Evaluation of IFN-γ, Inflammatory Marker and Infectious Etiology as Risk Factors for Coronary Artery Disease

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ABSTRACT

Background: We evaluated the role of Chlamydia pneumoniae (C. pneumoniae) and Cytomegalovirus (CMV) infection in coronary artery disease (CAD) patients. Interferon-γ (IFN-γ) levels in serum were assessed for ongoing inflammatory activity by these microorganisms.

Materials and Methods: Persistent and chronic infection of C. pneumoniae was assessed by the level of IgA and IgG antibodies as well as 16S rRNA gene of C. pneumoniae. Chronic CMV presence was detected by IgG antibodies. IFN-γ levels in serum were analysed by ELISA.

Results: Among the 63 CAD cases, C. pneumoniae IgG antibodies were present in 34 (53.97%; 95% CI: 41.66% to 66.28%) cases, C. pneumoniae IgA antibodies were detected in 42 (66.67%; 95% CI: 55.03% to 78.31%) cases (p<0.05). PCR for detection of C. pneumoniae was done on 40 cases, among which 13 (32.5%; 95% CI: 17.99% to 47.01%) of cases were positive for 16SrRNA gene of C. pneumoniae. CMV IgG was positive in 41 (65.08%; 95% CI: 53.31% to 76.85%) cases. On assessing the patients of CAD for major traditional risk factors, 35 (55.55%) had past history of CAD, 20 (31.75%) of dyslipidemia, 19 (30.16%) gave history of smoking and 18 (28.75%) were obese. On comparison of infectious etiology and traditional risk factors, it was clear that C. pneumoniae and CMV both were more (p<0.001) strongly associated with CAD than the traditional risk factors. Mean level of IFN-γ in these patients was 32.13 pg/ml and that of controls was 11.32 pg/ml. Raised IFN-γ was observed in 39 (61.9%) patients of which 14 (35.9%) were patients of primary hypertension, 7 (17.9%) were patients of angina and 18 (46.15%) were patients of MI.

Conclusions: C. pneumoniae and chronic CMV infection appear to have a role in the pathogenesis of CAD. Elevated IFN-γ levels suggest that these pathogens aggravate the inflammatory process of atherogenesis and hasten the disease process.

Keywords: Coronary Artery Disease, Chlamydia Pneumoniae, CMV, IFN-γ

Introduction

Coronary artery disease (CAD) is a major cause of morbidity and mortality worldwide. Asian Indians are reported to have three times higher rates of incidence and prevalence of CAD when compared with western populations,[1,2] leading to a greater social and economic impact.[3,4] Various factors, such as metabolic syndrome, diabetes, dyslipidemia and possible genetic predisposition are believed to contribute to this rising epidemic in India. [1] The known risk factors of CAD are not sufficient to explain all the epidemiological variables and fluctuations of the disease. These observations have fueled renewed interest in the link between CAD and infectious agents. The hypothesis of infectious etiology of atherosclerosis is one of the most interesting areas of vascular research, it states that more than one infectious agent could play a role in atherogenesis and atherothrombosis. This entire process may involve pro-inflammatory mechanisms like interleukins and cytokines.[5] Infectious agents are considered as significant factors in the pathogenesis of atherosclerosis because atherogenic processes resemble chronic inflammation—a process that may be promoted by microorganisms like Chlamydia pneumoniae (C. pneumoniae) and Cytomegalovirus (CMV). The hypothesis of infectious etiology is very attractive because it would provide us with a powerful platform for prevention and treatment of a disease which represents an escalating public health problem worldwide and even more so in India. Even after extensive research the role of infectious agents in CAD still remains an enigma.[3]

Till date the question of whether C. pneumoniae and CMV have a direct role in CAD or are innocent bystanders remains unresolved but it continues to be an ever intriguing possibility. If a temporal or casual relationship is established between C. pneumoniae and CMV with CAD it would bracket CAD in the rare category of “Preventable Chronic Diseases” — where a simple antibiotic or antiviral may go a long way in preventing this disease. The impact
on public health and preventive medicine in terms of mortality, morbidity, cost effectiveness and reduction in interventional cardiology procedures will be dramatic. Both C. pneumoniae and CMV are intracellular pathogens, we felt that IFN-γ would be the first key cytokine to rise that could in turn activate the chain of events that could unleash a “cytokine-storm” causing a worsening of atherogenesis. In chronic diseases such as in CAD, biochemical reactions induced by IFN-γ may have detrimental consequences for host cells. Therefore, the aim of this study was to study the prevalence and association of C. pneumoniae and Cytomegalovirus with CAD and evaluation of inflammatory marker (IFN-γ) as potential predictors of risk for cardiovascular events.

