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The journal has had 7 points in Ministry of Science and Higher Education parametric evaluation. Part B item 1223 (26/01/2017). 1223 Journal of Education, Health and Sport eISSN 2391-8306 7 © The Authors 2019; This article is published with open access at Licensee Open Journal Systems of Kazimierz Wielki University in Bydgoszcz, Poland Open Access. This article is distributed under the terms of the Creative Commons Artribution Noncommercial License Which permits any noncommercial use, distribution and reproduction in any medium, provided the original author (s) and source are credited. This is an open access article licensed under the terms of the Creative Commons Attribution Noncommercial license Share alike. (http://creativecommons.org/licenses/by-nc-sa/4.0) which permits unrestricted, non commercial use, distribution and reproduction in any medium, provided the work is properly cited. The authors declare that there is no conflict of interests regarding the publication of this paper. Received: 05.07.2019. Revised: 20.07.2019. Accepted: 24.07.2019.

## FEATURES OF MICROBIOLOGICAL DIAGNOSIS AND CHOICE OF SELECTIVE ANTIBIOTIC THERAPY IN PATIENTS WITH ACUTE CHOLECYSTITIS

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#### Abstract

**The objective:** to improve the results of destructive forms of acute cholecystitis (AC) surgery by the development of antibiotic therapy algorithm and taking into account the severity of the disease according to Tokyo guidelines (2013). Materials and methods: 266 patients having been operated on for AC made up a treatment group and 203 patients - the control one. In 234 treatment group patients microflora's bacteriological study was carried out and its selective sensitivity to antibiotics was determined. Results and discussion. The AC treatment group patients (n=266) were arranged according to Tokyo guidelines (2013) into 3 grades. Mild course (grade I) consisted of 29 patients (11%), microbiological studies were not conducted. Antibiotic prophylaxis with beta-lactam antibiotics was used. At moderately severe AC (grade II, n = 200, 75%) escalation antibiotic therapy was carried out. At severe disease (grade III, n = 37, 14%) de-escalation antibiotic therapy was performed. Bacteriological studies were conducted in 237 patients with destructive forms of AC, at that 531 microflora isolates were cultured. Intestinal group microorganisms were the most frequently cultured. Beta-lactam antibiotics' sensitivity was prevailing and constituted 83.3%. The number of suppurations in the treatment group decreased from 9.8% to 4.3%, and mortality rate decreased from 1.48% to 0.75% (p<0.05) as contracted with control group where empirical antibiotic therapy was used. Conclusions. 1. Intestinal group microflora prevailed in destructive forms of AC. 2. In mild course of cholecystitis antibiotic prophylaxis is consider to be method of choice, in moderately severe disease – escalating therapy, and in severe cases – de-escalating antibiotic therapy should be used, as that beta-lactam antibiotics had the highest selective sensitivity (83.3%). 3. AC is local intra-abdominal infection, where evidence-based antibiotic therapy has reduced postoperative complications to 4.3%, and mortality rate - to 0.75%.

Key words: acute cholecystitis, prophylactic, escalation, de-escalation selective antibiotic therapy, microbiological diagnostics.

The urgency of the problem. Acute cholecystitis (AC) is an inflammation of the gallbladder (GB) with extra-and intrahepatic bile ducts (2000  $m^2$ ) of the liver frequent involvement in the pathological process, pancreas with the threat of the development of biliary peritonitis, perivascular ulcer and abscess, cholangitis, choledocholithiasis and holodolar and cholangitis. sepsis [4, 5, 11].

The frequency of surgical interventions in AC is close to that in appendectomies, and in the patients over 50 y. o. surpasses it with relatively high mortality rate [7].

AC is a local intra-abdominal infection, where adequate antibiotic therapy significantly affects purulent complications and mortality. A number of works [1, 2, 3, 6, 7, 8, 10, 12, 13] are devoted to the problems of choosing antibiotics in the treatment of AC, differentiation of tactics according to the severity of the course, determination of microflora sensitivity.

**The objective:** to improve the results of destructive AC surgery by the development of antibiotic therapy algorithm and taking into account the severity of the disease according to Tokyo guidelines (2013) and determine the selective sensitivity of the microflora cultured to antibacterial drugs.

**Materials and methods**. 266 AC patients were examined. They have been operated on in the surgical department of Kiev Regional Clinical Hospital in 2016 – 2018. The control group included 203 AC patients operated on in 2013–2015, similar in sex, age, comorbidities, treatment and surgical tactics to each other, but with an empirical choice of antibiotic therapy. On presentation the material into the bacteriological laboratory, smears were made, Gram's stained and micro scoped. After microflora's identification antibiotic was empirically administered.

