

Ο Προεμφυτευτικός Έλεγχος στην Ανδρική Υπογονιμότητα



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SEmen ANALYSIS

Volume

According to WHO (Cooper et al., 2010)

Normal Volume \geq 1.5ml

Sperm count

Normal count $\geq 15 \times 10^6/\text{ml}$

Motility

Linear Progression $\geq 32\%$

Morphology

Normal Morphology $\geq 4\%$

pH

Liquefaction

Sperm DNA Fragmentation

Sperm Molecular Cytogenetic Analysis

Karyotype

Y chromosome microdeletions

CFTR mutations



Male infertility and sperm aneuploidy

1-2% of infertile males have Obstructive Azoospermia.

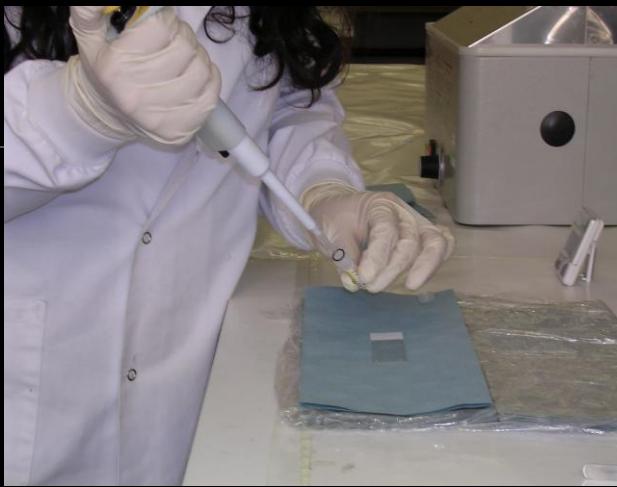
90% of cases are due to CFTR mutations. There are more than 1500 CFTR mutations worldwide >100 identified in the Greek population.

In general the lower the sperm count..... the higher the incidence of sperm aneuploidy.

Men with $<5 \times 10^6/\text{ml}$ are 10x more likely to have increased sperm aneuploidy.

Men with NOA are at a high risk of having sperm with increased aneuploidy (especially of sex chromosomes). *Clementini et al 2005 & Vincent et al 2002.*

Molecular cytogenetic analysis by FISH



Addition of
Fluorescent Probe



Placement of
Coverslip



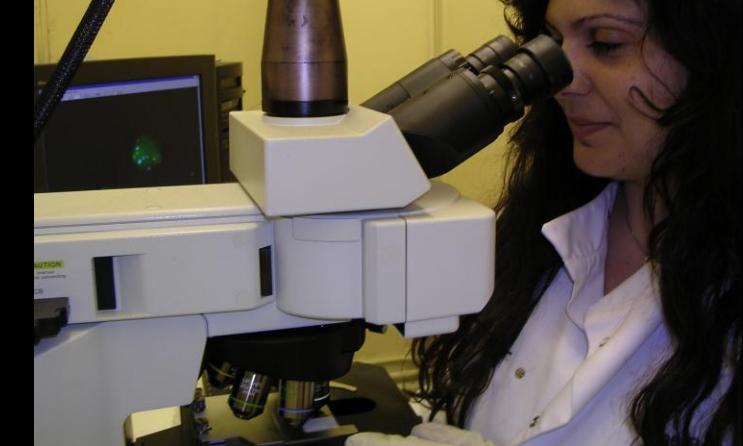
DNA Denaturation at 75°C
Hybridisation at 37°C



