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PROBLEMS OF INFECTIOUS AND PARASITIC DISEASES VOLUME 44, NUMBER 1/2016

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PHENOTYPIC DETECTION OF 16S RRNA METHYLTRANSFERASE-PRODUCING ENTEROBACTERIACEAE BY ROUTINE ANTIBIOGRAM

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ABSTRACT

16S rRNA methyltransferase (16S-RMTase)-mediated aminoglycoside resistance, formerly confined to aminoalvcoside producers as a mechanism of self-defence, has recently been identified in Gram-negative pathogens. The genes encoding these determinants are usually associated with extended-spectrum β -lactamases (ESBLs), but recently they have been found in association with carbapenemases resulting in difficult-to-treat multidrug-resistant bacteria. In this study, we analysed resistance phenotypes of 120 well-characterised, consecutive 16S-RMTase-producing Enterobacteriaceae, collected at a cancer hospital during an 11-years survey period (2004-2015). Our results suggest that concomitant use of gentamicin, amikacin, and apramycin provide a reliable screening for 16S-RMTase production in all Enterobacteriaceae. Moreover, the addition of meropenem as an indicator of carbapenemase production and the double-disc synergy test with cefepime for ESBL detection provided a good tool for further surveillance of 16S-RMTase-producing Enterobacteriaceae at the hospital level. In conclusion, concomitant high level resistance to gentamicin and amikacin, but susceptibility to apramycin could be a sensitive screening tool for rapid detection, improved surveillance, and appropriate treatment of 16S-RMTase-producing multidrug-resistant enterobacterial pathogens.

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Stefana Sabtcheva Laboratory for Clinical Microbiology Specialised Hospital for Active Treatment in Oncology 6 Plovdivsko pole Str 1756 Sofia, Bulgaria Tel. 8076293 E-mail: stefanasabtcheva@gmail.com Keywords: *Enterobacteriaceae*, aminoglycoside resistance, 16S rRNA methyltransferase

INTRODUCTION

Aminoglycoside along with β -lactam resistance raises clinical concerns and its occurrence could compromise the widely used combinations of aminoglycoside and b-lactam antibiotics for the treatment of serious infections caused by pathogenic bacteria. Although the inactivation by aminoglycoside-modifying enzymes remains the most prevalent and clinically relevant mechanism of aminoglycoside resistance, since 2003 an emerging mechanism - the alteration of ribosomal targets by methylation, has gradually increased (1). Unlike aminoglycoside-modifying enzymes that vary in their substrate ranges, the acquired 16S rRNA methyltransferase (16S-RMTase) confers high-level resistance to all 4,6-disubstituted 2-deoxystreptamine (gentamicin, sisomicin, isepamicin, kanamycin, amikacin, tobramycin, and arbekacin), but not to 4,5-disubstituted 2-deoxystreptamine (neomycin), streptomycin, and apramycin (1, 2).

Taking into account the peculiar enzymatic function of 16S-RMTases and the resulting extraordinary high-level resistance to all clinically useful aminoglycosides, we attempted to develop a cost-effective disc-based method to screen for 16S-RMTase producers within the routine antibiogram.

MATERIALS AND METHODS

From January 2004 to December 2015 all Enterobacteriaceae isolated at a 242-bed cancer hospital in Sofia were systematically tested against gentamicin, amikacin, tobramycin, netilmicin, kanamycin and apramycin with disc diffusion method according to Clinical and Laboratory Standards Institute (CLSI) guidelines. Isolates that exhibited concomitantly no inhibitory zone around all aminoglycosides tested, except apramycin, had already been confirmed by PCR analysis to harbour known16S-RMTase genes (3, 4, 5). The so created collection of well-characterised, consecutive 16S-RMTase-producing Enterobacteriaceae was analysed in an attempt to select the best phenotypic indicators able to detect the 16S-RMTase producers within routine susceptibility testing at the hospital.

RESULTS

Between January 2004 and December 2015, routinely performed disc diffusion susceptibility tests composed of the above mentioned aminoglycosides revealed the presence of a resistance phenotype compatible with 16S-RMTase production in 120 clinical *Enterobacteriaceae* (Fig.1).

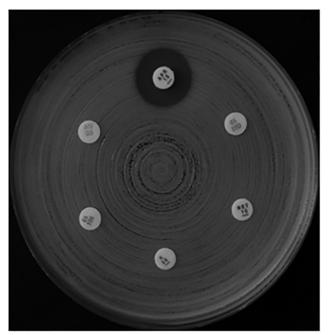


Figure 1: Typical image of 16S-RMTase-mediated resistance phenotype observed in clinical *Enterobacteriaceae* isolates at the cancer hospital. Clockwise: APR, apramycin; GM, gentamicin; NET, netilmicin; K, kanamycin; NN, tobramycin; AN, amikacin. The isolates exhibited concomitantly no inhibitory zone around all aminoglycosides tested, except apramycin.

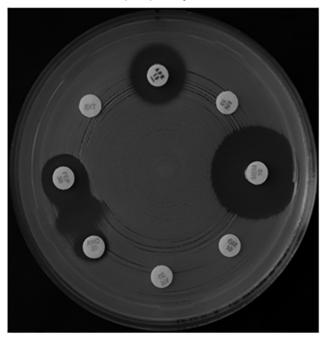


Figure 2: Antibiogram of *armA*-positive *Enterobacter cloacae* strain co-producing ESBL. Clockwise: APR, apramycin; NN, tobramycin; MEM, meropenem ; GM, gentamicin; AN, amikacin; AMC, amoxicillin/clavulanic acid; FEP, cefepime, SXT, trimethoprim/sulfamethoxazole.

All of them were confirmed to harbour 16S-RMTase-encoding genes in association with extended-spectrum β -lactamase (ESBL) genes as reported (3, 4, 5). In 16 of the 16S-RMTase-positive isolates co-production of metallo-β-lactamases (MBLs) was also determined as previously reported (3, 4, 5). Based on analysis of the observed resistance phenotypes and their corresponding genotypes we selected the following antibiotic disc indicators able to screen for 16S-RMTase and associated β-lactamase production that could be included in the routine antibiogram (Fig. 2). Both gentamicin and amikacin discs with no inhibitory zone, along with susceptibility to apramycin, proved indicative of 16S-RMTase production. Synergy between amoxicillin/clavulanic acid and cefepime proved indicative of ESBL production in all Enterobacteriaceae. Finally, meropenem disc was also included as an indicator of carbapenemase producion in accordance with the EUCAST guidelines for detection of resistance mechanisms (6).

DISCUSSION

16S rRNA methyltransferase-mediated aminoglycoside resistance, formerly confined to aminoglycoside producers as a mechanism of self-defence, has recently been identified in Gram-negative pathogens (1). In recent years, however, diverse 16S-RMTases have emerged as acquired resistance determinants in several genera of the Enterobacteriaceae and have been detected not only in humans, but also in animals (7). The genes encoding these determinants are usually associated with ESBLs but recently they have been found in association with carbapenemases resulting in difficult-to-treat multidrug-resistant pathogens (3, 4, 5, 7, 8). Thereby, it would be very important to continue monitoring the trend of 16S-RMTase producers under hospital and countrywide surveillance programs in both humans and animals to prevent their further global spread. In this connection, the development of practical screening techniques for detection of 16S-RMTase-producing Enterobacteriaceae will be of great assistance to the epidemiological study and rapid identification of such isolates in clinical microbiology laboratories.

In this study, we analysed resistance phenotypes of 120 well-characterised, consecutive 16S-RMTase-producing *Enterobacteriaceae*, collected at one hospital during an 11-years survey period (2004-2015) (3, 4, 5). Our results suggest that concomitant use of gentamicin, amikacin, and apramycin provide a reliable screening for 16S-RMTase production in all *Enterobacteriaceae*. Similar suggestion, as the two-step MIC determination of both gentamicin and amikacin, was proposed from other researchers (7). Moreover, the addition of meropenem as an indicator of carbapenemase production and the double-disc synergy test for ESBL detection provided a good tool for further surveillance of 16S-RMTase-producing Enterobacteriaceae at the hospital level. In conclusion, we have developed a cost-effective disc-based method to screen for 16S-RMTase producers and associated resistance markers within the routine antibiogram. Concomitant high level resistance to gentamicin and amikacin, but susceptibility to apramycin could be a sensitive screening tool for rapid detection, improved surveillance, and appropriate treatment of 16S-RMTase-producing multidrug-resistant enterobacterial pathogens.

ACKNOWLEDGEMENTS

We thank Mr. Stoyan Yordanov Atanasov for the precise photography of the antibiograms.

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COMPARISON OF TWO COMBINATION DISC TESTS FOR PHENOTYPIC DETECTION OF CARBAPENEMASE-PRODUCING ENTEROBACTERIACEAE

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ABSTRACT

Background: The prompt and accurate detection of carbapenemase-producing Enterobacteriaceae is essential for patient care and infection control procedures. Therefore, the introduction of a standardised method for routine carbapenemase detection in clinical diagnostic laboratories seems to be mandatory. The aim of the present study was to compare the performance of two commercially available combination disc tests for confirmation of carbapenemase production in Enterobacteriaceae, isolated in Bulgaria. Material and methods: Two commercial combination disc tests (CDTs): the MAST Carbapenemase Detection Set (MAST-CDS) and the KPC&M-BL&OXA-48 disc kit were evaluated in comparison with molecular detection of carbapenemase genes. Test strains comprised 58 well-characterised Enterobacteriaceae from the collection of the National Reference Laboratory for Control and Monitoring of Antibiotic Resistance. Of these, 40 Enterobacteriaceae were confirmed to

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Keywords: *Enterobacteriaceae*, carbapenemase, phenotypic detection, combination disc test.

