like F2) is less rational than simply restoring to olden day's time tested diets (F1 and F3) and realigning dietary habits with the physiological needs (O'Keefe et al., 2008).



Figure 4. Postprandial incremental plasma glucose AUC $_{0:120 \text{ min}}$ with different dietary formulations and soluble potato starch feeding in rats. Area under the curve (AUC) was calculated following trapezoidal rules applying formula: AUC $_{0:120 \text{ min}} = (BG_0 + BG_3) \times 0.5/2 + (BG_{30} + BG_{60}) \times 1.22$, where $BG_0 + BG_{30} \times 0.5/2 + (BG_{120} + BG_{60}) \times 1.22$, where $BG_0 + BG_{30} \oplus BG_{30} \oplus BG_{120}$ represent plasma glucose level at 0, 30, 60 and 120 min after soluble potato starch (control) or dietary formulations (F1, F2 and F3) feeding. ANOVA followed by Dunnett's multiple comparison tests was applied to find difference between the groups. *p < 0.05 when compared with control. Values represent mean \pm SD, n = 6.

Consideration of hyperglycemia rates among domesticated mammals suggests that ethnic and national heterogeneity in diabetes prevalence is rooted in more recent events (Gerstein and Waltman, 2006). Therefore, the rising rates of diabetes in traditional living population of developing countries may be due to the recent adoption of western culture and dietary habits (to which they were least exposed) leaving aside their traditional dietary practices to which they were raised for long time. Infact, for at least 300 years Europeans have enjoyed a relatively stable food supply and increasingly available laboursaving devices (Diamond, 2003). The populations of developing countries are newly exposed to such an environment therefore, may have little resistance to its diabetogenic effect (Gerstein and Waltman, 2006). The Mediterranean diet was first described in the 1960s by Ancel Keys based on his observation of food habits of some populations in the Mediterranean region (Keys et al., 1986). So far, Mediterranean diet is one of the best-known and studied diets for its beneficial effects on human health that may act beneficially against development of T2DM, including reduced oxidative stress and insulin resistance (Kastorini and Panagiotakos, 2010). However, Indian Ayurvedic dietary formulations being claimed as one of the ancient medical dietary prescriptions for the people suffering from diabetes could not receive global attention. These dietary compositions (Shastri et al., 1962; Shastri, 1949) do not differ from that of the Mediterranean diet. The prevalence of T2DM in rural as well as urban populations of developing countries of the South/East Asia is taking shape of epidemic. These countries share many of the common features in their dietary practices and pattern. Therefore, scientific exploration, design and development of dietary formulations mentioned in Ayurvedic texts for the prevention of development of T2DM epidemic in these regions may offer better therapeutic dietary substitute than the dietary prescriptions explored, studied, designed and developed to suite white Caucasian population.

Conclusion

The human genome and genetically determined biology of human was evolved and adapted to natural diet, environment, and hunter-gatherer life style before the dawn of Neolithic period. However, with the beginning of agricultural practices and domestication of animals during Neolithic period to the modern industrialized and globalized world, it got exposed to extreme changes in the lifestyle conditions and dietary patterns to which it is less adopted. Resultantly, human population around the world is facing epidemic of several metabolic disorders like T2DM, CVD, obesity, stress and cancer etc. Nevertheless, ancient civilizations in the world to which these disease symptoms were known, had developed several dietary and natural therapeutic formulations for prevention of the development and treatment of these diseases. Ayurveda, one of the oldest medical systems in the world, described barley-based dietary formulations beneficial for the people suffering from diabetes mellitus (Prameha and Madhumeha). The low-postprandial glycemic load as well as potent antioxidant activities present in Ayurvedic dietary formulations (F1 and F3) observed in this study when compared with modern high-postprandial glycemic value diets, approves their preventive as well as therapeutic effectiveness. Therefore, development and adoption of such diet may help prevent the prevalence of PPHG and epidemic of T2DM in the people of South/East Asian origin in particular where such diet was in traditional practice. To the best of our knowledge this is the first report of its kind that explored effect of Ayurvedic diet on postprandial glycemia and evaluated antioxidant properties present in such dietary formulations.

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