Healthy Baby Born to a Robertsonian Translocation Carrier following Next-Generation Sequencing-Based Preimplantation Genetic Diagnosis: A Case Report

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Abstract

Preimplantation genetic diagnosis (PGD) is well established method for treatment of genetic problems associated with infertility. Moreover, PGD with next-generation sequencing (NGS) provide new possibilities for diagnosis and new parameters for evaluation in, for example, aneuploidy screening. The aim of the study was to report the successful pregnancy outcome following PGD with NGS as the method for 24 chromosome aneuploidy screening in the case of Robertsonian translocation. Day 3 embryos screening for chromosomal aneuploidy was performed in two consecutive in vitro fertilization (IVF) cycles, first with fluorescent in situ hybridization (FISH), and then with NGS-based protocol. In each IVF attempt, three embryos were biopsied. Short duration of procedures enabled fresh embryo transfer without the need for vitrification. First IVF cycle with the embryo selected using PGD analysis with the FISH method ended with pregnancy loss in week 8. The second attempt with NGS-based aneuploidy screening led to exclusion of the following two embryos: one embryo with 22 monosomy and one with multiple aneuploidies. The transfer of the only euploid blastocyst resulted in the successful pregnancy outcome. The identification of the euploid embryo based on the NGS application was the first successful clinical application of NGS-based PGD in the case of the Robertsonian translocation carrier couple.

Keywords

► aneuploidy screening  
► next-generation sequencing  
► preimplantation genetic diagnosis  
► Robertsonian translocation

Robertsonian translocations are classified as the most common balanced structural chromosomal rearrangements in the human population occurring with the frequency of approximately 1 in 1,000 newborns.1 The majority of Robertsonian translocation cases involve two different acrocentric chromosomes. Carriers of heterologous Robertsonian translocations are diagnosed more frequently among the infertile couples (especially in oligospermic males) than in healthy population.2 Because of the well-known risk of unbalanced conceptions in carriers of Robertsonian translocations, the

ISSN 2157-7005.
male carriers are the candidates for intracytoplasmic sperm injection (ICSI)—IVF along with preimplantation genetic diagnosis (PGD).\textsuperscript{3,4}

Previously published cases of PGD concerning Robertsonian translocation carriers usually involved the diagnosis of ploidy status using the multicolor FISH method. However, the reports of imbalances of other embryo chromosomes in such cases\textsuperscript{5} indicated the need for broadening the analysis to include additional chromosome copy number abnormalities. The next-generation sequencing (NGS) has been used routinely in various PGD protocols at INVICTA Fertility Clinic since August 2013. This report presents NGS-based PGD application in the case of a Robertsonian translocation carrier.

**Patients and Methods**

In 2011, a couple with 5 years of primary infertility (females 31 and males 46) was seeking treatment at the INVICTA Fertility Clinic. In 2009, male infertility factor with severe oligospermia was diagnosed and determined as the reason of infertility.

The couple underwent standard clinical investigation and all the hormonal results were normal including AMH (anti-Mullerian hormone) at 2.4 ng/mL (AnshLabs I generation assay) and unchanged female karyotype. Male karyotype was abnormal with Robertsonian translocation 45.XY,der (14;15) (q10;q10). The couple’s decision to undergo IVF was followed by genetic counseling regarding PGD options and presentation of success rates, risk of misdiagnosis, and possible genetic, clinical, and social outcomes. Because of the low risk of father-to-child transmission in case of father being Robertsonian heterozygote, the couple decided not to perform PGD directed to 14 and 15 uniparental disomy.\textsuperscript{1}

Translocation carrier state is associated with high risk of aneuploidy in offspring, therefore, we proposed FISH analysis for fused chromosomes 14q and 15q. A written consent form, in which a possible risk of the in vitro procedure and PGD diagnosis of translocation and aneuploidy of all 24 chromosomes simultaneously with the use of new NGS-based PGD method. In September 2013, the ICSI procedure and the embryo culture with preparation for PGD were performed as described earlier. Three embryos obtained in second IVF cycle were biopsied on day 3. Cells were transferred into thin-walled 0.2 mL polymerase chain reaction (PCR) tubes containing sodium dodecyl sulphate (SDS) lysis buffer with proteinase K and frozen at −20°C. After that, cells were lysed by incubation for 20 minutes at 37°C and 15 minutes at 65°C.

PGD NGS method was designed at the INVICTA Fertility Clinic Molecular Laboratory Department. Whole genome amplification (WGA) was performed according to the manufacturer’s protocol (PicoPlex WGA kit, New England Biolabs, Ipswich, MA). Concentration of DNA after WGA was quantified with Qubit 2.0 Fluorometer and Qubit dsDNA HS Assay (Life technologies, CA). For DNA fragmentation, Ion Xpress Plus Fragment Library Kit, Life Technologies, according to manufacturer’s protocol for 10 to 100 ng of gDNA input was used. For barcoding, Ion Xpress Barcode Adapters 1–96 Kit (Life Technologies) were used. Ion Xpress Equalizer Kit (Life Technologies) was used for library input normalization, according to manufacturer’s protocol. Barcoded libraries were clonally amplified with The Ion PGM Template OT2 200 Kit (Life Technologies) using Ion One Touch 2 System. After chip loading, sequencing was performed using Ion PGM Sequencing 200 Kit v2 (Life Technologies) on Ion 314 and 316 chips. Preliminary analysis, for example, base calling and reads mapping against human genome reference sequence (Hg19) were performed with Ion Torrent Suite Software (Life Technologies, Carlsbad, CA). The INVICTA Bioinformatics Team Script was used for further computational calculations. Read coverage for each chromosome was corrected for guanosine-cytosine (GC)-bias, and aneuploidy detection was performed using sample results comparison to baseline values obtained from 72 males and 52 females euploid samples processed beforehand with established protocol described earlier. The INVICTA algorithm was introduced to eliminate the influence of sample-to-sample reads coverage variance on false-positive calls. Male control samples were processed together with probes from blastomeres and underwent the same computational analysis to exclude any performance malfunctions. Negative control sample was processed to exclude contaminations. The protocol was validated using cell lines and was accurate regarding detection of whole chromosome aneuploidies.
On day 5, the transfer of the single healthy, euploid blastocyst was performed. Pregnancy was confirmed and prenatal diagnosis was performed at week 15.

