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1 What antibiotics for what pathogens? The sensitivity spectrum of isolated strains in an 2 intensive care unit

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26 Abstract

Antibiotic sensitivity spectrum of isolated strains differs according to hospital departments, the 27 hospitals themselves, and countries. Discrepancies also exist in terms of antibiotic use and 28 dosage. The aim of the present study is to compare the antibacterial agents, the types of 29 infections, the number and type of pathogens, and the sensitivity to antibiotics used in the 30 Intensive Care Unit (ICU) of the Emergency Clinical County Hospital of Oradea, Romania. Over 31 32 a one-year period, data were gathered from the pharmacy computer system and medical records 33 of inpatients. WHO Anatomical Therapeutic Chemical (ATC) / defined daily doses (DDD) methodology was used to assess drug administration data, and antibiotic use was expressed as 34 35 DDD/1000 PD (patient days). The antibiotic susceptibility of isolated strains was expressed through the cumulative antibiogram. The overall consumption of antimicrobial agents was 36 1247.47 DDD/1000 PD. The most common drugs used were cephalosporins and 37 38 fluoroquinolones (52.97% of the total). Ceftriaxone was the most commonly used, followed by levofloxacin. Infections of the respiratory and urinary tract were the most frequently diagnosed 39 40 infections. The most commonly isolated bacteria type was Acinetobacter baumannii (22.12% overall), isolated especially from the respiratory tract and resistant to all the β -lactam antibiotics 41 including carbapenems. Antimicrobials intake at the ICU is much higher compared to medical 42 43 and surgical wards. After we tested the existence of a possible connection between antibiotic 44 consumption and antibiotic resistance of bacteria, it was revealed that on our sample exists a 45 poor positive association.

46 Key words: antibiotic; defined daily dose; infection; intensive care unit; bacteria; susceptibility.

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49 **1. Introduction**

Antimicrobial resistance is one of the major problems in global public health, and the main cause behind the increasing presence of resistant pathogens is the inappropriate and widespread use of antimicrobial agents. The outcome is increased morbidity and mortality, and rise in healthcare costs (Prestinaci et al., 2015).

The European Centre for Disease Prevention and Control (ECDPC) (2017) has confirmed that 54 55 many countries in Europe, including Romania, present high antimicrobial resistance rates. 56 However, the report does not include all hospitals. While the resistance of gram-positive bacteria has remained almost the same (ECDPC, 2017), there has been an upward trend in the resistance 57 58 of gram-negative species. This increase has been shown to be related above all to thirdgeneration cephalosporins, fluoroquinolones, carbapenems and aminoglycosides for Klebsiella 59 pneumoniae and to ceftazidime, piperacillin tazobactam and fluoroquinolones for Pseudomonas 60 61 aeruginosa. For instance, in 2017, rates of resistance of K. pneumoniae were reported to be 22.5% for carbapenems and 64.1% for third generation cephalosporins and fluoroquinolones 62 (ECDC, 2017). Acinetobacter spp., which has fewer therapeutic options (Lin and Lan, 2014), 63 presented a combined resistance rate of 81.3%. 64

Production and marketing of new antimicrobial agents directed against new bacterial targets or receptors seem unable to keep up with the development of antibiotic resistance (Fair and Tor, 2014) Experimental findings suggest that some vegetable extracts can exert some protection against bacterial infections, but they are not yet in use (Abdel-Daim et al., 2018).

69 Several studies have demonstrated a higher presence of antimicrobial resistance in ICUs than in 70 general patient-care departments (Brusselaers et al., 2011; Meyer et al., 2010). Other surveys 71 have additionally shown that monitoring of hospital data can change prescriptions in specific patient-care areas (Lai et al., 2016; Owens et al., 2004). One of the primary goals of these studies
was to understand the connection between antimicrobial use and the increasing resistance of
pathogens, as well as to define their relationship with individual patient-care areas.

Antimicrobial consumption in Romania is complex and still largely unexplored, and data from different studies are fragmented. Published or unpublished reports on antibiotic consumption have shown use that has greatly increased (is proportionally high) and excessive (Cioca and Munteanu, 2019). Increased consumption leads to dissemination of antibiotics in the environment, from excretion from the human body and through inappropriate disposal of medicinal waste (Bungau et al., 2015, Bungau et al., 2016).

81 Significant environmental and health impacts, difficult to control, occur due to urban waste streams with high concentrations in pharmaceutical products such as antibiotics. These 82 compounds are not completely removed in wastewater treatment plants, and their residues may 83 84 be released through effluents into the aquatic environment (Turkdogan and Yetilmezsov, 2009). Because they pollute soil and water, they have negative effects on organisms in the environment, 85 and they also generate negative effects on human health, one of the most serious being the 86 increase in antibiotic resistance and interference with the hormonal system. Increasing 87 concentrations in active substances are the cause of the increase of the negative effects (Bungau 88 et al., 2018). 89