INTRODUCTION

In Enterobacteriaceae resistance to carbapenems is mainly due to the production of carbapenemases of molecular class A, B, and D or, in rare cases, to the production of extended-spectrum β-lactamase (ESBL) and/or AmpC cephalosporinases combined with reduced permeability of the outer membrane due to loss or mutations in porins (1). Carbapenemases are β-lactamases that vary in their ability to hydrolyse all β -lactam antibiotics depending on the molecular class to which they belong (1, 2). The most common class A carbapenemases are the KPC-types initially confined to Klebsiella pneumoniae isolates, but afterwards continued to spread even outside the Enterobacteriaceae family. Carbapenemases of VIM-, IMP-, and NDM-types represent class B enzymes. Besides Enterobacteriaceae, they are frequently isolated from multidrug-resistant Pseudomonas spp. and Acinetobacter spp. as well. Class D enzymes are mainly represented by OXA-48-like producers identified mainly in Enterobacteriaceae. Genes encoding carbapenemases are often associated with mobile genetic elements, located on plasmids which also carry fluoroquinolone, aminoglycoside, and other resistance determinants (2). This facilitates their rapid dissemination and seriously compromises the treatment of life-threatening infections which is often limited to colistin and tigecycline (3). Therefore, rapid detection of carbapenemase-producers becomes essential for infection control purposes, successful treatment of patients and the preservation of carbapenem efficacy (1). Although the molecular identification of carbapenemase genes remains the "gold standard", phenotypic assessment using the routine antibiogram provided reliable and cost effective method for rapid detection of carbapenemase production (4). In Bulgaria, commercially available combination disc tests were recently introduced for phenotypic detection and differentiation of carbapenemases, based on specific inhibition of the hydrolytic activity of class A and B enzymes. As yet there is no specific inhibitor of class D enzymes, their differentiation is based on the established high level resistance to temocillin defined as phenotypic indicator of production of OXA-48-like carbapenemases (5).

The aim of the present study was to compare the performance of two commercially available combination disk tests for confirmation of carbapenemase production in *Enterobacteriaceae*, isolated in Bulgaria.

MATERIALS AND METHODS

The study was performed with 58 clinical Enterobacteriaceae isolates, with inhibition zone diameter around 10 µg meropenem disc <25 mm (4), subdivided in two groups: carbapenemase-positive and carbapenemase-negative. The first group included 40 strains with already confirmed genotypes by Real-time PCR and sequencing as follows: 17 $bla_{\rm KPC}$ -positive Klebsiella pneumoniae, 3 bla_{OXA-48}-positive K. pneumoniae, 11 bla_{NDM}-positive strains (4 Escherichia coli and 7 K. pneumoniae), and 9 bla_{vim}-positive strains (6 Proteus mirabilis, 2 Serratia marcescens, and 1 K. pneumoniae). Furthermore, 18 carbapenem-non-susceptible and carbapenemase-non-producing Enterobacteriaceae were also included as negative controls. They were confirmed biochemically by the Carba NP test (6).

Phenotypic detection and confirmation of carbapenemase production was performed in parallel with the MASTDISCS[™] ID carbapenemase (*Enterobacteriaceae*) detection disc set (MAST Diagnostics, Merseyside, UK) and the KPC&MBL&OXA-48 disc kit (Liofilchem, Roseto degli Abruzzi, Italy), following the manufacturer's instructions. Briefly, Mueller-Hinton II agar plate was inoculated with overnight culture suspension of the test organism equal to 0.5 McFarland standard. Disc meropenem 10 µg, disc meropenem 10 µg with phenylboronic acid as a KPC inhibitor, disc meropenem 10 µg with cloxacillin as an AmpC inhibitor, disc meropenem 10 µg with dipicolinic acid as a class B metallo-B-lactamase (MBL) inhibitor and disc temocillin 30 µg were placed on inoculated agar plate. Following incubation at 36°C for 18–24 hours, the differences between inhibition zones around the meropenem disc compared to the meropenem discs combined with inhibitors were determined in mm. The test was considered positive for KPC production when the diameter of the growth-inhibitory zone around the meropenem+phenylboronic acid disc was ≥4 mm larger than that around the meropenem disc alone. The test was considered positive for MBL production when the diameter of the growth-inhibitory zone around the meropenem+dipicolinic acid disc was ≥5 mm larger than that around the meropenem disc alone. Finally, the concomitant absence of synergy with any inhibitor and zone diameter <11 mm around the temocillin disc was interpreted as suspicious for OXA-48-like production. As MASTDISCS™ ID carbapenemase detection disc set does not include temocillin disc, it was additionally supplied by MAST in order to obtain comparable results.

RESULTS

Performance results for the two combination disk tests - the MAST Carbapenemase Detection Set (MAST-CDS) and the KPC&MBL&OXA-48 disc kit were easy to interpret and correctly classified all KPC producers to Ambler class A of enzymes, showing significant increase of the growth-inhibitory zone around the meropenem+phenylboronic acid discs compared to that around the meropenem-alone disc (Fig.1).

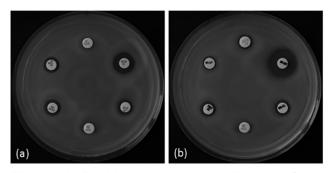


Figure 1: Positive combination disc test for a KPC-producing *K. pneumoniae* isolate (a) using MAST-CDS; (b) using KPC&MBL&OXA-48 disc kit. Clockwise: A/MEM, meropenem; C/MR+BO, meropenem with phenylboronic acid; D/MR+CL, meropenem with cloxacillin; TZP, piperacillin/ tazobactam; TEM/TMO, temocillin; B/MR+DP, meropenem with dipicolinic acid.

Regarding MBL producers some discrepancy was observed between the two combination disk tests. MAST-CDS correctly classified all NDM- and VIM-producing non-*Proteus* species to class B enzymes, whereas the KPC&MBL&OXA-48 disc kit was not able to confirm VIM-production in *S.marcescens* strains (Fig.2 and Fig.3). The both CDTs studied failed to detect VIM production in *P. mirabilis* strains (Fig.4).

Finally, the both CDTs correctly classified all OXA-48 producers to class D of enzymes based on the concomitant absence of synergy with any inhibitors and zone diameter 6 mm with temocillin disc (Fig5). At the same time, no synergy with any of the inhibitors was observed among the18 carbapenemase non-producers rending an overall specificity of 100% for the two combination disk tests studied.

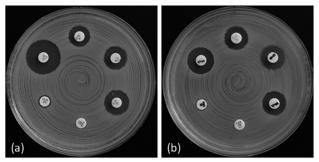


Figure 2: Positive combination disc test for a NDM-producing *K. pneumoniae* isolate (a) using MAST-CDS; (b) using KPC&MBL&OXA-48 disc kit. Clockwise: A/MEM, meropenem; C/MR+BO, meropenem with phenylboronic acid; D/MR+CL, meropenem with cloxacillin; TZP, piperacillin/ tazobactam; TEM/TMO, temocillin; B/MR+DP, meropenem with dipicolinic acid.

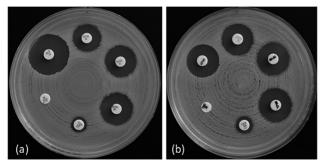


Figure 3: (a) Positive combination disc test for a VIM-producing *S. marcescens* isolate using MAST-CDS; (b) Negative combination disc test for a VIM-producing *S. marcescens* isolate using KPC&MBL&OXA-48 disc kit. Clockwise: A/ MEM, meropenem; C/MR+BO, meropenem with phenylboronic acid; D/MR+CL, meropenem with cloxacillin; TZP, piperacillin/tazobactam; TEM/ TMO, temocillin; B/MR+DP, meropenem with dipicolinic acid.

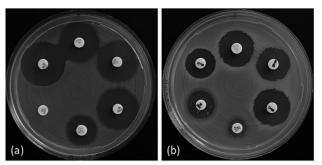


Figure 4: Negative combination disc test for a VIM-producing *P. mirabilis* isolate (a) using MAST-CDS; (b) using KPC&MBL&OXA-48 disc kit. Clockwise: A/MEM, meropenem; C/MR+BO, meropenem with phenylboronic acid; D/MR+CL, meropenem with cloxacillin; TZP, piperacillin/ tazobactam; TEM/TMO, temocillin; B/MR+DP, meropenem with dipicolinic acid.

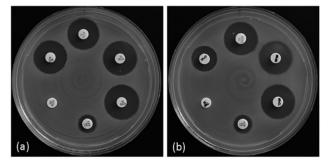


Figure 5: Positive combination disc test for a OXA-48-producing *K. pneumoniae* isolate (a) using MAST-CDS; (b) using KPC&MBL&OXA-48 disc kit. Clockwise: A/MEM, meropenem; C/MR+BO, meropenem with phenylboronic acid; D/MR+CL, meropenem with cloxacillin; TZP, piperacillin/tazobactam; TEM/TMO, temocillin; B/MR+DP, meropenem with dipicolinic acid.

DISCUSSION

Rapid and accurate detection of carbapenemase-producing *Enterobacteriaceae* is essential for patient treatment and infection control management. This necessitates the introduction of a standardised, reliable, and cost effective method for routine detection of carbapenemases in clinical microbiology laboratories. Unfortunately the process is seriously hampered by the low-level resistance to carbapenems, conferred by the corresponding genes, which also facilitates their rapid dissemination. The efforts of many researchers aim at developing efficient algorithms for phenotypic detection and differentiation of carbapenemase-producers.

In our previous study, following the characterisation of the first OXA-48-positive *K. pneumoniae* strain in Bulgaria, we developed an algorithm for routine identification of the difficult-to-detect OXA-48-positive/ESBL-negative phenotype (7, 8). In another study we evaluated the effectiveness of the KPC&MBL&OXA-48 disc kit (Liofilchem, Italy) for detection and differentiation of KPC-producing clinical *K. pneumoniae* strains, isolated in the country. We found that this test, like other commercial standardised tests, identifies KPC-producers with 100% sensitivity and specificity (9).

The current study focused on the comparative performance of two commercially available combination disc tests for confirmation of carbapenemase production in *Enterobacteriaceae* isolated in Bulgaria. In line with studies from other authors (4), KPC-producing *K. pneumoniae*, OXA-48-producing *K. pneumoniae*, and NDM-producing *Enterobacteriaceae* were correctly confirmed with 100% sensitivity and specificity. Unfortunately, our results support another trend concerning the difficulty in detecting IMPand VIM-producers as a whole and particularly in *P. mirabilis* strains (2, 10). The reason for this phenomenon is again indicative of the low-level resistance to carbapenems.

In conclusion, our results indicate that the MAST-CDS combined with temocillin disc and the KPC&MBL&OXA-48 disc kit provide reliable phenotypic confirmation for class A, B, and OXA-48 carbapenemases in non-*Proteus Enterobacteriaceae* species isolated in Bulgaria. For phenotypic detection of VIM-producing *P. mirabilis* strains another method should be considered.