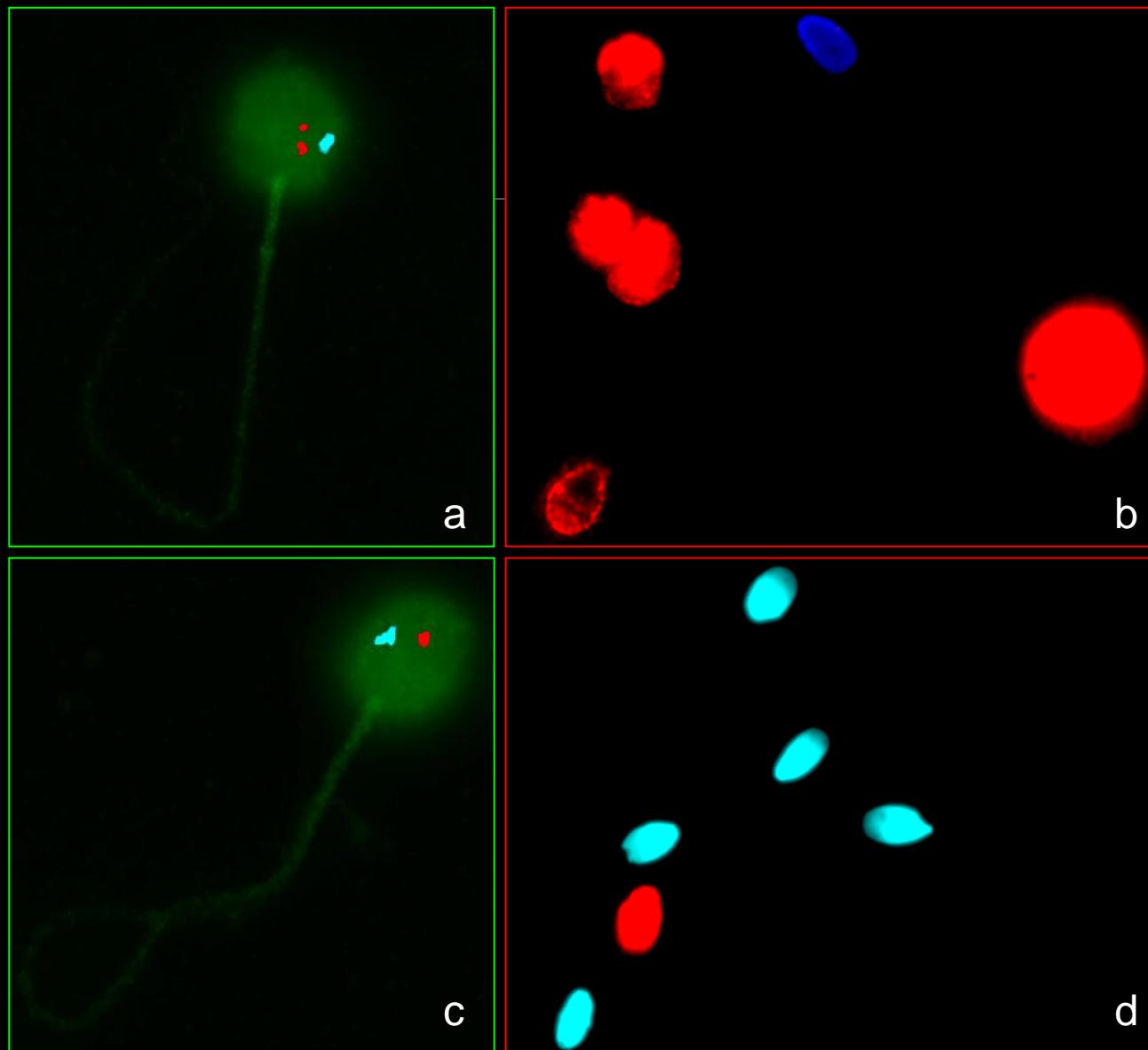
Washes



Fluorescence Microscopy



Molecular cytogenetic analysis by FISH and DFI by Tunnel



Photomicrographs showing sperm analysis by FISH and TUNEL in a patient with OAT and RIF (a-b) and a patient with normal sperm parameters (c-d). (a,c) sperm hybridised with a probe for chromosomes X-green, Y-red, 18-aqua showing disomy for chromosome Y (a) and normal haploid chromosomal constitution (c). (b, d) Photomicrographs showing Tunel labeled sperm. Note that in (b) there is only one normal spermatozoon (blue) while all the remaining are fragmented (red). In contrast, all the sperm except one (red) are normal in (d).

Molecular cytogenetic analysis by FISH

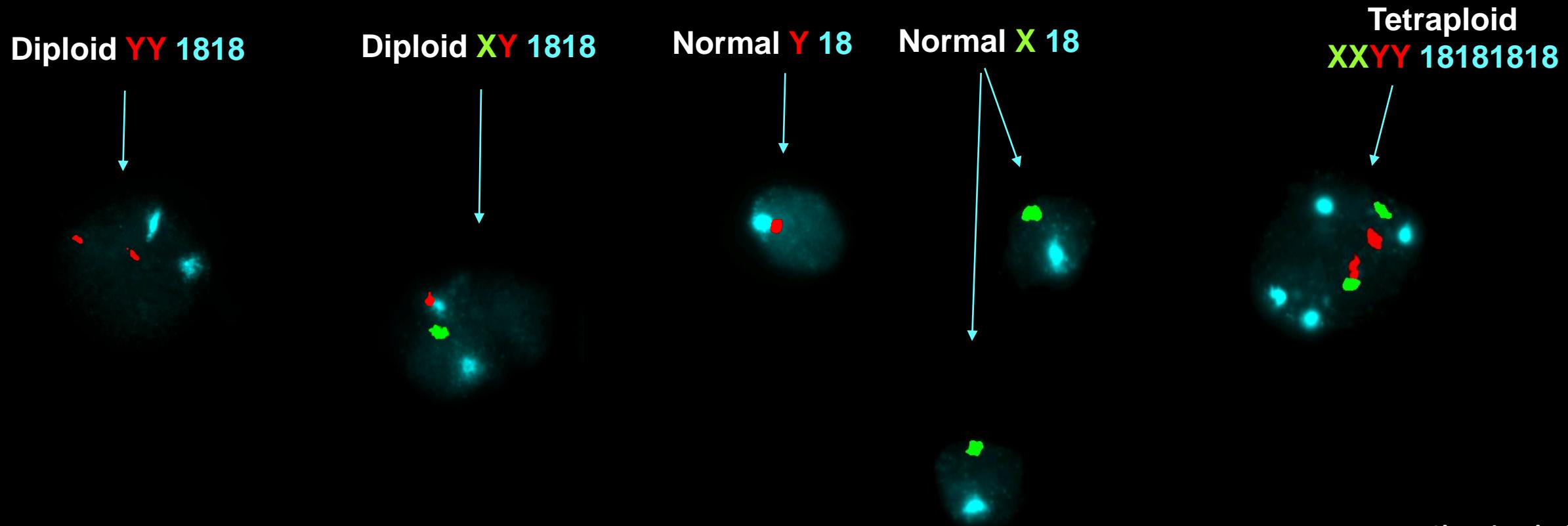
Probe for chromosomes X,Y,15

Diploid XX 1515



Dr K. Chatzimeletiou

Molecular cytogenetic analysis by FISH using a triple-colour probe for chromosomes X,Y,18



Dr K. Chatzimeletiou

Hydatidiform Moles

Pregnancies characterized by abnormal development of extraembryonic and embryonic tissues. Subclassified on the basis of ploidy, histopathology and clinical features as complete or partial.

