

# Noninvasive chromosome screening of human embryos by genome sequencing of embryo culture medium for in vitro fertilization

Juanjuan Xu<sup>a,1</sup>, Rui Fang<sup>b,1</sup>, Li Chen<sup>a,1</sup>, Daozhen Chen<sup>b</sup>, Jian-Ping Xiao<sup>b</sup>, Weimin Yang<sup>b</sup>, Honghua Wang<sup>b</sup>, Xiaoqing Song<sup>b</sup>, Ting Ma<sup>c</sup>, Shiping Bo<sup>c</sup>, Chong Shi<sup>c</sup>, Jun Ren<sup>c</sup>, Lei Huang<sup>d,e,f,g</sup>, Li-Yi Cai<sup>b,2</sup>, Bing Yao<sup>a,2</sup>, X. Sunney Xie<sup>d,g,h,2</sup>, and Sijia Lu<sup>c,2</sup>

<sup>a</sup>Reproductive Medical Center of Nanjing Jinling Hospital and the Collaborative Innovation Platform for Reproductive Biology and Technology, Nanjing University School of Medicine, Nanjing, Jiangsu 210002, China; <sup>b</sup>Reproductive Medicine Center, Wuxi Maternity and Child Health Hospital Affiliated to Nanjing Medical University, Wuxi, Jiangsu 214002, China; <sup>c</sup>Department of Clinical Research, Yikon Genomics Company, Ltd., Shanghai 201499, China; <sup>d</sup>Biodynamic Optical Imaging Center (BIOPIIC), School of Life Sciences, Peking University, Beijing 100871, China; <sup>e</sup>Department of Obstetrics, Gynecology, and Reproductive Biology, Brigham and Women's Hospital, Boston, MA 02115; <sup>f</sup>Harvard Medical School, Boston, MA 02115; <sup>g</sup>Beijing Advanced Innovation Center for Genomics, Peking University, Beijing 100871, China; and <sup>h</sup>Department of Chemistry and Chemical Biology, Harvard University, Cambridge, MA 01238

Contributed by X. Sunney Xie, August 10, 2016 (sent for review April 28, 2016; reviewed by Eva Hoffmann and John Rasko)

Preimplantation genetic screening (PGS) is widely used to select in vitro-fertilized embryos free of chromosomal abnormalities and to improve the clinical outcome of in vitro fertilization (IVF). A disadvantage of PGS is that it requires biopsy of the preimplantation human embryo, which can limit the clinical applicability of PGS due to the invasiveness and complexity of the process. Here, we present and validate a noninvasive chromosome screening (NICS) method based on sequencing the genomic DNA secreted into the culture medium from the human blastocyst. By using multiple annealing and looping-based amplification cycles (MALBAC) for whole-genome amplification (WGA), we performed next-generation sequencing (NGS) on the spent culture medium used to culture human blastocysts ( $n = 42$ ) and obtained the ploidy information of all 24 chromosomes. We validated these results by comparing each with their corresponding whole donated embryo and obtained a high correlation for identification of chromosomal abnormalities (sensitivity, 0.882, and specificity, 0.840). With this validated NICS method, we performed chromosome screening on IVF embryos from seven couples with balanced translocation, azoospermia, or recurrent pregnancy loss. Six of them achieved successful clinical pregnancies, and five have already achieved healthy live births thus far. The NICS method avoids the need for embryo biopsy and therefore substantially increases the safety of its use. The method has the potential of much wider chromosome screening applicability in clinical IVF, due to its high accuracy and noninvasiveness.

chromosomal abnormalities | PGS | IVF | WGA | MALBAC

Human embryos are prone to chromosomal abnormalities, mainly due to age-dependent chromosomal segregation errors during meiosis I (1). Chromosomal abnormalities could cause early pregnancy loss or severe chromosomal diseases such as Down and Patau syndrome among many others (2, 3). The occurrence of chromosomal abnormalities is substantially higher in patients of advanced maternal age, patients with recurrent pregnancy loss, or those who carry chromosomal aberrations such as translocations, all of which result in poor clinical outcome for reproduction.

