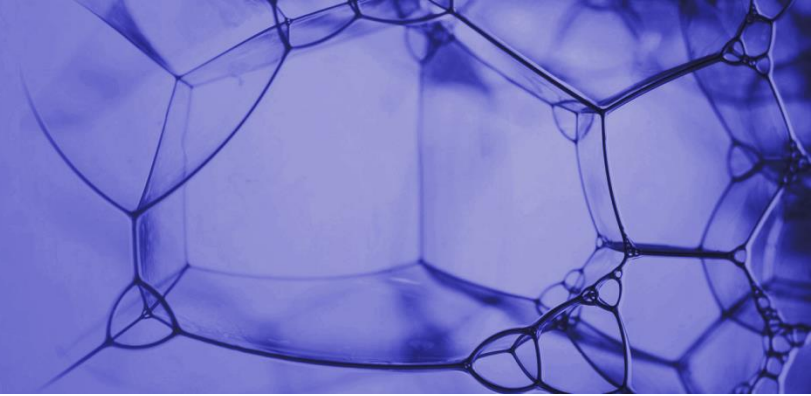


**LOSCHMIDT  
LABORATORIES**



## **Microfluidics – „Lab on a Chip“**

# Outline

- ❑ introduction to microfluidics
- ❑ physics of micro-scale
- ❑ lab on a chip applications
  - life and medical science
  - discovery of novel proteins
  - protein and metabolic engineering
- ❑ design and fabrication
- ❑ sensing and detection

# Lab on a Chip Concept

**incubation**



**pre-treatment**



**analysis**



**preparation**

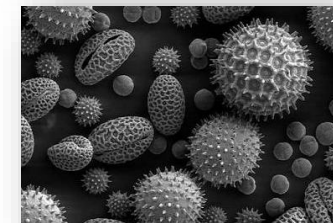
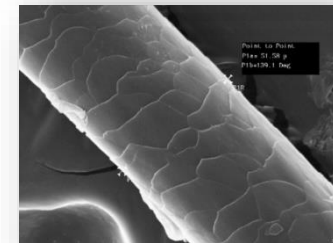
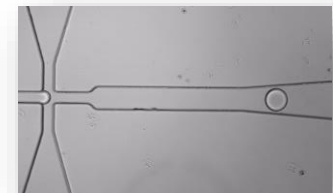


**collection**

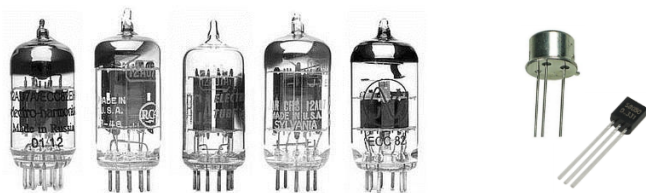


□ „behavior, control and manipulation of fluids geometrically constrained to a small dimensions“

- dimensions (1'-100'  $\mu\text{m}$ )
- volumes (nL, pL, fL)
- unrivalled precision of control
- (ultra)high analytical throughput
- reduced sample and power consumption
- facile process integration and automation

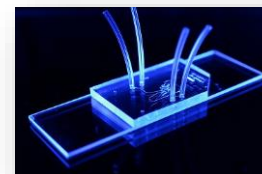
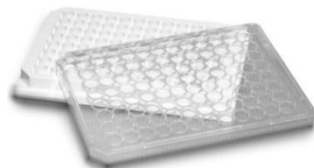


# Miniaturization & integration



	vacuum tubes	transistors	micro chips
size (mm)	100	10	0.000 01
price (USD)	10	1	0.000 000 1

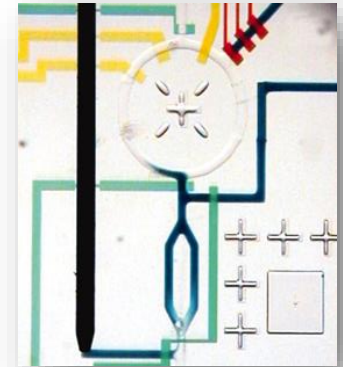
	test tube	microtiter plate	$\mu$ -fluidic chip
volume ( $\mu$ L)	1 000	10	0.000 001
throughput (assays/day)	10	1 000	1 000 000



# Concepts in microfluidics

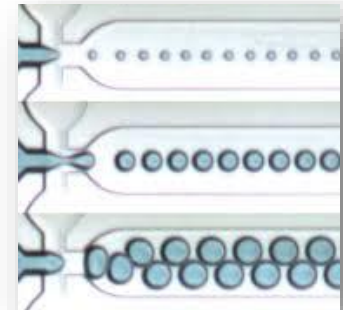
- ❑ **continuous-flow microfluidics**

manipulation of continuous liquid flow  
through micro-fabricated channels



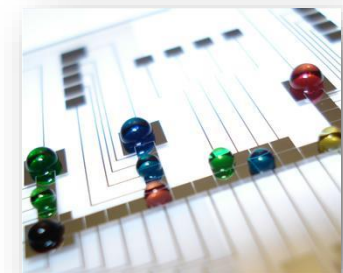
- ❑ **droplet-based microfluidics**

manipulating discrete volumes of fluids  
in immiscible phases



- ❑ **digital microfluidics**

droplets manipulated on a substrate  
using electro-wetting





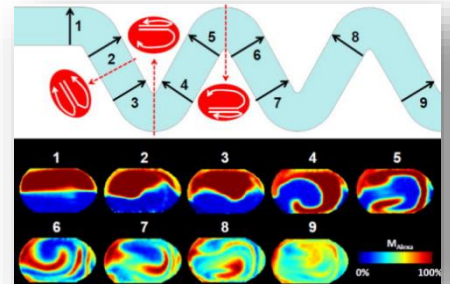
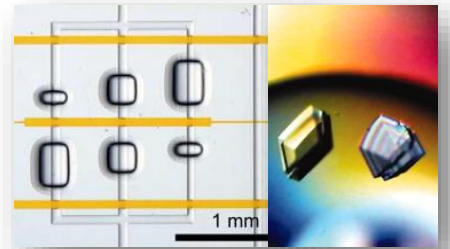
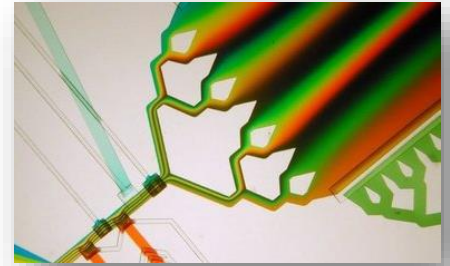
# Novel Physics of Micro-Scale

□ viscosity, surface tension and capillary forces dominate

▪ lack of turbulent phenomena

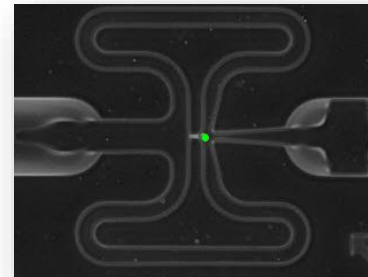
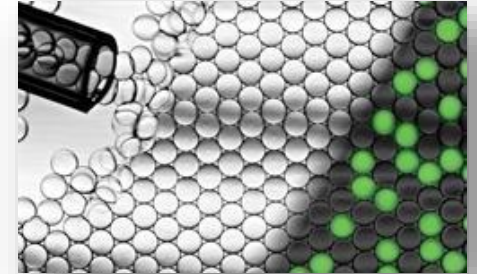
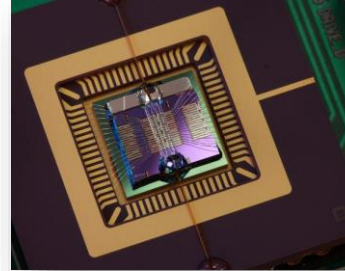
▪ absence of density-driven convection

▪ strong shearing forces



# Lab on a Chip applications

- ❑ analytics and chemistry
- ❑ PCR and sequencing
- ❑ point of care diagnostics
- ❑ pharmacology
- ❑ clinical studies
- ❑ single cell biology
- ❑ high throughput biology





# Polymerase chain reaction

## ❑ classical PCR

- slow heating/cooling cycles
- PCR tubes (strips), 96-well MTP
- volume 50 to 500  $\mu\text{L}$