ACKNOWLEDGEMENTS

We thank Mr. Stoyan Yordanov Atanasov for the precise photography of the antibiograms.

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PHENOTYPIC DETECTION OF AAC(6')-IB-CR-PRODUCING ENTEROBACTERIACEAE BY ROUTINE ANTIBIOGRAM

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ABSTRACT

Plasmid-mediated quinolone resistance (PMQR) in Enterobacteriaceae has increased in the last years, seriously compromising the management of life-threatening infections. PMQR genes (qnr, aac(6,)-Ib-cr, gepA and ogxAB) confer low-level quinolone resistance and are frequently co-transmitted with extended-spectrum β-lactamase genes. AAC(6)-lb-cr-mediated aminoglycoside and fluoroquinolone resistance was first detected in 2006, but is now recognised to be widely disseminated. In the present study, we attempted to develop a disc-based method to screen for AAC(6)-lb-cr producers, based on the peculiar substrate profile of the AAC(6')-lbcr enzyme. We analysed resistance phenotypes of 52 well-characterised aac(6,)-lb-cr-positive Enterobacteriaceae, collected at a cancer hospital. Our results show that resistance to tobramycin along with reduced susceptibility to amikacin and ciprofloxacin, but susceptibility to gentamicin, nalidixic acid and levofloxacin provide a reliable screening for AAC(6')-Ib-cr producion in Enterobacteriaceae. The proposed antibiotic disc configuration reflects the substrate profile of AAC(6')-lb-cr enzyme and could be useful for early detection of AAC(6)-lb-cr-positive/ quinolone mutation-negative isolates. We conclude that the developed disc-based method is a simple cost-effective screening tool for rapid detection and appropriate treatment of AAC(6')-Ib-cr-producing multidrug-resistant enterobacterial pathogens.

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INTRODUCTION

Plasmid-mediated quinolone resistance (PMQR) is an emerging healthcare concern, especially in *Enterobacteriaceae*, involving mechanisms of target protection (*qnr*), antbiotic inactivation (*aac*(6,)-*lb*-*cr*) or active efflux systems (*qepA* and *oqxAB*) (1). The PMQR determinants confer only low-level quinolone resistance, but the *aac*(6')-*lb*-*cr* carriage could facilitate the selection of higher level chromosomal resistance mutations when quinolones are used (2). To date, no specific phenotypic tests for detection of PMQR mechanisms are available.

In the present study, we attempted to develop a disc-based method to screen for

AAC(6)-Ib-cr producers, based on the peculiar substrate profile of the AAC(6')-Ib-cr enzyme.

MATERIALS AND METHODS

A total of 52 isolates of *Escherichia coli* (n=47), Citrobacter freundii (n=2), Klebsiella pneumoniae (n=1), Enterobacter aerogenes (n=1) and Morganella morganii (n=1), already confirmed to harbour the aac(6')-lb-cr gene by PCR and sequencing were studied (3, 4). The isolates were tested using disc diffusion according to European Committee on Antimicrobial Susceptibility Testing (EUCAST) recommendations (5) with ciprofloxacin 5 µg, norfloxacin 10 µg, levofloxacin 5 μ g, nalidixic acid 30 μ g, amikacin 30 μ g, tobramycin 10 µg, netilmicin 10 µg, kanamycin 30 µg and gentamicin 10 µg discs (Becton Dickinson, Sparks, MD). The results were interpreted according to EUCAST breakpoints (6), except for nalidixic acid and kanamycin zone diameters that were interpreted using Clinical and Laboratory Standards Institute (CLSI) breakpoints (7). E. coli ATCC 25922 was used as antibiotic-susceptible control.

RESULTS AND DISCUSSION

The common aminoglycoside acetyltransferase gene *aac(6')-lb* confers resistance to kanamycin, tobramycin, netilmicin and amikacin but not to gentamicin (2). According to EUCAST expert rules v. 2.0, rule no. 12.7., production of acquired AAC(6')-l-like enzyme confers phenotypic resistance to tobramycin, kanamycin, netilmicin, but may not confer phenotypic amikacin resistance despite modification of amikacin (8). Indeed, our susceptibility testing results showed that all isolates were resistant to tobramycin, kanamycin, netilmicin, but 75% (39/52) of the isolates exhibited only reduced susceptibility to amikacin as shown in Figure 1. In addition, the *cr* variant of the common acetyltransferase gene - *aac*(6')-*Ib*-*cr* confers low-level resistance to ciprofloxacin and norfloxacin but not to nalidixic acid and the other fluoroquinolones like levofloxacin, as indicated in Figure 2.

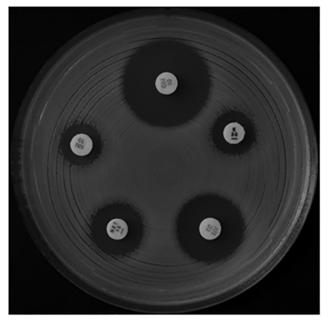


Figure 1: Typical image of aminoglycoside resistance phenotype due to AAC(6')-Ib production observed in clinical *Enterobacteriaceae* isolated at the cancer hospital. Clockwise: GM, gentamicin; K, kanamycin; AN, amikacin; NET, netilmicin; NN, tobramycin. The isolates exhibited resistance to tobramycin, kanamycin and netilmicin, reduced susceptibility to amikacin, and were susceptible to gentamicin.

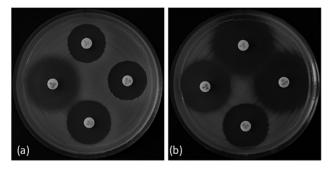


Figure 2: (a) Typical image of resistance phenotype due to AAC(6')-Ib-cr enzyme concerning quinolones. Clockwise: CIP, ciprofloxacin; NOR, norfloxacin; NA, nalidixic acid; LVX, levofloxacin. The isolate exhibited reduced susceptibility to both ciprofloxacin and norfloxacin, but remained susceptible to nalidixic acid and levofloxacin. (b) Typical image of quinolone "susceptible" resistance phenotype, shown for comparison.

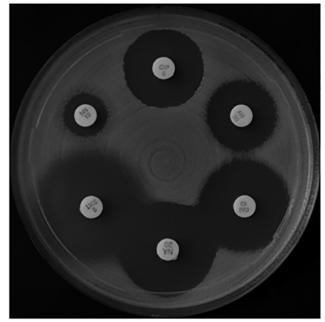


Figure 3: Antibiogram of *aac(6')lb-cr*-positive *E. aerogenes* strain. Clockwise: CIP, ciprofloxacin; AN, amikacin; GM, gentamicin; NA, nalidixic acid; LVX, levofloxacin;

NOR, norfloxacin; NN, tobramycin. Resistance to tobramycin along with reduced susceptibility to amikacin and ciprofloxacin, but susceptibility to gentamicin, nalidixic acid and levofloxacin are indicative of AAC(6')-Ib-cr production.

Based on analysis of the observed resistance phenotypes and their corresponding genotypes we selected tobramycin, amikacin, ciprofloxacin as antibiotic disc indicators along with gentamicin, nalidixic acid, levofloxacin as anti-indicators able to screen for AAC(6')-Ib-cr producion (Fig.3) Our results show that resistance to tobramycin along with reduced susceptibility to amikacin and ciprofloxacin, but susceptibility to gentamicin, nalidixic acid and levofloxacin provide a reliable screening for AAC(6')-Ib-cr producion in Enterobacteriaceae. The proposed antibiotic disc configuration reflects the substrate profile of the AAC(6')-lbcr enzyme and could be useful for early detection of AAC(6)-Ib-cr-positive/quinolone mutation-negative isolates.

We conclude that the developed disc-based method is a simple cost-effective screening tool for rapid detection and appropriate treatment of AAC(6')-Ib-cr-producing multidrug-resistant enterobacterial pathogens.

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COMPARISON OF EUROPEAN COMMITTEE ON ANTIMICROBIAL SUSCEPTIBILITY TESTING AND CLINICAL LABORATORY STANDARDS INSTITUTE CRITERIA FOR THE INTERPRETATION OF EXTENDED-SPECTRUM B-LACTAMASE-PRODUCING ENTEROBACTERIACEAE ISOLATED AT A CANCER HOSPITAL

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ABSTRACT

Background: In 2010 European Committee on Antimicrobial Susceptibility Testing (EUCAST) and Clinical and Laboratory Standards Institute (CLSI) eliminated the need for routine extended-spectrum β-lactamase (ESBL) detection and reporting β-lactam susceptibility testing results accordingly. The aim of this study was to determine how many ESBL-producing Enterobacteriaceae test susceptible using EUCAST and CLSI clinical breakpoints. Material and methods: One hundred consecutive Enterobacteriaceae with ESBL phenotype were collected at the Specialised hospital for active treatment in oncology between August 2015 and June 2016. ESBL production was systematically determined on the basis of observation of a synergy zone between

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Stefana Sabtcheva Laboratory for Clinical Microbiology Specialised Hospital for Active Treatment in Oncology 6 Plovdivsko pole Str 1756 Sofia, Bulgaria Tel. 8076293 E-mail: stefanasabtcheva@gmail.com amoxicillin/clavulanic acid and cefepime and/or ceftazidime and/or cefotaxime as described in the EUCAST guidelines for detection of resistance mechanisms. The collection comprised 40 Escherichia coli, 25 Klebsiella pneumoniae, 12 Enterobacter cloacae, 12 Serratia marcescens, 7 Klebsiella oxytoca, 2 Citrobacter freundii, and single isolates of Enterobacter aerogenes and Morganella morganii. Disc diffusion antimicrobial susceptibility testing was performed and interpreted according to EUCAST 2015 and CLSI 2015 guidelines. Results: All 100 ESBL-producing isolates were non-susceptible to ceftriaxone and cefotaxime if EUCAST and CLSI zone diameter clinical breakpoints were applied, whereas 6% of all (5/40 E. coli and 1/25 K. pneumoniae) were susceptible to cefepime according to both breakpoints. However, significant differences in the susceptibility rates of ceftazidime and aztreonam were demonstrated applying both guidelines. According to EUCAST 2015 breakpoints 4% of all ESBL-producers (3/40 E. coli and 1/25 K. pneumoniae) were susceptible to ceftazidime and aztreonam, whereas, according to CLSI 2015 clinical breakpoints 22% of all (18/40 E. coli, 2/12 S. marcescens, 1/25 K. pneumoniae, 1/1 M. morganii) were susceptible to ceftazidime and 19% (17/40 E. coli, 1/25 K. pneumoniae, 1/1 M. morganii) were susceptible to aztreonam. Conclusions: All ESBL-producing Enterobacteriaceae isolated at the cancer hospital would be reported to be non-susceptible to cefotaxime and ceftriaxone applying the EUCAST and CLSI 2015 clinical breakpoints, but a substantial number of ESBL-producing E. coli strains would be reported to be susceptible to ceftazidime and aztreonam, according to CLSI 2015 clinical breakpoints. Further harmonisation is needed to cease the controversial reporting of ceftazidime and aztreonam in therapy recommendations when using the EUCAST and CLSI guidelines.

