CASE REPORT

DOWN SYNDROME AND PATAU SYNDROME IN THE SAME SIBSHIP: RANDOM OR NOT?

Foong Eva¹, Hasliani Hassan², Azizah Othman², Ilunihayati Ibrahim³, Nazihah Mohd Yunus¹, Siti Mariam Ismail¹, Nik Mohd Zulfikri Bin Mat Zin¹, Asraihan Abdul Malik¹, Nurul Alia Nawi¹, Ravindran Ankathil¹, Zilfalil Bin Alwi¹

- 1. Human Genome Centre, School of Medical Sciences, Universiti Sains Malaysia, 16150, Kubang Kerian, Kelantan, Malaysia.
- 2. Department of Paediatrics, School of Medical Sciences, Universiti Sains Malaysia, 16150, Kubang Kerian, Kelantan, Malaysia.
- 3. Department of Haematology, Hospital Raja Perempuan Zainab II, Kota Bharu, Kelantan, Malaysia.

Abstract

Objectives: Chromosomal abnormalities especially aneuploidies are the most common etiology for pregnancy loss. Trisomy 13, trisomy 18 and trisomy 21 are the most common chromosome autosomal aneuploidies with trisomy 21 (Down syndrome) being the most common chromosomal abnormality among liveborn infants. In previous reports, we noted that the recurrence of these aneuploidies in some families may not occur by chance alone. Methods: Extraction of relevant data from review of medical case notes of a young couple with two offspring with Down syndrome (DS) and Patau syndrome. Results: A family history of DS is a predisposing factor for both DS and other types of aneuploidy. Certain instances of non-disjunction error are not random. Conclusion: As the maternal age was not advanced in both pregnancies, there is a possibility that the recurrent aneuploidy in this family may not be accounted by chance alone. The risk of having subsequent affected pregnancy cannot be ignored in this family and prenatal diagnosis is strongly recommended in the subsequent pregnancy.

Keywords: Recurrent Aneuploidy, Down Syndrome, Trisomy 21, Patau Syndrome, Trisomy 13 Corresponding Author: Dr Foong Eva, Human Genome Centre, School of Medical Sciences, Universiti Sains Malaysia, 16150, Kubang Kerian, Kelantan, Malaysia

Tel: +609-7676789 Fax: +609-7653370

Email: foongeva85@gmail.com

Introduction

Fifteen percent of clinically recognized pregnancies result in fetal death [1]. Cytogenetic abnormalities more are common in spontaneous abortions (50% of fetal deaths less than 20 weeks) than in stillbirths (6 to 13% of fetal deaths more than 20 weeks) [1]. A 1998 multicentre survey of 103 069 live births in the United identified major chromosomal abnormalities in 1 in 140 live births [2]. The

American College of Obstetricians and Gynaecologists also published similar incidence rate in which about 1 in 150 babies is born with a chromosomal abnormality [3-5]. Trisomy 13, trisomy 18 and trisomy 21, the most common chromosome autosomal aneuploidies. accounted for 73.6% of all clinically significant chromosomal abnormalities with a rate of 1.13% [6]. In an earlier report, the incidence of Down syndrome in Malaysia has been reported as 1 in 950 and little variation has been reported among the three largest ethnic groups (Malays 1:981, Chinese 1:940, Indians 1:860) [7.8]. Unfortunately, there are no reports on the recent incidence of Down syndrome in the Malaysian population [8]. This case report describes a family with Down syndrome and Patau syndrome in same sibship. The recurrence of these aneuploidies in this family may not occur by chance alone.

Case history

A borderline premature baby was born to a 31-year-old woman who was gravida 2 para 1. He was born vigorous but appeared dysmorphic. He had hypertelorism, depressed nasal bridge, low set ears and single palmar crease. He was clinically diagnosed to have Down syndrome and

cytogenetic analysis was sent. He had transient hypothyroidism and required L-thyroxine 25mcg up till 10 months of age. He also had congenital nasolacrimal duct obstruction, borderline hypermetropia and left alternating esotropia. Hearing assessment was normal but he required speech therapy. Otherwise he is well and attends kindergarten.