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To isolate aerobic bacteria pure cultures standardized nutrient media of industrial production namely 5% blood, gall-salt, Endo, enterococcus and chocolate agar were used. All of them are allowed for the use in Ukraine and have corresponding quality certificates.

Identification of isolates and determination of their sensitivity to antibacterial drugs was performed on a VITEK-2 Compact 15 microbiological analyzer (France).

Then the minimum inhibitory concentrations were determined and individually the treatment doses of drugs were calculated.

**Results and discussion.** The AC treatment group patients (n=266) were arranged according to Tokyo guidelines (2013) into 3 grades.

Grade I consisted of 29 patients (11%) with AC mild course. They had moderately promoted GB changes and none severe comorbidities. Microbiological studies were not conducted. Standard antibiotic prophylaxis (*Sulbactamum, Amoksiklav, Cephalosporins*) was used.

Grade II included 200 patients (75%) with moderately severe AC. GB was palpable and there was an infiltrate in the right hypochondrium. In this group before plated microflora sensitivity determination, the escalation antibiotic therapy was used (cepahalosporins of the III<sup>rd</sup> generation)

Grade III (n = 37, 14%) included patients with circumscribed, rarer generalized peritonitis. Patients with multisystem failure symptoms were not enrolled. De-escalation antibiotic therapy (carbapenems) was performed in this group.

All 266 patients were operated on under general endotracheal anesthesia using muscle relaxants. 253 patients (95%) were operated on with laparoscopic techniques with 6% conversions; 13 patients (5%) with laparotomy access.

Bacteriological studies were conducted in 237 (89%) of patients with phlegmonous and gangrenous AC forms and perivesical abscesses. Material for the study was taken from the abdominal cavity exudates, GB content, perivesical abscess. The investigation was conducted in dynamics, up to 3 times.

Both in monoculture and microbial associations 531 microflora isolate were excreted.

Table 1 presents the results of bacteriological studies in patients with destructive forms of AC.

# Table 1

| <b>Bacteriological studies of 237</b> | nationts with destructive | forms of acute cholecystitis |
|---------------------------------------|---------------------------|------------------------------|
| Dacteriological studies of 257        | patients with acouldent   | forms of acute choice ystins |

| Bacteriological studies of 257 patients with destr                  |        |               |
|---|--------|---------------|
| Microorganism   | Number | <u>%</u>      |
| Achromobacterxylosoxidansss. Xylosoxidans<br>Acinetobacterbaumannii | 10     | $\frac{0}{2}$ |
|   | 10     |               |
| Actinomycesnaeslundii   | 1      | 0             |
| Bacillus cereus   | 2      | 0             |
| Bacillus sp.  | 1      | 0             |
| Burkholderiacepacia   | 1      | 0             |
| Candida albicans  | 5      | 1             |
| Candida parapsilosis  | 1      | 0             |
| Candida sp.   | 2      | 0             |
| Chryseomonasluteola   | 1      | 0             |
| Citrobacterfreundii   | 3      | 1             |
| Citrobactersedlakii   | 2      | 0             |
| Citrobacter sp.   | 2      | 0             |
| Enterobacteraerogenes   | 26     | 5             |
| Enterobacter cloacae  | 16     | 3             |
| Enterococcus avium  | 1      | 0             |
| Enterococcus casseliflavus  | 1      | 0             |
| Enterococcus faecalis   | 17     | 3             |
| Enterococcus faecium  | 29     | 5             |
| Enterococcus gallinarum   | 1      | 0             |
| Escherichia coli  | 66     | 12            |
| Escherichia vulneris  | 1      | 0             |
| Gemellahaemolysans  | 1      | 0             |
| Gram positive bacteria  | 1      | 0             |
| Gram positive cocci   | 1      | 0             |
| Klebsiellaoxytoca   | 3      | 1             |
| Klebsiellapneumoniae ss. Pneumoniae                                 | 50     | 9             |
| Klebsiella sp.  | 1      | 0             |
| Kocuriakristinae  | 3      | 1             |
| Lactococcusgarvieae   | 2      | 0             |
| Leuconostocmesenteroides ss. Mesenteroides                          | 1      | 0             |
| Morganellamorganii ss. Morganii                                     | 1      | 0             |
|   | 127    | 24            |
| No growth   | 2      | 0             |
| Normal flora  |        |               |
| Peptostreptococcus sp.  | 1      | 0             |
| Proteus mirabilis   | 5      | 1             |
| Proteus rettgeri  | 4      | 1             |
| Proteus vulgaris  | 5      | 1             |
| Pseudomonas aeruginosa  | 17     | 3             |
| Raoultellaplanticola  | 1      | 0             |
| Serratiafonticola   | 3      | 1             |
| Serratiamarcescens  | 1      | 0             |
| Staphylococcus aureus ss. Aureus                                    | 38     | 7             |
| Staphylococcus epidermidis  | 54     | 10            |
| Staphylococcus haemolyticus   | 2      | 0             |
| Staphylococcus hominis ss. Hominis                                  | 2      | 0             |
| Staphylococcus saprophyticus ss. saprophyticus                      | 1      | 0             |
| Streptococcus pneumoniae  | 2      | 0             |
| Streptococcus pyogenes  | 2      | 0             |
| Streptococcus sp.   | 4      | 1             |
| N = 531   |        |               |