Keywords: *Enterobacteriaceae*, extended-spectrum β-lactamase, interpretive criteria

INTRODUCTION

Enterobacteriaceae, especially those producing extended-spectrum β -lactamases (ESBLs), can cause various nosocomially acquired life-threatening infections (1). Until 2009, antimicrobial susceptibility testing (AST) guidelines recommended routine ESBL detection and reporting *in vitro* susceptible and intermediate AST results for penicillins, cephalosporins, and monobactams in ESBL-producing isolates as resistant (2, 3). In 2010 European Committee on Antimicrobial Susceptibility Testing (EUCAST) and Clinical and Laboratory Standards Institute (CLSI) eliminated the need for routine ESBL testing and allowed the treatment of ESBL-producing isolates with cephalosporins which tested susceptible according to the revised versions of the EUCAST and CLSI guidelines.

The aim of this study was to determine how many recent ESBL-producing clinical *Enterobacteriaceae* isolates test susceptible using the current EUCAST and CLSI clinical breakpoints.

MATERIALS AND METHODS

The study was carried out at the Specialised hospital for active treatment in oncology, Laboratory for Clinical Microbiology between August 2015 and June 2016. During this period. 100 consecutive clinical Enterobacteriaceae isolates with ESBL phenotype were collected. The ESBL production was routinely determined on the basis of observation of a synergy zone between amoxicillin/clavulanic acid and cefepime and/or ceftazidime and/or cefotaxime as described in the EUCAST guidelines for detection of resistance mechanisms (4). Identification to the species level was determined by Vitek-2 (bioMérieux, Marcy l'Etoile, France). Disc diffusion antimicrobial susceptibility testing was performed on Mueller-Hinton II agar plates (Becton Dickinson, Sparks, MD, USA) using overnight cultures with a turbidity equivalent to 0.5 McFarland standard followed by incubation at 35°C for 18h. Antibiotic discs supplied by Becton Dickinson were used. Results were interpreted according to the EU-CAST 2015 and CLSI 2015 guidelines (5, 6).

RESULTS

The 100 ESBL-producing isolates were recovered from the following specimens: wound (n=56), urine (n=34), respiratory fluid (n=6), and peritoneal fluid (n=4). ESBLs were found in eight different species. The major ESBL-producers were Escherichia coli (40%) and Klebsiella pneumoniae (25%), followed by Enterobacter cloacae (12%), Serratia marcescens (12%), Klebsiella oxytoca (7%), Citrobacter freundii (2%), and single isolates of Enterobacter aerogenes and Morganella morganii. In Table 1 are summarised the comparative inhibition zone interpretation results of cephalosporins and aztreonam for 100 ESBL-producers according to EUCAST 2015 and CLSI 2015 breakpoints. Equal patterns of susceptibility to ceftriaxone and cefotaxime were detected applying the EUCAST 2015 and CLSI 2015 guidelines. None of the ESBL-producers were interpreted as "susceptible". Equal rates of susceptibility (6% of all) were demonstrated for cefepime applying both guidelines. Cefepime "susceptible" isolates comprised 5/40 E. coli and 1/25 K. pneumoniae, but none of the chromosomal AmpC Enterobacteriaceae. However, ESBL-producing isolates had significantly different susceptibility rates to ceftazidime and aztreonam when comparing EU-CAST 2015 with CLSI 2015 breakpoints (4% versus 22% ceftazidime "susceptible", and 4% versus 19% aztreonam "susceptible", respectively). In detail, 3/40 E. coli and 1/25 K. pneumoniae tested susceptible for ceftazidime and aztreonam according to EUCAST interpretive criteria, whereas 18/40 E, coli, 2/12 S. marcescens, 1/25 K. pneumoniae, and 1/1 M. morganii were susceptible to ceftazidime and 17/40 E. coli, 1/25 K. pneumoniae, 1/1 M. morganii were susceptible to aztreonam according to CLSI 2015 clinical breakpoints. Of note, the discrepancy between EUCAST and CLSI susceptibility testing results were observed mainly among ESBL-producing E. coli isolates.

DISCUSSION

In 2010 EUCAST and CLSI eliminated the need for routine ESBL detection and reporting β -lactam susceptibility testing results accordingly. The treatment of ESBL-producing isolates with cephalosporins and aztreonam is allowed depending on the individual susceptibility testing categorisation. However, few data are available showing antibiotic susceptibility patterns for defined populations of ESBL-producing isolates according to revised EUCAST and CLSI guidelines (7, 8).

This study describes various resistance patterns for ESBL phenotype/species combinations if EUCAST 2015 and CLSI 2015 zone diameter breakpoints are applied (Table 1). Lack of susceptibility to cefotaxime and ceftriaxone was always found for all ESBL-producing isolates, whereas ceftazidime was categorised as susceptible in 4% and 22%, respectively, if EUCAST 2015 and CLSI 2015 breakpoints were applied. Similarly, aztreonam was categorised as susceptible in 4% and 19%, respectively, according to EUCAST 2015 and CLSI 2015 breakpoints, whereas cefepime was categorised as susceptible only in 6% of ESBL-producers if both 2015 guidelines were applied. Our results are in concordance with other studies and reflect the current worldwide dominance of CTX-M-type enzymes (8, 9, 10). The variation in susceptibility patterns of cef-

		EUCAST 2015			CLSI 2015		
Drug/ interpretation (%)		<i>E.coli</i> n=40	<i>Klebsiella</i> spp. n=32	Chromosomal AmpC <i>Enterobacteriaceae</i> ^a n=28	<i>E.coli</i> n=40	<i>Klebsiella</i> spp. n=32	Chromosomal AmpC <i>Enterobacteriaceae</i> ª n=28
Ceftriaxone	R	38	32	28	38	32	28
	I	2	0	0	2	0	0
	S	0	0	0	0	0	0
Cefotaxime ^b	R	38	32	28	38	32	28
	I	2	0	0	2	0	0
	S	0	0	0	0	0	0
Ceftazidime⁵	R	32	31	25	20	26	25
	I	5	0	3	2	5	0
	S	3	1	0	18	1	3
Cefepime	R	25	31	26	24	31	24
	I	10	0	2	11	0	4
	S	5	1	0	5	1	0
Aztreonam	R	23	31	27	22	31	25
	I	14	0	1	1	0	2
	S	3	1	0	17	1	1

Table 1. Comparison of antibiotic susceptibility profiles of 100 ESBL-producing clinical *Enterobacteriaceae* isolates according to the EUCAST 2015 and CLSI 2015 breakpoints.

R, resistant; I, intermediate; S, susceptible

^a Isolates include: 12 *E. cloacae*, 12 *S. marcescens*, 2 *C. freundii*, 1 *E. aerogenes*, and 1 *M. Morganii*. ^b For these drugs, EUCAST and CLSI use different disc contents: cefotaxime (EUCAST 5 μg/disc, CLSI 30 μg/disc), ceftazidime (EUCAST 10 μg/disc, CLSI 30 μg/disc). Isolates were tested with both disc contents in parallel and interpretation was carried out accordingly.

tazidime could be attributed to the lower antibiotic disc content recommended by EUCAST while retaining almost the same breakpoint values as those in CLSI. Differences in aztreonam categorisation presumably arise due to setting the resistance breakpoint in EUCAST at a value corresponding to the susceptibility breakpoint in CLSI.

A limitation of this study was the local origin of the clinical strains and the absence of ESBL genotype determination. However, even suggested from the phenotype, our results underline that different ESBL/species combinations produce distinct antibiotic susceptibility patterns. Thus, further studies are needed to characterise the relationship between ESBL production, antimicrobial susceptibility and clinical outcome.

In conclusion, all ESBL-producing *Enterobacteriaceae* isolated at the cancer hospital would be reported to be non-susceptible to cefotaxime and ceftriaxone applying the EUCAST and CLSI 2015 clinical breakpoints, but a substantial number of ESBL-producing *E. coli* strains would be reported to be susceptible to ceftazidime and aztreonam, according to CLSI 2015 clinical breakpoints. Further harmonisation is needed to cease the controversial reporting of ceftazidime and aztreonam in therapy recommendations when using the EUCAST and CLSI guidelines.

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SUCCESSFUL **TREATMENT OF PERITONITIS CAUSED BY GLYCOPEPTIDE-**RESISTANT **ENTEROCOCCUS** FAECIUM, AND **EXTENDED-SPECTRUM B-LACTAMASE-**PRODUCING ESCHERICHIA COLI. **ENTEROBACTER** CLOACAE, KLEBSIELLA PNEUMONIAE, AND **KLEBSIELLA OXYTOCA. CASE REPORT**

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ABSTRACT

Background: Considering the high burden of morbidity and mortality, postoperative peritonitis is a feared complication of intra-abdominal surgery. Typically this is a polymicrobial infection and the most common isolates include aerobic Gram-negative bacilli, anaerobes, and enterococci. Here, we describe a case of postoperative complication involving multidrug-resistant pathogens in a cancer patient. **Case presentation:** 79-year-old male patient diagnosed with

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Laboratory for Clinical Microbiology, Specialised hospital for active treatment in oncology, 6 Plovdivsko Pole Str, Sofia 1756, Bulgaria e-mail: bojana_e@abv.bg cancer of the bladder and prostate was admitted to the urology ward at a 242-bed oncology hospital in Sofia, Bulgaria, for radical cystoprostatectomy. Due to small bowel adhesion to the pelvis and ileus causing peritonitis, two re-operations ensued. Empirical therapy was initiated with piperacillin/tazobactam, teicoplanin, and metronidazole. The first positive abdominal drainage fluid cultures yielded ESBL-producing Klebsiella pneumoniae and K. oxytoca followed by isolation of glycopeptide-resistant Enterococcus faecium, and ESBL-producing Escherichia coli, and Enterobacter cloacae. Piperacillin/tazobactam was replaced with administration of meropenem and linezolid, together with metronidazole until clinical improvement and negative cultures. Conclusion: We report a clinical case of successful treatment of peritonitis caused by glycopeptide-resistant E. faecium, and ESBL-producing E. coli, E. cloacae, K. pneumoniae, and K. oxytoca. Revealing the evolution of underlying resistance mechanisms and changes in microbial etiology during the course of therapy will improve the management of patients undergoing surgical interventions and suffering from comorbidities.

