

Association of Cytomegalovirus (CMV) Infection with Guillain-Barré Syndrome (GBS) In Tertiary Care Hospital (BSMMU) of Bangladesh

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Abstract:

Background: Guillain-Barré syndrome (GBS) usually preceded by infections, in particular cytomegalovirus (CMV). It may occur by primary infection, reinfection or by reactivation of CMV. **Objective:** The aim of the present study was to evaluate the association of Guillain-Barré syndrome (GBS) with Cytomegalovirus (CMV) infection. **Methodology:** This case control study was carried out in the indoor and outpatient Department of Neurology at Bangabandhu Sheikh Mujib Medical University (BSMMU), Dhaka from 1st January 2010 to 31st December 2011 for the duration of two years. All patients with GBS, who attended in neurology OPD or inpatient department at BSMMU during the study period, fulfilling the inclusion and exclusion criterias were included in this study. Age & sex matched volunteers, patients attendants, patients other than GBS who were non-diabetic, had no renal or hepatic diseases or family history of polyneuropathy were included in control group. **Results:** A total number of 78 respondents of which 39 patients were taken as cases and rest 39 were taken as controls who appeared in neurology OPD or inpatient department at BSMMU during the study period, fulfilling the inclusion and exclusion criterias were included in this study. The mean age \pm SD of case and control groups were 30.82 ± 12.56 and 31.00 ± 12.77 years respectively ($p=0.950$). In case group the history of respiratory tract infection was present in 46.2% cases and absent in control group ($p=0.001$). In case group the history of gastroenteritis was present in 28.2% cases and absent in control group ($p=0.001$). In case group the history of fever was present in 30.8% cases and absent in control group ($p=0.001$). Anti-CMV IgM antibody was positive in 5.1% cases. Four fold rise of IgG in case group was present in 10.3% cases and absent in control group ($p=0.040$). Confirmed CMV infected GBS cases were 15.4% and absent in control group ($p=0.011$). **Conclusion:** The findings of this study permit to conclude that there is a significant association of Guillain-Barré syndrome (GBS) with Cytomegalovirus (CMV) infection.

Keywords: Cytomegalovirus (CMV), Guillain-Barré Syndrome (GBS).

Introduction:

Guillain-Barré syndrome (GBS) now ranks as the most frequent cause of acute flaccid paralysis since the near-elimination of poliomyelitis

throughout the world and its median annual incidence is 1 to 2 per 100,000 populations¹. According to an epidemiologic survey, the average annual incidence of GBS in the United States is

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3.0 cases per 100,000 populations. With poliomyelitis under control in developed countries, GBS is the most important cause of acute flaccid paralysis.

GBS remains a diagnosis made primarily through the assessment of clinical history and findings².

In epidemiologic surveys, the overall death rate related to GBS ranges from 2-12% of patients. Deaths usually occur in ventilator-dependent patients, resulting from such complications as pneumonia, sepsis, adult respiratory distress syndrome, and less frequently, autonomic dysfunction³. Although the classic description of GBS is that of a demyelinating neuropathy with ascending weakness, many clinical variants have been well documented in the medical literature.

GBS is a post-infectious, immune-mediated disease. Cellular and humoral immune mechanisms probably play a role in its development. Most patients report an infectious illness in the weeks prior to the onset of GBS. Many of the identified infectious agents are thought to induce antibody production against specific gangliosides and glycolipids, such as GM₁, GM₂ and GD_{1a} etc, distributed throughout the myelin in the peripheral nervous system⁴. The favored hypothesis is that the immune response to certain infective agents in some people may trigger cross reactive immunity, with initially one or more myelin or axonal antigens leading to an autoimmune attack on the nerve tissue. Antiglycolipid antibodies have often been found in affected patients⁵.

The pathophysiologic mechanism of an antecedent illness and of GBS can be typified by *Campylobacter jejuni* infections⁶.

A preceding CMV infection with high titres of IgM antibody has been implicated in 10-15% of the patients with GBS⁷.

Cytomegalovirus is a member of a herpes virus group. It is a DNA virus having double stranded DNA. It causes primary infection, reactivation or reinfection. Route of transmission are by breast milk, saliva, sexual transmission, blood transmission, organ transplantation and droplet infection etc⁸.

In primary infection IgM against CMV develop and persist for 3-4 months, but in case of reinfection or reactivation IgM is not usually found, IgG is found.