*Escherichia coli* (12%), *Staphylococcus epidermidis* (10%), *Klebsiella pneumonia* (9%), *Staphylococcus aureus* (7%), *Enterobacter aerogenes* and *Enterococcus faecium* (5% each) were cultured more frequently. In 127 AC patients (24%) there was no growth of aerobic microflora.

Table 2 presents the selective sensitivity of the microflora cultured in AC destructive forms patients to antibiotics.

Table 2

| Антибіотик                           | Кількість | %R     | %I    | %S   |
|--------------------------------------|-----------|--------|-------|------|
| 1                                    | 2         | 3      | 4     | 5    |
| Beta-lactamase                       | 18        | 16.7   |       | 83.3 |
| ESBL                                 | 16        | 50.0   |       | 50.0 |
| PEN.ND10 Penicillin G                | 31        | 67.7   | 0.0   | 32.3 |
| PEN_NM Penicillin G                  | 12        | 75.0   | 0.0   | 25.0 |
| AMP_ND10 Ampicillin                  | 84        | 83.3   | 1.2   | 14.3 |
| AMP_NM Ampicillin                    | 33        | 81.8   | 0.0   | 18.2 |
| AMX_NM Amoxicillin                   | 1         | 100.0  | 0.0   | 0.0  |
| AMX_ND25 Amoxicillin                 | 66        | 47.0   | 12.1  | 40.9 |
| AZL_ND75 Azlocillin                  | 35        | 77.1   | 0.0   | 22.9 |
| PIP_ND100 Piperacillin               | 66        | 60.6   | 7.6   | 31.8 |
| PIP_NM Piperacillin                  | 21        | 57.1   | 4.8   | 33.3 |
| TIC_ED75 Ticarcillin                 | 27        | 81.5   | 0.0   | 18.5 |
| OXA_ED1 Oxacillin                    | 3         | 100.0  | 0.0   | 0.0  |
| OXA_NM Oxacillin                     | 12        | 50.0   | 0.0   | 50.0 |
| AMC_ED25 Amoxicillin/Clavulanic acid | 24        | 50.0   | 8.3   | 41.7 |
| CCV_ND30 Ceftazidime/Clavulanic acid | 7         | 28.6   | 0.0   | 71.4 |
| CSL_ND30 Cefoperszcne/Sulbactam      | 63        | 41.3   | 19.0  | 39.7 |
| SAM_ND10 Ampicillin/SulDactam        | 43        | 32.6   | 14.0  | 53.5 |
| SAM_NM Ampicillin/Sulbactam          | 23        | 43.5   | 4.3   | 52.5 |
| TCC_ND75 Ticarcillin/Clavulanic acid | 8         | 50.0   | 0.0   | 50.0 |
| TZP_NDiOOPineracil'in/T azobactam    | 89        | 31.5   | 14.6  | 53.9 |
| TZP_NM Piperacillin, Tazobactam      | 33        | 45.5   | 21.2  | 30.3 |
| CZO_NM Cefazolin                     | 21        | 47.6   | 0.0   | 52.4 |
| CZO_ND30 Cefazolin                   | 88        | 58.0   | 3.4   | 38.6 |
| CFPJMD75 Cefoperazone                | 36        | 47.2   | 11.1  | 41.7 |
| CXM_ND30 Cefuroxime                  | 114       | 66.7   | 3.5   | 29.8 |
| CXM_NM Cefuroxime                    | 15        | 73.3   | 0.0   | 26.7 |
| MAN_ND30 Cefamandole                 | 29        | 44.8   | 6.9   | 48.3 |
| CAZ_ND30 Ceftazidime                 | 133       | 60.2   | 12.0  | 27.8 |
| CRO_ND30 Ceftriaxone                 | 61        | 59.0   | 4.9   | 36.1 |
| CRO_NM Ceftriaxone                   | 32        | 56.2   | 0.0   | 43.8 |
| CTX_ND30 Cefotaxime                  | 71        | 62.0   | 12.7  | 25.4 |
| FEP_NM Cefepime                      | 52        | 57.7   | 5.8   | 36.5 |
| FEP_ND30 Cefepime                    | 111       | 65.8   | 9.0   | 19.8 |
| FOX_ND30 Cefoxitin                   | 98        | 45.9   | 3.1   | 51.0 |
| FOX_NM Cefoxitin                     | 7         | 28.6   | 0.0   | 71.4 |
| CFM_NM Cefixime                      | 12        | 75.0   | 0.0   | 25.0 |
| CFM_ND5 Cefixime                     | 26        | 61.5 ' | 11.5  | 26.9 |
| CPD_ND10 Cefpodoxime                 | 1         | 0.0    | 100.0 | 0.0  |