90 Recent data on use in hospitals or the primary-care sector show that the variation in resistance to 91 different antibiotics between different departments can be explained by the variation of selective 92 pressure on resistance to this type of drug (Oz et al., 2014). In Romania, data on antibiotic use or 93 susceptibility of isolated strains are not very well characterized, due to few studies or reports 94 having evaluated the subject. The aim of this study was to evaluate the antibiotic consumption 95 for one year and the variability in the present pathogens, to compare the antimicrobial 96 prescriptions with the pathogen sensitivity spectrum, and to suggest possible future interventions. 97

98 **2. Methods**

99 This study was carried out using data available at the ICU of the Emergency Clinical County 100 Hospital of Oradea, Romania. It is a clinical university hospital, with 917 acute-care beds, 6.5% 101 belonging to the ICU. For this study, data were collected over the period between January 1 and 102 December 31, 2017. The information used was selected from the medical records of the patients, 103 InfoWorld and the WHONET database on the number and type of pathogens, respecting each 104 patient's privacy.

Matrix-assisted laser desorption/ionization-time of flight (MALDI-TOF) was used for the 105 identification of isolates (Clark et al., 2013); it is a three-step process, as follows: 1. mixture of a 106 107 portion of a colony (the sample) with a matrix material and applying to a target plate; 2. pulsed laser irradiation of the sample, causing the ablation and the desorption of the matrix material and 108 the sample; 3. finally, ionization of the molecules of the analyte (by protonation / deprotonation 109 in the hot plume of ablated gases); then, they can be accelerated into whichever mass 110 spectrometer. The mass spectra are generated and analyzed by a special software, being 111 compared with the stored profiles providing a faster species diagnosis. 112

113 The VITEK 2 system and Kirby–Bauer disk diffusion method were used for the antibiotic 114 susceptibility testing. Suspensions were prepared by emulsifying the bacterial isolates in 0.45 % 115 saline to the equivalent of a 0.5 McFarland turbidity standard. The isolate was classified as 116 susceptible, intermediate, or resistant based on the Clinical Laboratory Standards Institute 117 (CLSI) criteria (2018). *E. coli* ATCC 25922, *S. aureus* ATCC 29213, *P. aeruginosa* ATCC 27853, *E. faecalis* ATCC 29212 were used as quality control strains to check the quality of culture media, and antimicrobial cards and disks. The pharmacy has a computerized record of each antibiotic prescribed, and antimicrobial data were extracted from the system.

Evaluation of the prescribed antimicrobials was carried out using the DDD methodology, which 122 was derived from the ATC/DDD Index 2018 made by the WHO Collaborating Centre for Drug 123 124 Statistics Methodology (2018), the reference standard for drug utilization methodology. WHO promotes the use of the DDD methodology for each drug and route of administration. The 125 methodology defines the DDD as the assumed average maintenance adult dose per day for the 126 127 main direction of use of the agent and the maintains updates. The WHO DDD has also been used to demonstrate a quantitative, ecological relationship between antimicrobial use and resistance in 128 hospitals (Muller et al., 2003; 2004; Westh et al., 2004; Kern et al, 2005). 129

Hospital consumption of antimicrobials was calculated by adding the total number of grams of each antimicrobial used during the period of interest (one year). The result is divided by the WHO-assigned DDD resulting number of DDD (DDDs). To control the population number, the antimicrobial use density was determined, expressed as DDD / 1000 PD, for each antibiotic and route of administration. The annual number of PD was provided by the hospital's admission department.

In this research, antibiotics were defined as antibacterial for systemic use or group J01 of the WHO ATC classification system. The antibiotics excluded from this study were: antibacterial against tuberculosis (rifampicin), as this pathology is not treated in our institution; rifaximin used in diarrheal disorders without a specific etiology; topical antibiotics (parenterally administered prevalent); oral colistin and vancomycin (not used); metronidazole and antifungals (used without having a sensitivity test); those not present in this study hospital (fosfomycin, ticarcillin,
ticarcillin + tazobactam, tobramycin).

The analyzed classes of antibiotics were cephalosporins, fluoroquinolones, aminoglycosides and 143 carbapenems prescribed in therapy and prophylaxis of hospital infections produced by gram-144 negative and gram-positive bacteria. Almost all antibiotics were administered parenterally. The 145 positive pathological samples were sputum, fluids (pleura, peritoneal, pericardial), trachea-146 147 bronchi aspirates, secretions of lesions, wounds, catheters, urine and blood. Antibiotic sensitivity 148 analysis was performed using a cumulative antibiogram, and only the percentage of strains considered susceptible was reported. Descriptive analysis was performed. Values are expressed 149 150 as numbers with percentages; WHO DDD/1000 PD are calculated as described above.

To seek potential connection between antibiotic consumption and antibiotic resistance of bacteria, we performed a correlation analysis using the SPSSv19 software and for the Scatter Plot – the Microsoft Excel 2013 software. The Pearson coefficient, the determination coefficient and the p value (obtained from the correlation model and considering α =0.05 as the confidence level) were also calculated.

