

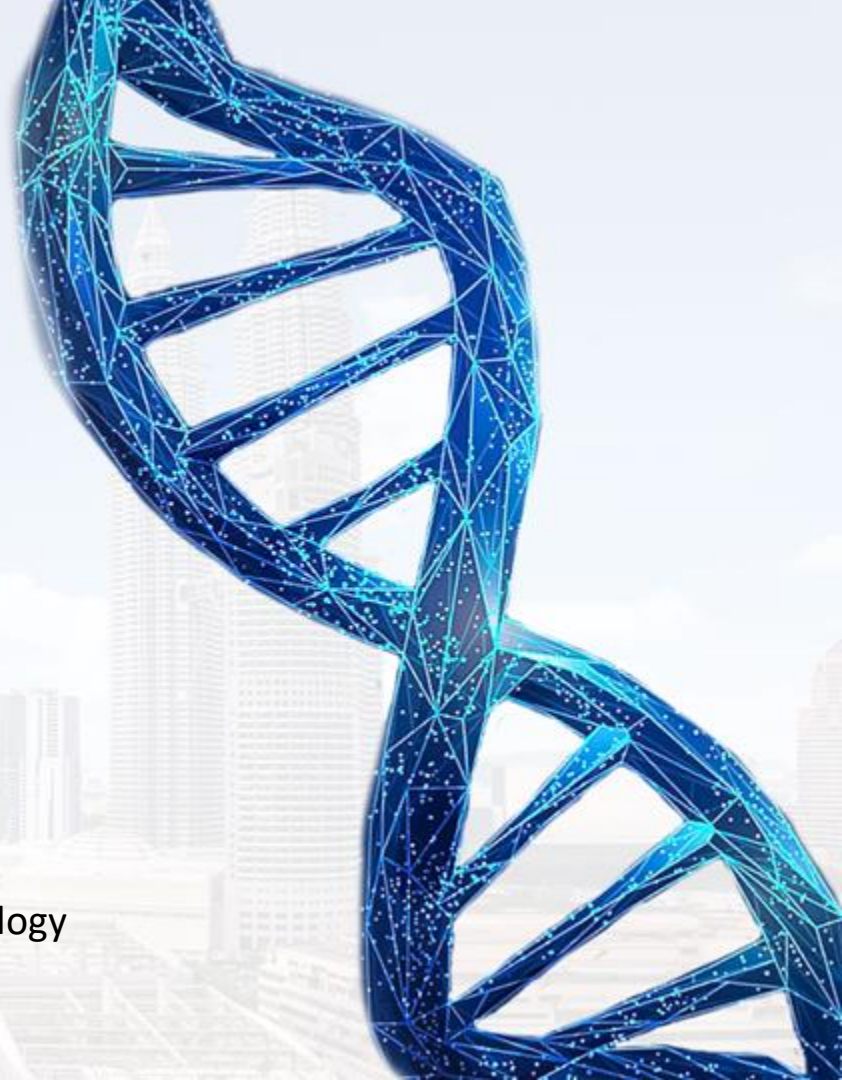
Analysis of spent culture medium

Richard CHOY, PhD

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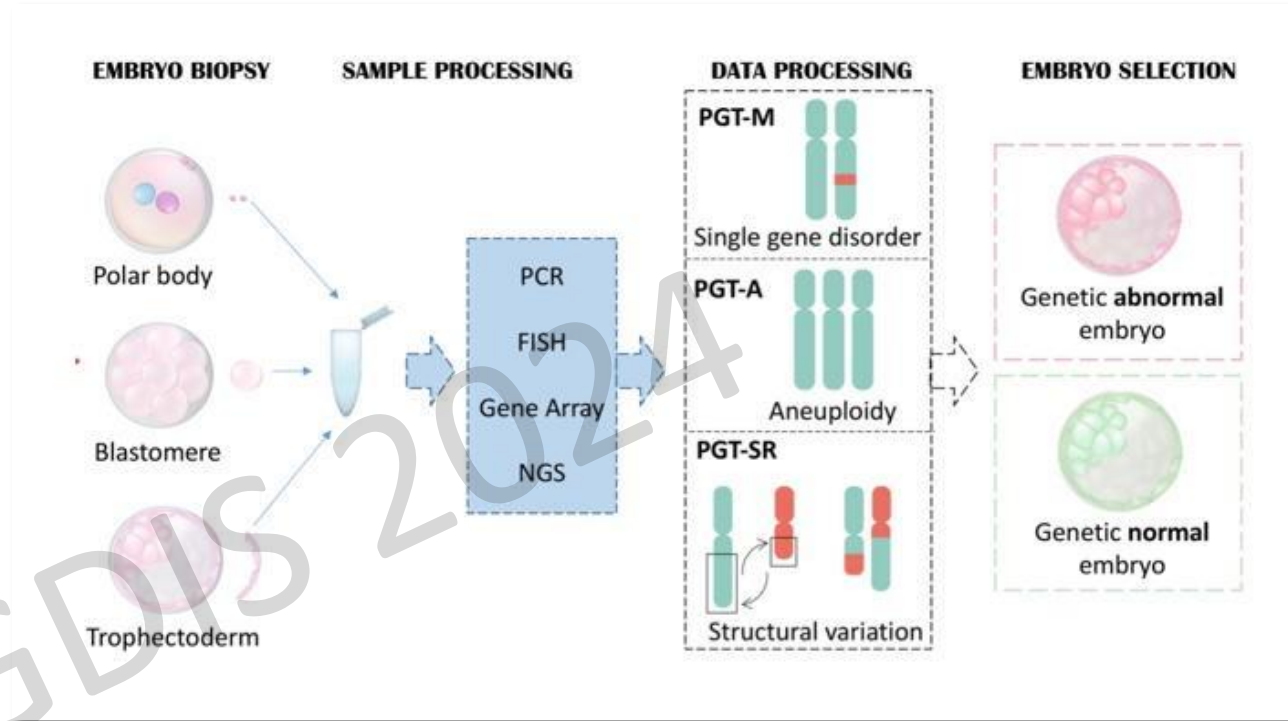


Outline

- Why analysis spent culture medium
- Part I: Non-invasive PGT-A (niPGT-A)
- Part II: Metabolic profiling



Biopsy hurts the embryo



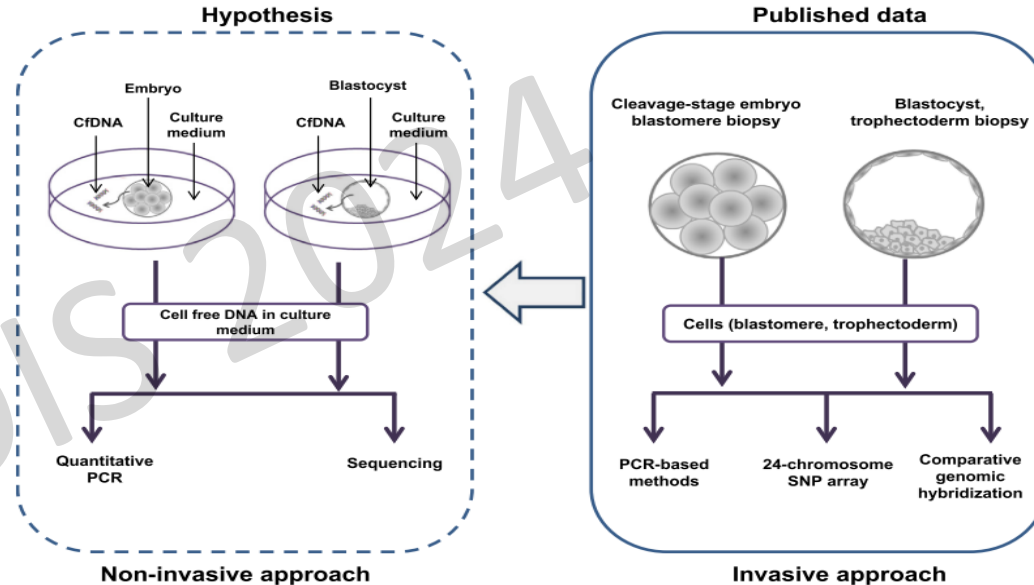
What are the down sides of PGT?

