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**ASSOCIATION OF INTERFERON-GAMMA AN INFLAMMATORY MARKER AND FASTING SERUM BILIRUBIN AN OXIDATIVE STRESS MARKER IN ESTABLISHED CASES OF METABOLIC SYNDROME AROUND KOLKATA AND ITS NEIGHBORHOODS**

\*Sandip Ghosh<sup>1</sup>, Debjyoti Bhattacharya<sup>2</sup>, Chittaranjan Maity<sup>1</sup> and Gora Chand Bhattacharya<sup>3</sup>

<sup>1</sup>Department of Biochemistry, KPC Medical College, Jadavpur, Kolkata, India

<sup>2</sup>Department of Biochemistry, Calcutta National Medical College, Kolkata, India

<sup>3</sup>Department of Biochemistry, MGM Medical College & LSK Hospital, Kishanganj 855107, Bihar, India

\*Author for Correspondence

**ABSTRACT**

Subjects of Metabolic Syndrome are prone to develop risk factors for type 2 diabetes mellitus and cardiovascular diseases. . Inflammatory process and atherosclerosis are associated with the disorder. In this study, serum interferon gamma a potent inflammatory marker and serum total bilirubin an antioxidant, have been measured in 150 subjects of Metabolic Syndrome and compared with 150 controls. We observed an inverse relationship between interferon gamma and total bilirubin. The subjects of metabolic syndrome were diagnosed as per NCEP (ATP III) criteria BMI, waist –hip ratio, lipid profile and HOMA- IR of the test and control groups have been considered for diagnostic criteria.

**Keywords:** *Metabolic Syndrome, Gamma Interferon, Serum Bilirubin*

**INTRODUCTION**

Metabolic syndrome or Syndrome –X is recognized as one of the most serious risk factors for developing type II diabetes, as well as cardiovascular disease (Wilson *et al.*, 2005). Criteria for diagnosis of metabolic syndrome are the criteria of the National Cholesterol Education Programme (NCEP) Adult Treatment Panel III (ATP-III) (Grundy, 2008; Eckel *et al.*, 2005; NCEP, 2001), with combination of body mass index (BMI) and waist circumference. The key points of this criteria are: 1) triglyceride levels  $\geq 150$  mg/dl ;2) HDL – C levels  $< 40$  mg/dl in men or  $< 50$  mg/dl in women; 3) Fasting plasma glucose levels  $\geq 110$  mg/dl or taking an anti diabetic medication; 4) systolic blood pressure  $\geq 130$  mmHg or diastolic blood pressure  $\geq 85$  mmHg (or receiving drug therapy for hypertension); and 5) BMI  $> 25$  Kg/Meter <sup>2</sup> (Ishizaka *et al.*, 2005).

The exact mechanism underlying metabolic syndrome has not yet been elucidated completely. Many cross- sectional or longitudinal studies have shown that metabolic syndrome is strongly associated with inflammation (Wang *et al.*, 2004; Rutter *et al.*, 2004; Festa *et al.*, 2000), insulin sensitivity (Lann and LeRoith, 2007), endothelial dysfunction (Gideon *et al.*, 2007). As a result, several biological markers have been proposed as risk factors for metabolic syndrome that are associated with an increase in the risk of metabolic syndrome: homeostasis model assessment insulin resistance index (Festa *et al.*, 2000); (HOMA-IR; as a marker of insulin resistance), homocysteine (Gideon *et al.*, 2007) (as a marker of endothelial dysfunction). Gamma interferon was described by E. F. Wheelock as a product of human leukocytes when stimulated with phytohemagglutinin an extract of the kidney beans *Phaseolus vulgaris* (Wheelock, 1965), and by others as a product of antigen-stimulated lymphocytes (Green *et al.*, 1969) or tuberculin-sensitized mouse peritoneal lymphocytes (Milstone and Waksman, 1970); the resulting supernatants were shown to inhibit growth of vesicular stomatitis virus. Those reports also contained the basic observation underlying the now widely employed interferon gamma release assay used to test for tuberculosis. In humans, the IFN $\gamma$  protein is encoded by the *IFNG* gene (Naylor *et al.*, 1983). IFN $\gamma$  was formerly known as immune IFN, macrophage activating factor, Tcell IFN, and type II IFN. IFN $\gamma$  is synthesized by activated T cells (CD4Th0, CD4Th1 and CD8) and NK cells. IFN $\gamma$  is produced by CD4 Th1 cells following antigen activation (Mytar *et al.*, 1995). Interferon gamma (IFN $\gamma$ ) is a dimerized soluble cytokine that is the only member of the type II class of interferons (Gray and

