A patient with unusual features and a 69.5 Mb duplication from a *de novo* extra der (9): A case report

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Abstract. Partial trisomy 9 is a common autosomal trisomy, which is characterized by non-specific psychomotor delay, mental retardation and moderately abnormal characteristic facial features. Generally, partial trisomy 9 leads to variable phenotypes depending on the size and position of the duplicated region. However, a precise genotype/phenotype map has not been determined. The present study reports the case of a 3-year-old female with certain typical features of trisomy 9p syndrome, who presented with a number of the distinctive symptoms, as well as sensorineural hearing loss, which has not previously been associated with this trisomy. Karyotype, M-FISH and OaCGH analysis were performed on the patient and her parents. The final karyotype was determined to be 47, XX, +mar.ish der (9) (wcp9+). arr cgh 9pterq21.12 (DOCK8→LOC138225)x3. Cytogenetic results showed a de novo extra der (9) with 69.5 Mb duplication. Although the molecular mechanism underlying the hearing loss is unclear, it was proposed that the $9q13 \rightarrow 9q21$ region may be critical for hearing.

Introduction

Partial trisomy 9 is the fourth most common autosomal trisomy after trisomies 21, 18 and 13. Since Rethoré *et al* (1) reported the first identified case of partial trisomy 9 as a chromosomal anomaly, >150 cases have been described. In addition to non-specific psychomotor delay and mental

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retardation, common clinical features, including moderately abnormal characteristic facial features, clinodactyly of the 5th fingers, shortened digits, hypoplastic nails, abnormal dermatoglyphics and hypoplastic brain association with Dandy-Walker malformation are observed (2). Trisomy for 9pter-p21 is hypothesized to be responsible for the majority of these features (3). Intrauterine growth retardation, cleft lip/palate, skeletal anomalies and heart defects are more common with trisomic segments extending through 9q22-9q32 (4-7). In general, partial trisomy 9 leads to variable phenotypes dependent upon the size and position of the duplicated region (8). However, a precise genotype/phenotype map has not yet been proposed. The present study describes the case of a 3-year-old female with a number of the typical features of trisomy 9p syndrome, as well as distinctive features that include sensorineural hearing loss and mild body asymmetry. Cytogenetic results showed the presence of a de novo extra der (9) with 69.5 Mb duplication.

Case report

Case presentation and analysis. A 3-year-old Chinese female was referred to us for further investigation for mental retardation and hearing loss. The girl was born full-term with uneventful gestation by elective cesarean as the first child of nonconsanguineous parents. The mother and father were 28 and 27 years old, respectively, at her birth. Family history was negative, meaning the other families in this pedigree exhibited no similar ilness. The girl had a birth weight of 3,900 g (95th centile), length of 50 cm (50th centile) and head circumference was 35 cm (50th centile), as well as 1 min and 5 min Apgar scores were of 10, respectively. Her psychomotor development was substantially delayed with severe speech retardation. The patient spoke at the age of 3-years and walked without assistance at the age of 2-years. On examination, the girl had a height of 98.5 cm (77th centile) and weight of 16.5 kg (80th centile). The patient presented with a characteristic face with an antimongoloid slant of palpebral fissures, a broad and prominent nasal bridge, low-set and forwardly-rotated auricles, large poorly lobulated ears and downturned corners of the mouth (Fig. 1). A short neck, clinodactyly of both of the 5th fingers, a bilateral simian crease, joint hyperlaxity and hypoplasia of the toenails were also observed. In addition to

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Figure 1. Facial features of a patient with duplication of $9pter \rightarrow 9q21.12$.

the phenotypes of typical trisomy 9p, the patient presented with distinctive features, including the left side of the body slightly smaller than the right with ptosis and strabismus of left eye and sensorineural hearing loss (left ear at 100 decibels, right at 40 decibels). Cerebral computerized tomography showed enlargement of the lateral ventricles (Fig. 2A), 3rd, 4th ventricles and basal cistern, with a mild agenesis of the cerebellar tonsil. Roentgenograms of the skeleton demonstrated hypoplastic pubic bones (Fig. 2B), and bilateral hypoplastic distal phalanges of the feet, pes valgus and bilateral clinodactyly of both 5th fingers (Fig. 2C and D). Cardiac and renal ultrasound findings were normal. This study was approved by the ethics committee of Jinling Hospital, Nanjing University School of Medicine (Nanjing, China), and written informed consent was obtained from the parents.