Microflora selective sensitivity in AC patients

| 1                                    | 2   | 3     | 4    | 5     |
|--------------------------------------|-----|-------|------|-------|
| CXA NM Cefuroxime axetil             | 15  | 73.3  | 0.0  | 26.7  |
| ATM_ND30 Aztreonam                   | 48  | 64.6  | 20.8 | 14.6  |
| ATM_NM Aztreonam                     | 20  | 60.0  | 0.0  | 40.0  |
| DORJMD10 Doripenem                   | 17  | 58.8  | 5.9  | 35.3  |
| ETP_ND10 Ertapenem                   | 2   | 0.0   | 50.0 | 50.0  |
| IPM_ND10 Imipenem                    | 74  | 35.1  | 6.8  | 58.1  |
| IPMJMM Imipenem                      | 21  | 33.3  | 28.6 | 38.1  |
| MEM_NM Meropenem                     | 41  | 31.7  | 17.1 | 51.2  |
| MEM_ND10 Meropenem                   | 96  | 41.7  | 13.5 | 44.8  |
| AMK_ND30 Amikacin                    | 103 | 46.6  | 7.8  | 45,6  |
| AMK_NM Amikacin                      | 29  | 20,7  | 3,4  | 72,4  |
| GEN_NM Gentamicin                    | 48  | 43,8  | 4,2  | 52,1  |
| GEN_ND10 Gentamicin                  | 77  | 39,0  | 5,2  | 55,8  |
| KAN_ND30 Kanamycin                   | 76  | 63,2  | 15,8 | 21,1  |
| NET.ND30 Netilmicin                  | 146 | 21,2  | 2,1  | 76,7  |
| TOB_ND10 Tobramycin                  | 190 | 31,1  | 4,7  | 64,2  |
| TOB_NM Tobramycin                    | 21  | 28,6  | 9,5  | 57,1  |
| RIFJMD5 Rifampin                     | 12  | 16,7  | 8,3  | 75,0  |
| CIP_ED10 Ciprofloxacin               | 58  | 50,0  | 1,7  | 48,3  |
| CIP_NM Ciprofloxacin                 | 46  | 65,2  | 0,0  | 34,8  |
| GAT_ND5 Gatifloxacin                 | 64  | 43,8  | 3,1  | 51,6  |
| LOM_ND10 Lomefloxacin                | 1   | 100,0 | 0,0  | 0,0   |
| LVX_ND5 Levofloxacin                 | 103 | 54,4  | 2,9  | 42,7  |
| LVX_NM Levofloxacin                  | 51  | 56,9  | 0,0  | 43,1  |
| MFX_NM Moxifloxacin                  | 35  | 57,1  | 2,9  | 40,0  |
| MFX_\'D5 Moxifloxacin                | 66  | 51,5  | 6,1  | 42,4  |
| NOR_ND10 Norfloxacin                 | 71  | 64,8  | 4,2  | 31,0  |
| NOR_MM Norfloxacin                   | 11  | 90,9  | 0,0  | 9,1   |
| OFX_ND5 Ofloxacin                    | 123 | 47,2  | 6,5  | 46,3  |
| PEF_ED5 Pefloxacin                   | 2   | 100,0 | 0,0  | 0,0   |
| SXT_NM Trimethoprim/Sulfamethoxazole | 46  | 37,0  | 0,0  | 21,7  |
| TMP_ND5 Trimethoprim                 | 36  | 66,7  | 0,0  | J3,3  |
| TMP_NM Trimethoprim                  | 14  | 50,0  | 0,0  | 50,0  |
| FO3_N320 Fosfomycin                  | 14  | 50,0  | 7,1  | 42,9  |
| FOS_ED20 Fosfomycin                  | 5   | 80,0  | 0,0  | 20,0  |
| FOS_NM Fosfomycin                    | 11  | 45,5  | 0,0  | 45,5  |
| CLI_NM Clindamycin                   | 21  | 61,9  | 0,0  | 38,1  |
| CLI_ND2 Clindamycin                  | 96  | 44,8  | 8,3  | 46,9  |
| LIN_ED15 Lincomycin                  | 6   | 50,0  | 33,3 | 16,7  |
| COL_ND10 Colistin                    | 52  | 5,8   | 1,9  | 92,3  |
| COL_NM Colistin                      | 18  | 11,1  | 5,6  | 83,3  |
| AZM_ND15 Azithromycin                | 68  | 70,6  | 7,4  | 22,1  |
| ERY_NM Erythromycin                  | 21  | 66,7  | 0,0  | 33,3  |
| ERY_ND15 Erythromycin                | 68  | 48,5  | 26,5 | 25,0  |
| LNZ_NM Linezolid                     | 15  | 0,0   | 0,0  | 100,0 |
|                                      | 15  | 0,0   | 0,0  | 100,0 |

