

INRA

BIO CELLPHE CELLular protein PHEnotyping at single-cell level Engineered E. coli to produce Raman Reporters (RaRs) that can be detected by Surface-enhaced Raman Scattering (SERS)

Ultrasensitive **BIO**sensing platform for multiplex

Hadi JBARA¹, Saptarshi GHOSH¹, Gustavo BODELON^{2*}, Jean-Loup FAULON^{1*}

¹ Université Paris-Saclay, INRAE, AgroParisTech, Micalis Institute, 78352, Jouy-en-Josas, France ² Departamento de Química Física, Universidade de Vigo, 36310 Vigo, Spain *Corresponding authors: gbodelon@uvigo.es jean-loup.faulon@inrae.fr



- BIOCELLPHE project proposes the generation of an engineered *E.coli* bacteria that binds to circulating tumor cells (CTCs).
- The binding transduces the signal to activate the biosynthesis of small diffusible metabolites (i.e., RaRs) and can be detected by surface-enhaced Raman scattering

Selection RaRs output multiplex SERS

collected a set of 113 molecules



(SERS).

- Three criteria were applied in order to select appropriate RaRs:
- The molecules must display high SERS activity.
- II. The molecules must be unambiguously identified in mixtures.
- III. The biosynthetic pathway must be expressed and optimized in *E.coli*.
- SERS advantages:
- Highly specific and sensitive.
- Achieve single molecule detection level under optimal conditions.
- The narrow bandwidths of Raman spectra facilitate the simultaneous detection of up to 100 distinct molecules.



• After shortlisting, the promising metabolites were analyzed in order to identify those with:

- detectable by SERS from various literature sources [1] [2] [3].
- We ran a machine learning programs trained on selected bacterial metabolites and retrosynthesis software from the Galaxy-SynBioCAD portal [4] for engineering and optimizing their biosynthesis pathways in *E.coli*.
- We built a negative set by randomly drawing molecules from HMBD [5].

Homocystine	0.9±0.05
L-Tryptophanamide	0.9±0.05
4-Imidazoleacetic Acid	0.89±0.03
Indole-3-Acetic Acid	0.88±0.11
1-Methylnicotinamide	0.88±0.05
peonidin	0.87±0.04
Methylguanidine	0.87±0.08
Dimethyl-1,4-Phenylenediamine	0.84±0.08
malvidin	0.83±0.06
Dopamine	0.82±0.05
2-Quinolinecarboxylic Acid	0.81±0.1
Glutathione	0.79±0.11
Pipecolate	0.79±0.08
Octopamine	0.77±0.07
Lumichrome	0.74±0.09
rosinidin	0.74±0.05
N-Methyl-D-Aspartic Acid	0.74±0.07
europinidin	0.74±0.05
Kynurenine	0.73±0.09
caffeine	0.72±0.09
N-Acetyl-L-Cysteine	0.69±0.07
Lipoamide	0.68±0.1
N-Methyltryptamine	0.68±0.08
3-Methoxytyramine	0.66±0.06
scytonemin	0.64±0.08
Dethiobiotin	0.63±0.08
pyocyanin	0.62±0.07
Riboflavin	0.6±0.07
L-Methionine Sulfoximine	0.59±0.07
Biliverdin	0.58±0.08
Mandelic Acid	0.57±0.06

 Table 1: Metabolic pathways producing
SERS detectable molecules in *E.coli* DH5α

Generation of biosynthesis pathways for output RaRs

Higher SERS activity Ι.

- simultaneous detection based on Amenable their Raman to fingerprints.
- candidates were selected and analyzed by SERS following published procedures developed by University of Vigo (UVIGO).









 From the histogram it is evident that RBS-3 has the highest yield of PDV (60 mg/ml) and DV (35,89 mg/ml).

- By using machine learning approach, RaRs are shortlisted from a large pool of candidates and then screened by retrosynthesis software (table 1).
- UVIGO showed that 5 molecules are SERS-active (Figure 1) and four of them (prodeoxyviolacein, desoxyviolacein, violacein and pyocyanin) can de distinguished in mixtures by PCA analysis.
- After the selection of the RaRs, the next step is to create the pathways for these molecules in *E.coli*

References

[1]

[2]

[3]

[4]

[5]

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