156

157 **3. Results**

The overall consumption of antimicrobial agents (expressed as the total number of DDD of antibiotics – ATC group J01, antibacterial for systemic use) was 114,269, and the overall DDD per 1000 PD was 2180.4 in 2017. **Table 1** summarizes the antibiotic use for the whole hospital as well as by department. Most of these were used in the ICU, followed by the surgical and medical wards. 163 The parenteral administration of antibiotics represented almost 94.54%, while oral forms 164 represented only 5.46% of total antibiotic intake. Dividing antibiotic use by surgical/medical 165 wards and ICU, it was found that more than a half of the total antimicrobials were consumed in 166 the latter.

General use for the prescribed antibiotics classes is presented in **Table 2**. The most prescribed antibiotics were cephalosporins in all departments (51.55% of the total), followed by fluoroquinolones (10.91%), aminoglycosides (7.46%), penicillin and beta-lactamase inhibitor combinations (6.36%), carbapenems (5.66%), penicillin (3.84%) and cefoperazone + sulbactam (3.71%) (Table 2). Less frequently prescribed were linezolid, macrolides and tetracyclines.

The total consumption of antimicrobials in the ICU was 1247.4 DDD/1000 PD in 2017 (**Table** 1). Distribution of the top ten antibiotics was: ceftriaxone (J01DD04, 39.65%), levofloxacin (J01MA12, 7.02%), cefuroxime (J01DC02, 6.30%), cefoperazone + sulbactam (J01DD62, 4.98%), meropenem (J01DH02, 4.79%), colistin (J01XB01, 4.43%), amikacin (J01GB06, 4.13%), piperacillin + tazobactam (J01CR05, 3.43%), ertapenem (J01DH03, 3.04%), ciprofloxacin (J01MA02, 2.37%).

Less frequently prescribed were the following antibiotics: oxacillin (J01CF04), ampicillin +
sulbactam (J01CR01), doxycycline (J01AA02), ceftaroline fosamil (J01DI02), cefixime
(J01DD08), clarithromycin (J01FA09), azithromycin (J01FA10), benzylpenicillin (J01CE01),
ampicillin (J01CA01), cefaclor (J01DC04).

The total number of isolated pathogens in the hospital was 2750, and the most diagnosed infections were those of the urinary tract (1031), followed by wounds (947) and respiratory infections (651). Urinary and respiratory tract infections thus dominate (78.61%), followed by wounds (8.99%) (**Table 4**). The most commonly isolated pathogen was *Escherichia coli* (26.07%), which is responsible for most of the urinary tract infections in the entire hospital (**Table 5**). This strain showed good sensitivity rates to all antibiotics tested, except for ampicillin (**Figure 1A**).

The second most frequently isolated pathogen was *Staphylococcus aureus* (13.85%), which was the common cause of wound and respiratory tract infections; its antibiotic sensitivity patterns is presented in **Figure 1B**. *Enterobacter* spp. was the third most frequently isolated pathogen from respiratory and urinary tract infections. *Enterobacter* spp. showed lower sensitivity rates especially to fluoroquinolones and Gentamycin. Its resistance rates were higher : 77.4% for ceftriaxone, 74.6% for ceftazidime, and 65.62 % for cefoperazone (**Figure 1C**).

Enterococcus faecalis and *Proteus* spp. were isolated from urinary tract infections and wounds at almost the same ratio (8.18% and 8.58% respectively). Isolated *Proteus spp.* strains also showed high resistance rates, and they represented 8.53% overall, but therapeutic options were more varied and included aminoglycosides, cefoperazone + sulbactam, piperacillin + tazobactam, and carbapenems. Isolated strains showed low sensitivity rates to cephalosporins and fluoroquinolones despite their frequent prescription, as represented in **Figure 1D**.

Over the same period, a total of 868 strains of pathogens were isolated in the ICU. The most common was *Acinetobacter baumannii* (22.12%), especially from the respiratory tract. This multidrug-resistant pathogen showed sensitivity to colistin (99.03%), ampicillin + sulbactam (62%), and cefoperazone + sulbactam (41.7%), as presented in **Figure 1E**. *E. coli* and *Enterobacter* spp. were isolated at equal proportions (14.41%). Those two strains were the second-most frequently isolated in the ICU, followed by *Proteus* spp. (8.53%), *Pseudomonas aeruginosa* (7.72%), *Klebsiella* spp. (5.30%) and *Serratia* spp. (5.18%). *Morganella* spp., 208 *Citrobacter* spp., *Providencia stuartii*, *Burkholderia complex* and *Stenotrophomonas maltophylia*209 were isolated in small numbers (less than 1% for each).

The fourth etiological agent of the infectious disease in the ICU was *Pseudomonas aeruginosa*, which showed increased resistance to the tested antibiotics. In this case, the therapeutic options were colistin, cefoperazone-sulbactam, and ciperacillin + tazobactam (**Figure 1F**). Half of the tested strains were resistant to ceftazidime and 62.8% to cefepime, but both antibiotics were relatively rarely administered.

A total of 128 *Klebsiella* spp. strains were isolated in the evaluated period in the ICU, representing 5.30%; these strains exhibited low rates of sensitivity to the cephalosporins (26% in the case of ceftriaxone) and fluoroquinolones (**Figure 1G**), which were frequently prescribed.

