



From Nicotine-Containing Waste to Value-Added Chemicals: The Case of *Paenarthrobacter nicotinovorans*

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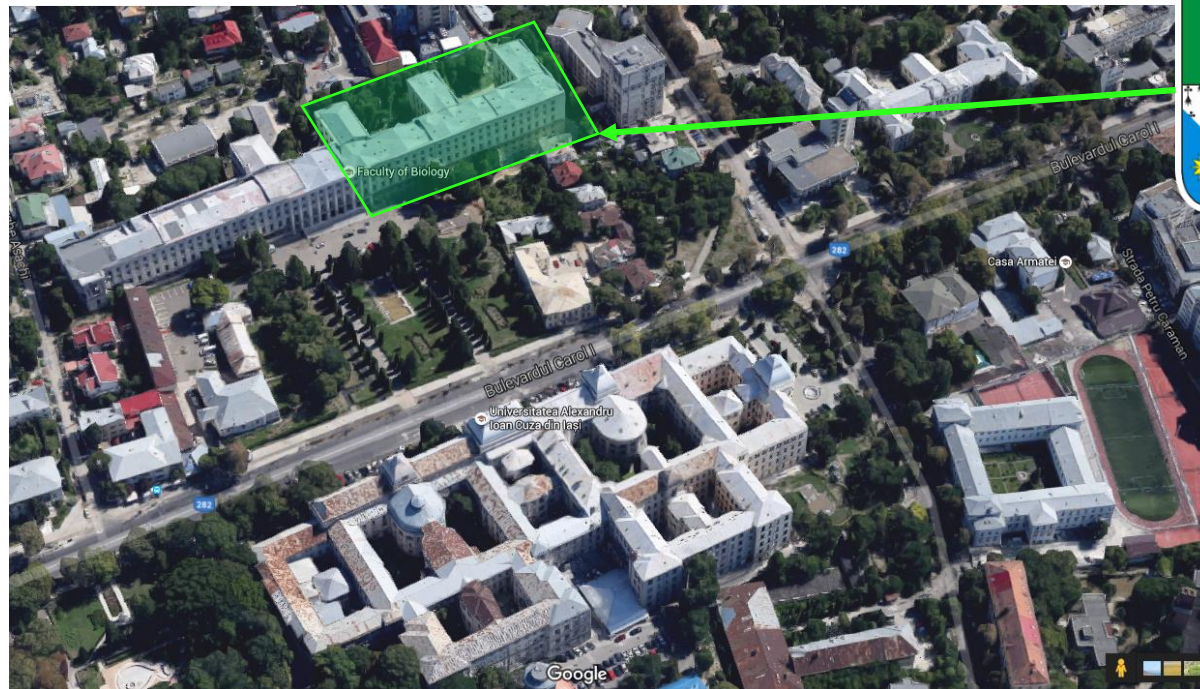
<http://cercetare.bio.uaic.ro/grupuri/bioactive/index.html>



A bit about us and our work in Iași

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www.bio.uaic.ro

BioActive research group

- Isolation
- Identification
- Characterization
- Biological effects (neurological effects, cytotoxicity, oxidative stress, antimicrobial activity)

of biological active molecules with potential applications in biotechnology.



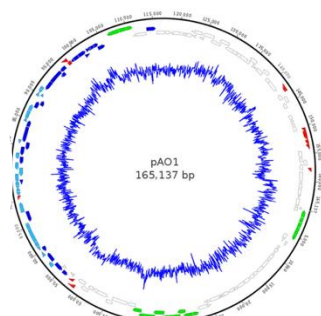
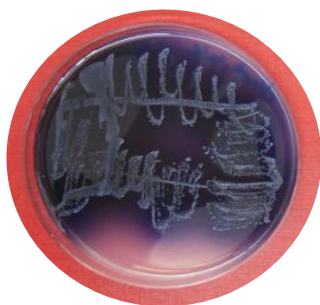


A bit about us and our work in Iași

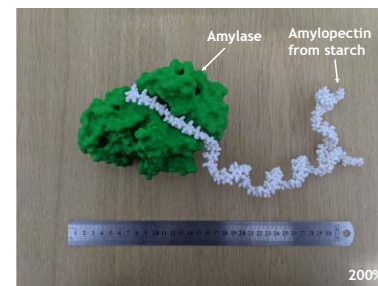
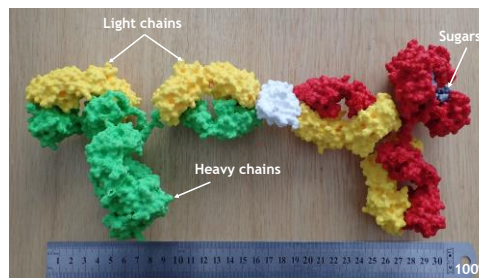
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Main research subjects:

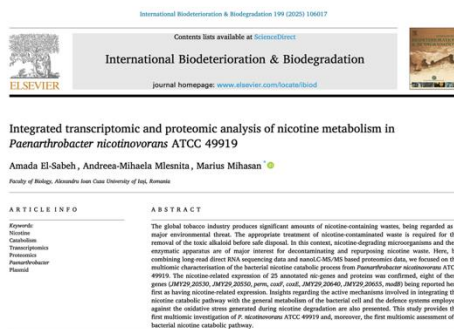
- molecular biology of pAO1 megaplasmid related to nicotine catabolism, stress induced by nicotine degradation and biotechnological applications..



- using 3D printing for creating teaching materials to support molecular bioscience education.



Latest paper:



ICMB2025

Wed 05.11, 18:10

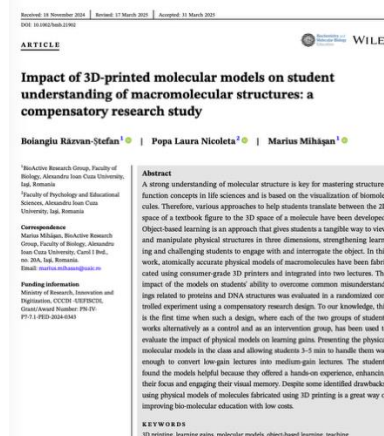
Tiberius Iustin Munteanu

Thu 6.11, 9:55 – 10:15

Andreea Mihaela Mlesnita

[10.1016/j.ibiod.2025.106017](https://doi.org/10.1016/j.ibiod.2025.106017)

Latest paper:



Abstract

A strong understanding of molecular structure is key for mastering structure-function concepts in life sciences and is based on the visualization of biomolecules. Therefore, various approaches to help students translate between the 2D space of a textbook figure to the 3D space of a molecule have been developed. Object-based learning is an approach that gives students a tangible way to view and manipulate physical structures in three dimensions, strengthening learning and challenging students to engage with and interrogate the object. In this work, anatomically accurate physical models of macromolecules have been fabricated using consumer-grade 3D printers and integrated into two lectures. The impact of the models on students' ability to overcome common misunderstandings related to proteins and DNA structure was evaluated in a randomized controlled experiment using a compensatory research design. To our knowledge, this is the first time when such a design, where each of the two groups of students works alternatively as a control and as an intervention group, has been used to evaluate the impact of physical models on learning gains. Presenting the physical molecular models in the class and allowing students 3–5 min to handle them was enough to convert low-gain lecture into medium-gain lectures. The students found the models helpful because they offered a hands-on experience, enhancing their focus and engaging their visual memory. Despite some identified drawbacks, using physical models of molecules fabricated using 3D printing is a great way of improving bio-molecular education with low costs.

KEYWORDS

3D printing, learning gains, molecular models, object-based learning, teaching

[10.1002/bmb.21902](https://doi.org/10.1002/bmb.21902)

Developed tools:



Jmol/JSmol script for automatic generation of 3D printable molecular models from structural data.

ICMB2025

Fri 07.11, 11:10



Paenarthrobacter nicotinovorans

Biology Department, Alexandru Ioan Cuza University of Iași, Romania

> [J Biol Chem](#). 1959 Jan;234(1):151-5.

The bacterial oxidation of nicotine. I. Nicotine oxidation by cell-free preparations

> [J Biol Chem](#). 1972 Dec 10;247(23):7628-33.

The bacterial oxidation of nicotine. 8. Synthesis of 2,3,6-trihydroxypyridine and accumulation and partial characterization of the product of 2,6-dihydroxypyridine oxidation

> [J Bacteriol](#). 2003 Mar;185(6):1976-86. doi: 10.1128/JB.185.6.1976-1986.2003.

