Development of a medium-to-high throughput organoid and bacteria co-culture platform for the assessment of host pathogen interaction



<u>Eider Valle-Encinas</u>¹, Roshni Nair ¹, Mayke Doorn¹, Katerina Pisa¹, Farzin Pourfarzad¹, Lani San Mateo², Nicole Desch², Prashanth Gokare², David Pocalyko², Sylvia F. Boj¹, and Carla Verissimo¹ ¹HUB Organoids (HUB), Yalelaan 62, ²Janssen R&D

Introduction

HUB's patient derived organoids, (HUB Organoids[®] or PDOs) are self-organizing epithelial cell structures with near-physiological features, extensively used to model aspects of cancer initiation and progression. Microinjection of colibactin-producing pks+ *E. coli* into the lumen of PDOs results in the appearance of two co-occurring mutational signatures identified in a subset of colorectal cancer (CRC) patients¹, demonstrating that pks+ *E. coli* plays a causative role in CRC development. However, the scalability of bacteria microinjection in PDOs is limited and represents a bottleneck in the screening of preventive therapies for patients. Here we developed a bacteria and PDO co-culture system (PDO fragment exposure model), alternative to PDO microinjection, that is compatible with medium-to-high throughput screening methods. We validated the genotoxicity of colibactin-producing bacteria and showed the potential of the PDO fragment exposure model for the screening of drugs targeting colibactin-dependent genotoxicity.

Fig 4. Colibacting-producting bacteria induces dna damage in the PDO fragment exposure model

Co-culture of PDOs fragments with *pks+ E. coli* (colibactin-producing bacteria) resulted in increased levels of yH2AX (proxy for DNA damage) in live single cells analyzed by flow cytometry. PDOs survive following co-culture with colibactin-producing bacteria.