Chromosome analysis

Karyotype analysis. Karyotyping was performed on peripheral blood lymphocytes from the patient and her parents. Peripheral blood lymphocyte cultures were cultivated using RPMI media supplemented with 10% fetal calf serum (Lai Fu institute of biotechnology, Qing Dao, China). Metaphase chromosomes were GTG-banded using standard procedures.

Multiplex fluorescence in situ hybridization (M-FISH) analysis. M-FISH was performed on the metaphase spreads using Spectra Vysion WCP probe (Vysis, Inc., Downers Grove, IL, USA) according to manufacturer's procedures. Images were captured with Olympus BX51 microscope (Olympus, Tokyo, Japan) and analyzed with the Cytovision 3.0 (Applied Imaging, Sunderland, UK) image analyses software.

OaCGH analysis. In order to investigate the extent of duplication on molecular level, analysis of using a genomic-wide high density oligo array (OaCGH244 K) was conducted according to Agilent manufacturer's procedures and statistical algorithms (www.agilent.com.chem/gocgh) (9).

Chromosomal analysis. showed a female non-mosaic karyotype with an extra chromosome in all metaphases analyzed (Fig. 3A). M-FISH analysis using the Spectra Vysion WCP



Figure 2. Patient analysis. (A) Cerebral computerized tomography showed enlargement of the lateral ventricles. (B) Roentgenogram of the hypoplastic pubic bones. Roentgenogram of the (C) left hand and (D) right hand, shows the left is smaller than the right and there is clinodactyly of both 5th fingers.



Figure 3. Chromosome analysis. (A) An extra der (9) (9pter \rightarrow 9q21) (arrowhead) by high-resolution G-banding (400-band level). (B) The extra chromosome from chromosome 9 (arrowhead) was confirmed by multiplex fluorescence *in situ* hybridization.



Figure 4. A 69.5 Mb duplication segment at genomic position 273,048-72,521,148 bp in the 9pter-q21.12 region was confirmed (Left). The duplicate region spanned 148 annotated genes in which 28 genes are expressed in the cochlea (Right). Dashed circles highlight the location of TMC1 and TJP2 on the map.

Probe (Vysis, Downers Grove, IL, USA) confirmed the extra chromosome from chromosome 9 (Fig. 3B). A 69.5 Mb duplication segment at genomic position 273,048 bp \rightarrow 72,521,148 bp in the 9pter \rightarrow q21.12 region was confirmed (Fig. 4). The final karyotype was interpreted to be 47, XX, +mar.ish der (9) (wcp9+). arr cgh 9pterq21.12 (DOCK8 \rightarrow LOC138225)x3. The duplicate region spanned 148 annotated genes in which 28 genes are expressed in the cochlea (Fig. 4). Chromosome analysis of the parents showed normal karyotype, indicating a *de novo* extra chromosome.

Discussion

To date, 65 genes for non-syndromic hearing loss have been identified (http://hereditaryhearingloss.org/) (10). However, to the best of our knowledge, hearing loss with isolated partial trisomy 9 (9pter \rightarrow q21.12) has not been previously reported. The functions of the 28 genes identified in the chromosomal anal-

ysis, which are expressed in the cochlea, are mostly unknown. Reviewing the literature, cases of two males with partial trisomy 9, including duplication of 9per \rightarrow q21 was reported by Morrissette et al (11) and Centerwall et al (12), respectively; however, the patients succumbed to the disease at four weeks following birth and thus it was uncertain whether or not hearing loss occurred. Comparing our case with other cases in the literature (2-8,13-18) it was found that the patients without hearing loss have overlapping regions of 9pter→9q13 or 9q22-9q32. On the basis of these data, it was hypothesized that 9q13-q21 may be a critical region for hearing. Recently, mutations of two genes in the region of 9q13-9q21.1 were confirmed to be responsible for deafness. For example, transmembrane channel-like gene 1 (TMC1, MIM 606706, GenBank ID NT_023935 position 4301249-4615799), mutations are identified by Kurima et al (19) as a cause of autosomal dominant (#MIM 606705) and autosomal recessive non-syndromic hearing loss (#MIM 600974). The association between mutations in the gene with hearing