Sequence of the 165-kilobase catabolic plasmid pAO1 from *Arthrobacter nicotinovorans* and identification of a pAO1-dependent nicotine uptake system

> [J Mol Evol](#). 2013 Aug;77(1-2):22-30. doi: 10.1007/s00239-013-9576-x.

pAO1 of *Arthrobacter nicotinovorans* and the spread of catabolic traits by horizontal gene transfer in gram-positive soil bacteria

Marius Mihasan¹, Roderich Brandsch

Affiliations + expand

PMID: 23884627 DOI: [10.1007/s00239-013-9576-x](#)





Nicotine - a lead compound for developing drugs

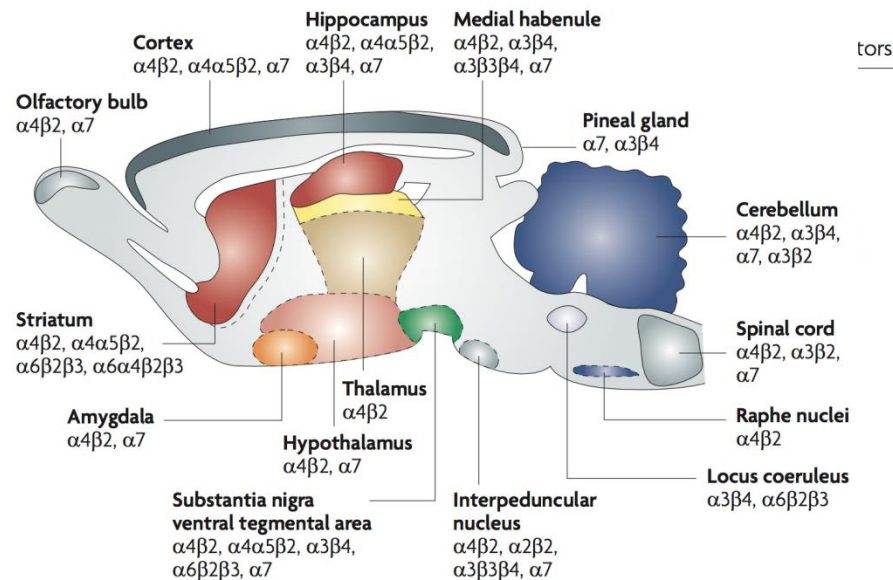
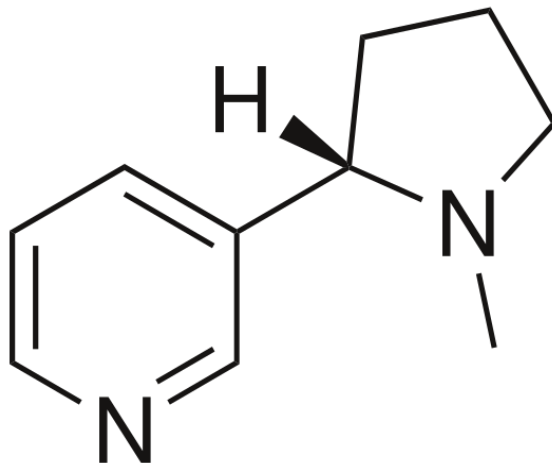
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Nicotine

- Produced by the tobacco plant to prevent consumption by herbivores
- **Responsible for smoking addiction** - by releasing dopamine
- Potentially lethal to humans (LD₅₀ 0.5 - 1.0 mg/kg)
- **Improves: cognition, alertness, memory**



Nicotiana tabacum



nAChRs - Nicotine acetylcholine receptors

ligand-gated ion channel superfamily of neurotransmitter receptors

Nicotine is a lead compound used for developing drugs for cognitive dysfunctions

Levin ED and Rezvani AH. Nicotinic treatment for cognitive dysfunction. Curr DrugTargets CNS Neurol Disord. 2002;1(4):423-31.

Taly et. al. Nicotinic receptors: allosteric transitions and therapeutic targets in the nervous system Nat. Rev. Drug Discov. 2009, 8, 733-750.



6-hydroxy-L-nicotine – a neuroprotective agent

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Nicotine binds to nAChRs subtypes $\alpha 4\beta 2$ -

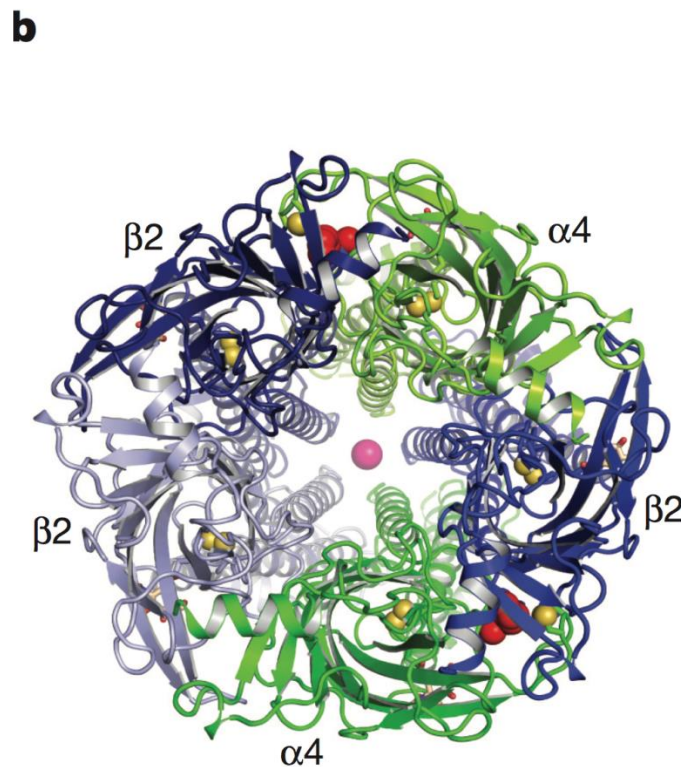
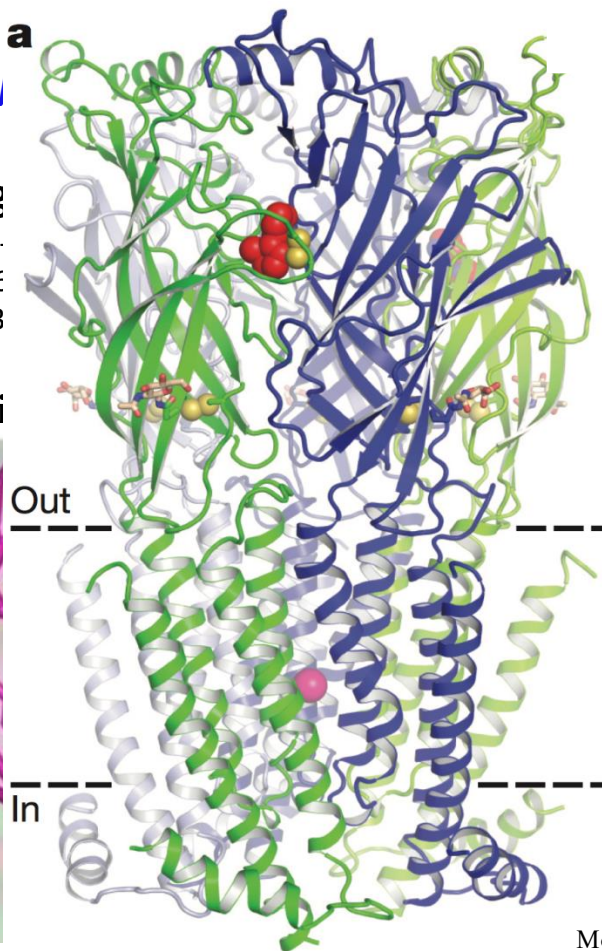
Murray and Abeles, *Aging & Mental Health*. 2002, 6, 129–138.