Complete hydatidiform moles

No foetal tissues
Focal trophoblastic hyperplasia
with villous hydrops

Diploid (46 chromosomes)
but entirely androgenetic

They result from fertilisation of an inactivated [egg](#) by two spermatozoa, or more frequently by a single diploid sperm, or a haploid sperm which has undergone chromatid endoreduplication.

Partial hydatidiform moles (HM)

Identifiable foetal tissues
Focal trophoblastic hyperplasia
with villous hydrops

triploid (69 chromosomes)
23 maternal + 46 paternal

They arise usually from fertilization of a normal egg by two spermatozoa or by a diploid spermatozoon.

2430.e1 Fertility and Sterility® Vol. 95, No. 7, June 2011

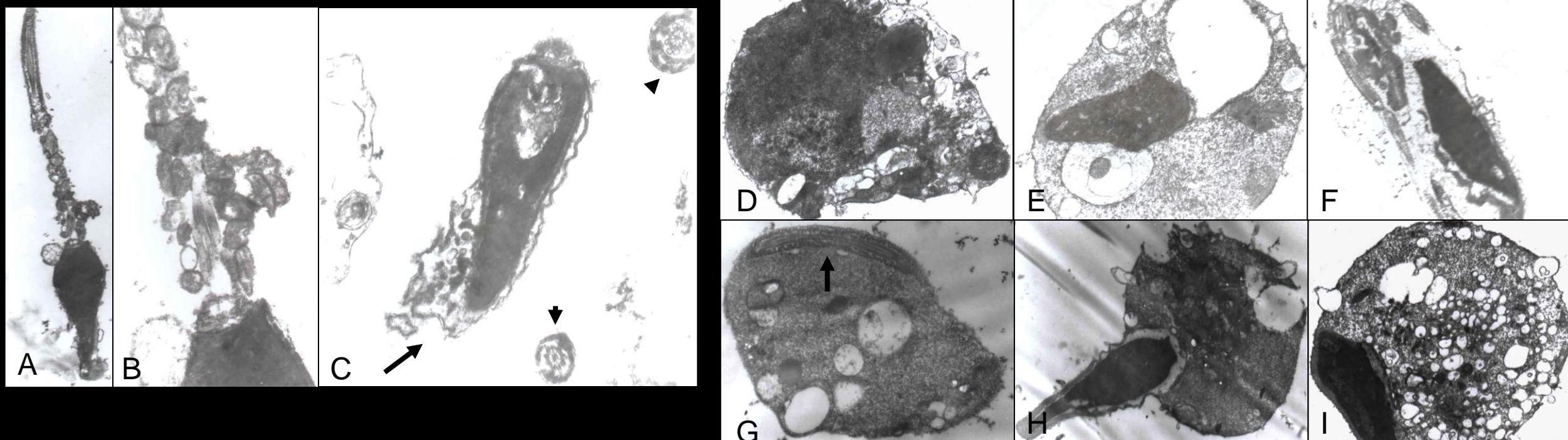
Semen analysis by electron and fluorescence microscopy in a case of partial hydatidiform mole reveals a high incidence of abnormal morphology, diploidy, and tetraploidy

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| Standard semen analysis. | |
|------------------------------|-------------------------------|
| Measurement | Value |
| Volume | 3 mL |
| pH | 8.5 |
| No. | $61.37 \times 10^6/\text{mL}$ |
| Total number | 184.125/ejaculation |
| Motility, % | |
| Linear progression | 31 |
| No progression - tail moving | 30 |
| Immotile | 39 |
| Morphology | |
| Normal, % | 4 |
| Abnormal, % | 96 |
| Big head | 3 |
| Small head | 3 |
| Long head | 4 |
| Pear-shaped head | 27 |
| Round head | 1 |
| Amorphous head | 18 |
| Vacuoles | 36 |
| Small acrosome | 3 |
| Short tail | 1 |
| Double tail | 1 |
| Fourchette | 3 |
| Broken tail | 1 |
| Spiral tail | 9 |
| Asymmetric tail extrusion | 1 |
| Broken neck | 12 |
| Cytoplasmic droplet | 14 |
| Thick midpiece | 34 |
| White blood cells | 5% |
| Round spermatids | 8% |