In the third pregnancy, the maternal age was 35 years old. A baby boy was born and subsequently diagnosed to have Patau syndrome. The labour was uneventful but the baby was born with Appar score of 6 at 1 minute and 9 at 5 minutes and he required resuscitation. Upon examination, the baby appeared dysmorphic and had cyanosis. He had a broad forehead, depressed nasal hypertelorism, receding bridge, chin. webbed neck, trisomy fingers, micropenis, joint laxity of both ankle, bilateral rocker bottom feet and cutis aplasia congenita. Echocardiography of the heart showed Tetralogy of Fallot with severe pulmonary hypertension. Cranial ultrasound revealed corpus dysgenesis. Cytogenetic analysis was sent. The findings were consistent with a clinical diagnosis of Patau Syndrome. Family counselling was performed and the child was managed conservatively; he died at 6 months of age.

2 5

7 years old

9 years old

9 years old

Normal female

Normal male

Down syndrome

Patau syndrome

Figure 1. Family tree of the family. One of the paternal brother has a daughter with Down syndrome

Cytogenetics

Chromosomal analysis was performed in the two probands and their parents. Chromosomal preparations obtained from Phytohaemagglutinin (PHA) – stimulated peripheral blood cultures, were subjected to Giemsa-Trypsin-Giemsa (GTG) banding and karyotyping was done according to ISCN 2009 and ISCN 2014 Chromosomal

analysis (using ISCN 2009) of first proband revealed 47,XY,+21 [Figure 2]. Chromosomal analysis (using ISCN 2014) of second proband revealed 47,XY,+13 [Figure 3]. Fluorescent in situ hybridization (FISH) analysis was also performed on the second proband and revealed 3 signals for chromosome 13 in 200 interphase nuclei examined. Parental karyotypes according to ISCN 2014 were normal [Figures 4 and 5].

Figure 2. Cytogenetic analysis of the male infant with Down Syndrome



Figure 3. Cytogenetic analysis of the male infant with Patau Syndrome

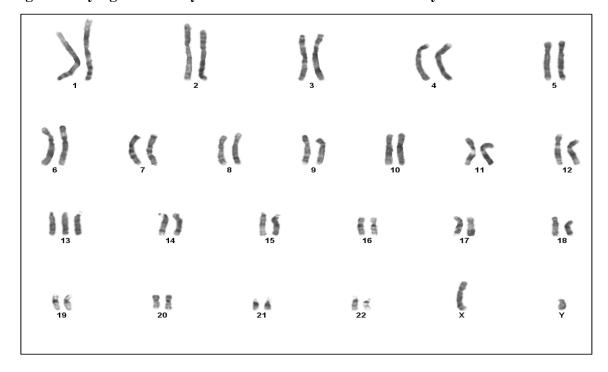


Figure 4. Cytogenetic analysis of the father showing a karyotype of 46,XY

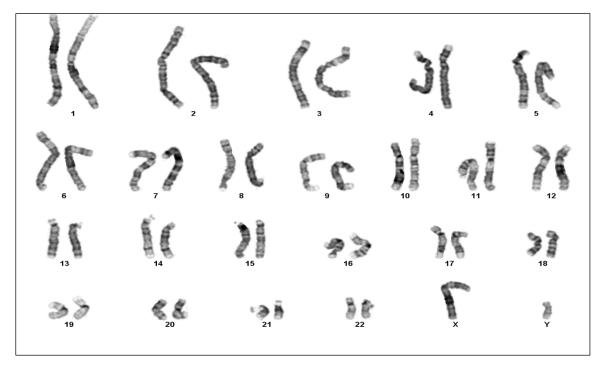
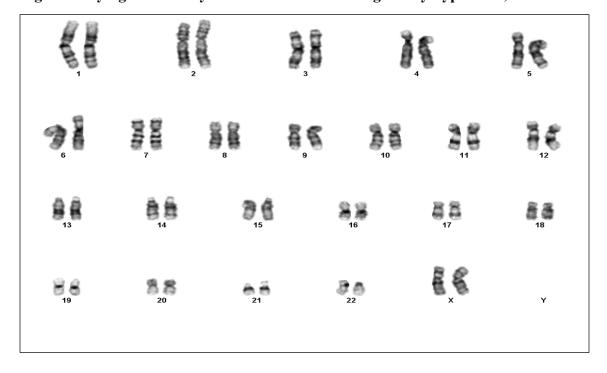