### **Research Article**

Goeddel, 1982). IFN $\gamma$  is also produced as an aberrant expression of class II histocompatibility antigens by tissue cells (such as thyroid cells) that do not normally express them leading to autoimmune disease (Dorland, 2003). It is an important cytokine produced by body in altered adipose tissue metabolism (Viviane *et al.*, 2008). IFN $\gamma$  and growth factors (such as FGF) belong to distinct groups regarding their regulation by heparan sulfate/heparin-like molecules (Sadir *et al.*, 1998). C terminal residues of IFN $\gamma$  binds to heparan sulfate by electrostatic interactions between lysine or arginine charged side chains and heparan sulfate sulphate groups and it also requires an octasaccharide for effective binding (Vanhaverbeke *et al.*, 2004). As there is no available data for study in Indian population, the object of present study is to establish any co-relation between interferon-gamma an inflammatory marker and fasting serum bilirubin an oxidative stress marker in established cases of metabolic syndrome around Kolkata and neighborhoods in India.

As bilirubin acts as a potent antioxidant (Stocker *et al.*, 1987), the object of study is to establish any correlation between gamma IFN interferon and fasting serum bilirubin in proved cases of metabolic syndrome among the patients visiting outpatient departments of Calcutta National Medical College, Kolkata, India. Informed consent was duly taken from each subject under study, and the entire procedure was done as per the Institutional ethical permission.

**Inclusion Criteria:** In this hospital based cross sectional study, subjects of metabolic syndrome were diagnosed with high BMI, waist-hip ratio, HOMA-IR, hypertension, known type-2 diabetes mellitus. Healthy controls were chosen from the spouse of metabolic syndrome patients, medical students, teaching and non-teaching staffs of the Calcutta National Medical College, Kolkata.

**Exclusion criteria:** Subjects with chronic diseases, known tuberculosis, hepatitis due to any cause, Cushing's syndrome, pregnancy, chronic alcoholism, renal dysfunction, long treated diabetes mellitus, taking treatment for thyroid disorders, recent history of fever and infection were excluded from the study.

### **MATERIALS AND METHODS**

Blood samples were collected from the patients visiting the outpatient department of Calcutta National Medical College, Kolkata, India. Necessary clearance from Institutional ethical Committee was obtained. Blood was collected from ante cubital vein. Cells were separated by centrifugation at 5000 rpm for 5 minutes. Blood glucose, serum bilirubin were estimated immediately. Serum was separated and stored in -20°C for future use. Patients Body Mass Index was measured using standard measuring tape and weighing machine. Height was measured against wooden vertical scale. Body mass index was measured using the following formula mass in Kg/ height in meter<sup>2</sup>. Fasting plasma glucose and total bilirubin was estimated using autoanalyser (A125-Transasia). HbA1c was estimated using Bio-Rad D-10. Fasting insulin was estimated using ELISA kit of Monobind Inc. waist -hip ratio was measured using standard measuring tape in the outpatient department. Insulin resistance was estimated by calculating HOMA-IR originally described by Mathew *et al.*, (1985) (Homeostatic model assessment- Insulin resistance). HOMA-IR was calculated using the following formula:

$$\text{HOMA-IR} = \frac{\text{fasting glucose (mg\%)} \times \text{fasting insulin (\mu U/ml)}}{405}$$