Chromosomal abnormalities can be prevented in in vitro fertilization (IVF) by performing preimplantation genetic screening (PGS) of all 24 chromosomes. There are various PGS methods for comprehensive chromosome screening currently in clinical use, including comparative genomic hybridization (array-CGH) (4, 5), single-nucleotide polymorphism (SNP) arrays (6–9), multiplex quantitative PCR (10), and next-generation sequencing (NGS) (11, 12). Multiple clinical trials have confirmed the clinical efficacy of PGS, including increasing implantation and clinical pregnancy rates, as well as decreasing miscarriage rates (13–16). However, the applicability of PGS has been limited for a number of reasons:

(i) PGS requires invasive embryo biopsy, which has been shown to decrease embryo quality after cleavage-stage biopsy (17); (ii) long-term biosafety of embryo biopsy in humans has not been evaluated, whereas animal studies have shown negative influences on neural and adrenal development (18–20); (iii) it involves technical expertise, requiring special training and experienced embryologists to perform the biopsy, which significantly increase the overall costs of clinical PGS cycles. Therefore, a noninvasive and easy-to-perform screening tool would greatly facilitate the widespread performing of chromosome screening before embryo implantation, thereby improving success rates.

Efforts have been made to develop noninvasive approaches for PGS (21, 22). Palini et al. (23) reported the observation of the existence of DNA in the blastocoele fluid, Gianaroli et al. (24) performed a pilot study on chromosome screening using blastocentesis, and Stigliani et al. (25, 26) observed genomic and mitochondria DNA contents in the culture medium, which were correlated with embryo quality. Wu et al. (27) reported the PCR detection of the secreted genomic DNA in the culture medium for preimplantation

## Significance

In in vitro fertilization (IVF), current methods of diagnosing chromosome abnormality and screening for viability of transfer require biopsy of embryos, which affects embryo quality, awaits long-term biosafety test, and requires specialized skills. We demonstrate the principle of noninvasive chromosome screening (NICS), which is based on sequencing the genomic DNA secreted into the culture medium from the embryo, avoiding the need for embryo biopsy and substantially increasing the safety. By characterizing its precision and demonstrating successful live births, we validate that NICS offers the potential of significantly improving the clinical outcome of IVF.

Author contributions: J.X., R.F., L.C., D.C., J.-P.X., L.-Y.C., B.Y., X.S.X., and S.L. designed research; J.X., R.F., L.C., J.-P.X., W.Y., H.W., X.S., T.M., S.B., J.R., L.H., L.-Y.C., B.Y., and S.L. performed research; R.F., L.C., T.M., S.B., C.S., J.R., L.H., L.-Y.C., X.S.X., and S.L. analyzed data; and J.X., R.F., S.B., C.S., L.H., L.-Y.C., B.Y., X.S.X., and S.L. wrote the paper.

Reviewers: E.H., University of Copenhagen; and J.R., Royal Prince Alfred Hospital, Sydney Local Health District.

Conflict of interest statement: X.S.X. and S.L. are cofounders of Yikon Genomics Company, Ltd.

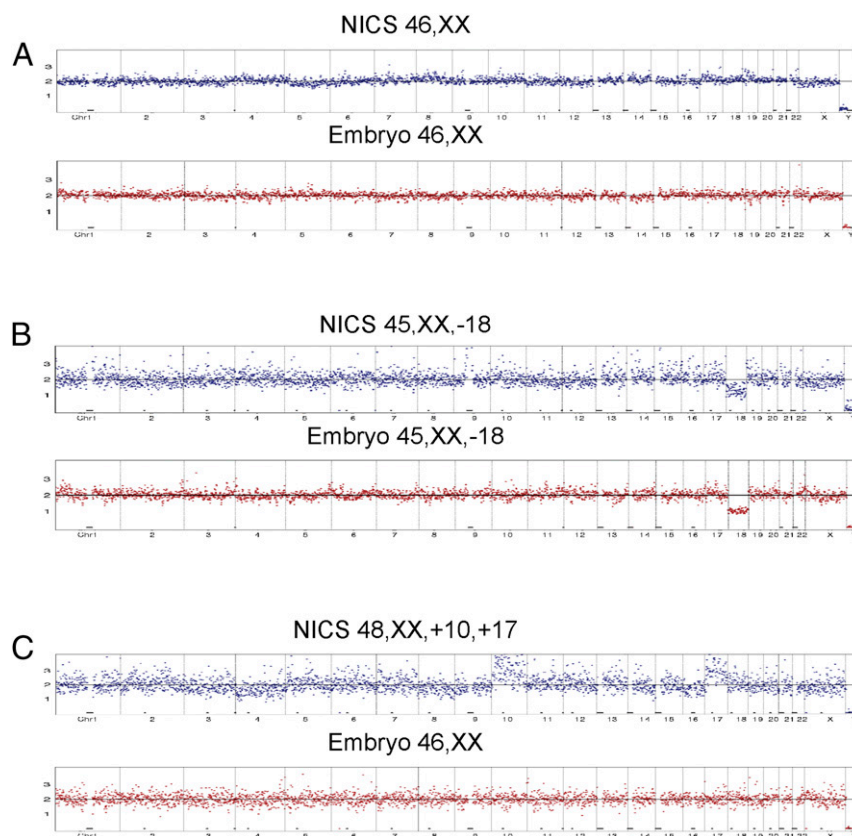
Data deposition: The data reported in this paper have been deposited in the Gene Expression Omnibus (GEO) database, [www.ncbi.nlm.nih.gov/geo](http://www.ncbi.nlm.nih.gov/geo) (accession no. SRP089980).