IgG is persist for life long⁷. It has also suggested raised concentrations of antibodies to ganglioside GM2 in patients with GBS after cytomegalo virus (CMV) infection⁹. The association between anti-ganglioside antibody responses and Guillain-Barré syndrome (GBS) after a recent cytomegalovirus (CMV) infection Khalili SA et al.¹⁰, conducted a study. They concluded that antibodies to ganglioside GM2 are often associated with GBS after CMV infection, but their relevance is not known. It is unlikely that CMV infection and anti-ganglioside GM2 antibodies are solely responsible and an additional factor is required to elicit GBS¹⁰.

Guillain-Barré syndrome may occur by primary infection or by reinfection or reactivation of CMV¹¹. To study the association of cytomegalo virus infection with Guillain-Barre syndrome needs diagnosis of CMV infection which required at least one of the following laboratory method:- serology, specific intrathecal antibody production, virus isolation, direct detection of CMV PP65 antigen in blood, CMV culture, biopsy, positive specific immunohistochemical staining, polymerase chain reaction (PCR) assay etc⁸.

Serological studies indicating an acute CMV infection includes:

1. The presence of positive IgM anti CMV antibodies with undetectable CMV specific IgG antibodies, or
2. Presence of CMV specific IgG antibodies of low avidity in the presence or absence of virus specific IgM antibodies¹¹.
3. Presence of increase in the titre of IgG anti CMV antibodies in paired sample obtained during the infection⁸.

This study was evaluated by serological test. The proposed study was evaluated the relationship of CMV infection with GBS. This study was focused on new insights into the epidemiology and information concerning the relationship between CMV infection and GBS.

Methods:

This is a case-control study. The study was carried out in the Department of Neurology, Bangabandhu Sheikh Mujib Medical University (BSMMU), Dhaka. This study was conducted from 1st January 2010 to

31st December 2011 for the duration of two years. All patients with GBS, who attended in neurology department, BSMMU during the study period, fulfilling the inclusion and exclusion criteria were included in this study. Age & sex matched volunteers, patient's attendants who were non-diabetic, had no renal or hepatic diseases or family history polyneuropathy were included in control group. A total number of 78 study subject, 39 patients presented with Guillain-Barre syndrome and 39 controls were enrolled in this study by purposive sampling.

All data were compiled and edited meticulously by thorough checking and rechecking. All omissions and inconsistencies were corrected and were removed methodically.

All data were recorded systematically in preformed data collection form (questionnaire) and quantitative data was expressed as mean and standard deviation, and qualitative data was expressed as frequency distribution and percentage. Statistical analysis was performed by using Statistical Package for Social Science (SPSS) for windows version 12.0. 95% confidence limit was taken. Probability value <0.05 was considered as level of significance.

Results and Observations:

A total 78 number of study subjects 39 patients with GBS as cases and 39 volunteers, were taken as control who were attended in neurology department at BSMMU during the study period, fulfilling the inclusion and exclusion criteria were included in this study.

Table-I

Distribution of the study subjects by IgG more than 4 fold rise (CMV positive) in the 2nd sample (n=78) . Sample – Blood

4 fold rise of IgG	Group		p value
	Case (n=39)	Control (n=39)	
Positive	4 (10.3) [#]	0 (0.0)	0.124
Negative	35 (89.7)	39(100.0)	
Total	39(100)	39(100)	

*Chi-square test (after Yates correction) was done to measure the level of significance.

[#]Figure within parentheses indicates in percentage.

4 fold or more rise of IgG titre in 2nd sample considered as CMV positive.

Table I shows the distribution of the study subject by IgG more than 4 fold rise (CMV positive) in the 2nd sample. In the case group 4 (10.3%) cases showed 4 fold rise of IgG and the rest 35 (89.7%) cases showed negative results. In control group 4 fold rise of IgG was absent in all controls which was 39 (100.0%). The difference between case and control was not statistically significant (p=0.124).

In first case serum anti CMV IgG level in 1st sample was 85.1AU/ml and in 2nd sample was 854.1 AU/ml.

In second case serum anti CMV IgG level in 1st sample was 418AU/ml and in 2nd sample was 1219 AU/ml.

In third case serum anti CMV IgG level in 1st sample was 30.7AU/ml and in 2nd sample was 270AU/ml.

