



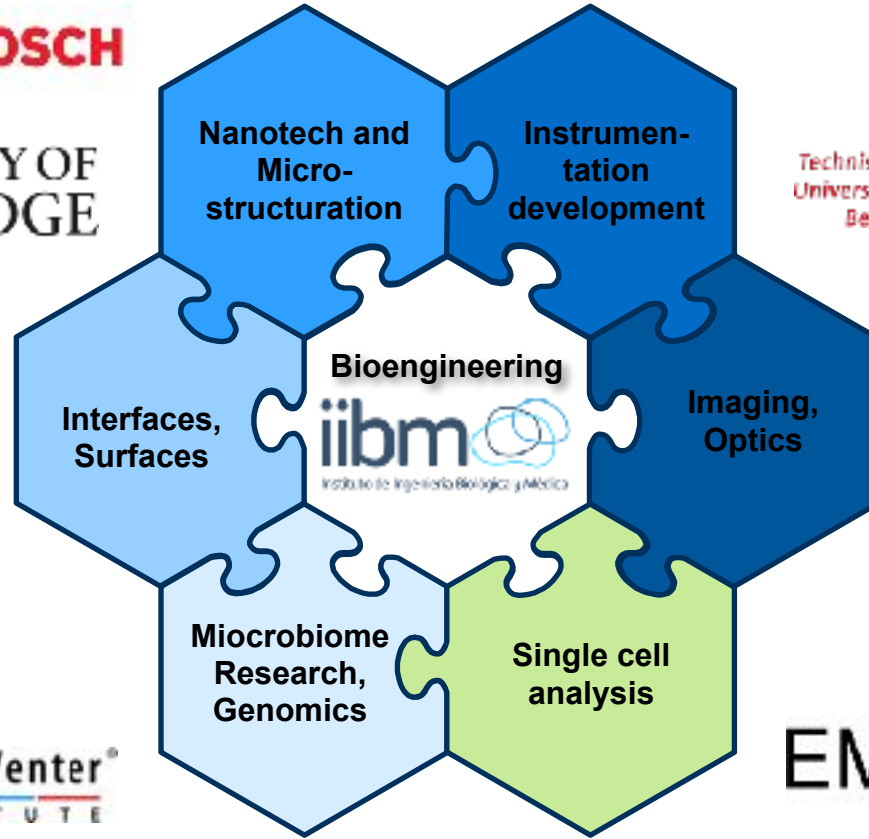
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Instituto de Ingeniería Biológica y Médica

Microfluidic method development for Single-cell microbiota analysis

Asst. Prof. Dr. Tobias Wenzel



The (human) microbiome as research target

- > 50% of cells in a human are bacteria (10^{13} cells in gut)
- 99% of genes in a human are bacterial genes
- 100+ disease associations with gut microbiome composition
- Opportunity to tackle **resistant pathogens** with **microbiome interactions**

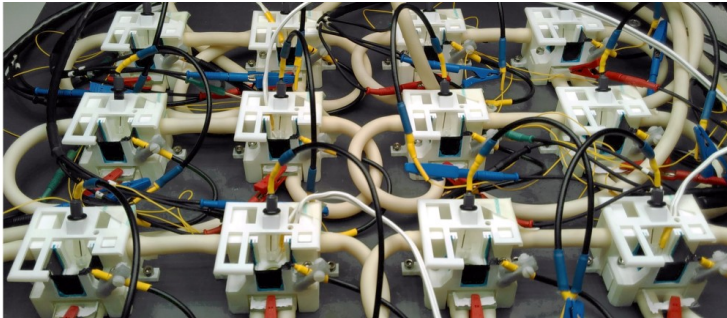


Image source: UC San Diego Health for Guardian article

Physics/Engineering perspective:

- Novel technological approaches to get data using **nano- and micro-fabrication**
- More novel data: ultra-high **throughput** experiments with **microfluidics**