Could
Paenarthro

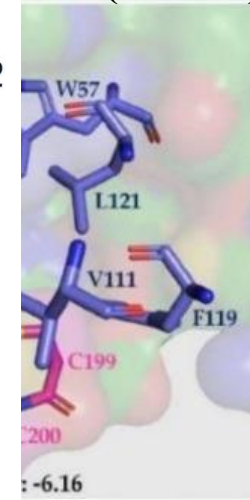
In silico - Target

- rigid receptor - $\alpha 4$
- targeted region - ϵ
- flexible ligands - 3

(S)-Nicotine



otine(docked)



Morales-Perez C et al., *Nature*, 2016, 538 (7625): 411–415

[10.3390/antiox10020212](https://doi.org/10.3390/antiox10020212)



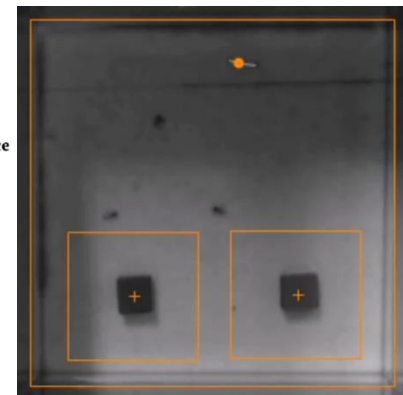
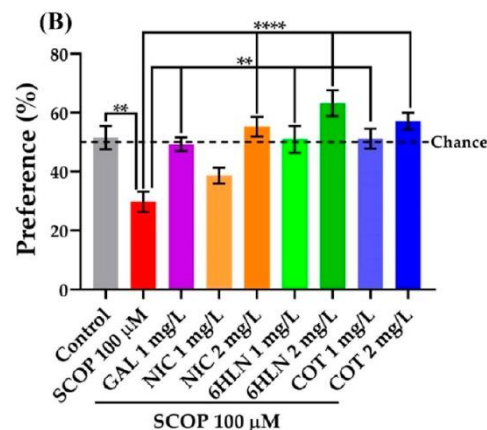
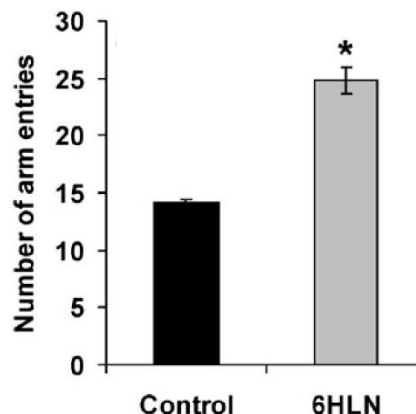
6-hydroxy-L-nicotine – a neuroprotective agent

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In-vivo tests

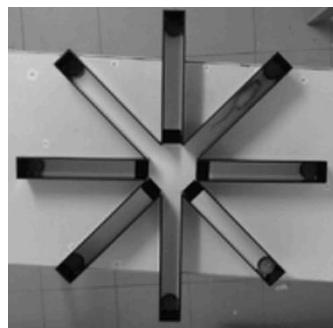
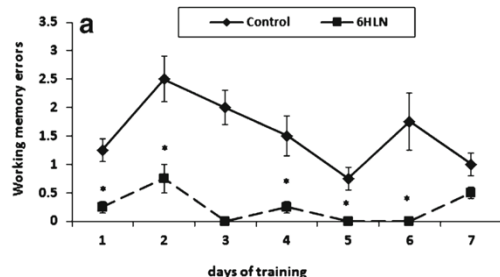
● male Wistar rats (3-4 months old)

● zebrafish (*Danio rerio*), both sexes (50:50)



Y-maze task

Values are mean $\pm$ S.E.M. (n=10 animals per group), *p<0.0001 vs. control group



Radial arm-maze task

Values are mean $\pm$ S.E.M. (n=10 animals per group), *p<0.0001 vs. control group

[10.1016/j.biopha.2016.12.008](https://doi.org/10.1016/j.biopha.2016.12.008)

Object Discrimination Test

- Values are mean $\pm$ S.E.M. (n=10 animals per group), F(8,81) = 8.078, p < 0.0001. For Tukey posthoc analyses—**** p < 0.0001, and ** p < 0.001



Prof. Lucian H., PhD



Radu I.



Razvan B.

[10.3390/antiox9080768](https://doi.org/10.3390/antiox9080768)



Nicotine-containing waste need to be decontaminated

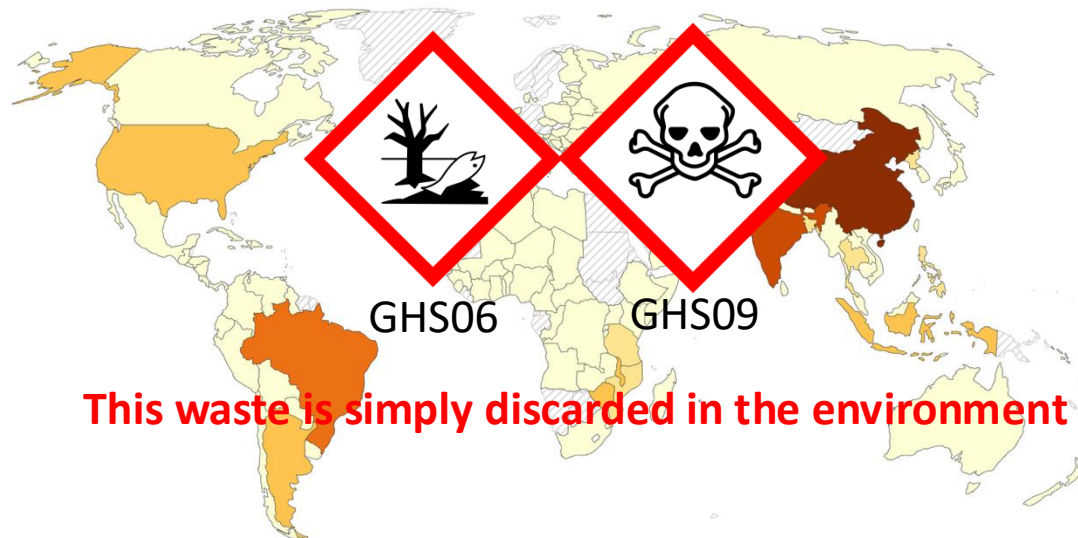
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Tobacco industry produces 300 000 tons of non-recyclable nicotine-containing waste per year, 18 g nicotine/per kg

- According to EPA the waste is **Toxic Release Inventory (TRI)** chemicals
- According to the European Union Regulations (EUR) - “**toxic and hazardous**”(when nicotine content is over 0.05% w/w)

Tobacco production, 2021

Tobacco production is measured in tonnes.



Why not using *Paenarthrobacter nicotinovorans* to decontaminate this waste and convert it to useful chemicals?



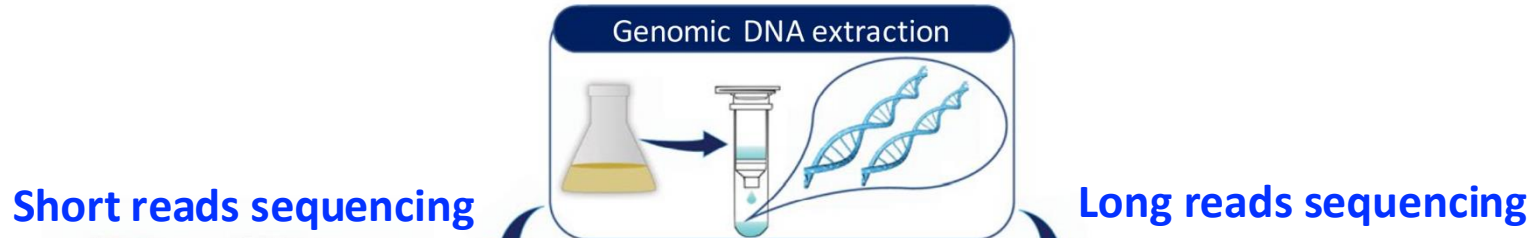
Source: Food and Agriculture Organization of the United Nations