In fourth case serum anti CMV IgG level in 1st sample was 36.1AU/ml and in 2nd sample was 320 AU/ml.

Table-II

Distribution of the study subjects by Anti-CMV IgM (n=78). Sample – Blood

Anti CMV IgM	Group		p value
	Case (n=39)	Control (n=39)	
Positive	2(5.1)	0(0.0)	0.474
Negative	37(94.9)	39(100.0)	
Total	39(100.0)	39(100.0)	

*Chi-square test (after Yates correction) was done to measure the level of significance.

[#]Figure within parentheses indicates in percentage.

Serum anti CMV IgM positive considered as CMV positive.

Table II shows the distribution of the study subjects by anti CMV IgM. In case group anti CMV IgM was positive in 2 (5.1%) cases and the rest 37 (94.9%) cases were anti CMV IgM negative. In control group anti CMV IgM was negative in all 39 (100.0%) controls. The difference between case and control was not statistically significant (p=0.474).

Table-III
Distribution of the Study subjects by CMV detection (n=78)

CMV detection	Group		p value
	Case (n=39)	Control (n=39)	
Positive	6(15.4)#	0(0.0)	0.034
Negative	33(84.6)	39(100.0)	
Total	39(100)	39(100)	

*Chi-square test (after Yates correction) was done to measure the level of significance.

#Figure within parentheses indicates in percentage.

CMV positive (by IgM = 02 and by 4 fold rise of IgG = 04) = 06.

Table III shows the distribution of the study subjects by CMV detection. In case group CMV was positive in 6 (15.4%) cases and the rest 33 (84.6%) cases were CMV negative. In control group CMV was negative in all 39 (100.0%) controls. The difference between case and control groups was statistically significant (p=0.034).

Table-IV
Distribution of the study subjects by Anti-CMV IgG Sample-1 (n=78). Sample – Blood

Anti-CMV IgG (Sample-1)	Group		p value
	Case (n=39)	Control (n=39)	
Positive	33(84.6)	37(94.9)	0.135
Negative	6(15.4)	2(5.1)	
Total	39(100.0)	39(100.0)	

*Chi-square test was done to measure the level of significance.

Figure within parentheses indicates in percentage.

Table IV shows the distribution of the study subjects by Anti-CMV IgG Sample-1. In case group Anti-CMV IgG (Sample-1) was positive in 33(84.6%) cases and negative in 6(15.4%) cases. In control group Anti-CMV IgG (Sample-1) was positive in 37(94.9%) controls and negative in 2(5.1%) controls. The difference between case and control was not statistically significant (p=0.135).

Table-V
Distribution of the study subjects by Anti-CMV IgG Sample-2 (n=78). Sample – Blood

Anti-CMV IgG (Sample-2)	Group		p value
	Case (n=39)	Control (n=39)	
Positive	35(89.7)	38(97.4)	0.165
Negative	4(10.3)	1(2.6)	
Total	39(100.0)	39(100.0)	

*Chi-square test was done to measure the level of significance.

Figure within parentheses indicates in percentage.

Table V shows the distribution of the study subjects by Anti-CMV IgG Sample-2. In case group Anti-CMV IgG (Sample-2) was positive in 35(89.7%) cases and negative in 4(10.3%) cases. In control group Anti-CMV IgG (Sample-2) was positive in 38(97.4%) controls and negative in 1(2.6%) control. The difference between case and control was not statistically significant (p=0.165). Two sample of blood for serum anti CMV IgG antibody were taken to see the rising titre of IgG.

Table-VI
Distribution of the study subjects by CMV, IgG (n=78). Sample – Blood

Anti-CMV, IgG by MEIA	Group		p value
	Case (Mean ± SD)	Control (Mean ± SD)	
1 st Sample	141.03 ± 268.80	61.99 ± 40.51	0.073
2 nd Sample	195.69 ± 266.12	61.22 ± 36.15	0.003

*t test was done to measure the level of significance.

Figure within parentheses indicates in percentage

Table VI shows the distribution of the study subjects by CMV IgG. In case group the mean value of Anti-CMV, IgG by MEIA in 1st sample was 141.03 ± 268.80 and in control group was 61.99 ± 40.51. The difference between case and control was not statistically significant (p=0.073). In case group the mean value of Anti-CMV, IgG by MEIA in 2nd sample was 195.69 ± 266.12 and in control group was 61.22 ± 36.15. The difference between case and control was statistically significant (p=0.003).

