Myllia Company Presentation

Austrian Life Science Day

Thomas Moser, PhD | CEO May 2023

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Myllia Biotechnology - Executive Summary

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- Myllia Biotechnology has been founded in Feb 2018 in Vienna, Austria by experienced serial entrepreneurs
- Myllia capitalizes on one of the most important trends in modern life sciences. The
 single-cell analysis revolution raises biology to the next level, transforming it into a
 data-intensive industry with extremely high throughput.
- Myllia combines advanced single-cell transcriptomics read-out with challenging, physiologically relevant models like primary T-cells or iPSCs
- One of the largest teams worldwide focussing exclusively on next-generation singlecell CRISPR screens
- **Fully integrated workflow** from design and implementation of the screen to full bioinformatics analysis and visualization, all performed in-house.
- Conclusion of several large partnerships with big-pharma companies in US, UK and JP, typically in the area of target discovery and MoA analysis
- Business model focussed on revenue-generating partnerships and collaboration deals.



The Team myllia.



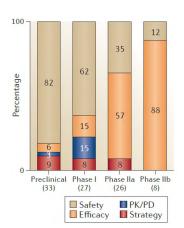
Team of 40+ Scientists

Dedicated wet-lab and computational biology teams

The quest for better drug targets

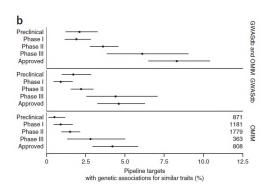
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Drugs often fail for lack of efficacy



Cook, D., Brown, D., Alexander, R. et al. Lessons learned from the fate of AstraZeneca's drug pipeline: a five-dimensional framework. *Nat Rev Drug Discov* **13**, 419–431 (2014). https://doi.org/10.1038/nrd4309

Drugs hitting genetically validated targets are more successful



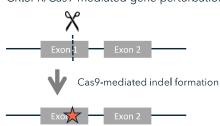
Nelson, M., Tipney, H., Painter, J. et al. The support of human genetic evidence for approved drug indications. *Nat Genet* **47**, 856–860 (2015). https://doi.org/10.1038/ng.3314

What is a CRISPR screen? myllia.

Gene products are drug targets



CRISPR/Cas9-mediated gene perturbation



CRISPR screen: Unbiased gene perturbation





Positive Selection Resistance Screens



Negative Selection Dropout Screens



CRISPR screening with single-cell RNA sequencing read-out (CROP-Seq)

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Cancer Cell Line



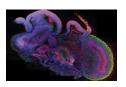
iPSC



Primary T cell



iPSCderived cell



Organoid

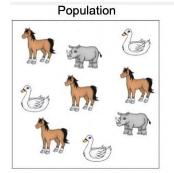
Complexity of the cellular model

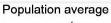


Bulk Screens



Single-cell Screens







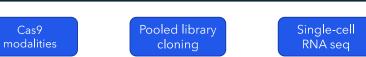
Meshal Ansari and Fabian Theis

The Myllia technology platform!













panels





Drug target identification and validation



Mode of action analysis



Identification of disease-associated gene function



Genetic screening for phenotypes of primary T cells

Our Location myllia.





Vienna Biocenter Campus

3 world-class academic institutes (IMP, IMBA, MPL) 35+ Biotech Companies 1.500 Biomedical Scientists Access to Core Facilities (NGS and FACS)

Headquarter at DOCK27

~600m2 lab space

~400m2 office space

Newly furnished and newly equipment Close to city centre (15min), airport (20min) and Vienna Biocenter (10min)

Founding Member of



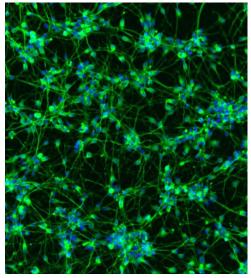
GSK has entered into a collaboration with Myllia Biotechnology, applying Myllia's single-cell CRISPR screening platform for drug target discovery in the following areas:

- 1. CRISPR screens in primary human T cells to uncover novel targets in autoimmune disease and immuno-oncology
- 2. CRISPR screens to uncover novel targets in solid tumours
- 3. CRISPR interference screens to modulate target gene expression



Our first child: bit.bio discovery!





Key Facts

- Joint Venture between Myllia Biotechnology (based in Vienna/ Austria) and bit.bio (based in Cambridge/ UK)
- Focus on induced pluripotent stem cells and cells obtained by directed differentiation
- Combines unbiased CRISPR screens with the best human cellular models available.

Unique selling points

- Best-in-class cellular models
- Custom isogenic disease models
- Unprecedented robustness and consistency
- CRISPR screens with single-cell resolution

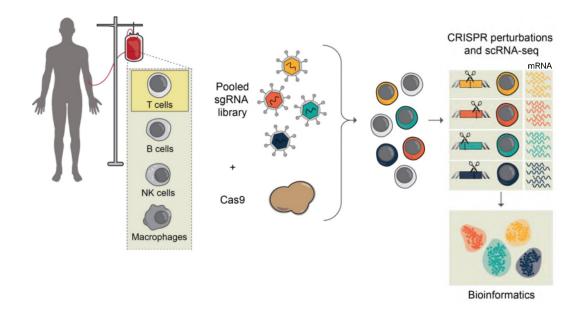


Case Study:

TCR signalling in primary human T cells

CROP-Seq screens in human primary T cells

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CROP-seq experiment in pan T-cells

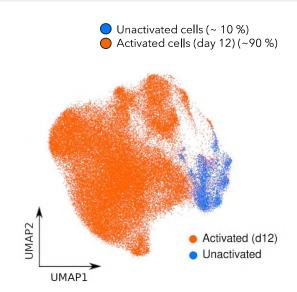
- Unactivated cells
- Cells activated with CD3/CD28-Dynabeads for 12 days

Number of perturbed genes

- 42 genes
- 188 sgRNAs (4 sgRNAs/ gene)

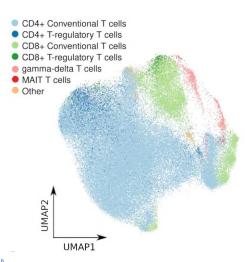
Read-out by targeted single-cell RNA sequencing of 300 mRNAs $\,$

- T cell activation markers
- Markers for cell types
- Cell cycle markers
- Immune checkpoint genes



A CROP-seq screen in 100,000 primary human T cells

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Which T cell subsets do we see?

Which effector types do we see?