Materials and Methods
A prospective study was conducted at the Departments of Microbiology and Medicine, Jawaharlal Nehru Medical College, AMU, Aligarh, from November 2008 to September 2009. Study group consisted of 63 randomly selected patients with CAD. A written and informed consent was obtained from all subjects. Ethical clearance was obtained by ethical committee of JNMC. Control arm consisted of 30 age and sex matched healthy volunteers. Controls were confirmed as having coronary artery disease by testing for cardiac injury markers such as creatinine phosphokinase, Troponin-T, Troponin-I, coronary angiography and stress testing. Exclusion criteria were as follows: patients with secondary hypertension (renal causes, endocrine causes, drug induced); patients with pulmonary involvement (e.g. chronic obstructive pulmonary disease [COPD], pneumonia, etc.) or with any other septic foci; and patients with history of antibiotic intake in the previous one month.

After obtaining informed consent, a detailed clinical history and medical examination was performed. Blood pressure was measured for all subjects in a quiet room using a sphygmomanometer and according to standard methods. A blood pressure reading higher than 140/90 mm Hg on two or more occasions measured at least eight hours apart was considered as hypertension.

A sample of venous blood was also collected from each subject in the study after a 10-hour overnight fast for determination of their fasting total cholesterol and blood sugar using the enzymatic colorimetric method. Hypercholesterolemia was defined as a total cholesterol level of ≥ 6.2 mmol/L. Blood sugar estimation was also performed in all patients and non-diabetic subjects were given 75 g of glucose after an overnight fast for an oral glucose tolerance test. A case of diabetes mellitus was defined as a subject with a known history or with a two-hour plasma glucose concentration ≥ 11.1 mmol/L by the oral glucose tolerance test. Electrocardiograph (ECG) was performed in all cases.

Collection of Sample for Serology and Amplification of 16SrRNA: Taking all aseptic precautions, 10 ml blood was obtained by venipuncture, 5 ml in an EDTA vial and 5 ml in plain vial. Serum was separated from the samples in a plain vials by centrifugation and stored at -20°C. Samples in the EDTA vials were used for DNA extraction by the phenol chloroform extraction method.

Serology: IgA antibodies to C. pneumoniae were detected by Sero CP™ IgA (Savlon Diagnostics, St. Ashdod, Israel). IgG antibodies to C. pneumoniae was detected by sero CP™ IgG obtained from SAVYON DIAGNOSTICS, Israel. IgG antibodies to Cytomegalovirus were detected by kit obtained from Calbiotech Diagnostics, USA. The human IFN-γ ELISA was used for the in-vitro quantitative determination of interferon gamma (IFN-γ) in human serum. The kit was obtained from Diamclone, Stamford, Connecticut, USA.

Amplification: PCR for detection of the 16SrRNA gene of C. pneumoniae was performed for 40 cases and 11 controls. Primers were ordered from Fermentas, Life Sciences, Maryland, USA. The sequences for primer
pairs of 16SrRNA were as follows: 5' GTG GAT G 3', 832 bp 5' CGA CAC GGA TGG GTG TG 3' using Master Mix from Fermentas, Life sciences, Maryland USA.

Statistical Analysis: The data were entered and analyzed using the Statistical Package for the Social Sciences version 17.0 (SPSS, Chicago, USA). Continuous variables were presented using mean ± standard deviation while categorical variables were presented as proportions. Categorical comparisons were made using Chi-square/Kruskal-Wallis test where appropriate and a P value of less than 0.05 was considered significant.