SEmen ANALYSIS BY TRANSMISSION ELECTRON MICROSCOPY

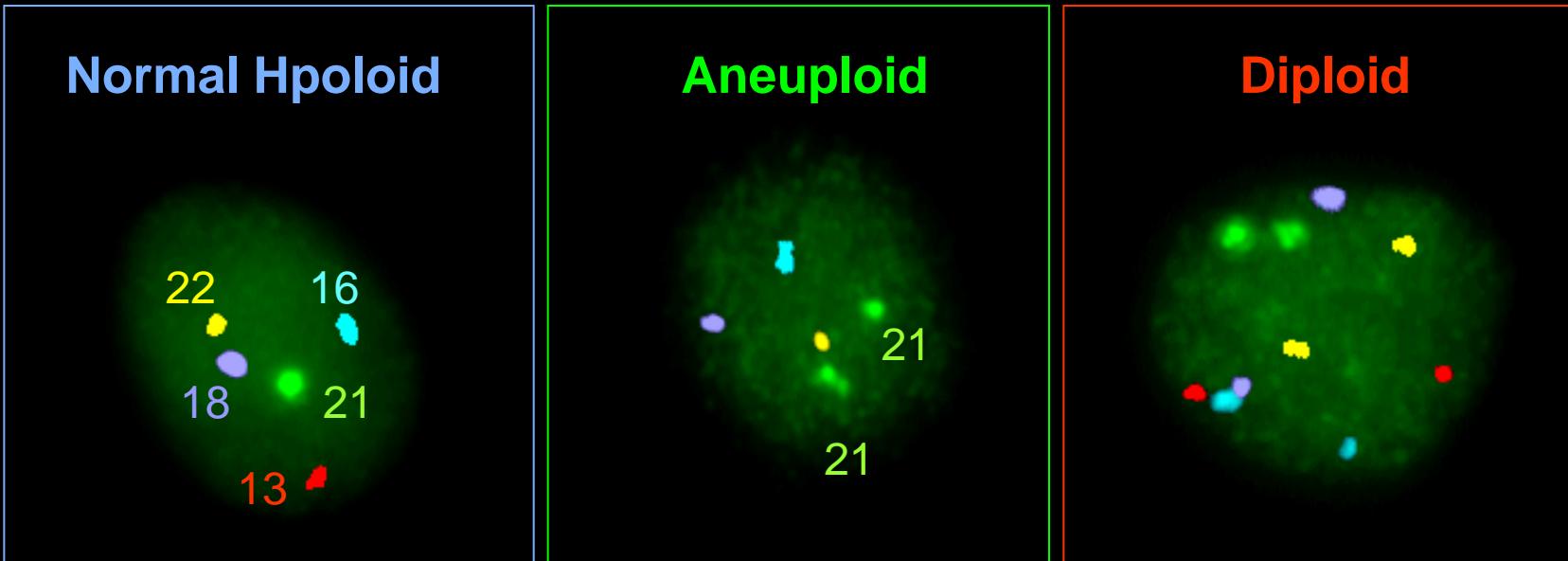


Chatzimeletiou et al 2011 Fertility and Sterility 95(7):2430.e1-5.

Incidence of chromosomal abnormalities in sperm assessed by FISH (total number of sperm analyzed in each probe set used = 400).

| | Haploid, n (%) | Aneuploid, n (%) | Diploid, n (%) | Tetraploid, n (%) |
|---|----------------|------------------|---------------------------|-------------------|
| Probe: Multivysion PB Chromosomes 13, 16, 18, 21, 22 | 373 (93.25) | 5 (1.25) | 21 (5.25) | 1 (0.25) |
| Probe: CEPX/Y/18 Chromosomes X, Y, 18 | 374 (93.5) | 4 (1.0) | 6 (XY), 14 (XX,YY) (5.0) | 2 (0.5) |
| Probe: CEPX/Y/15 Chromosomes X, Y, 15 | 375 (93.75) | 2 (0.5) | 7 (XY), 15 (XX, YY) (5.5) | 1 (0.25) |

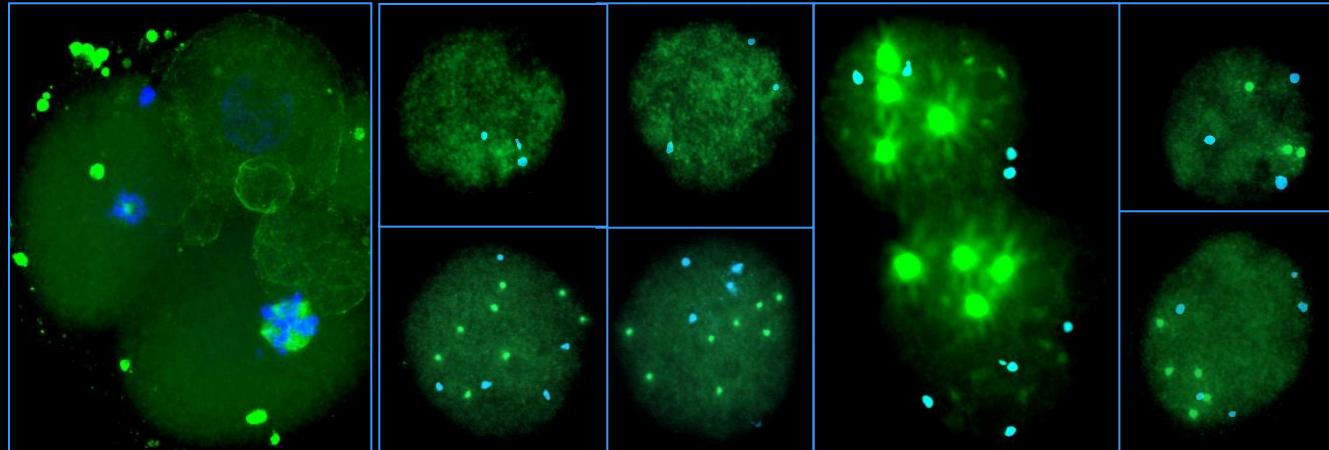
Chatzimeletiou. Semen analysis in a case of partial mole. Fertil Steril 2011.



Chatzimeletiou et al 2011 Fertility and Sterility 95(7):2430.e1-5.

