



Contents lists available at ScienceDirect

Taiwanese Journal of Obstetrics & Gynecology

journal homepage: www.tjog-online.com

Research Letter

Pregnancy with *de novo* 9q34.3 microdeletion and Kleeftstra syndrome in the fetus may be associated with an abnormal maternal serum screening resultChih-Ping Chen^{a, b, c, d, e, f, *}, Shuan-Pei Lin^{b, g, h, i}, Hui-Bo Li^j, Yen-Ni Chen^a, Wayseen Wang^{b, k}^a Department of Obstetrics and Gynecology, Mackay Memorial Hospital, Taipei, Taiwan^b Department of Medical Research, Mackay Memorial Hospital, Taipei, Taiwan^c Department of Biotechnology, Asia University, Taichung, Taiwan^d School of Chinese Medicine, College of Chinese Medicine, China Medical University, Taichung, Taiwan^e Institute of Clinical and Community Health Nursing, National Yang-Ming University, Taipei, Taiwan^f Department of Obstetrics and Gynecology, School of Medicine, National Yang-Ming University, Taipei, Taiwan^g Department of Medicine, Mackay Medical College, New Taipei City, Taiwan^h Department of Pediatrics, Mackay Memorial Hospital, Taipei, Taiwanⁱ Mackay Medicine, Nursing and Management College, Taipei, Taiwan^j Department of Obstetrics and Gynecology, Chung Shan Hospital, Taipei, Taiwan^k Department of Bioengineering, Tatung University, Taipei, Taiwan

ARTICLE INFO

Article history:

Accepted 30 April 2015

A 32-year-old, primigravid woman underwent amniocentesis in the second trimester due to an abnormal maternal serum screening result of a Down syndrome risk of 1/44 calculated from a level of 0.78 multiples of the median (MoM) of α -fetoprotein (AFP), a level of 3.05 MoM of free β -human chorionic gonadotrophin (β -hCG), a level of 2.66 MoM of inhibin A, and a level of 0.98 MoM of unconjugated estriol (E3) at 15 weeks of gestation. Amniocentesis revealed a karyotype of 46,XY. Prenatal ultrasound findings were unremarkable, and the pregnancy was uneventful. A male baby was delivered smoothly at term with a body weight of 3674 g and a body length of 49 cm. The infant postnatally manifested characteristic facial appearance of Kleeftstra syndrome, intellectual disability, and developmental delay. Array comparative genomic hybridization (aCGH) analysis of the peripheral blood revealed an 8.08-kb 9q34.3 microdeletion or arr 9q34.3 (140,687,823–140,695,906) \times 1 encompassing an OMIM gene of *EHMT1* consistent with the diagnosis of Kleeftstra syndrome.

The present case had an 8.08-kb 9q34.3 microdeletion encompassing the *EHMT1* gene and manifested characteristic Kleeftstra syndrome and 9q subtelomeric deletion syndrome. Kleeftstra

syndrome (OMIM 610253) is caused by haploinsufficiency or heterozygous mutations of the *EHMT1* gene (OMIM 607001) and is characterized by distinct facial dysmorphism of arched eyebrows, midface hypoplasia, upturned nares, full everted lower lip, cupid bowed upper lip and prognathia, congenital heart defects (primarily ventricular septal defect and atrial septal defect) in half of the affected patients, intellectual disability, developmental delay and childhood hypotonia, and minor features of genitourinary defects, seizures, and behavior problems [1–7].

We previously reported prenatal diagnosis of 9q34.3 microdeletion with Kleeftstra syndrome and 3q26.31–q29 duplication in a fetus with abnormal first-trimester maternal serum screening result at 12 weeks of gestation: an elevated level of 4.04 MoM of maternal serum free β -hCG, a level of 1.069 MoM of maternal serum pregnancy-associated plasma protein-A (PAPP-A), a Down syndrome risk of 1/8 and a trisomy 18 risk of 1/136 [8]. We additionally also reported prenatal diagnosis of mosaic ring chromosome 9 or r(9)(p24q34.3) in a pregnancy with an abnormal second-trimester maternal serum screening result at 15 weeks of gestation: a level of 0.63 MoM of maternal serum AFP, a level of 1.15 MoM of maternal serum free β -hCG level, and a Down syndrome risk of 1/57 [9]. The present case provides additional evidence that pregnancy with 9q34.3 microdeletion and Kleeftstra syndrome in the fetus may be associated with an abnormal maternal serum screening result but without abnormal fetal sonographic findings. The aCGH analysis has the advantage of detecting microdeletion syndromes in the fetuses with a normal karyotype by conventional cytogenetic analysis [10,11]. In this regard, the Kleeftstra syndrome may be detected by prenatal application of aCGH in addition to conventional cytogenetic analysis in the pregnancy with an abnormal

* Corresponding author. Department of Obstetrics and Gynecology, Mackay Memorial Hospital, 92, Section 2, Chung-Shan North Road, Taipei, Taiwan.

