

# Transformational technologies drive the next generation of PGT-A

**Christopher Weier, PhD** 

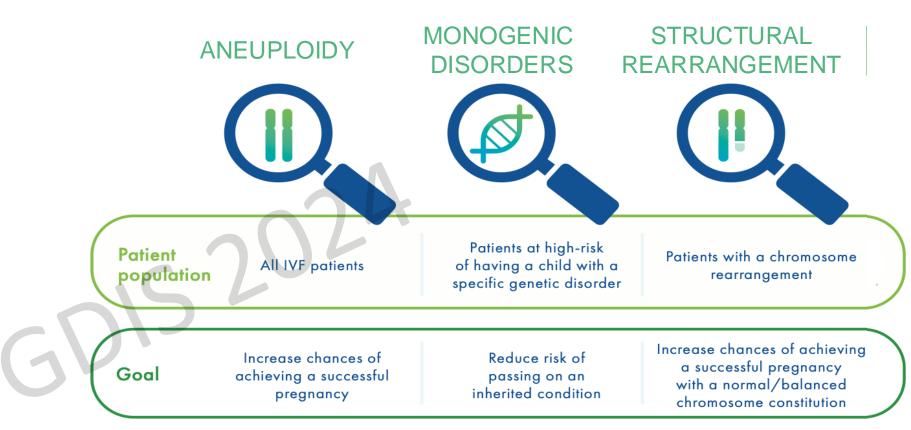
**CooperSurgical Research and Development** Senior Scientist

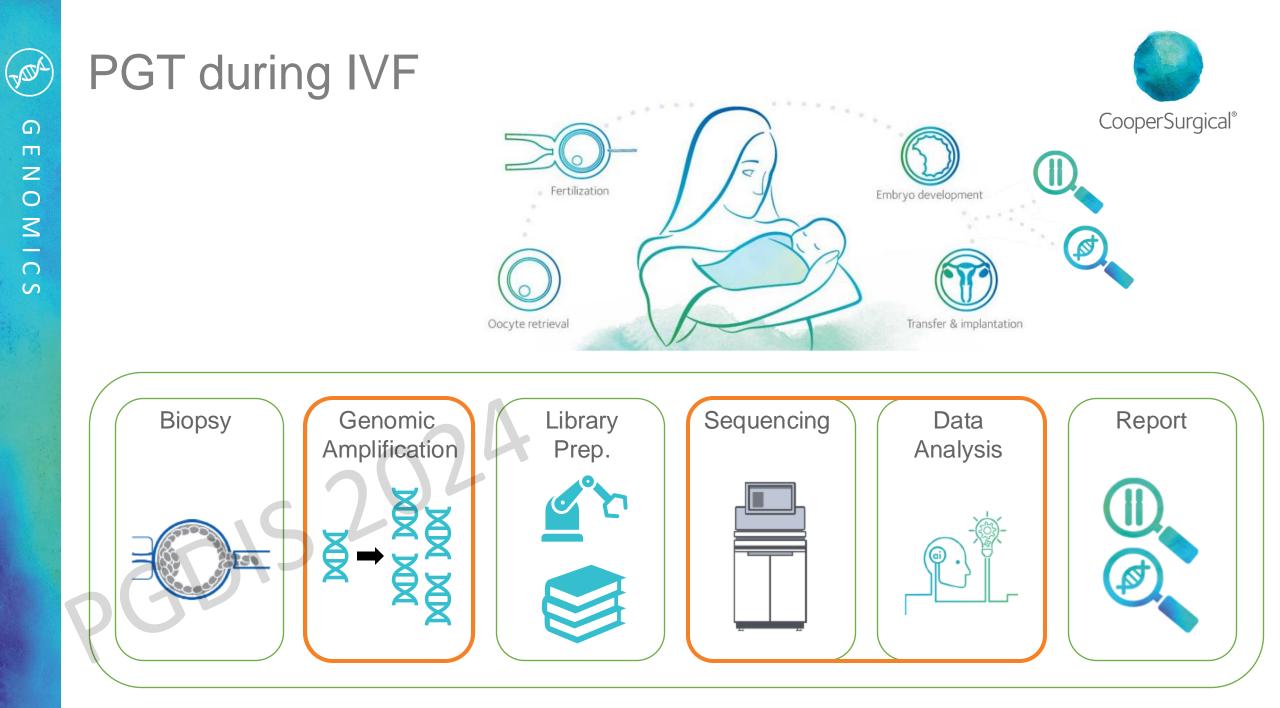
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Preimplantation Genetic Testing – PGT Laboratory tests to look at the health of the developing embryo