Key words: peritonitis, glycopeptide-resistant *E. faecium*, ESBL-producing *Enterobacteriaceae*

INTRODUCTION

Clinical presentations of intra-abdominal infections can range from localised peritonitis to diffuse inflammation of the abdominal cavity. Bowel injuries, such as perforation, strangulation, or infection are a common cause of intraperitoneal infections manifested as peritonitis (1). With its high burden of morbidity and mortality, postoperative peritonitis is a feared complication of intra-abdominal surgery. Typically this is a polymicrobial infection and the most common isolates include aerobic Gram-negative bacilli, anaerobes, and enterococci (2, 3). Empirical antimicrobial therapy relies on broad-spectrum antibiotics until culture results are obtained. Precise performance of antibiotic susceptibility assays determines the de-escalation or escalation of antibiotics (1, 4). Our aim was to present a clinical case of successful treatment of peritonitis caused by glycopeptide-resistant Enterococcus faecium, and extended-spectrum β-lactamase-producing Escherichia coli, Enterobacter cloacae, Klebsiella pneumoniae, and Klebsiella oxytoca in a cancer patient in parallel with the changing microbial etiology in the course of therapy.

CASE PRESENTATION

79-year-old male patient diagnosed with cancer of the bladder and prostate was admitted to the urology ward at a 242-bed oncology hospital in Sofia, Bulgaria, for radical cystoprostatectomy. The patient had a history of undergoing left nephrectomy and three assigned surgeries for the bladder cancer in other healthcare facilities. The radical cystoprostatectomy was performed by laparotomy along with lymph node dissection followed by difficult postoperative period. On the 6th day after the cystoprostatectomy the patient was re-operated due to complications arising from small bowel adhesion to the pelvis and ileus causing peritonitis. Six days later laparotomy was performed for the third time with anterior abdominal wall reconstruction and creation of ileostomy. In accordance with the hospital guidelines, a combination of ampicillin/sulbactam and metronidazole was administered as surgical prophylaxis during the first laparotomy. Along with the development of complications, empirical therapy was initiated with piperacillin/tazobactam, teicoplanin, and metronidazole. Two days after the second laparotomy abdominal drainage fluid cultures yielded K. pneumoniae and K. oxvtoca isolates with multidrug-resistant phenotype. Species identification of isolates was determined by Vitek-2 (bioMérieux, Marcy l'Etoile, France). Susceptibility to antimicrobials was determined by the disc diffusion method following the guidelines of the European Committee on Antimicrobial Susceptibility Testing (EUCAST) (5). Results were interpreted according to EUCAST breakpoints (6). Suggestive evidence of Extended-spectrum β-lactamase (ESBL) production was revealed by the double-disk synergy test (7). The Klebsiella strains were categorised as ESBL producers and susceptible to piperacillin/ tazobactam. Abdominal drainage fluid cultures and peritoneal fluid collected on the day of the third laparotomy, yielded glycopeptide-resistant E. faecium, E.coli, and E. cloacae besides the already detected Klebsiella isolates. E. coli and E. cloacae were also ESBL-positive. Identical in susceptibility profile E. faecium strain was also recovered from stool and urine cultures. Stool sample culture yielded also significantly high Candida albicans counts. After obtaining the antimicrobial susceptibility results, piperacillin/tazobactam was replaced with administration of meropenem, linezolid, and fluconazole together with metronidazole until clinical improvement and negative cultures. As a usual

consequence of a prolonged treatment with carbapenem antibiotics, sixteen days after the third laparotomy abdominal drainage fluid and urine cultures were positive for *Stenotrophomonas maltophilia*, treated subsequently with levofloxacin. The patient was discharged on hospital day 38 in satisfactory general condition.

DISCUSSION

Carcinoma of the bladder and prostate ranks among the most common cancers in men and its incidence increases with age. A large percentage of new cases are registered in individuals aged 75–84 years. Radical cystoprostatectomy is a complex surgical procedure and a standard treatment but it is associated with complications, morbidity, prolonged recovery time, and extended hospital stays (8, 9). Further entanglement arises from the fact that patients with this disease are typically older and often have a history of comorbidities. Therefore treatment of elderly patients with urinary bladder cancer and prostate cancer poses challenges to urologists (9).

Enterococci and members of the Enterobacteriaceae are part of the normal intestinal flora in humans, but they are also involved in urinary infections, bacteraemia, intra-abdominal, and nosocomial infections. Glycopeptide-resistant enterococci present a serious problem in hospital settings with very limited treatment options. In Bulgaria the first vancomycin-resistant enterococcal infections were reported in 2005 from the University Hospital in Stara Zagora (10). Cases of glycopeptide-resistant enterococci causing peritonitis successfully treated with linezolid and achieving eradication of the infection were reported in studies from Korea (11, 12). Polymicrobial infections involving ESBL-producers are well known as further impeding the antimicrobial treatment. Previous reports from Bulgarian hospitals identify different families of ESBL enzymes in Klebsiella and other members of the Enterobacteriaceae, with CTX-M-like and SHV-like often found as the most prevalent (13, 14).

In conclusion we report a clinical case of successful treatment of peritonitis caused by glycopeptide-resistant *E. faecium*, and ESBL-producing *E. coli*, *E. cloacae*, *K. pneumoniae*, and *K. oxytoca* in a cancer patient. Investigating the evolution of microbial etiology in the course of therapy and the underlying mechanisms of resistance will improve the management of patients undergoing surgical interventions and suffering from comorbidities.

SUCCESSFUL TREATMENT OF PERITONITIS CAUSED BY GLYCOPEPTIDE-RESISTANT ENTEROCOCCUS FAECIUM...

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MULTIPLE VIRAL PATHOGENS CAUSE "UNDIFFERENTIATED" CNS INFECTIONS

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ABSTRACT

In an attempt to elucidate etiology, we tested 110 serum samples from patients with "undifferentiated" viral CNS infections for antibodies against herpes simplex virus (HSV), varicella zoster virus (VZV), Epstein-Barr virus (EBV), Coxsackie B viruses (CVB), influenza viruses, parainfluenza viruses, tick-borne encephalitis virus (TBE), and West Nile fever virus (WNV). A total of 21 (19.1%) of the samples were reactive showing presence of IgM antibodies against at least one of the tested viruses. The most common reactivity was found against either EBV, Coxsackie B viruses or influenza viruses - 11 (10%) of the samples reacted with one of the three viruses, followed by HSV-2, TBEV, VZV, and parainfluenza viruses - 6 (5.5%) of the samples were reactive to one of the latter four viruses. Cases of HSV-1 and WNV infections were not detected. The vast majority of the annually recorded viral meningitis and encephalitis cases in Bulgaria are diagnosed only on clinical manifestations and the diagnosis is usually not supported by detection of virus species as the etiological agent. In this study, we succeeded to identify a possible viral etiological agent in one fifth of the cases, showing that investigation for multiple virus species should be pursued in an at-

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Iva Christova, Head, National reference laboratory of vector-borne infections, leptospirosis and listeriosis, NCIPD, 26 Yanko Sakazov blvd, Sofia 1504, Bulgaria tempt to confirm the viral etiology of the CNS infection.

Key words: meningitis, CSF, enteroviruses, influenza viruses, TBE

INTRODUCTION

Hundreds of "undifferentiated" viral meningitis and encephalitis cases are reported every year to the Bulgarian Ministry of Health. The term encompasses all officially registered CNS infections with clinical manifestations suggesting viral etiology but with no causative agent detected.

In an attempt to elucidate the missing etiology, a collection of serum samples from patients with "undifferentiated" viral CNS infections was tested for the presence of antibodies against herpes simplex virus (HSV), varicella zoster virus (VZV), Epstein-Barr virus (EBV), Coxsackie B viruses (CVB), influenza viruses, parainfluenza viruses, tick-borne encephalitis virus (TBE), and West Nile fever virus (WNV). Usually, herpes simplex virus infection is the most common cause of infectious encephalitis in humans (1). HSV-1 causes up to 90% of HSV encephalitis cases. HSV-2 infection is a rare one and is implicated mostly in meningitis cases. On the other hand, recurrence of meningitis is typical for HSV-2 infection.

Varicella zoster virus (VZV) is a neurotropic virus with life-long latency in sensory ganglia. VZV can cause encephalitis, myelitis, and acute myeloradiculitis. In immunocompromised patients, VZV can induce multifocal encephalitis and necrotising myelitis.

Epstein-Barr virus (EBV) causes infectious mononucleosis. Less than 5% of patients develop meningitis, encephalitis or polyradiculomyelitis before, during or after suffering infectious mononucleosis.

All members of the *Herpesviridae* family are double-stranded DNA viruses. Herpesviruses are divided into 3 subfamilies. Alpha-herpesviruses include HSV-1, HSV-2, and VZV. Typically, they have short reproductive cycle and establish latency in sensory ganglion neurons. Beta-herpesviruses include human cytomegalovirus and the roseola viruses HHV-6 and HHV-7. They replicate relatively slowly and establish latency in multiple cell types and organs. Gamma-herpesviruses represented by EBV and HHV-8, are oncogenic. They also replicate relatively slowly but establish latency in restricted cell types such as lymphocytes. All herpesviruses establish a life-long latency and reactivate intermittently.

Enteroviruses cause more than 90% of viral meningitis cases and only occasionally are implicated as the cause of encephalitis (2). They affect mainly children under 10 years of age. The Enterovirus genus belongs to the Picornaviridae family which consists of small, non-enveloped single-stranded RNA viruses. The multitudinous genus of enteroviruses includes 7 species that are pathogenic to humans: enterovirus A, B, C, and D and rhinovirus A, B, and C. Among them, enteroviruses A, B, C, D are those that can cause meningitis, myelitis, encephalomyelitis, Guillen-Barre syndrome, etc. The notorious EV-71 as well as several coxsackie A viruses, are included in species EV-A: Coxsackie B viruses. Coxsackie virus A9, and echoviruses belong to species EV-B; species EV-C includes polioviruses 1-3, several Coxsackie A viruses and several newer enteroviruses assigned a number; and species EV-D includes 5 serotypes, among them EV-68 (2). Enteroviral meningitis is caused primarily by echoviruses and Coxsackie B viruses. Enterovirus infection should be suspected in children with neurologic symptoms.

