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INVESTIGATION OF MENTAL RETARDATION ETIOLOGY IN ROMANIAN CHILDREN USING CLINICAL, CYTOGENETIC AND ARRAY-CGH DIAGNOSTIC TECHNIQUES

M. Budisteanu<sup>1,2</sup>, A. Arghir<sup>2</sup>, S.M. Chiriac<sup>2</sup>, A. Tutulan-Cunita<sup>2</sup>, C. Burloiu<sup>3</sup>, C. Iliescu<sup>3</sup>, D. Craiu<sup>3</sup>, D. Barca<sup>3</sup>, B. Budisteanu<sup>4</sup>, I. Minciu<sup>3</sup>, S. Magureanu<sup>3</sup>, A. Lungeanu<sup>2</sup>

<sup>1</sup>Pediatric Neurology, 'Prof. Dr. Alex. Obregia' Clinical Hospital of Psychiatry, <sup>2</sup>Laboratory of Medical Genetics, "Victor Babes" National Institute of Pathology, <sup>3</sup>Pediatric Neurology, <sup>4</sup>Child and Adolescent Psychiatry, "Prof. Dr. Alexandru Obregia" Clinical Hospital of Psychiatry, Bucharest, Romania

**Introduction:** Mental retardation (MR) is the most common developmental disability, affecting 2-3% of the general population. A major challenge in both clinical practice and research in the field of MR is to identify the underlying causes: genetic, chromosomal and environmental factors that have an influence on a person's development and behavior.

**Objective:** We present the results of our study regarding genetic abnormalities associated with mental retardation in children.

**Methods:** A total of 180 children were studied using a diagnostic protocol based on dysmorphic and clinical assessment. A disease, familial and personal history were noted. All patients were evaluated by clinical and paraclinical exams (including dysmorphic features, psychological tests, neurological features, neuroimaging studies). Genetic investigations included a karyotype with GTG banding, FISH and array-CGH.

**Results:** A specific causes for the mental handicap was identified in 80 children (44%). These included a chromosomal abnormality in 32 cases (17%), microdeletion syndromes in 25 children(14%), recognizable syndromes in 23 (13%). Array CGH identified a 22q11 deletion in a girl with unusual phenotype for DiGeorge syndrome, a Xp21 duplication in a girl with severe phenotype (including severe mental retardation, epilepsy, dysmorphic features, genital anomalies, glaucoma, dental anomalies), and a 4p14 deletion in a girl with moderate mental retardation, dysmorphic features, diparesis, congenital heart malformation.

**Conclusions:** While clinical diagnosis and conventional techniques form the mainstay of investigation of children with mental retardation, array CGH proved important diagnostic tool.

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