Advancing High-Throughput Organoid Drug Screening: Automated Live-Cell Imaging and the Orbits Image and Data Analysis Pipeline

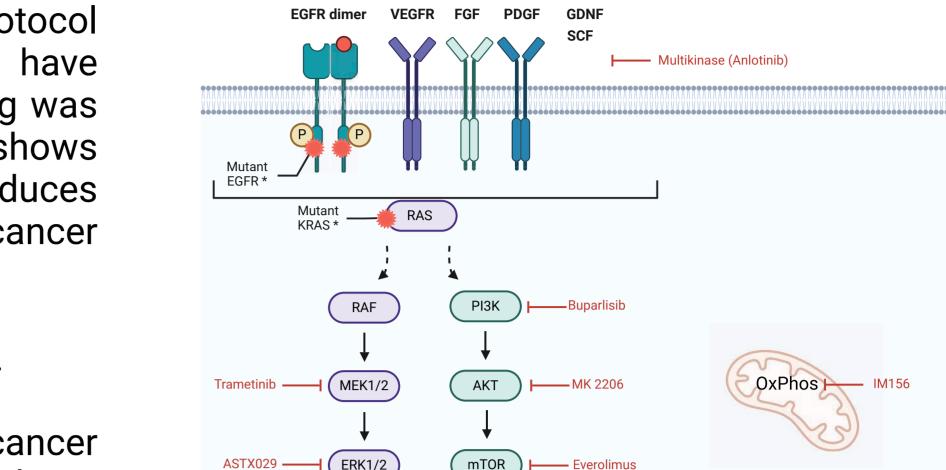
Christophe Deben^a, Edgar Cardenas De La Hoz^b, Maxim Le Compte^a, Abraham Lin^a, Felicia Rodrigues Fortes^a, Sofie Seghers^a, Geert Roeyen^{c,d}, Filip Lardon^a

a. Center for Oncological Research (CORE), University of Antwerp, Belgium / b. Industrial Vision Lab, University of Antwerp, Belgium / c. Department of Oncology, Antwerp University Hospital, Belgium / d. Department of Hepatobiliary Transplantation and Endocrine Surgery, University Hospital Antwerp, Belgium

INTRODUCTION

The dynamic and complex nature of biological systems has sparked a transition away from traditional flat cell cultures toward 3D organoid models. Although organoids better represent in vivo conditions, their high-throughput screening poses challenges, primarily due to intricate morphology and non-homogeneous responses. To address this, we've innovated a protocol for high-throughput drug screening with widefield microscopes, distinguishing cytostatic and cytotoxic responses at the organoid level¹⁻³.

OBJECTIVES



To test the high-throughput compatibility of our protocol and image and data analysis platform, we have performed a drug screening for Auranofin. The drug was initially approved for rheumatoid arthritis and shows promise as an anti-cancer agent. Auranofin induces oxidative stress and disrupts the redox balance in cancer cells, leading to cell death. In this study we aimed to:

- Identify the best organoid drug response metric.
- Study the selectivity of Auranofin towards cancer

To overcome the challenges related to high-throughput image and data analysis, we have developed 'Orbits', an innovative platform that integrates label-free Al image analysis with in-house developed drug response metrics tailored to high-throughput organoid drug screening and prediction of patient therapy response. cells by including normal epithelial lung organoids.

 Identify the most promising combination strategies based on selectivity and synergy. And overview of the selected compounds is shown in Figure 1.

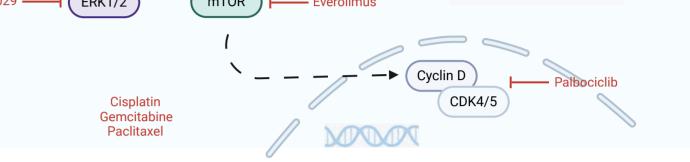


Figure 1: Overview of the compounds that were combined with Auranofin.

