Complex Genetic Disorders, Genetic Susceptibility to Infections

## AB090. Preliminary results of array CGH test in Vietnamese children with autism spectrum disorder

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**Background:** Autism spectrum disorder (ASD) is highly prevalent accounting for around 1% of children worldwide. Chromosomal microarray is widely recommended as the first-tier test for autistic children with diagnostic yield of around 10%. However, this test is still new with limited clinical study and practice in Vietnam. Our study aims to identify genetic patterns of ASD in the Vietnamese children by using Microarray-based Comparative Genomic Hybridization (aCGH).

**Methods:** We recruit 100 children with ASD and their parents at Vinmec International Hospital in Hanoi. Children are diagnosed with ASD by the Diagnostic and Statistical Manual V (DSM-V), Autism Diagnostic Observation Schedule (ADOS) and Childhood Autism Rating Scale (CARS). Rett and Fragile X tests are applied to exclude probands with Rett and Fragile X syndromes. We perform aCGH test on probands and their parents to detect copy number of variants (CNVs).

**Results:** Here we report our current results on 33 probands. We identified 14 CNVs containing autism-related genes in 11 probands. Results from parents of these 11 probands shows that 4 CNVs on three probands are *de novo* mutations (one proband has two *de novo* CNVs). In addition, one proband has a loss CNV consisting of *NPHP1* gene that is related to ASD in the autosomal recessive inheritance model. The initial findings suggest the specific etiology of ASD in the Vietnamese children with a diagnostic yield of 12%.

**Conclusions:** This is the first empirical study on the Vietnamese children with ASD using aCGH. This work will be expanded to whole exome sequencing and promising findings would be expected.

**Keywords:** Autism Spectrum Disorder (ASD); Microarray-based Comparative Genomic Hybridization (aCGH); Vietnam

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