Result
Among the 63 CAD cases, highest number of patients [29(44.4%)] were of MI followed closely by primary hypertension (HTN) 21(33.3%) and angina 14(22%) (Table 1). Males 38(60%) outnumbered females 25(40%). No difference was found in hypertensives, but males were significantly more in angina patients (p < 0.01) and MI patients (p < 0.05). The mean age distribution was 52.7 ± 8.9 years in cases and 54.01 ± 8.1 years in controls.

While C. pneumoniae IgG antibodies were detected in 34(53.97%; 95% CI: ±12.31, range 41.66%-66.28%) cases, C. pneumoniae IgA antibodies were detected in a larger no. 42(66.67%; 95% CI: ±11.64, range 55.03%-78.31%) cases (p<0.05). PCR for detection of C. pneumoniae was done on 40 cases and 11 controls. 13 (32.5%; 95% CI: ±14.51, range 17.99%-47.01%) of cases were positive for 16 SrRNA gene of C. pneumoniae. None of the controls were positive. CMV IgG was detected in 41(65.08%; 95% CI: ±11.71, range 53.31%-76.85%) cases.

On comparing C. pneumoniae IgA and IgG seropositivity, it was observed that IgA antibodies were seen in a larger percentage of patients with MI (71.43%) and HTN (61.9%) as compared to IgG prevalence in MI (53.47%) and HTN (42.5%). However C. pneumoniae IgG antibodies predominated in patients with angina (71.43%) as compared to IgA (64.29%) Table 1. The sensitivity (Sn), specificity (Sp), positive predictive value (PPV), negative predictive value (NPV) and odds ratio (OR) for serological tests are depicted in Table 2.

C. pneumoniae IgA antibodies were detected in 42(66.66%) cases and 12(41.37%) controls. This test had a Sn of 66.66%, Sp of 51.72%, PPV of 63.63%, NPV of 41.67% and OR of 2:1. Association of C. pneumoniae IgA with CVD was statistically significant (p < 0.05). Among these angina patients had higher sensitivity (64.29%) and NPV (77.27%), whereas MI cases had higher PPV (56.25%) and an odds ratio of 3.54.

C. pneumoniae IgG antibodies were detected in 34(53.97%) cases and 14(48.28%) controls. This test had a Sn of 53.97%, Sp of 51.72%, PPV of 70.83%, NPV of 31.09% and an OR of 1.26. However the rate of positivity in cases and controls was statistically not significant. Among C. pneumoniae IgG seropositives, MI patients had higher Sn (53.57%) and PPV (51.72%), however angina patients showed higher NPV (78.95%) and an OR of 2.86. Out of 63 cases 41(65.08%) were positive for IgG antibodies against CMV and 8(27.58%) of the controls were IgG positive (p<0.01). Sn of this test was 65.08%, Sp was 72.41%, PPV (83.67%), NPV (48.84%), and an OR of 4.89. Among CMV IgG seropositives, MI patients showed higher Sn (67.86%), PPV (70.37%) and OR (5.54) whereas higher NPV (80.77%) was observed in angina patients.

High OR associated with C. pneumoniae IgA antibodies in patients of CAD suggests their role in its pathogenesis. Infact OR was highest in MI (3.54) and lowest in hypertensive (2.3) patients which is again pointing to a spectrum of diseases. Similar association was observed in CMV, however OR were far higher in all three disease groups, especially so in MI patients (5.54).

16SrRNA Gene of C. Pneumoniae: PCR for detection of C. pneumoniae was performed for 40 cases (15 of MI & HTN each and 10 of angina) and 11 controls. Thirteen (32.5%) of the cases and none of the controls were positive on PCR. 6 (40%) of 15 hypertensives 5(33.3%) of 15 patients of MI and 2(20%) of 10 angina patients were PCR positive. Among 13 PCR positive cases, majority (38.46%) demonstrated C. pneumoniae IgA antibodies and only 5(38.4%) were positive for C. pneumoniae IgG (Figure 1).

Assessment of Infection and Traditional Risk Factors for Development of CAD: On assessing the patients of CAD for major traditional risk factors, 35 (55.5%) had past history of CAD, 20 (31.75%) of dyslipidemia, 19 (30.16%) gave history of smoking and 18 (28.75%) were obese.

