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## Case Report

# Familial Transmission of a Robertsonian Translocation rob(21;22): A Case Report

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## Abstract

**Introduction:** Robertsonian Translocations (RT) are the most common balanced rearrangements. However, double Robertsonian translocations, in which two balanced RT occur simultaneously in the same carrier, are extremely rare conditions.

**Objective:** to study the segregation of an RT rob(21;22) in the offspring of a phenotypically normal couple who were found to carry this RT.

**Case report:** The first daughter of the couple has Down Syndrome and double RT, her karyotype was 45, XX, rob(21;22)2,+21. The couple's karyotypes reveal RT in both parents involving chromosomes 21 and 22. Prenatal diagnosis in the second pregnancy revealed a double RT: 44, XX, rob(21;22) x2; the second child was phenotypically normal. The family pedigree also included other cases of Down syndrome.

**Conclusion:** The consanguinity seems to be a risk factor not only for Mendelian disorders but also for chromosomal rearrangement.

## Introduction

Robertsonian Translocations (RT) are translocations involving two acrocentric chromosomes (13,14,15,21, or 22) by centromeric fusion and loss of the two short arms [1]. They are theoretically not accompanied by phenotypic manifestations. The two chromosomes involved can be two homologous or two non-homologous chromosomes, which accounts for approximately 90% of RTs [1]. RT are more common than translocations between other autosomes and occur with an incidence approximately of 1/1000 [2-4].

The RT rob(13;14) is the most common, with an incidence of 1/1500 in live births [5] and 60 to 70% of RT. The RT rob(14;21) is the second most frequent with a proportion of 10% [6,7]. The occurrence of RT in the homozygous condition is an extremely rare event [8-16], which supposes its inheritance from both parents simultaneously; however, consanguinity could increase this probability [17]. Another situation could be the transmission of a RT from one of the two parents and the

appearance of a second RT *de novo* which represents an even rarer situation, this situation has not yet been reported for the same translocation, but has been reported for two different RT.

RT does not provide specific phenotypic manifestations; however, it increases the risk of fetal losses, infertility, uniparental disomy, and aneuploidy in offspring through the genesis of unbalanced gametes [18-20].

We report, for the first time, the observation of a couple of relatives and their descendants in whom RT rob(21;22) segregates.

## Case presentation

A Tunisian couple (f = 1/16), with no significant pathological history, aged 32 for the husband and 27 for the wife at the time of the first consultation (subject III3 and III4 of Figure 1). The family history is marked by the existence of two cousins of the couple with Down syndrome (subjects III-6 and IV-3 in Figure 1). The couple has a first child with confirmed Down syndrome.

Informed consent was obtained from the parents before the study of the R-band karyotype on circulating blood lymphocytes that showed the following formula: 45, XX, rob(21;22)(q10;q10)\*2,+21 (Figure 2). This result confirms the diagnosis of Down syndrome and permits clarification of its mechanism. This is a trisomy 21 by RT involving a chromosome 21 and a chromosome 22. It is associated with a second RT involving the two other chromosomes, 21 and 22.

The study (after informed consent) of the blood karyotype of the two parents showed that each of them had the same translocation with the following formula: 45, XY, rob(21;22)(q10;q10) in the father and 45,XX,rob(21;22)(q10;q10) in the mother.

During the second pregnancy, the study of the fetal karyotype is proposed from the first trimester. After medical counseling, the couple opted to begin with screening, the nuchal translucency measurement found a value of 1.3 mm for an LCC of 54 mm.

The study of the fetal karyotype is carried out at 16 weeks on amniocytes and finds the following formula: 44, XX, rob(21;22)(q10;q10)\*2 (Figure 3).

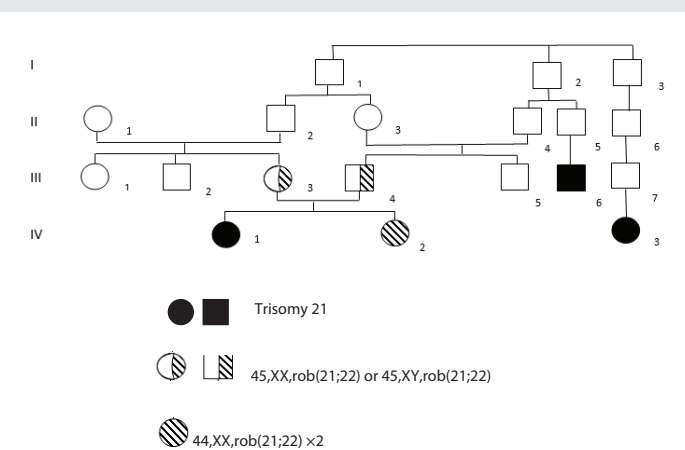


Figure 1: Pedigree of the family.