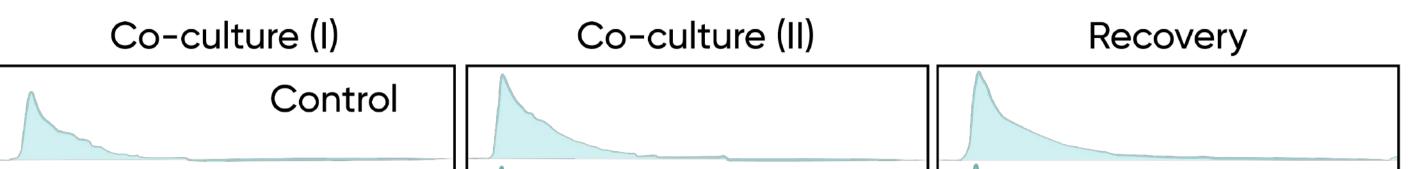


Fig 1. Development of the first PDO

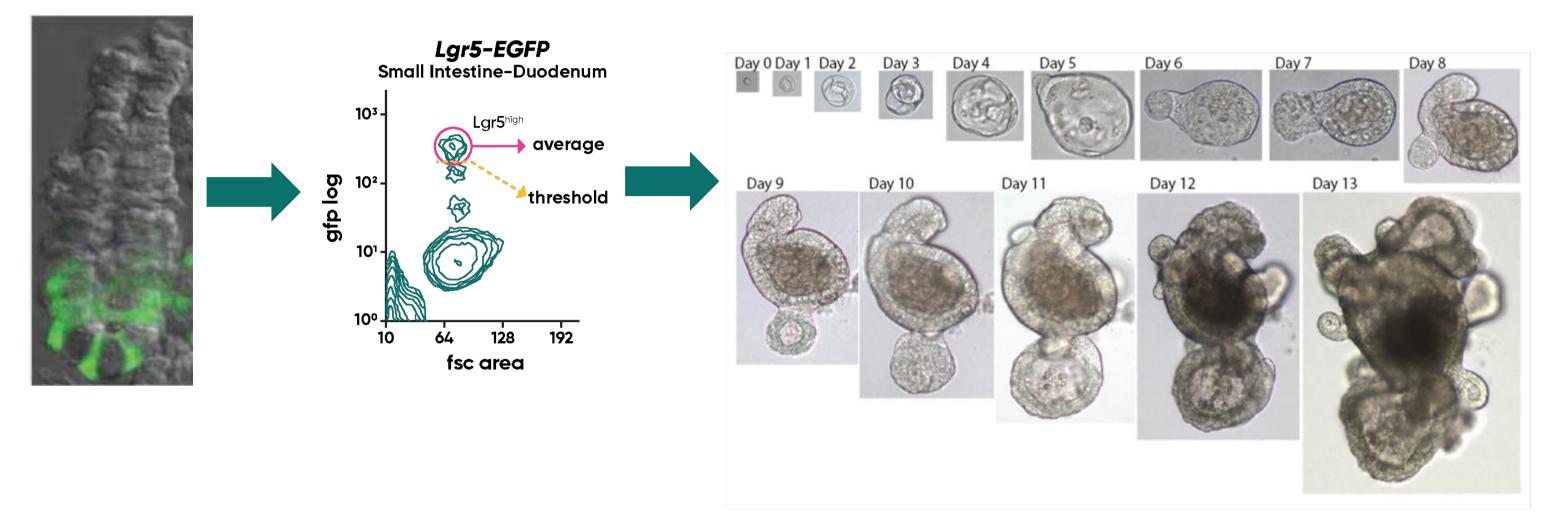


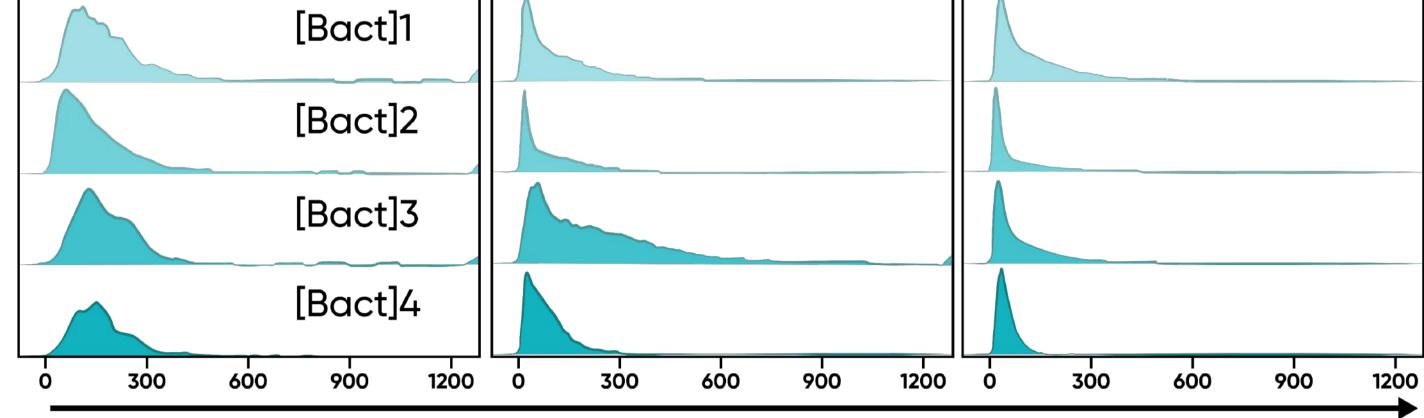
Fig 2. HUB Organoids for the development of therapies targeting pathogen-host interactions

Correlation between microbiome profiles and human disease



Complex microbiome ecosystem





DNA damage (yH2AX)