On comparison of infectious etiology (as putative risk factors) and traditional risk factors, it was clear that C. pneumoniae and CMV both were more (p<0.001) strongly associated with CAD than the traditional risk factors (Table 3).

Estimation of Interferon-γ (IFN-γ) Levels: IFN-γ was raised in most of the patients with CVD. The mean level of IFN-γ in these patients was 32.13pg/ml and that of controls was 11.32pg/ml. Raised IFN-γ was observed in 39(61.9%) patients of which 14(35.9%) were patients of primary hypertension, 7(17.9%) were patients of angina and 18
(46.15%) were patients of MI. The mean level of IFN-γ in hypertensives was 3.39 pg/ml, in patients of angina was 18.5 pg/ml and on MI patients was 23.89 pg/ml. Of these, 30 (76.92%) were C. pneumoniae IgA seropositive, 27 (69.23%) were C. pneumoniae IgG seropositive and 26(66.7%) CMV IgG seropositive. Correlation of Interferon-γ with C. pneumoniae and Cytomegalovirus seropositivity in different study groups is given in Figure 2.

Table 1: Prevalence of C. pneumoniae and CMV in different disease groups

<table>
<thead>
<tr>
<th>STUDY GROUP</th>
<th>C. pneumoniae IgG n=34</th>
<th>C. pneumoniae IgA n=42</th>
<th>C. pneumoniae PCR positivity n=13</th>
<th>Cytomegalovirus IgG n=41</th>
</tr>
</thead>
<tbody>
<tr>
<td>Primary Hypertension</td>
<td>9(42.85%)</td>
<td>13(61.90%)</td>
<td>6(40%)</td>
<td>13(61.90%)</td>
</tr>
<tr>
<td>Angina n=14</td>
<td>10(71.43%)</td>
<td>9(64.29%)</td>
<td>2(20%)</td>
<td>9(64.29%)</td>
</tr>
<tr>
<td>Myocardial Infarction n=28</td>
<td>15(53.57%)</td>
<td>20(71.43%)</td>
<td>5(33%)</td>
<td>19(67.86%)</td>
</tr>
<tr>
<td>CONTROLS n=29</td>
<td>14(48.27%)</td>
<td>12(41.37%)</td>
<td>0</td>
<td>8(27.5%)</td>
</tr>
</tbody>
</table>

Table 2: Comparison of performance of different serological tests used. (Cases=63, Controls=30)

<table>
<thead>
<tr>
<th>Serology</th>
<th>Sensitivity</th>
<th>Specificity</th>
<th>Positive Predictive value</th>
<th>Negative Predictive value</th>
<th>Odds ratio</th>
</tr>
</thead>
<tbody>
<tr>
<td>C.pn IgG</td>
<td>53.97%</td>
<td>51.72%</td>
<td>70.83%</td>
<td>31.09%</td>
<td>1.26</td>
</tr>
<tr>
<td>HTN</td>
<td>42.86%</td>
<td>51.72%</td>
<td>39.13%</td>
<td>55.55%</td>
<td>0.80</td>
</tr>
<tr>
<td>Ang</td>
<td>71.43%</td>
<td>51.72%</td>
<td>41.66%</td>
<td>78.95%</td>
<td>2.86</td>
</tr>
<tr>
<td>MI</td>
<td>53.57%</td>
<td>51.72%</td>
<td>51.72%</td>
<td>53.57%</td>
<td>1.24</td>
</tr>
<tr>
<td>C. pn IgA</td>
<td>66.6%</td>
<td>58.62%</td>
<td>63.63%</td>
<td>41.67%</td>
<td>2.14</td>
</tr>
<tr>
<td>HTN</td>
<td>61.9%</td>
<td>58.62%</td>
<td>52%</td>
<td>68%</td>
<td>2.3</td>
</tr>
<tr>
<td>Ang</td>
<td>64.29%</td>
<td>58.62%</td>
<td>42.86%</td>
<td>77.27%</td>
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</tr>
<tr>
<td>MI</td>
<td>71.43%</td>
<td>58.62%</td>
<td>56.25%</td>
<td>68%</td>
<td>3.54</td>
</tr>
<tr>
<td>CMV IgG</td>
<td>65.08%</td>
<td>72.41%</td>
<td>83.67%</td>
<td>48.84%</td>
<td>4.89</td>
</tr>
<tr>
<td>HTN</td>
<td>61.90%</td>
<td>72.41%</td>
<td>61.90%</td>
<td>72.41%</td>
<td>4.27</td>
</tr>
</tbody>
</table>

Table 3: Evaluation of traditional risk factors and C. pneumoniae & CMV infection as risk factor in CAD.