Table-VII

Distribution of the study subjects by history of respiratory tract infection (n=78) preceding illness (1-4 weeks before)

History of Respiratory Tract Infection	Group		p value
	Case (n=39)	Control (n=39)	
Present	18 (46.2)	0 (0.0)	<0.001
Absent	21(53.8)	39(100.0)	
Total	39(100.0)	39(100.0)	

Chi square test (after Yates correction) was done to measure the level of significance.
Odd ratio (95%CI) = 2.86 (2.02-4.03)

Patients with history of respiratory tract infection had 2.86 times more chance to develop GBS than that of control.

Table VII shows the distribution of the study subjects by history of respiratory tract infection. In case group respiratory tract infection was present in 18 (46.2%) cases; 4 (10.2%) cases were CMV positive and absent in 21(53.8%) cases respectively. In control group respiratory tract infection were absent in all 39(100.0%) controls. The difference between case and control was statistically significant (p=<0.001).

Table-VIII

Distribution of the study subjects by history of gastroenteritis (n=78) preceding illness (1-4 weeks before)

History of Gastro-enteritis	Group		p value
	Case (n=39)	Control (n=39)	
Present	11(28.2)	0 (0.0)	<0.001
Absent	28(71.8)	39(100.0)	
Total	39(100.0)	39(100.0)	

Chi square test (after Yates correction) was done to measure the level of significance.
Odd ratio (95%CI) = 2.39 (1.80 - 3.17)

Patients with history of gastroenteritis had 2.39 times more chance to develop GBS than that of control.

Table VIII shows the distribution of the study subjects by history of gastroenteritis. In case group gastroenteritis was present in 11(28.2%) cases; 2(5.1%) cases were CMV positive and absent in 28(71.8%) cases respectively. In control group

gastroenteritis was absent in all 39(100.0%) controls. The difference between case and control was statistically significant (p=<0.001).

Table-IX

Distribution of the study subjects by past history of fever (n=78) preceding illness (1-4 weeks before)

History of Fever	Group		p value
	Case (n=39)	Control (n=39)	
Present	12(30.8)	0(0.0)	<0.001
Absent	27(69.2)	39(100.0)	
Total	39(100.0)	39(100.0)	

Chi square test (after Yates correction) was done to measure the level of significance.
Odd ratio (95%CI) = 2.44 (1.83 - 3.27)

Patients with history of fever had 2.44 times more chance to develop GBS than that of control.

Table IX shows the distribution of the study subjects by history of fever. In case group fever was present in 12(30.8%) cases and absent in 27(69.2%) cases 01(2.5%) cases was CMV positive. In control group fever was absent in all 39(100.0%) controls. The difference between case and control was statistically significant (p=<0.001).

Table-X

Distribution of the study subject by age (n=78)

Age in years	Group		p value
	Case (n=39)	Control (n=39)	
<20	8 (20.5)	8 (20.5)	
20 - 29	13 (33.3)	13 (33.3)	
30 - 39	5 (12.8)	5 (12.8)	
40 - 49	10 (25.6)	10 (25.6)	
50 - 59	3 (7.7)	3 (7.7)	
Total	39(100.0)	39(100.0)	
Mean±SD (Max-min)	30.82±12.56 (58-12)	31.00±12.77 (59-12)	0.950*

*t test was done to measure the level of significance.
Figure within parentheses indicates in percentage

Table X shows the distribution of the study subjects by age. In both case and control group all were equal in number in the age group of 20 – 29 years, 40 – 49 years, less than 20 years, 30 – 39 years and 50 – 59 years which were 13 (33.3%) cases,

10 (25.6%) cases, 8 (20.5%) cases, 5 (12.8%) and 3(7.7%) cases respectively. The mean \pm SD of case and control groups were 30.82 ± 12.56 and 31.00 ± 12.77 respectively. It was not statistically significant ($p=0.950$).

Table-XI

Distribution of the study subjects by sex (n=78)

Sex	Group	p value
Fever	Case (n=39) Control (n=39)	
Male	25(64.1) 25(64.1)	1.000
Female	14(35.9) 14(35.9)	
Total	39(100.0) 39(100.0)	

*Chi-square test was done to measure the level of significance.

Figure within parentheses indicates in percentage.

