METHOD OF NON-TUBERCULOSIS MYCOBACTERIA SPECIES IDENTIFICATION IN THE PRESENCE OF ACID-RESISTANT MYCOBACTERIA IN THE SPUTUM OF THE PATIENT

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Abstract

The *aim of study* was to improve the efficiency and standardization the methods of laboratory diagnostics of mycobacteriosis by developing a method for the specific accelerated identification of non-tuberculosis Mycobacterium tuberculosis (NTMB) at bacteriological laboratories.

Materials and methods. The object of the study was sputum from patients with suspected tuberculosis/mycobacteriosis.

For the phenotypic identification of NTMB, two approaches were used: the classical – using biochemical and cultural identification, and the modern – using the BD MGIT TBc ID or STANDARD Q TB MPT64 Ag Test identification immunochromatographic test (manufactured by SD Biosensor INC., Korea) to determine the MPT64 antigen specific for the complex *M. tuberculosis*.

Genotypic species identification of NTMBs was performed using DNA strip technology by Hain Lifescience (Germany) using the GenoType® Mycobacterium CM kit.

Results. A method has been developed for the specific identification of NTMBs in the presence of acid-resistant mycobacteria in the sputum of the patient, in which, using modern genetic methods, a reduction in the period of their identification was achieved. To solve the problem, two molecular genetic systems were used - GeneXpert Ultra and GenoType Micobacterium CM. The use of the GeneXpert Ultra system makes it possible to detect only tuberculosis complex mycobacteria very quickly within 1.5 hours. In the presence of acidresistant mycobacteria in the test sediment, according to the results of light microscopy and a negative test result in the GeneXpert Ultra system, inoculation of sputum sediment in the Middlerooc 7H9 liquid nutrient medium in the WASTEC MGIT system to obtain a pure mycobacterium culture followed by examination in the GenoType Micobacterium CM system allowed the identification of DNA probes linear analysis method (LIPA), which was designed for DNA by amplification and subsequent hybridization on nylon membranes, the so-called DNA-strips with marker oligonucleotide probes. The application of the diagnostic systems, proposed in this study algorithm can significantly reduce the study time - up to 2 working days and identify NTMB types that are frequently encountered and clinically significant.

Conclusion. A method for the specific identification of NTMB in the presence of acid-resistant mycobacteria in the sputum of the patient has been developed. It allows to shorten the identification period of NTMB from 4 weeks to 2 working days.

Key words: mycobacteriosis, diagnostics, non-tuberculous mycobacteria, identification, GenoType Mycobacterium CM/AS, DNA-strips.

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