

Transformational technologies drive the next generation of PGT-A

Christopher Weier, PhD

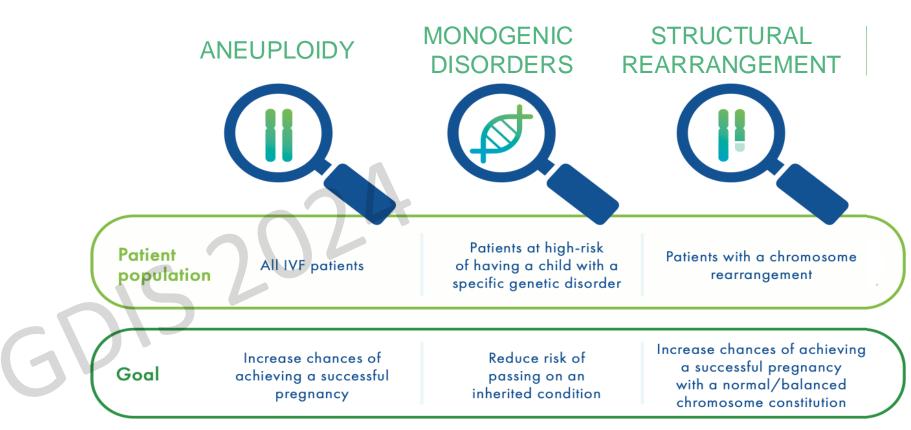
CooperSurgical Research and Development Senior Scientist

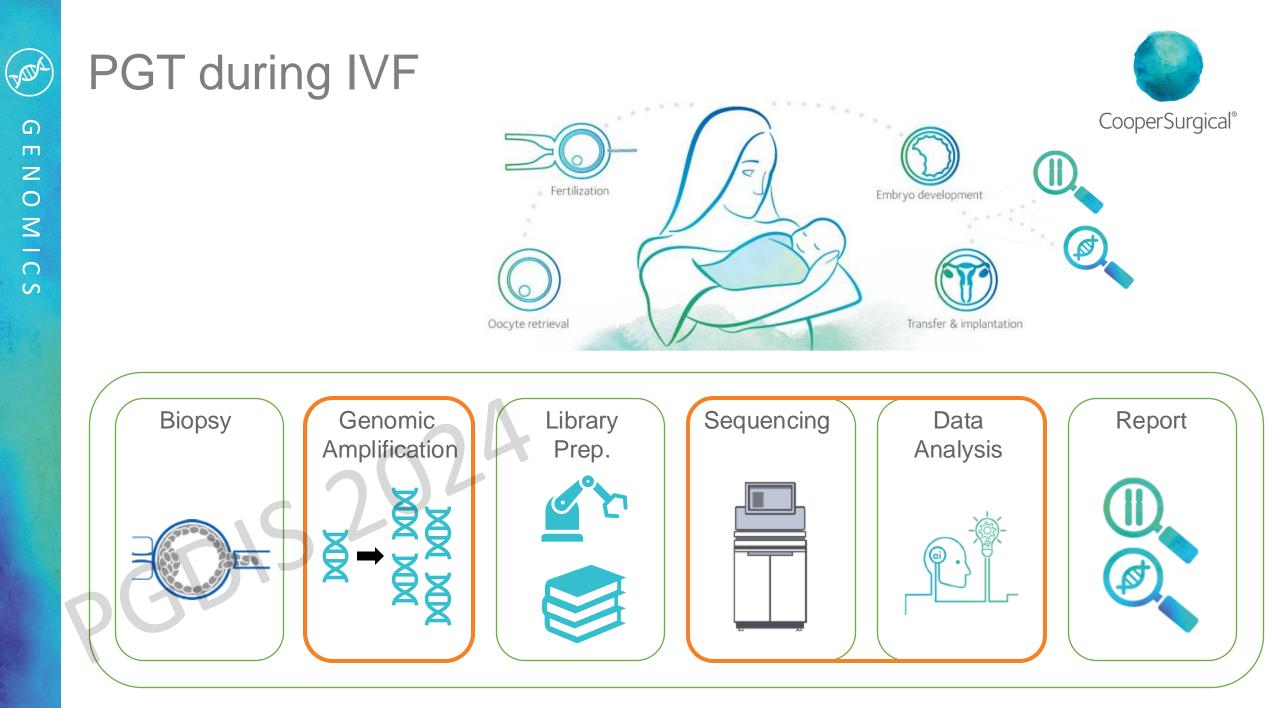
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Preimplantation Genetic Testing – PGT Laboratory tests to look at the health of the developing embryo