Table XI shows the distribution of the study subject by sex. In both case and control groups male and female were equal in number which were 25(64.1%) and 14(35.9%) respectively. The difference between case and controls was not statistically significant ($p=1.000$).

Table-XII

Distribution of the cases by CSF study (n=39)

CSF Study	Mean \pm SD	Min- Max
CSF cell	1.95 ± 2.28	0.00 - 10.00
CSF sugar	3.76 ± 1.35	0.00 - 7.30
CSF protein	78.11 ± 94.84	0.49 - 390.00

Table XII shows the distribution of the cases by CSF study. The Mean \pm SD of CSF cell, CSF sugar and CSF protein were 1.95 ± 2.28 mg/dl, 3.76 ± 1.35 cells/cmm and 78.11 ± 94.84 mmol/L respectively.

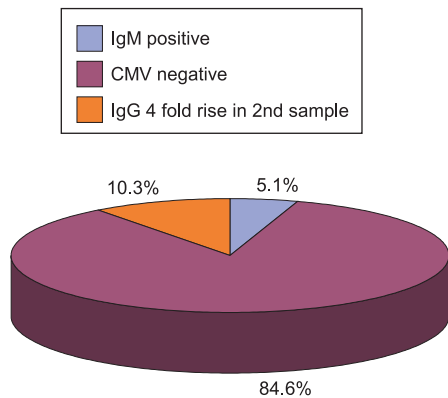


Fig.-1: Distribution of cases by CMV (n=39)

Figure 1: Shows the distribution of cases by CMV. CMV positive cases by IgM positivity (10.3%), 2 (5.1%) by IgG 4 fold rise in the 2nd sample positive and rest 33 (84.6%) cases were CMV Negative.

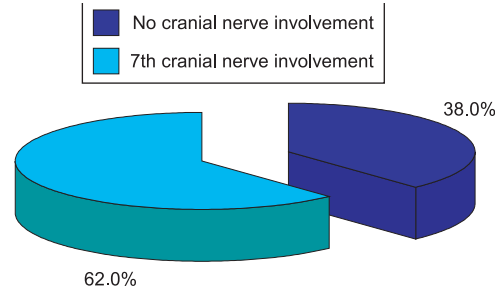


Fig.-2: Distribution of cases by cranial nerves involvement (n=39).

Figure 2: Shows the distribution of cases by cranial nerve involvement. In 24 (62.0%) cases 7th cranial nerve was involved and in 15(38.0%) cases no cranial nerve was involved.

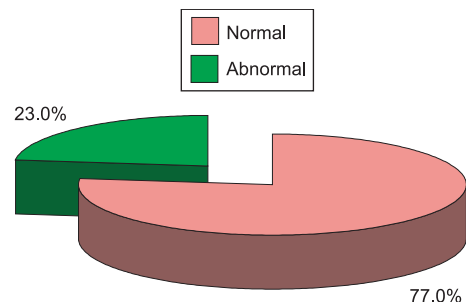


Fig.-3: Distribution of cases by ECG finding (n=39)

Fig. 3: depicts that near one-quarter 9 cases (23%) had abnormal ECG (6 had sinus tachycardia, 3 had sinus bradycardia) and the rest 30 cases (77%) had normal ECG.

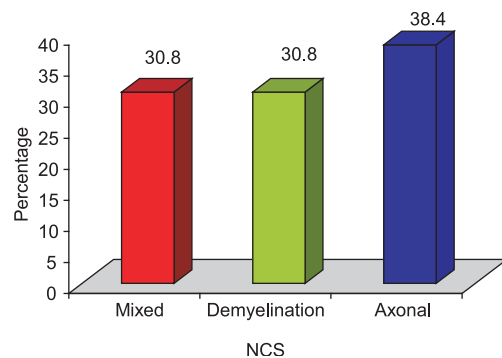


Fig.-4: Distribution of cases by NCS findings (n=39)

Fig. 4: display the distribution of cases by NCS findings. 12 cases (30.8%) shows demyelination and 15 cases (38.4%) shows axonal and rest 12(30.8%) cases shows mixed demyelinating and axonal type of NCS.

Discussion:

Guillain-Barre' Syndrome (GBS) is the most common cause of acute flaccid paralysis¹². Annual incidence of GBS is approximately 1-3 cases per 100,000 persons in Europe, US and Australia¹³.

