This means that in hospital setting in case of infections caused by *Klebsiella* spp. a preferred choice would be either piperacillin/tazobactam or cefoperazone/sulperazone, or tigecycline. Employing carbapenems, ofloxacin/levofloxacin or doxycycline could allow us counting on a proper effect in slightly more than 50% of cases.

**Conclusion.** Developing the resistance in hospital environment microorganisms would have a significant impact on the treatment outcomes regarding patients with nosocomial infections. A bacteriologic monitoring of such microorganisms evaluating their resistance to a wide range of anti-microbial medications should be a mandatory activity not only in particular inpatient units yet in the region as a whole. This work would contribute both to the development of general standard requirements for empirical therapy of infections in hospitals and to establishing individual approaches when shaping the therapeutic tactics for particular patients.

**References**


** EVALUATION OF KLEBSIELLA SPP. AND ACINETOBACTER SPP. ANTIBIOTIC RESISTANCE IN HOSPITAL ENVIRONMENT (STAVROPOL, RUSSIA) **


Key words: *Klebsiella* spp., Acinetobacter spp., nosocomial infections

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**OЦЕНКА АНТИБИОТИКРЕЗИСТЕНТНОСТИ KLEBSIELLA SPP. И ACINETOBACTER SPP. В СТАЦИОНАРАХ Г. СТАВРОПОЛЯ **

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**Ключевые слова:** Klebsiella spp., Acinetobacter spp., нозокомиальные инфекции

**GENE POLYMORPHISM OF LIPID METABOLISM MARKERS IN CALCIFIC AORTIC VALVE DISEASE**

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Calcfic aortic valve disease (CAVD) (senile, degenerative aortic stenosis) is the process of thickening and calcification of aortic valve (AV) leaflets in the absence of rheumatic heart disease. In the cases of AV sclerosis (calcification) thickened leaflets don’t impact normal intracardiac hemodynamics. Aortic stenosis is characterized by obstruction of the left ventricle outflow tract. Pathogenesis of calcific aortic valve disease involves lipid dismeabolism. [4]. A thorough analysis of the lipid profile in the CAVS patient population revealed increased levels of total cholesterol and athe-
rigenic lipid fractions. It is interesting to note that the total cholesterol in these patients was higher than that of patients with coronary artery disease who underwent coronary bypass surgery [3]. However, only 40% of patients with calcific aortic stenosis (carotid angioplasty with stenting) have hemodynamically significant stenosis of the coronary arteries, and on the contrary in only 2% of patients with coronary artery disease, severe aortic stenosis is detected [1]. Apolipoprotein E (apoE) and paraoxonase 1 (PON1) regulate blood lipid levels being the mediators of lipids hydrolysis and oxidation. Increase of plasma levels of the above substances was observed in aortic valve calcinosis [5]. Perhaps allelic polymorphism of apoE and PON1 genes is responsible for CAVD formation.

The aim of the study was to study the contribution of Leu28Pro polymorphism of apoE gene (rs429358) and Gln192Arg polymorphism of PON1 gene (rs662) to CAVD pathogenesis.

Material and Methods. Open non-randomized comparative case-control trial was performed. Objects were 100 patients (mean age 72.5±7.5 years, 48.5% – men) with CAVD identified. Aortic stenosis was diagnosed in accordance with the recent international guidelines [2]. 46 controls without aortic valve calcification but with the similar profile of cardiovascular background, match for age, sex, the presence of coronary artery disease and treatment were included. The serum apoE and PON1 levels were determined by ELISA using AssayMax Human Apolipoprotein E ELISA Kit (Assaypro, USA) and Human serum paraoxonase 1 (PON1) ELISA Kit (Aviscera Bioscience Inc., USA) in accordance with the manufacturer’s protocols. DNA was isolated from whole blood leukocytes using a DNA-sorb-C kit (Central Research Institute of Epidemiology, Federal Service on Customers’ Rights Protection and Human Wellbeing Surveillance, Russia). In order to detect the Leu28Pro mutation in the apoE (rs429358) gene and the Gln192Arg mutation in the PON1 (rs662) gene, we used the corresponding reagent kits for revealing polymorphisms in the human genome by PCR with SNP-EXPRESS electrophoresis detection scheme (Research and Production Company Litekh, Russia).

Statistical analysis was performed using IBM SPSS Statistics 21 for Windows (IBM SPSS Inc., USA).

Results. The majority of patients in both groups were homozygous for the allele 28Leu of apoE (93% – in CAVD and 96% – in controls, p>0.05), 1 patient with CAVD was homozygous for allele 28Pro, the rest were heterozygous. There was no difference in the prevalence of PON1 allele polymorphisms in the studied groups: Arg192Arg genotype was revealed in 60% of CAVD patients and 50% of controls, Gln192Arg genotype was revealed in 31% of CAVD cases and 34% in controls, and Gln192Gln genotype – in 9% in 7%, respectively.

Allele 28Pro of apoE gene was associated with increased concentration of low density lipoproteins in patients with CAVD (3.9±1.05 vs 3.13±1.08 mmol/l, p<0.02) and total cholesterol in controls (6.2; 6.5 vs 5.11±0.89 mmol/l, p<0.05). PON1 genotype did not influence lipid metabolism parameters in CAVD. Controls with 192Gln allele demonstrated considerably decreased blood levels of total cholesterol (4.86±0.86 vs 5.5±0.87 mmol/l, p<0.05), triglycerides (1.41±0.69 vs 1.86±0.6 mmol/l, p<0.01), very low density lipoproteins (0.64±0.31 vs 0.85±0.27 mmol/l, p<0.01) and apoE (0.02–0.03 vs 0.05(0.02–0.06) mg/ml, p<0.01) and also increased PON1 serum concentration (4.1 (3.2–7.4) vs 3.3 (2.7–3.9) mg/ml, p<0.01).

Conclusion. No association between Gln192Arg and Leu28Pro polymorphisms of PON1 and apoE genes respectively and CAVD was revealed. 28Pro allele of apoE gene was related to the increased blood levels of total cholesterol and low density lipoproteins in CAVD.

References