<sup>1</sup>J.X., R.F., and L.C. contributed equally to this work.

<sup>2</sup>To whom correspondence may be addressed. Email: [caili76@hotmail.co.jp](mailto:caili76@hotmail.co.jp), [yaobing@nju.edu.cn](mailto:yaobing@nju.edu.cn), [xie@chemistry.harvard.edu](mailto:xie@chemistry.harvard.edu), or [lusijia@yikongenomics.com](mailto:lusijia@yikongenomics.com).

This article contains supporting information online at [www.pnas.org/lookup/suppl/doi:10.1073/pnas.1613294113/-DCSupplemental](http://www.pnas.org/lookup/suppl/doi:10.1073/pnas.1613294113/-DCSupplemental).





**Fig. 2.** Examples of validation of results from the comparison of NICS versus the whole-blastocyst embryos. (A and B) Equivalent karyotype results obtained from NICS and the corresponding blastocyst embryo. Fig. 2A shows consistent results from a normal/transferrable embryo, and Fig. 2B shows consistent results from an embryo with a chromosome loss in chr18. (C) An example of inconsistent results obtained from NICS and the blastocyst embryo, with the embryo showing balanced chromosomal composition and the NICS assay identifying chromosome gains of chr10 and chr17.

**Embryo Selection by NICS on the First Patient with Balanced Translocation.** With the NICS assay validated by comparison with the voluntarily donated and IRB-approved embryos, we first applied our NICS method on a patient with a balanced translocation. IRB approval (Wuxi Maternity: 2014-04-0515-02) and informed consent were obtained before applying the NICS assay on the embryos. Karyotype analysis of the patient showed a balanced translocation  $t(14;15)(q22;q24)$ . We obtained a total of three blastocysts from this patient, and we performed NICS on D3–D5 culture medium of all three embryos. Chromosomal abnormalities were detected with the NICS assay in two of them, and therefore those could not be used for transfer (Fig. 3A and B). To confirm these results, the two embryos were collected and lysed for chromosome screening using the whole embryo, and the same results were obtained, confirming the NICS analyses (Fig. S2). Only one out of the three blastocysts showed a normal karyotype with NICS, and therefore that one was selected for transfer (Fig. 3C). A successful pregnancy resulted from this selected embryo, and a karyotype of the developing embryo was obtained by performing amniocentesis at 19 wk of gestation, confirming the karyotype results previously obtained with the NICS assay. The patient's pregnancy resulted in the live birth of a chromosomally normal and healthy baby boy on March 5, 2016.

**NICS Clinical Application Results in Successful Pregnancies.** After careful and systematic validation of the NICS assay, we have performed NICS on six more patients, in addition to the translocation patient described above. Single-blastocyst transfer was performed on all six patients, and five of these patients achieved successful pregnancies; five of these have already delivered chromosomally normal, healthy newborns, and we continue to follow up on the last currently ongoing pregnancy. Only one patient

failed to implant. The clinical indications and outcomes are summarized in Table 2.

## Discussion

**Blastocyst Biopsy and the Controversy of Extensive PGS for Patients Under IVF/ICSI Treatment.** Blastocyst trophoctoderm biopsy has been increasingly used and widely accepted in the PGS field due to its relatively low invasiveness, compared with performing blastomere biopsy at the cleavage stage. Chromosome screening on all 24 chromosomes has been mostly used on patients with advanced maternal age, recurrent pregnancy loss, repeated implantation failure, as well as on patients with abnormal karyotype such as balanced translocation and Robertsonian translocation (32–37). Extensive use of PGS on all IVF/ICSI cycles has been hotly debated in the past few years (38–41) due to the invasiveness of the biopsy procedure itself, particularly regarding the potential harm on the trophoctoderm and possible compromise of implantation potential, as well as potential concerns on long-term effects on the offspring, which are very difficult to assess. In addition, the procedure of performing blastocyst-stage biopsy requires considerable training and expertise to perform the sophisticated embryo manipulation, increasing the costs of performing PGS. Of note, the procedure we report here, which simply involves collecting embryo culture medium, requires no special expertise in embryo manipulation and therefore can be potentially used on all IVF/ICSI cycles, thereby holding promise to improve overall clinical success rates.