GBS is an autoimmune disorder of the peripheral nervous system (PNS) with a range of presentations from mild to life-threatening paralysis¹⁴. The etiology of GBS is unknown, yet several studies link some common exposures as precipitating factors, many of which are commonly seen in the primary care setting. Vaccinations, viral infections, and certain type of food poisoning are examples of antecedent factors of GBS¹⁴. Due to severity of complications, practitioners need to be aware of what can trigger GBS, who is at risk, how to recognize early signs/symptoms, what possible prevention exists, and how to educate patients¹⁴.

Gullian Barre syndrome may occur by primary infection or by reinfection or reactivation of CMV¹¹. The association of cytomegalo virus infection with Guillian-Barre syndrome needs diagnosis of CMV infection which required at least one of the following laboratory method like serology, specific intrathecal antibody production, virus isolation, direct detection of CMV PP65 antigen in blood, CMV culture, Biopsy, positive specific immunohistochemical staining, Polymerase chain Reaction (PCR) assay etc⁸.

A total number of 39 patients with GBS as cases and 39 volunteers, patient's attendants were taken as control who were attended in neurology department at BSMMU during the study period, fulfilling the inclusion and exclusion criteria were included in this study.

The distribution of the study subject by IgGs (Fig.-1) more than 4 fold rise in the 2nd sample was recorded in this study. In the case group 4 (10.3%) cases showed 4 fold rise of IgG and the rest 35

(89.7%) cases showed negative results. In control group 4 fold rise of IgG was absent in all controls which was 39 (100.0%).

The distribution of the study subjects by Anti-CMV IgM was recorded in this study. Anti-CMV IgM antibody was positive in 2 (5.1%) cases and negative in 37 (94.9%) cases. Anti CMV IgM antibody was negative in all 39(100.0%) controls. The difference between case and control was not statistically significant ($p=0.474$). Similar result was reported by Kimoto K et al. (2006)¹⁵ and added that CMV infections had a role in the development of GBS. Andary (2011)¹⁶ reported that CMV could occur after upper respiratory and flu like illness and it was the most common viral trigger of GBS with the presence of Anti-CMV IgM which was consistent with the present study. Jacobs et al., (1996)¹⁷ found that in CMV-associated GBS, antibodies were common following CMV infection.

The distribution of the patients by CMV positive was recorded in this study. In case group CMV was overall positive in 6 cases of which IgM was positive in 2 cases and 4 fold rise of IgG was positive in 4 cases, rest 33(84.6%) cases were CMV negative. In control group, CMV was negative in all 39(100%) cases. That indicated the relationship between CMV and GBS, which was significantly associated. The difference between case and control group was statistically significant ($p=0.034$). Similar result was reported by Andary MT (2011)¹⁶ and added that Cytomegalo virus (CMV) infections were the second most commonly reported infections preceding GBS.

The distribution of the study subjects by Anti-CMV IgG Sample-1 was recorded. In case group Anti-CMV IgG (Sample-1) was positive in 33(84.6%) cases and negative in 6(15.4%) cases. In control group Anti-CMV IgG (Sample-1) was positive in 37(94.9%) controls and negative in 2(5.1%) controls. The difference between case and control was not statistically significant ($p=0.135$). The distribution of the study subjects by Anti-CMV IgG Sample-2 was recorded in this study. In case group Anti-CMV IgG (Sample-2) was positive in 35(89.7%) cases and negative in 4(10.3%) cases. In control group Anti-CMV IgG (Sample-2) was positive in 38(97.4%) controls and negative in 1(2.6%) control. The

difference between case and control group was not statistically significant ($p=0.165$). The distribution of the study subjects by CMV IgG was recorded in this study. In case group the mean value of Anti-CMV IgG by MEIA in 1st sample was 141.03 ± 268.80 and in control group was 61.99 ± 40.51 . The difference between case and control group was not statistically significant ($p=0.073$). In case group the mean value of Anti-CMV, IgG by MEIA in 2nd sample was 195.69 ± 266.12 and in control group was 61.22 ± 36.15 . The difference between case and control group was statistically significant ($p=0.003$).