The gut microbiome - complex, important

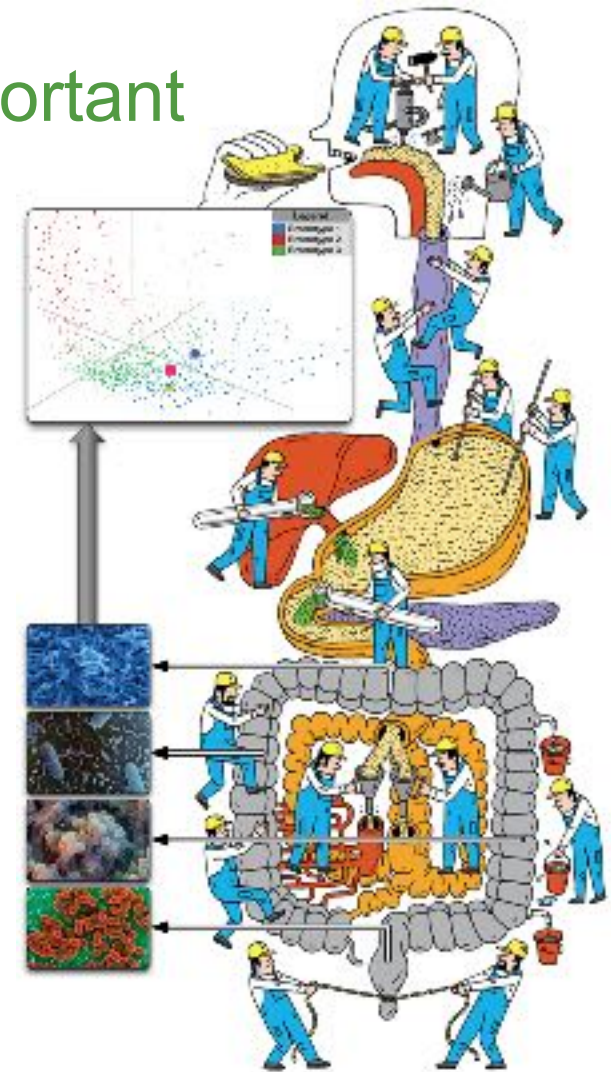