**False Positives and False Negatives in NICS.** We observed two false negatives (2 of 17) and four false positives (4 of 25) with the NICS assay, converting to a false-negative rate and a false-positive rate of

**Table 1. Summary of the 42 samples profiled by NICS versus their corresponding blastocysts**

21 Normal embryos (NICS and biopsy consistent)			15 Abnormal embryos (NICS and biopsy consistent)		
Sample ID	NICS	Biopsy	Sample ID	NICS	Biopsy
EM02	46,XX	46,XX	EM01	50,XX,+1,+5,+10,+13	47,XX,+4
EM04	46,XX	46,XX	EM10	45,XX,-18	45,XX,-18
EM06	46,XY	46,XY	EM11	46,XY,+5q	46,XY,-5(p12→qter,~135M,mos ~30%)
EM07	46,XY	46,XY	EM13	46,XX,-1p(pter→p21.1)	46,XX,-1p(pter→p21.1,~103M), +18q(q12.3→qter,~31M, mos)
EM09	46,XY	46,XY	EM14	45,XY,-18	46,XY,+1(p21.1→qter,~142M), -18(pter→q21.31,~56M)
EM16	46,XX	46,XX	EM15	46,XX,+1p(pter→p21.2), -18(q21.32→qter)	46,XX,+1p(pter→p21.2,~97M), -18q(q21.32→qter,~21M)
EM21	46,XY	46,XY	EM17	55,XY,+5,+6,+8,+11,+17, +19,+20,+21,+22	46,XY,+1(p21.1→qter,~143M), -18(pter→q21.31,~58M)
EM22	46,XY	46,XY	EM18	46,XX,+1p(pter→p21.1), -18(q21.32→qter)	46,XX,+1p(pter→p21.3,~100M), -18q(q21.31→qter,~22M)
EM23	46,XY	46,XY	EM19	46,XX,+14q(q23.3→qter), -15q(q26.1→qte)	46,XX,+14q(q23.1→qter,~45M), -15q(q26.1→qter,~12M)
EM24	46,XY	46,XY	EM20	46,XY,-1,+15	46,XY,-14,+15
EM25	46,XX	46,XX	EM33	52,XX,+4,+6,+9,+10,+14,+17	44,XX,-4,-14,+15(mos),-18(mos)
EM26	46,XY	46,XY	EM35	45,XX,-16	45,XX,+14(q32.12→qter,~13M),-16
EM27	46,XY	46,XY	EM37	45,XY,-22	45,XY,-22
EM28	46,XY	46,XY	EM41	50,XX,+15,+17,+18,+20	46,XX,-X(mos),+5(p12→q13.1,~23M,mos), -5q(q13.1→qter,~112M,mos)
EM29	46,XX	46,XX	EM42	46,XX,-1p,+18q	46,XX,-1p,+18q
EM30	46,XY	46,XY	4 False-positive embryos (NICS abnormal, biopsy normal)		
EM31	46,XX	46,XX	EM03	48,XX,+10,+17	46,XX
EM32	46,XX	46,XX	EM08	48,XY,+6,+18	46,XY
EM34	46,XY	46,XY	EM12	45,XY,-18	46,XY
EM36	46,XY	46,XY	EM39	47,XY,+22	46,XY
EM38	46,XX	46,XX	2 False-negative embryos (NICS normal, biopsy abnormal)		
			EM05	46,XX	45,XO
			EM40	46,XX	49,XY,+6,+8,+14

chromosomal abnormalities identification of 11.8% and 16.0%, respectively. The two false negatives might have resulted from contamination from the cumulus cells, which are maternal in origin and normally have a balanced chromosomal content (Table 1). In the future, the false-negative rate could be further minimized by carefully and thoroughly removing all cumulus cells before embryo culture.