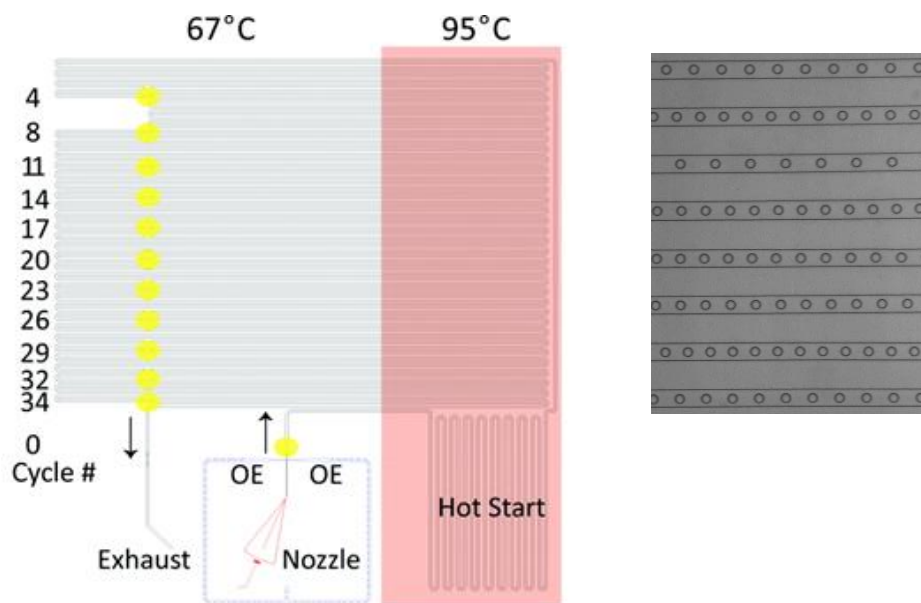
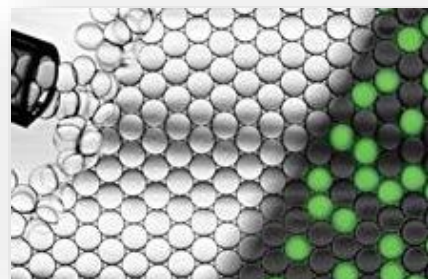
**Kary Mullis**

Nobel Prize in 1993

# Digital polymerase chain reaction

## □ digital PCR (single molecule)

- 1 nanoliter droplets
- 20 000 droplets per run
- fast heat transfer



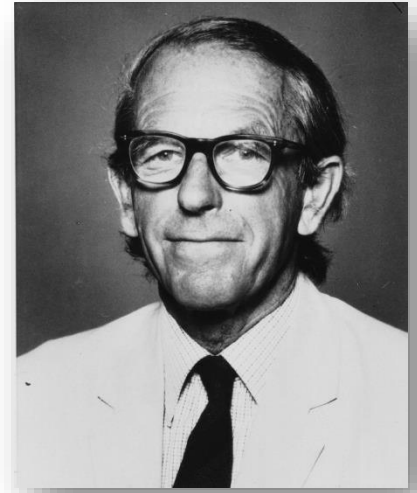
# Next-generation sequencing

- parallelization of single molecule pyrosequencing

- 454 Pyrosequencing (Roche)

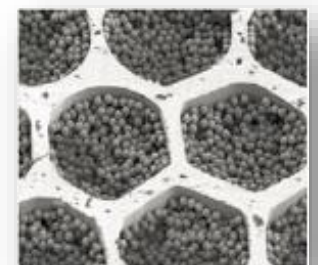
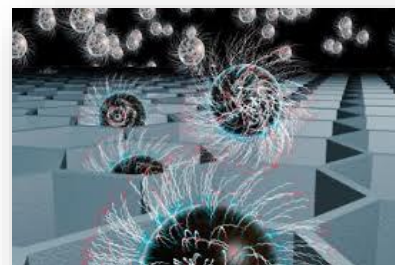
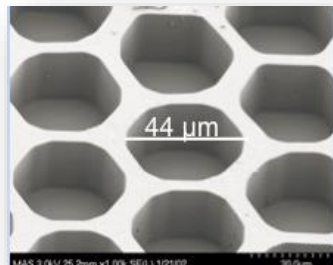
water in oil droplets 1 picoliter ( $10^{-12}$  liters)

1 mil. reads/run, 10 USD/Mbase

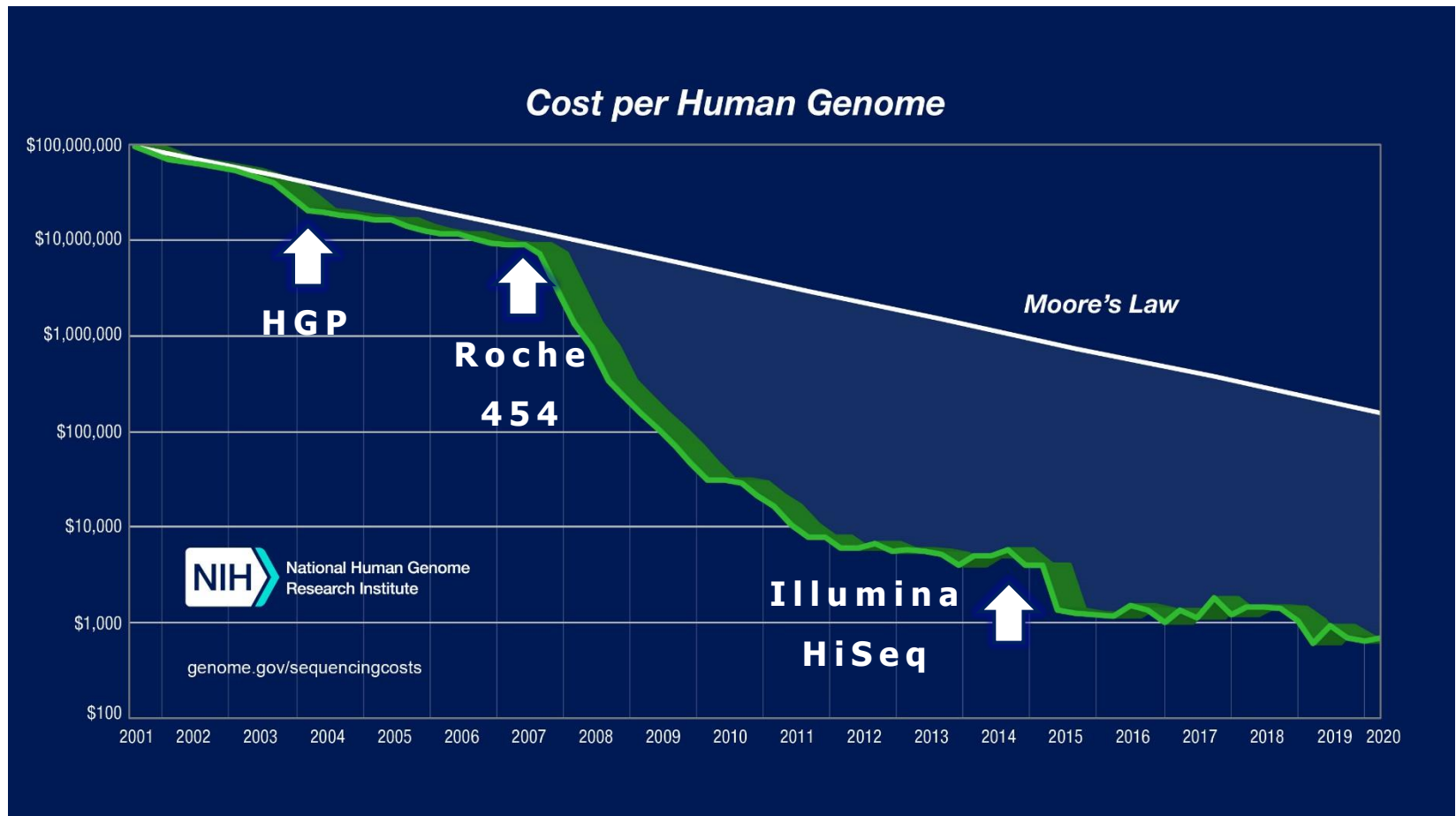


**Frederick Sanger**

Nobel Prize in 1980



# Revolution in DNA analysis

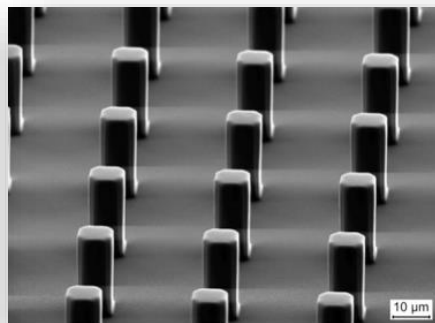
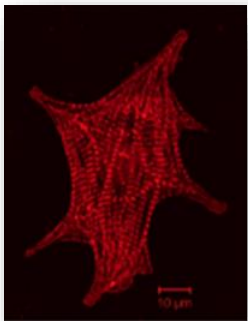


- ❑ 2003: 13 years, 3 billion USD
- ❑ 2018: days, < 1,000 USD

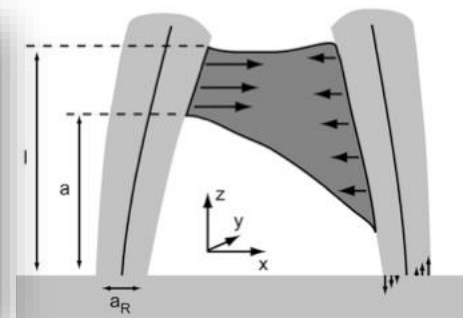
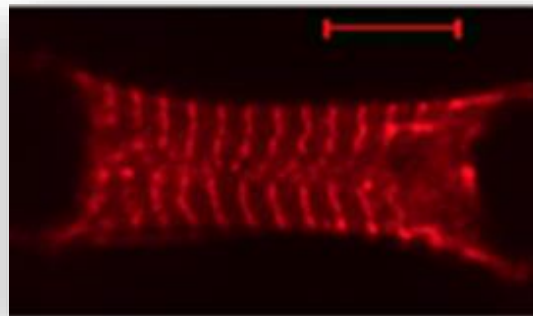
# Organ(oid)s on chip