Figure 5. Cytogenetic analysis of the mother showing a karyotype of 46,XX



Discussion

To date. chromosomal abnormalities especially aneuploidies are the most common etiology for pregnancy loss. Trisomy 21 (Down syndrome-DS) remains the most common chromosomal abnormality among liveborn infants [6,9-11]. In our case report, this family had a son with Down syndrome during the second pregnancy. A family history of DS is a predisposing factor for both DS and other types of aneuploidy [12-16]. Some of these recurrences may occur by chance alone but this cannot account for most of them. Familial concentrations of patients with different cytogenetic abnormalities has supported that certain instances of non-disjunction error are not random [17-25]. As seen in our case report, this family also had a son with Patau syndrome. As the maternal age was not advanced in both pregnancies, there is a possibility that the recurrent aneuploidy in

this family may not be accounted by chance alone

Table 1 shows the second trimester amniocentesis results in four studies of women who had the procedure because of a previous pregnancy with trisomy Overall, of the 4953 pregnancies, 42 (0.85%) had DS, a highly statistically significant excess of 27 (0.54%) compared with the number expected on the basis of maternal age alone (chi-squared=46; p \leq 0.0001) [26]. The studies in Table 1 also documented aneuploidies other than DS among these group of women. There were affected pregnancies (0.61%),statistically significant excess of 12 (0.25%) compared with the number expected from the maternal age alone (chi-squared=8.3; p <0.005). Of the 30 affected pregnancies, three have Patau syndrome. This clearly supports that some of the recurrences of cytogenetic abnormalities are not explained by chance alone.

Table 1. Risk of DS and other aneuploidy in 4 studies of women having amniocentesis because of previous DS pregnancy, Arbuzova S et al, 2001 [26]

Study (reference)	Down syndrome						Other
(1010101100)	Maternal age					Total	aneuploidy*
	<25	25-29	30-34	35-39	40+		
Canada (22)	0/51	0/96	1/64	0/24	1/7	2/242	2
Europe I (8)	2/199	1/452	1/418	3/244	0/75	7/1388	10
Europe II (9)	3/331	7/826	2/734	6/343	1/119	19/2353	13
Japan (10)	0/41	5/301	3/394	5/195	1/39	14/970	5
Total	5/622	13/1675	7/1610	14/806	3/240	42/4953	30
Rate (%)	0.80	0.78	0.43	1.74	1.25	0.85	0.61
Expected (%)**	0.10	0.13	0.20	0.59	2.20	0.31	0.36
Excess (%)	0.70	0.65	0.23	1.15	-0.75	0.54	0.25

*Klinefelter (6), Edwards (4), Turner (4), Patau (3), XYY (3), XXX (2), +fragment, +marker and 5 others of unspecified karyotype.

**Based on prevalence rates for each chromosomal abnormality among livebirths in 5-year maternal age intervals, adjusted for the estimated rate of fetal loss from the time of amniocentesis (1).

Arbuzova S et al., [26] also emphasized that familial aggregation of DS and other aneuploidies is not attributable to chance alone and cannot be satisfactorily explained by parental mosaicism. Evidence from several sources strongly suggests the involvement of mitochondrial (mtDNA) in the aetiology of DS and other aneuploidies [26-31]. Mutations in mtDNA bring about an increase in the generation of free radicals and reduce ATP levels, and thereby may affect the synaptonemal complex, chromosomal segregation and division spindle, alter recombination (the enzymes participating in recombination and DNA repair are ATP dependent [32, 33]) and thus lead to aneuploidy. It is well established that the number of mtDNA mutations increases with age in different cells, particularly in oocytes [28], as does the risk of trisomy 21 [6, 27, 30, 31], trisomy 13 and trisomy 18 [6]. However, in our case report, as maternal age was not advanced, involvement of mtDNA mutation may not be the aetiology of her recurrent aneuploidy.

