A study on bacterial infections and their antibiotic susceptibility pattern in decompensated liver disease patients in a tertiary care Hospital

Rajeswari Jayakumar
Assistant Professor, Dept. of Microbiology, Chengalpattu Medical College, Chengalpattu, Tamil Nadu

*Corresponding Author:*
Email: rajeswari1962@yahoo.co.in

**Abstract**

**Background & Objectives:** Decompensated liver disease (DCLD) is defined as irreversible chronic injury of the hepatic parenchyma. Bacterial infections are more common in decompensated liver disease and causes 30%-50% of deaths. Therefore, this study was done to determine the various bacterial agents causing infections in decompensated liver disease patients and to determine the drug susceptibility and resistance pattern.

**Methods:** A prospective study was conducted over a period of one year in a tertiary care hospital. Ascitic fluid, urine, sputum, blood and wound swab were collected. All the samples were processed through Gram’s stain and culture. The organisms were identified by standard protocols and antibiotic susceptibility testing.

**Results & Conclusion:** Out of 150 samples, culture positivity seen in 81(54%). In 81 culture positive isolates, 63(78%) were Gram Negative bacilli (GNB) and 18 (22%) were Gram Positive cocci (GPC). Among Gram negative bacilli, *Escherichia coli* and in Gram positive cocci, *Staphylococcus aureus* was the most common isolates. The most common infections were spontaneous bacterial peritonitis (27%) followed by urinary tract infections (26%), Spontaneous bacteraemia (19%), pneumonia (16%), and skin and soft-tissue infections (12%).

**Keywords:** Decompensated Liver Disease, Spontaneous Bacterial Peritonitis, *Escherichia coli*, *Staphylococcus aureus*.

**Introduction**

Liver failure leading to cirrhosis is one of the most common causes of death in our country.¹ ¹ Cirrhosis is a chronic progressive liver disorder caused by alcoholic liver diseases, viral hepatitis (HBV and HCV) and cryptogenic causes which can lead to liver failure and death.⁴ According to the stages of liver injury, signs and symptoms and survival rate, cirrhosis is classified into compensated and decompensated liver disease.⁵

 Decompensated liver disease (DCLD) is defined as irreversible chronic injury of the hepatic parenchyma and extensive fibrosis in association with the formation of regenerative nodules and leading to loss of liver function.⁶

Bacterial infections are more common in Patients with decompensated liver disease is due to altered and impaired immunity and causes 30%-50% of deaths.⁷ In decompensated liver disease patients, the spontaneous bacterial peritonitis (SBP) is a serious common bacterial infection, followed by urinary tract infections (UTI), spontaneous bacteraemia, pneumonia, and skin infections.⁸ The common causative organisms for bacterial infections in DCLD patients are *Enterobacteriaceae*, nonfermentable Gram-negative bacilli and Gram positive cocci and most of them are multidrug resistant.⁹ ¹⁰ The prognosis of these patients is closely related to a prompt and accurate diagnosis and appropriate treatment decreases the mortality rates.

**Materials & Methods**

A prospective study was conducted over a period of one year. About 150 patients (≥18 yrs.), admitted in various wards with signs & symptoms suggestive of bacterial infections in DCLD patients are included in the study. Ascitic fluid, urine, sputum, blood and wound swab were collected. All the samples were processed through Gram’s stain and inoculated onto Blood agar plate, Chocolate agar and MacConkey agar. The inoculated culture plates were incubated overnight at 37°C in an incubator. A Gram’s stain was done the next day from the growth and examined. The organisms were identified by standard protocols and antibiotic susceptibility of recommended drugs (CLSI guidelines) was performed by using Kirby Bauer disc diffusion and Minimum inhibitory concentration (MIC) method.

**Results**

Out of 150 samples, culture positivity seen in 81(54%). In 81 culture positive isolates, 63(78%) were Gram Negative bacilli and 18 (22%) were Gram Positive cocci. Among Gram negative bacilli, *Escherichia coli* was the most common isolates and in Gram positive cocci, *Staphylococcus aureus* was the most common isolates (Table 1). The most common infections were spontaneous bacterial peritonitis (27%) followed by urinary tract infections (26%), Spontaneous bacteraemia (19%), pneumonia (16%), and skin and soft-tissue infections (12%). (Chart 1)
A study on bacterial infections and their antibiotic susceptibility pattern in DCLD patients

Fig. 1: Distribution of culture positivity in various types of clinical samples in DCLD patients