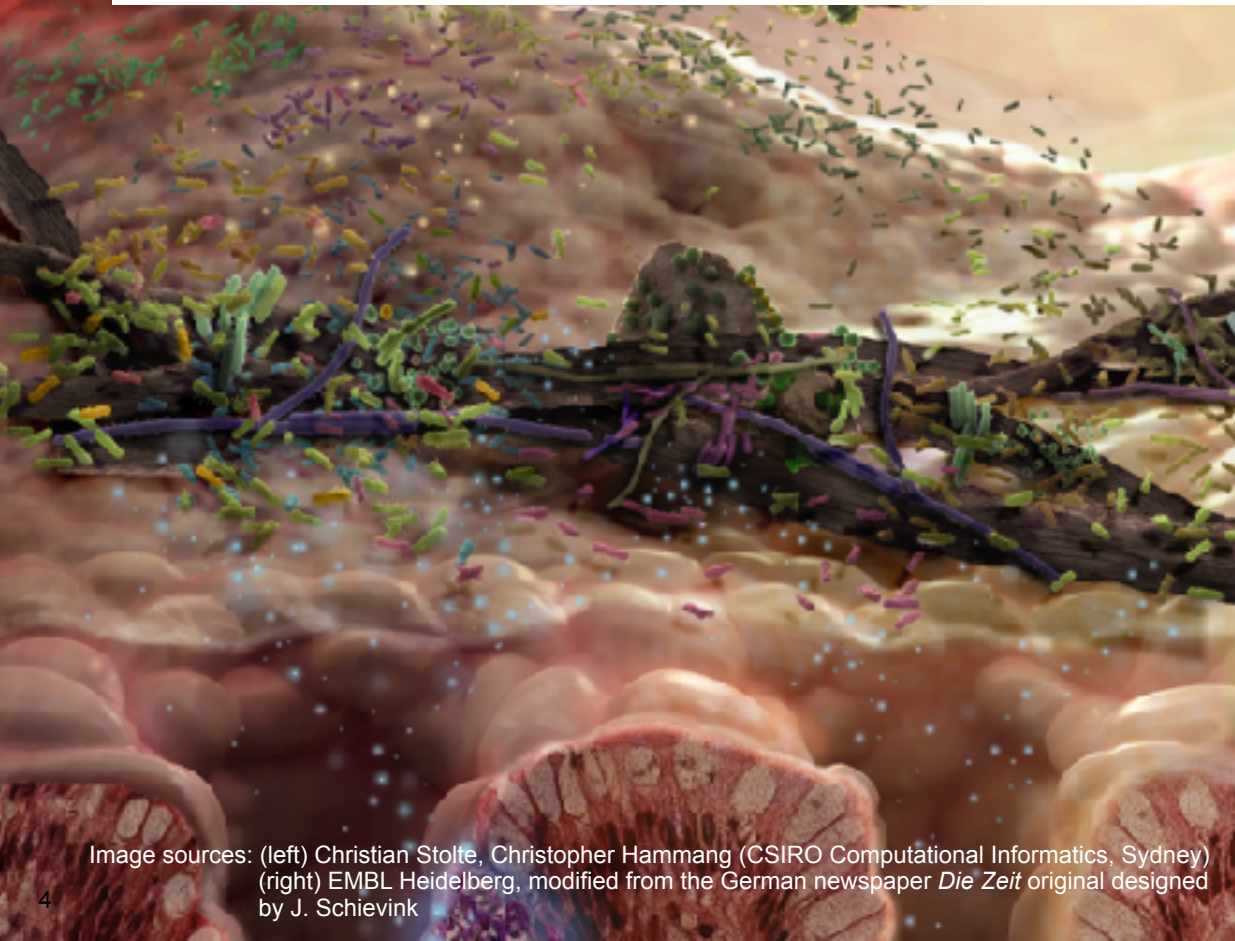
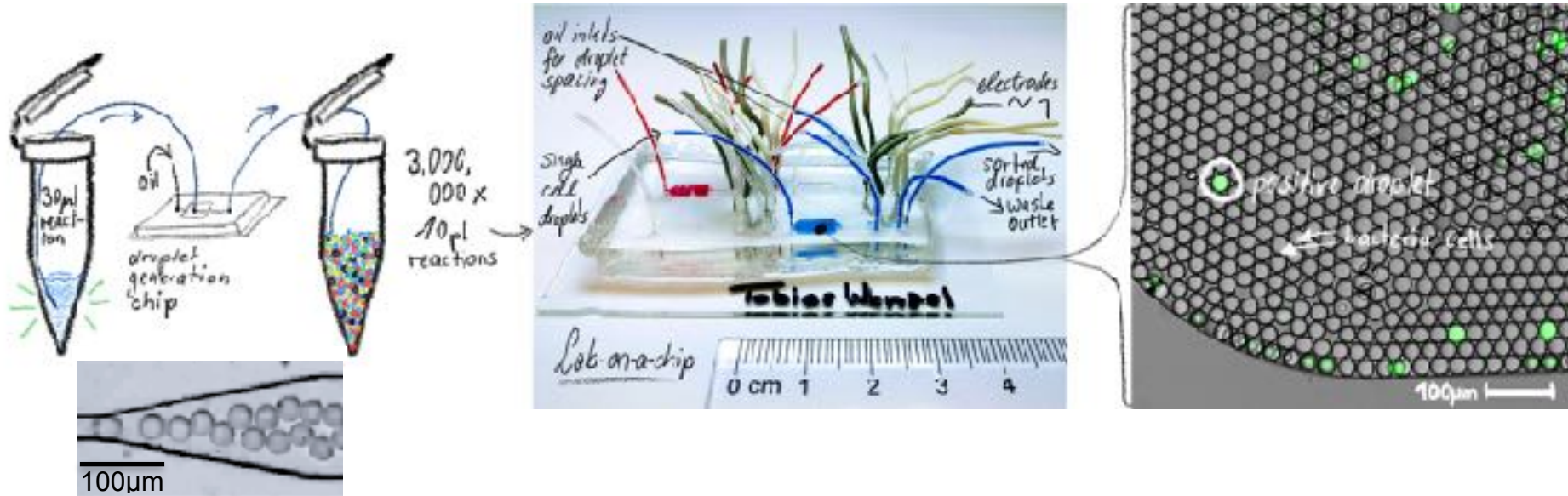