- ❑ 3D chips mimicking human's physiological responses  
(e.g., pathological, pharmacokinetic, toxicological)
- ❑ realistic *in vitro* model closer to *in vivo* cell environment  
(e.g., mechanical strain, patterning, fluid shear stresses)

## flat surface



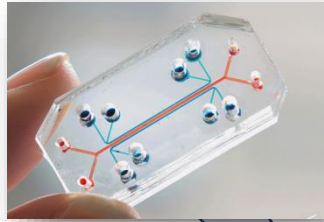
## micropillar



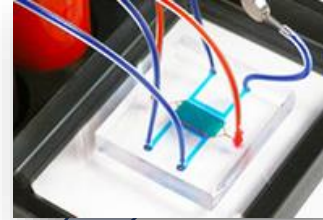


# Organs on chip

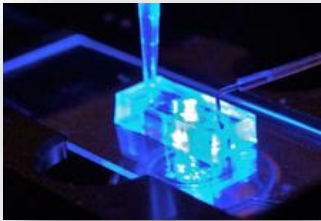
**Lung**



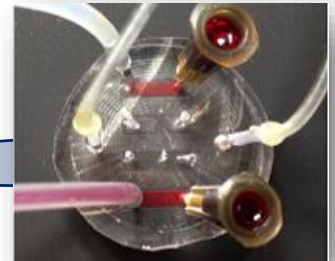
**Neurovascular**



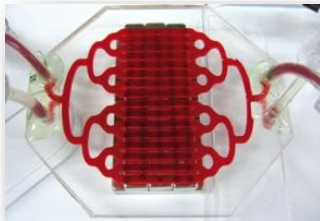
**Heart**



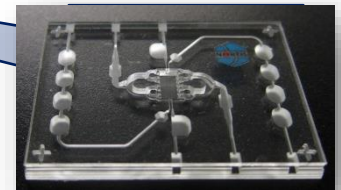
**Artery**



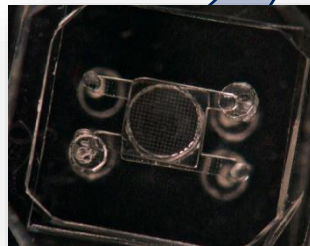
**Spleen**



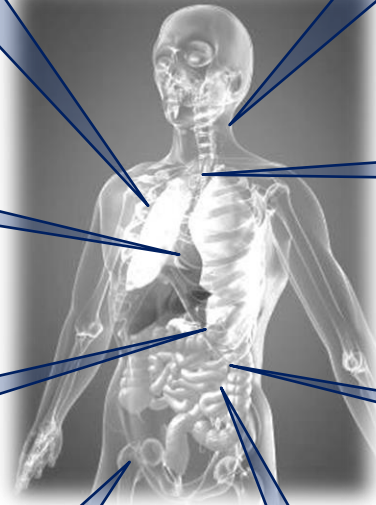
**Kidney**



**Bone**



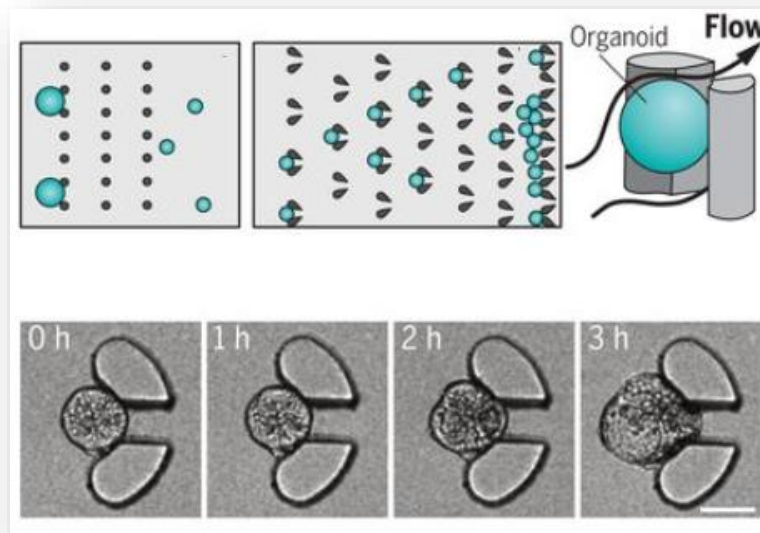
**Intestine**





# Organ(oid)s on chip

- ❑ **3D chips mimicking human's physiological responses**  
(e.g., pathological, pharmacokinetic, toxicological)
- ❑ **realistic *in vitro* model closer to *in vivo* cell environment**  
(e.g., mechanical strain, patterning, fluid shear stresses)



- ❑ **can replace expensive and controversial animal testing**

# Protein Discovery and Engineering

**ENZYME MINER** v1.0  
Automated mining of soluble enzymes with diverse structures, catalytic properties and stabilities

Submit new job | Help | Example | Acknowledgements

**ABOUT**

EnzymeMiner identifies putative members of enzyme families and facilitates the selection of promising targets for experiments. The server mines sequences that are likely to show the desired catalytic activity. Key selection criteria are: (i) predicted soluble expression in *Escherichia coli*, (ii) sequence identity, and (iii) deposit date. The search query can be a sequence from the Swiss-Prot database or a custom sequence with specified essential residues. The output is an interactive selection table and a sequence similarity network.

User guide | Example results Hide

**JOB INPUT**

Swiss-Prot sequences | Custom sequences

1.1.1.1 - Alcohol dehydrogenase. (240) Load example

Select sequences from table (max. 40) | Select sequences from similarity network (max. 40)

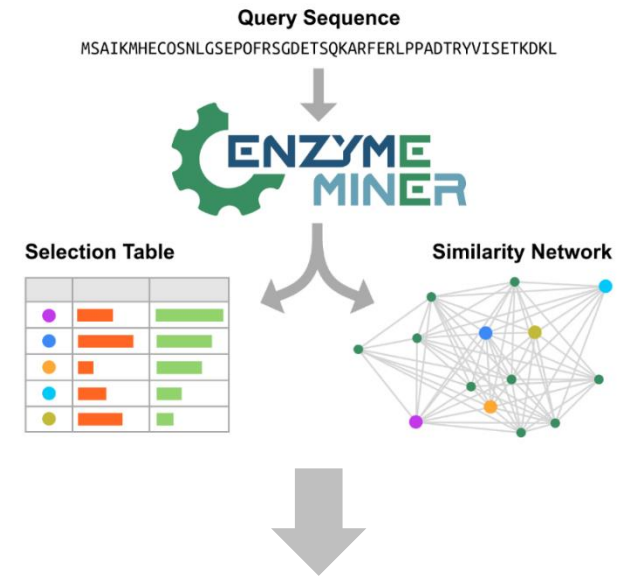
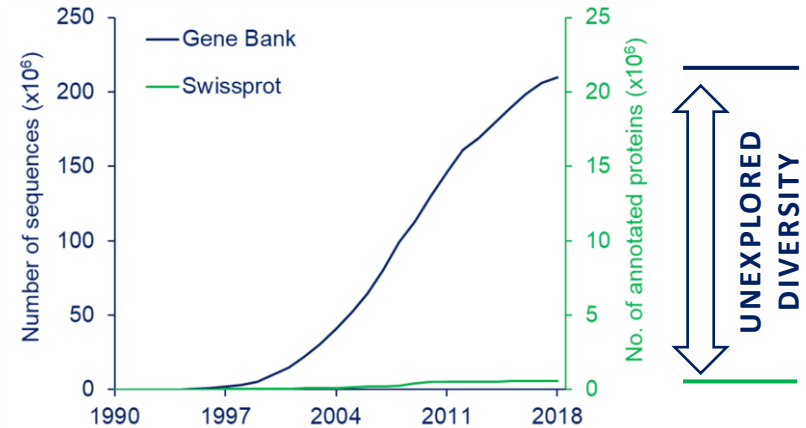
Accession	ER	Length	Sequence plot
<input type="checkbox"/> A0A075TMP0	9	340	
<input type="checkbox"/> A1A835	7	369	
<input type="checkbox"/> A1CFL1	7	388	
<input type="checkbox"/> A1L4Y2	14	394	
<input type="checkbox"/> A2XAZ3	14	381	
<input type="checkbox"/> A5IYX5	1	309	
<input type="checkbox"/> A6ZTT5	0	382	
<input type="checkbox"/> A7ZIA4	7	369	
<input type="checkbox"/> A7ZX04	7	369	
<input type="checkbox"/> B1J085	7	369	
<input type="checkbox"/> B1LIP1	7	369	
<input type="checkbox"/> B4M8Y0	4	254	
<input type="checkbox"/> E1ACQ9	8	339	
<input type="checkbox"/> F2Z678	10	351	
<input type="checkbox"/> F8DVL8	0	383	

Selected 0 of 240 (max. 40) | 1 to 15 of 240 | Page 1 of 16

Filter sequences by Pfam domains...