Table 1: Bacteria isolated from the various samples in DCLD patients

<table>
<thead>
<tr>
<th>Organism</th>
<th>Ascitic fluid</th>
<th>Urine</th>
<th>Sputum</th>
<th>Skin</th>
<th>Blood</th>
<th>No. of isolates</th>
<th>% of isolates</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Escherichia coli</em></td>
<td>7</td>
<td>8</td>
<td>1</td>
<td>1</td>
<td>2</td>
<td>19</td>
<td>23.5</td>
</tr>
<tr>
<td><em>Klebsiella pneumoniae</em></td>
<td>3</td>
<td>3</td>
<td>3</td>
<td>1</td>
<td>4</td>
<td>14</td>
<td>17</td>
</tr>
<tr>
<td><em>Klebsiella oxytoca</em></td>
<td>1</td>
<td>6</td>
<td>6</td>
<td>-</td>
<td>-</td>
<td>13</td>
<td>16</td>
</tr>
<tr>
<td><em>Proteus mirabilis</em></td>
<td>1</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>1</td>
<td>1.2</td>
</tr>
<tr>
<td><em>Proteus vulgaris</em></td>
<td>-</td>
<td>-</td>
<td>3</td>
<td>-</td>
<td>-</td>
<td>3</td>
<td>3.7</td>
</tr>
<tr>
<td><em>Enterobacter cloacae</em></td>
<td>1</td>
<td>1</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>2</td>
<td>2.5</td>
</tr>
<tr>
<td><em>Citrobacter koseri</em></td>
<td>1</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>1</td>
<td>1.2</td>
</tr>
<tr>
<td><em>Pseudomonas aeruginosa</em></td>
<td>1</td>
<td>-</td>
<td>3</td>
<td>1</td>
<td>2</td>
<td>7</td>
<td>9</td>
</tr>
<tr>
<td><em>Acinetobacter baumannii</em></td>
<td>1</td>
<td>2</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>3</td>
<td>3.7</td>
</tr>
<tr>
<td><em>Staphylococcus aureus</em></td>
<td>4</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>2</td>
<td>3</td>
<td>9</td>
</tr>
<tr>
<td><em>Enterococcus faecalis</em></td>
<td>2</td>
<td>1</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>3</td>
<td>3.7</td>
</tr>
<tr>
<td><em>Staphylococcus epidermidis</em></td>
<td>-</td>
<td>-</td>
<td>2</td>
<td>2</td>
<td>2</td>
<td>4</td>
<td>5</td>
</tr>
<tr>
<td><em>Streptococcus viridans</em></td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>2</td>
<td>-</td>
<td>2</td>
<td>2.5</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>22</strong></td>
<td><strong>21</strong></td>
<td><strong>13</strong></td>
<td><strong>10</strong></td>
<td><strong>15</strong></td>
<td><strong>81</strong></td>
<td><strong>100</strong></td>
</tr>
</tbody>
</table>

Most of the organisms were 75% sensitive to amino glycosides and 50% sensitive to fluoroquinolones. All the GNB were 100% sensitive to carbapenem except one carbapenem resistant isolates, *Klebsiella oxytoca* was isolated from sputum sample.

In 81 culture-positive infections, 33(41%) drug resistant bacterial infections were identified: 27 (81%) Extended spectrum Beta Lactamases (ESBL), 4(10%) Methicillin resistant Staphylococcus aureus (MRSA), 1(2.3%) Vancomycin resistant *Enterococci* (VRE) and 1(2.3%) Metallo Beta lactamases (MBL). Of the culture-positive infections, these drug resistant bacterial infections occurred in 11 of 21 (52%) of the UTIs, 8 of 22 (36%) of the SBP, 3 of 15 (20%) of the spontaneous bacteraemia cases, 7 of 13 (54%) of the pneumonia and 4 of 10 (40%) of the skin and soft tissue infection cases.

Fig. 2: Double disk diffusion synergy test for detection of ESBL

Fig. 3: Modified Hodge Test (MHT)
Isolate 1-Meropenem sensitive – MHT Negative
Isolate 2- Meropenem resistant – MHT Positive

Table 2: Minimum inhibitory concentration (MIC) for detecting drug resistance by macro broth dilution method

<table>
<thead>
<tr>
<th>Drugs</th>
<th>Organisms</th>
<th>Observed MIC value</th>
<th>Interpretation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Vancomycin</td>
<td>Staphylococcus aureus (MRSA)</td>
<td>≤2μg/l</td>
<td>Sensitive (≤ 2μg/ml – Susceptible</td>
</tr>
<tr>
<td></td>
<td>(Isolates no=4)</td>
<td></td>
<td>4–8μg/ml – Intermediate</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>≥16μg/ml – Resistant.)</td>
</tr>
<tr>
<td>Vancomycin</td>
<td>Enterococcus faecalis (Isolate no=1)</td>
<td>64μg/ml</td>
<td>Resistant (≤ 4μg/ml – Susceptible</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>8–16μg/ml – Intermediate</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>≥32μg/ml – Resistant)</td>
</tr>
<tr>
<td>Meropenem</td>
<td>Klebsiella oxytoca (Isolate no=1)</td>
<td>≥16μg/ml</td>
<td>Resistant (≤ 1μg/ml – Susceptible</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>2μg/ml – Intermediate</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>≥4μg/ml – Resistant)</td>
</tr>
</tbody>
</table>