The distribution of the study subjects by history of respiratory tract infection was recorded in this study. In case group respiratory tract infection was present in 18(46.2%) cases; 4(10.2%) cases were CMV positive and absent in 21(53.8%) cases. In control group respiratory tract infection were absent in all 100.0% cases. The difference between case and control group was statistically significant ($p<0.001$). The distribution of the study subjects by history of gastroenteritis was recorded in this study. In case group the Gastroenteritis was present in 11(28.2%) cases; 2(5.1%) cases was CMV positive and negative in 28(71.8%) cases. In control group gastroenteritis was absent in all 39(100.0%) controls. The difference between case and control group was statistically significant ($p<0.001$). The distribution of the study subjects by past history of fever was recorded in this study. In case group fever was present in 12(30.8%) cases; 1(2.5%) case was CMV positive and negative in 27(69.2%) cases. In control group fever was absent in all 39(100.0%) controls. The difference between case and control group was statistically significant ($p<0.001$). Similar result was reported by Andary MT (2011)¹⁶ and mentioned that GBS was considered to be a postinfectious, immune-mediated disease targeting peripheral nerves. Baravelli M et al., (2009)¹⁸ added that up to two thirds of patients report an antecedent bacterial or viral illness prior to the onset of neurologic symptoms. Similarly Nelson L et al., (2009)¹⁹ also reported that respiratory tract infections were most frequently reported, followed by gastrointestinal infections which was consistent with the present study. In another similar study it was mentioned that other systemic illnesses which

was manifested by fever have also been associated with GBS¹⁶.

The distribution of the study subjects by age was recorded in this study. In both case and control groups all were equal in number in the age group of 20–29 years, 40–49 years, less than 20 years, 30 – 39 years and 50 – 59 years which were 13 (33.3%) cases, 10 (25.6%) cases, 8 (20.5%) cases, 5 (12.8%) and 3(7.7%) cases respectively. The mean \pm SD of case and control groups were 30.82 ± 12.56 and 31.00 ± 12.77 respectively which was not statistically significant ($p=0.950$). Similar result was reported by Jiang GX²⁰ and mentioned that GBS had been detected in all age groups, with the syndrome occurring at any time between infancy and old age. In the United States, the syndrome's age distribution seemed to be bimodal, with a first peak in young adulthood (age 15-35 years) and a second, higher one in elderly persons (age 50-59 years). Infants appeared to have the lowest risk of developing GBS²⁰.

The distribution of the study subjects by sex was recorded in this study. In both case and control groups male and female was equal in number which were 25(64.1%) and 14(35.9%) respectively which was not statistically significant ($p=1.000$). Similar result was reported by Andary MT (2011)¹⁶ and mentioned that GBS had a male-to-female ratio of 1.5:1; male preponderance was seen especially in older patients. However, a Swedish epidemiologic study reported that GBS rates decrease during pregnancy and increase in the months immediately following delivery²¹.

The distribution of the cases by ECG (Fig.-3) was recorded in this study. Near the one-quarter cases 9 (23%) has abnormal ECG (6 had sinus tachycardia, 3 had sinus bradycardia) and the rest 30 (77%) cases had normal ECG. Distribution of cases by cranial nerve involvement which shows (Fig.-2). 15 (38%) had no cranial nerve involvement and 24 (62.0%) had 7th cranial nerve involvement. Distribution of cases by NCS (Fig.-4) in this study shows 12 (30.8%) were demyelinating, 15(38.4%) were axonal and rest 12 (30.8%) cases were mixed type. The distribution of cases by CSF study shows the mean \pm SD of CSF cell, CSF sugar and CSF

protein were 1.95 ± 2.28 cells/cmm, 3.76 ± 1.35 mg/dl and 78.11 ± 94.84 respectively.

However, although numerous studies had led to an accurate description of the GBS related to C. jejuni (Cj-GBS), the GBS associated with primary CMV infection (CMV-GBS) remained poorly documented²². Current data were available from just a few studies, most of which had included only a small number of CMV-GBS cases¹¹. In these studies, recent primary CMV infection was clearly defined and the presence or absence of CMV DNA in the blood was not documented²². Because of the small number of patients studied, epidemiological characteristics and specific prognostic features were not specified, and the risk of developing GBS following primary CMV infection was not determined. Visser et al., (1996)⁷ in another study had mentioned that cytomegalovirus (CMV) infection accounts for the most common viral triggers of GBS.

This present study strongly showed that Guillain-Barré syndrome (GBS) had great association with cytomegalovirus (CMV) infection.

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