Gamma interferon was estimated using ELISA kit of Ray Biotech in TECAN ELISA reader, Imperial Biotech. Subjects were divided in two groups - control and test. Control group had 150 healthy individuals and test group had 150 proved cases of metabolic syndrome. Test group subjects were selected using standard criteria for metabolic syndrome. Individuals with chronic disease manifestation were excluded from study. All reagents and samples were brought to room temperature (18 - 25°C) before use. A standard graph was prepared with standards provided with the kit. Standards were diluted by serial dilution method. In case of controls eight obese and eight non obese subjects were tested with duplicates. 100  $\mu$ l of each standard and sample into appropriate wells. Wells were covered and incubated for 2.5 hours at room temperature with gentle shaking. The solution was then discarded and washed four times by automated ELISA plate washer with 1X wash solution. After washing traces of wash solution was removed by gentle tapping against blotting paper towel. 100  $\mu$ l of 1X prepared biotinylated antibody was added to each well and incubated for 1 hour at room temperature with gentle shaking in a motorized

**Research Article**

shaker. Solution was discarded. Wash procedure was repeated as before. 100 µl of prepared Streptavidin solution was added to each well and incubated for 45 minutes at room temperature with gentle shaking. The solution was discarded and the wash procedure was repeated as before. 100 µl of TMB One-Step Substrate Reagent was added to each well and incubated for 30 minutes at room temperature in the dark with gentle shaking. 50 µl of Stop Solution was added to each well. ELISA plate was read at 450 nm immediately using plate reader.

Data was analyzed in Microsoft Excel and SPSS software. Value of individual parameter was expressed as mean ± standard deviation. Significance of difference of the means within the groups was tested by unpaired Student’s t- test. Every where P<0.05 was considered as significant.

**RESULTS AND DISCUSSION**

**Results**

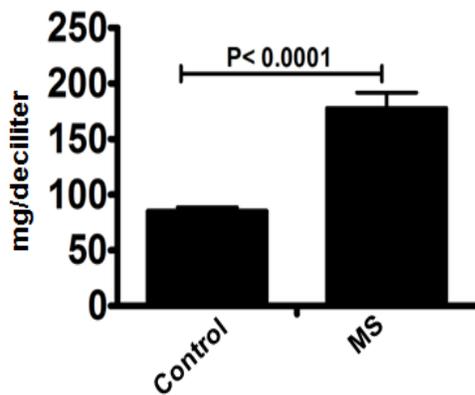
Analysed data are furnished in the tables below. Table-1 shows the physical parameters including HOMA-IR of control and test group subjects. All the subjects were euthyroids and were not reported to have history of any chronic disease. Obesity markers were also estimated which are not included in this article. Table -1 shows clear evidence that there is no significant difference in the average age of the subjects. Male and female ratios of the subjects are also close among the groups.

**Table 1: Age and gender distribution in sample population with physical parameters**

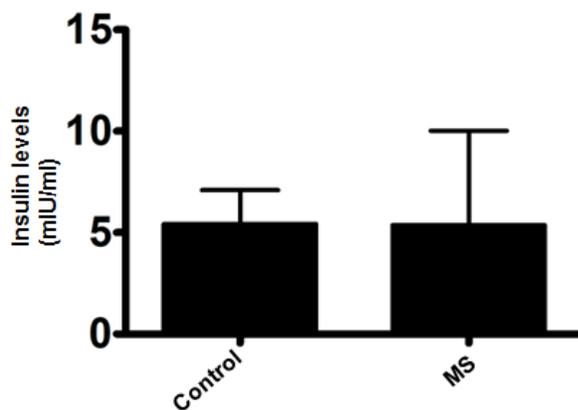
Parameter	Control Group N=150	Test Group n=150
Age in years	43.42±12.84	44.71±11.2
Male	88	82
Female	62	68
Waist/Hip ratio	0.82±0.03	0.95±0.05
HOMA-IR Score	1.148±0.35	2.34±2.0