Control
[Bact]1
[Bact]2
[Bact]3
[Bact]4

Image: Image

Fig 5. DNA damage detected in PDO fragment exposure model is

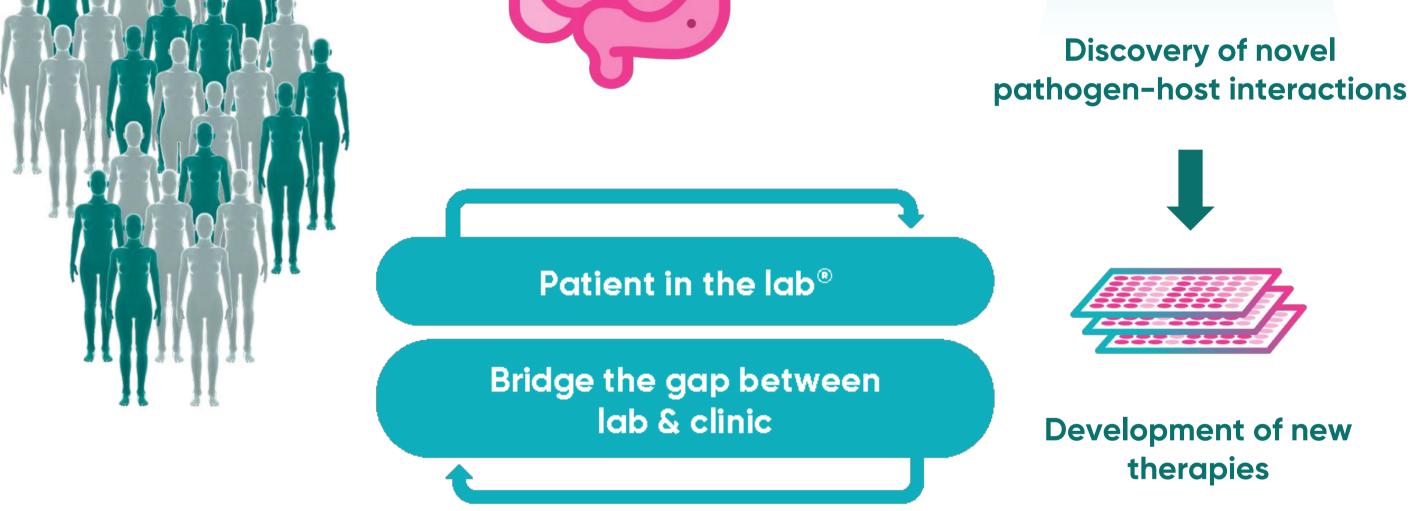
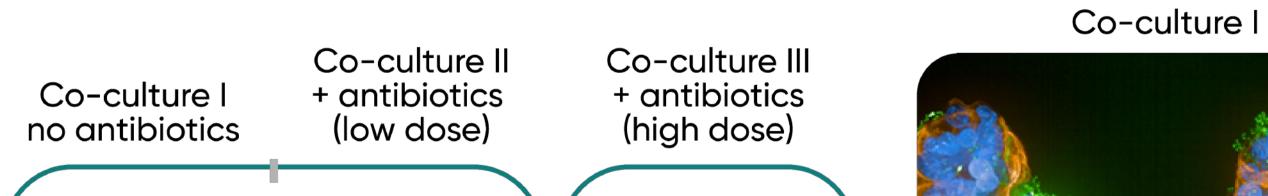


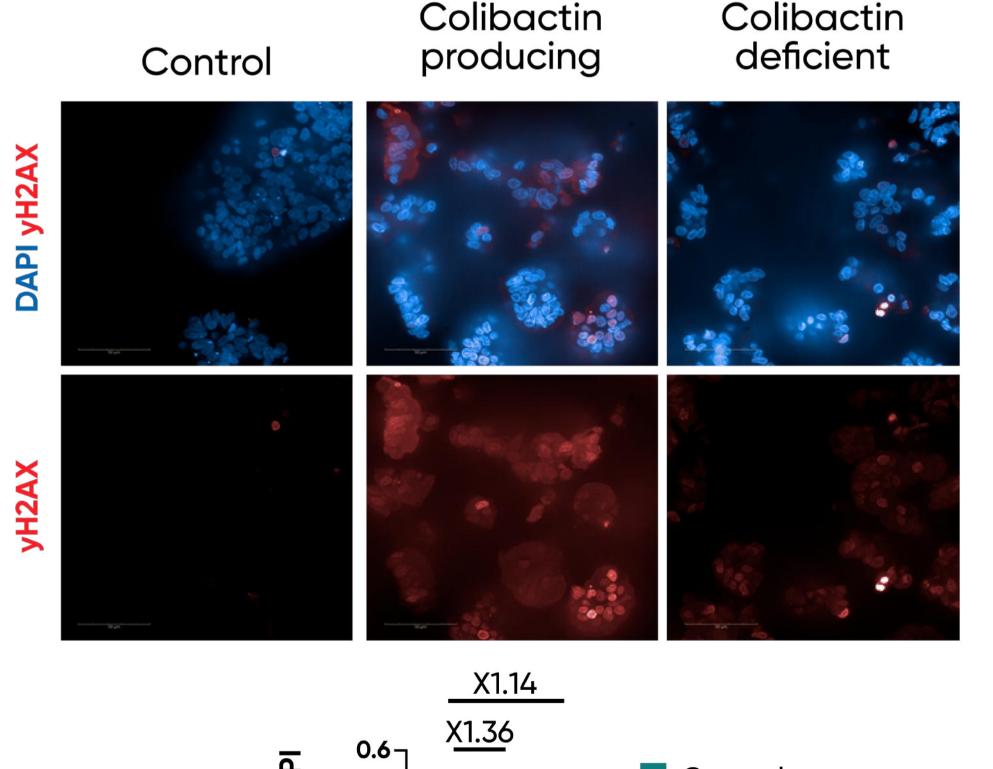
Fig 3. Strategy for PDO fragments and colibactin-producing bacteria co-culture