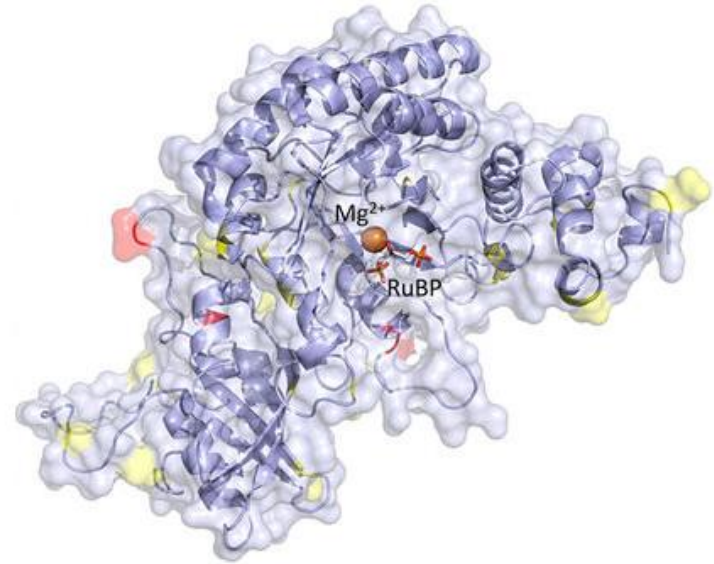
Select all | Deselect all | Show selected only

Advanced options



# Protein Discovery and Engineering

1-MDQSSRYVNLALKEEDLIAGGEHVLCAIYIMKPKAGYGYVATAAHFAAESS-50  
51-TGTNVEVCTTDDFTRGV<sup>D</sup>DALVYEVDEARE<sup>E</sup>LTKIAYPVALFDRN<sup>I</sup>TDGK<sup>A</sup>M-100  
101-IAS<sup>F</sup>LTLTMGNNQGMGDVEYAK<sup>M</sup>HDFYVPEAYRALFDG<sup>S</sup>VNISALWKVL-150  
151-GRPEVDGGLVVGT<sup>I</sup>IKPKLGLRPKPF<sup>A</sup>EACH<sup>H</sup>AFWLGGDFIKNDEPQGNQP-200  
201-FAPLRD<sup>T</sup>IALVADAMRRAQDETGEAKLFSANITADDPFEI IARGEYVLET-250  
251-FGENASHVALLVDGYVAGAA<sup>A</sup>AITTARRRFPDNFLHYHRAGHGAVTS<sup>F</sup>QSK-300  
301-RGYTAFVHCK<sup>M</sup>ARL<sup>Q</sup>GASGIHTGTMGFGKMEGES<sup>S</sup>DRAIAYMLT<sup>Q</sup>DEAQG-350  
351-PFYRQSWGGMK<sup>A</sup>CTPIISGGMNALR<sup>M</sup>PC<sup>G</sup>FFENLGNANVILTAGGGAFGHI-400  
401-DGPVAGARSLRQAWQAWRDGVP<sup>V</sup>L<sup>D</sup>YAREHKELARAFESFPGDAD<sup>Q</sup>I<sup>I</sup>YPG-450  
451-WRKALGV<sup>E</sup>DT<sup>R</sup>SALPA-466



No.	Coverage (95%)
1	94
2	3 066
3	98 163
4	3 141 251
5	100 520 093

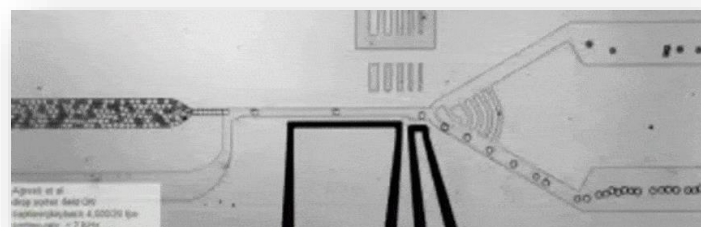
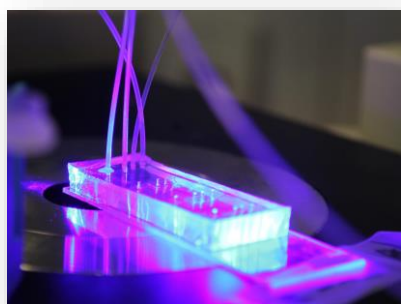
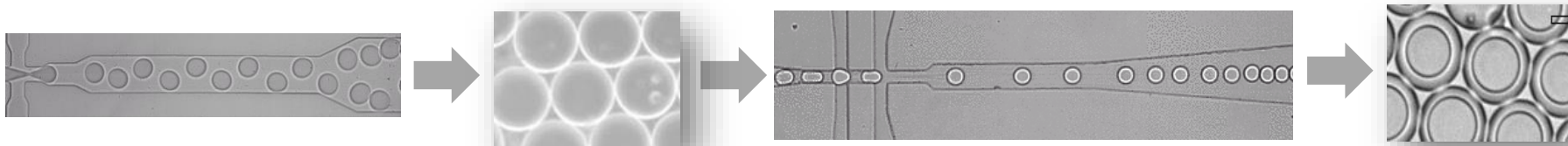


SCREENING AND IDENTIFICATION  
OF POSITIVE HITS

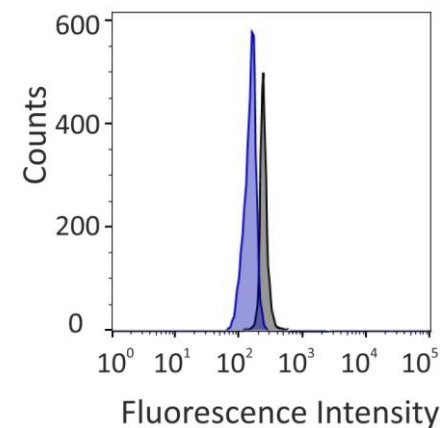


FUNCTIONAL CHARACTERISATION

# (Ultra)High-throughput screening

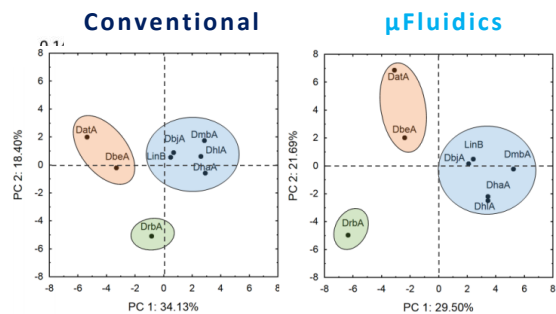
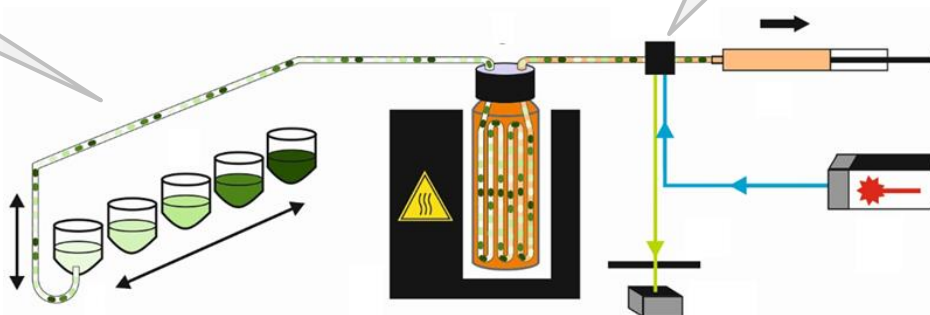
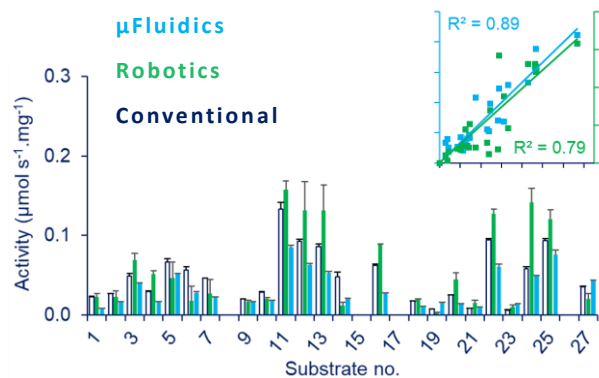
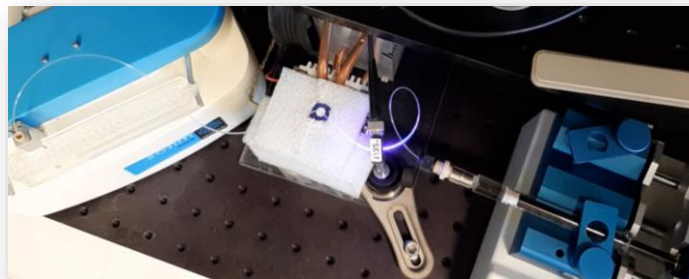
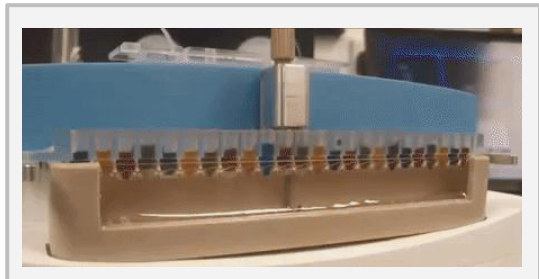