**Table 2: Special parameters of metabolic syndrome and control subjects (n=150)**

Parameter	Unit	Control		Test		P value
		Mean	±sd	Mean	±sd	
Fasting Plasma Glucose	mg/deciliter	85.59	2.9	178.6	13.84	<0.0001
Fasting Insulin	mIU/milliliter	5.43	1.67	5.37	4.65	0.8683
Body Mass Index	Kg/meter <sup>2</sup>	23.29	3.9	36.31	4.5	<0.0001
HbA1c	NGSP (%)	5.7	0.14	10.43	0.87	<0.0001
Serum Gamma Interferon	picogram/milliliter	6.38	2.03	11.19	0.86	<0.0001
Serum Total Bilirubin	mg/deciliter	0.74	0.11	0.38	0.07	<0.0001

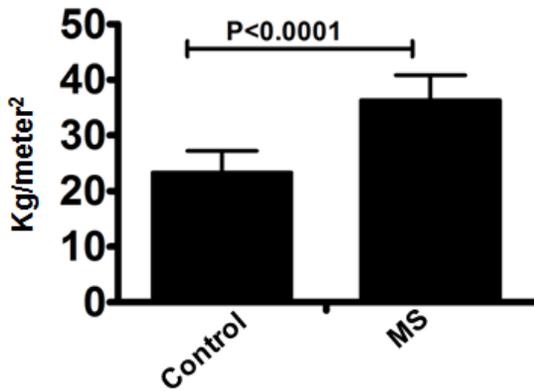


**Figure 1: Fasting Plasma Glucose levels**

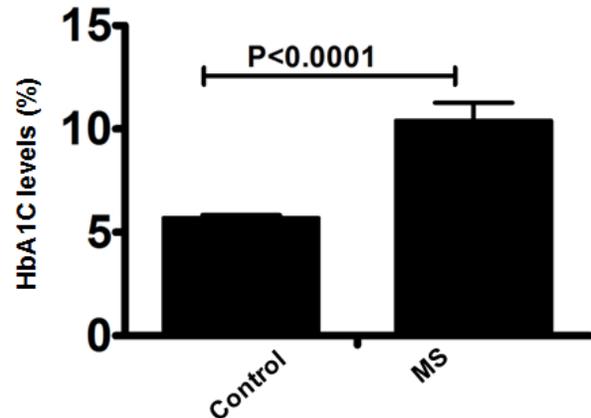


**Figure 2: Fasting Insulin Levels**

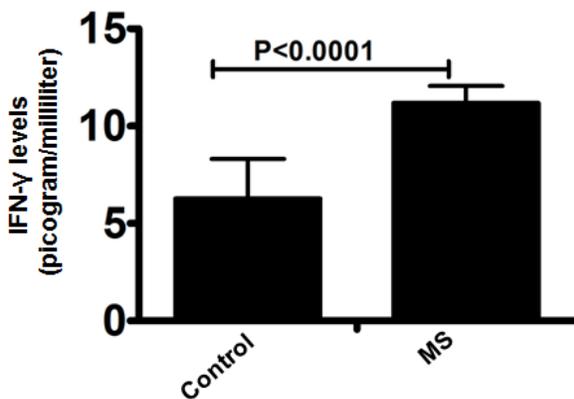
**Research Article**



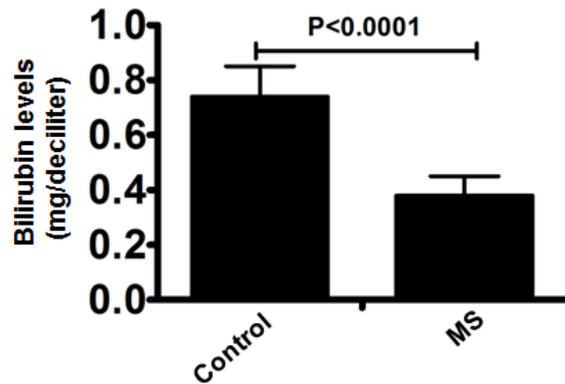
**Figure 3: Body Mass Index**



**Figure 4: Plasma HbA1C levels**



**Figure 5: IFN-γ levels**



**Figure 6: Fasting Total Bilirubin levels**

**Waist –hip Ratio:** waist –hip ratio is an important parameter for assessment of obesity (WHO Report, 2012). WHR was found to be  $0.82 \pm 0.03$  and  $0.95 \pm 0.05$  in control and metabolic syndrome groups respectively (Table 1).