Gram-positive microorganisms were susceptible to linezolid (100%), teicoplanin (97.01%), and vancomycin (96.81%), in descending order. The slight reduction in susceptibility to vancomycin was found only for *Enterococcus* spp. Among these antimicrobials, the most prescribed was teicoplanin, followed by vancomycin. Antibiotic susceptibility to gram positive strains can be observed in **Figure 2**.

By analyzing the cumulative antibiogram results for both gram positive and negative microorganisms, the best sensitivity rates were for linezolid (J01XX08, 100%), colistin (J01XB01, 99.03%), teicoplanin (J01XA02, 97.01%), and vancomycin (J01XA01, 96.81%), followed by amoxicillin + clavulanic acid (J01CR02, 67.18%), ertapenem (J01DH03, 66.26%), tetracycline (J01AA07, 63.85%), cefoperazone + sulbactam (J01DD62, 63.76%), piperacillin + tazobactam (J01CR05, 59.75%), amikacin (J01GB06, 56.8%), meropenem (J01DH02, 55.16%), trimethoprim + sulfamethoxazole (J01EE01, 52.83%), and ampicillin + sulbactam (J01CR01, 230 50.85%). The sensitivity rates of the tested antibiotics (according to the cumulative antibiogram)
231 are presented in Table 6.

For the correlation model, the data presented in Table 3 and Table 6 were used. As Table 3 232 contains 32 entries and Table 6 contains 33 entries, we analyzed only the 27 common entries 233 (Table 7 which presents values obtained from calculating the percentage of antibiotic use and the 234 sensitivity values of the tested antibiotics). For these two variables, a correlation model was run 235 236 and a positive insignificant correlation (the Pearson coefficient r=0.21; determination coefficient 237 $R^2=0.04$; p value returned from the correlation model p=0.28) was obtained. Our sample thus shows a poor association between the variables, but this relation can't be generalized for the 238 239 whole population. In order to have a better image of this relation, the association was plotted using a Scatter Plot chart, and a line chart as well – in order to highlight the dynamics of our data 240 241 (Figure 3 and 4).

242

243 4. **Discussion**

Antimicrobials used in hospitals vary widely, a fact that can be explained by differences in patient and hospital characteristics, antibiotic policies, physician education, and healthcare systems. There is also great variability in the antibiotic sensitivity spectrum of isolated strains (Leekha et al., 2011).

The present study highlights the combination of antibiogram reports with antibacterial consumption in the ICU, which provides potential measures for lowering morbidity and mortality. The prescribed daily dose for most of the antibiotics corresponded to the WHO defined daily dose, reflecting adherence to international recommendations. Antimicrobials intake at the ICU is much higher than that at the medical and surgical wards, and they are prescribed mainly for respiratory and urinary tract infections in a manner similar to other data reported in literature, in accordance with the type of patient population and the geographical location studied (Bitterman et al., 2016; Gianino et al, 2019). Antimicrobial consumption, sensitivity rates and spectrum of pathogens isolated in the ICU and medical/surgical departments showed disputable differences.

The most frequently prescribed antibacterial was ceftriaxone (39.65%), a third-generation 258 259 cephalosporin that is active on Enterobacterales, except Enterobacter spp. In the present study, 260 E. coli, Klebsiella spp., Proteus spp., Citrobacter spp. and Serratia spp. were isolated in 33.53% of cases, but annual sensitivity rates to ceftriaxone were low (28.35%), except for E. coli. The 261 262 high consumption of ceftriaxone despite the limited sensitivity of isolated strains could be explained by its use in empirical therapy, and it could be a reasonable therapeutic option because 263 the most commonly isolated pathogen is Escherichia coli from urinary tract infections (which 264 265 demonstrate good sensitivity rates to all antibiotics tested, except for ampicillin). This contrasts with other studies, which showed an increase in the prevalence of extended-spectrum β-266 lactamase - producing Escherichia coli (ESBL - producing E. coli) worldwide (Alqasim et al., 267 268 2018).

As mentioned previously, *Enterobacter* spp. has lower sensitivity rates to fluoroquinolones and gentamycin. According to the ECDPC (2016), resistance to third-generation cephalosporins was reported in 44% of cases of *Enterobacter* spp. isolates. In this study, the resistance rates were higher: 77.4% for ceftriaxone, 74.6% for ceftazidime, and 65.62 % for cefoperazone.

Levofloxacin and cefuroxime, the second- and third-most used in the ICU (7.02% and 6.3%
respectively), can be used for both gram-negative and positive strains, showing sensitivity rates
of 33.8% and 45% respectively. Their consumption could be justified, as *E. coli* was the second

276 greatest etiological agent of infections in ICU and has good sensitivity to levofloxacin and 277 cefuroxime, which furthermore are used in treating half the cases of *Staphylococcus aureus* (the 278 fourth etiological agent in the ICU). For *Enterococcus faecalis* strains, 71.42% were sensitive to 279 Levofloxacin, but they represent only 2.53% of all strains. Sensitivity rates to Cefuroxime, the 280 second etiological agent of infections, were 53.1% for *Staphylococcus aureus* strains and 62.5% 281 for *E. coli* strains.