<table>
<thead>
<tr>
<th>Risk Factors</th>
<th>Positive Cases n=63</th>
</tr>
</thead>
<tbody>
<tr>
<td>Traditional Risk Factors</td>
<td></td>
</tr>
<tr>
<td>Smoking</td>
<td>19(30.16%)</td>
</tr>
<tr>
<td>Diabetes</td>
<td>5(7.94%)</td>
</tr>
<tr>
<td>Dyslipidemia</td>
<td>20(31.75%)</td>
</tr>
<tr>
<td>Obesity</td>
<td>18(28.75%)</td>
</tr>
<tr>
<td>Family History</td>
<td>12(19.05%)</td>
</tr>
<tr>
<td>Past History</td>
<td>35(55.55%)</td>
</tr>
<tr>
<td>Infectious Etiology as Risk Factor</td>
<td></td>
</tr>
<tr>
<td>C.pneumoniae IgA</td>
<td>42(66.66%)</td>
</tr>
</tbody>
</table>
Discussion

Chlamydia pneumoniae. C. pneumoniae is a gram-negative intracellular bacterium. An association of coronary heart disease and infections with C. pneumoniae, an important respiratory pathogen, has been initially found by sero-epidemiologic studies. The presence of C. pneumoniae has been shown in atheromatous plaques of coronary and carotid arteries [6,7,8]. An immune reaction to chlamydial Hsp60 has been shown in coronary atheroma tissue, especially in macrophages and foam cells [9]. In human carotid artery plaques, C. pneumoniae reactive T lymphocytes have been identified [10]. A further possible role of C. pneumoniae in the pathogenesis of arteriosclerosis could be that chronic or acute chlamydial infection anywhere in the body activates arteriosclerotic plaques. An association between immunoglobulin A (IgA), but not IgG, antibodies to C. pneumoniae and the subsequent risk of death from ischemic heart disease, the risk of ischemic

Fig. 1: C. pneumoniae PCR positivity according to study Group and seropositivity (n=13)

Fig. 2: Correlation of Interferon-γ with Chlamydia pneumonia and cytomegalovirus seropositivity in different study groups
stroke, and the severity of arteriosclerosis of carotid or femoral arteries has been found[11]. In patients undergoing carotid endarterectomy, the plaques were investigated for C.pneumoniae and the blood was examined for antichlamydial antibodies. Associated with symptomatic disease was a high level of antichlamydial IgA, whereas no association between C. pneumoniae presence in the plaque and symptomatic disease could be detected[12]. Furthermore, there are indications that seropositivity for C. pneumoniae might enhance the atherogenic effects of other vascular injuries. A case-control study suggests that the proatherogenic effects of lipoprotein may be enhanced or partly mediated through the formation of circulating immune complexes containing C. pneumoniae specific IgG antibodies[13]. The simultaneous presence of high levels of antibodies to C. pneumoniae and of antibodies to human Hsp60 have been found to substantially increase the risk for coronary arteriosclerosis[14].

Cytomegalovirus (CMV): CMV infection of endothelial cells may increase cellular proliferation and inhibit apoptosis of infected smooth-muscle cells, thereby contributing to an increase in the mass of arteriosclerotic lesions. Furthermore, individuals infected with CMV have impaired endothelium-mediated coronary vasodilator responses[15]. Two studies in patients undergoing coronary angiography have shown that CMV elicits a subclinical inflammatory response, but only in certain individuals, and that individuals with an inflammatory response appear to be susceptible to the atherogenic effects of CMV, whereas those without such a response appear to be resistant[16,17]. Postulating that sex might have an effect on patterns of inflammatory and immune responses to CMV infection, a study in patients evaluated for coronary artery disease found that, in men, CMV appears to contribute to coronary artery disease risk, in so far as it predisposes to inflammation. In women, other mechanisms, possibly related to the type of immune response generated by the host, appear to be responsible for the proatherogenic effects of CMV[18].