**HOMA IR Score:** control and metabolic syndrome groups were found to have significantly different HOMA IR score i.e.,  $1.148 \pm 0.35$  and  $2.34 \pm 2.0$  respectively. The  $p$  value is  $<0.0001$  (Table 1).

**Fasting Plasma Glucose:** Mean and SD of fasting plasma glucose of control group was  $85.59 \pm 2.9$  mg/deciliter. Mean and SD of fasting plasma glucose of metabolic syndrome group was  $178.6 \pm 13.84$  mg/deciliter. The  $p$  value is  $<0.0001$  (Table 2, figure 1).

**Fasting Insulin:** Mean and SD of fasting insulin of control group was  $5.43 \pm 1.67$  mIU/milliliter. Mean and SD of fasting insulin of metabolic syndrome group was  $5.37 \pm 4.65$  mIU/milliliter. The  $p$  value is  $<0.8683$  (Table 2, figure 2). HOMA-IR is better parameter than fasting insulin for detection of metabolic syndrome cases (Matsuba et al., 2012).

**Body Mass Index:** Mean and SD of body mass index of control group was  $23.29 \pm 3.9$  Kg/Meter<sup>2</sup>. Mean and SD of body mass index of metabolic syndrome group was  $36.31 \pm 4.5$  Kg/Meter<sup>2</sup>. The  $p$  value is  $<0.0001$  (Table 2, figure 3).

**HbA1c:** Mean and SD of HbA1c of control group was  $5.7 \pm 0.14$  %. Mean and SD of HbA1c of metabolic syndrome group was  $10.43 \pm 0.87$  %. The  $p$  value is  $<0.0001$ . HbA1c or A1c is a diagnostic as well as prognostic parameter for of duration of hyperglycemic state of an individual (American Diabetes Association, 2011) (Table 2, figure 4).

**Serum Gamma Interferon:** Mean and SD of Serum Gamma Interferon of control group was  $6.38 \pm 2.03$  picogram/milliliter. Mean and SD of Serum Gamma Interferon of metabolic syndrome group was  $11.19 \pm 0.86$  picogram/milliliter. The  $p$  value is  $<0.0001$  (Table 2, figure 5).

### **Research Article**

**Serum Total Bilirubin:** Mean and SD of Serum Total Bilirubin of control group was  $0.74 \pm 0.11$  mg/deciliter. Mean and SD of Serum Total Bilirubin of metabolic syndrome group was  $0.38 \pm 0.07$  mg/deciliter. The *p* value is  $<0.0001$  (Table 2, figure 6).

### **Discussion**

According to NCEP (ATP 111) protocol individuals of study group are diagnosed as metabolic syndrome subjects. There is clear evidence that two groups of subjects are significantly distinct. Fasting plasma glucose, HOMA-IR, waist-hip ratio, BMI, HbA1c, interferon gamma and fasting serum total bilirubin levels are significantly distinct between control and study group each parameter is having *p* value  $<0.05$ . There is a report of inverse relationship of fasting serum bilirubin and metabolic syndrome in Korean population (Hwang and Kim, 2010; Choi et al., 2013). Our study also includes a new finding that gamma interferon, a potent inflammation marker has an inverse relationship with fasting serum total bilirubin. Metabolic syndrome is a condition in which there is increase of oxidative stress in the affected subjects. Researchers from other part of the globe also reported increased oxidative stress in metabolic syndrome (Ishizaka et al., 2005; Rutter et al., 2004; Festa et al., 2000; Gideon et al., 2007). Bilirubin functions as an anti-oxidant in association with glutathione (Sedlak et al., 2009). We propose that decrease in fasting serum total bilirubin may be caused by increased oxidative stress in metabolic syndrome.

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### **Research Article**

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