Indications for Preimplantation Genetic Testing for aneuploidy (PGT-A) due to male factor

1. Previous history of aneuploid pregnancies due to increased sperm aneuploidy
2. Previous history of molar pregnancies due to increased sperm diploidy
3. Some cases of severe male infertility (OAT)



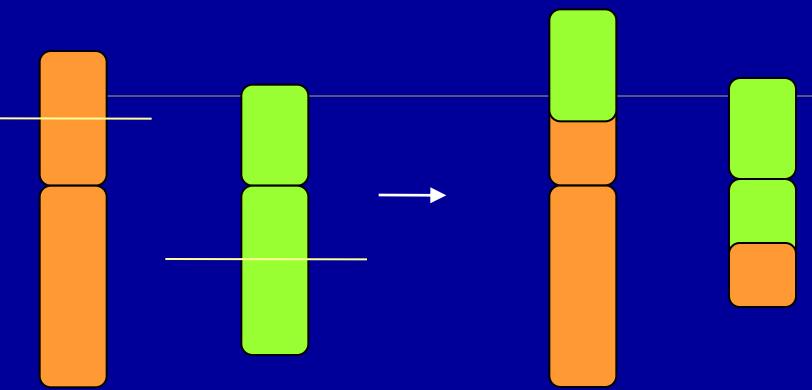
Chatzimeletiou et al., 2007 Zygote 15: 81-90

Chatzimeletiou et al., 2005 Hum. Reprod. 20 (3): 672-682

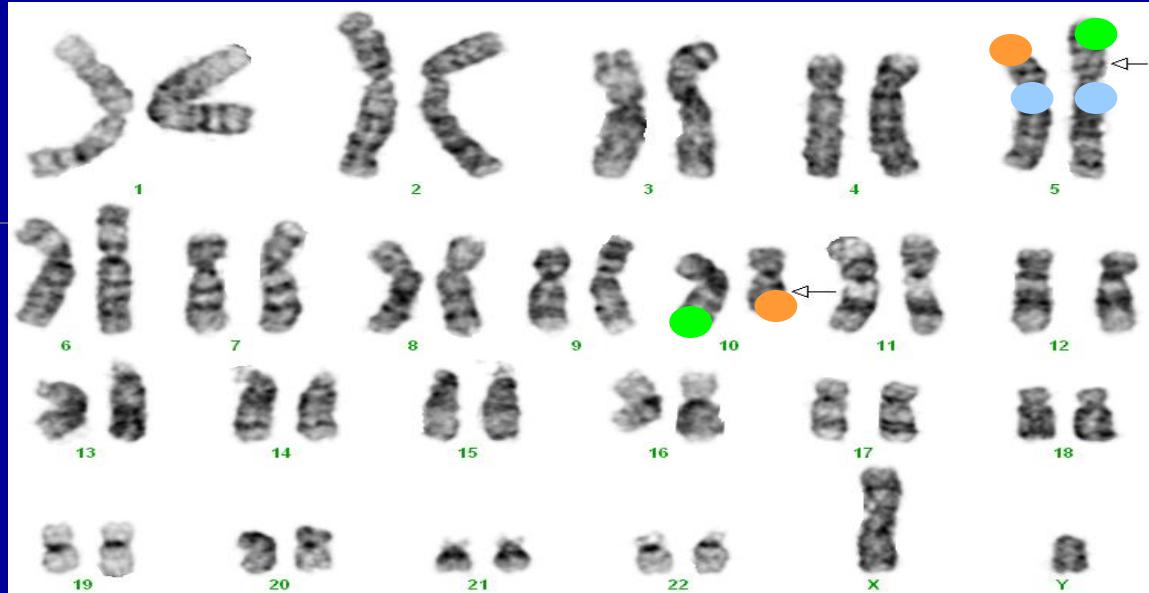
4. Males with Robertsonian or Reciprocal Translocations
5. Males with XXY or XYY Karyotype

Balanced translocations

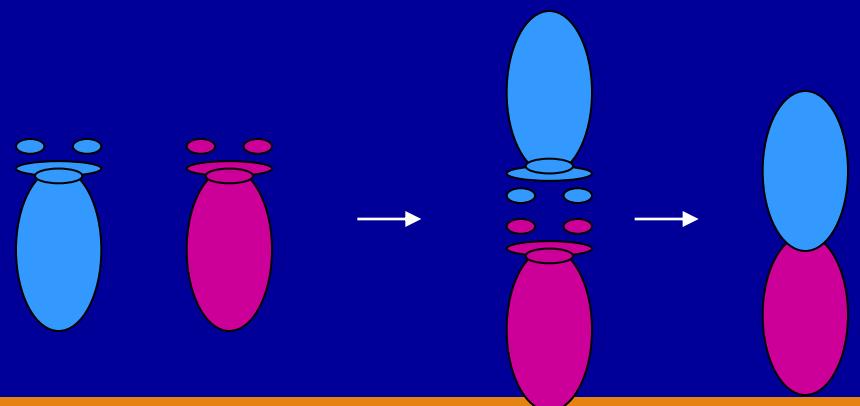
Reciprocal translocations 1:600 livebirths