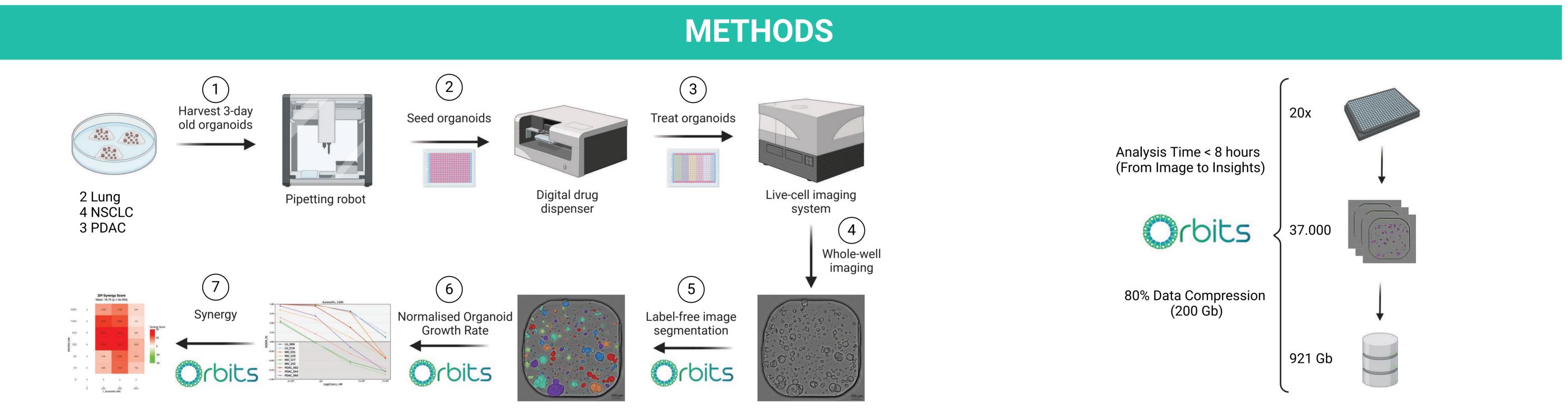


Figure 2: Workflow overview. Le Compte et al., JoVE, 2022.

Figure 3: Image and data analysis pipeline.