**Results**

Three blastomeres biopsied from three embryos on day 3 of the culture were analyzed in both cycles. The embryological and PGD results for this couple are shown in Table 1.

**Discussion**

Originally, the main reason to introduce PGD was to avoid pregnancy termination in case of single gene diseases with severe outcome. Soon thereafter, the indications for the PGD were expanded to add late-onset diseases with high predisposition for malignancy, human leukocyte antigen typing and chromosomal aberrations. Owing to improvements in embryos selection methods, screening for aneuploidies related to age of parents became one of the main reasons for PGD along with desire to avoid transmission of inherited genetic abnormalities. Similar approach of screening embryos not only for the derivatives of abnormal segregation of translocation but also for the risk of aneuploidies of other chromosomes was proposed by some authors. The interchromosomal effect in Robertsonian carriers reproductive cells was supported by some authors, and regarded as doubtful with the need for further studies by others. Nevertheless, in the case of the male carriers with oligospermia, it seems to be crucial to analyze both aspects of the observed higher aneuploidy rates.

The first molecular, cytogenetic technique—FISH—was introduced as an easy and simple PGD procedure. Even though it has some limitations, for example, defective nucleus fixation, abnormal patterns of hybridization, and the intensity of fluorescence from simultaneously hybridized probes, it is, currently, the most frequently used screening methods. However, those examples suggest that FISH is not a completely reliable method. That is the reason why the interpretation of the results is often doubtful and requires rehybridization. Also, it may cause delay of the transfer decision both with overestimation of aneuploid blastomeres.

To perform genome-wide scans, the researchers developed comparative genomic hybridization methods. However, the technological advances directed the laboratory efforts toward the golden standard of the molecular genetics sequencing. NGS is a rapidly developing technology which produces enormous amount of data with the wide range of applications. The development of the different NGS platforms and diminishing costs enabled their introduction into the preimplantation diagnostics. The semiconductor sequencers are another step forward allowing analysis of embryos in fresh IVF cycles achieved by the faster turnaround time. In cases of screening for the abnormal segregation of the translocation, it was used to count multiple amplicons from each chromosome and compared with the relative ratio of fragments obtained from the healthy individuals. NGS-based PGD allows the precise analysis of number of all human chromosome copies in a probe by achieving high coverage of the

Table 1 The embryological and PGD results for investigated couple

<table>
<thead>
<tr>
<th></th>
<th>PGD FISH program</th>
<th>PGD NGS program</th>
</tr>
</thead>
<tbody>
<tr>
<td>Start date</td>
<td>January 2012</td>
<td>September 2013</td>
</tr>
<tr>
<td>Antral follicles count</td>
<td>17</td>
<td>10</td>
</tr>
<tr>
<td>Stimulation duration (d)</td>
<td>9</td>
<td>10</td>
</tr>
<tr>
<td>hMG total dosage (IU)</td>
<td>2,025</td>
<td>2,250</td>
</tr>
<tr>
<td>Number cumulus oocyte complexes</td>
<td>12</td>
<td>15</td>
</tr>
<tr>
<td>MII</td>
<td>6</td>
<td>11</td>
</tr>
<tr>
<td>2PN</td>
<td>4</td>
<td>5</td>
</tr>
<tr>
<td>Number of embryos biopsied on day 3</td>
<td>3</td>
<td>3</td>
</tr>
<tr>
<td>Embryos PGD results</td>
<td>1: (14q D14S1419 × 2) (D15Z4 × 2)</td>
<td>1: normal, 46, XX</td>
</tr>
<tr>
<td></td>
<td>2: (14q D14S1419 × 0) (D15Z4 × 1)</td>
<td>2: unbalanced, 45 (−22)</td>
</tr>
<tr>
<td></td>
<td>3: (14q D14S1419 × 3) (D15Z4 × 2)</td>
<td>3: unbalanced, 44, (−7, −11, −17, +21)</td>
</tr>
<tr>
<td>Total reads</td>
<td>1 embryo: 4</td>
<td>1 embryo: 51,810</td>
</tr>
<tr>
<td></td>
<td>2 embryo: 1</td>
<td>2 embryo: 73,847</td>
</tr>
<tr>
<td></td>
<td>3 embryo: 5</td>
<td>3 embryo: 70,246</td>
</tr>
<tr>
<td>Embryos transferred</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>4 wk 2 d hCG level (mIU/mL)</td>
<td>94</td>
<td>322,3</td>
</tr>
<tr>
<td>Ongoing pregnancy</td>
<td>No (miscarriage in wk 10)</td>
<td>Healthy baby born</td>
</tr>
</tbody>
</table>

Abbreviations: FISH, fluorescent in situ hybridization; hCG, human chorionic gonadotropin; hMG, human menopausal gonadotropin; MII, metaphase II; NGS, next-generation sequencing; PGD, preimplantation genetic diagnosis.
human genome sequence. Concluding, we confirm the positive impact of NGS-based PGD in the case of the Robertsonian translocation carrier. We believe that this first report of the healthy baby birth achieved after the application of new NGS technology is just a first step leading to further studies reaffirming reliability of NGS-based PGD in screening for the embryonic ploidy status.

References