OurWorldInData.org/agricultural-production • CC BY



Sequencing the *Paenarthrobacter nicotinovorans* genome

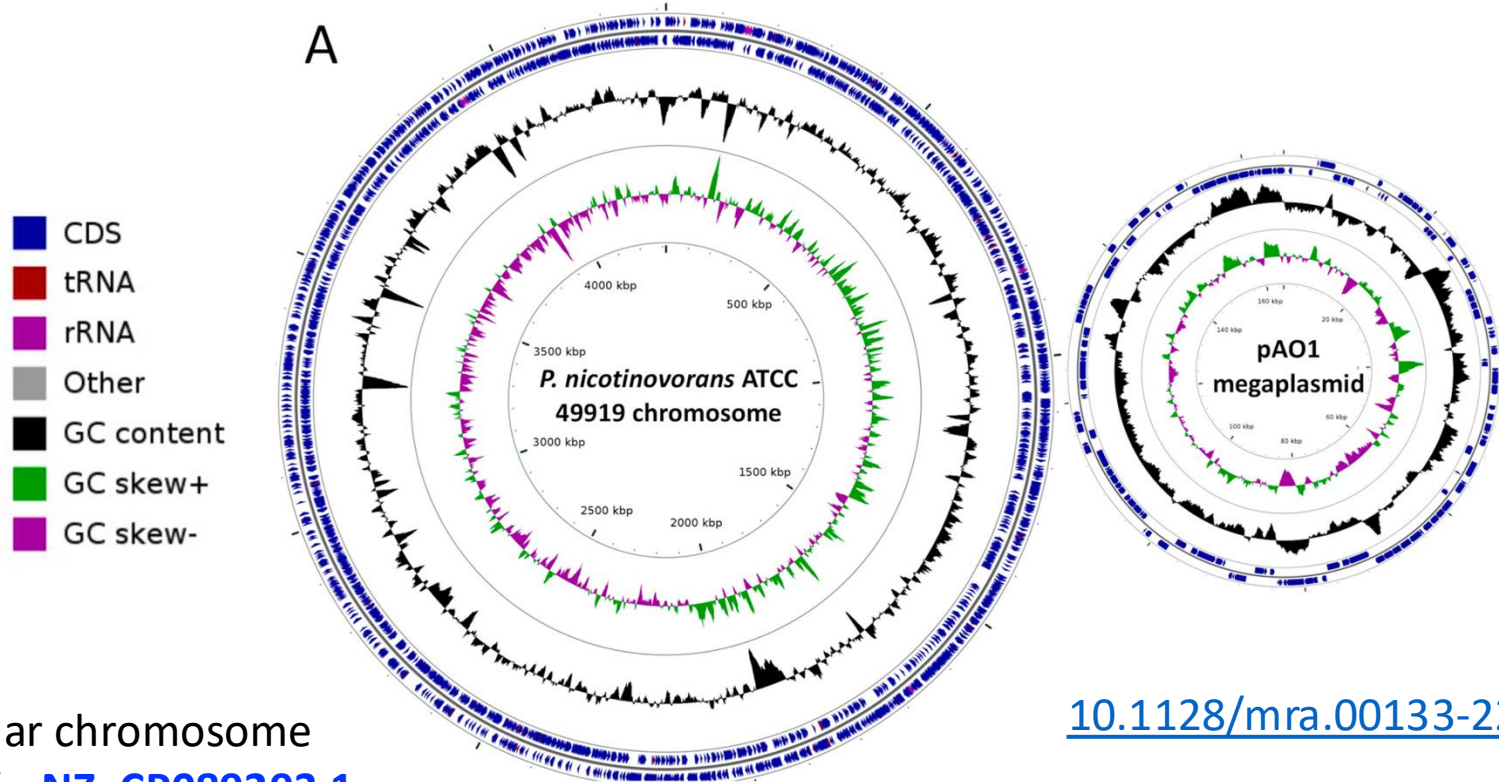
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Sequencing the *Paenarthrobacter nicotinovorans* genome

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[10.1128/mra.00133-22](https://doi.org/10.1128/mra.00133-22)

4 316 184-bp circular chromosome
GC content - 63.2% - [NZ_CP089293.1](https://doi.org/10.1128/mra.00133-22)
165 141-bp circular megaplasmid
GC content - 59.7% - [NZ_CP089294.1](https://doi.org/10.1128/mra.00133-22)



Amada ES.

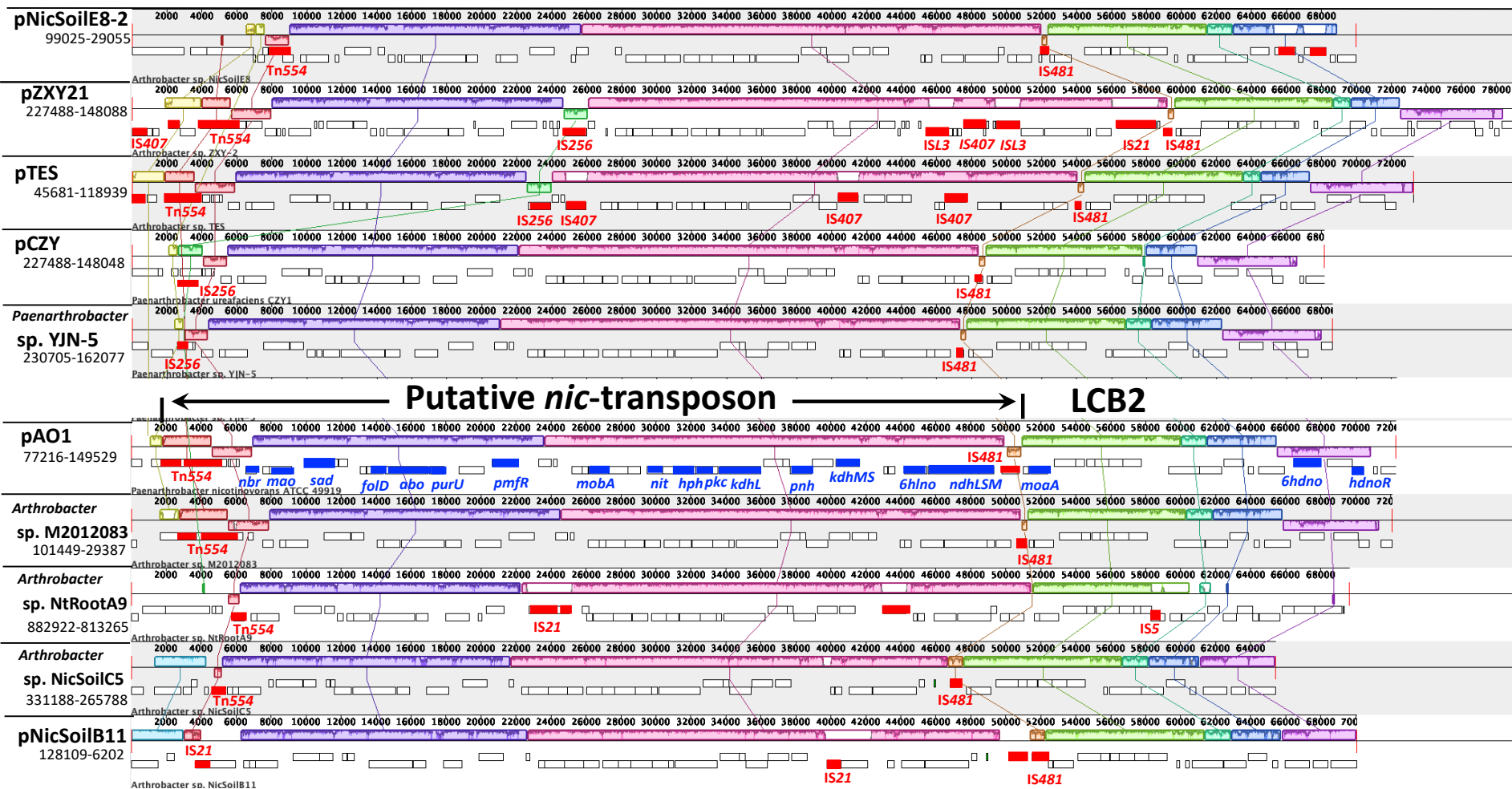
CheckM analysis
Completeness: 85.21%
Contamination: 10.14%





nic-genes cluster as a putative catabolic transposon

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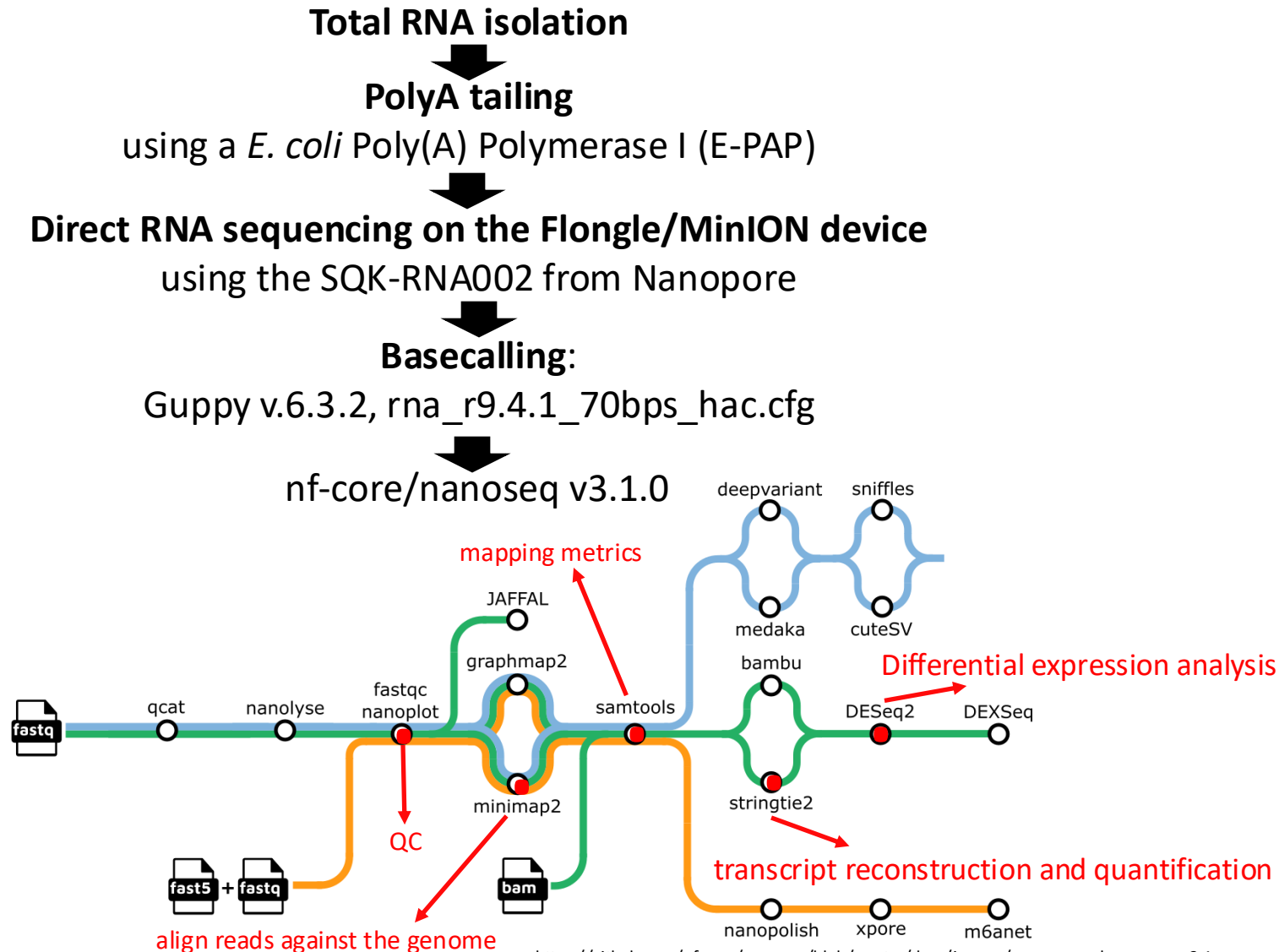


