Aaltonen LA, Ponder BA, Baylin SB, Herman JG (2001*b*) DNA methylation patterns in hereditary human cancers mimic sporadic tumorigenesis. Hum Mol Genet 10:3001– 3007

- Grady WM, Willis J, Guilford PJ, Dunbier AK, Toro TT, Lynch H, Wiesner G, Ferguson K, Eng C, Park JG, Kim SJ, Markowitz S (2000) Methylation of the CDH1 promotor as the second genetic hit in hereditary diffuse gastric cancer. Nat Genet 26:16–17
- Gruis NA, van der Velden P, Sandkuijl LA, Prins DE, Weaver-Feldhaus J, Kamb A, Bergman W, Frants RR (1995) Homozygotes for CDKN2 (p16) germline mutation in Dutch familial melanoma kindreds. Nat Genet 10:351–353
- Herman JG, Graff JR, Myohänen S, Nelkin BD, Maylin SB (1996) Methylation-specific PCR: a novel PCR assay for methylation status of CpG islands. Proc Natl Acad Sci USA 93:9821–9826
- Lal G, Liu L, Hogg D, Lassam NJ, Redston MS, Gallinger S (2000) Patients with both pancreatic adenocarcinoma and melanoma may harbor germline CDKN2A mutations. Genes Chrom Cancer 27:358–361
- Lynch HT, Brand RE, Hogg D, Deters CA, Fusaro RM, Lynch JF, Li L, Knezetic J, Lassam NJ, Goggins M, Kern S (2002) Phenotypic variation in eight extended CDKN2A germline mutation familial atypical multiple mole melanoma-pancreatic carcinoma prone families: the familial atypical mole melanoma-pancreatic carcinoma syndrome. Cancer 94:84–96
- Poi MJ, Yen T, Li J, Song H, Lang JC, Schuller DE, Pearl DK, Casto B, Tsai MD, Weghorst CM (2001) Somatic INK4a-ARF locus mutations: a significant mechanism of gene inactivation in squamous cell carcinomas of the head and neck. Mol Carcinog 30:26–36
- Sambrook J, Fritsch EF, Maniatis T (2nd ed) (1989) Molecular cloning—a laboratory manual. Cold Spring Harbor Laboratory, Cold Spring Harbor, NY
- Schneider-Stock R, Walter H, Haeckel C, Radig K, Rys J, Roessner A (1998) Gene alterations at the CDKN2A (p16/MTS1) locus in soft tissue tumors. Int J Oncol 13:325–329
- van der Velden PA, Sandkuijl LA, Bergman W, Hille ETM, Frants RR, Gruis NA (1999) A locus linked to p16 modifies melanoma risk in Dutch familial atypical multiple mole melanoma (FAMMM) syndrome families. Genome Res 9:575– 580
- van der Velden PA, Sandkuijl LA, Bergman W, Pavel S, van Mourik L, Frants RR, Gruis NA (2001) Melanocortin-1 receptor variant R151C modifies melanoma risk in Dutch families with melanoma. Am J Hum Genet 69:774–779
- Vasen HFA, Gruis NA, Frants RR, van der Velden PA, Hille ETM, Bergman W (2000) Risk of developing pancreatic cancer in families with familial atypical multiple mole melanoma associated with a specific 19 deletion of p16 (p16-Leiden). Int J Cancer 87:809–811

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Another Case of Imprinting Defect in a Girl with Angelman Syndrome Who Was Conceived by Intracytoplasmic Sperm Injection

To the Editor:

Intracytoplasmic sperm injection (ICSI) has been established as an efficient treatment for male infertility and also as a supplement to in vitro fertilization (IVF) without obvious male infertility. ICSI is now regarded as a procedure that is safe overall, and no increase in developmental delay was found in a follow-up of 221 ICSIconceived children in the 2nd year of life (Sutcliffe et al. 2001). However, the possibility of an increased risk of imprinting defects has been raised (Manning et al. 2000). Two children conceived by ICSI who had Angelman syndrome (AS [MIM 105830]) due to a presumably sporadic imprinting defect have recently been reported (Cox et al. 2002).