Besides respiratory complications, influenza viruses can also cause infections of the CNS. Development of CNS diseases is the most common extra-respiratory complication of infection with influenza A viruses. Meningitis, encephalitis, encephalopathies, myelitis can be caused by influenza A viruses. Influenza viruses belong to the Orthomyxoviridae family of enveloped negative-strand RNA viruses. Influenza A viruses are subtyped based on their surface glycoproteins, hemagglutinin, and neuraminidase. Seasonal, pandemic, and zoonotic influenza A viruses affect humans. Seasonal influenza A viruses cause yearly epidemics (H3N2, H1N1). Pandemic influenza viruses are a result of cross-species transmission and subsequent efficient transmission among humans (H1N1, H2N2). Zoonotic influenza virus infections are a result of zoonotic transmission with inefficient transmission among people (H7N9, H5N1).

Some viruses of the *Paramyxoviridae* family can also cause a wide range of neurological manifestations ranging from acute encephalitis to severe long-term infections. These include measles, mumps, Hendra, and Nipah viruses. Viruses in this virus family are enveloped and contain single-stranded negative-sense RNA. Some flaviviruses typically cause infections of

CNS. Tick-borne encephalitis virus and West Nile virus, even rare in this country, should also be considered. The Flavivirus genus of the Flaviviridae family includes arboviruses that cause encephalitis. Flaviviruses are single-stranded positive-sense RNA viruses. TBE viruses are classified into three subtypes -European, Siberian, and Far-Eastern subtype. TBE virus is transmitted by Ixodes ricinus ticks. Neurological manifestations of TBE include meningitis (50%), meningoencephalitis (40%) or meningoencephalomyelitis (10%) (3). West Nile virus is subdivided in seven genetic lineages, of which lineages 1 and 2 cause the epidemics. WNV is transmitted by Culex mosquitoes in an enzootic cycle between birds. Small outbreaks of WNV infections appear yearly in countries neighbouring Bulgaria and the first WNV cases were described in Bulgaria last year. It is estimated that 1 in 140 infected people develops meningoencephalitis, ranging from 1 in 50 in patients over 65 years to 1 in 300 in patients younger than 65 years.

Viruses belonging to all of the above described families can cause infections of the CNS and are likely to be found as etiological agents of "undifferentiated" viral meningitis and encephalitis cases. In the present study, serum samples from such patients were tested by ELI-SA to detect specific IgM antibodies against possible viral etiological agents in order to establish the extent to which each of these virus species could contribute to the etiology of the observed CNS infection.

MATERIALS AND METHODS Patients

A total of 110 patients with clinically diagnosed viral meningitis (n=78), encephalitis (n=23) or meningoencephalitis (n=9) of unknown etiology were enrolled in the study. Serum samples from the patients were collected in The Clinical Microbiology Laboratory of University Hospital "Tsaritsa Yoanna – ISUL" (n=41), The National Reference Laboratory of Vector-Borne Infections (n=49), and The National Reference Laboratory of Enteroviruses (n=20). The serum samples were collected in the period between 2012 and 2015. The age of the patients varied between 3 and 88, mean 42.

ELISAs

All 110 serum samples from the patients were tested for antibodies against 9 different virus species using commercially available ELISA tests (4). Test procedures and interpretation of results were carried out according to manufacturer's instructions. The serum samples were tested for infection with four herpesviruses - HSV-1, HSV-2, VZV, and EBV, infection with Coxsackie B viruses representing one of the enteroviruses, namely EV-B, for infection with two respiratory viruses - influenza A and parainfluenza viruses, and for two flavi arboviruses - TBEV and WNV. The ELISA tests used in the survey included: Anti-HSV-1 IgM ELISA with antigens of glycoprotein C1 (Euroimmun, Germany); Anti-HSV-2 IgM ELISA with antigens of glycoprotein G2 (Euroimmun, Germany); Anti-VZV Glycoprotein IgM ELI-SA (Euroimmun, Germany); Anti-EBV Capsid antigens IgM ELISA (Euroimmun, Germany); Anti-Coxsackie B virus IgM ELISA (Euroimmun, Germany); Anti-Influenza A and B virus pool IgM ELISA (Euroimmun, Germany); Anti-Parainfluenza viruses pool of types 1 to 4 IgM ELISA (Euroimmun, Germany); Anti-TBE Virus IgM ELISA (Euroimmun, Germany); and Anti-West Nile Virus IgM ELISA (Euroimmun, Germany). In addition, IgM reactive sera for TBE and WNV were tested for IgG antibodies.

RESULTS

Serum samples (n=110) collected within 4 years between 2012 and 2015 from patients with undifferentiated viral meningitis, encephalitis or meningoencephalitis were tested by ELISAs (one sample per patient) for infection with 9 virus species known to cause acute CNS infections.

A total of 21 (19.1%) of the samples were reactive showing presence of IgM antibodies against at least one of the tested viruses. Of them, 17 (15.5%) samples were clearly IgM positive and other 4 were IgM borderline (3.6%). One of the samples showed reactivity against 3 viruses (positive for VZV and WNV,

Virus infections tested	Number of positive samples	Number of borderline samples	Total number of reactive samples (%)
HSV-1	0	0	0
HSV-2	1	1	2 (1.8%)
VZV	1	0	1 (0.9%)
EBV	4	0	4 (3.6)
Coxsackie B viruses	2	1	3 (2.7%)
Influenza virus	3	1	4 (3.6)
Parainfluenza virus	1	0	1 (0.9%)
ТВЕ	3	0	3 (2.7%)
WNV	4	2	6 (5.5%)
TOTAL	19	5	24* (21.8%)

Table 1. Specific IgM antibodies found in serum samples from patients with acute undifferentiated viral CNS infections against 9 different virus species known to cause neurological infections.

* Sum of positively reacted serum samples exceeds the actual number of positive samples because two of the serum samples have reacted with multiple viral pathogens. and borderline for HSV-2) and another one showed reactivity to two of the tested viruses (EBV and Influenza virus). Summarised results are presented in Table 1. The sum of positively reacted serum samples exceeds the actual number of positive samples because two of the serum samples have reacted with multiple viral antigens.

In addition, all samples that were found IgM positive for TBE or WNV were tested for IgG antibodies to prove specificity of the reaction since IgG antibodies appear very early in the course of flavivirus infections. IgG antibodies against TBE virus were confirmed in 2 of the 3 IgM reactive sera. IgG antibodies against WNV were found in none of the 6 IgM reactive serum samples.

The most common reactivity was found against either EBV, Coxsackie B viruses or influenza viruses – 11 (10%) of the samples reacted with one of the three viruses, followed by HSV-2, TBEV, VZV, and parainfluenza viruses – 6 (5.5%) of the samples were reactive to one of these four viruses. Cases of HSV-1 and WNV infections were not detected.

All patients with reactive samples detected, were clinically diagnosed as viral meningitis except the two patients with TBEV infection detected, who were diagnosed as viral encephalitis and meningoencephalitis.

DISCUSSION

The vast majority of annually registered viral meningitis and encephalitis cases in Bulgaria are diagnosed only on clinical manifestations and lack a confirmed etiological agent. In this study, we have succeeded to identify the possible viral etiological agent in one fifth of the cases, revealing that investigation for multiple virus species as the probable cause of the disease should be pursued in an attempt to confirm the viral etiology of the CNS infection concerned.

The present study reveals that investigated "undifferentiated" viral meningitis and encephalitis cases are mostly due to complications of Ebstein-Barr, Coxsackie B, and influenza virus infections. Less common reaction is detected against HSV-2, VZV, and parainfluenza viruses.

Most of the tested serum samples originate from patients with meningitis and this is the most probable reason why HSV-2 is suspected and not HSV-1 which is known to cause encephalitis. Coxsackie B virus infections when symptomatic are mainly associated with meningitis. Development of CNS diseases is the most common extra-respiratory complication of infection with influenza viruses.

One of the serum samples reacted with VZV, HSV-2, and WNV and most probably was infected with VZV because reactivity to HSV-2 was borderline on one hand, and on the other hand, reactivity to WNV was not confirmed by detection of specific IgG antibodies. Another serum sample showed reactivity to both EBV and influenza virus and was probably infected with EBV and cross-reacted with influenza virus.

TBEV infection is limited to certain geographical regions and is a rare one in Bulgaria but still should be considered in differential diagnosis. Detection of IgG antibodies in addition to IgM antibodies against TBEV in 2 of the 3 samples has confirmed the diagnosis and clearly has shown that TBE is not so unusual in this country. The results of the present study also reveal that TBE infections are manifested as encephalitis or meningoencephalitis while the rest of the detected viral infections presented as meningitis.

Although WNV infections are extremely rare in the country, they should also be considered during the corresponding seasons when mosquitoes are active.

Using PCR to test CSF samples of patients with neurological disorders, Kleines et al (5) detected EBV in 1.6%, VZV in 1.3%, HSV in 1.24%, and EV in 0.4% of the patients. In patients with infectious central nervous system disorders, HSV, VZV, and EV prevailed (5).

Similar design of the study, but focused on patients over 65 years, was conducted by Parisi et al (6). They detected viral RNAs in 2.3% of the samples – HSV in 35.4%, EV in 23.1%, EBV in 21.5%, VZV in 18.5%, and CMV in 1.5% of the positives.

When most of the patients are children, enteroviruses considerably prevail. For example, in the study of Akhvlediani et al (7) EV was detected in 26 patients, VZV in 4, and HSV-1 in 2 patients out of 140 patients with infectious CNS disorders tested by multiplex PCR of their CSF samples. In this study, 58% of the patients were children.

The results of our study reveal that herpesviruses, enteroviruses, and influenza viruses are the most common causes of CNS infections. Appropriate testing is needed to properly identify the etiological agent in each case.