Figure 2: Karyotype of the first child of the couple showing trisomy 21 and double RT 45, XX, rob(21;22)(q10;q10)\*2,+21.



Figure 3: Fetal balanced karyotype: 44, XX, rob(21;22)(q10;q10)\*2.

## Discussion

This is the first observation reporting the homozygous transmission of the Robertsonian translocation involving chromosomes 21 and 22. The RT rob(21;22) is one of the rarest TRs involving two non-homologous acrocentric chromosomes with a frequency of 1.9% [6,7].

Carriers of balanced chromosomal rearrangements are couples at high risk of having descendants carrying chromosomal aberrations and a high risk of infertility, particularly through recurrent miscarriages [19,20]. The production of gametes balanced for RT is higher during male meiosis, reaching 82% compared to 60% during female meiosis [21–24]. This difference is explained by the blocking of male meiosis during the checkpoints of metaphase I leading to apoptosis of spermatocytes with chromosomal imbalances; only those with balanced content survive without abnormalities or with balanced abnormalities survive [24,25] while the occurrence of RT is more frequent during oogenesis especially *de novo* and without blocking the gametogenesis thus the proportion of balanced gametes will be lower since there are also unbalanced ones [26].

Homozygous transmission of RT in phenotypically healthy subjects is a rare condition. Around twenty observations have focused on the most frequent Robertsonian translocation rob(13;14) for which cases of homozygosity have been reported, particularly in consanguineous subjects [27,28]. The RT (14;21) in the homozygous state has also been reported in an Indian patient with primary amenorrhea [29]. The couple reported in our observation is also consanguineous. The observations, already reported, show, for the most part, a history of fetal loss [8,28], which is not the case for our couple.

Two prenatal observations with a karyotype with 44 chromosomes have also been reported, it is about a rob(14;21) in the homozygous state and the postnatal follow-up found a normal child [9,10].

Even more exceptional are the 44-chromosome karyotypes with two different Robertsonian translocations. This is a situation reported only once [30]. Theoretically, the phenotype

should be normal since the two rearrangements are balanced. However, the effect on the possible combinations at the level of gametes at the time of meiosis makes the situation complex [30].

In our observation, the parents are related and carry the same translocation in the heterozygous state. This is probably a translocation inherited from a common ancestor, according to its low incidence and the probability of finding it in the same couple is low [6,7]. Studying the karyotype of other family members was not possible.

The two other related children, cousins of our couple, carrying Down syndrome, would very likely have the same RT at least in the heterozygous state.

We provided adequate genetic counseling for the couple and advised them to carry out a fetal karyotypic study for each pregnancy (preimplantation diagnosis is currently not available in Tunisia).

Most carriers of RT in the homozygous state are healthy and do not present particular phenotypic traits with preserved fertility; their descendants are in all cases carriers of the same RT in the heterozygous state [15].

The detection of RT in the homozygous state makes it possible to find heterozygous descendants who present a high risk of having children with unbalanced chromosomal rearrangements and enables the proposal of prenatal diagnosis for their offspring or better a preimplantation diagnosis which avoid pregnancy termination and its psychological damages [10].

No difference in the production of unbalanced gametes depending on the sex of the carrier parents has been reported [24].

## Conclusion

This case adds further evidence that people with 45 chromosomes are mostly healthy and free of dysmorphic features. Identification of a RT makes it possible to find heterozygous parents who are at risk of having children with imbalanced chromosomal rearrangements, and then prevent the birth of abnormal offspring. People with 44 chromosomes are extremely rare conditions that must lead to more familial investigations.

