

First Familial Occurrence of Prader-Willi Syndrome in China: Two Cases and Family Studies

Chao Yunqi, Zou Chaochun

the Children's Hospital of Zhejiang University school of medicine

Introduction

• To present the first reported two cases of familial PWS in China.

Results

1, Case 1: The proband was a 1-day-old boy. He and his sister presented the characteristic features of PWS. They have one suspicious paternal cousin with some autistic type behaviours. SNP array and MS-MLPA detected a paternally transmitted submicrodeletion , 417 kilobase pairs (kbp) in size, at 15q11.2-q13 region: array15q11.2(24,963,375-25,380,656)x1, which was verified to pass through the paternal line from the patients' father and paternal grandmother.

Conclusions

 Molecular genetics investigations are the gold standards for the molecular diagnosis of PWS.

• To carry out familial studies to analyze different underlying cytogenetics and molecular genetics mechanisms and to formulate a comprehensive summary of this rare condition.

Methods

 Detailed clinical features of the probands and the clinical history of other



 The familial occurrence of PWS suggests a wide clinical variability of severity within an affected family and a recessive mode of inheritance. The awareness of familial PWS is of great value for early and accurate diagnosis and administration of appropriate therapy. Conducting relevant family researches is necessary for understanding the mechanism of familial inheritance, estimating the recurrence risks to provide more corresponding genetic counseling on birth guidance and prenatal diagnosis.

affected family members were observed and described.

• The genomic DNA isolation and purification from whole blood was applied to the cytogenetic studies and molecular genetics investigationshigh-resolution microarray analysis-Single Nucleotide Polymorphism (SNP) Array and Methylation Specific-Multiplex Ligation Probe Amplification(MS-MLPA) 2、Case 2: The affected girl is 32 months old, and she had one sister-both of whom fulfilled the diagnostic criteria for PWS. Cytogenetic studies of karyotype analysis revealed that the patients and their unaffected mother shared the same chromosome abnormality-45,XX,rob(15;15)(q10;q10) due to a translocation involving maternal chromosome 15 and hence effective maternal-origin uniparental disomy for the PWS region.



References

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Poster Code: P2 - 135