PREIMPLANTATION GENETIC TESTING









CooperSurgical[®]

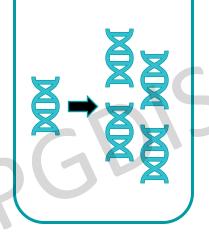


5-6 Trophectoderm Cells ~50 picograms genomic DNA

~1000 ng Amplified Product



Genomic Amplification



✓ Robust

high consistency of producing results

Accurate

high fidelity = very few errors

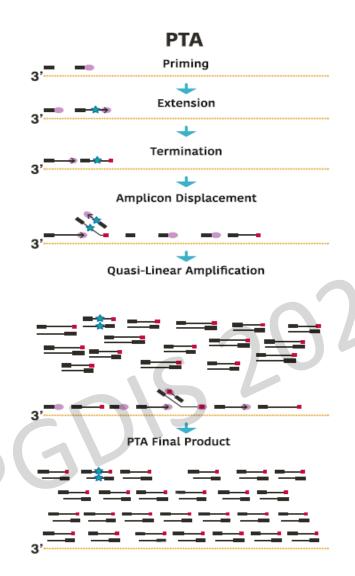
✓ Complete

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

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0

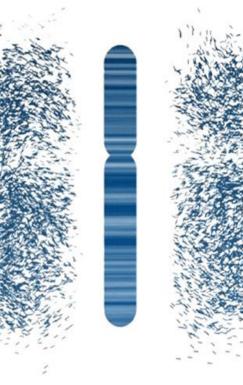
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Non-PTA

Random Priming Amplification Methods

PTA

Primary Template Directed Amplification



True whole genome amplification

G E Z

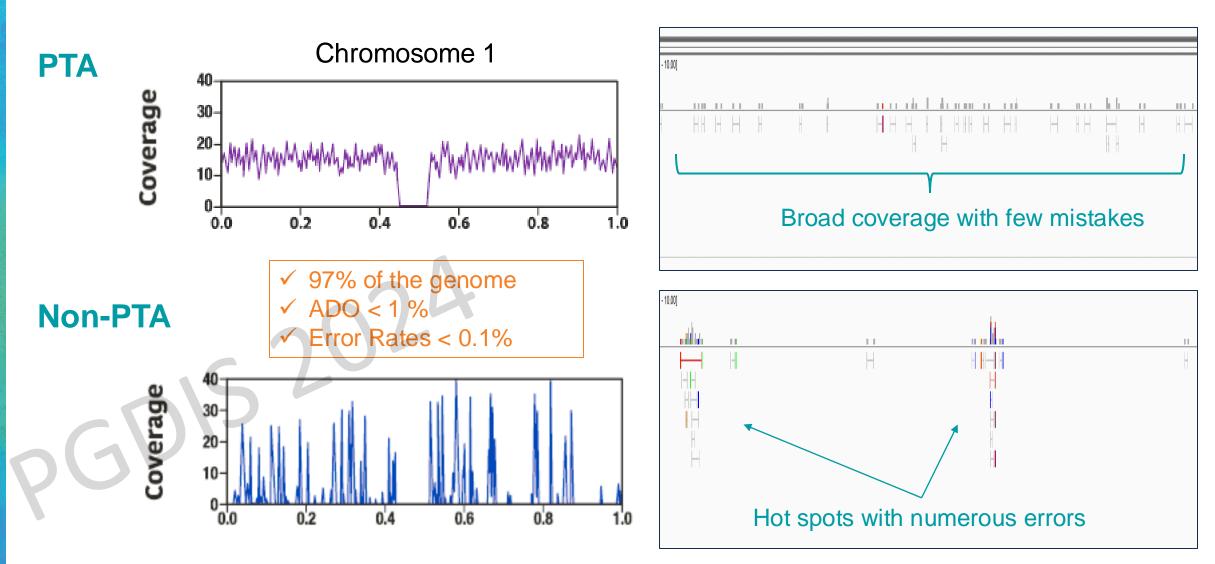
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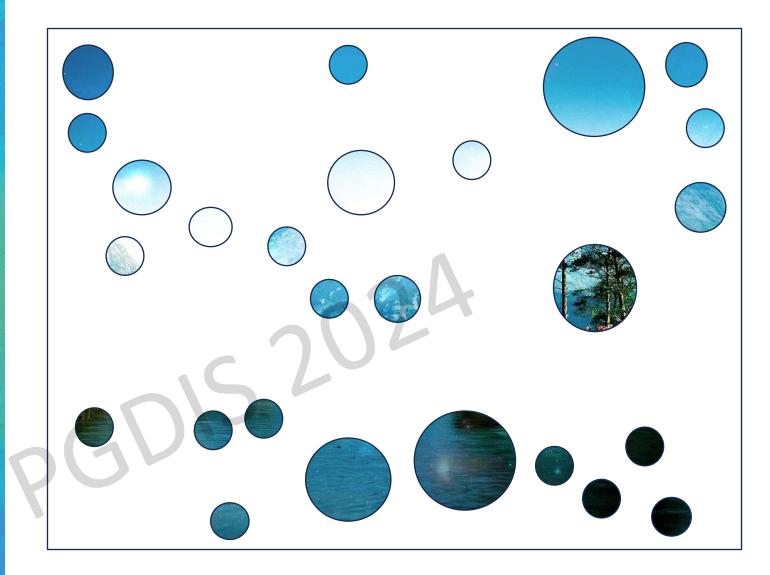
Accurate, unbiased distribution of data across the entire genome CooperSurgical®