## References

1. Sullivan BA, Wolff DJ, Schwartz S. Analysis of centromeric activity in Robertsonian translocations: implications for a functional acrocentric hierarchy. *Chromosoma*. 1994;103:459–67. Available from: <https://doi.org/10.1007/bf00337384>
2. Hamerton JL, Canning N, Ray M, Smith S. A cytogenetic survey of 14,069 newborn infants. I. Incidence of chromosome abnormalities. *Clin Genet*. 1975;8(4):223–43. Available from: <https://doi.org/10.1111/j.1399-0004.1975.tb01498.x>
3. Nielsen J, Wohler M. Chromosome abnormalities found among 34,910 newborn children: results from a 13-year incidence study in Arhus, Denmark. *Hum Genet*. 1991;87(1):81–3. Available from: <https://pubmed.ncbi.nlm.nih.gov/2090319/>

4. Hochstenbach R, van Binsbergen E, Engelen J, Nieuwint A, Polstra A, et al. Array analysis and karyotyping: workflow consequences based on a retrospective study of 36,325 patients with idiopathic developmental delay in the Netherlands. *Eur J Med Genet*. 2009;52(4):161–9. Available from: <https://doi.org/10.1016/j.ejmg.2009.03.015>
5. Page SL, Shin JC, Han JY, Choo KH, Shaffer LG. Breakpoint diversity illustrates distinct mechanisms for Robertsonian translocation formation. *Hum Mol Genet*. 1996;5(9):1279–88. Available from: <https://doi.org/10.1093/hmg/5.9.1279>
6. Schoemaker MJ, Jones ME, Higgins CD, Wright AF, United Kingdom Clinical Cytogenetics Group, Swerdlow AJ. Mortality and cancer incidence in carriers of balanced Robertsonian translocations: a national cohort study. *Am J Epidemiol*. 2019;188:500–8. Available from: <https://doi.org/10.1093/aje/kwy266>
7. Zhao WW, Wu M, Chen F, Jiang S, Su H, et al. Robertsonian translocations: an overview of 872 Robertsonian translocations identified in a diagnostic laboratory in China. *PLoS One*. 2015;10(5):e0122647. Available from: <https://doi.org/10.1371/journal.pone.0122647>
8. Martinez-Castro P, Ramos MC, Rey JA, Benitez J, Sanchez Cascos A. Homozygosity for a Robertsonian translocation (13q14q) in three offspring of heterozygous parents. *Cytogenet Cell Genet*. 1984;38(4):310–2. Available from: <https://doi.org/10.1159/000132080>
9. Rockman-Greenberg C, Ray M, Evans JA, Canning N, Hamerton JL. Homozygous Robertsonian translocations in a fetus with 44 chromosomes. *Hum Genet*. 1982;61(3):181–4. Available from: <https://doi.org/10.1007/bf00296437>
10. Dallapiccola B, Ferranti G, Altissimi D, Colloridi F, Paesano R. First-trimester prenatal diagnosis of homozygous (14;21) translocation in a fetus with 44 chromosomes. *Prenat Diagn*. 1989;9(8):555–8. Available from: <https://doi.org/10.1002/pd.1970090804>
11. Morgan R, Bixenman H, Hecht F. Human chromosome variation with two Robertsonian translocations. *Hum Genet*. 1985;69:178–80. Available from: <https://doi.org/10.1007/bf00293293>
12. Eklund A, Simola KO, Rynänen M. Translocation t(13;14) in nine generations with a case of translocation homozygosity. *Clin Genet*. 1988;33:83–6. Available from: <https://doi.org/10.1111/j.1399-0004.1988.tb03415.x>
13. Rajangam S, Michaelis RC, Velagaleti GVN, Lincoln S, Hegde S, Lewin S, et al. Down syndrome with biparental inheritance of der(14q21q) and maternally derived trisomy 21: confirmation by fluorescence in situ hybridization and microsatellite polymorphism analysis. *Am J Med Genet*. 1997;70:43–7. Available from: [https://doi.org/10.1002/\(sici\)1096-8628\(19970502\)70:1%3C43::aid-ajmg9%3E3.0.co;2-s](https://doi.org/10.1002/(sici)1096-8628(19970502)70:1%3C43::aid-ajmg9%3E3.0.co;2-s)
14. Omrani MD, Gargari SS. Uniparental disomy resulting from heterozygous Robertsonian translocation (13q14q) in both parents. *J Res Med Sci*. 2007;12:100–3. Available from: <http://jrms.mui.ac.ir/index.php/jrms/article/view/483>
15. Malekpour N, Kormi SMA, Azadbakht M, Yousefi M, Hasanzadeh-Nazar Abadi M. The survey of double Robertsonian translocation 13q;14q in the pedigree of 44;XX woman: a case report. *Int J Mol Cell Med*. 2017;6(4):243–8. Available from: <https://doi.org/10.22088/bums.6.4.243>
16. Sahraeean S, Jebelli A, Shahbazi Z, Piryaee F. Homozygosity for Robertsonian translocation (14q;15q) in a newborn with a familial history of recurrent abortion and newborns affected by hepatosplenomegaly: a case report. *J Reprod Infertil*. 2023;24(4):301–5. Available from: <https://doi.org/10.18502/jri.v24i4.14158>
17. Abdalla EM, Kholeif SF, Elshaffie RM. Homozygosity for a Robertsonian translocation (13q;14q) in an otherwise healthy 44,XY man with a history of repeated fetal losses. *Lab Med*. 2013;44:254–7.