Recently, total number of deaths due to CAD has increased globally to 17.5 million from 14.4 million in 1990. More than 80 percent of the deaths occurred in low and middle income countries. [19,20,21] WHO estimates there will be about 20 million CAD deaths in 2015, accounting for 30 percent of all deaths worldwide.[22] There are well known risk factors for CAD such as hypertension, diabetes mellitus and obesity.[23] However in approximately 50% cases of CAD no identifiable risk factors are observed.[24] Consequently, despite advances in the pharmacological control of hypertension and diabetes, as well as mounting efforts in discouraging smoking, the decrease in the clinical complications of atherosclerosis has been modest, indicating that there are as yet unknown contributory factors in the pathogenesis of atherosclerosis. This has spurred passionate search for new risk factors such as infectious agents, inflammatory markers and coagulation factors. [25] In a study from India, C. pneumoniae was detected in 73.7% of CAD patients.[26]

In this we study we assessed the role of C. pneumoniae and CMV in pathogenesis of CAD. C. pneumoniae IgA antibodies were detected in 42 (66.66%) cases and 12 (41.37%) controls. Primary C. pneumoniae infection is characterized by a significant immunoglobulin (IgM) antibody response, a delayed IgG titer and a low IgA level. Secondary infection with C. pneumoniae infection is notable for an absence of IgM antibody and elevated IgA and IgG levels. [27] It has been suggested that serum IgA antibody titers to C. pneumoniae may be more predictive of coronary heart disease than are IgG titres because IgA antibodies have different biological characteristics(such as shorter half-life) and may be better indicators of persistent chlamydial infection in certain chronic diseases. [28] This test had a Sn of 66.66%, Sp of 51.72%, PPV of 63.63%, NPV of 41.67% and an OR of 2:1.4.Detection of C. pneumoniae in CAD cases was statistically significant (p < 0.05). A few small studies have reported stronger associations of C. pneumoniae IgA titers with MI [29] or stroke.[28] C. pneumoniae IgG antibodies were detected in a lower no. of 34(53.97%) cases and 14(48.28%) controls. This test had a Sn of 53.97%, Sp of 51.72%, PPV of 70.83%, NPV of 31.09% and an OR of 1.26. However the rate of positivity in cases and controls was statistically not significant. Maximum IgG seropositives 15 (53.57%) were among MI patients followed by 10(71.43%) angina patients and 9(42.85%) in hypertensives. No statistical difference was found in comparing seropositivity in different study groups within themselves and with healthy controls. Our findings are in contrast to another study where the investigators showed that cases had a higher frequency of IgG seropositivity than did non cases.[29] However another study showed no evidence of association between C. pneumoniae IgG seropositivity and risks of future MI. [30] A great majority of cross-sectional and retrospective studies suggest that a positive association for C. pneumoniae have relied primarily or exclusively on serology to determine exposure status.[31] Previous studies have used the presence of IgG and IgA antibody to C. pneumoniae as a proxy for chronic infection, but the validity of this approach is uncertain. C. pneumoniae infection generally starts in the respiratory tract and probably disseminates systemically in the blood stream within alveolar macrophages. In both laboratory animals and human subjects, the persistence of
vascular *C. pneumoniae* DNA has been associated with the presence of viable organism.[32] We therefore assessed the prevalence of *C. pneumoniae* DNA in the blood mononuclear cells of 40 patients of CVD and correlated the results with clinical and other parameters. PCR for detection of *C. pneumoniae* was done on 40 cases and 11 controls. 13(32.5%) cases were positive by PCR (p<0.05).None of the controls were positive. Boman et al, working in Sweden first demonstrated the detection of *C. pneumoniae* DNA in PMBC of cardiac patients.[33] They found a prevalence of *C. pneumoniae* 59% in patients. The largest of these nine studies examined 1,205 British patients undergoing coronary angiography and was conducted by Wong et al. [14] They found a low prevalence of 8.7% in patients with angiographic ally proven CAD, versus 7.2% in those with normal coronary arteries. There is limited epidemiological evidence relating CMV specifically to primary coronary atheroma in immune-competent individuals. [35]