	Robotic	$\mu$ Fluidic
Reaction volume	100 $\mu$ L	5 pL
Reactions / day	50 000	$1 \cdot 10^8$
Total time	5 years	3 days
Total volume	5 000 L	150 mL
No. of plates / devices	250 000	2.0
No. of tips	28 000 000	10

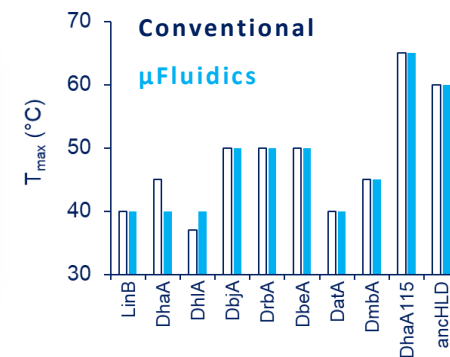




# Activity and specificity in $\mu$ -droplets

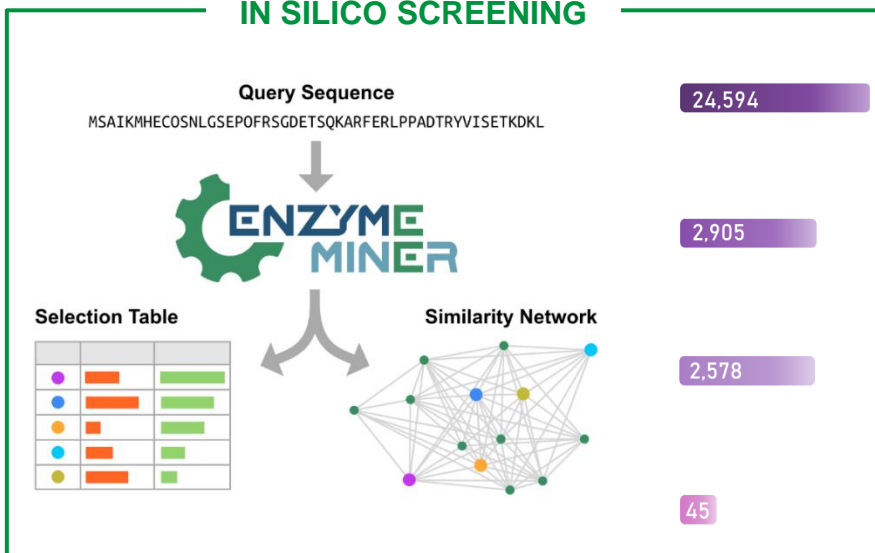


	Test tube	Robotic	$\mu$ Fluidic
Reaction volume (mL)	10	1	0.000 15
Total enzyme (mg)	540	54	0.5
Total time (days)	100	30	5

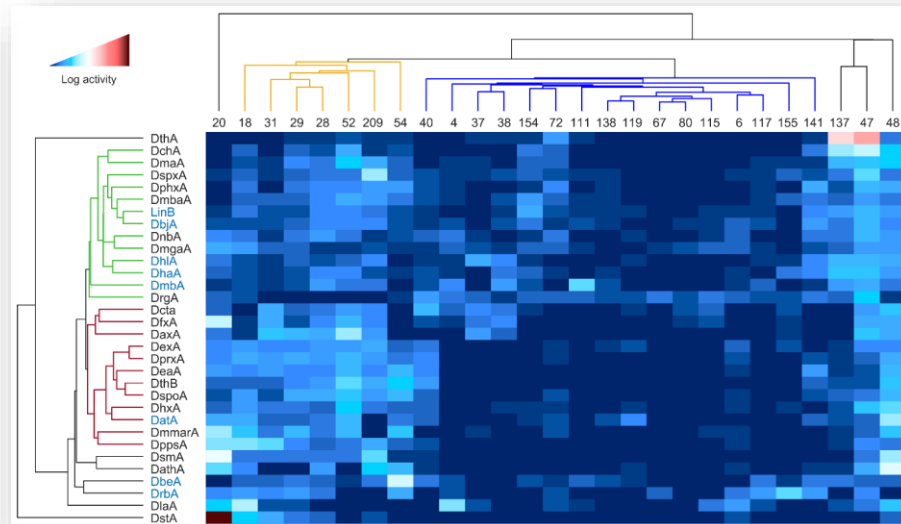
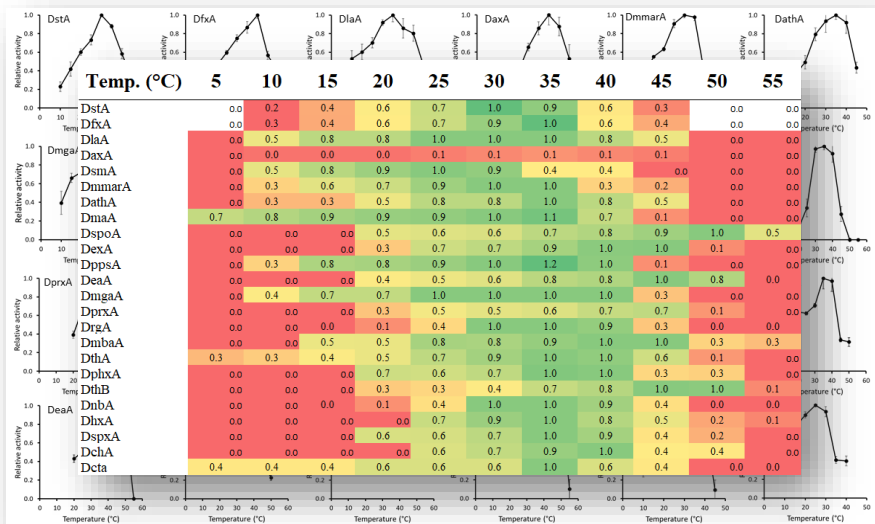
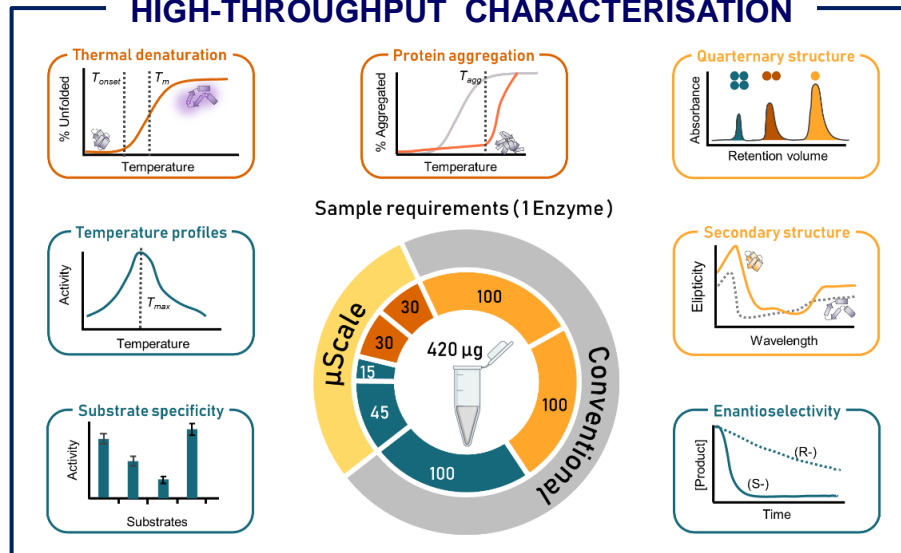