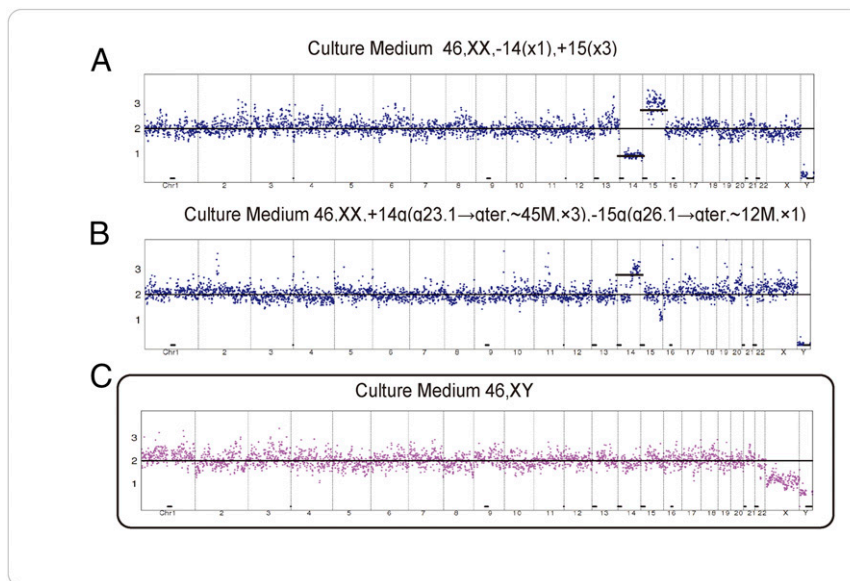
The false positives most likely arose from mosaicism. It was previously hypothesized that, during embryo development, embryos tend to exclude those cells with mitotic errors to the exterior of the embryo (42). It is therefore possible that the false-positive rate of 16.0% arose from the debris in the culture medium, originating primarily from cells eliminated from the embryo.

Although the false-positives and false-negatives could come from measurements, we note that MALBAC provides better WGA evenness and hence higher precision for CNV determination compared with degenerate oligonucleotide-primed PCR and multiple displacement amplification methods (43).

The NPV of chromosomal abnormalities with the NICS assay is 91.3%, which is substantially higher than the PPV (78.9%) of the assay. The high NPV suggests that the assay is more effective in selecting normal and transferrable embryos than identifying embryos with chromosomal abnormalities. Single-embryo transfer has been increasingly used due to its effectiveness in decreasing multiple pregnancy and miscarriage rates (15, 44). As such, we propose NICS as an additional, and risk-free, procedure for single-embryo transfer.

Regarding the clinical implications of the false positives and false negatives, we note that, in the case of all embryos detected to exhibit aneuploidy by NICS, false positives can in principle be verified by further sequencing a blastocyst biopsy on the sixth day. We are currently developing a system in which NICS can be done rapidly and results can be obtained before embryos are frozen so that a cell or two can be extracted for verification. False negatives, on the other hand, usually do not result in successful pregnancies, and hence are less problematic. Even in the case of leading to pregnancy, they can be detected and avoided by noninvasive prenatal test.

**Genome Coverage and Estimated DNA Amount in Spent Culture.** The genome coverages of the spent culture media for the three samples were determined to be 24%, 33%, and 65% by high-sequencing depth of 30× reads. It has been shown that the normal genome coverage for single diploid human cells is about ~72% with high-sequencing depth (30×) (43). Despite of our low coverage, the CNV results matched exactly their corresponding blastocyst biopsies (a few cells) on the fifth day, in either the normal or aneuploid samples (Fig. S1). Larger cell numbers usually result in saturation of the genome coverage, approaching unity. Our coverage results suggest that the loss or degradation of DNA fragments in the spent culture must occur randomly along the genome and does not affect the inference of the copy number pattern in the embryo. The fact that we observed identical copy number patterns in the spent culture samples and their corresponding embryo biopsies suggests that, under our experimental conditions, the total amount of DNA in each spent culture is equivalent to that of a fraction of a single cell.



**Fig. 3.** Embryo screening and selection using NICS from a patient carrying a balanced translocation of chr14/15. A total of three embryos successfully developed to the blastocyst stage, and D3–D5 culture medium from each embryo was collected for the NICS assay to screen for chromosomal abnormalities. (A and B) Embryos showed chromosomal abnormalities with chr14/15 and therefore could not be transferred. (C) An embryo showed balanced chromosomal composition and was therefore transferred into the uterus of the patient, resulting in a successful pregnancy and a healthy live birth.

**Limitations of the NICS Assay for Extensive Chromosome Screening in IVF/ICSI Patients.** We think the limitations of NICS could primarily derive from two aspects: (i) the requirement to be scrupulous in the removal of all cumulus-corona radiata cells (which are of maternal origin and usually with normal chromosomal composition) before performing ICSI. If such removal is not complete, residual cells may release DNA during embryo development, thereby potentially being the cause of false-negative detection. (ii) Similar to PGS, the NICS procedure would be highly recommended to be performed in conjunction with ICSI due to the difficulty of ensuring removal of any supernumerary sperm attached to the zona pellucida. Although culture medium is replaced on D3, which may decrease the likelihood of contamination due to residual cumulus cells and supernumerary sperm, all precautions should be made to reduce such contamination to a minimum if NICS is used routinely in clinical IVF. Of note, in the current stage of technological development, we consider NICS to be a screening assay for chromosomal-level abnormalities instead of being a diagnostic assay for segmental aneuploidies. More validation research is needed to identify segmental aneuploidies with the NICS assay.