[10.1186/s12864-023-09644-3](https://doi.org/10.1186/s12864-023-09644-3)



But when and which genes are expressed when nicotine is present?

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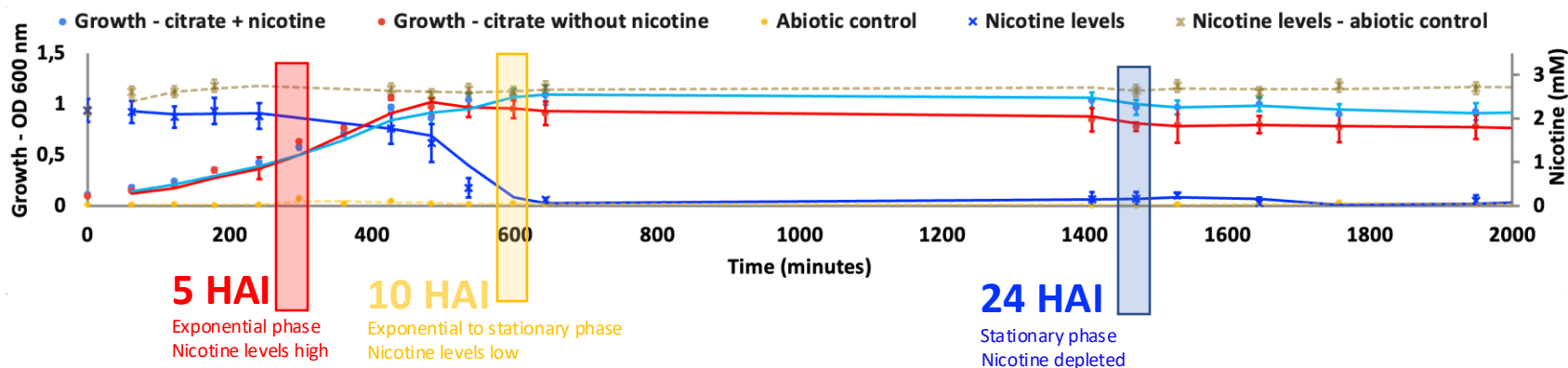


https://github.com/nf-core/nanoseq/blob/master/docs/images/nanoseq_subwaymap_v3.1.png



Nicotine-related transcriptome of *Paenarthrobacter nicotinovorans*

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Differentially expressed genes (citrate vs citrate + nicotine)

pADJ < 0.1; log2FoldChange > 1	pAO1 nic-genes cluster			Chromosome encoded		
	5 HAI	10 HAI	24 HAI	5 HAI	10 HAI	24 HAI
	<i>nbr, mao, sad, abo, coxF, coxD, nit, coxG, hph, pkc, hyp, kdhL, pnh, kdhSM, 6-hln, ndhLSM, moaA, moaE, modA, modB, modC</i>	<i>nbr, mao, sad, abo, coxD, nit, coxG, hph, pkc, hyp, kdhL, pnh, kdhSM, 6-hln, ndhLSM, moaE, modA, modB, modC</i>	nd	site-specific integrase; argininosuccinate synthase; helix-turn-helix transcriptional regulator; Fis family transcriptional regulator; argB (acetylglutamate kinase); acetylornithine transaminase; argF (ornithine carbamoyltransferase); arginine repressor; ATP-dependent chaperone ClpB; RNAs and ribosomal proteins	site-specific integrase; argininosuccinate synthase; helix-turn-helix transcriptional regulator; Fis family transcriptional regulator; universal stress protein; tripartite tricarboxylate transporter permease; tripartite tricarboxylate transporter TctB family protein; tripartite tricarboxylate transporter substrate binding protein;	nd

Bold – genes not previously shown to have nicotine related expression;

Red – genes differentially expressed between 5 HAI and 10 HAI.



P. nicotinovorans – nicotine induced proteome

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Growth, harvest and cell lysis

0.05% nicotine No nicotine



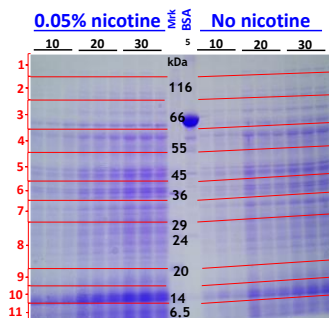
Lysozyme
SDS
DTT

12,000 g
30 min

Soluble proteins

Protein preparation

Protein fractionation
SDS-PAGE maxi
9-16% gradient gels



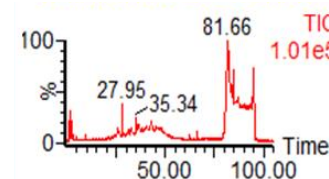
In-gel digestion

Peptide mixture

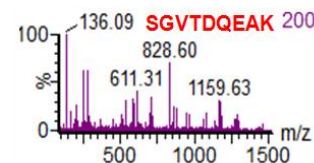


Identification and data analysis

Nano-liquid
chromatography tandem
mass spectrometry



Protein identification by mass
spectrometry



Data analysis: MASCOT;
SCAFFOLD

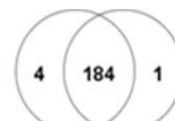
Chromosome

Nic + -



pAO1

+ - Nic



Darie Costel, PhD
Clarkson University
Potsdam, NY, USA



Devika C.



Emma D.



Roshi A.





P. nicotinovorans – nicotine induced proteome

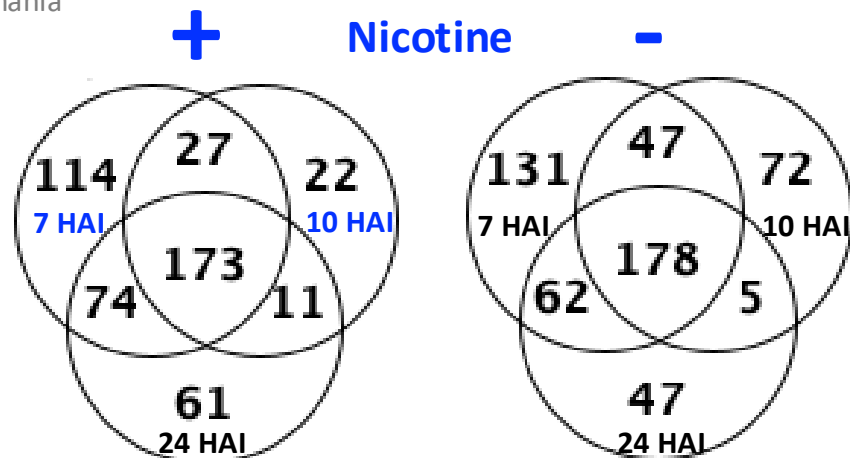
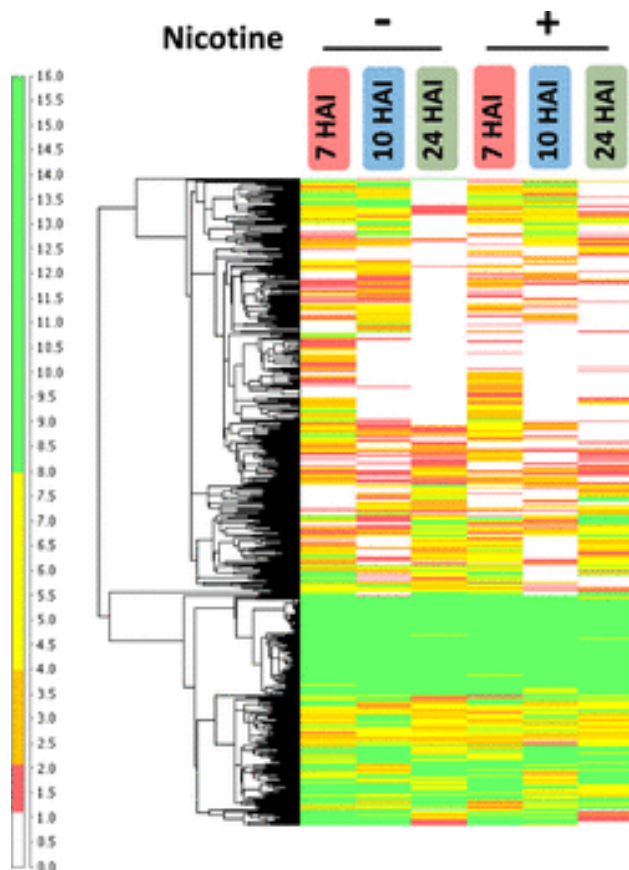
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44136 MS/MS spectras assigned