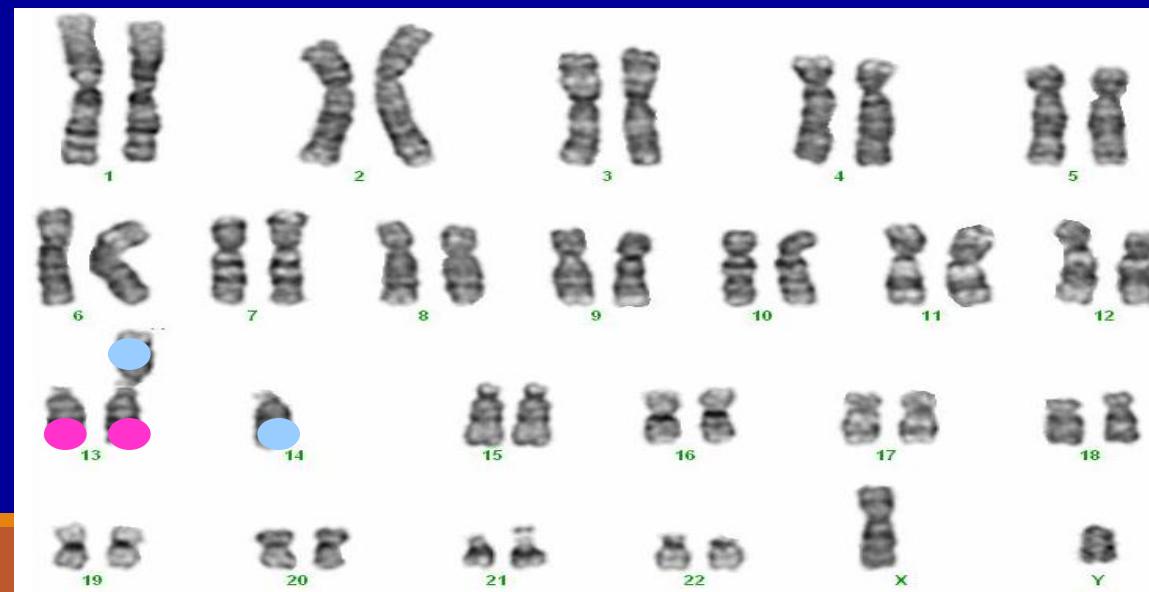
46,XY,t(5;10)(p14.2;q22.1)



Robertsonian translocations 1:1000 livebirths

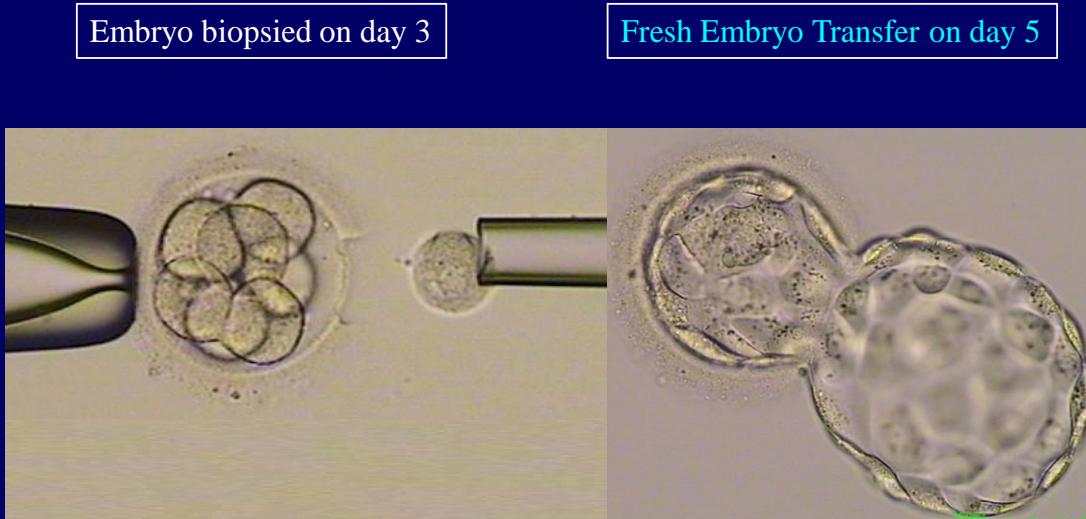


45,XY der (13;14)(q10;q10)