# Activity and specificity in $\mu$ -droplets

## IN SILICO SCREENING

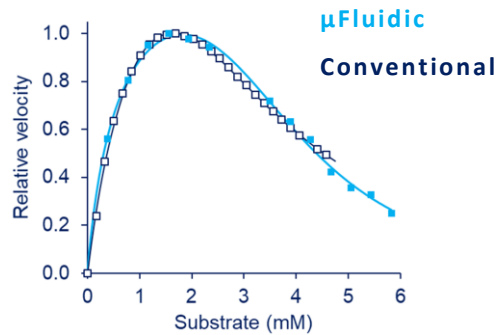
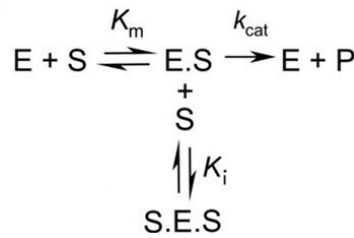
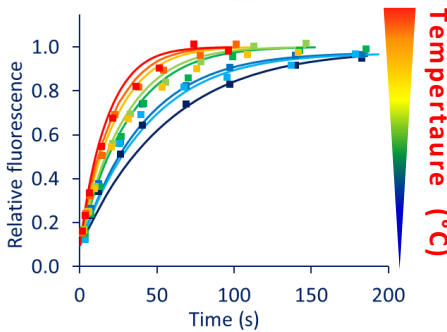
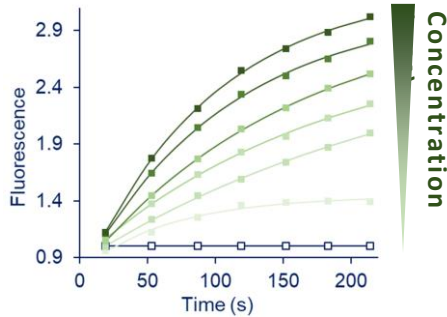
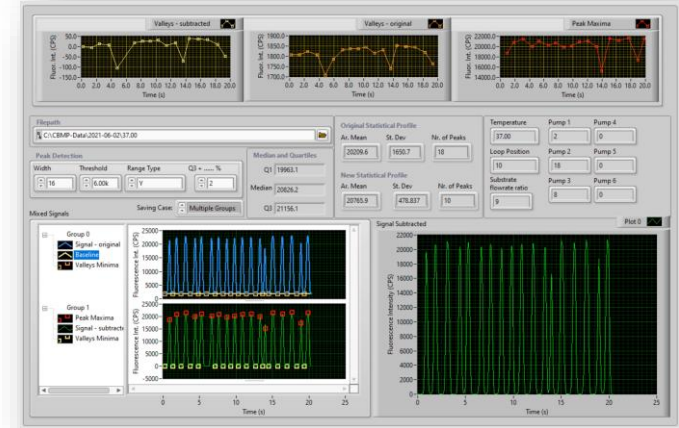
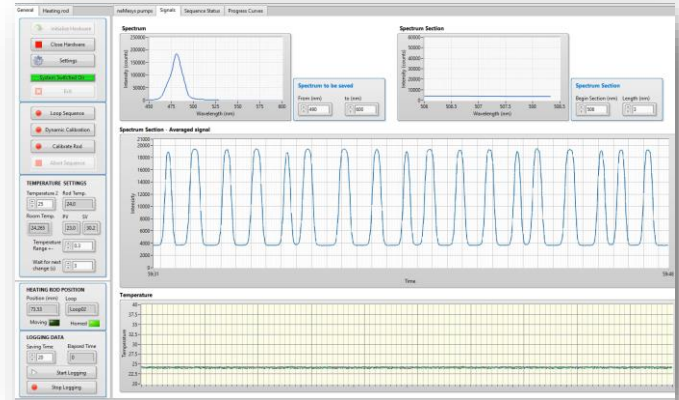
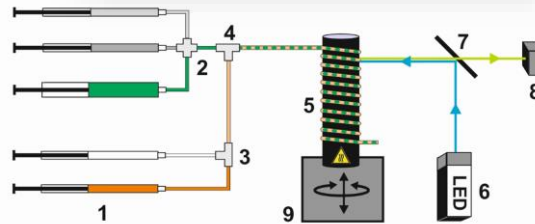
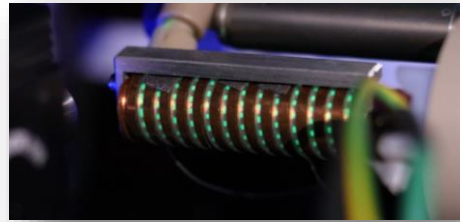
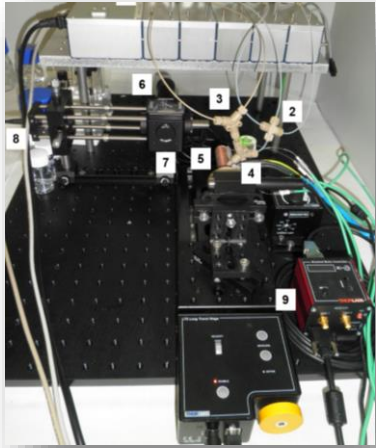


## HIGH-THROUGHPUT CHARACTERISATION



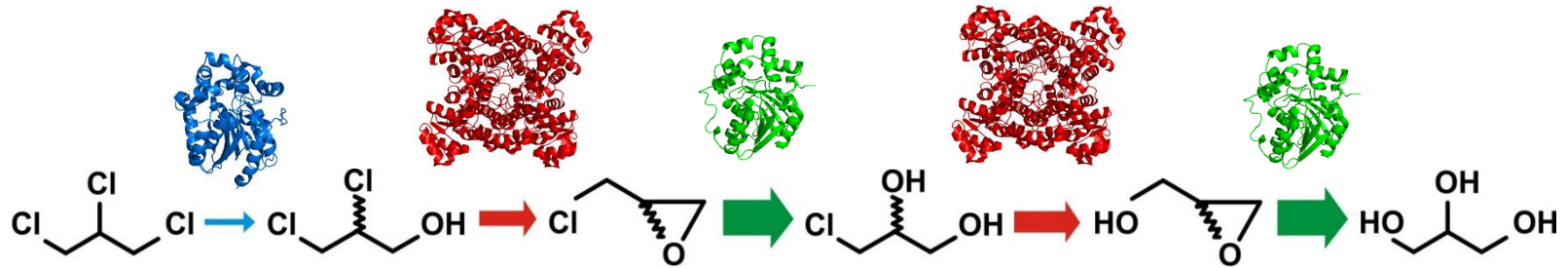


# Steady-state kinetics in $\mu$ -droplets

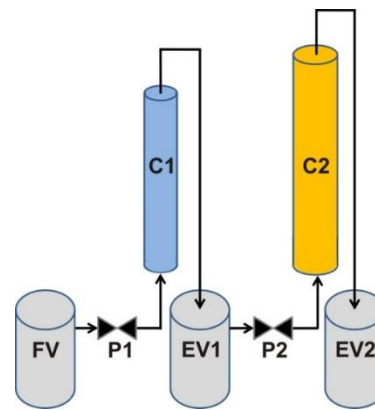
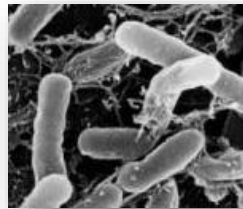
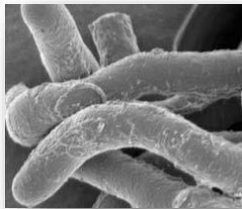


	Test tube	$\mu$ Fluidic
Reaction volume (mL)	10	0.000 1
Total enzyme (mg)	1	0.01
Throughput per hour	50	10 000

# Synthetic biology



*Rhodococcus*      *Agrobacterium*



$$\frac{dc_{TCP}}{dt} = \frac{k_{cat,TCP,(R)-DCP} \times c_{DhaA} \times c_{TCP}}{c_{TCP} + K_{m,TCP}} - \frac{k_{cat,TCP,(S)-DCP} \times c_{DhaA} \times c_{TCP}}{c_{TCP} + K_{m,TCP}}$$

$$\frac{dc_{(R)-DCP}}{dt} = \frac{k_{cat,TCP,(R)-DCP} \times c_{DhaA} \times c_{TCP}}{c_{TCP} + K_{m,TCP}} - \frac{k_{cat,(R)-DCP} \times c_{HheC} \times c_{(R)-DCP}}{c_{(R)-DCP} + K_{m,(R)-DCP}}$$

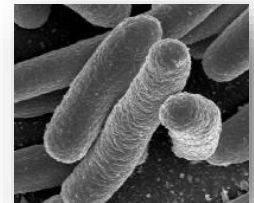
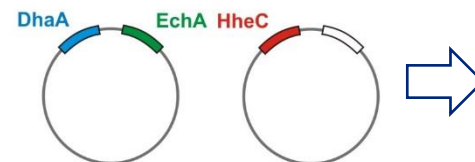
$$\frac{dc_{(S)-DCP}}{dt} = \frac{k_{cat,TCP,(S)-DCP} \times c_{DhaA} \times c_{TCP}}{c_{TCP} + K_{m,TCP}} - \frac{k_{cat,(S)-DCP} \times c_{HheC} \times c_{(S)-DCP}}{c_{(S)-DCP} + K_{m,(S)-DCP}}$$

$$\frac{dc_{ECH}}{dt} = \frac{k_{cat,(R)-DCP} \times c_{HheC} \times c_{(R)-DCP}}{c_{(R)-DCP} + K_{m,(R)-DCP}} + \frac{k_{cat,(S)-DCP} \times c_{HheC} \times c_{(S)-DCP}}{c_{(S)-DCP} + K_{m,(S)-DCP}} - \frac{k_{cat,ECH} \times c_{EchA} \times c_{ECH}}{c_{ECH} + K_{m,ECH}}$$

