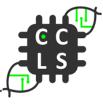
Francisco (Paco) Barona-Gómez, Associate professor IBL (Full professor Cinvestav, Mexico)



Langebio, Cinvestav-IPN, Irapuato, Mexico. (2008 - 2022)

Institute of Biology Leiden University, the Netherlands. (2022 -)



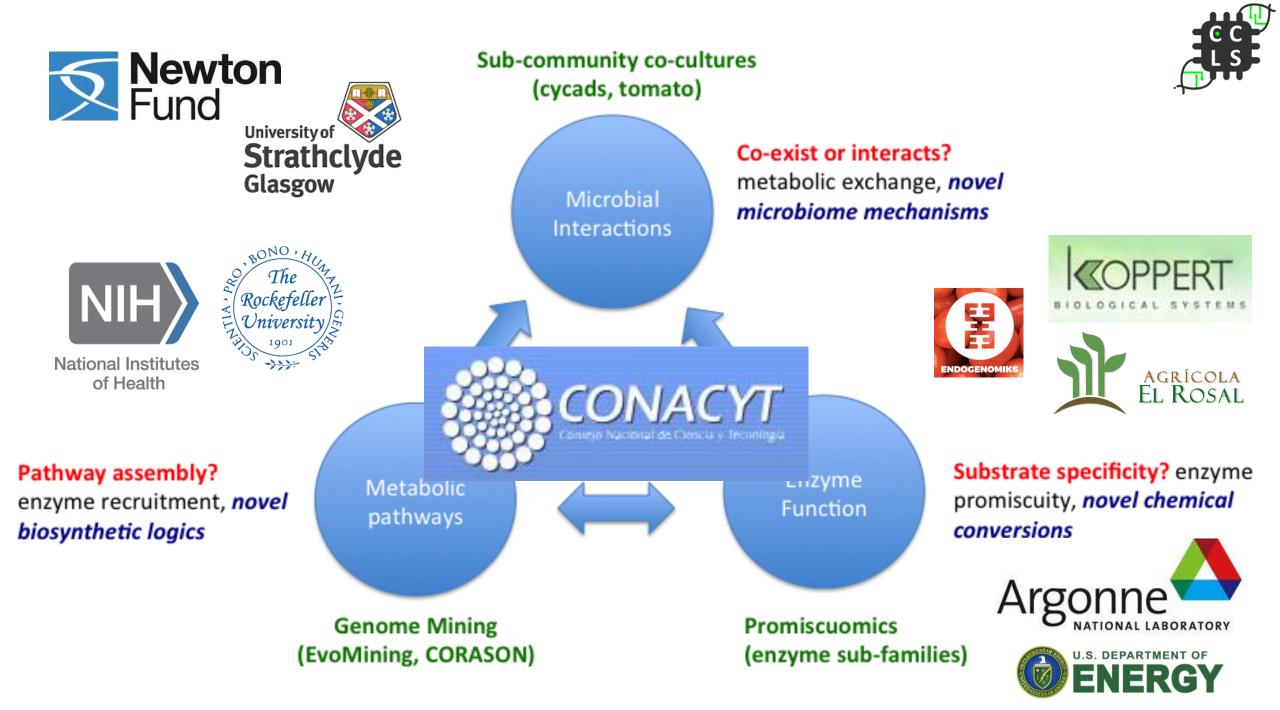
I am interested in deciphering the **evolutionary mechanisms**, at different scales (from atoms to microbial communities) underlying the evolution of **metabolism** during bacterial adaptation.

We investigate keystone bacterial taxa and their specialized metabolism emerging after ecological **hostmicrobiome interactions** studies, mainly in plants and insects.

We dissect the roles played by natural products in mediating **microbial interactions**, as well as deciphering novel **biosynthetic logics** leading to unprecedented **chemical scaffolds**.

Based in this basic knowledge, we develop **bioinformatics tools for genome mining** of natural products and **biotechnological applications**, mainly for agriculture.

Microbial Biodiversity and Specialized Metabolism Laboratory



Natural Product Reports



View Article Online

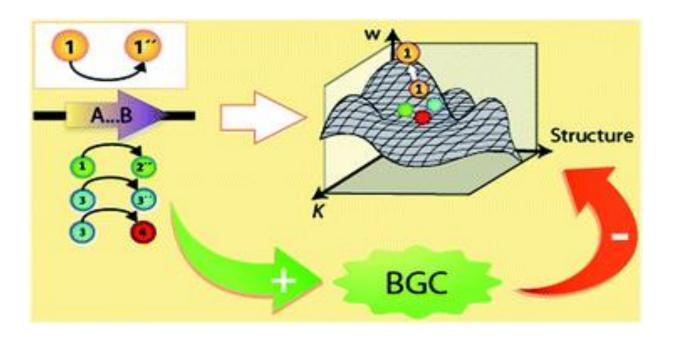


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REVIEW

Evolutionary dynamics of natural product biosynthesis in bacteria

Marc G. Chevrette, ^(D)^a Karina Gutiérrez-García,^b Nelly Selem-Mojica, ^(D)^b César Aguilar-Martínez,^b Alan Yañez-Olvera,^b Hilda E. Ramos-Aboites,^b Paul A. Hoskisson ^{(D) c} and Francisco Barona-Gómez ^{(D) *b}