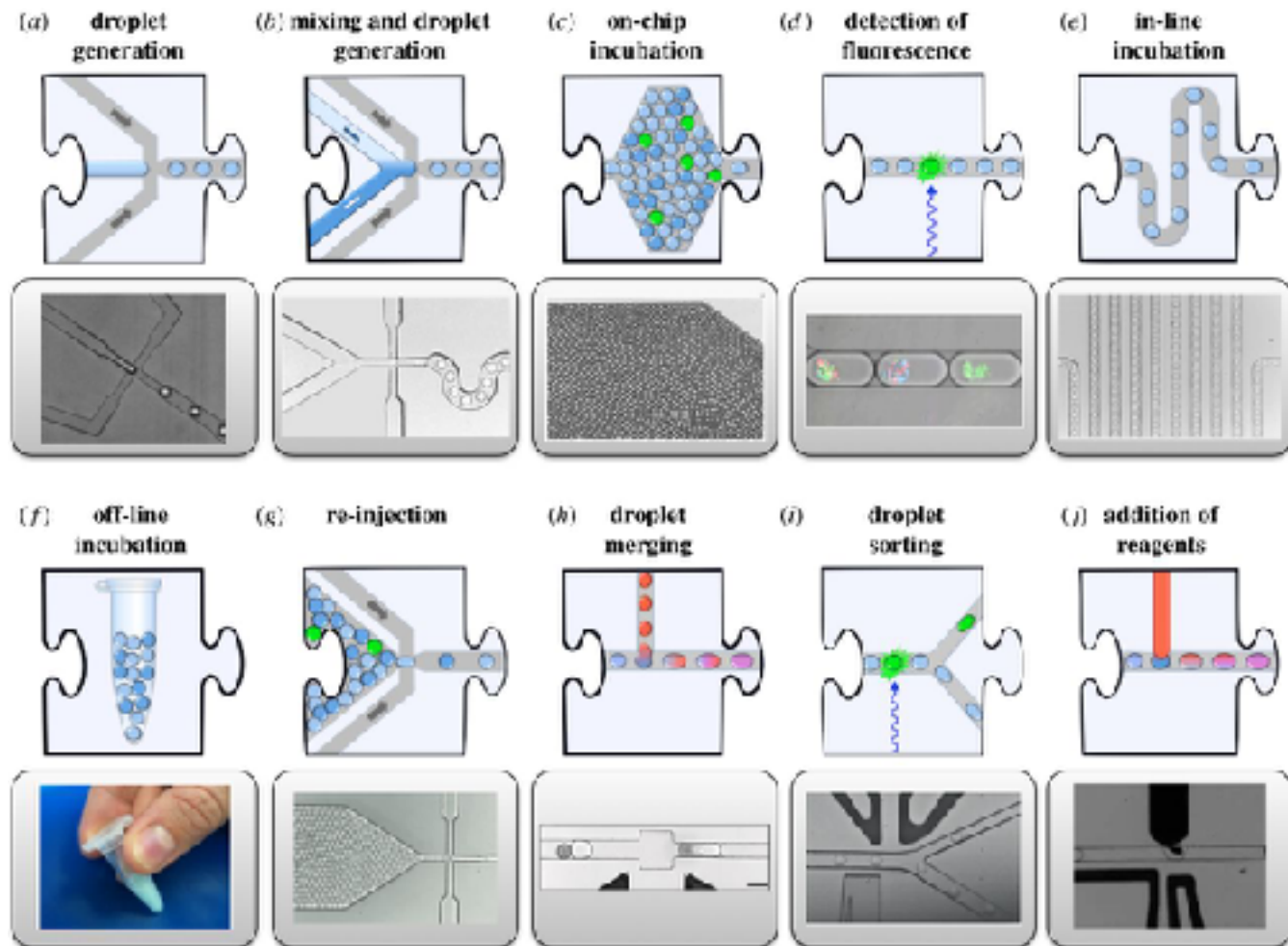
Image sources: (left) Christian Stolte, Christopher Hammang (CSIRO Computational Informatics, Sydney)
(right) EMBL Heidelberg, modified from the German newspaper *Die Zeit* original designed by J. Schievink