(Interpretation according to CLSI guidelines)

Discussion

In 150 DCLD patients, males 145 (97%) were predominant group when compared to females 5(3%). This predilection of higher frequency rates among male is attributed towards the presence of underlying risk factors like alcoholism.(11,12)

Out of 150 patients of DCLD, 81(54%) were culture positive. In 81 culture positive isolates, 63(78%) were Gram Negative bacilli and 18 (22%) were Gram Positive cocci, which was correlated significantly (P value = 0.005). In decompensated liver disease (DCLD) patients, the most common isolates were Gram negative bacilli which may be due to translocation of normal flora (most of the normal flora in the GIT are GNB) from the gastrointestinal tract. Among bacterial infections, Escherichia coli were the most common pathogen (24%).(13)

Among culture positive infections, spontaneous bacterial peritonitis (27%) was the most common infection due to translocation of enteric organisms from the intestine to the peritoneum and diagnostic and therapeutic paracentesis were predisposed to bacterial infections, followed by urinary tract infection (26%), spontaneous bacteraemia (19%), pneumonia (16%) and skin and soft tissues infection (12%).(11,14,15)

In patients with spontaneous bacterial peritonitis (SBP) (27%) the most frequently isolated organisms was Escherichia coli (31.82%) as it is the commonest enteric pathogen, followed by Staphylococcus aureus (18.18%), Klebsiella pneumonia (13.36%) and Enterococcus faecalis (9.09%).(19,20)

In this study, among 44 urine samples, 21 (26%) were culture positive. Of which Escherichia coli (38.09%) were the most common isolates followed by Klebsiella oxytoca (28.57%), Klebsiella pneumonia (14.28%), Acinetobacter baumanii (9.52%) and Enterococcus faecalis (4.76%). The incidence of urinary tract infection (UTI) is higher in decompensated cirrhotic patients with indwelling urinary catheters.(16)

Among 150 blood samples, 15 (19%) samples were culture positive. Of which Klebsiella pneumoniae (26.33%) was the most common isolate followed by Staphylococcus aureus (20.00%), Escherichia coli, Pseudomonas aeruginosa, Staphylococcus epidermidis and Streptococcus viridians. The Porto systemic shunt
circulation in DCLD patients will favour the organisms to escape from phagocytosis by hepatic reticuloendothelial system, there by establishing systemic bacteremia.\(^{(17)}\)

In the sputum samples, 13(16%) samples were culture positive. Klebsiella oxytoca (46.14%) was the most common isolates followed by Klebsiella pneumoniae (23.07%), Pseudomonas aeruginosa (23.07%) and Escherichia coli (7.69%). Some procedures like tracheal intubation, oesophageal tamponade and clinical conditions like hepatic encephalopathy, alcoholism were clearly predisposing factors for pneumonia in cirrhotic patients.

Out of 18 wound swabs, 10(12%) were culture positive. Proteus vulgaris (30%) was the main isolate followed by Staphylococcus aureus (20%), Staphylococcus epidemidis (20%), Escherichia coli and Klebsiella pneumoniae. Lymphangitis of the lower extremities and abdominal wall are frequent in cirrhotic patients with oedema or ascites which will leads to skin and soft tissue infections in DCLD patients.\(^{(17,18)}\)

Most of the organisms were 75% sensitive to amino glycosides and 50% sensitive to fluoroquinolones. All the GNB were 100% sensitive to carbapenem except one carbapenem resistant isolates, Klebsiella oxytoca was isolated from sputum sample.

In 81 culture-positive isolates, 31 were drug resistant bacterial infections were identified: 81% (25of 31) were ESBL, 13% (4/31) were Methicillin resistant Staphylococcus aureus, 3% (1/31) were vancomycin resistant Enterococci (VRE), and 3% (1/31) were MBL (3%). Of the culture-positive isolates, these drug resistant bacterial infections occurred in 11 of 21 (52%) of the urine tracts infections (UTIs), 7 of 22 (32%) of the spontaneous bacterial peritonitis (SBP), 3 of 15 (20%) of the spontaneous bacteraemia cases, 6 of 13 (46%) of the pneumonia and 4 of 10 (40%) of the skin and soft tissue infection cases.

**Conclusion**

The prognosis of DCLD patients is closely related to identify the definitive etiologic diagnosis with its antimicrobial susceptibility and resistant pattern. Antibiotic prophylaxis must be restricted to selected patients and encouraging the use of first line antibiotics and to avoid unnecessary use of higher antibiotics. This will help to lower the occurrence of new resistant strains, which can significantly reduce hospital stay and morbidity and improve survival rate.

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