#### **PREIMPLANTATION GENETIC TESTING**









CooperSurgical<sup>®</sup>

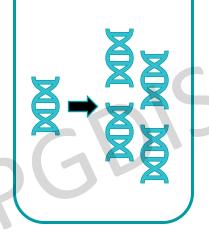


5-6 Trophectoderm Cells ~50 picograms genomic DNA

~1000 ng Amplified Product



Genomic Amplification



✓ Robust

high consistency of producing results

#### Accurate

high fidelity = very few errors

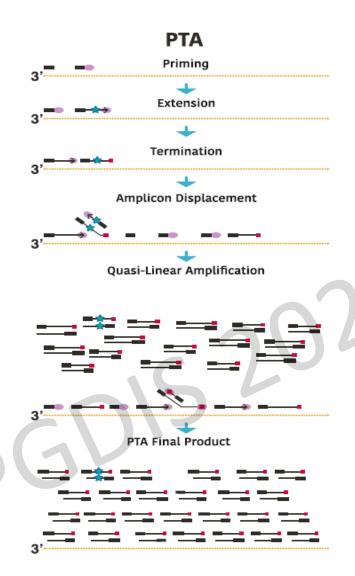
#### ✓ Complete

low bias = nearly whole genome & both alleles

# True whole genome amplification



Accurate, unbiased distribution of data across the entire genome CooperSurgical®



VARA

G E N

0

 $\leq$ 

0

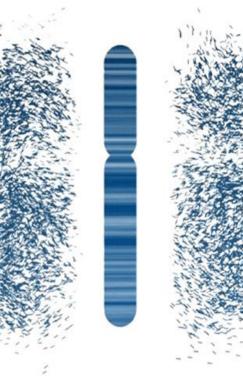
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#### **Non-PTA**