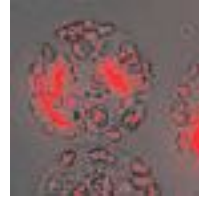
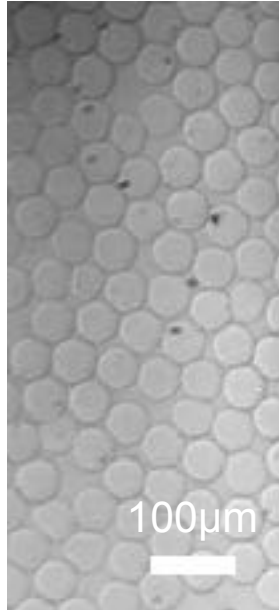
- One droplet = one experiment, e.g. with one cell each
- Experiments with millions of droplets per hour



Droplet Microfluidics

- Single cell control
- Interactions
- Ultra-high throughput
- Less contamination
- Versatile
- Reagent efficient

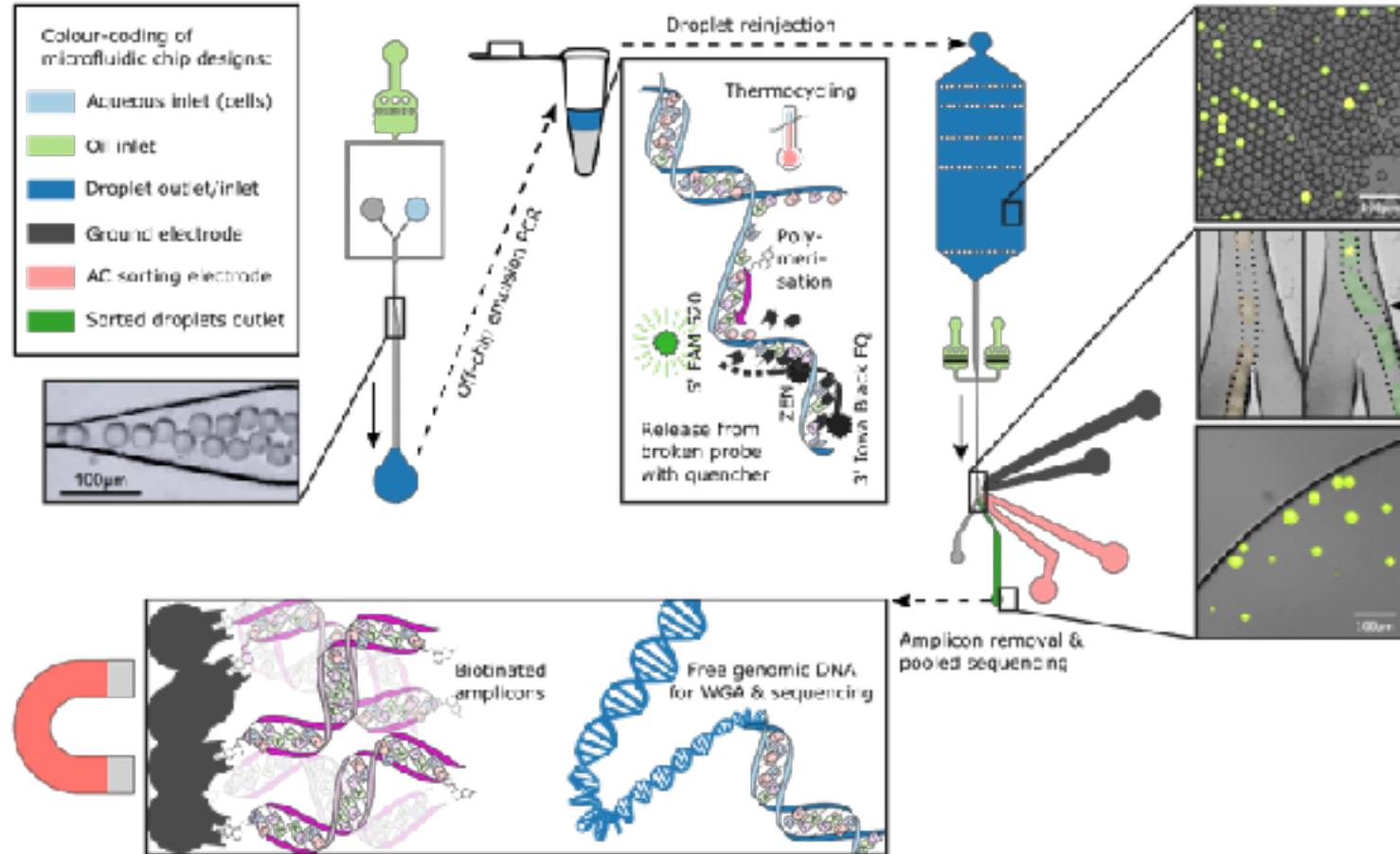




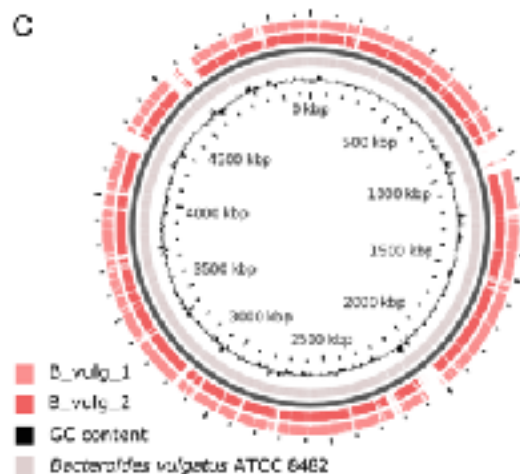
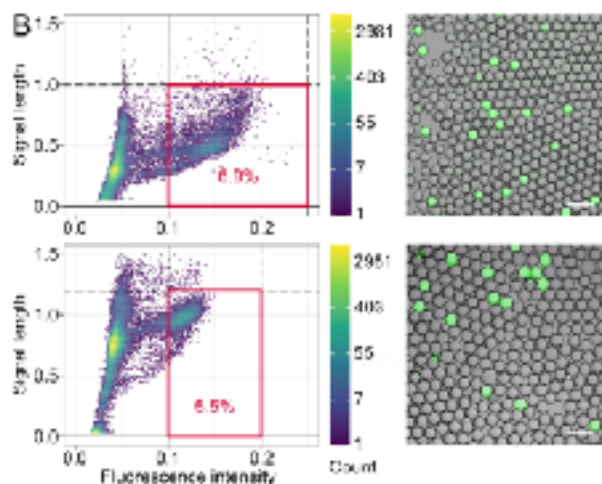
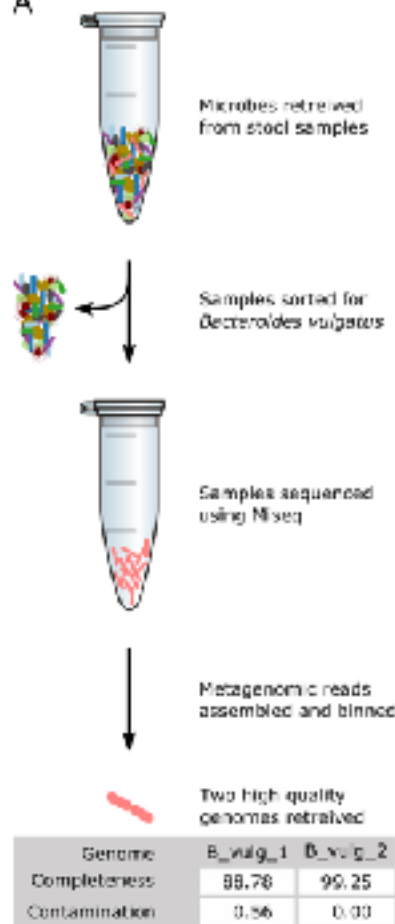
- Community functionality (resilience) - is more than individual strain properties
 - protection mechanisms, accumulation, intermediate metabolites, emergence
- Interactions are essential for cultivability
- Novelty: no systematic knowledge and use

Tackling antibiotic resistance with microbiota interactions