Preimplantation Genetic Testing for aneuploidy (PGT-A) due to male factor

Day 3 Embryo Biopsy

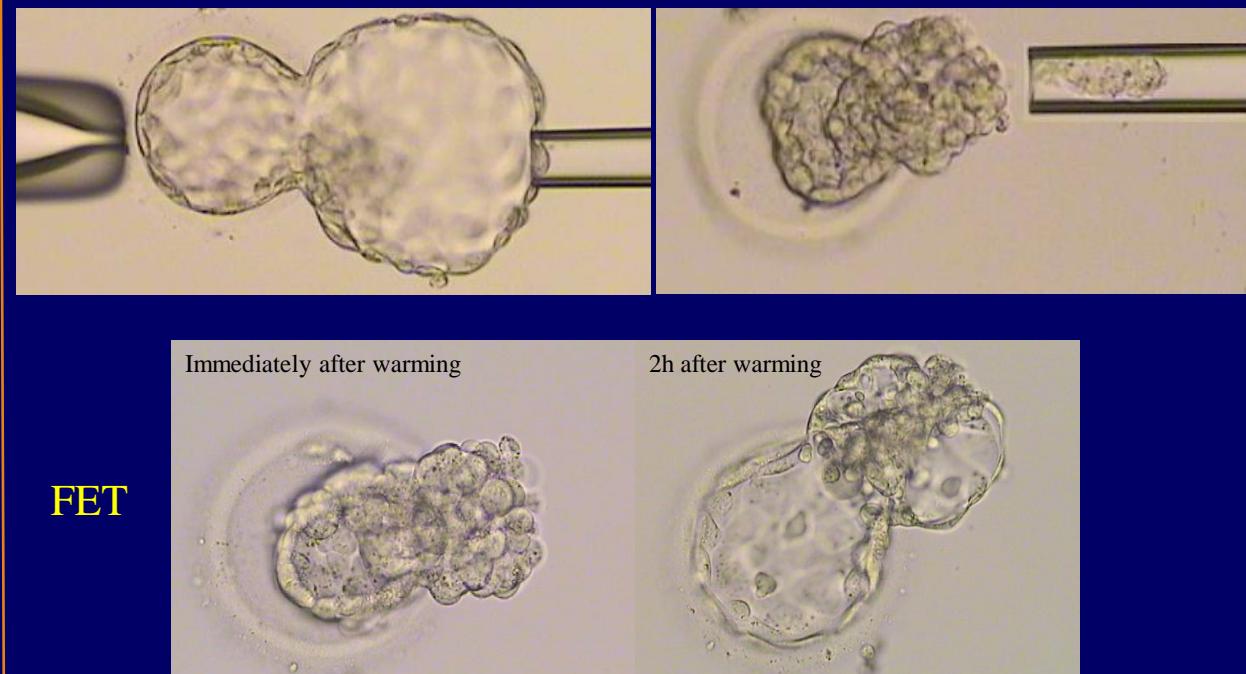


K. Chatzimeletiou Biogenesis IVF Unit

Day 5 Blastocyst Biopsy

Embryo biopsied on day 5

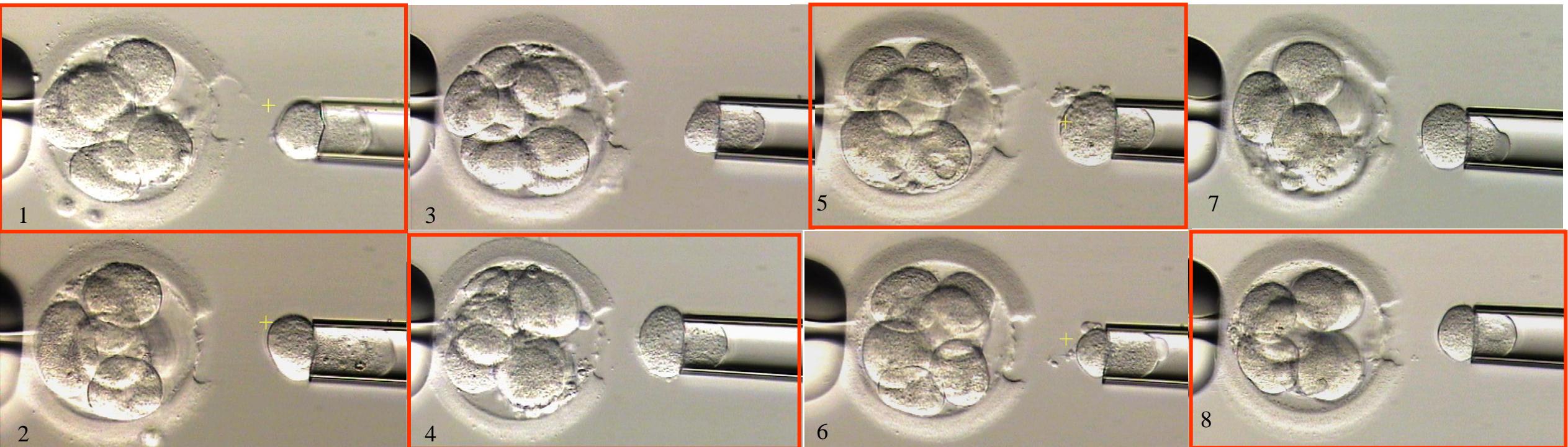
Vitrification on day 5



K. Chatzimeletiou Biogenesis IVF Unit

Preimplantation Genetic Testing for aneuploidy (PGT-A)

For the indication of Paternal Robertsonian Translocation 45,XY der (13;14)(q10;q10)

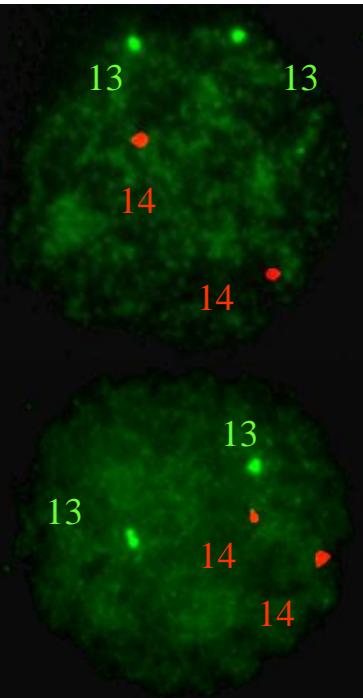


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PREIMPLANTATION GENETIC DIAGNOSIS

For the indication of Paternal Robertsonian Translocation 45,XY der (13;14)(q10;q10)

2 normal embryos were transferred



↓
Twin Pregnancy



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PREIMPLANTATION GENETIC DIAGNOSIS

For the indication of Paternal Reciprocal Translocation 46,XY t (3;16)(p28.2;p26.4)