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MEDITERRANEAN SPOTTED FEVER (MSF) WITH UNUSUAL PORTAL OF ENTRY – CASE REPORT

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Abstract

Background: Mediterranean spotted fever (or Boutonneuse fever) is re-emerging tick-borne infectious disease, caused by Rickettsia conorii. The trend of disease distribution in Bulgaria is increasing in the last two decades. Pleven region is not endemic and only sporadic cases have been registered. The aim was to report a case of R. conorii infection with unusual portal of entry. Case presentation: 58-years-old woman crushed a tick manually followed by conjunctival exposure of the left eve to the tick's blood. Six days later there was a sudden onset of fever and shivering. Intensive conjunctival hyperaemia, swollen eyelids, and worsened vision ensued. On the 5th day after hospital admission generalised papulous rash developed involving the palms and the soles. The fever persisted, the woman was in a state of nervous excitement with headache and hepatomegaly; there were no neurological signs. Laboratory investigations revealed normocytosis with granulocytosis (WBC 6.9 x 10⁹/L, granulocytes 0.76, respectively), C-reactive protein 16.1 mg/dL, fibrinogen level 5.14 g/L, and normal liver biochemical tests. Later on, the degree of leucocytosis and granulocytosis increased and the C-reactive protein rose up to 90.2 mg/l. Haemoculture was negative and serological investigation was positive for R. conorii. Complex etiologic and supportive treat-

ADDRESS FOR CORRESPONDENCE:

Tsetsa Georgieva Doichinova Clinic of Infectious Diseases Department of Infectious Diseases, Epidemiology, Parasitology, and Tropical Medicine at Faculty of Public Health, Medical University – Pleven, Bulgaria 8a Georgi Kochev str. 5800 Pleven, Bulgaria Phone: +35964886439, mobile +359888729532 e-mail: doichinova@abv.bg ment was administered, including ciprofloxacin, lincomycin, and metronidazole intravenously, eye-drops, and symptomatic drugs. The patient improved and discharged after twelve days of hospital treatment. **Conclusions:** The case is interesting because of the unusual portal of entry. An increased awareness of MSF could eliminate the risk of delayed and incorrect diagnosis and also improve the prognosis.

Key words: *Rickettsia conorii*, Mediterranean spotted fever, intraocular inflammation

Mediterranean spotted fever (MSF) or Boutonneuse fever is an acute systemic re-emerging infectious disease first described in 1910 in Tunisia by Conor and Bruch. It is transmitted to humans by tick bites. The brown dog tick Rhipicephalus sanguineus is the usual vector. The etiologic agent is Rickettsia conorii, a small intracellular organism belonging to the spotted fever group of the Rickettsia family (1). MSF is an endemic disease occurring during spring and summer in Asia, Africa, and Mediterranean countries. MSF is the most common endemic tick-transmissible rickettsiosis in Bulgaria during the last decade and the trend of distribution is rising. Pleven region is not endemic, only sporadic cases have been registered. Our aim was to describe a case of MSF with unusual portal of entry. We used the records in the patient's hospital documentation.

CASE REPORT

Fifty-eight-year-old woman was admitted on 8 May, 2014 to the Clinic of Infectious Diseases at the University Hospital – Pleven. Six days prior to admission she crushed a big tick with fingers followed by exposure of her left eye to the sudden bleeding of tick blood. Immediately, hyperaemia of the conjunctiva appeared followed by swelling of the eyelids and visual worsening. On the next day, fever of 38°C with shivering, weakness, and edema of the cervical region developed. The patient reported a past ischemic brain stroke with left-side hemiparesis and hypertonic disease. She had no contact with pets.

On physical examination, the patient was afebrile, conscious, but in a state of nervous excitement and unstable walking. The skin was without rash. The left eye was with intensive hyperaemia of the conjunctiva, suffusion on the sclera and swollen eyelids. Enlarged cervical and inguinal lymph nodes were found. She presented also with a coated tongue and mild sore throat. There were no pulmonary auscultation findings. The heart rate was normal and the blood pressure was 140/90. The abdomen was painless. Slightly enlarged liver was found without splenomegaly. There were no signs of meningeal irritation.

Laboratory tests revealed erythrocytes 4.1: 3.89: 4.44 x 10¹²/L; haemoglobin 119; 111; 123 g/L; white-blood cell count 6.9; 7.8; 12.9 x 10⁹/L; neutrophils 0.78; 0.80; 0.59; platelet count 225; 240; 506 x 10¹²/L; aspartate aminotransferase 18; 33 IU/L; alanine aminotransferase 13, 35 IU/L; alkaline phosphatase 102 IU/L: gamma-glutamyl transferase 89 IU/L; fibrinogen 5.14 g/L; C-reactive protein 16.1; 90.2; 0.74 mg/L; total protein 64 g/L; albumins 39 g/L; blood urea nitrogen 3.7 mmol/L; creatinine 75 µmol/L; Na⁺ 136; K⁺ 4,0; Cl⁻ 100 mmol/L. Uroculture and haemoculture were negative. Serological investigation by ELISA was positive for MSF: IgM antibodies titer was 27 (reference value <11) and IgG antibodies 4 (reference value < 9).

The patient had consultation with ophthalmologists who recommended a local therapy.

Empirical treatment was initiated intravenously with ceftiaxone but on the fourth hospital day the characteristic generalised maculopapular rash developed, involving the palms and soles. Antimicrobial therapy was changed to ciprofloxacin and metronidazole intravenously, and local ocular treatment was performed.

The patient improved and was discharged after twelve days of hospital treatment. She was afebrile, with resolution of the rash, and the left eye was without hyperaemia and swelling.

Discussion

MSF is known as an acute disease with benign prognosis but the patients' quality of life worsens due to fever and multiple organ disorders. The disease could be life-threatening for the elderly, immune-compromised, and patients with co-morbidity (2).

Biology of the agent, epidemiology, pathogenesis, and immunogenesis are still being studied. The clinical symptoms and organ damage are continuously investigated.

Systemic disorders involve cardiovascular system – myocarditis (3), heart-rhythm disorders (4), phlebitis (5), gastrointestinal tract (granulomatous hepatitis) and bleeding (6), kidneys – tubulointerstitial nephritis (7, 8), nervous system – encephalitis, neuritis (9, 10), arthritis and myositis (11), respiratory system – bronchopulmonary symptoms, exudative pleuritis (12).

The aim of this report was to focus the attention on affecting the sense organs, and in particular, the visual sense. The prevalence of these complications is not high but with various localisations and degree of severity. There are reports on changed visual ability (13), uveitis (14, 15), optic nerve neuritis (16), keratitis (17, 18), and posterior segment damage (19).

Popivanova N (2006) performed the most extended study on MSF in Bulgaria, including variability of the clinical forms and systemic disorders. Her special attention was focused on affecting the sense organs. The author described cases with conjunctivitis and reversible visual disorders. All patients crushed ticks with fingers followed by contamination of the eyes, i.e. the conjunctiva was a portal of entry. Fever and maculopapular rash always presented without eschar ("black spot") (20). In the case we report, the patient also developed the typical rash and fever, "black spot" was absent but the infection was transmitted by direct inoculation of tick blood in the eye. We did not find in the current literature R. conorii entering the human body in a similar mode and this fact makes the reported case unique.

Laboratory investigations revealed the typical changes associated with moderately severe clinical form of MSF – discrete leucocytosis with granulocytosis, normal haemoglobin level, erythrocytes, and platelet number, normal liver biochemical parameters, slightly elevated fibrinogen and C-reactive protein levels (after treatment – normal), normal blood urea nitrogen, creatinine, and electrolytes.

Empirical treatment of the patient was started with ceftriaxone due to unclear diagnosis. Following the appearance of the typical rash, therapy was changed to ciprofloxacin intravenously – a consideration in accordance with the literature (20).

Conclusions: The case is interesting because of the unusual portal of entry. An increased awareness of MSF could eliminate the risk of delayed and incorrect diagnosis and also improve the prognosis.

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CASE REPORT WITH PULMONARY AND NEURAL TUBERCULOSIS

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ABSTRACT

Background: Neurotuberculosis is the most hazardous type of systemic tuberculosis (TB) because of high mortality and possible serious neurological complications. Early diagnosis and prompt treatment are crucial for favourable outcome. Case report: A 36-yearold female patient was admitted to suburban Infectious Diseases Ward with a one-week history of fever, headache, and vomiting. Primary diagnosis was "viral encephalitis". Four days later, she became unconscious and was transported to the Clinic of Infectious Diseases at the University Hospital – Pleven. On the physical examination, there was syndrome of meningeal irritation, depressed tendon reflexes, and a positive Babinski's sign bilaterally. Investigation of cerebrospinal fluid (CSF) revealed increased protein level (3.15 g/L), leucocytes count 80/µL (30% neutrophils and 70% mononuclears), decreased glucose level (0.61 mmol/L). Mycobacterium tuberculosis was confirmed by culture of CSF. CT-scan revealed brain edema and subarachnoid cyst suboccipitally. The first X-ray of the lungs was considered as "negative", the second as "pleuropneumonia". Tuberculostatic (streptomycin, isoniazid, rifampicin, pyrazinamide) and supportive treatment was performed but the patient's condition worsened, oculomotor and abducens nerves were involved, respiratory disorders appeared requiring mechanic ventilation. The patient died on the 20th day after

ADDRESS FOR CORRESPONDENCE:

Tsetsa Georgieva Doichinova Clinic of Infectious Diseases Department of Infectious Diseases, Epidemiology, Parasitology, and Tropical Medicine at Faculty of Public Health, Medical University – Pleven, Bulgaria 8a Georgi Kochev str. 5800 Pleven, Bulgaria Phone: +35964886439, mobile +359888729532 e-mail: doichinova@abv.bg admission. On autopsy, infiltrative-pneumonic TB, fibrinous-purulent pneumonia, and bronchiolitis were found. Subsequently tuberculous meningoencephalitis was found, visualised morphologically by lymphocytic basal meningitis and parenchymal vessel vasculitis with microthrombosis. Cortical and basal multifocal ischemia and pulmonary disorders were the direct cause of the lethal outcome. **Conclusion:** The globally increased incidence of TB and co-existence of extra-neural tuberculosis and neurotuberculosis require diagnostic improvement and specific therapy even in suspected cases.