In this case report, we postulate that the recurrence of trisomy 13 after trisomy 21 in the same siblingship may not be purely by chance. Firstly, maternal age was not advanced in both pregnancies. Secondly, maternal factor as a cause of recurrent aneuploidy is questionable as there is family history of Down syndrome within the paternal family. The risk of having subsequent affected pregnancy cannot be ignored in this family and prenatal diagnosis is strongly recommended in subsequent pregnancy. These instances of multiple

aneuploidy within families may have resulted from common factors producing repeated meiotic errors possibly due to mutation of gene involved in meiosis. Hence further study is needed to determine factors which influence non-disjunction.

For this family, we have counselled the couple on the nature, consequences and general management of children with Down syndrome and Patau syndrome. We also informed the couple on the probability and risk of occurrence of each disorder in their future pregnancy. We advised them on proper family planning. The importance of early planning is emphasized if they wish to have more children as the likelihood of recurrent fetal aneuploidy is increased not only due to genetic factors but also nongenetic factors such as advanced maternal nutritional status and radiation exposure. Information and importance of fetal aneuploidy testing to screen for common aneuploidies in future pregnancy was offered to this couple. Information about community resources and support groups were provided to the family. All of these were done to facilitate the process of informed choices.

Acknowledgements

We would like to thank the Director of Human Genome Centre, Associate Professor Dr. Sarina Sulong and all staff of Cytogenetic Laboratory of the Human Genome Centre, School of Medical Sciences, Universiti Sains Malaysia for their assistance in providing the cytogenetic analysis in this family. We would also like to thank the family who kindly consented to this case report.

References

- [1] Reddy UM, Page GP, Saade GR. The role of DNA microarrays in the evaluation of fetal death. Prenat Diagn. 2012; 32:371.
- [2] Hsu LYF. Prenatal diagnosis of chromosomal abnormalities through amniocentesis. In: Genetic disorders and the fetus, 4th ed, Milunsky A (Ed), The Johns Hopkins University Press, Baltimore 1998. p.179.
- [3] American College of Obstetricians and Gynecologists. Prenatal diagnosis of fetal chromosomal abnormalities. ACOG Prac Bull. 2001; 27: 12.
- [4] American College of Obstetricians and Gynecologists. Your pregnancy and birth (4th ed.). Washington, DC: ACOG. 2005.
- [5] Carey JC. Chromosomal disorders in Rudolph and Rudolph (Eds.), Rudolph's Pediatrics. New York: McGraw Hill, 2003; 731-41.
- [6] Zhu Y, Lu S, Bian X, Wang H, Zhu B, Wang H, et al. A multicenter study of fetal chromosomal abnormalities in Chinese women of advanced maternal age. Taiwanese Journal of Obstetrics and Gynecology.2016; 55: 379-84.
- [7] Hoe TS, Boo NY, Clyde MM. Incidence of Down's syndrome in a large Malaysian maternity hospital

- over an 18 month period. Singapore Med J. 1989; 30:246-8.
- [8] Azman BZ, Ankathil R, Siti Mariam I, Suhaida MA, Norhashimah M, Tarmizi AB et al. Cytogenetic and clinical profile of Down syndrome in Northeast Malaysia. Singapore Med J. 2007; 48(6):550-4.
- [9] Centers for Disease Control and Prevention (CDC). Improved national prevalence estimates for 18 selected major birth defects-United States, 1999-2001. MMWR Morb Mortal Wkly Rep. 2006; 54:1301.
- [10] Collins VR, Muggli EE, Riley M, et al. Is Down syndrome a disappearing birth defect? J Pediatr. 2008; 152:20.
- [11] Weijerman ME, van Furth AM, Vonk Noordegraaf A, et al. Prevalence, neonatal characteristics, and first-year mortality of Down syndrome: a national study. J Pediatr. 2008; 152:15.
- [12] Hook EB. Chromosomal abnormalities: prevalence, risks and recurrence. In: Brock DJH, Rodeck CH, Ferguson-Smith MA, eds. Prenatal Diagnosis and Screening. Edinburgh: Churchill Livingstone, 1992; 351–92.
- [13] Mikkelsen M, Stene J. Previous child with Down syndrome and other chromosome aberration. In: Murken J, Stengel-Rutkowski S, Schwinger EW, eds. Prenatal Diagnosis. Proceedings of the Third European Conference on Prenatal Diagnosis of Genetic Disorders. Stuttgart: Enke, 1979; 22–29.