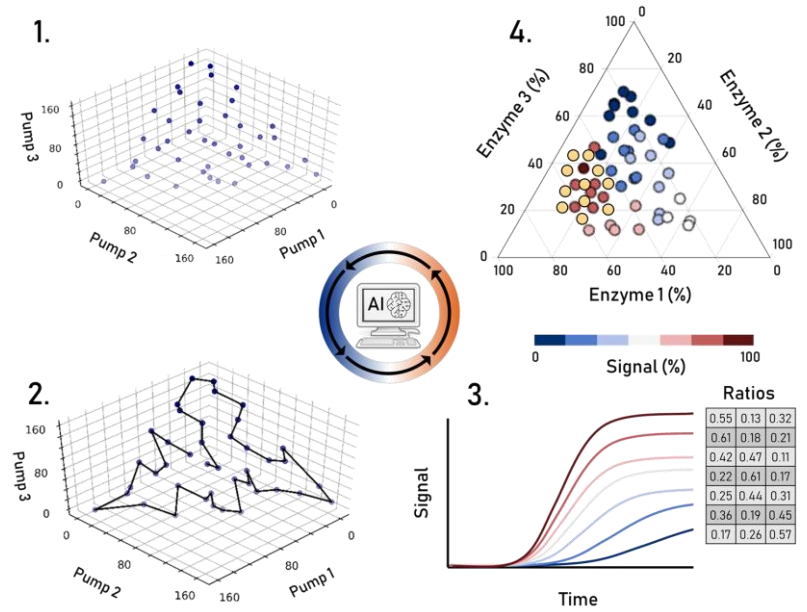
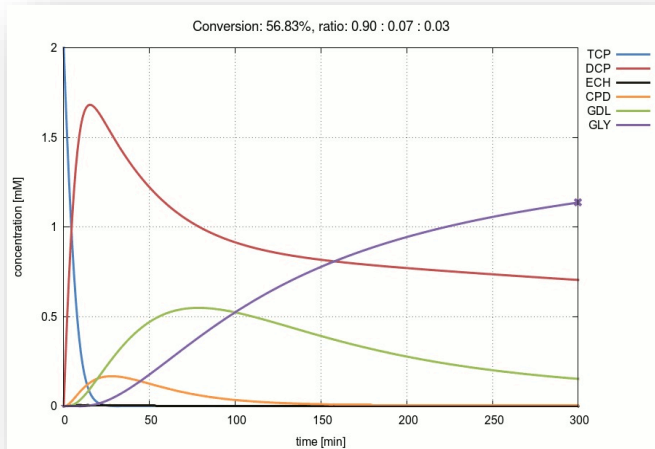
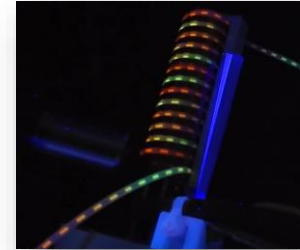
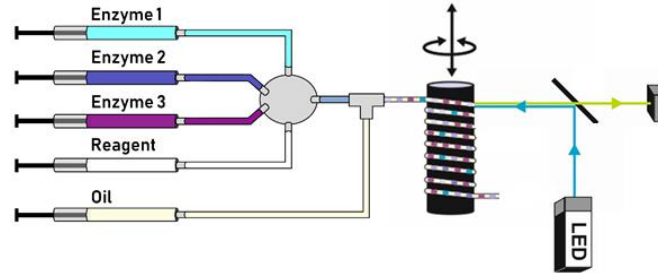
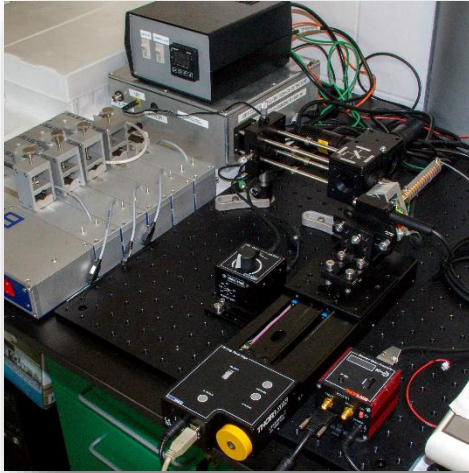
$$\frac{dc_{CPD}}{dt} = \frac{k_{cat,ECH} \times c_{EchA} \times c_{ECH}}{c_{ECH} + K_{m,ECH}} - \frac{k_{cat,CPD} \times c_{HheC} \times c_{CPD}}{c_{CPD} + K_{m,CPD}}$$

$$\frac{dc_{GDL}}{dt} = \frac{k_{cat,CPD} \times c_{HheC} \times c_{CPD}}{c_{CPD} + K_{m,CPD}} - \frac{k_{cat,GDL} \times c_{EchA} \times c_{GDL}}{c_{GDL} + K_{m,GDL} \times \left(1 + \frac{c_{GLY}}{K_i} + \frac{c_{TCP}}{K_c}\right)}$$

$$\frac{dc_{GLY}}{dt} = \frac{k_{cat,GDL} \times c_{EchA} \times c_{GDL}}{c_{GDL} + K_{m,GDL} \times \left(1 + \frac{c_{GLY}}{K_i} + \frac{c_{TCP}}{K_c}\right)}$$



# Enzyme cascades in $\mu$ -droplets



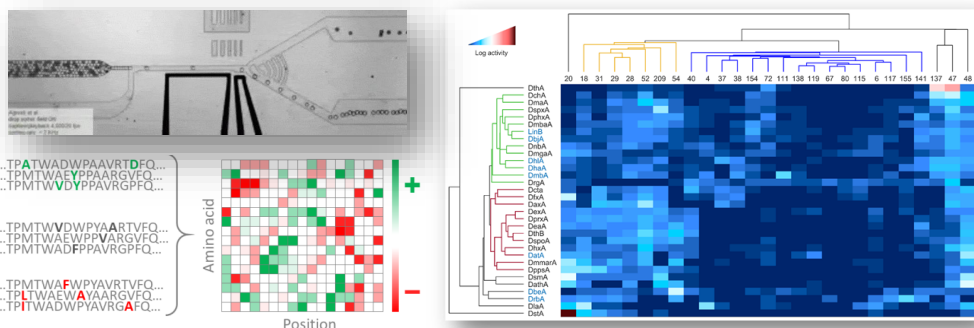


# AI in protein engineering

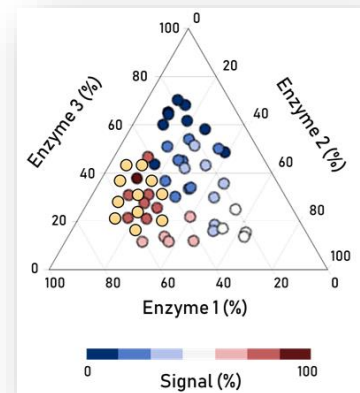
## Machine Learning in Enzyme Engineering

Stanislav Mazurenko,<sup>\*,†,⊕</sup> Zbynek Prokop,<sup>†,‡,⊕</sup> and Jiri Damborsky<sup>†,‡,⊕</sup>

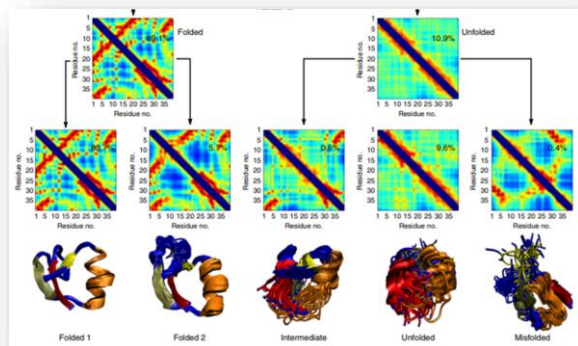
### SEQUENCE BASED PREDICTION supervised learning



### ITERATIVE OPTIMIZATION Bayesian optimization



### MOLECULAR DYNAMICS unsupervised learning

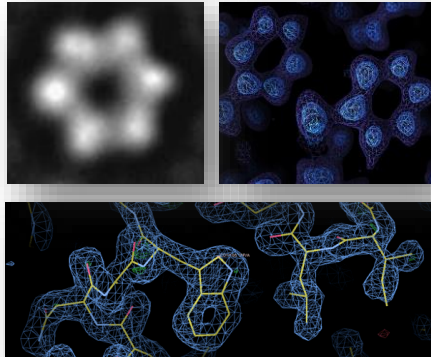


### STRUCTURE PREDICTION deep learning

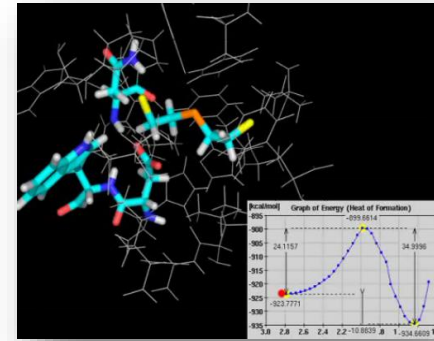
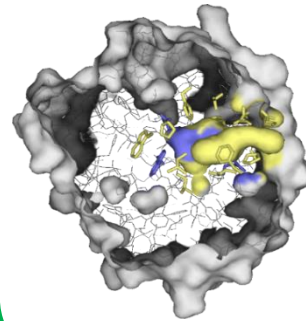


# Multidisciplinary science

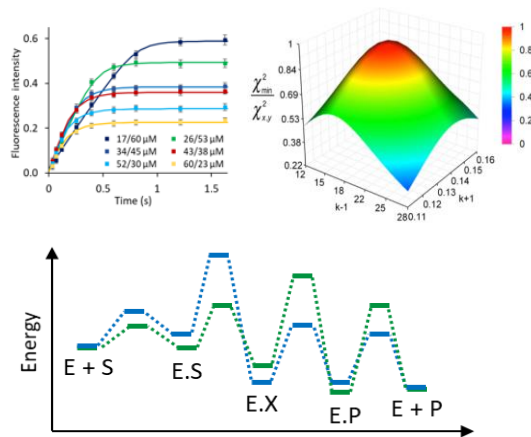
## STRUCTURAL BIOLOGY



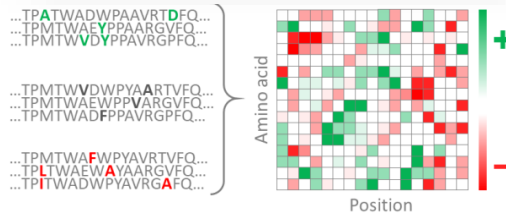
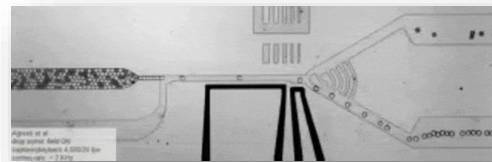
## COMPUTATIONAL DESIGN