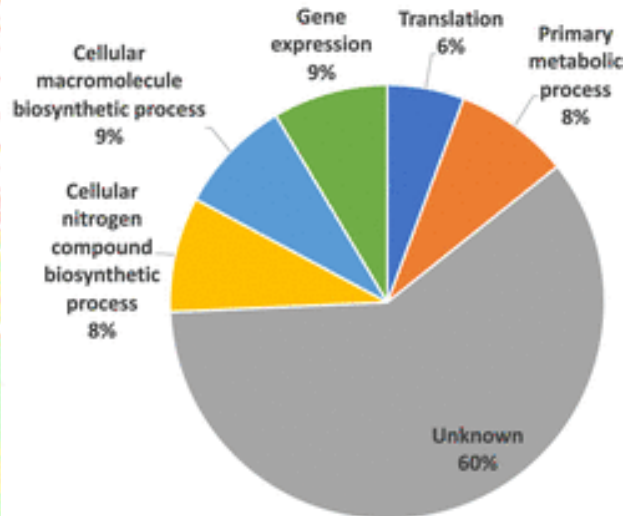
2 peptides minimum

607 proteins at 1% FDR

ProteomeXchange: PXD012577



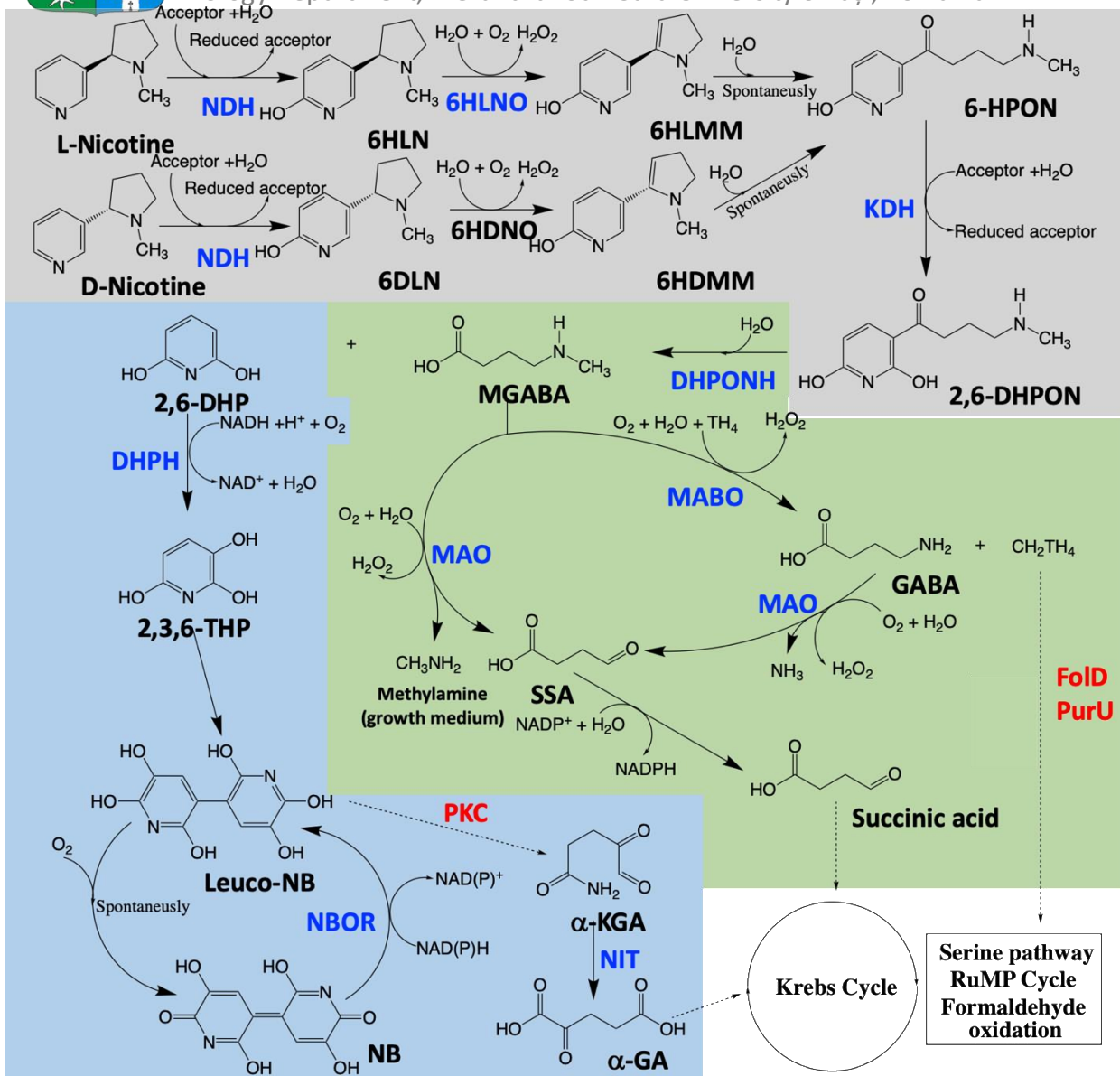
Percent of proteins with significant fold change





Proteomic and transcriptomic view over the nicotine degradation pathway

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Detected by MS/MS and transcriptomics, known

to be involved in nicotine metabolism

Detected by MS/MS and transcriptomics, putatively involved

in nicotine metabolism

Detected by MS/MS and transcriptomics,

CoxF

CoxD

CoxG

MoaA

MoaE

ModA

ModB

ModC

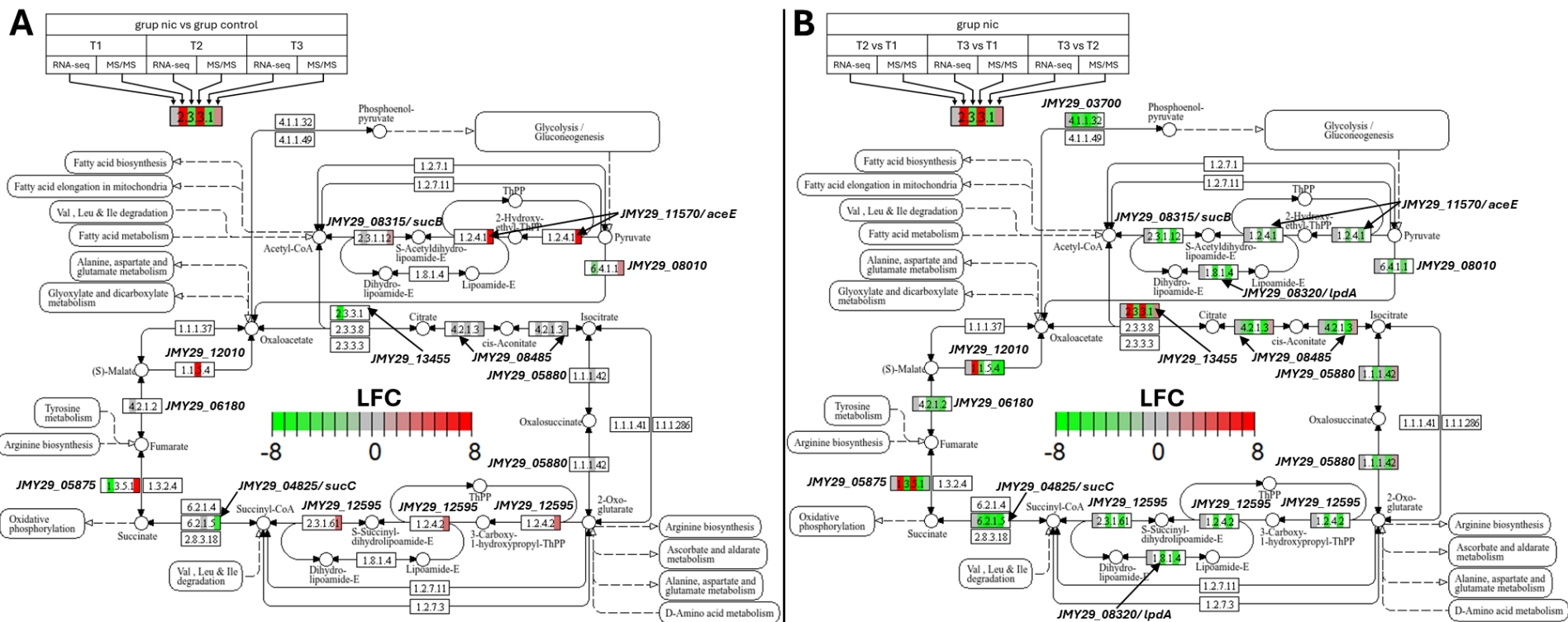
Molybdenum intake / cofactor / multimeric enzyme assembly