In summary, our validation data and initial clinical applications strongly suggest that the NICS assay could help to improve the clinical outcome of IVF embryo selection with ploidy information, in a noninvasive manner. We envision that randomized clinical trials will be designed and performed in the near future, using the NICS assay with single-embryo transfer to evaluate the clinical effectiveness of the assay in different patient groups.

## Materials and Methods

**Embryo Preparation for the NICS Assay.** We recruited 17 patients from the Reproductive Medicine Centre of Wuxi Maternity and Child Health Hospital and the Reproductive Medical Center of Nanjing Jinling Hospital. IRB approvals (Nanjing Jinlin: 2014NZKY-005; Wuxi Maternity: 2014-04-0515-02) and informed consent were obtained before applying the NICS assay on the embryos. All donated embryos were in excess of clinical needs, and consents on donated D3 embryos were obtained for use in the comparison study. All embryos were fertilized by (ICSI). Donated D3 embryos were previously frozen by vitrification (Cryotop Safety Kit; KITAZATO BioPharma) according to the manufacturer's instructions and stored in liquid nitrogen. Vitrified embryos were warmed using warming media (Vitrification Thaw Solution; KITAZATO BioPharma). Briefly, the Cryotop strip was quickly immersed into Thawing Solution (containing 1 M sucrose) for 1 min at 37 °C. Embryos were then transferred into dilution solution (containing 0.5 M sucrose) and incubated for 3 min, followed by incubation in two 80- $\mu$ L droplets of Washing Solution for 3 min each at room temperature (25 °C).

**Blastocyst Culture and Transfer.** Warmed D3 embryos were placed in 30- $\mu$ L droplets of Quinn's Advantage Protein Plus blastocyst medium (SAGE) containing washed and pregassed mineral oil (SAGE), and they were then further cultured to the blastocyst stage under 5.5% CO<sub>2</sub>, 5% O<sub>2</sub>, and balance N<sub>2</sub> at 37 °C in Labotect C16 incubators (Labotect). After 2 d in culture, the development and quality of the blastocysts were evaluated according to the blastocyst scoring system. A single blastocyst was selected for transfer to each patient based on the NICS results.

**Sample Collection.** To prevent media cross-contamination, different Pasteur pipettes were used for each embryo. Five to 20  $\mu$ L of blastocyst medium from each embryo was transferred into RNase–DNase-free PCR tubes containing

**Table 2.** Clinical outcome of the first seven patients subjected to NICS

Patient no.	Maternal age	Clinical indications	Transfer cycles	Clinical outcome
P01	30	Reciprocal translocation 46,XY,t(14;15)	1	Singleton pregnancy—live birth
P02	28	Azoospermia	1	Singleton pregnancy—live birth
P03	34	Inversion 46,XY,inv(9)	1	Singleton pregnancy—live birth
P04	32	Reciprocal translocation 46,XX,t(1;18)	2	Implantation failure
P05	26	Recurrent pregnancy loss	1	Singleton pregnancy—live birth
P06	32	47,XY	2	Singleton pregnancy—live birth
P07	29	Recurrent implantation failure	1	Singleton pregnancy—following up

5  $\mu$ L of cell lysis buffer (Yikon Genomics). As a negative control, we collected the same amount of blastocyst culture medium but without its being used for embryo culture. All collected samples were frozen immediately in liquid nitrogen and stored at  $-80^{\circ}\text{C}$  until subjected to the NICS assay.

**Embryo Selection by NICS on a Patient with Balanced Translocation.** The patient couple for NICS treatment in Wuxi Maternity reproduction center had 3 y of infertility history. Their chromosome examination showed that the male partner carries a t(14;15)(q22;q24) translocation, and the female has a normal karyotype. We collected five eggs in the IVF cycle (*SI Materials and Methods*); four MII oocytes were fertilized by ICSI, and three of them developed to the blastocyst stage. All blastocysts were frozen with vitrification freezing protocol after performing NICS (*SI Materials and Methods*). We obtained normal karyotype from one embryo with the NICS assay (Fig. 3C). We therefore transferred this normal embryo and confirmed clinical pregnancy with enhanced  $\beta$ -HCG followed up by ultrasound. The result of fetal chromosome examination in the amniotic fluid at 19 wk of gestation confirmed the balanced chromosome results previously obtained by NICS. After obtaining informed consent from the patients, the two embryos with chromosome imbalance, according to our NICS results, were lysed and sequenced for validation purposes (Fig. 1).