- Increases costs IVF
- Need elective cryopreserving embryos, delay ET until results back
- Invasive procedure
 - Embryo biopsy may affect embryo development
 - Lead to potential loss of embryo
 - Long –term effect of embryo biopsy not defined at this time
 - Reported to have 5% relative reduction in live birth rate due to damage and also false positive results from inherent technical errors



Hypotheses to analysis spent culture medium

- Embryos release DNA, RNA, and protein molecules into its surroundings (handling and/or culture media)
- The **identities** and **quantities** of these molecules reflect the **health** and **birth potential** of the embryo



Human embryo reveals nuclear DNA

Please cite this article in press as: Domingo-Muelas et al., Human embryo live imaging reveals nuclear DNA shedding during blastocyst expansion and biopsy, Cell (2023), <https://doi.org/10.1016/j.cell.2023.06.003>

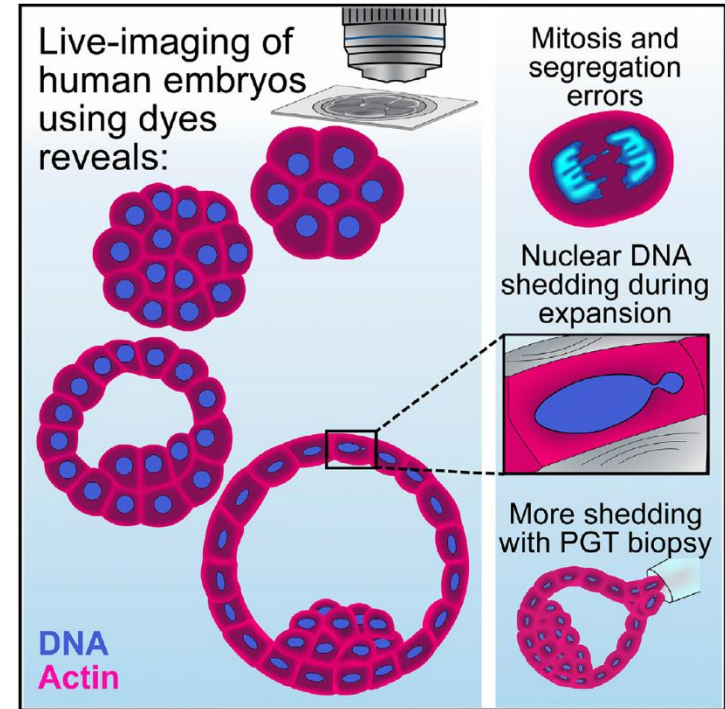
Cell

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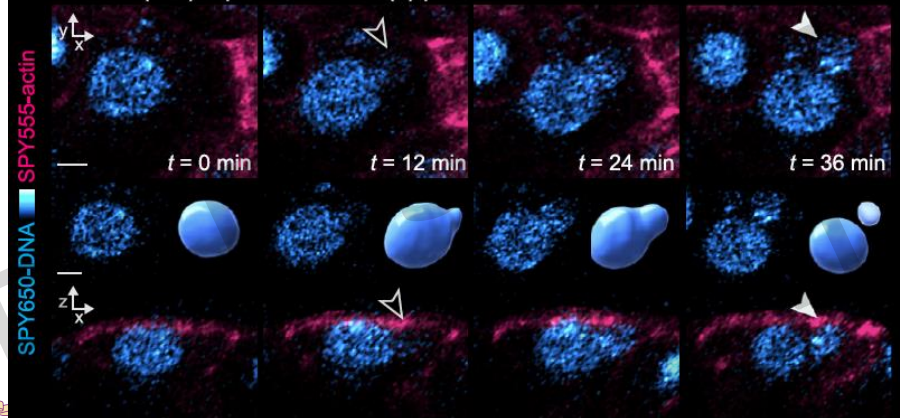
Article

Human embryo live imaging reveals nuclear DNA shedding during blastocyst expansion and biopsy

Ana Domingo-Muelas,^{1,2,10} Robin M. Skory,^{1,3,10} Adam A. Moverley,^{1,4} Goli Ardestani,⁵ Oz Pomp,¹ Carmen Rubio,⁶ Piotr Tetlak,¹ Blake Hernandez,¹ Eric A. Rhon-Calderon,¹ Luis Navarro-Sánchez,⁶ Carmen M. García-Pascual,⁶ Stephanie Bissiere,¹ Marisa S. Bartolomei,¹ Denny Sakkas,^{5,*} Carlos Simón,^{2,7,8,9,*} and Nicolas Plachta^{1,11,*}



Human (live) 1 μm thick axial (z) plane



Analysis spent culture medium

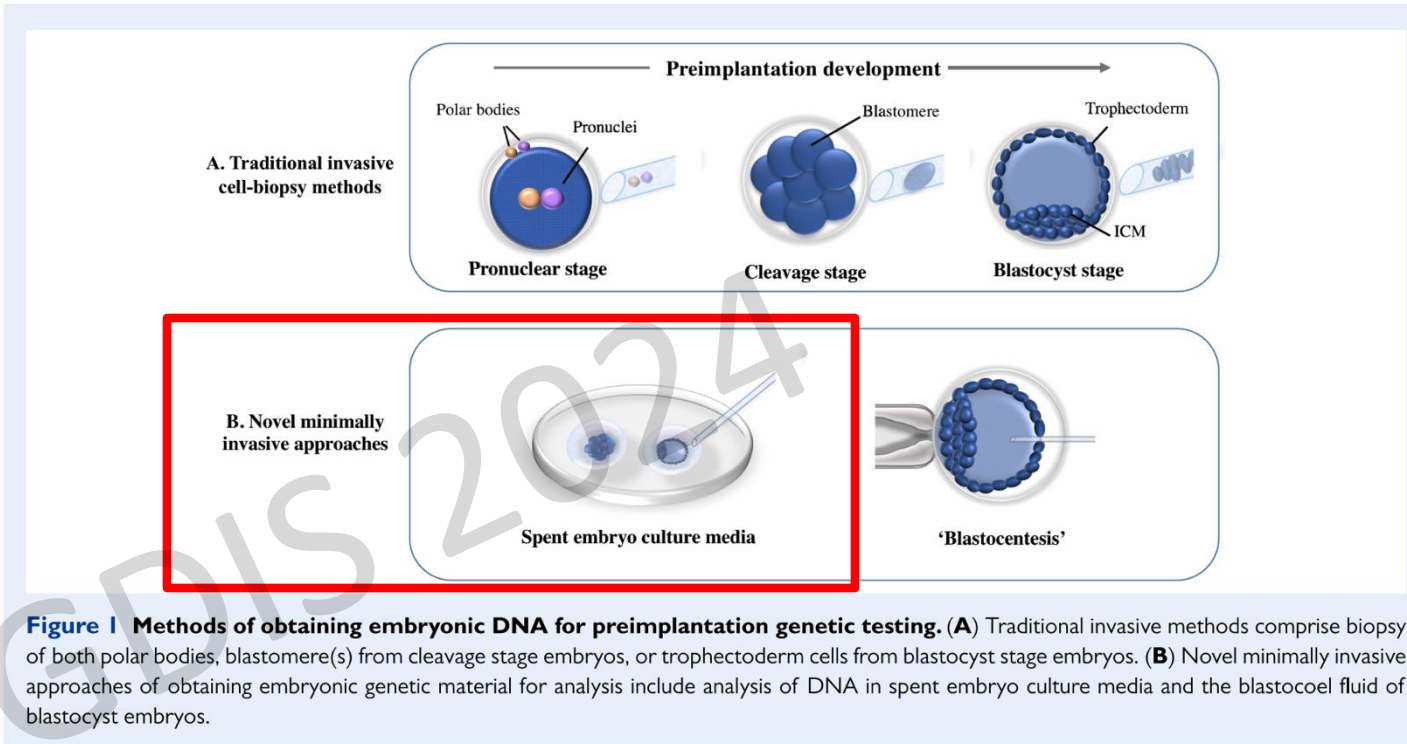


Figure 1 Methods of obtaining embryonic DNA for preimplantation genetic testing. **(A)** Traditional invasive methods comprise biopsy of both polar bodies, blastomere(s) from cleavage stage embryos, or trophectoderm cells from blastocyst stage embryos. **(B)** Novel minimally invasive approaches of obtaining embryonic genetic material for analysis include analysis of DNA in spent embryo culture media and the blastocoel fluid of blastocyst embryos.