18. Yamazawa K, Ogata T, Ferguson-Smith AC. Uniparental disomy and human disease: an overview. *Am J Med Genet C Semin Med Genet.* 2010;154C(3):329–34. Available from: <https://doi.org/10.1002/ajmg.c.30270>
19. Fryns JP, Van Buggenhout G. Structural chromosome rearrangements in couples with recurrent fetal wastage. *Eur J Obstet Gynecol Reprod Biol.* 1998;81:171–6. Available from: [https://doi.org/10.1016/s0301-2115\(98\)00185-7](https://doi.org/10.1016/s0301-2115(98)00185-7)
20. Keymolen K, Van Berkel K, Vosselmanns A, Staessen C, Liebaers I. Pregnancy outcome in carriers of Robertsonian translocations. *Am J Med Genet A.* 2011;155:2381–5. Available from: <https://doi.org/10.1002/ajmg.a.33941>
21. Ferlin A, Garolla A, Foresta C. Chromosome abnormalities in sperm of individuals with constitutional sex chromosomal abnormalities. *Cytogenet Genome Res.* 2005;111:310–6. Available from: <https://doi.org/10.1159/000086905>
22. Sarrate Z, Blanco J, Anton E, Egozcue S, Egozcue J, et al. FISH studies of chromosome abnormalities in germ cells and its relevance in reproductive counseling. *Asian J Androl.* 2005;7:227–36. Available from: <https://doi.org/10.1111/j.1745-7262.2005.00061.x>
23. Tempest HG. Meiotic recombination errors: the origin of sperm aneuploidy and clinical recommendations. *Syst Biol Reprod Med.* 2011;57:93–101. Available from: <https://doi.org/10.3109/19396368.2010.504879>
24. Dang T, Xie P, Zhang Z, Hu L, Tang Y, et al. The effect of carrier characteristics and female age on preimplantation genetic testing results of blastocysts from Robertsonian translocation carriers. *J Assist Reprod Genet.* 2023;40(8):1995–2002. Available from: <https://doi.org/10.1007/s10815-023-02853-5>
25. Eaker S, Pyle A, Cobb J, Handel MA. Evidence for meiotic spindle checkpoint from analysis of spermatocytes from Robertsonian-chromosome heterozygous mice. *J Cell Sci.* 2001;114:2953–65. Available from: <https://doi.org/10.1242/jcs.114.16.2953>
26. Bandyopadhyay R, Heller A, Knox-DuBois C, McCaskill C, Berend SA, et al. Parental origin and timing of de novo Robertsonian translocation formation. *Am J Hum Genet.* 2002;71(6):1456–62. Available from: <https://doi.org/10.1086/344662>
27. Miryounesi M, Diantpour M, Motevaseli E, Ghafouri-Fard S. Homozygosity for a Robertsonian translocation (13q;14q) in an otherwise healthy 44,XX female with a history of recurrent abortion and a normal pregnancy outcome. *J Reprod Infertil.* 2016;17(3):184–7. Available from: <https://pubmed.ncbi.nlm.nih.gov/27478773/>
28. O'Neill ID. Homozygosity for constitutional chromosomal rearrangements: a systematic review with reference to origin, ascertainment and phenotype. *J Hum Genet.* 2010;55(9):559–64. Available from: <https://doi.org/10.1038/jhg.2010.80>
29. Kopakka N, Dalvi R, Shetty DL, Das BR, Mandava S. Balanced autosomal translocation and double Robertsonian translocation in cases of primary amenorrhea in an Indian population. *Int J Gynaecol Obstet.* 2012;116(3):253–7. Available from: <https://doi.org/10.1016/j.ijgo.2011.09.029>
30. Pierron L, Irrmann A, de Chalus A, Bloch A, Heide S, Rogers E, et al. Double chromosomal translocation in an infertile man: one-step FISH meiotic segregation analysis and reproductive prognosis. *J Assist Reprod Genet.* 2019;36(5):973–8. Available from: <https://doi.org/10.1007/s10815-019-01430-z>

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