Random Priming Amplification Methods

#### PTA

Primary Template Directed Amplification



### True whole genome amplification

G E Z

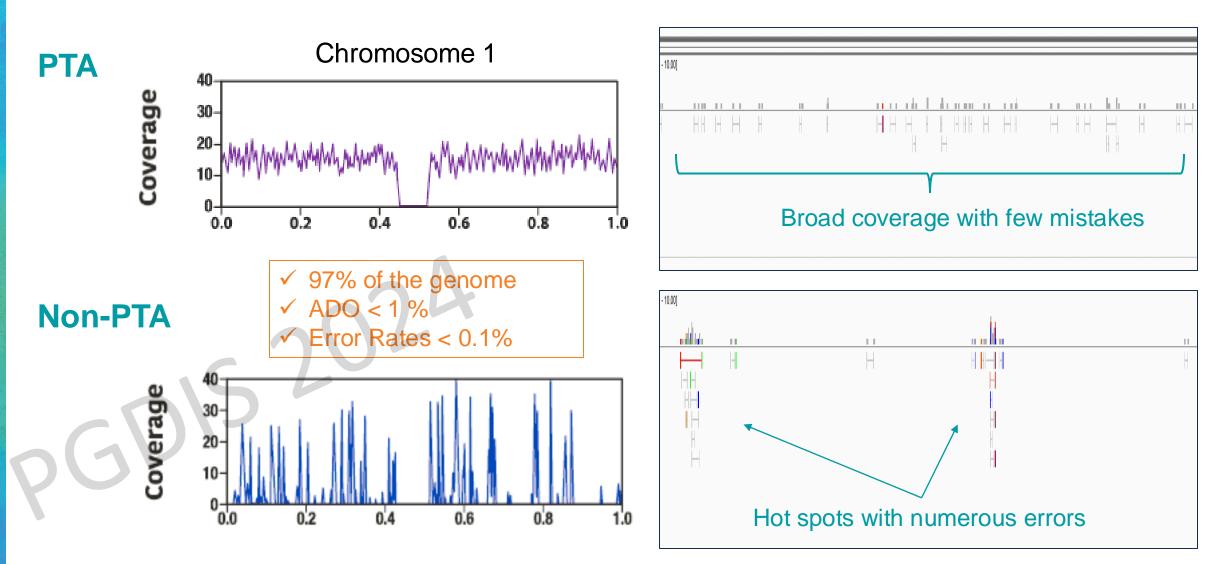
0

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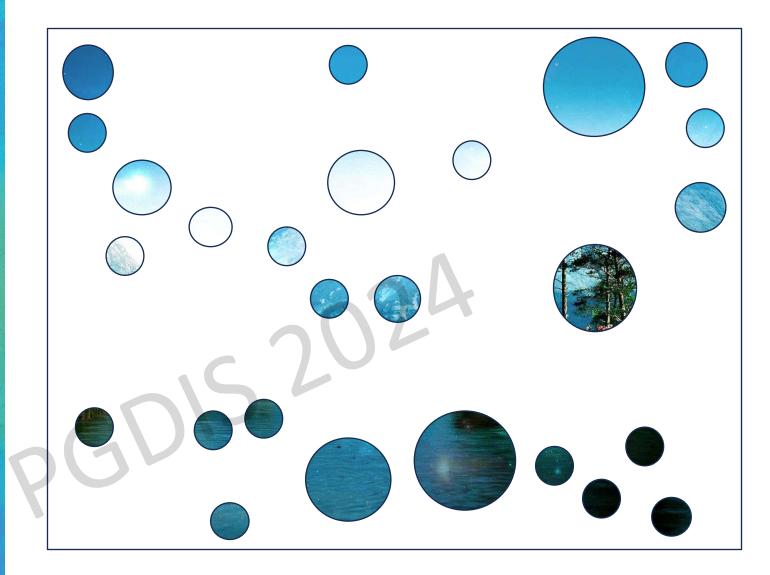
Accurate, unbiased distribution of data across the entire genome CooperSurgical®





### Filling in the gaps with PTA Our most complete and accurate basis for PGT





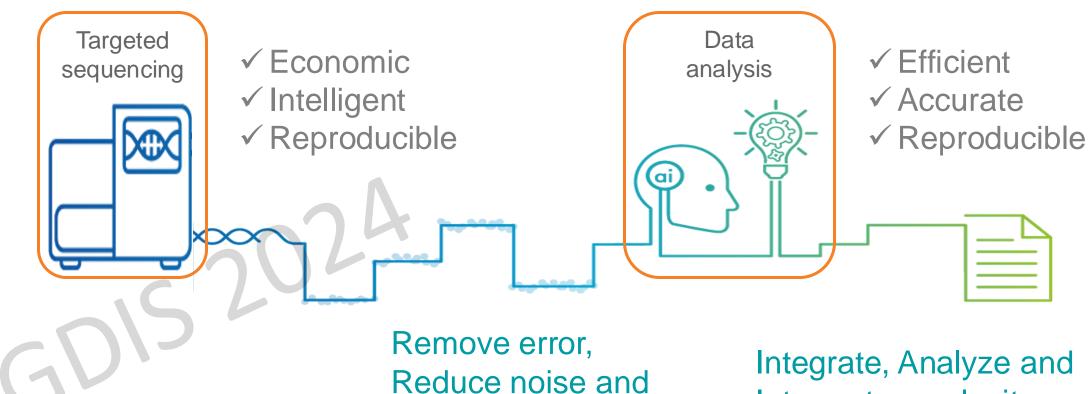
How do we <u>most responsibly</u> use this technology to the benefit of patients?