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





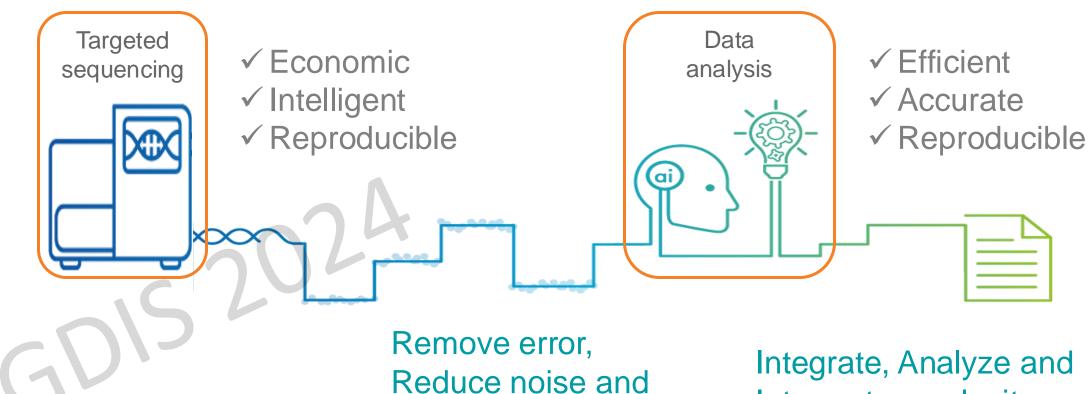
How do we <u>most responsibly</u> use this technology to the benefit of patients?



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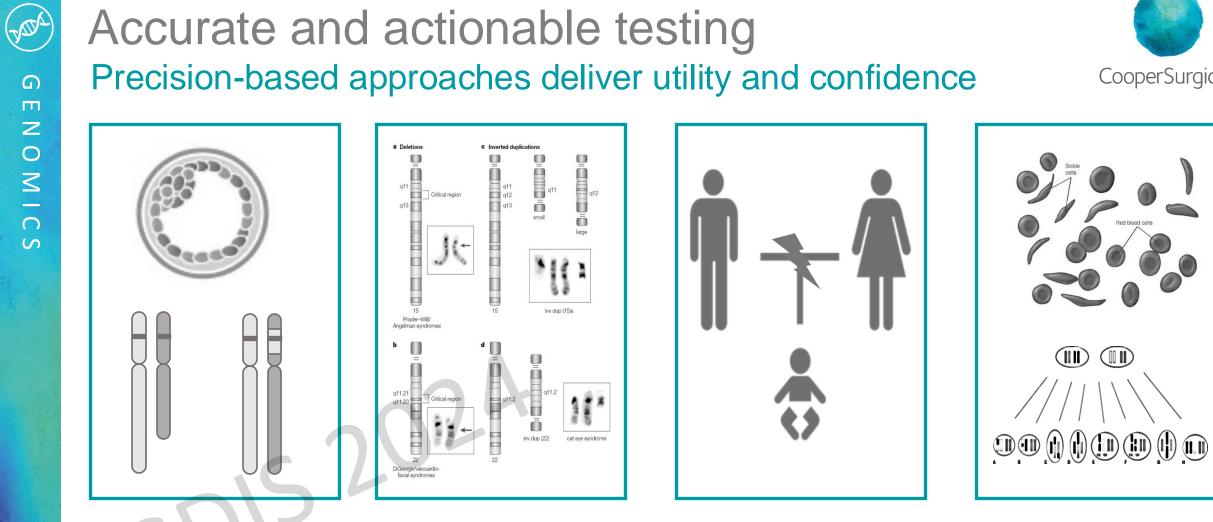
Accurate and actionable testing Precision-based approaches deliver utility and confidence