Out of 63 cases 41(65.08%) were positive for IgG antibodies against CMV and 8(27.58%) of 29 controls were IgG positive. Detection of CMV was significantly more (p<0.01) in cases than controls. Sn of this test was 65.08%,Sp was 72.41%,PPV(83.67%),NPV(48.84%) and an OR of 4.89. Among CMV IgG seropositives, MI patients showed higher Sn (67.86%),PPV(70.37%) and OR(5.54) whereas higher NPV (80.77%) was observed in angina patients. The results of a study by Grayston JT et al [36] demonstrated a similar prevalence of seropositivity in patients with coronary artery disease and in an age and sex matched control group; however, anti-CMV IgG antibody titers were higher in patients with coronary artery disease (52% of the patients had ≥ 1:800 anti-CMV titres, whereas only 16% of the control group had high antibody titers).

A quantitative analysis if IFN-γ levels in CVD patients was done using ELISA. A total of 39 patients were assessed, out of which 38.9% were primary hypertensives, 17.95% were angina patients and majority 43.59 were MI patients. Most of these patients were *C. pneumoniae* or CMV serology positive. The mean value of IFN-γ in CAD cases was 15.3±51.2 pg/ml. The mean value of IFN-γ in hypertensives was 3.39pg/ml, of Angina patients was 18.5pg/ml and of MI patients was 23.89pg/ml. The mean of controls was 5.15±4.16 pg/ml. A study found that mononuclear leukocytes from patients with stable as well as unstable angina pectoris secrete significantly more IFN-γ and TNF-α than do the mononuclear cells from healthy but elderly control subjects.[37,38] An interesting trend of increasing value of IFN-γ with the severity and chronicity of CVD is clearly visible in our study. This may suggest an increasing role of inflammation and inflammatory markers not only in the initiation but also in the progression of CVD. But in contrast to our study, Hem C.J. reported IFN-γ was significantly lower (P=0.006) in *C. pneumoniae* IgA negative patients.[39,40,41]

Although there were 30(76.92%) patients who had *C. pneumoniae* IgA and IFN-γ elevated and 27(69.23%) patients who had *C. pneumoniae* IgG and IFN-γ elevated there was considerable overlapping, so there were actually only 22(56.41%) patients had *C. pneumoniae* IgA and IgG with IFN-γ elevated. Only 15(38.46%) patients had *C. pneumoniae* IgA, IgG and CMV IgG elevated with IFN-γ. So contrary to our expectation number of patients co-infected with *C. pneumoniae* and CMV with raised IFN-γ were not more. Thus we re-evaluated our understanding of the inflammatory potential of *C. pneumoniae*. Patients positive for *C. pneumoniae* IgA were more susceptible to elevated Th1 response due to increased levels of IFN-γ. We hypothesize that such patients ( *C. pneumoniae* IgA positive with raised IFN-γ) risk of adverse clinical outcome.

**Conclusion**

In this study *C. pneumoniae* and CMV both appear to have a role in pathogenesis of CAD. However it would be more sapien to take them as added risk factors in causation of CAD. In fact compared to traditional risk factors, *C. pneumoniae* and CMV were more significant risk factors in the present study. Moreover, their potential to trigger an inflammatory response superimposed on the natural atherogenetic process may hasten the pathogenesis of CAD. On analyzing association of infection with IFN-γ, we observed that persistent infection with raised interferon correlated well with chronicity and severity of disease. Large scale prospective studies with longer follow-up and confirmation of disease status by imaging techniques like angiography are needed to further elucidate the role of these agents. If their role is proven without doubt then we would have in our hands a “major modifiable risk factor”. This would radically change the way we prevent and treat cardiovascular diseases. A window for research for similar “modifiable risk factors” in other chronic diseases will also be thrown open.

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