Integration of transcriptomics and proteomics data

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PATHVIEW



Differentially expressed genes (DEGs) and proteins (DEPs) involved in TCA:

A. nicotine vs control;

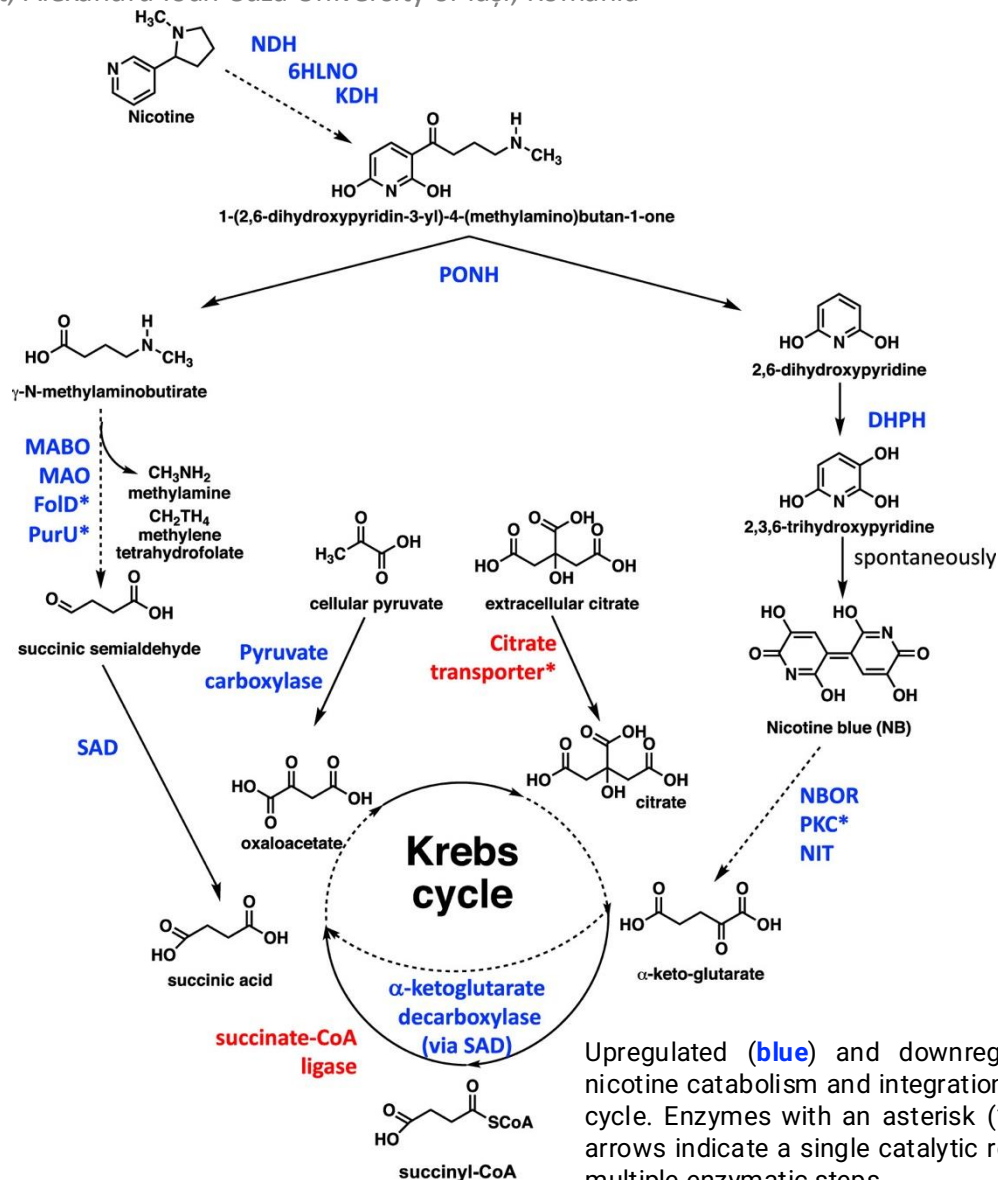
B. time course analysis in the nicotine-induced cells; LFC: log2FoldChange;

[10.1016/j.ibiod.2025.106017](https://doi.org/10.1016/j.ibiod.2025.106017)



Nicotine metabolism integration

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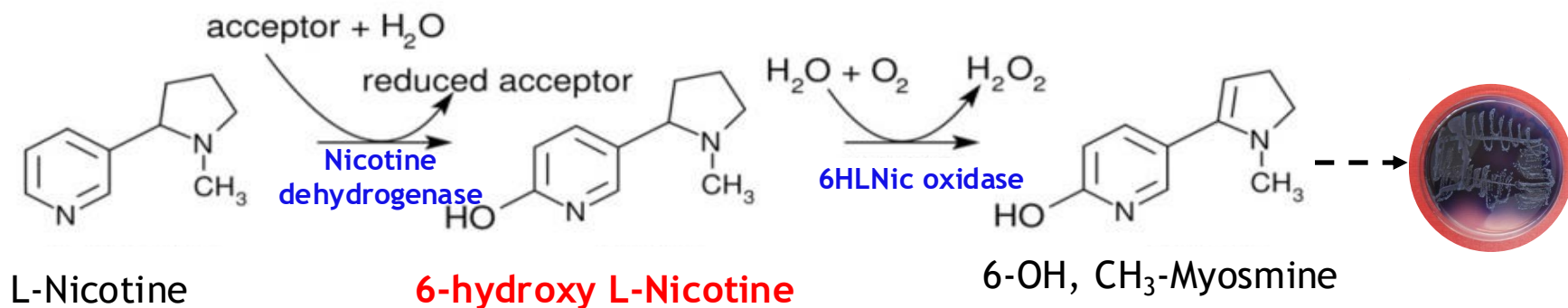


Upregulated (blue) and downregulated (red) proteins involved in nicotine catabolism and integration of the end products into the Krebs cycle. Enzymes with an asterisk (*) have putative functions. The full arrows indicate a single catalytic reaction; the dashed arrows indicate multiple enzymatic steps.



A *P. nicotinovorans* based biotechnology to produce 6HLNic

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Diana M., B.Sc.



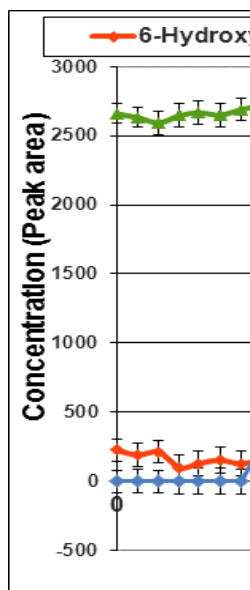
Ana I., B.Sc.



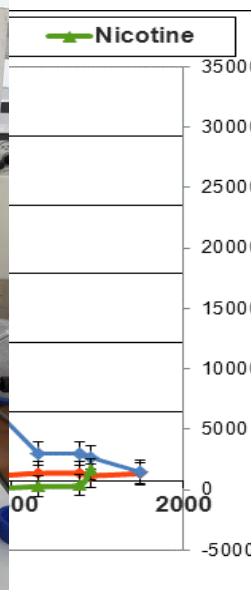
Andreea P., B.Sc.



Carmen C., B.Sc.



Patent pending



Razavn B., B.Sc.

Doina G. B.Sc.

Andreea A., M.Sc.



Madalina M., B.Sc.