Part I: Non-invasive PGT-A (niPGT-A)

PNAS

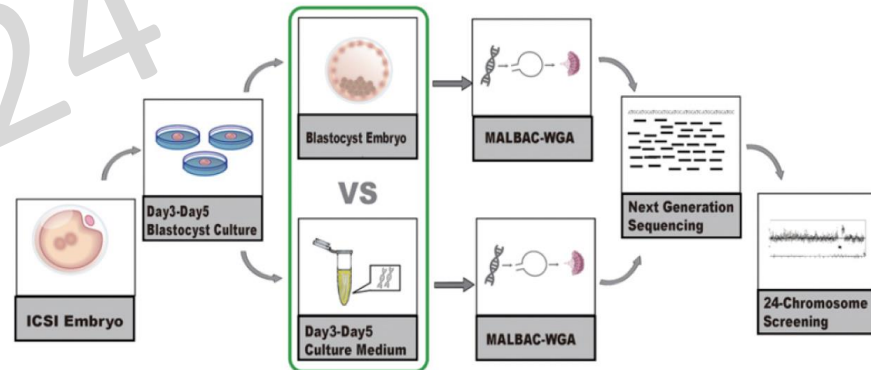
Noninvasive chromosome screening of human embryos by genome sequencing of embryo culture medium for in vitro fertilization

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

^aReproductive 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; ^bReproductive Medicine Center, Wuxi Maternity and Child Health Hospital Affiliated to Nanjing Medical University, Wuxi, Jiangsu 214002, China; ^cDepartment of Clinical Research, Yikon Genomics Company, Ltd., Shanghai 201499, China; ^dBiodynamic Optical Imaging Center (BIOPIC), School of Life Sciences, Peking University, Beijing 100871, China; ^eDepartment of Obstetrics, Gynecology, and Reproductive Biology, Brigham and Women's Hospital, Boston, MA 02115; ^fHarvard Medical School, Boston, MA 02115; ^gBeijing Advanced Innovation Center for Genomics, Peking University, Beijing 100871, China; and ^hDepartment of Chemistry and Chemical Biology, Harvard University, Cambridge, MA 01238

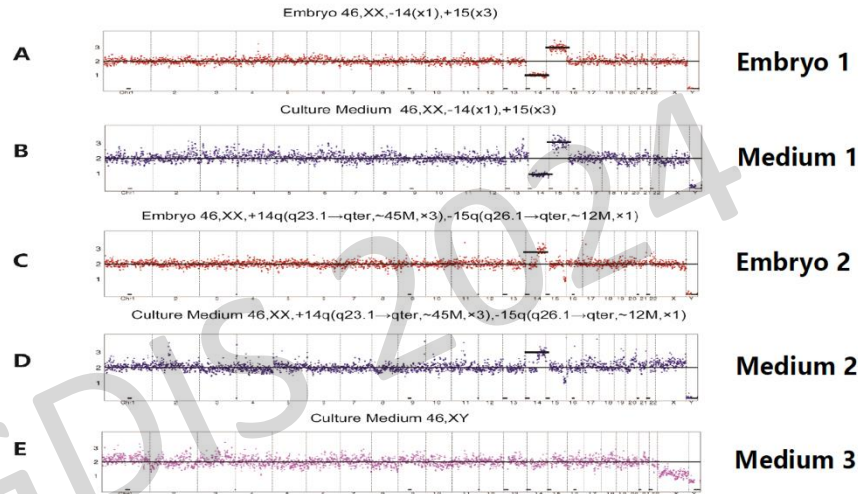
	Number	Ratio(%)
Sensitivity	15/17	88.2%
Specificity	21/25	84.0%
PPV	15/19	78.9%
NPV	21/23	91.3%

PPV: pos. predictive value
NPV: neg. predictive value



First clinical case

IVF Cycles	Wife	Husband	No. of Blastocyst Embryos	No. of Transferrable Embryos	Clinical Outcome
3	Premature ovarian failure	t(14;15)(q22;q24)	3	1	Healthy boy was born at 38 ⁺⁴ wk



Xu J, et al., PNAS, 2016 Oct 18;113(42):11907-1191.



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Non-invasive PGT-A (niPGT-A)

Spent Embryo
culture medium (SCM)

Noninvasive preimplantation genetic testing for
aneuploidy in spent medium may be more reliable
than trophectoderm biopsy

Lei Huang^{a,b}, Berhan Bogale^b, Yaqiong Tang^{c,d}, Sijia Lu^e, Xiaoliang Sunney Xie^{a,c,d,1}, and Catherine Racowsky^{b,1}

Journal of Assisted Reproduction and Genetics
<https://doi.org/10.1007/s10815-019-01517-7>

PNAS

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ASSISTED REPRODUCTION TECHNOLOGIES

A prospective study of non-invasive preimplantation genetic testing
for aneuploidies (NiPGT-A) using next-generation sequencing (NGS)
on spent culture media (SCM)



Queenie S. Y. Yeung¹ • Ying Xin Zhang² • Jacqueline P. W. Chung¹ • Wai Ting Lui² • Yvonne K. Y. Kwok² •
Baoheng Gui^{3,4} • Grace W. S. Kong¹ • Ye Cao² • Tin Chiu Li¹ • Kwong Wai Choy^{2,3}