RESULTS

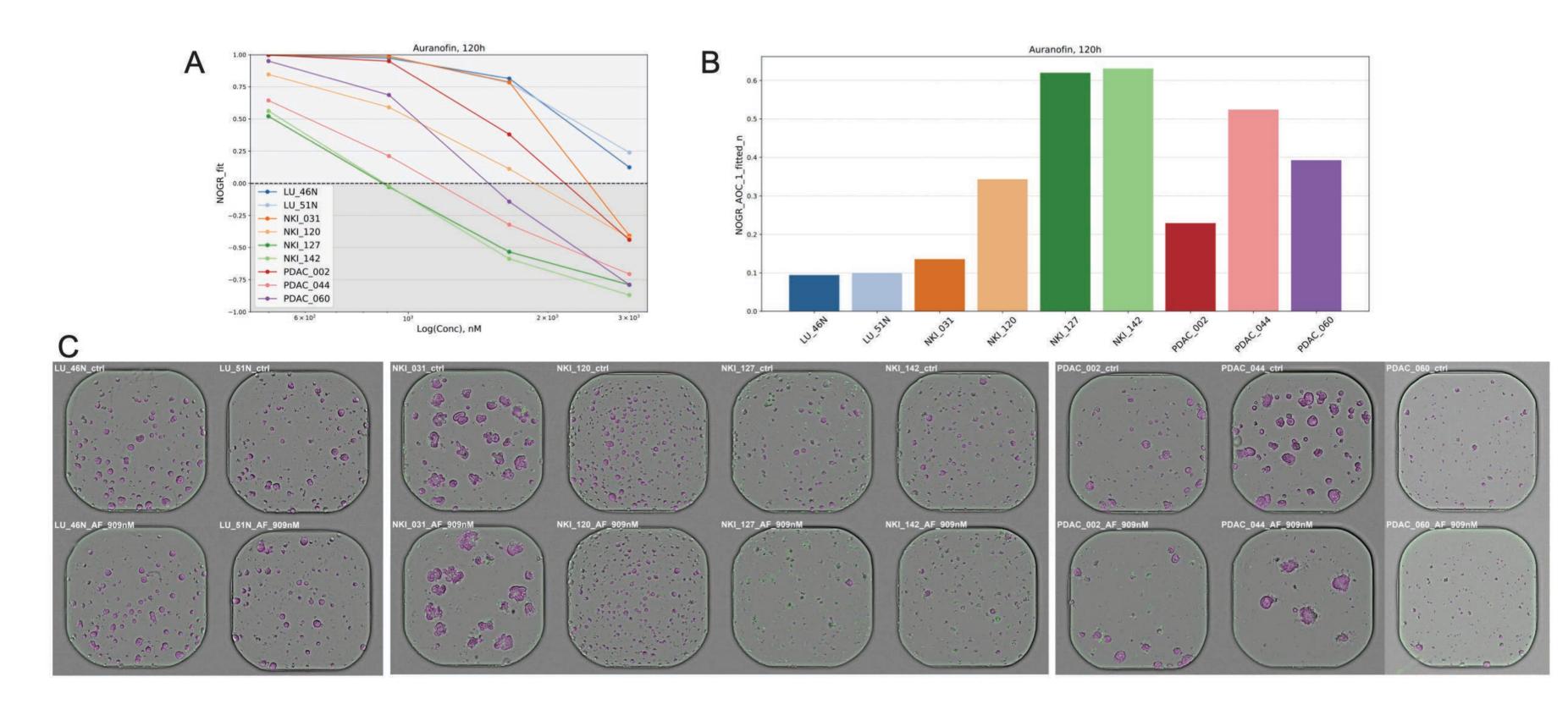


Figure 4. (A) Fitted dose-response curves of the Normalised Organoid Growth Rate (NOGR) for Auranofin monotherapy. (B) Area Over the Curve (AOC) of the fitted dose-response curves. (C) Representative images of organoids treated with vehicle or 909 nM Auranofin for 120h. Magenta shows the label-free image segmentation by Orbits. LU_: normal lung organoids. NKI_: non-small cell lung cancer organoids, kindly provided by E. Voest, Netherlands Cancer Institute. PDAC_: pancreatic ductal adenocarcinoma organoids.

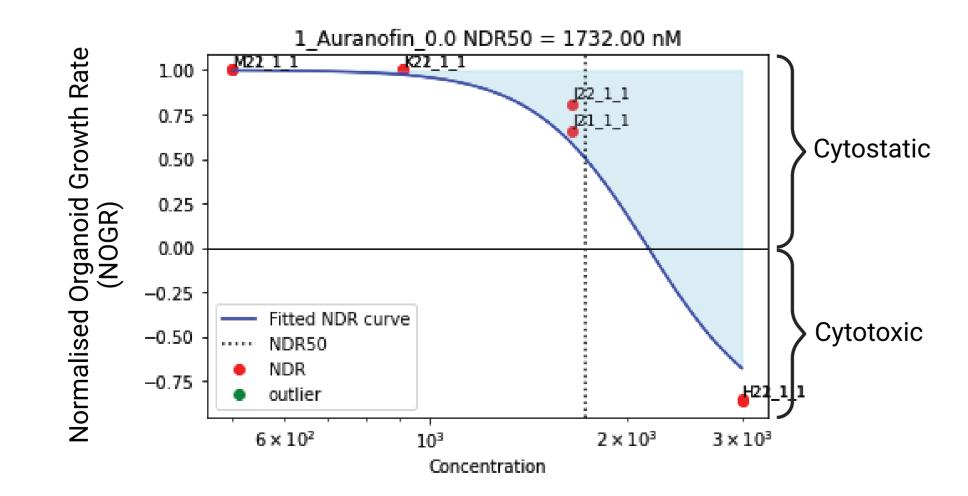
CONCLUSIONS

Drug Screening Pipeline:

- Established a major advancement in organoidbased drug discovery.
- Offered a robust protocol for high-throughput screening and automated data analysis.
- The Orbits Normalised Organoid Growth Rate (NOGR) significantly improves the identification of organoid drug responses and synergistic interactions from live-cell imaging.

Auranofin:

- Selective towards cancer cells compared to normal epithelial lung cells.
- Highly synergistic with the AKT inhibitor MK2206 in multiple tumor organoid lines.
- Additional organoid dependent synergistic interactions.



Auranofin: MK2206 909 nM

Figure 5: Normalised Organoid Growth Rate (NOGR). Blue area represents the Area Over the Curve (AOC).

Figure 6: Bubble plot visualizing the ZIP synergy score (bubble size) and NOGR value (colour heatmap) for the combination of Auranofin and MK2206 in all organoid lines.

CONTACT

Prof. Dr. Christophe Deben University of Antwerp, Belgium christophe.deben@uantwerpen.be

Orbits image and data analysis platform: www.orbits-oncology.com

Organoid drug screening services: www.drugvision.ai