loss were further confirmed in other studies (20-23). Between 2002 and 2008, a total of 2 dominant and 18 recessive TMC1 mutations were reported as the cause of hearing loss in 34 families (24). Additionally, Hilgert et al (24) found the other six families with non-syndromic hearing loss were associated with mutations in DFNA36 and DFNB7/11, rather than mutations in TMC1, which implied at least one additional deafness-causing gene at loci DFNA36 and DFNB7/11. Another candidate gene, tight junction protein 2 (TJP2, MIM 607709), was considered a good candidate due to its function as a tight junction protein and its expression in the cochlea. Hilgert et al (24) reported a Guatemalan family with autosomal dominant nonsyndromic hearing loss. In exon 19 of the gene, a novel sequence variant, The mutation, c.2971A>T, was identified in the girl with the hearing loss phenotype, and this lead to an amino acid change from proline to valine at codon 924 (P924V). This aspartic acid residue is a member of a conserved acidic domain of the protein. The mutation was predicted to cause decreased stability by bioinformatic analysis. However, our hypothesis remains to be proven.

In addition to the typical clinical features of partial trisomy 9, the present case presented a group of distinctive phenotypes: The left side of the body was slightly smaller than the right one; left hearing loss was more severe than right; ptosis and strabismus of the left eye, all of which were not previously associated with partial trisomy 9. Body asymmetry is a complex developmental malformation and has already been described in syndromes, such as Beckwith-Wiedemann Syndrome (MIM 147470), Silver-Russell Syndrome (MIM 180860), Proteus syndrome (MIM 176920) and Klippel-Trenaunay-Weber syndrome (MIM 149000). Reviewing the literature, only one case of mosaic tetrasomy 9p with this anomaly was found (25). Considering the malformations are rare, it is uncertain whether the distinctive features were associated with partial trisomy 9 or not. However, the unusual clinical features with a detailed molecular karyotyping may provide information on this phenotype and expand existing knowledge.

In conclusion, the patient carrying a segmental duplication of 9pter-q21.12 exhibits distinctive phenotypes, such as sensorineural hearing loss. Although the molecular mechanism underlying the hearing loss is not clear, it was proposed that the region of $9q13 \rightarrow 9q21$ may be critical for hearing.

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References

 Rethoré MO, Larget-Piet L, Abonyi D, *et al*: 4 cases of trisomy for the short arm of chromosome 9. Individualization of a new morbid entity. Ann Genet 13: 217-232, 1970.