282 According to our results, the most commonly isolated pathogen in ICU was Acinetobacter 283 baumannii, which has sensitivity to colistin, ampicillin + sulbactam, and cefoperazone + sulbactam. All three of these antimicrobials were relatively rarely prescribed, although they can 284 285 eradicate the infection, usually respiratory infections, especially in combination with carbapenems (Lenhard et al., 2017; Altun, et al., 2014). Acinetobacter baumannii was the only 286 pathogen that was consistently isolated in the ICU (22.12% of total pathogens) in a way similar 287 288 to other studies. All the Acinetobacter baumannii isolates were resistant to all the β -lactam antibiotics, including the carbapenems, as other studies have described (Handal et al., 2017). 289

According to ECDPC (2016), the most frequently isolated microorganism was *Pseudomonas aeruginosa* in ICU-acquired pneumonia episodes. Moreover, the same study indicated that rates of resistance of *Klebsiella* spp. isolates were 38% to third generation cephalosporins and 11% to carbapenem. In contrast, isolated *Klebsiella spp* strains showed good sensitivity to Amikacin (88.6%), as demonstrated by other studies as well (Baicus et al., 2018).

Fluoroquinolones were the second therapeutic option, in both hospitals and ICUs: ciprofloxacin, although quite commonly used, and moxifloxacin (less often prescribed) showed annual sensitivity of only 38.17% and 12.76% respectively to the strains tested. In the same situation, the following were also prescribed quite often but with relatively low sensitivity: clindamycin,ceftazidime, cefoperazone and cefepime.

In the case of carbapenems, the most prescribed was meropenem, which showed a sensitivity rate of 55.16%. ertapenem, which is prescribed less often than meropenem, presented better sensitivity in 66.26% of the tested strains. Both are useful in the treatment of pneumonia, complicated urinary tract infections (including pyelonephritis), complicated intra-abdominal infections, and complicated skin infections. This difference in sensitivity rate was clinically significant and indicated underutilization of ertapenem and overutilization of meropenem. Imipenem, although less, used showed sensitivity in 42.3% of the tested gram-negative strains.

As for aminoglycosides, 56.8% of the strains tested were sensitive to amikacin, which was
predominantly prescribed. This figure was 43.29% for gentamycin, which was prescribed less
often.

Combinations (cefoperazone + sulbactam, ampicillin + sulbactam, amoxicillin + clavulanate,
 piperacillin + tazobactam) showed good annual sensitivity rates. Cefoperazone + sulbactam and
 piperacillin + tazobactam were more frequently prescribed than amoxicillin + clavulanate.

For the *Staphylococcus* spp. strains, 35% were methicillin sensitive. As a result, they should have been treated with oxacillin, but this antibiotic has not been prescribed to the same extent. The most frequently isolated microorganisms were coagulase-negative staphylococci in ICU-acquired bloodstream or catheter infections.

Some of the isolated strains were tested for susceptibility to tetracycline and trimethoprim +
sulfamethoxazole and showed good sensitivity rates (63.85% and 52.83% respectively).
Doxycycline was rarely administered to patients included in our study, and trimethoprim +
sulfamethoxazole was not administered.

According to the data available in Romania, statistics on antibiotic use or susceptibility of isolated strains are not very clear or well centralized; only very few research studies evaluated and presented valuable conclusions on these subjects. Even under these circumstances, published reports show increased antibiotic resistance in diverse healthcare settings (Bedreag et al., Penes et al., 2017; Timofte et al., 2016).

The analysis of antimicrobial consumption is an important component of medical audits for 326 327 monitoring, evaluation, and making the necessary changes in physicians' prescribing patterns in 328 order to achieve rational and cost-effective medical care. Yet, antibiotic resistance is a major worldwide concern, and selective use of antimicrobial drugs is by far the most important driving 329 330 force behind the development of resistance. As a result, stewardship initiatives should direct educational efforts to shortening durations of antimicrobial exposure and review the current 331 approach to empirical prescribing (Davey et al., 2017). Antimicrobial stewardship programs 332 333 should be implemented to control microbe resistance in inpatients, and surveillance of antimicrobial use with observation of bacterial resistance rates can help control resistance in 334 general. Each hospital must enact priority measures to control the emergence of antibiotic-335 resistant bacteria, including control of overuse (Doron and Davidson, 2011). 336

Optimal management of antimicrobial use implies that physicians should prescribe patients the least harmful treatment, using the cheapest antibiotic, for the shortest possible time needed to cure or prevent an infection. Literature data recommend four possible strategies to ensure that this goal is achieved: rational use of all antimicrobials; selective control and restriction of antimicrobial agents; cyclic or rotational antimicrobial use; and the use of combined antimicrobial therapy, where appropriate, to prevent the emergence of resistance (Shales et al., 1997). 344 Considerable efforts have been made in recent years and should be continued and intensified, to educate physicians, pharmacists, and the population about the importance of minimizing the 345 unnecessary use of antimicrobials and the inappropriate disposal of waste in the form of expired 346 or unused medicinal products, in order to reduce negative impact on the health of the population 347 and the environment (Bungau et al., 2018; Tit et al., 2016). Romania is among the top four 348 countries of Europe in terms of the total amount of antibiotics used (ECDPC, 2017). This is due 349 350 to insufficient information on the effect of antibiotics and insufficient education on their role and 351 effects. The consequences of abusive use of antibiotics are precisely increases in antibiotic resistance and the amount of antibiotic-derived waste (Bungau et al., 2018). 352