Received: 10 February 2019 / Accepted: 30 June 2019

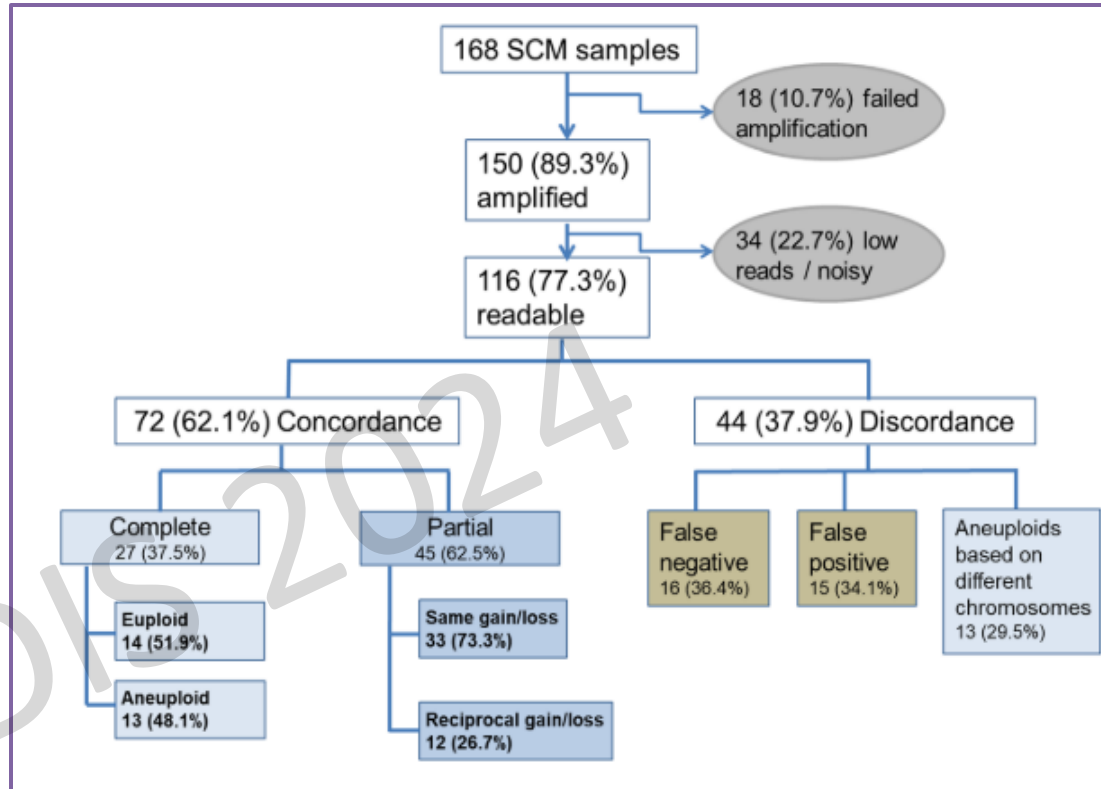


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PGT-A vs niPGT-A concordance rate



PGT-A concordance 62% between SCM and trophectoderm (TE)

Overview of current niPGT-A application

Table IV niPGT-A analysis of SCM.

Study	Sample details		Analysis				Concordance (%)		
	Media analyzed	Sample number	Amplification method	Amplification rate (%)	Cytogenetic method	Sequencing platform	Overall ploidy (aneuploid or euploid)	Full karyotype	Per single chromosome
Spent media only									
Shamoni et al. (2016)	D3-5/6	57	Repli-G (Qiagen)	97	aCGH (Agilent Technologies)	Not applicable	—	—	—
Xu et al. (2016)	D3-5	42	MALBAC (Yikon Genomics)	100	NGS (NEB Ultra DNA Kit, NEB)	HiSeq 2500 (Illumina)	Normal (euploid) vs 'abnormal' (inc. mosaicism and segmentals): WE: 86	—	—
Liu et al. (2017)	D1-5	88	MALBAC (Yikon Genomics)	91	NGS (Library prep method not specified)	HiSeq 2500 (Illumina)	Normal (euploid) vs 'abnormal' (inc. mosaicism and segmentals): TE: 84	Including >40% mosaicism and large copy number variation: TE: 65. Cells of arrested/degenerated embryos (which didn't reach blastocyst stage): 44	—
Feichtinger et al. (2017)	D1-5	22	SurePlex (Illumina)	82	aCGH (24Sure, Illumina)	Not applicable	Euploid vs aneuploid: PB: 72	—	PB: 49% of single chromosomal aneuploidies concordant with SCM
Ho et al. (2018)	D1-3 D1-5	41	PicoPLEX (Rubicon Genomics), using 20 cycles instead of 14	D1-3 = 39 D1-5 = 80	NGS	Ion S5 Sequencer (Life Technologies)	Euploid vs aneuploid: D1-3 SCM vs WE: 56 D1-5 SCM vs WE: 46	—	—
Vera-Rodriguez et al. (2018)	D3-5	56	SurePlex (Illumina), then a second round using Ion Reproseq (ThermoFisher)	91	NGS (Ion Reproseq, ThermoFisher)	Ion PGM instrument (ThermoFisher)	Both aneuploid: TE: 30	Including segmental aneuploidies and mosaicism: TE: 16	Whole chromosome aneuploidies only: TE: 5
Fang et al. (2019)	D3-5/6	170	MALBAC (Yikon Genomics)	97	NGS (NEB Ultra DNA Kit, NEB)	HiSeq 2500 (Illumina)	—	—	—
Huang et al. (2019)	D5-6 and D6/7	52	NICSwift—modified MALBAC (Yikon Genomics)	92	NGS (NEBNext Ultra II DNA kit, NEB)	HiSeq 2500 (Illumina)	Euploid vs aneuploid: WE: 94	Including segmental aneuploidies and mosaicism: WE: 83	—
Yeung et al. (2019)	D3-5/6	168	SurePlex (Illumina)	89	NGS (VeriSeq, Illumina)	MiSeq (Illumina)	Euploid vs aneuploid: TE: 73	Autosomal chromosomes: TE: 62 Sex chromosomes: TE: 82	—
Rubio et al. (2019)	D4-5/6/7	115	Modified version of IonReproseq (ThermoFisher)	95	NGS (Ion Reproseq, ThermoFisher)	Ion S5TM XL system (ThermoFisher)	Euploid vs aneuploid: D4-5 SCM vs TE = 63 D5-6/7 SCM vs TE = 84 Overall vs TE = 79	Including segmental aneuploidies: D4-5 SCM vs TE = 41 D5-6/7 SCM vs TE = 72 Overall vs TE = 64	—

- Currently by niPGT-A, not all groups can achieve 100% amplification rate.
- The highest sensitivity is 88.6% and 87.5% by Kuznyetsov, V. et al.
- An estimated >20% to 80% of instances involve contamination with maternal genetic material

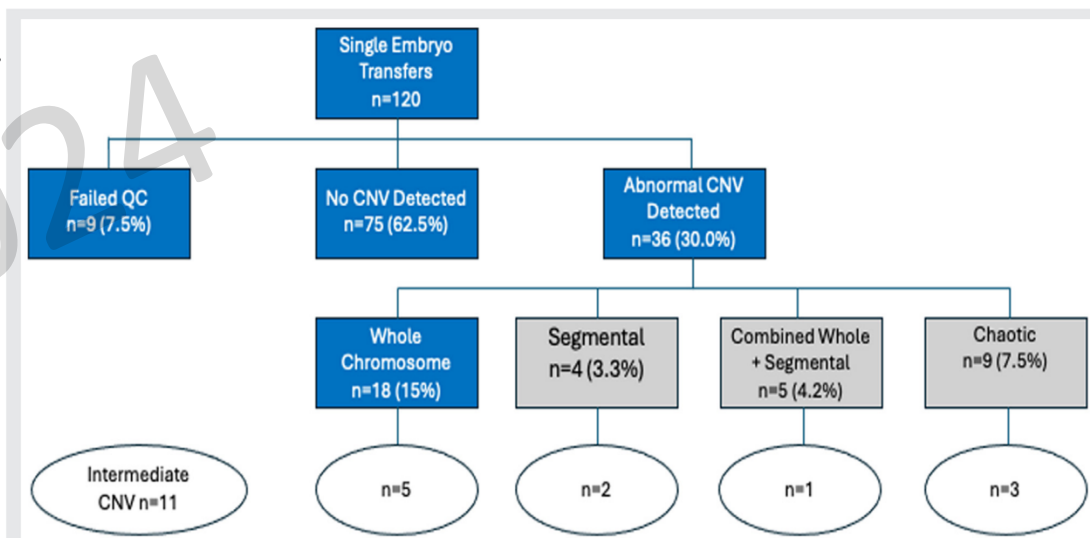


A pilot study to investigate the clinically predictive values of copy number variations detected by next-generation sequencing of cell-free deoxyribonucleic acid in spent culture media

Gary Nakhuda, M.D.,^a Sally Rodriguez, Sc.M., C.G.C.,^b Sophia Tormasi, B.Sc. and Catherine Welch, M.B.A., T.S.^b

-NOT all CNV abnormalities apparent on NGS profiles are equally predictive of nonviability.
-Sex chromosome: consistent results in 74.1% (n = 40/54).