ΓΕΝΙΚΟ ΠΕΡΙΦΕΡΕΙΑΚΟ ΝΟΣΟΚΟΜΕΙΟ ΠΑΠΑΓΕΩΡΓΙΟΥ

ΕΡΓΑΣΤΗΡΙΟ ΠΡΟΕΜΦΥΤΕΥΤΙΚΗΣ ΓΕΝΕΤΙΚΗΣ ΔΙΑΓΝΩΣΗΣ
ΜΟΝΑΔΑ ΑΝΘΡΩΠΙΝΗΣ ΑΝΑΠΑΡΑΓΩΓΗΣ
Α' ΜΑΙΕΥΤΙΚΗ & ΓΥΝΑΙΚΟΛΟΓΙΚΗ ΚΛΙΝΙΚΗ Α.Π.Θ.
Διευθυντής Κλινικής: Β. Κ. Ταρλαζής Υπεύθυνη Εργαστηρίου: Κ. Χατζημελετίου
Τηλ. 231332 3827, 231332 3377

Προεμφυτευτική Γενετική Διάγνωση

'Όνοματεπώνυμο ♀: [REDACTED] Νο♀: [REDACTED]

ΗΩ: [REDACTED] Ηλικία♀: 33 Ένδειξη: 46,XY t(3 ; 16)(p ; p)

Ανιχνευτές: CEP 16 (aqua), Telomeric 16p (green) , 3p (orange) (Abbott, Cytocell)

Ωάρια : 5 MII Ωάρια : 5 2pn Ζυγωτά : 3 Διαιρούμενα έμβρυα: 3

Τρία (3) φυσιολογικά γονιμοποιημένα (2pn) έμβρυα, διαιρέθηκαν και υποβλήθηκαν σε βιοψία την ημέρα 3 μετά τη γονιμοποίηση, στο στάδιο των 8 κυττάρων (ε2 grade 2.1, ε4 grade 2.2, ε5 grade 2.2).
Ένα βλαστομερίδιο απομονώθηκε από κάθε έμβρυο, και πυρήνες ανιχνεύτηκαν σε όλα τα βλαστομερίδια.
Τα αποτελέσματα της μοριακής κυτταρογενετικής ανάλυσης έδειξαν ότι:
Το έμβρυο 2 ήταν ισοζυγισμένο (2x3p, 2x16cep, 2x16p).
Το έμβρυο 4 ήταν ανισόζυγο/ανευπλοιοειδές (2x3p, 3x16cep, 3x16p).
Το έμβρυο 5 ήταν ισοζυγισμένο (2x3p, 2x16cep, 2x16p).
Προτείνονται τα έμβρυα 2 και 5 για μεταφορά στη μήτρα.

| Έμβρυο Νο | ΧΡΩΜΟΣΩΜΙΚΗ ΚΑΤΑΣΤΑΣΗ (3p, 16cep, 16p) | ΕΜΒΡΥΟΜΕΤΑΦΟΡΑ |
|-----------|---|----------------|
| 2 | Ισοζυγισμένο | ΝΑΙ |
| 4 | Ανισόζυγο | ΟΧΙ |
| 5 | Ισοζυγισμένο | ΝΑΙ |
| | | |

Έμβρυολόγος- Γενετιστής: Κατερίνα Χατζημελετίου B.Sc. (Hons), M.Sc., Ph.D.



2/11/2012

Development of the 2 normal embryos transferred



↓
Twin Girls born
free of translocation

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Summary of PGT-A results for male factor at Papageorgiou General Hospital

| | |
|-----------------------------|-------|
| Total No of eggs collected | 229 |
| No of MII eggs | 188 |
| No of 2PN zygotes | 138 |
| No of embryos biopsied | 103 |
| No of embryos analysed | 100 |
| No of normal embryos | 41 |
| No of abnormal embryos | 59 |
| +ve hCG/ ET | 73.3% |
| Clinical pregnancy rate/ ET | 60.0% |

Next generation sequencing (NGS) for preimplantation genetic screening (PGS) improves pregnancy outcomes compared with array comparative genomic hybridization in single thawed euploid embryo transfer (STEET) cycles

Friedenthal et al (2018) Fertil Steril. 109(4):627-632

548 STEET cycles using NGS vs 368 STEET cycles using array-CGH

The implantation rate was significantly higher in the NGS group compared with the aCGH group (71.6% vs. 64.6%). The OP/LBR was also significantly higher in the NGS group (62% vs. 54.4%), and there were significantly more biochemical pregnancies in the aCGH group compared with the NGS group (15.1% vs. 8.7%).

ΣΥΜΠΕΡΑΣΜΑΤΑ

Ο Προεμφυτευτικός Γενετικός Έλεγχος χρησιμοποιείται σε συγκεκριμένες περιπτώσεις ανδρικού παράγοντα υπογονιμότητας:

- 1. Αυξημένη Ανευπλοειδία ή/και διπλοειδία στο σπέρμα**
- 2. OAT**
- 3. Καρυότυπο με Robertsonian και Reciprocal Μεταθέσεις**
- 4. Καρυότυπο με XXY και XYY**

ΣΥΜΠΕΡΑΣΜΑΤΑ

Ο Προεμφυτευτικός Γενετικός Έλεγχος συγκαταλέγεται σήμερα στις σημαντικότερες τεχνολογικές εξελίξεις στον τομέα της ανθρώπινης αναπαραγωγής συμβάλλοντας ουσιαστικά στην πρόληψη των γενετικών και χρωμοσωμικών ασθενειών και στη γέννηση νεογνών απαλλαγμένων από χρωμοσωμικές ανωμαλίες και από το βαρύ κληρονομικό ιστορικό που φέρουν οι γονείς τους.

ΣΑΣ ΕΥΧΑΡΙΣΤΩ !



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