- Tonni G, Lituania M, Chitayat D, Bonasoni MP, Keating S, Thompson M and Shannon P: Complete trisomy 9 with unusual phenotypic associations:Dandy-Walker malformation, cleft lip and cleft palate, cardiovascular abnormalities. Taiwan J Obstet Gynecol 53: 592-597, 2014.
- Lewandowski RC Jr, Yunis JJ, Lehrke R, O'Leary J, Swaiman KF and Sanchez O: Trisomy for the distal half of the short arm of chromosome 9. A variant of the trisomy 9p syndrome. Am J Dis Child 130: 663-667, 1976.
- 4. Hou JW and Wang TR: Molecular c1993netic studies of duplication 9q32->q34.3 inserted into 9q13. Clin Genet 48: 148-150, 1995.
- Naritomi K, Izumikawa Y, Goya Y, Gushiken M, Shiroma N and Hirayama K: Trisomy 9q3 syndrome: a case report and review of the literature. Clin Genet 35: 293-298, 1989.
- Kleczkowska A, Fryns JP, Lemay P and Van den Berghe H: The characteristic phenotype of distal 9q3 trisomy is due to duplication of band 9q32. Genet Couns 4: 217-221, 1993.
- Temtamy SA, Kamel AK, Ismail S, et al: Phenotypic and cytogenetic spectrum of 9p trisomy. Genet Couns 18: 29-48, 2007.
- Wilson GN, Raj A and Baker D: The phenotypic and cytogenetic spectrum of partial trisomy 9. Am J Med Genet 20: 277-282, 1985.
- Fan YS, Jayakar P, Zhu H, et al: Detection of pathogenic gene copy number variations in patients with mental retardation by genomewide oligonucleotide array comparative genomic hybridization. Hum Mutat 28: 1124-1132, 2007.
- Ganapathy A, Pandey N, Srisailapathy CR, *et al*: Non-syndromic hearing impairment in India: High allelic heterogeneity among mutations in TMPRSS3, TMC1, USHIC, CDH23 and TMIE. PLoS One 9: e84773, 2014.
- Morrissette JJ, Laufer-Cahana A, Medne L, *et al*: Patient with trisomy 9p and a hypoplastic left heart with a tricentric chromosome 9. Am J Med Genet A 123A: 279-284, 2003.
- Centerwall WR, Mayeski CA and Cha CC: Trisomy 9q-. a variant of the 9p trisomy syndrome. Humangenetik 29: 91-98, 1975.
- Haddad BR, Lin AE, Wyandt H and Milunsky A: Molecular cytogenetic characterisation of the first familial case of partial 9p duplication (p22p24). J Med Genet 33: 1045-1047, 1996.
- Sanlaville D, Baumann C, Lapierre JM, et al: De novo inverted duplication 9p21pter involving telomeric repeated sequences. Am J Med Genet 83: 125-131, 1999.
- Centerwall WR, Miller KS and Reeves LM: Familial 'partial 9p' trisomy: six cases and four carriers in three generations. J Med Genet 13: 57-61, 1976.
- Sutherland GR, Carter RF and Morris LL: Partial and complete trisomy 9: delineation of a trisomy 9 syndrome. Hum Genet 32: 133-140, 1976.
- Smart RD, Viljoen DL and Fraser B: Partial trisomy 9-further delineation of the phenotype. Am J Med Genet 31: 947-951, 1988.
- Teraoka M, Narahara K, Yokoyama Y, Ninomiya S, Mizuta S, Une T and Seino Y: Maternal origin of a unique extra chromosome, der (9)(pter->q13::q13->q12:) in a girl with typical trisomy 9p syndrome. Am J Med Genet 102: 25-28, 2001.
- Kurima K, Peters LM, Yang Y, *et al*: Dominant and recessive deafness caused by mutations of a novel gene, TMC1, required for cochlear hair-cell function. Nat Genet 30: 277-284, 2002.
- 20. Meyer CG, Gasmelseed NM, Mergani A, *et al*: Novel TMC1 structural and splice variants associated with congenital nonsyndromic deafness in a Sudanese pedigree. Hum Mutat 25: 100, 2005.
- 21. Santos RL, Wajid M, Khan MN, *et al*: Novel sequence variants in the TMC1 gene in Pakistani families with autosomal recessive hearing impairment. Hum Mutat 26: 396, 2005.
- 22. Kitajiri Sİ, McNamara R, Makishima T, *et al*: Identities, frequencies and origins of TMC1 mutations causing DFNB7/B11 deafness in Pakistan. Clin Genet 72: 546-550, 2007.
- Tlili A, Rebeh IB, Aifa-Hmani M, et al: TMC1 but not TMC2 is responsible for autosomal recessive nonsyndromic hearing impairment in Tunisian families. Audiol Neurootol 13: 213-218, 2008.
- Hilgert N, Alasti F, Dieltjens N, *et al*: Mutation analysis of TMC1 identifies four new mutations and suggests an additional deafness gene at loci DFNA36 and DFNB7/11. Clin Genet 74: 223-232, 2008.
- Čazorla Calleja MR, Verdú A and Félix V: Dandy-Walker malformation in an infant with tetrasomy 9p. Brain Dev 25: 220-223, 2003.