-woman of age 35 years or younger at the time of oocyte retrieval



NGS interpretations for 120 single embryo transfers. CNV, copy number variation; NGS, next-generation sequencing; QC, quality control.

Nakhuda. Study of predictive values of niPGT-A. Fertil Steril 2024.



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Noninvasive preimplantation genetic testing for aneuploidy in spent culture medium as a substitute for trophectoderm biopsy

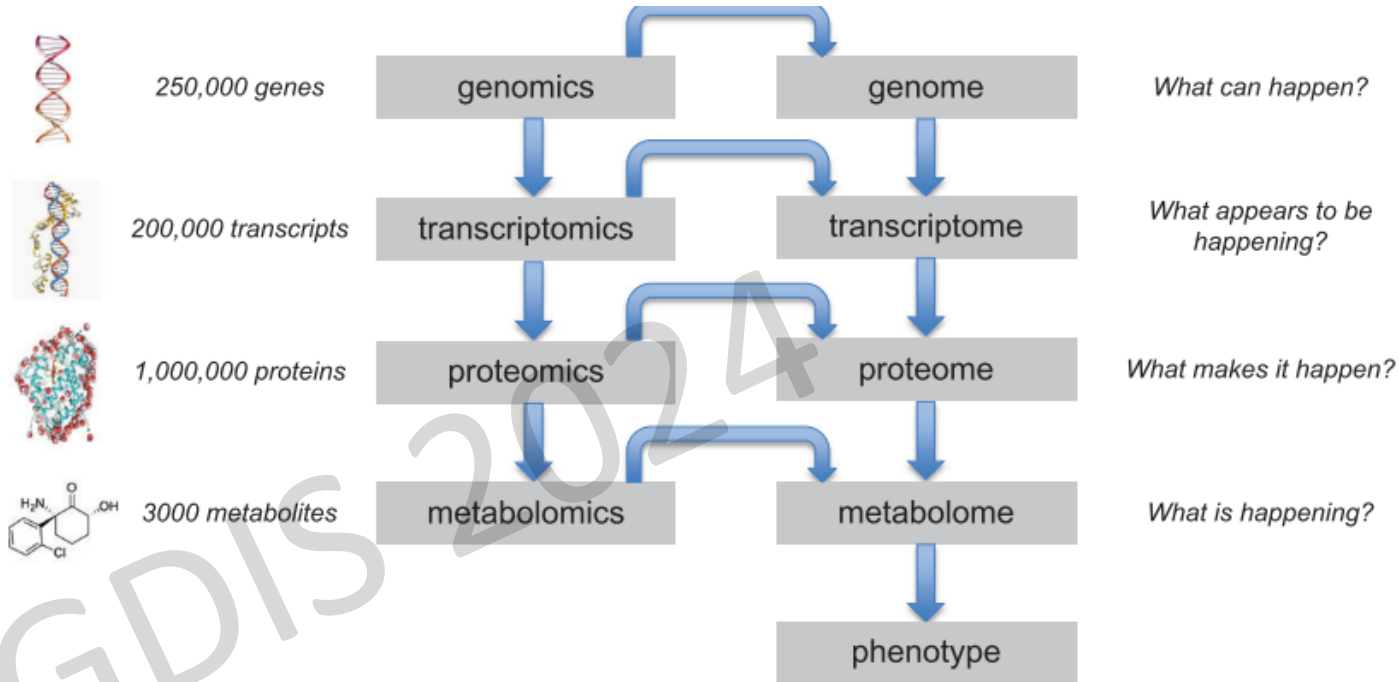
Carmen Rubio, Ph.D.,^{a,b} Catherine Racowsky, Ph.D.,^c David H. Barad, M.D., M.S.,^{d,e}
Richard T. Scott Jr., M.D., H.C.L.D.,^{f,g} and Carlos Simon, M.D., Ph.D.^{h,i}

Pros and cons:

1. How reliable is embryo cfDNA versus teDNA as a representative of embryo chromosomal constitution?
2. Is embryo cfDNA secretion related to embryo chromosomal self-correction and/or apoptosis?
3. What are the reasons that the noninvasive model will prevail or not over the invasive model?
4. **niPGT-A still an expensive approach**



The Multi-omics approach?



Application to predict Repeated Implantation Failure (RIF)

compared between embryos from RIF patients (n=35) and oocyte donors as controls (n=15)

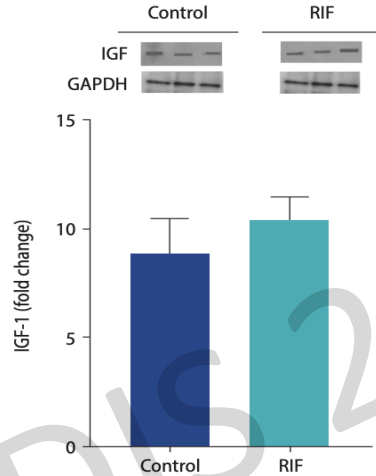


Figure 1. The insulin-like growth factor 1 (IGF-1) protein expression in culture medium from the repeated implantation failure (RIF) and control groups. Values are presented as mean \pm standard deviation of at least three replicates. GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

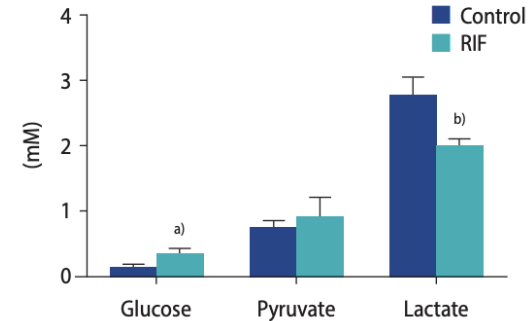


Figure 2. Glucose, pyruvate, and lactate concentrations in the repeated implantation failure (RIF) and control groups. Values are presented as mean \pm standard deviation of three replicates. ^{a)} $p<0.05$; ^{b)} $p<0.01$.



Metabolic profiling-(niPGT?)

Hypothesis

- Embryo secretes DNA, RNA and protein into the spent culture medium
- The identities and quantities of these molecules reflect the **health** and **birth potential** of the embryo

PNAS

Proceedings of the
National Academy of Sciences
of the United States of America

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

Juanjuan Xu, Rui Fang, Li Chen, Daozhen Chen, Jian-Ping Xiao, Weimin Yang, Honghua Wang, Xiaoqing Song, Ting Ma, Shiping Bo, Chong Shi, Jun Ren, Li-Yi Cai, Bing Yao, X. Sunney Xie, and Sijia Lu

PNAS October 18, 2016 113 (42) 11907-11912; published ahead of print September 29, 2016
<https://doi.org/10.1073/pnas.1613294113>

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

Noninvasive metabolomic profiling of embryo culture media using Raman and near-infrared spectroscopy correlates with reproductive potential of embryos in women undergoing in vitro fertilization

Emre Seli, M.D.,^a Denny Sakkas, Ph.D.,^a Richard Scott, M.D.,^b Shing C. Kwok,^c Scott M. Rosendahl,^c and David H. Burns, Ph.D.^c

^aDepartment of Obstetrics, Gynecology, and Reproductive Sciences, Yale University School of Medicine, New Haven, Connecticut; ^bReproductive Medicine Associates, Morristown, New Jersey; and ^cDepartment of Chemistry, McGill University, Montreal, Quebec, Canada



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Seli et al., 2007 Fertil Steril. 88(5):1350-1357.
Xu et al., 2016 Proc Natl Acad Sci. 113(42):11907-11912.