- Battaglia DE, Goodwin P, Klein NA, Soules MR (1996) Influence of maternal age on meiotic spindle assembly in oocytes from naturally cycling women. *Hum Reprod* 11(10):2217–2222.
- Munné S (2006) Chromosome abnormalities and their relationship to morphology and development of human embryos. *Reprod Biomed Online* 12(2):234–253.
- Korenberg JR, et al. (1994) Down syndrome phenotypes: The consequences of chromosomal imbalance. *Proc Natl Acad Sci USA* 91(11):4997–5001.
- Hellani A, Abu-Amero K, Azouri J, El-Akoum S (2008) Successful pregnancies after application of array-comparative genomic hybridization in PGS-aneuploidy screening. *Reprod Biomed Online* 17(6):841–847.
- Gutiérrez-Mateo C, et al. (2011) Validation of microarray comparative genomic hybridization for comprehensive chromosome analysis of embryos. *Fertil Steril* 95(3):953–958.
- Thornhill AR, et al. (2015) Karyomapping—a comprehensive means of simultaneous monogenic and cytogenetic PGD: Comparison with standard approaches in real time for Marfan syndrome. *J Assist Reprod Genet* 32(3):347–356.
- Natesan SA, et al. (2014) Live birth after PGD with confirmation by a comprehensive approach (karyomapping) for simultaneous detection of monogenic and chromosomal disorders. *Reprod Biomed Online* 29(5):600–605.
- Treff NR, et al. (2011) Single nucleotide polymorphism microarray-based concurrent screening of 24-chromosome aneuploidy and unbalanced translocations in preimplantation human embryos. *Fertil Steril* 95(5):1606–1612.e1,2.
- Natesan SA, et al. (2014) Genome-wide karyomapping accurately identifies the inheritance of single-gene defects in human preimplantation embryos in vitro. *Genet Med* 16(11):838–845.
- Treff NR, et al. (2012) Development and validation of an accurate quantitative real-time polymerase chain reaction-based assay for human blastocyst comprehensive chromosomal aneuploidy screening. *Fertil Steril* 97(4):819–824.
- Martin J, et al. (2013) The impact of next-generation sequencing technology on preimplantation genetic diagnosis and screening. *Fertil Steril* 99(4):1054–1061.e3.
- Hou Y, et al. (2013) Genome analyses of single human oocytes. *Cell* 155(7):1492–1506.
- Scott RT, Jr, et al. (2013) Blastocyst biopsy with comprehensive chromosome screening and fresh embryo transfer significantly increases in vitro fertilization implantation and delivery rates: A randomized controlled trial. *Fertil Steril* 100(3):697–703.
- Forman EJ, et al. (2013) In vitro fertilization with single euploid blastocyst transfer: A randomized controlled trial. *Fertil Steril* 100(1):100–107.e1.
- Yang Z, et al. (2012) Selection of single blastocysts for fresh transfer via standard morphology assessment alone and with array CGH for good prognosis IVF patients: Results from a randomized pilot study. *Mol Cytogenet* 5(1):24.
- Keltz MD, et al. (2013) Preimplantation genetic screening (PGS) with comparative genomic hybridization (CGH) following day 3 single cell blastomere biopsy markedly improves IVF outcomes while lowering multiple pregnancies and miscarriages. *J Assist Reprod Genet* 30(10):1333–1339.
- Cimadomo D, et al. (2016) The impact of biopsy on human embryo developmental potential during preimplantation genetic diagnosis. *BioMed Res Int* 2016:7193075.
- Wu Y, et al. (2014) Blastomere biopsy influences epigenetic reprogramming during early embryo development, which impacts neural development and function in resulting mice. *Cell Mol Life Sci* 71(9):1761–1774.
- Zhao H-C, et al. (2013) Aberrant epigenetic modification in murine brain tissues of offspring from preimplantation genetic diagnosis blastomere biopsies. *Biol Reprod* 89(5):117.
- Zeng Y, et al. (2013) Preimplantation genetic diagnosis (PGD) influences adrenal development and response to cold stress in resulting mice. *Cell Tissue Res* 354(3):729–741.
- Assou S, Ait-Ahmed O, El Messaoudi S, Thierry AR, Hamamah S (2014) Non-invasive preimplantation genetic diagnosis of X-linked disorders. *Med Hypotheses* 83(4):506–508.
- Galluzzi L, et al. (2015) Extracellular embryo genomic DNA and its potential for genotyping applications. *Future Science OA* 1(4):FSO62.