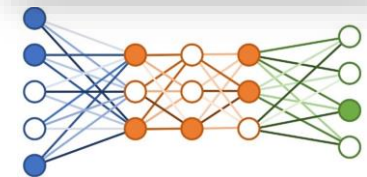
## ENZYME KINETICS



## MICROFLUIDICS

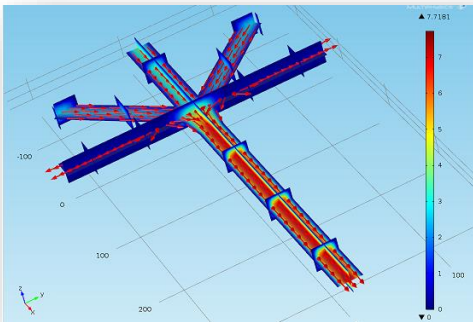
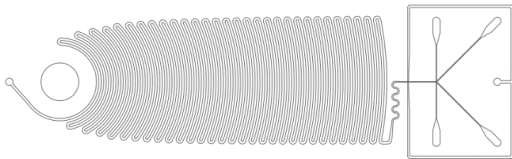


## ARTIFICIAL INTELLIGENCE



- **soft lithography** originates from semiconductor industry

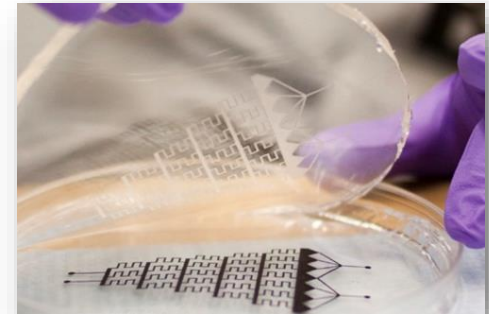
## DESIGN / MODELING



## MASK / MOLD



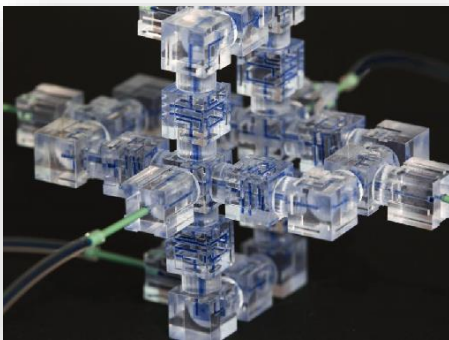
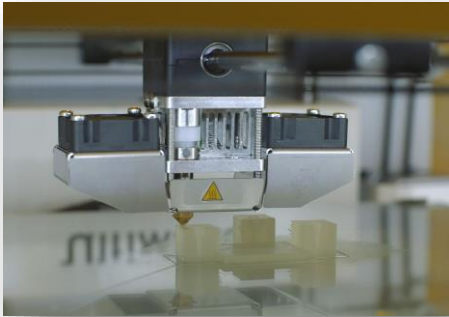
## CASTING / BONDING



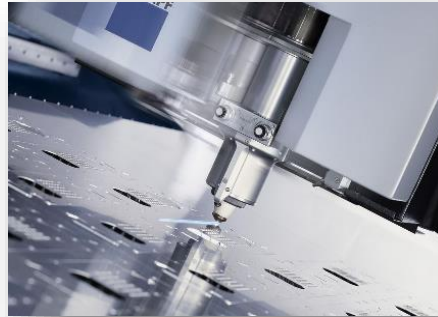


## □ direct fabrication methods

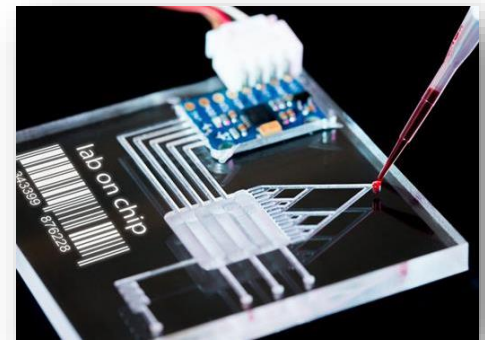
### 3D PRINTING



### LASER CUTTING



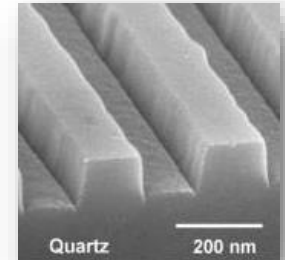
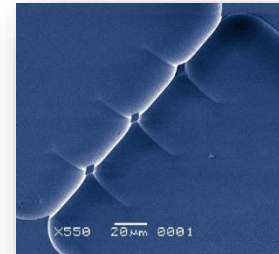
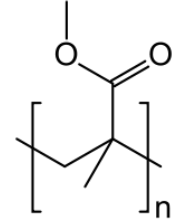
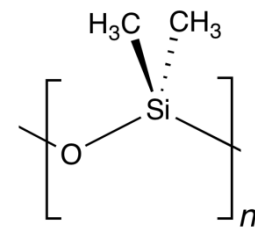
### CNC $\mu$ -MILLING



# Design and fabrication

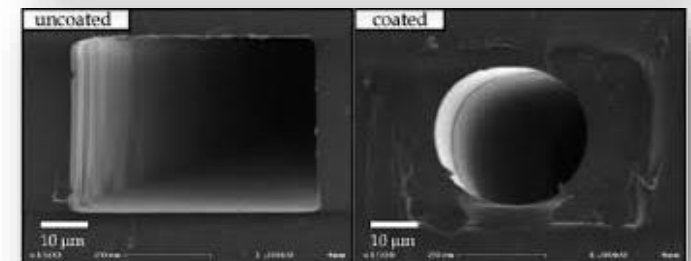
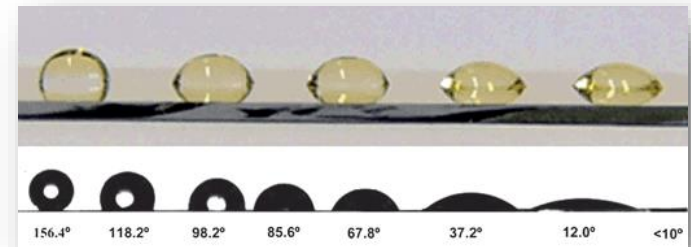
## □ materials

- inert and transparent
- PDMS - poly(dimethyl siloxane)
- PMMA - poly(methyl methacrylate)
- fused silica, quartz and glass



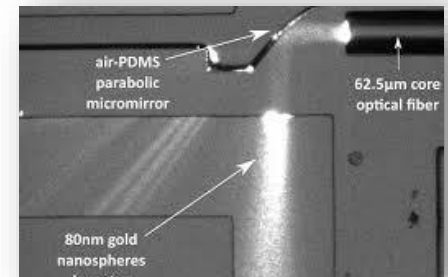
## □ surface modification

- plasma treatment
- silanization
- sol-gel coating

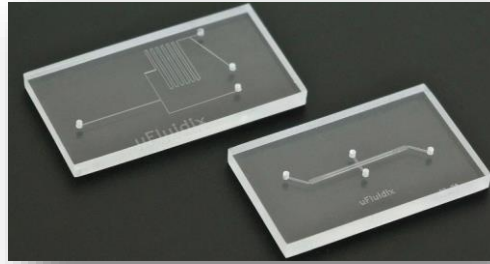
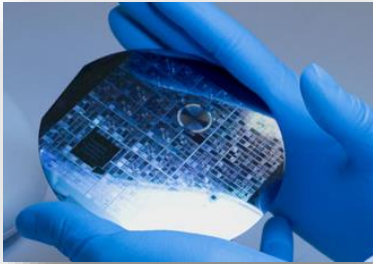


# Sensing and detection

- ❑ processing of **small reagent volumes**
- ❑ **analytical timescale** and performance
- ❑ **on chip detection**
  - fluorescence (LSM, FCS, FLIM)
  - UV/VIS absorbance
  - IR spectroscopy
  - Raman scattering
  - (chemo/electro) luminescence
  - thermal conductivity
  - RI variation
- ❑ **off chip detection**
  - GC, HPLC, MS
  - NMR, X-ray



- customized design and fabrication



- entire technologies



# Conclusions

- ❑ reduced sample/reagent/power consumption
- ❑ superior performance and novel physics
- ❑ applications in life and medical sciences
- ❑ in-house as well as commercial technologies

**microfluidics revolutionize science & technology**



- ❑ Yum, K., 2014: **Physiologically relevant organs on chips**. *Biotechnol. J.* 2014, 9, 16–27
- ❑ 2. *Key elements of microenvironments* (page 18-22)

Review

## Physiologically relevant organs on chips

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<sup>1</sup> Department of Bioengineering, University of California, Berkeley, CA, USA

<sup>2</sup> Department of Materials Science and Engineering, University of Texas, Arlington, TX, USA