Precision Genomics: Enriching gut microbiome strains for cultivation-free genome sequencing using droplet microfluidics



A



ARTICLE IN PRESS

Enrichment of gut microbiome strains for cultivation-free genome sequencing using droplet microfluidics

Anna Physick ¹, Tobias Venzel ², Kiley West-Beltz ³, Falk Hildebrand ⁴, Eyal Kariel ⁵

Marco Raffaele Cosentino ⁶, Vladimir Biondo ⁷, Peter Bork ^{8,9}, Christoph A. Wenzel ^{1,2,3,4,5,6,7,8,9}, Sheng Cao ¹⁰, Shoufeng Chen ¹¹

Open Access | Published: December 28, 2021 | DOI: <https://doi.org/10.1016/j.crmeth.2021.100137>

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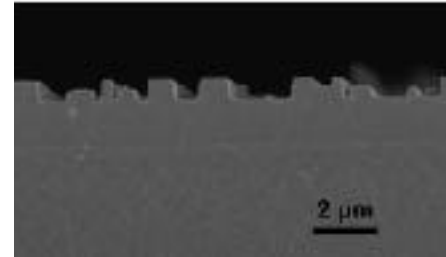
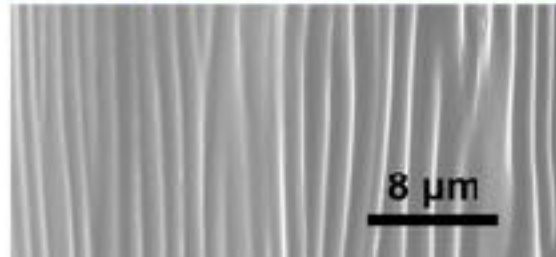
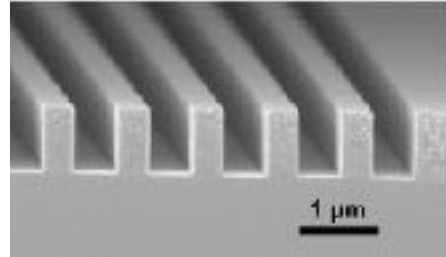
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Microfabrication and Bio-Nanotechnology