- Palini S, et al. (2013) Genomic DNA in human blastocoele fluid. *Reprod Biomed Online* 26(6):603–610.
- Gianaroli L, et al. (2014) Blastocentesis: A source of DNA for preimplantation genetic testing. Results from a pilot study. *Fertil Steril* 102(6):1692–1699.e6.
- Stigliani S, Anserini P, Venturini PL, Scaruffi P (2013) Mitochondrial DNA content in embryo culture medium is significantly associated with human embryo fragmentation. *Hum Reprod* 28(10):2652–2660.
- Stigliani S, et al. (2014) Mitochondrial DNA in day 3 embryo culture medium is a novel, non-invasive biomarker of blastocyst potential and implantation outcome. *Mol Hum Reprod* 20(12):1238–1246.
- Wu H, et al. (2015) Medium-based noninvasive preimplantation genetic diagnosis for human  $\alpha$ -thalassemias-SEA. *Medicine (Baltimore)* 94(12):e669.
- Zong C, Lu S, Chapman AR, Xie XS (2012) Genome-wide detection of single-nucleotide and copy-number variations of a single human cell. *Science* 338(6114):1622–1626.
- Huang J, et al. (2014) Validation of multiple annealing and looping-based amplification cycle sequencing for 24-chromosome aneuploidy screening of cleavage-stage embryos. *Fertil Steril* 102(6):1685–1691.
- Yan L, et al. (2015) Live births after simultaneous avoidance of monogenic diseases and chromosome abnormality by next-generation sequencing with linkage analyses. *Proc Natl Acad Sci USA* 112(52):15964–15969.
- Huang J, et al. (2016) Validation of a next-generation sequencing-based protocol for 24-chromosome aneuploidy screening of blastocysts. *Fertil Steril* 105(6):1532–1536.
- Gui B, et al. (2016) Chromosomal analysis of blastocysts from balanced chromosomal rearrangement carriers. *Reproduction* 151(4):455–464.
- Schoolcraft W, et al. (2012) Comprehensive chromosome screening (CCS) with vitrification results in improved clinical outcome in women >35 years: A randomized control trial. *Fertil Steril* 98(3):51.
- Hodes-Wertz B, et al. (2012) Idiopathic recurrent miscarriage is caused mostly by aneuploid embryos. *Fertil Steril* 98(3):675–680.
- Margalioth EJ, Ben-Chetrit A, Gal M, Eldar-Geva T (2006) Investigation and treatment of repeated implantation failure following IVF-ET. *Hum Reprod* 21(12):3036–3043.
- Frydman N, et al. (2001) Assisting reproduction of infertile men carrying a Robertsonian translocation. *Hum Reprod* 16(11):2274–2277.
- Fragouli E, et al. (2010) Comprehensive chromosome screening of polar bodies and blastocysts from couples experiencing repeated implantation failure. *Fertil Steril* 94(3):875–887.
- Chen M, Wei S, Hu J, Quan S (2015) Can comprehensive chromosome screening technology improve IVF/ICSI outcomes? A meta-analysis. *PLoS One* 10(10):e0140779.
- Twisk M, et al. (2008) No beneficial effect of preimplantation genetic screening in women of advanced maternal age with a high risk for embryonic aneuploidy. *Hum Reprod* 23(12):2813–2817.
- Hardarson T, et al. (2008) Preimplantation genetic screening in women of advanced maternal age caused a decrease in clinical pregnancy rate: A randomized controlled trial. *Hum Reprod* 23(12):2806–2812.
- Schoolcraft WB, Katz-Jaffe MG, Stevens J, Rawlins M, Munne S (2009) Preimplantation aneuploidy testing for infertile patients of advanced maternal age: A randomized prospective trial. *Fertil Steril* 92(1):157–162.
- Taylor TH, et al. (2014) The origin, mechanisms, incidence and clinical consequences of chromosomal mosaicism in humans. *Hum Reprod Update* 20(4):571–581.
- Huang L, Ma F, Chapman A, Lu S, Xie XS (2015) Single-cell whole-genome amplification and sequencing: Methodology and applications. *Annu Rev Genomics Hum Genet* 16:79–102.
- Ryan GL, et al. (2007) A mandatory single blastocyst transfer policy with educational campaign in a United States IVF program reduces multiple gestation rates without sacrificing pregnancy rates. *Fertil Steril* 88(2):354–360.
- Bolger AM, Lohse M, Usadel B (2014) Trimmomatic: A flexible trimmer for Illumina sequence data. *Bioinformatics* 30(15):2114–2120.
- Li H, Durbin R (2009) Fast and accurate short read alignment with Burrows–Wheeler transform. *Bioinformatics* 25(14):1754–1760.