ArgenTAG - IDEA² Progress Report: Oct/Nov 2022

Goals

- Use case description with a higher level slide
- Target market redefinition
- Fundraising

Potential customers to try tech - Business

- Achieved our goal of 10 LOIs
- Preparing 3 collaborations. Working with legal departments of: NYU, Weill Cornell, Mt Sinai

Fundraising

~90 mostly US biotech investors, ~20 were interested and asked for more info ~10 of them arrived at the Data Room stage ~4 conversations on going Est, closing date: End of January

Proof of Concept

70% progress

Early adopter program

3 projects, 30% progress

Acknowledgments: Francisca Mulero & Arthur J. Hiller, IDEA² 2022 advisors

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Next-generation single-cell sequencing.



Cells are the basic unit of life

- Each of us is a set of **37 trillion** cells.
- Understanding how cells work could reveal hidden mechanisms of critical human diseases.





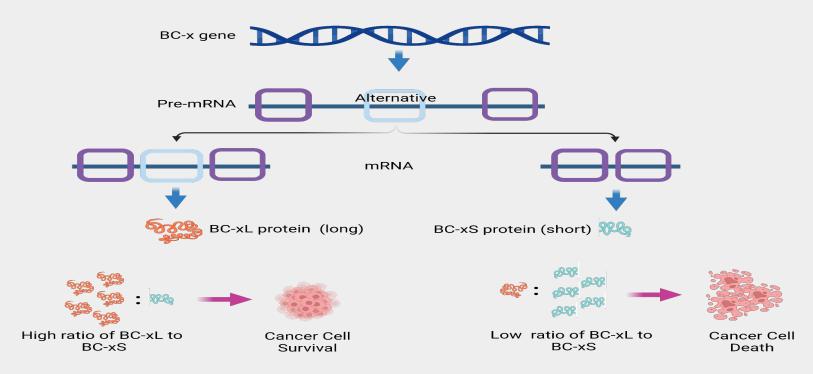
Single-cell methods show the state of cells

- Cells turn genes *on* and *off* in a combinatorial way to achieve diversity (≅ 200 cell types).
- Genes on make RNA molecules (≅ 300,000/cell).
- RNAs are associated with cell functions.
- RNAs are 'read' by DNA sequencing machines.





A human gene can make more than one RNA. So, genes may have more than one function

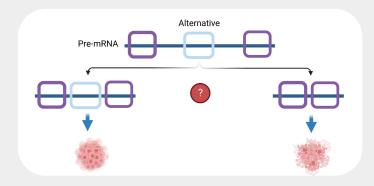


Stevens, M., & Oltean, S. (2019). Modulation of the apoptosis gene Bcl-x function through alternative splicing. Frontiers in Genetics, 10, 804.

Current single-cell methods are *low resolution*, they read only the s*tart* or *end* of RNAs

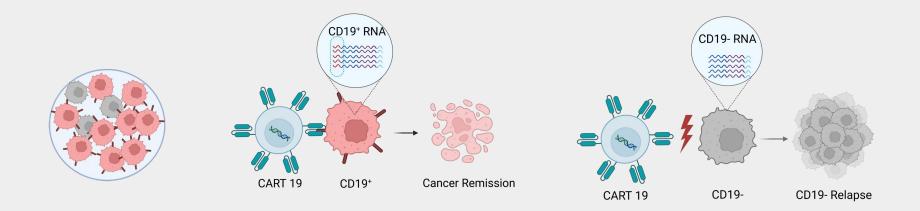
- Technically, 80% of human genes make more than one type of RNA but current methods only read one type of RNA per gene.
- New single-cell methods based on long read sequencers could read the complete RNA landscape...





Case study: Cancer relapse prediction in CAR T-cell therapy is a challenging task

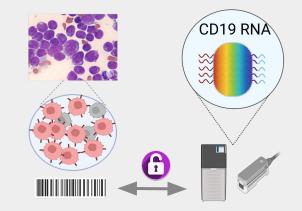
B-cell Acute Lymphoblastic Leukemia: 60-70% remission followed by 30-50% relapse.



Rabilloud T. et al. Single-cell profiling identifies pre-existing CD19-negative subclones in a B-ALL patient with CD19-negative relapse after CAR-T therapy. Nat Commun. 2021 Feb 8;12(1):865.

Use case: Improve cancer relapse prediction in CART 19 treatments with a high resolution single-cell method

Challenge: design millions of robust DNA barcodes to identify RNAs of cells to be sequenced together with noisy long-read sequencers.



ArgenTAG high-resolution single-cell kits unlock the hidden 80% of human RNA landscape

Powered by a <u>proprietary barcoding</u> technology and a <u>device-free</u> protocol for long-read sequencers.



ArgenTAG patent application protects our **barcoding technology**

- DNA barcode composition follows Digital Communication principles to withstand long-read sequencing noise (e.g., Nanopore, PacBio).
- DNA barcode design & deconvolution are scalable enabling highly multiplexed long-read applications beyond single-cell.



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Ezpeleta, J. *et al.* **Robust and scalable barcoding for massively parallel long-read sequencing**. *Sci Rep* **12**, 7619 (2022).

Initial market efforts will focus on single-cell long-read opportunity

- ArgenTAG will enter long-read sequencing market with a high-resolution single-cell sequencing kit
- 2. Priority will be to consolidate long-read seq with protocol automation
- 3. Longer term vision will be to expand long-read sequencing market towards precision medicine applications.