353 It is increasingly recognized and affirmed the role of the environment in the global spread of antibiotic clinical resistance (Singer et al., 2016). Also, bacterial resistance to antibiotics is a 354 natural phenomenon found in various and numerous locations in the environment, apparently 355 356 without antibiotics (Pallecchi et al., 2007) or with trace/low levels of antibiotics of natural synthetic origin (Raaijmakers and Mazzola). Although antibiotic concentrations found in the 357 environment are not very high, in the order of μg or ng/L, due to the cumulative effect of the 358 intensified synthesis of new antibiotics and their continuous release into ecosystems, the new 359 mechanisms of resistance to these antibiotics have adapted and grown; therefore, bacteria have 360 developed resistance to numerous antibiotics (Ding and He, 2010). Research has shown that 361 362 antibiotic resistance genes (ARGs) have ecological origins, but the release and accumulation of additional antimicrobials in the environment facilitated their spread (Baquero et al., 2008). 363 Consequently, ARGs are found in all environments and are currently considered becoming 364 visible pollutants (Pruden et al, 2006). 365

366 There are few studies in Romania that quantify the existence of antibiotics in an ARB or ARG environment. The presence of antibiotics in the classes of penicillin, cephalosporin and 367 tetracycline has been investigated by Opris et al. (2013) in wastewater and effluent samples 368 collected at a waste water treatment plant. The results indicated the presence of ceftriaxone 334 369 µg/L, tetracycline 146 µg/L and doxycycline 110 µg/L in residual water; these antibiotics were 370 not detected in water samples taken from effluents. Ceftriaxone was determined in the largest 371 372 amount, correlated with the results of our study showing that ceftriaxone is the most commonly prescribed antibacterial agent. 373

The impact and presence of waste resulting from or generated by antibiotics were investigated as 374 375 well (Szekeres et al., 2017). This research also evaluated the diversity of numerous antibiotic resistance genes (ARGs) along with the composition of the bacterial community that can be 376 found in the wastewater effluents discharged from different Romanian hospitals. The results 377 378 pointed to increased concentrations of certain β -lactam antibiotics including trimethoprim and glycopeptides. Moreover, ARGs collected from contaminated effluents were diverse, exhibiting 379 an increased relative abundance concomitantly to the dominance of the genera Acinetobacter, 380 Enterococcus, Pseudomonas and members of the Enterobacteriaceae family. The community 381 structure of these potential sources of resistance genes as demonstrated in our survey (Szekeres 382 et al., 2017) varied from one hospital to another. 383

It is noteworthy that the treatment of wastewater installed by authorities removed only partially these pollutants (Szekeres et al., 2017), which thus infers that residual waters in hospitals are important contributing factor in the dissemination of resistant bacteria and resistance genes. This is also supported by other recent findings evidencing the presence of isolated bacterial species in the environment (Hrenovic et al., 2014, Khan et al., 2018, Marinescu et al., 2015). 389 The present study on antimicrobials provides many important characteristics of antibacterial prescription. The antibiotic classes prescribed in more than 50% of cases were cephalosporins, 390 fluoroquinolones, aminoglycosides, and beta lactam/beta lactamase inhibitors. Cephalosporins 391 continued to be a main prescription of antibiotic therapy in the hospital because of their broad 392 spectrum of activity and clinical efficiency. The most commonly prescribed agent was 393 Ceftriaxone, the utilization pattern being similar to the one described by Anand et al (2016). The 394 395 preference of fluoroquinolones over aminoglycosides points towards a trend of using less toxic 396 antimicrobial classes and the broader spectrum of action including both gram-positive and negative. Many studies in this field demonstrated that the antibiotic resistance proportion of 397 398 bacterial strains presents large variations in different countries, most probably due to the differences between the consumption of antibiotics and hospital infection control. Proper and 399 effective implementation of prevention and control programs regarding antimicrobial resistance 400 401 is imperative at every hospital and pharmacy level (Davey et al., 2017; Lee et al., 2013).

Testing in our research the possible connection between antibiotic consumption and antibiotic resistance of bacteria, pointed out a weak positive association r=0.21; thus, it can be admitted that if the sensitivity of an antibiotic is increasing, the use of that antibiotic is increasing as well. Because of this poor association, medicine uses a wide variety of different types of antibiotics because there is no particular drug / treatment that works every time in every patient, which is easy to understand given the complexity of the human body.

Limitations of this study can be considered to be the collection of data from only one ICU, in a single hospital, and the short observation period. In addition, it was not possible to differentiate every patient and their comorbidities. The majority of the patients whose data were registered in the hospital were elderly, suffering chronic diseases (such as diabetes and hypertension). Another 412 (potential) limit of the study is that it may underestimate year-over-year changes in413 susceptibility.

414

415 Conclusions

This research shows differences between antibacterial use and the sensitivity spectrum of isolated strains. Among the isolated pathogens, the preponderance of gram-negative bacteria *Acinetobacter* spp. strains stands out, which can mean insufficient measures to control nosocomial infections.