The highest sensitivity of microflora isolated in destructive forms of AC patients (83.3%) is marked for beta-lactam antibiotics - penicillins, cephalosporins, monobactams, and carbapenems.

In the group of patients with AC mild forms antibiotic prophylaxis was used. Escalation antibacterial therapy was used in 200 mean heavy variant of AC patients (75%) and then a follow-up transition to selective antibiotics according to the results of microbiological sensitivity. In 37 AC severe form patients (14%) de-escalation antibiotic therapy (carbapenems) was used immediately

Before the results of selective sensitivity were obtained imipenem 0.5 gr i.v. 4 times per day or meropenem 0.5 gr i.v. 4 times per day or ertapenem 1 gr once per day were considered to be therapy of choice for the patients with severe course of AC.

Alternative therapy included: cefoperazon / sulbactam 2-4 gr i.v. twice per day or cefoperazon 2-4 gr i.v. twice-thrice per day or cefotaxim 1-2 gr i.m. thrice per day or levofloxacin 0.5 gr i. v. once –twice per day or ciprofloxacin 0.5 gr i.v. twice per day with metronidazole 0.5 gr i.v. thrice per day.

If microflora cultured was antibiotics-resistant (lack of growth) their change was carried out according to clinical signs.

The duration of antibiotic therapy was determined clinically. Decrease in body temperature below 37.5° C, normalization of leukocytes number in peripheral blood and the number of immature forms in the leukocyte formula, good condition of the postoperative wound was taking into account.

The duration of antibiotic therapy in AC destructive forms patients averaged  $13.4 \pm 0.5$  (minimum 6, maximum 24 days of continuous therapy).

Compared with the control group of 203 AC patients, 266 treatment group patients had decreased number of suppurations (from 9.8% to 4.3%; p<0.01), and decreased mortality rate (from 1.48% to 0.75%; p<0.05).

### **Conclusions:**

1. Severity of illness was determined according to Tokyo Agreement (Tokyo guidelines, 2013). For the patients of the first grade (light course), we recommend the use of antibiotic prophylaxis. Under moderate course escalation antibiotic therapy and under severe course of the AC de-escalation antibiotic therapy, followed by the use of selective antibacterial agents should be administered.

2. The highest selective sensitivity of microflora cultured in the patients operated on for AC destructive forms was revealed to beta-lactam antibiotics (83.3%).

3. The use of differentiated antibiotic regimens based on the severity of the disease according to Tokyo guidelines (2013) and the identification of the selective sensitivity of microflora with the optimal therapeutic and surgical tactics were able to reduce the number of purulent complications in the control group from 9.8% to 4.3% (p<0.01) and mortality rate from 1.48% to 0.75% (p<0.05).

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