~\$ 100 B

Life Science research tools, 2030, CAGR 11%

\$ 11,1 B

TAM > Consumables/reagents in the Global DNA sequencing market, 2030, CAGR 18%

\$ 2,3 B SAM, Long-read sequencing customers * \$ ArgenTAG kit * frequency, 2030

\$ 135 M SOM - Capture 20% of US long-read sequencing users, 2030

https://www.decibio.com/market-reports/next-generation-sequencing-ngs-market-assessment-report https://www.grandviewresearch.com/industry-analysis/dna-sequencing-market

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Source:

A family of kits will be offered to a full range of customers

> Pharmas, Government

Purpose: Cell therapies, population genomics Avg. expenditure: US\$ 729,640

Total customers 2021: 782 (PacBio + Nanopore)

US\$ 10,000 Production Large-scale Kit

> Biotechs, Core facilities

Purpose: Drug discovery Avg. expenditure: US\$ 78,416

Total customers 2021: 430 (Nanopore)

US\$ 5,000

Expert Medium-scale Kit Collaborations, Custom projects, Services

> Academic Labs, Pls

Purpose: Basic Research Ava. expenditure: US\$ 6,728 Total customers 2021: 5,501 (Nanopore) **US\$ 2,500** Exploratory Small-scale Kit

ArgenTAG is the first-to-market native solution for single-cell high resolution

	ArgenTAG	Parse Bio	10x Genomics		
Product	Kits	Kits	Instrument		
High Resolution	Yes	No	No		
Scalable	Yes	Yes	No		
Automation	No	No	Yes		
Quick results	Yes	No	No		
Price	\$5 to \$15k per kit, then a long-read seq run	\$5 to \$15k per kit, then a short-read seq run	Capex: \$35k to \$260k, \$30 to 50k in kits to run experiments. A service		

We are addressing a significant unmet need in the market



- Signed 10 LOIs with academic research centers and biotechs.
- Received +25 inbound and organic messages asking for services

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	 P	thir	Briefly tell us about your project: We are a lab						
	Р	of y also	that does a lot of 10	x Genomics	s based paired B				
		lot	cell sequencing. Will	your techn	ology work for				
			our purpose?						

Preparing strategic collabs and building relations with leaders

Potential collabs	Building relationships			
Icahn School of Medicine at Mount Sinai	ThermoFisher SCIENTIFICPacBiSCIENTIFICTWIST BIOSCIENCE			
Patch Biosciences	64x Bio Cellino			
Singler®n STAMM	UNOVARTIS BIOMAKERS [®]			

Working with the legal departments of Mt Sinai and Weill Cornell Medicine

Harvard Single-cell core facility and Patch Biosciences waiting to try our kits

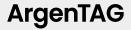
Early conversations with Singleron, Stamm, Thermo Fisher for co-development.

We met with C-level / Director of key companies in the US

Met with Nanopore CEO and he offered us native support and potential opportunity to commercialize through them

Native support





A team of engineers, biotechnologists and molecular biologist with +20y of experience pushing the limits of biotechnology



CFO

ELIZABETH TAPIA, Ph.D.

Engineering

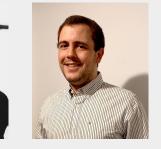


CTO

Engineering

COO PILAR BULACIO, Ph.D. LEANDRO CIAPPINA, MBA

Business



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Dir. Process Dev

SOFIA LAVISTA, Ph.D.

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ADVISORS

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Scientific Advisor LUCIANO MARTELOTTO, Ph.D. Head Single-cell Adelaide University

Scientific Advisor **GEORGE CHURCH, Ph.D.** Lead, Wyss Institute Harvard University

INVESTORS

IndieBio / SOSV **Draper University Ventures** Grid Exponential

Seed round to kick off collabs, deliver the first kit to early adopters and achieve product-market fit

	2021 - 22	2023			2024				
	2021 - 22	lQ	2Q	3Q	4Q	1Q	2Q	3Q	4Q
Sw Dev		Development and testing of single-cell kits			Early Access Phase				
Kit Dev	-Darwin Software. Fast barcoding analysis	Developer Phase (Achieve 1st MVP to collect data)			Early Access Phase - US customers				
Collabs	 Published our core tech in Scientific Reports. ArgenTAG's validation. Signed 10 LOIs 	dation. 1 to 3 collaborations to publish (Weill Cornell, Mt Sinai, HMS, NYU, Adelaide)				1 partnership with a key player (Nanopore, PacBio)			
Co-dev	- Proprietary LR barcoding protocol in real cells. 60% progress.								
Pilot Test	- UMI synthesis on Oligo-Mix. - Taggy Software. Scalable barcode			5 to 10 pilot testers - send a kit - to collect data (Patch, Quantumcyte, ImYoo)					
Sales	- Taggy Software. Scalable barcode design and deconvolution for LRs. - License granted, IP process initiated			pre-orders		US Paying Early Adopters Target 45 kits + Booked revenue with distributors 500 kits			500 kits
R&D	in Europe and US,	Single-cell Instrument Project							
	\$ 575,000 funding	SEED ROUND				Series A			

ArgenTAG

> First high-resolution single-cell solution that will enable scientist to reveal the 80% hidden RNA landscape of human cells - to positively impact new applications in oncology, immunology, and neuroscience.