Previous Embryonic Metabolomic Profiling

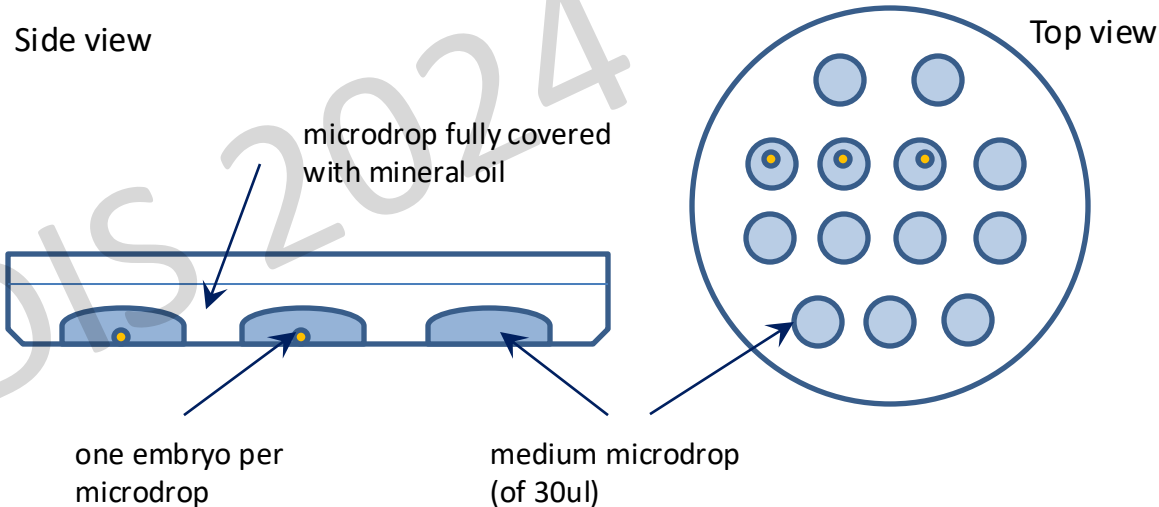
A list of the 92 metabolites found in the spent media of day 3 human embryos.

Super pathway	Metabolite	Fold change T vs. N	P value
Peptide	Glycylglutamine	1.01	.2668
Carbohydrate	Sucrose	1.67	.6042
	Glucose	1.31	.5945
	Pyruvate	0.96	.9970
	Lactate	1.01	.3787
Energy	Citrate	0.76	.5763
Lipid	Linolenate [alpha or gamma; (18:3n3 or 6)]	0.95	.7597
	Caproate (6:0)	1.14	.0382
	Heptanoate (7:0)	0.99	.8784
	Caprylate (8:0)	1.02	.1597
	Pelargonate (9:0)	0.98	.8312
	Caprate (10:0)	0.95	.8317
	Laurate (12:0)	1.04	.4924
	Myristate (14:0)	0.92	.0672
	Myristoleate (14:1n5)	1.05	.8060
	Oleate (18:1n9)	0.93	.1038
	Linoleate (18:2n6)	0.92	.0609
	Dihomo-linoleate (20:2n6)	0.93	.1673
	Arachidonate (20:4n6)	0.93	.2915
	Choline	1.01	.6761
	Glycerophosphorylcholine	1.01	.5283
	Dehydroisoandrosterone sulfate (DHEAS)	1.02	.7595
	Epiandrosterone sulfate	1.00	.5858
	Androsterone sulfate	1.06	.0229
	4-androsten-3beta, 17beta-diol disulfate 1	1.01	.4111
	4-androsten-3beta, 17beta-diol disulfate 2	1.05	.2181
	Pregnen-diol disulfate	1.01	.8393
	Pregn steroid monosulfate	0.97	.2769

Metabolomic profiling by liquid chromatography mass spectroscopy of 15 SCMs from Trisomy 21 Day 3 embryos and 15 controls.

Methods & Materials

- SCM sample collection
 - Single embryo culture
 - Sequential (G-2) or One-step (G-TL) medium
 - Collect the culture medium and store at -20 °C

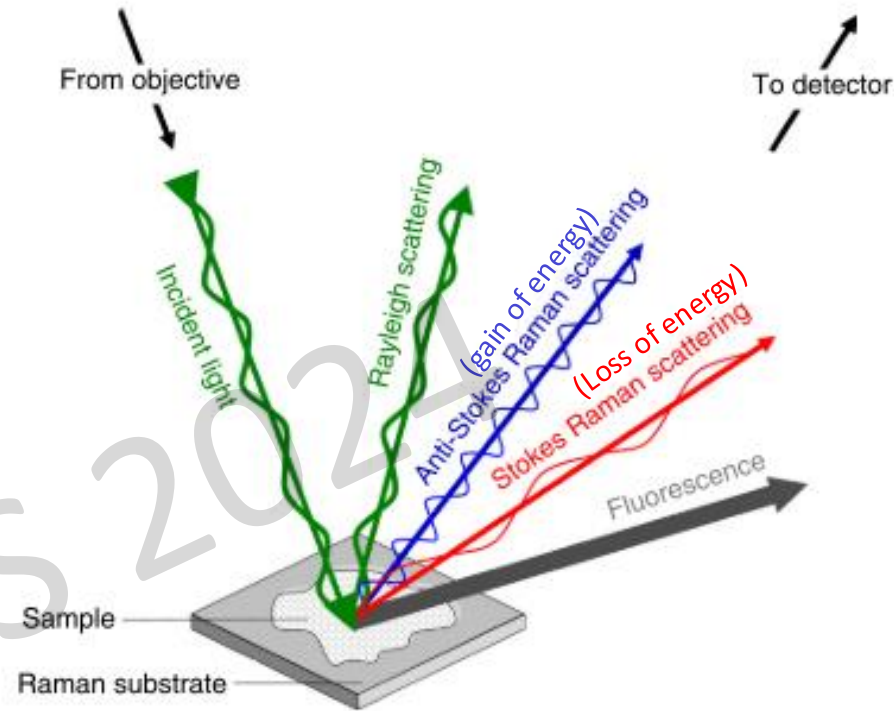


Hypothesis

- Non-invasive Preimplantation Genetic Testing
 - Spent Culture Med

To assess the Aneuploidies or implantation potential of an embryo by Metabolomic Profiling by Raman spectroscopy?

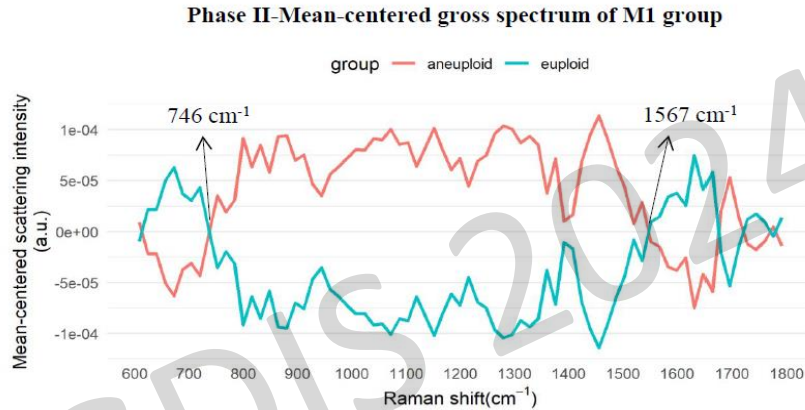
What is Raman spectroscopy?



Obtain instant result in
A few seconds

Objectives

Is Raman spectroscopy combining machine learning applicable for different types of culture media as a first-tier non-invasive screening test for aneuploidies?