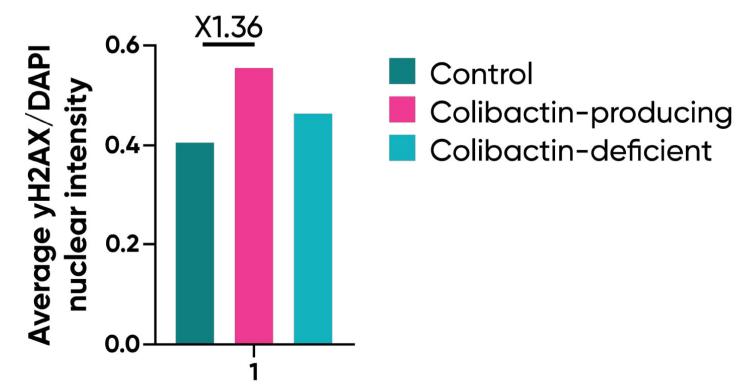
The PDO fragment exposure model consists of three phases: (a) Co-culture I or 'acute' DNA damage phase, where the bacteria and PDOs interact in suspension and the growth of the bacteria is not restricted; (b) Co-culture II or 'sustained' DNA damage phase, where PDOs and bacteria are cultured in hydrogels and bacteria growth is controlled by the addition of low concentrations of antibiotics; (c) Recovery phase, where the bacteria is killed by higher dose of antibiotics and PDOs continue to grow in hydrogels.

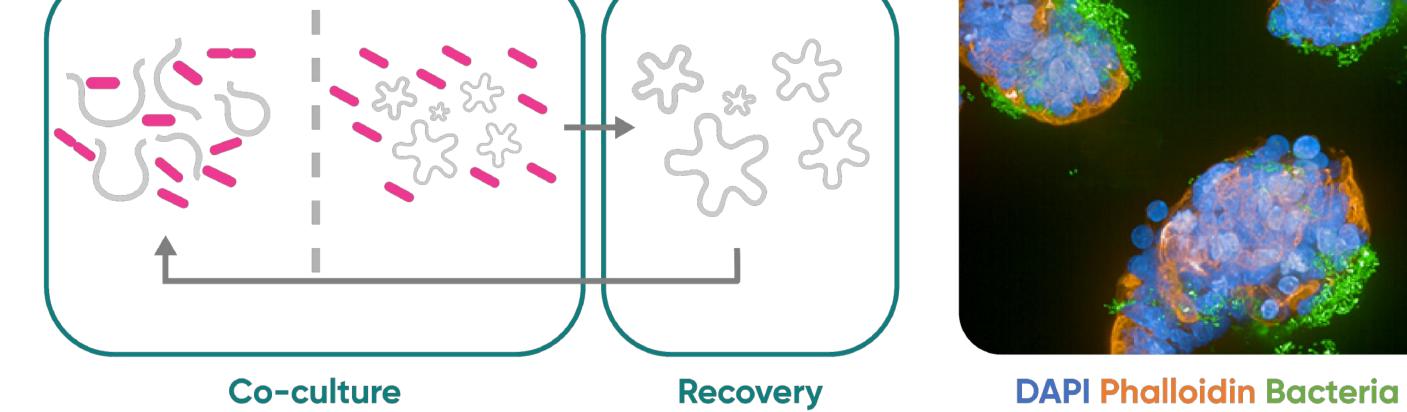


dependent on colibactin production

PDO fragments were incubated with either colibactin-producing bacteria or an isogenic mutant strain lacking the machinery for colibactin production. Colibactin-producing bacteria induced higher levels of DNA damage (yH2AX) compared to colibactin-deficient bacteria, as determined by IF.







Conclusions

We have developed a PDO and bacteria co-culture system compatible with mediumto-high throughput screening readouts (IF, flow cytometry and comet chip). This coculture system is currently tailored for the modeling of colibactin genotoxic effects in the gut epithelium but can be potentially extended as a discovery platform to identify targetable, novel, and complex interactions between host and pathogen.

References: ¹Pleguezuelos-Manzano *et al.* (2020)

© Copyright 2022. HUB Organoids (HUB). All rights reserved.