E-mail address: cpc_mmh@yahoo.com (C.-P. Chen).

maternal serum screening result. We suggest an application of aCGH on uncultured amniocytes at prenatal diagnosis of microdeletion syndrome in the following pregnancies if necessary.

Conflicts of interest

The authors have no conflicts of interest relevant to this article.

Acknowledgments

This work was supported by research grants NSC-101-2314-B-195-011-MY3 and MOST 103-2314-B-195-010 from the Ministry of Science and Technology and MMH-E-104-04 from Mackay Memorial Hospital, Taipei, Taiwan.

References

- [1] Ogawa H, Ishiguro K, Gaubatz S, Livingston DM, Nakatani Y. A complex with chromatin modifiers that occupies E2F- and Myc-responsive genes in G0 cells. *Science* 2002;296:1132–6.
- [2] Kleefstra T, Smidt M, Banning MJG, Oudakker AR, Van Esch H, de Brouwer AP, et al. Disruption of the gene euchromatin histone methyl transferase1 (EHMTase1) is associated with the 9q34 subtelomeric deletion syndrome. *J Med Genet* 2005;42:299–306.
- [3] Kleefstra T, Brunner HG, Amiel J, Oudakker AR, Nillesen WM, Magee A, et al. Loss-of-function mutations in euchromatin histone methyl transferase 1 (EHMT1) cause the 9q34 subtelomeric deletion syndrome. *Am J Hum Genet* 2006;79:370–7.
- [4] Kleefstra T, van Zelst-Stams WA, Nillesen WM, Cormier-Daire V, Houge G, Foulds N, et al. Further clinical and molecular delineation of the 9q subtelomeric deletion syndrome supports a major contribution of *EHMT1* haploinsufficiency to the core phenotype. *J Med Genet* 2009;46:598–606.
- [5] Kleefstra T, Nillesen WM, Yntema HG. Kleefstra Syndrome. In: Pagon RA, Adam MP, Ardinger HH, Wallace SE, Amemiya A, Bean LJH, et al., editors. *GeneReviews* [Internet]. Seattle (WA): University of Washington, Seattle; 1993. Available at: <http://www.ncbi.nlm.nih.gov/books/NBK47079/>. Oct 5, 2010. [accessed 30.03.15].
- [6] Stewart D, Kleefstra T. The chromosome 9q subtelomere deletion syndrome. *Am J Med Genet C Semin Med Genet* 2007;145C:383–92.
- [7] Willemsen MH, Vulto-van Silfhout AT, Nillesen WM, Wissink-Lindhout WM, van Bokhoven H, Philip N, et al. Update on Kleefstra syndrome. *Mol Syndromol* 2012;2:202–12.
- [8] Chen CP, Lin CJ, Chen YY, Wang LK, Chern SR, Wu PS, et al. 3q26.31-q29 duplication and 9q34.3 microdeletion associated with omphalocele, ventricular septal defect, abnormal first-trimester maternal serum screening and increased nuchal translucency: prenatal diagnosis and aCGH characterization. *Gene* 2013;532:80–6.
- [9] Chen CP, Lin CL, Chen LL, Lee CC, Wang W. Prenatal diagnosis of mosaic ring chromosome 9. *Prenat Diagn* 2006;26:870–1.
- [10] Chen CP, Chang SD, Wang TH, Wang LK, Tsai JD, Liu YP, et al. Detection of recurrent transmission of 17q12 microdeletion by array comparative genomic hybridization in a fetus with prenatally diagnosed hydronephrosis, hydro-ureter and multicystic kidney, and variable clinical spectrum in the family. *Taiwan J Obstet Gynecol* 2013;52:551–7.
- [11] Chen CP, Lin CJ, Chern SR, Liu YP, Kuo YL, Chen YN, et al. Prenatal diagnosis and molecular cytogenetic characterization of a 1.07-Mb microdeletion at 5q35.2-q35.3 associated with *NSD1* haploinsufficiency and Sotos syndrome. *Taiwan J Obstet Gynecol* 2014;53:583–7.