Eliminate subjectivity

Interpret complexity



CooperSurgical®

Precision testing for common disorders and rare conditions

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

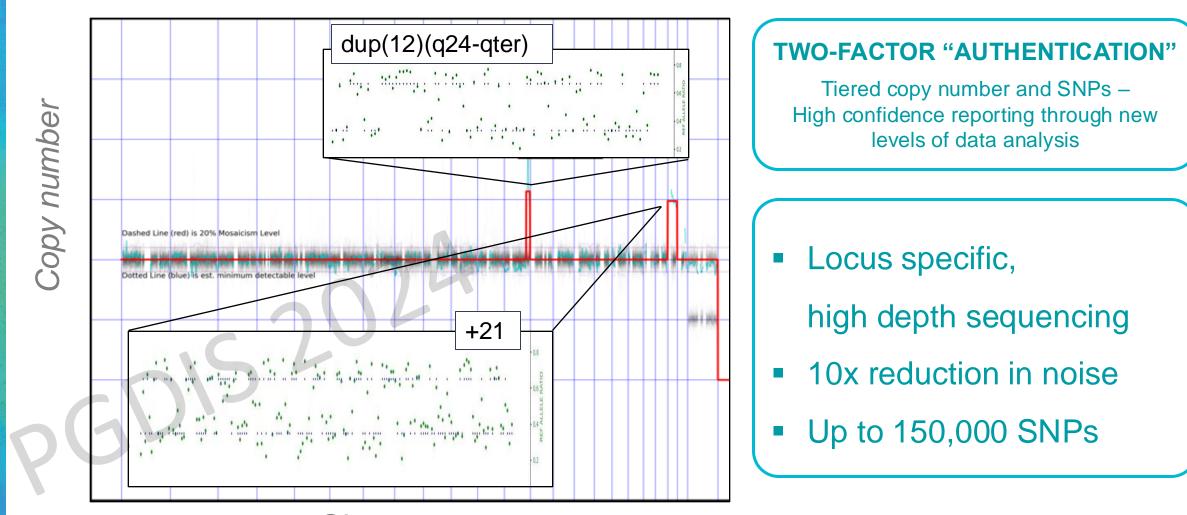
Deletion and Duplication Syndromes

Screening for *de novo* mutations and predisposition genes



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



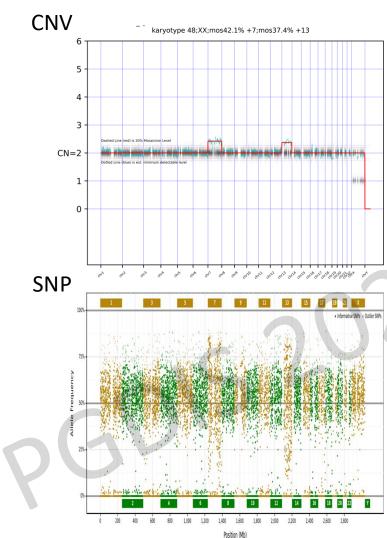


Chromosomes

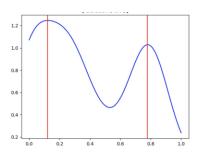


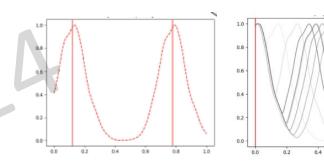
Clarifying mosaic embryos CNV + SNP + Analysis