Based on the record data, priority measures should be implemented to control the resistance of
microbes in inpatients and the surveillance of antimicrobial use, in particular by controlling overuse. Future studies should be undertaken to verify these findings in different settings and among

- 423 diverse inpatient populations.
- 424
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- 596 List of figures
- 597 Figure 1. Antibiotic sensitivity patterns of: A. E. coli isolates; B. Staphylococcus aureus; C.
- 598 Enterobacter spp.; D. Proteus spp.; E. Acinetobacter baumannii; F. Pseudomonas aeruginosa;
 599 G. Klebsiella spp.
- 600 Legend for figures 1.A.-G.: AK Amikacin; AMC Amoxicillin + Clavulanate; ; AM -
- 601 Ampicillin; CAZ Ceftazidime; CES Cefoperazone + Sulbactam; CFM Cefixime; CFP -
- 602 Cefoperazone; CIP Ciprofloxacin; CLR Clarithromycin; CN Gentamycin; CRO -
- 603 Ceftriaxone; CT Colistin; CXM Cefuroxime; DA Clindamycin; E Erythromycin; ETP –
- 604 Ertapenem; FEP Cefepime; IPM Imipenem; LEV Levofloxacin; LNZ Linezolid; MEM
- 605 Meropenem; MXF Moxifloxacin; OFX Ofloxacin; OX Oxacillin; P Benzylpenicillin;
- 606 SAM Ampicillin + Sulbactam; SXT Trimethoprim + Sulfamethoxazole; TE Tetracycline;
- 607 TEC Teicoplanin; TZP Piperacillin + Tazobactam; VA Vancomycin.
- 608 Figure 2. Antibiotic sensitivity patterns of gram-positive strains
- 609 Figure 3. The association between the antibiotic use values and the sensitivity values
- 610 Figure 4. The dynamics between the antibiotic use values and the sensitivity values



Figure 1.



Figure 2.



Figure 3.



Figure 4.

Table 1. Overall antibiotic use

Department	No. of PD	No. of DDDs	DDD/1000 PD
Intensive Care Unit	17,053	21,272	1247.4
Surgical wards	110 <mark>,</mark> 530	53,387	483.0
Medical wards	87 <mark>,</mark> 859	39,610	450.8
TOTAL	215,442	114,269	2180.4

	DDD/1000 PD				
Antibacterial	Surgical	Intensive	Medical	Total	
	warus	care unit	warus	No. of doses	%
Cephalosporins	31	61.99	15.53	108.52	51.55
Fluoroquinolones	3.69	12.74	6.54	22.97	10.91
Aminoglycosides	2.79	6.331	6.581	15.70	7.46
Penicillins and beta- lactamase inhibitor	1.08	5.9	6.403	13.38	6.36
Carbapenems	0.62	11.1	0.19	11.91	5.66
Penicillins	5.6	0.96	1.524	8.084	3.84
Cefoperazone + sulbactam	0.29	6.12	1.4	7.81	3.71
Polymyxins (Colistin)	0.04	5.45	0.07	5.57	2.65
Lincosamide (Clindamycin)	2.42	2.54	0.46	5.42	2.57
Glycopeptides	0.11	4.78	0.041	4.93	2.34
Tigecycline	0.01	2.59	0.009	2.61	1.24
Linezolid	0.029	1.93	0.01	1.97	0.94
Macrolides	0.17	0.33	0.69	1.19	0.57
Tetracycline (Doxycycline)	0.01	0.22	0.17	0.41	0.20

Table 2. General use for the prescribed antibiotics classes

A	ATC	DDD/1000 PD	
Antibacterial	code	No.	%
Ceftriaxon	J01DD04	48.76	39.65
Levofloxacin	J01MA12	8.64	7.02
Cefuroxime	J01DC02	7.75	6.30
Cefoperazone + sulbactam	J01DD62	6.12	4.98
Meropenem	J01DH02	5.89	4.79
Colistin	J01XB01	5.45	4.43
Amikacin	J01GB06	5.07	4.13
Piperacillin + tazobactam	J01CR05	4.22	3.43
Ertapenem	J01DH03	3.74	3.04
Ciprofloxacin	J01MA02	2.91	2.37
Teicoplanin	J01XA02	2.76	2.24
Tigecycline	J01AA12	2.59	2.11
Clindamycin	J01FF01	2.54	2.07
Ceftazidime	J01DD02	2.04	1.66
Vancomycin	J01XA01	2.02	1.64
Linezolid	J01XX08	1.93	1.58
Cefepime	J01DE01	1.91	1.55
Imipenem cilastin	J01DH51	1.47	1.20
Amoxicillin + clavulanate	J01CR02	1.41	1.15
Gentamycin	J01GB03	1.26	1.03
Moxifloxacin	J01MA14	1.19	0.97
Cefoperazone	J01DD12	1.13	0.92
Oxacillin	J01CF04	0.57	0.46
Ampicillin + sulbactam	J01CR01	0.27	0.22
Ceftaroline fosamil	J01DI02	0.24	0.20
Doxycycline	J01AA02	0.22	0.18
Clarithromycin	J01FA09	0.22	0.18
Benzylpenicillin	J01CE01	0.21	0.17
Ampicillin	J01CA01	0.18	0.15
Azthromycin	J01FA10	0.11	0.09
Cefixime	J01DD08	0.1	0.08
Cefaclor	J01DC04	0.06	0.05