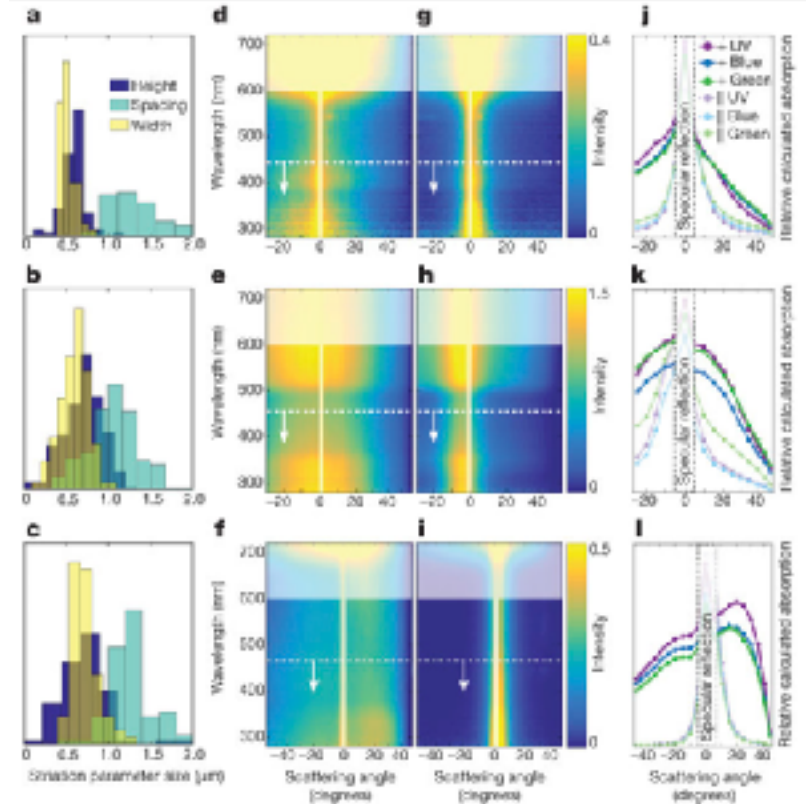
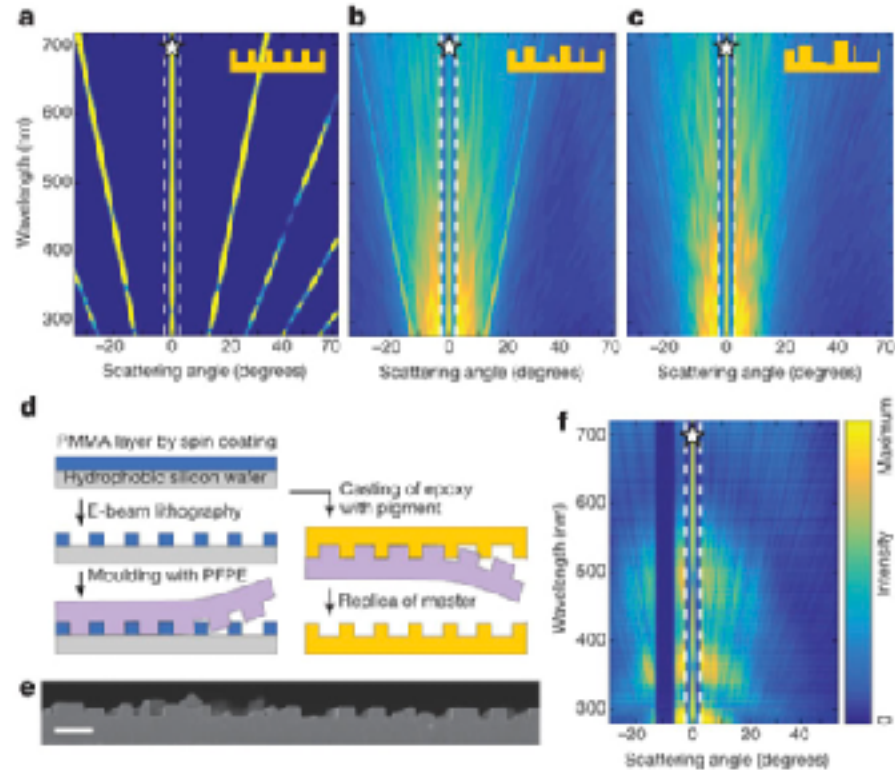
Understanding of natural "engineering principles" biomimicry