S

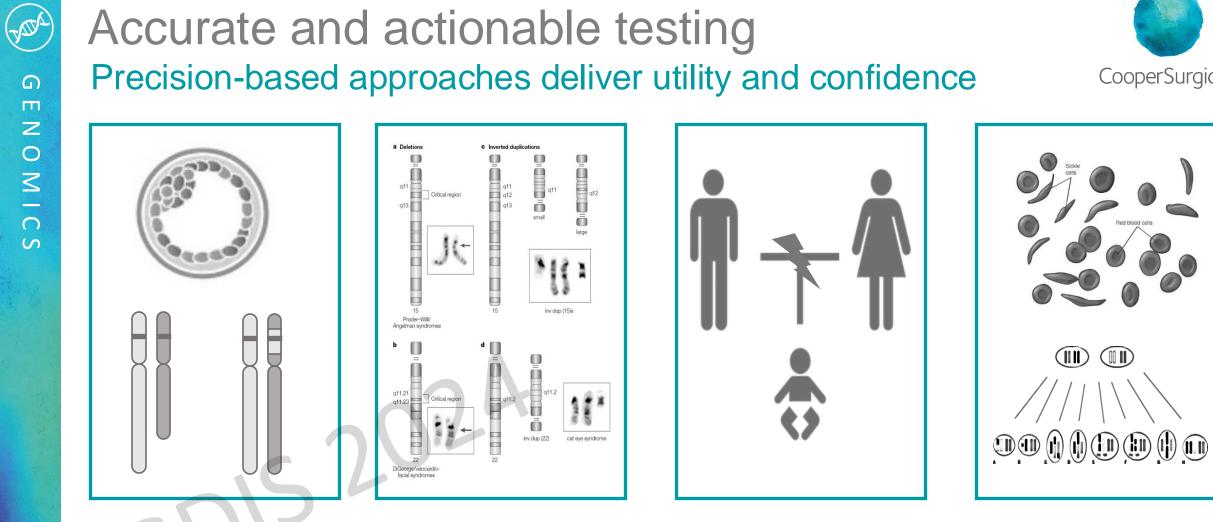
### Accurate and actionable testing Precision-based approaches deliver utility and confidence





Eliminate subjectivity

Interpret complexity



CooperSurgical®

Precision testing for common disorders and rare conditions

Segmental resolution, Mosaicism, and Loss of Heterozygosity (LoH)