Differential scattering intensity across Raman spectrum, classified by PGT-A result on trophectoderm biopsies

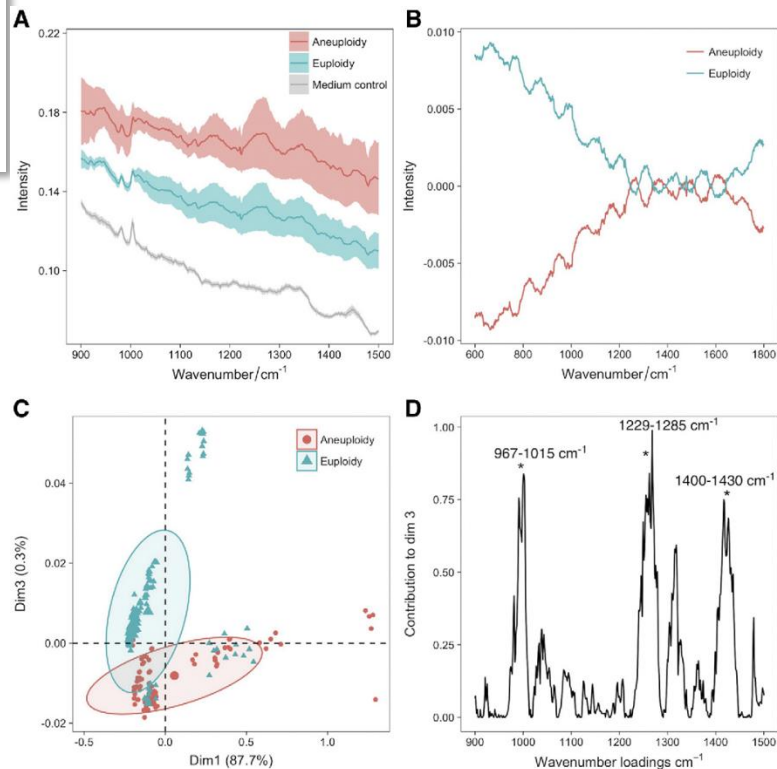


Raman 200 machine

Raman profiling of embryo culture medium to identify aneuploid and euploid embryos

Unique embryonic metabolomic profiling

Mean-centered Raman spectra demonstrated differential intensities of SCMs from euploid and aneuploid among Day 5 embryos



Message from this paper:

- Proof-of-concept of Raman combing machine learning for ploidy prediction
- One type of medium
- Validation size need to be expanded
- Reproducibility

Confusion matrix and performance evaluation of kNN, RF, XGB, and stacking classification models for an independent testing set of 222 Raman spectra.

Model	Actual class	Confusion matrix		Performance evaluation			
		Predicted euploidy	Predicted aneuploidy	Precision	Sensitivity	F1 score	Accuracy
kNN	Euploidy	117	8	96.7%	93.6%	0.951	94.6%
	Aneuploidy	4	93	92.1%	95.9%	0.939	
RF	Euploidy	72	53	78.2%	57.6%	0.664	67.1%
	Aneuploidy	20	77	59.2%	79.4%	0.678	
XGB	Euploidy	116	9	92.8%	92.8%	0.928	91.9%
	Aneuploidy	9	88	90.7%	90.7%	0.907	
Stacking	Euploidy	121	4	96.0%	96.8%	0.964	95.9%
	Aneuploidy	5	92	95.8%	94.9%	0.953	

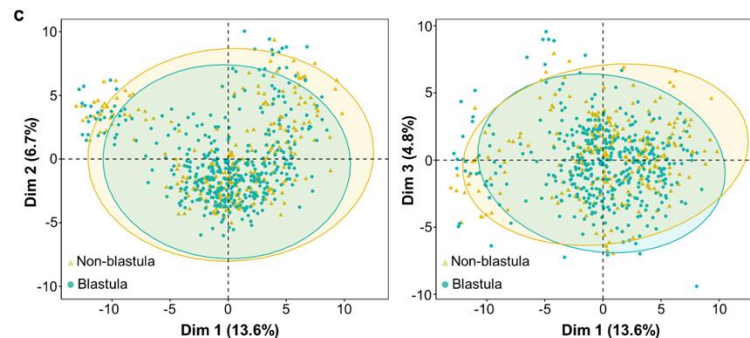
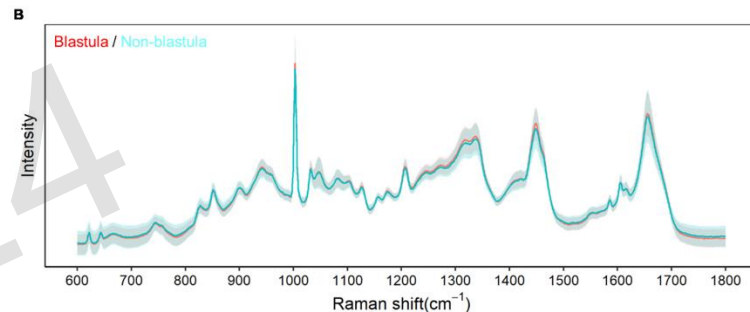
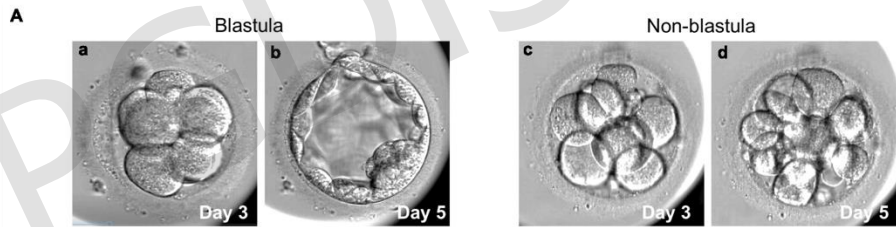
Note: Models were trained from a training set of 885 Raman spectra. kNN = k-nearest neighbors; RF = random forests; XGB = extreme gradient boosting. Stacking analysis is based on a first layer of kNN, RF, and XGB and a second layer of XGB.



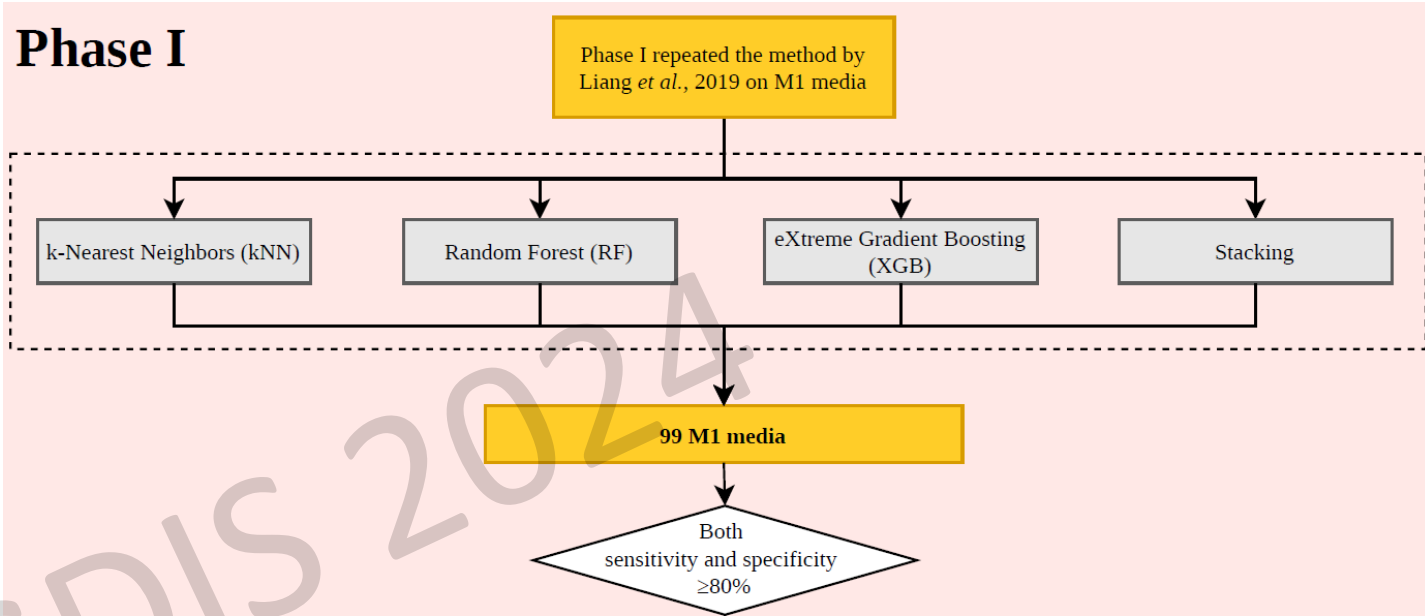
Metabolic profiling predicts the blastocyst development potential