- [14] Stene J, Stene E, Mikkelsen M. Risk for chromosome abnormality at amniocentesis following a child with a non-inherited chromosome aberration. Prenat Diagn. 1984;4: 81–95.
- [15] Uehara S, Yaegashi N, Maeda T et al. Risk of recurrence of fetal chromosomal aberrations: analysis of trisomy 21, trisomy 18, trisomy 13, and 45,X in 1,076 Japanese mothers. J Obstet Gynaecol Res. 1999; 25 (6): 373–79.
- [16] Stinissen P, Van Roy B, Van Camp G et al. Study of the origin of nondisjunction in a family with two cases of Down syndrome using cytogenetic and molecular polymorphisms. Am J Med Genet Suppl. 1990; 7: 133–6.
- [17] Fitzpatrick DR, Boyd E. Recurrences of trisomy 18 and trisomy 13 after trisomy 21. Hum Genet. 1989; 82: 301.
- [18] Hecht F, Bryant JS, Gruber D, Townes PL. The nonrandomness of chromosomal abnormalities. Association of trisomy 18 and Down's syndrome. New Engl J Med. 1964; 271: 1081–6.
- [19] Miller OJ, Breg WR, Schmickel RD, Tretter W. A family with an XXXXY male, a leukemic male, and two 21-trisomic Mongoloid females. Lancet. 1961;2: 78.
- [20] Therman E, Patau K, Smith DW, Demars RI. The D trisomy syndrome and XO gonadal dysgenesis in two sisters. Am. J. Human Genet. 1961; 13: 193.

- [21] Wright SW, Day RW, Mosier HD, Koons A, Mueller H. Klinefelter's syndrome, Down's syndrome (Mongolism), and twinning in same sibship. J Pediat. 1963; 62: 217–24.
- [22] Benirschke K, Brownhill L, Hoefnagel D, Allen FH. Langdon Down anomaly (mongolism) with 21/21 translocation and Klinefelter's syndrome in same sibship. Cytogenetics. 1962; 1: 75–89.
- [23] David TJ, Jones AJ. Trisomy 21 and trisomy 18 in halfsiblings. Hum Genet. 1975; 27: 351–2.
- [24] Pangalos CG, Conover Talbot C, Lewis JG et al. DNA polymorphism analysis in families with recurrence of free trisomy 21. Am J Hum Genet. 1992; 51: 1015–27.
- [25] Johnston A W, Petrakis J. K. Mongolism and Turner's syndrome in the same sibship. Annals of Human Genetics. 1963; 26 (4): 407–14.
- [26] Arbuzova S, Cuckle H, Mueller R, Sehmi I. Familial Down syndrome: evidence supporting cytoplasmic inheritance. Clin Genet. 2001; 60: 456–62.
- [27] Arbuzova S. About the role of the mitochondrial DNA in the etiology of the regular trisomy 21. Cytol Genet. 1995; 29 (3): 77–80 (in Russian).
- [28] Keefe DL, Niven-Fairchild T, Powell S, Buradagunta S. Mitochondrial deoxyribonucleic acid deletions in oocytes and

- reproductive aging in women. Fertil Steril. 1995; 64 (3): 577–83.
- [29] Beerman F, Hummler E, Franke U, Hansmann I. Maternal modulation of the inheritable meiosis I error Dipl I in mouse oocytes is associated with the type of mitochondrial DNA. Hum Genet. 1988; 79 (4): 338–40.
- [30] Arbuzova S. Why it is necessary to study the role of the mitochondrial genome in trisomy 21 pathogenesis? Down's Syndrome: Res Practice. 1998; 5 (3): 26–9.

- [31] Arbuzova S, Hutchin T, Cuckle H. New mtDNA mutations in Down's syndrome. DSNews. 2000; 7 (2): 27.
- [32] Strick TR, Croquette V, Bensimon D. Single-molecule analysis of DNA uncoiling by a type II topoisomerase. Nature. 2000; 404 (6780): 901–4.
- [33] Schar P, Herrman G, Daly G, Lindahl T. A newly identified DNA ligase of *Saccharomyces cerevisiae* involved in RAD52-independent repair of DNA double-strand breaks. Genes Dev. 1997; 11 (15): 1912–24.