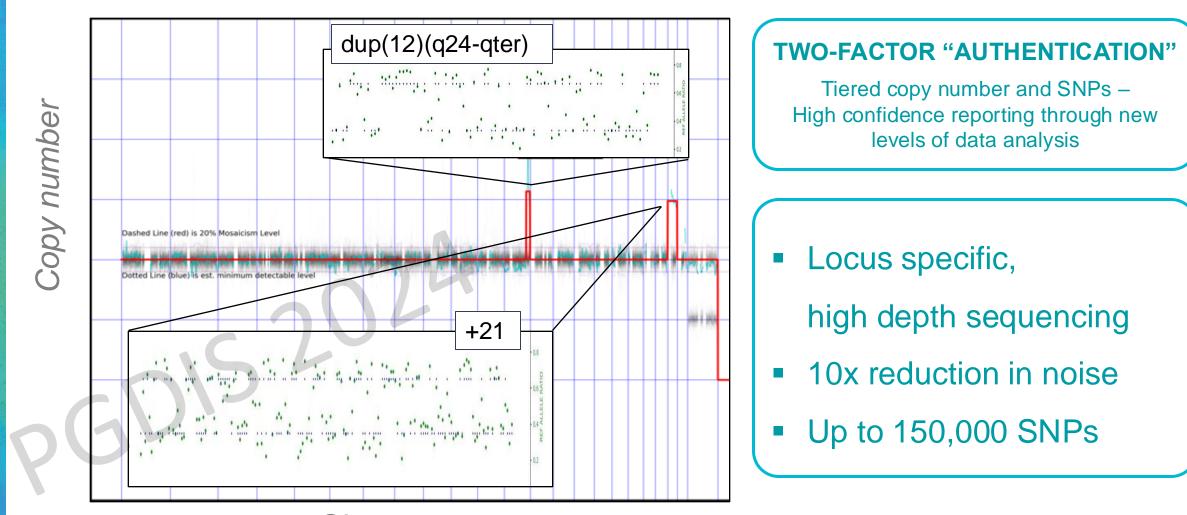
**Deletion and** Duplication Syndromes

Screening for *de novo* mutations and predisposition genes



### Whole chromosome and segmental CNV High resolution copy number variation + allele analysis



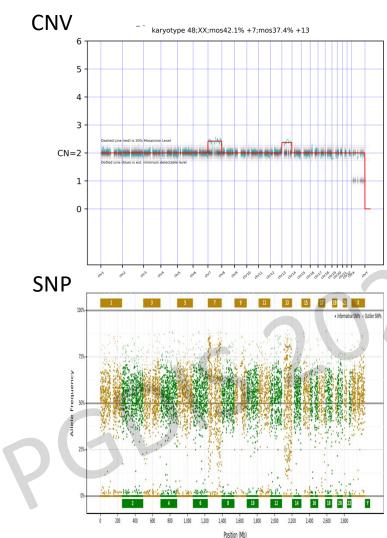


Chromosomes

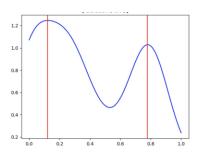


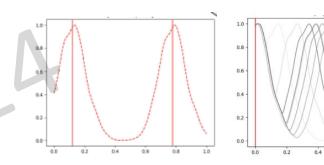
### Clarifying mosaic embryos CNV + SNP + Analysis



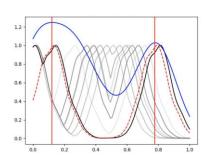


Mosaic Resolution by Decomposition





Clinical and Theoretical Decomposition to estimate mosaicism



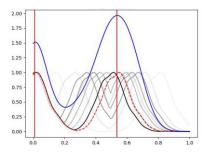
0,6

Allele Clusters

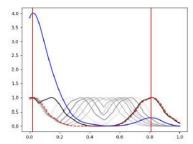
SNP density

estimated from

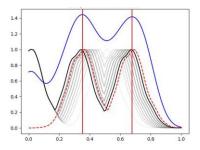
Diploid



Haploid



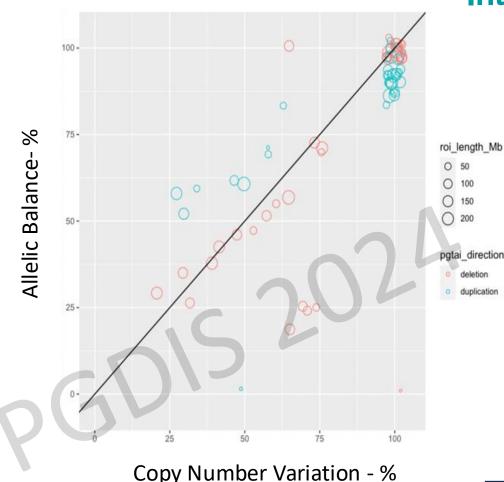
Triploid





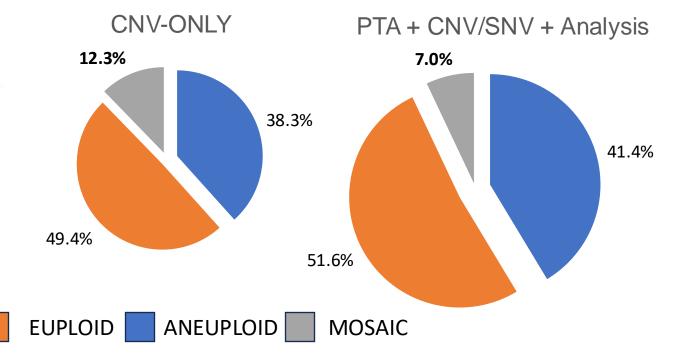
### Clarifying mosaic embryos CNV + SNP + Analysis





#### Integrated CNV/SNV analysis of 327 mosaics:

- ✓ 67.6% mosaics are confirmed
- ✓ 19% of low-level mosaics revert to euploid
- ✓ 13% of high-level mosaics are aneuploid