E. Moyroud* & T. Wenzel*,
et al.; *Nature* 550, 469-474
October 2017

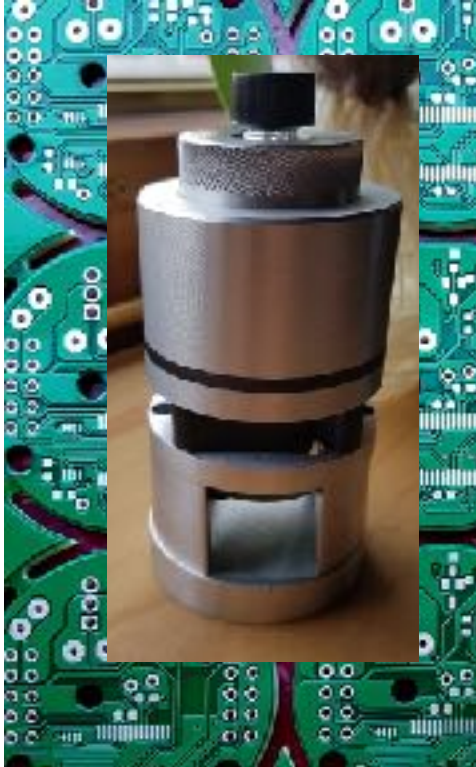


Coding for super-computer simulations

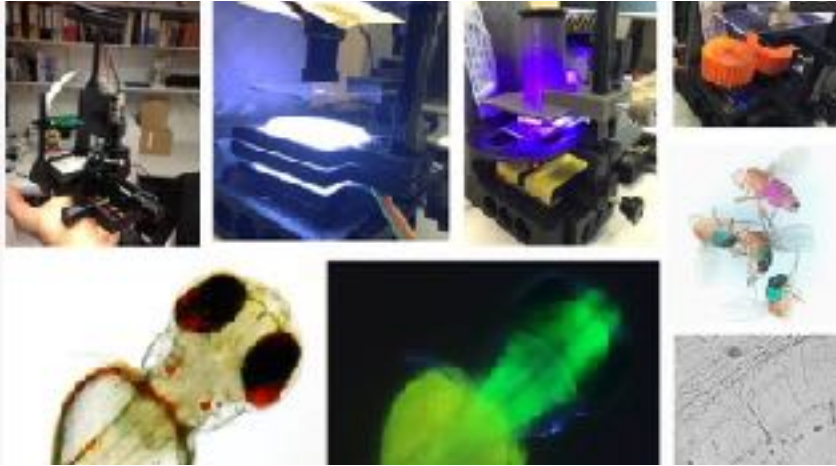


Upcoming: open source microfluidic workstation

3D printed fluorescence microscope

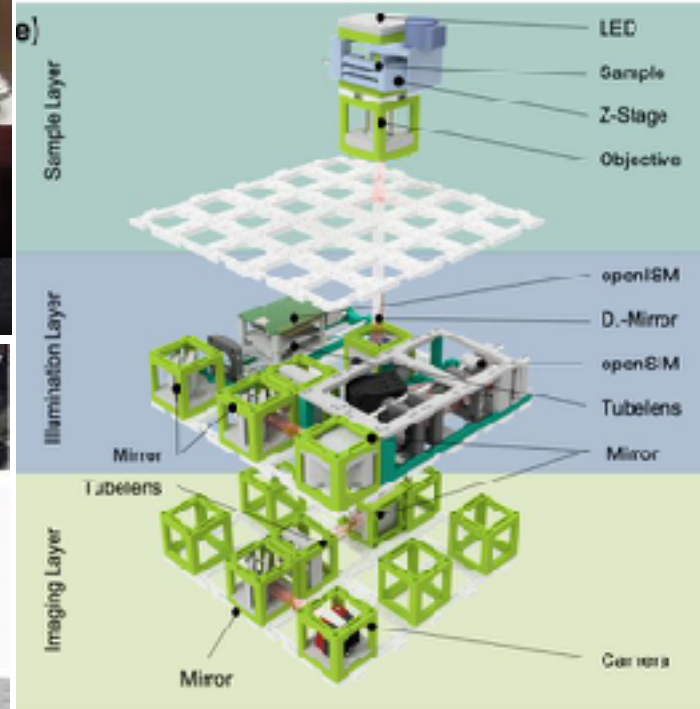


Low-cost high-end instrumentation access + automation = reproducibility



<- OpenFlexure

| UC2



<- FlyPi

“If I have seen further it is by standing on the
shoulders of Giants”

Sir Issac Newton

- Build creative tools that address new custom problems
- Reduction of cost
- Maintaining the equipment in-house
- Easy to automate and work with useful data formats
- Better collaborations - reproducibility, agile exchange



¡Gracias! Let's discuss...

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