Non-invasive Metabolomic Profiling of Embryo Culture Medium Using Raman Spectroscopy With Deep Learning Model Predicts the Blastocyst Development Potential of Embryos

prediction sensitivity 73.35%

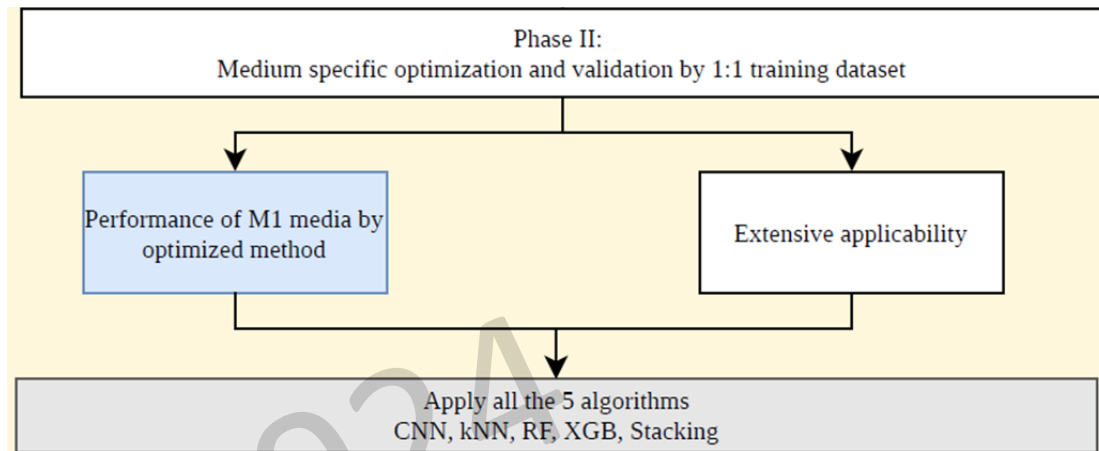


Reproducibility study



Phase I completely repeat the method by Liang *et al.* to test the reproducibility.

Aim: Optimization and Validation Study



Optimization by:

1. Testing wet lab improved quality control standard
2. Algorithms optimized by re-seeking hyperparameters
3. Euploid & aneuploid for training re-shuffled and retrained
4. Added Convolutional Neural Networks (CNN)

Phase II – Optimization and Validation

Re-trained & tested M1 media

Phase II – M1 Training (30 euploid vs. 30 aneuploid)

	CNN	kNN	RF	XGB	Stacking
Sensitivity	86.7% (26/30)	93.3% (28/30)	76.7% (23/30)	86.7% (26/30)	93.3% (28/30)
Specificity	83.3% (25/30) ✓	76.7% (23/30)	76.7% (23/30)	100.0% (30/30) ✓	83.3% (25/30) ✓

Stacking out

Phase II – M1 Testing (43 euploid vs. 56 aneuploid)

	CNN	kNN	RF	XGB	Stacking
Sensitivity	75.0% (42/56)	73.2% (41/56)	69.6% (39/56)	76.8% (43/56)	80.4% (45/56) ✓
Specificity	83.7% (36/43)	83.7% (36/43)	81.4% (35/43)	100.0% (43/43)	81.4% (35/43)



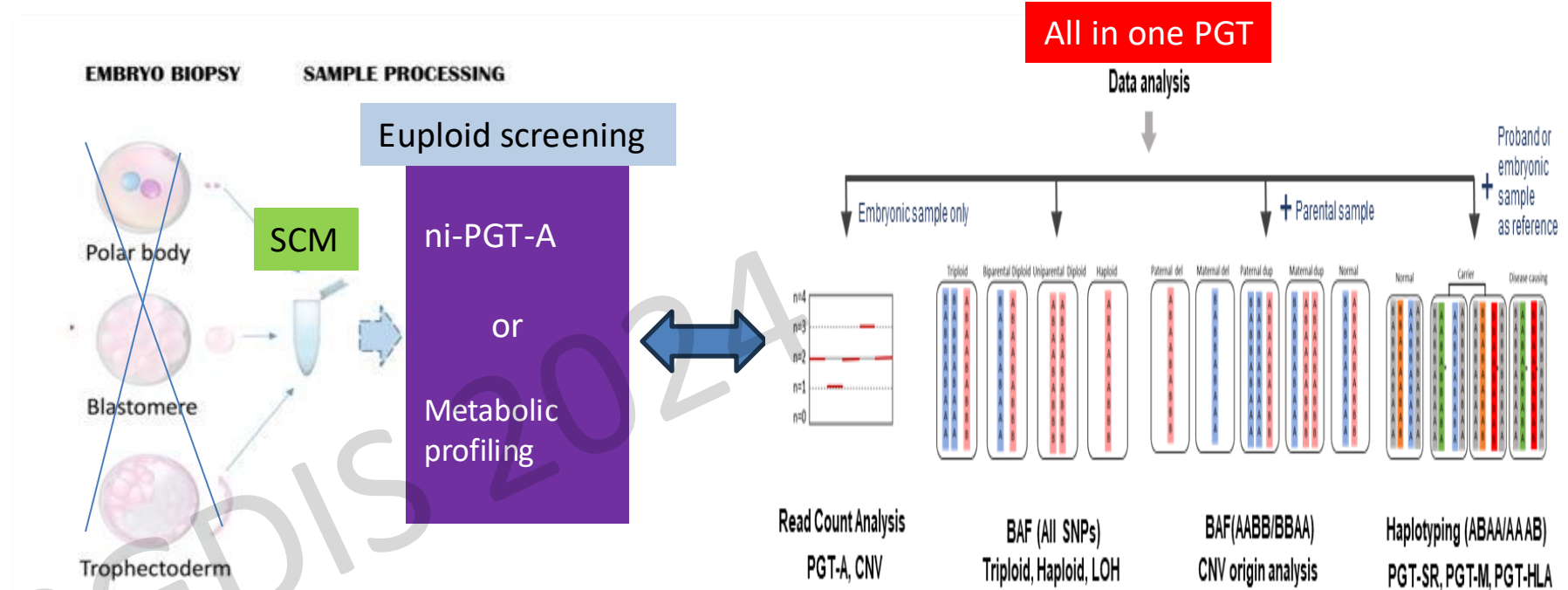
Table 3. Performance of Raman spectroscopy pregnancy outcome.

	Euploid transfers by TE biopsy (N=102)	Euploid transfers by Raman* (hypothetical, N=97)		
	n (%)	n (%)	Sensitivity	Specificity
Implantation failure	32 (31.4%)	30 (30.9%)	93.8%	95.7%
Biochemical pregnancy	8 (11.4%)	8 (11.9%)	72.7%	100%
Miscarriage (per clinical pregnancy)	12 (19.4%)	11 (18.6%)	91.7%	96.0%
Ongoing pregnancy/Live birth	50 (49.0%)	48 (49.5%)	96.0%	100%

*: Stacking algorithm was adopted here considering the best performance shown in our current study.



New algorithm for PGT-A



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General Research Fund (GRF)
Collaborative Research Fund


UGC 大學教育資助委員會
University Grants Committee

Thank You!!



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Genetic Counselling Society of Malaysia

IMPORTANT DATES

Abstract submission ✓

by May 20, 2024

Early bird registration ✓

by Jun 20, 2024



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**PGT and
BEYOND...**