Key words: tuberculosis, meningoencephalitis, pulmonary tuberculosis, tuberculostatic therapy

INTRODUCTION

The global increase in the incidence of tuberculosis (TB) is a health issue of universal concern. The acquired immunodeficiency syndrome (AIDS) and the problem of multidrug-resistant TB (MDRTB) are among the factors that have contributed to this increase (1, 2, 3). Advanced age, alcoholism, drug abuse, poverty, malnutrition, transmigration, lymphoma, and immunosuppressive medication also contribute to increased susceptibility (4). A primary TB manifests as pulmonary TB, miliary TB, and CNS TB (5, 6). Involvement of the central nervous system (CNS) by TB is the most hazardous type of systemic TB because of its high mortality rate and possible serious neurological complications and sequels. Co-existence of extra-neural tuberculosis is reported among 50% of cases of neurotuberculosis in the literature which may be a clue to the diagnosis of CNS TB (7, 8). A tuberculous infection of CNS can present either as basal exudative meningitis or as localised tuberculoma.

Tuberculous meningitis (TBM) is the most common presentation of CNS TB accounting for 70-80% of cases (4). The granulomatous infection of the leptomeninges is characterised by a thick exudate affecting the basal parts of the brain. Probably, early meningeal exudate arises from rupture into the subarachnoid space of a microscopic granuloma. This lesion is due to haematogenous dissemination of distant tuberculous focus or is a result of rupture of tuberculoma or miliary tubercle. It can become active even years after initial infection (7).

TBM most often presents with fever, headache, decreased level of consciousness, and meningeal signs such as neck stiffness, photophobia, and vomiting (9, 10). The disease usually evolves gradually over two to six weeks. Convulsions are possible during the course of the illness. Involvement of cranial nerves is seen in 17-70% of patients with TBM (4). Cranial nerves II, III, IV, VI, VII, and VIII are most frequently affected either due to ischemia of the nerve or its nucleus, or due to compression from basal exudates. Without treatment, death usually occurs in three to six weeks and with treatment, mortality rates range from 6-27% (7).

We report a case with co-existence of extra-neural tuberculosis (pulmonary TB) and neurotuberculosis (tuberculous meningoencephalitis).

CASE REPORT

Thirty-six-year-old female patient with a oneweek history of fever, headache, and vomiting was admitted to suburban Infectious Diseases Ward with diagnosis "viral encephalitis" after obtained informed consent. On admission, she was without mental alterations and the complaint was strong occipital headache. Four days later, she became disoriented and was transported to the Clinic of Infectious Diseases at the University Hospital because of worsening. Six months prior to admission, she had contact with relatives suffered from TB and three months later she was treated for "pneumonia".

On the physical examination she demonstrated marked stiffness, positive Kernig's sign, depressed patellar and Achilles' reflexes, and positive Babinski's sign. Cranial nerves were intact. On auscultation, decreased breathing without crackles was found with normal heart frequency and blood pressure.

Laboratory investigations revealed white-cells count in blood 11.800/mm³ and neutrophils 76%. Lumbar puncture was performed and CSF investigation revealed protein level 3.15 g/L and leucocytes 80/µL (30% neutrophills and 70% mononuclears). Glucose in CSF was 0.61 mmol/L. Both, direct microscopy and culture of CSF were positive for Mycobacterium tuberculosis. CT scan of the head revealed brain edema and subarachnoid cyst suboccipitally. According to the neurosurgeon this was congenital cyst in cysterna magna and the left cerebellar hemisphere that did not need surgical treatment. The first X-ray of the lungs was concluded as "negative" (Figure 1), and the second – as "pleuropneumonia" (Figure 2).

The treatment started with quadruple tuberculostatic combination of streptomycin, isoniazid,

rifampicin, and pyrazinamide. Supportive treatment was performed with dexamethasone, mannitol, furosemide, hepatoprotectors, infusion of fluids and blood products. Nevertheless, the patient's condition worsened, cranial nerves were involved (oculomotor nerve and abducens nerve palsies), respiratory disorders appeared requiring mechanic ventilation. The patient died on the 20th day after admission to the hospital.

On autopsy, infiltrative-pneumonic TB in the three right pulmonary lobes, as well as fibrinous-purulent pneumonia and purulent bronchiolitis in the medial and inferior lobe were found and confirmed histologically. Subsequently, tuberculous meningoencephalitis was found, morphologically visualised by lymphocytic basal meningitis and parenchymal vessel vasculitis with microthrombosis. The direct cause of the lethal outcome was multifocal cortical and basal ischemia with severe pulmonary disorders.

DISCUSSION

We present here a case of severe tuberculous meningoencephalitis co-existing with extra-neural pulmonary TB. The sudden onset of strong headache, vomiting, and fever is not typical for neurotuberculosis. The rapid unfavourable course of the case also is not characteristic of CNS TB – within four days following admission the patient became unconscious.

Suggestive for diagnosis is the fact that six months prior to admission, the patient had contact with relatives suffered from TB and three months later she was treated for "pneumonia". Probably at that time was the onset of pulmonary TB.

The diagnosis of tuberculous meningitis is difficult. According to Thwaites et al (2002), there are five diagnostic criteria such as patient's age, duration of complaints, number of WBC count, number of leucocytes in CSF, and percentage of neutrophils in CSF. Total diagnostic index (DI) is calculated according to the formula: DI (age) + DI (blood white-cell count) + DI (history of illness) + DI (cerebrospinal fluid white-cell count) + DI (% neutrophils in cerebrospinal fluid). Therefore the suggested diagnostic rule is: if the patient has a total diagnostic index score of 4 or less, he or she has tuberculous meningitis, and if the patient has a score of more than 4, he or she has bacterial meningitis (11). In the case we report here, the total diagnostic index score was -3 and was suggestive for tuberculous meningitis (Table 1).

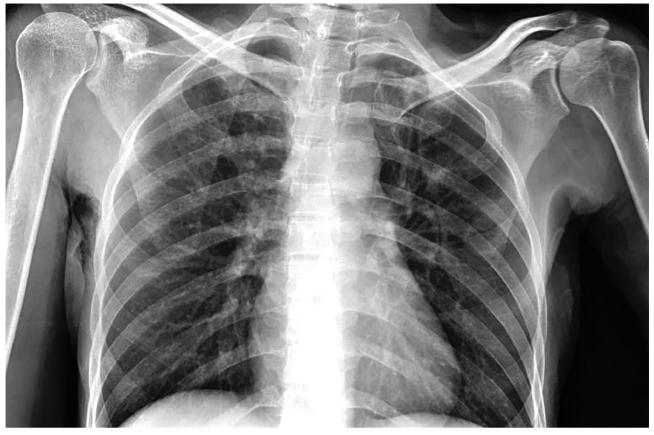


Figure 1. Pulmonary X-ray of the reported case on admission.

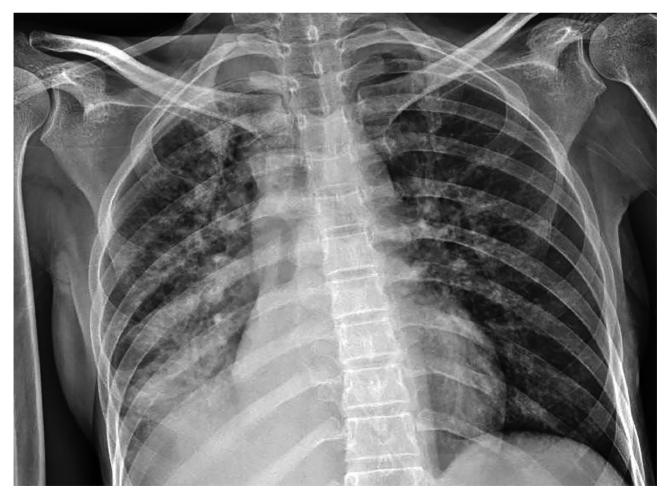


Figure 2. Pulmonary X-ray of the reported case thirteen days after admission.

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Characteristic	Value	Index	Patient's characteristic	Patient's index
Age	≥36 <36	2 0	36	2
WBC	>15,000/mm³ <15,000/mm³	4 0	11,800/mm ³	0
History duration	<6 days ≥6 days	0 -5	7 days	-5
Leucocytes in CSF	>900/mm ³ <900/mm ³	3 0	80/mm ³	0
Percentage of neutrophils in CSF	≥75 <75	4 0	30	0
Total DI				-3*

Table 1. Diagnostic assessment of the reported case according to Thwaites' diagnostic index (6).

WBC – white blood cells; DI – diagnostic index

*Total diagnostic index less than 4 is suggestive for tuberculous meningitis

Qamar et al (2013) considered that features predictive of tuberculous meningitis diagnosis are as follows: protein/glucose ratio of ≥ 2 , pleocytosis < 800, and presence of hydrocephalus (12). In the case we report, mild leucocytosis was found and CSF changes suggested typical tuberculous constellation - markedly elevated protein level contrasted with mild pleocytosis, neutrophils/mononuclears ratio of 3/7, and low glucose level. The ratio CSF protein/ CSF glucose level was 5.16 and in accordance with the consideration of Qamar et al (2013) (12). Dendane et al (2013) considered six features predictive of tuberculous meningitis diagnosis: female gender, duration of symptoms, presence of localising signs, blood leucocytes count, level of serum sodium, and pleocytosis in CSF (13).

The subarachnoid cyst found by CT also provoked diagnostic difficulties. We did not find reports on the simultaneous presence of such formation and tuberculous infection of CNS. In our case CT findings should be interpreted in a complex approach with results from other investigations – an opinion mentioned by Botha et al. (2012) (14). At the same time, pulmonary X-ray investigation is helpful for diagnosis but in the first X-ray there were no characteristic tuberculosis findings. The second X-ray investigation revealed typical pulmonary TB but the patient's condition irreversibly worsened irrespective of initiated treatment with four tuberculostatic drugs.

The treatment was in concordance with contemporary principles of antituberculous therapy mentioned in the WHO Global Tuberculosis Programme (2010) (15). Unfortunately, simultaneous presence of pulmonary TB and tuberculous meningoencephalitis worsened the prognosis of the presented case. Delayed diagnosis of previously untreated pulmonary TB also contributed to the lethal outcome.

Conclusion: The globally increased incidence of TB and co-existence of extra-neural tuberculosis and neurotuberculosis require diagnostic improvement and specific therapy even in suspected cases.

CONFLICT OF INTEREST STATEMENT (AUTHORS):

We certify that there is no conflict of interest with any financial organisation regarding the material discussed in the manuscript.

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