Table 3. Antibiotic use expressed as DDD/1000 PD in the ICU

Type of infection	Total	Intensive care unit $(n = 879, 100\%)$	
	(n = 28/0)	No.	%
Respiratory tract	651	586	66.67
Urinary tract	1031	105	11.95
Wounds	947	79	8.99
Blood	58	38	4.32
Catheter	46	40	4.55
Fluids (pleural, pericardial, peritoneal)	137	31	3.53

Table 4. Number and frequency (%) of infection types

Strains	Total	Intensive care unit $(n = 868, 100\%)$	
	(n = 2750)	No.	%
Staphylococcus aureus	381	65	7.49
Coagulase negative staphylococcus	119	47	5.41
Enterococcus faecalis	225	22	2.53
Enterococcus faecium	56	27	3.11
Streptococcus pneumoniae	2	1	0.12
Streptococcus pyogenes	3	0	0
Streptococcus spp. (B, F, G, viridans)	33	14	1.61
Escherichia coli	717	125	14.40
Enterobacter spp.	283	125	14.40
Proteus spp.	236	74	8.53
Klebsiella spp.	128	46	5.30
Acinetobacter spp.	240	192	22.12
Pseudomonas aeruginosa	195	67	7.72
Serratia spp.	80	45	5.18
Morganella spp.	32	4	0.46
Citrobacter spp.	5	1	0.12
Providencia stuartii	7	5	0.58
Burkholderia complex	7	7	0.81
Stenotrophomonas maltophylia	1	1	0.12

Table 5. Number and frequency (%) of the pathogens isolated

Antibacterial name	ATC code	Sensitivity rates %
Linezolid	J01XX08	100
Colistin	J01XB01	99.03
Teicoplanin	J01XA02	97.01
Vancomycin	J01XA01	96.81
Amoxicillin + Clavulanic acid	J01CR02	67.18
Ertapenem	J01DH03	66.26
Tetracycline	J01AA07	63.85
Cefoperazone + Sulbactam	J01DD62	63.76
Piperacillin + Tazobactam	J01CR05	59.75
Amikacin	J01GB06	56.8
Meropenem	J01DH02	55.16
Trimethoprim + Sulfamethoxazole	J01EE01	52.83
Ampicillin + Sulbactam	J01CR01	50.85
Ampicillin	J01CA01	45.2
Cefuroxime	J01DC02	45
Clarithromycin	J01FA09	43.75
Oxacillin	J01CF04	43.75
Gentamicin	J01GB03	43.29
Imipenem	J01DH51	42.3
Erythromycin	J01FA01	41.37
Amoxicillin	J01CA04	40.62
Ofloxacin	J01MA01	39.57
Cefoperazone	J01DD12	39.53
Ciprofloxacin	J01MA02	38.17
Ceftazidime	J01DD02	37.03
Cefixime	J01DD08	35.32
Clindamycin	J01FF01	35.04
Levofloxacin	J01MA12	33.84
Azithromycin	J01FA10	33.33
Benzylpenicillin	J01CE01	32.08
Ceftriaxone	J01DD04	28.35
Cefepime	J01DE01	26
Moxifloxacin	J01MA14	12.76

Table 6. Sensitivity rates of tested antibiotics (according to cumulative antibiogram)

Antibiotic	Antibiotic use	Sensitivity values
Cefixime	0.08	0.35
Azthromycin	0.09	0.33
Ampicillin	0.15	0.45
Benzylpenicillin	0.17	0.32
Clarithromycin	0.18	0.44
Ampicillin + sulbactam	0.22	0.51
Oxacillin	0.46	0.44
Cefoperazone	0.92	0.40
Moxifloxacin	0.97	0.13
Gentamycin	1.03	0.43
Amoxicillin + clavulanate	1.15	0.67
Imipenem cilastin	1.20	0.42
Cefepime	1.55	0.26
Linezolid	1.58	1.00
Vancomycin	1.64	0.97
Clindamycin	2.07	0.35
Tigecycline	2.11	0.64
Teicoplanin	2.24	0.97
Ciprofloxacin	2.37	0.38
Ertapenem	3.04	0.66
Piperacillin + tazobactam	3.43	0.60
Amikacin	4.13	0.57
Colistin	4.43	0.99
Meropenem	4.79	0.55
Cefoperazone + sulbactam	4.98	0.64
Cefuroxime	6.30	0.45
Levofloxacin	7.02	0.34

Table 7. The relation between the antibiotic use expressed as DDD/1000 PD in the ICU and the sensitivity values of tested antibiotics (according to cumulative antibiogram)



Among the isolated pathogens, the preponderance of gram-negative bacteria and the statistically significant isolation of *Acinetobacter* spp. strains stand out, which can mean insufficient measures to control nosocomial infections.