CooperSurgical internal data



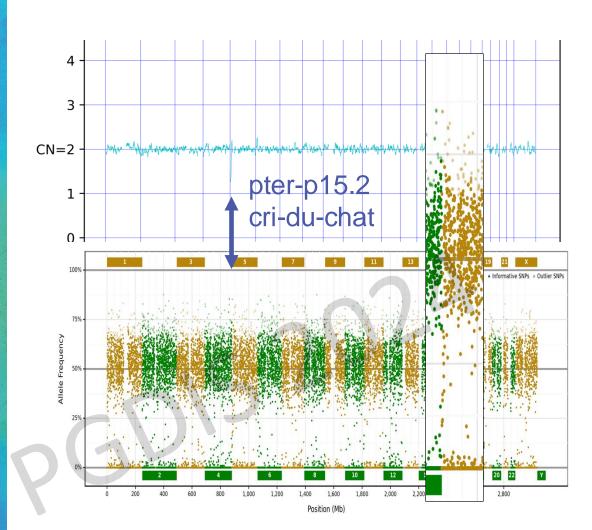
GENO

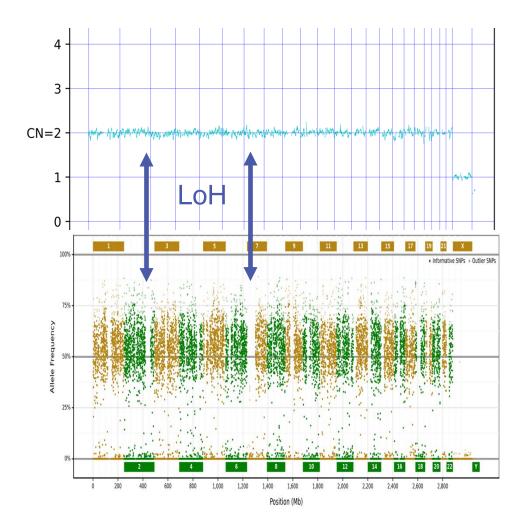
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### Minor variation with major clinical impact Deletions/Duplication syndromes and Loss of Heterozygosity







CooperSurgical internal data



### Direct mutation analysis

## Screening embryos for *de novo* and inherited mutations

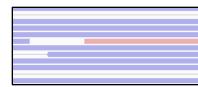


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0	Gene	Condition
MIC	BRCA 1/2	Hereditary Breast and Ova
	CFTR	Cystic Fibrosis
S	GJB2	Non-Syndromic Hearing Lo
	HBB	Sickle Cell Anemia
	SDHB	Her. Paraganglioma-Pheocl
	APC	Familial Adenomatous Poly
	TP53	Li-Fraumeni syndrome
	PMM2	Congenital Disorder of Gly
	MEFV	Familial Mediterranean Fev
	GALC	Krabbe Disease

Gene Condition		Concordant / Embryos Tested	
BRCA 1/2	Hereditary Breast and Ovarian Cancer	44/44	
CFTR	Cystic Fibrosis	21/21	
GJB2	Non-Syndromic Hearing Loss	21/21	
HBB	Sickle Cell Anemia	16/16	
SDHB	Her. Paraganglioma-Pheochromocytoma	12/12	
APC	Familial Adenomatous Polyposis	9/9	
TP53	Li-Fraumeni syndrome	6/6	
PMM2	Congenital Disorder of Glycosylation, T1a	6/6	
MEFV	Familial Mediterranean Fever	6/6	
GALC	Krabbe Disease	6/6	
ATM	ATM-Associated Cancer Susceptibility	6/6	
GAA	Glycogen Storage Disease, Type 2	5 / 5	
HBA1/HBA2	Alpha-Thalassemia	5 / 5	

### Total Concordant 200 / 200

#### Embryo 1 – Paternal Carrier



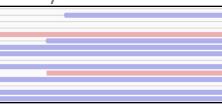


#### Embryo 2 – Maternal Carrier



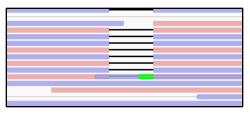
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#### Embryo 3 – Unaffected



#### Embryo 3 – Affected





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