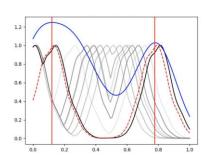


Mosaic Resolution by Decomposition





Clinical and Theoretical Decomposition to estimate mosaicism



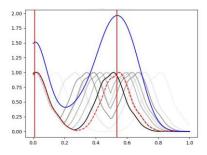
0,6

Allele Clusters

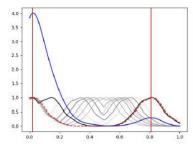
SNP density

estimated from

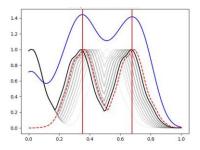
Diploid



Haploid



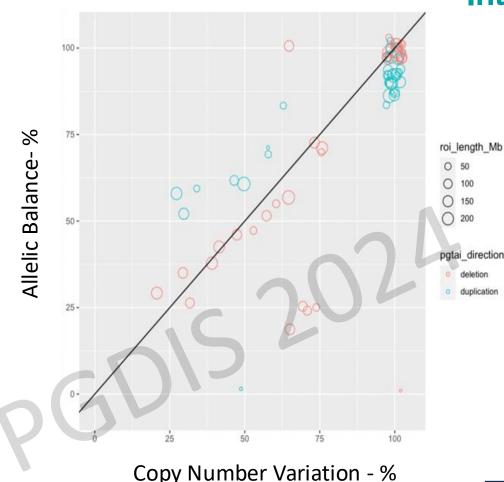
Triploid





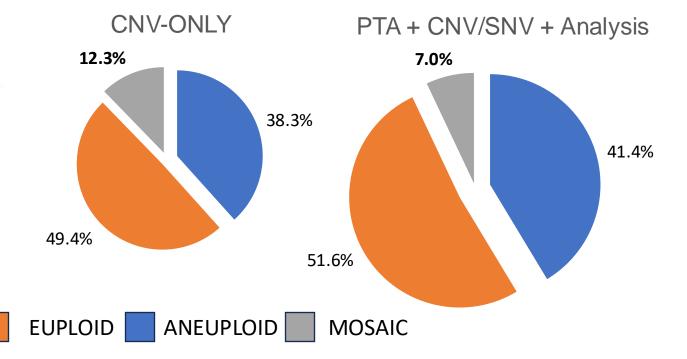
Clarifying mosaic embryos CNV + SNP + Analysis





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



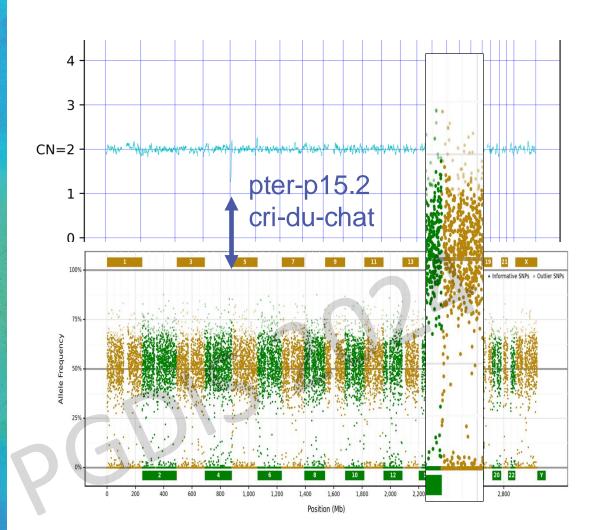
GENO

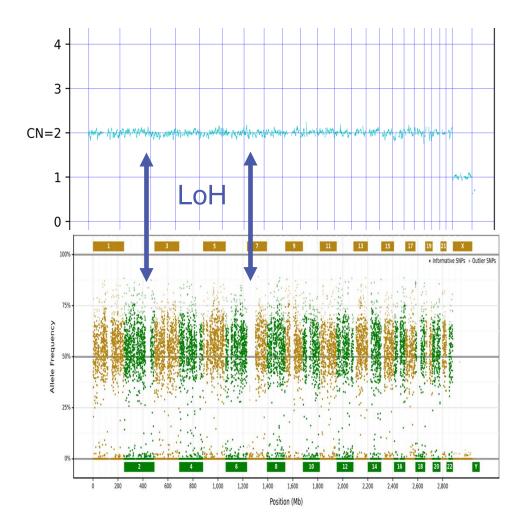
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Minor variation with major clinical impact Deletions/Duplication syndromes and Loss of Heterozygosity







CooperSurgical internal data



Direct mutation analysis

Screening embryos for *de novo* and inherited mutations

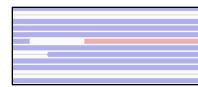


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0	Gene	Condition
MIC	BRCA 1/2	Hereditary Breast and Ova
	CFTR	Cystic Fibrosis
S	GJB2	Non-Syndromic Hearing Lo
	HBB	Sickle Cell Anemia
	SDHB	Her. Paraganglioma-Pheocl
	APC	Familial Adenomatous Poly
	TP53	Li-Fraumeni syndrome
	PMM2	Congenital Disorder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



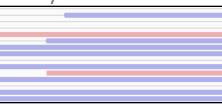


Embryo 2 – Maternal Carrier



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Embryo 3 – Unaffected



Embryo 3 – Affected





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