We here report a 3.5-year-old girl with AS due to a sporadic imprinting defect, born of a pregnancy that was also the result of ICSI. The girl was the first child of a 35-year-old mother and a 36-year-old father. The father has a healthy daughter by another partner, and sperm analysis was normal on three different occasions. The mother had one spontaneous abortion and two extrauterine pregnancies before treatment with IVF. Traditional IVF did not result in fertilized eggs, and ICSI was therefore performed in spite of the normal sperm analysis of the father. The first ICSI pregnancy resulted in another spontaneous abortion, whereas the second ICSI procedure resulted in a normal pregnancy. Birth was at term, birth weight was 3,760 g, length was 54 cm, and head circumference was 36 cm (75th percentile). Development was considered normal for the first 3-4 mo, after which she started to have infections. She walked at age 2 years. She had no epilepsy but had an abnormal electroencephalogram with large-amplitude slow-spike waves. There was no language development. Chromosomes, including subtelomeres, were normal. At age 3 years, her height and weight were at the 50th percentile, whereas her head circumference was 1 cm below the 2.5th percentile. She was mentally retarded and atactic. She was dysmorphic, with a square face, deep-set eyes, and a protruding tongue.

FISH analysis using the *SNRPN* probe (MIM 182279), as well as microsatellite studies, revealed normal chromosomes 15 of biparental origin. A common large deletion of 15q11-q13 and uniparental paternal disomy could therefore be excluded. Methylation-specific Southern blot analysis and methylation-specific PCR (Zeschnigk et al. 1997) for the *SNRPN* locus showed the presence of a normal unmethylated paternal band and the complete ab-

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sence of a methylated maternal band, indicating that the patient had an imprinting defect. Quantitative Southern blot analysis of the critical AS imprinting center (IC) region (Buiting et al. 1999) showed a normal dosage; therefore, an IC deletion was unlikely. This result was confirmed by sequence analysis of the 880-bp AS-IC element, where the patient was heterozygous for three different SNPs. Both parents had normal chromosomes and a normal methylation pattern. These findings suggest that the patient belongs to the group of patients with a sporadic imprinting defect (Buiting et al. 1998).

Both patients reported by Cox et al. (2002) had fathers with sperm abnormalities, and the possibility that the imprinting defect could be related to male infertility was discussed. The father of the child reported here had normal sperm, and a relationship to male infertility is therefore unlikely. However, similar to the mother, the maternal grandmother also had a history of reproductive difficulties. In addition to three healthy children, she had four spontaneous abortions and one daughter who was stillborn at term. Since both the maternal grandmother and the mother had reproductive difficulties, a maternal oogenesis defect cannot be excluded.

A sporadic imprinting defect is a very rare cause of AS, and Cox et al. (2002) therefore considered a relationship to the ICSI procedure to be likely. The report of a third patient with this rare disorder further supports the assumption that ICSI can lead to an increased risk for imprinting defects.

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Electronic-Database Information

Accession numbers and the URL for data presented herein are as follows:

Online Mendelian Inheritance in Man (OMIM), http://www .ncbi.nlm.nih.gov/Omim/ (for the AS gene [MIM 105830] and the SNRPN probe [MIM 182279])

References

- Buiting K, Dittrich B, Groß S, Lich C, Färber C, Buchholz T, Smith E, et al (1998) Sporadic imprinting defects in Prader-Willi syndrome and Angelman syndrome: implications for imprint-switch models, genetic counseling, and prenatal diagnosis. Am J Hum Genet 63:170–180
- Buiting K, Lich C, Cottrell S, Barnicoat A, Horsthemke B (1999) A 5-kb imprinting center deletion in a family with Angelman syndrome reduces the shortest region of deletion overlap to 880 bp. Hum Genet 105:665–666
- Cox GF, Bürger J, Lip V, Ulrike A, Mau UA, Sperling K, Wu BL, Horsthemke B (2002) Intracytoplasmic sperm injection may increase the risk of imprinting defects. Am J Hum Genet 71:162–164
- Manning M, Lissens W, Bonduelle M, Camus M, De Rijcke M, Liebaers I, Van Steirteghem A (2000) Study of DNA-methylation patterns at chromosome 15q11-q13 in children born after ICSI reveals no imprinting defects. Mol Hum Reprod 6: 1049–1053
- Sutcliffe AG, Taylor B, Saunders K, Thornton S, Liebermann BA, Grudzinskas JG (2001) Outcome in the second year of life after in-vitro fertilisation by intracytosplamic sperm injection: a UK case-control study. Lancet 357:2080–2084
- Zeschnigk M, Lich C, Buiting K, Horsthemke B, Dörfler W (1997) A single-tube PCR test for the diagnosis of Angelman and Prader-Willi syndrome based on allelic methylation differences at the *SNRPN* locus. Eur J Hum Genet 5:94–98

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