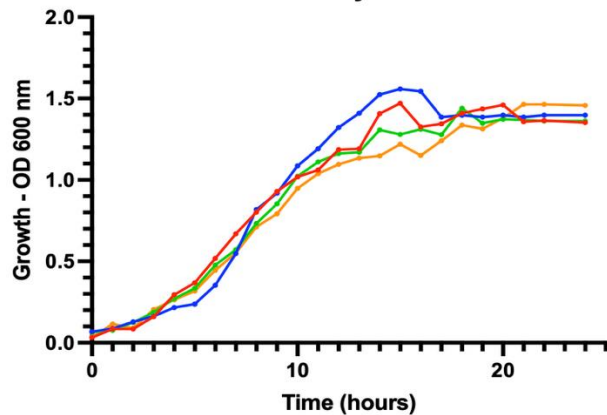
Loredana C., B.Sc.



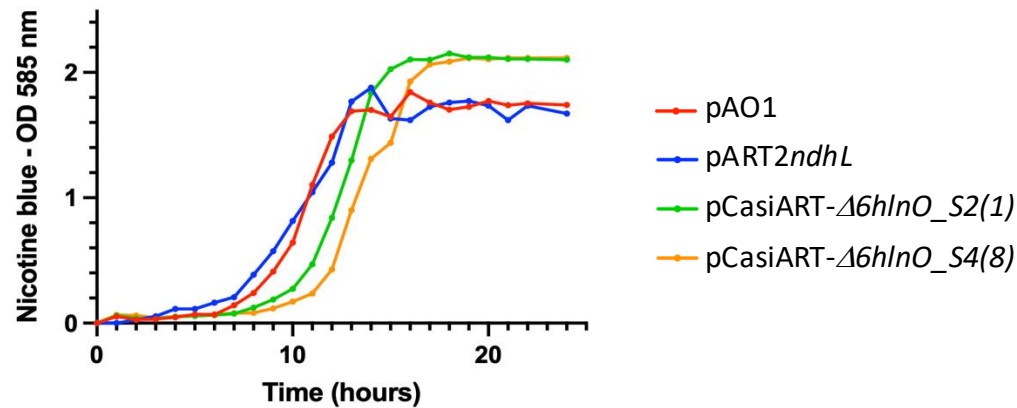
A *P. nicotinovorans* based biotechnology to produce 6HLNic

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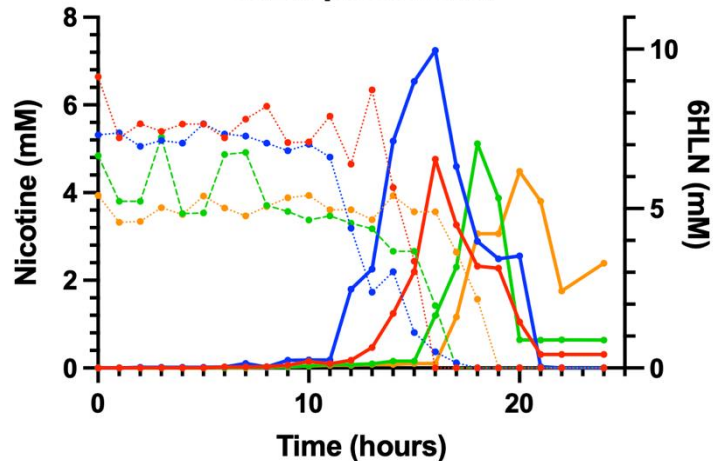
Growth dynamics



Nicotine blue acumulation



6HLN production



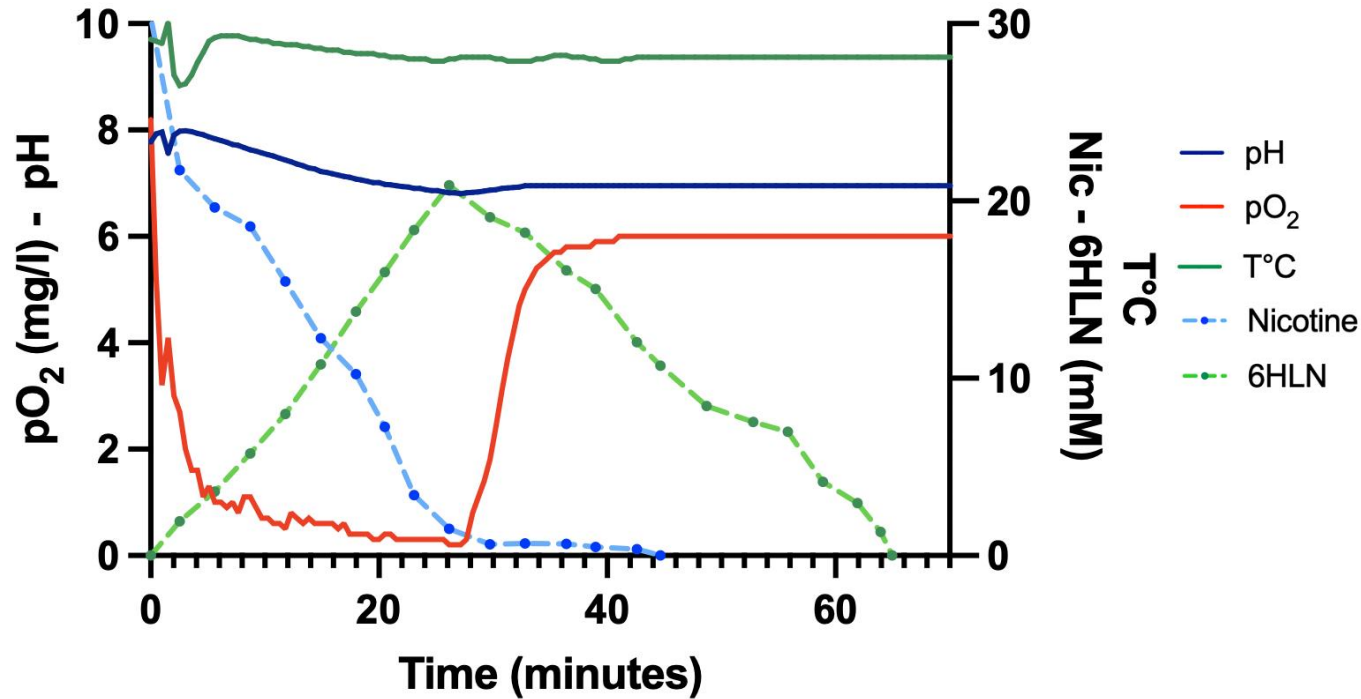
Values	pAO1	pART2ndhL	pCasiART- $\Delta 6hInO_S2(1)$	pCasiART- $\Delta 6hInO_S4(8)$
$\mu_{max_OD\ 600}$ (μ/h^{-1})	0.29	0.41	0.27	0.27
Lag_OD 600 (h^{-1})	5.25	3.2	5.22	4.25
$\mu_{max_OD\ 585}$ (μ/h^{-1})	0.52	0.50	0.56	0.66
Lag_OD 585 (h^{-1})	6.66	6.11	8	10.02
Nic_t½ (mM/h ⁻¹)	15.50	13.87	15.42	18.62
6HLN Titres (g/L)	1.16	1.77	1.25	1.09
STY (g/L/h ⁻¹)	0.068	0.104	0.066	0.052



A *P. nicotinovorans* based biotechnology to produce 6HLN

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pART2ndhL



Tiberius I. M.



Summary

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Paenarthrobacter nicotinovorans can **break-down nicotine** and use it for growth

The genome of *Paenarthrobacter nicotinovorans* was sequenced *and* contains **one chromosome** and **one plasmid**

The **nicotine proteome and transcriptome were solved**, and we now have a better idea how nicotine catabolism is regulated and integrated into the general metabolic pathways.

We are on our way to modify the bacteria so that it will **convert nicotine to** useful compounds such as **6-hydroxy-L-nicotine** more efficiently



Thanks

Biology Department, Alexandru Ioan Cuza University of Iași, Romania

Collaborators:



Prof. Darie Costel, PhD
Lead, Biochemistry & Proteomics Group
Department of Chemistry & Biomolecular Science,
Clarkson University, Potsdam, NY, USA
- *nicotine induced proteome of P. nicotinovorans*



apl. Prof. Roderich Brandsch, PhD
Institute of Biochemistry and Molecular Biology,
Freiburg i. Br., Germany
- *nicotine metabolism and pAO1 molecular organization*



BioActive research group

<https://www.facebook.com/BioActive.bio.uaic.ro>



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din IAȘI



CommScie

POSDRU/159/1.5/S/133652



PN-II-RU TD 236/2007; PN-II-RU PD 337/2010;
PN-II-RU-TE-2014-4-0106; PN-II 50BM/2016
PN-III-P2-2.1-PED-2016-0177; PN-III-P1-1.1-TE-2016-0367;
PN-III-P4-ID-PCE-2020-0656

GI-2014-02