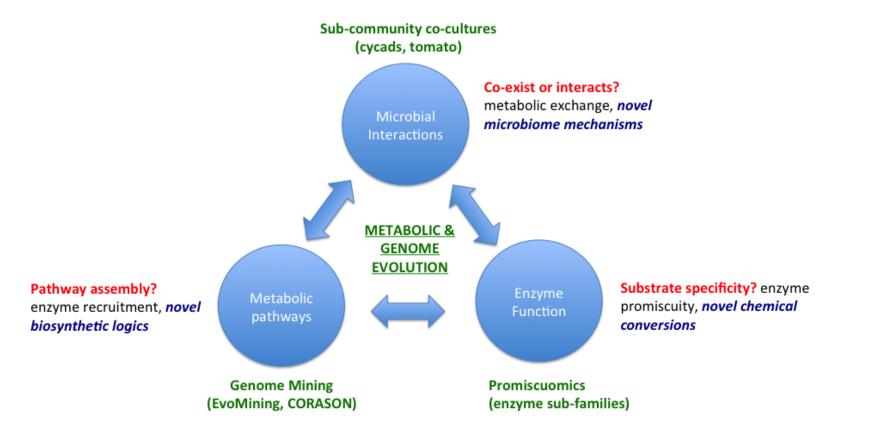
The Dynamic Chemical Matrix Evolutionary (DCME) hypothesis

"The chemical matrix hypothesis stems from the idea that secondary / specialized metabolism evolves from increased metabolic diversity driven by the concerted action of promiscuous enzymes.... BGCs are therefore evolutionary unstable and only the output of this process. Thus, fitness landscapes, defined by the relationship between relative fitness (W), chemical structure (S) and biomolecular activity (F) are a more appropriate representation of the evolution of natural products biosynthesis"

https://jgi.doe.gov/natural-prodcast-episode-13-francisco-barona-gomez/



- 1. Expansion-and-recruitment of promiscuous enzymes is the raw (and constant) material for the evolution of NPs (**EvoMining**)
- 2. NPs evolve after a dynamic gene clustering that increases chemical diversity (CORASON)

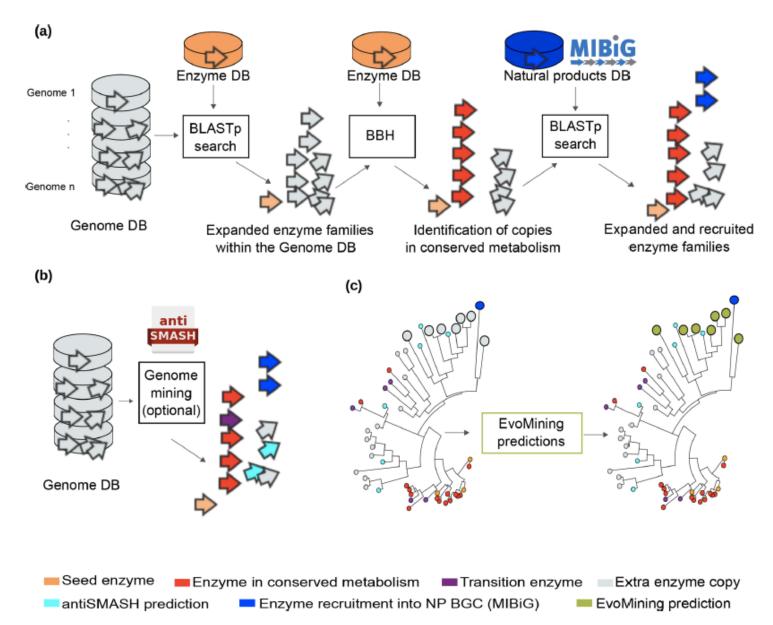


3. NPs are responsive at the population level to environmental conditions and the composition of the community (EcoMining – Experimental. PopGenNP - computational)

EvoMining 1.0: Cruz-Morales et al (2016) *Genome Biology & Evolution*.
EvoMining 2.0: Selem-Mojica et al (2019) *Microbial Genomics*.
EvoMining DB (*ActDES 1.0*) Schniete, Selem-Mojica et al (2021) *Microbial Genomics*



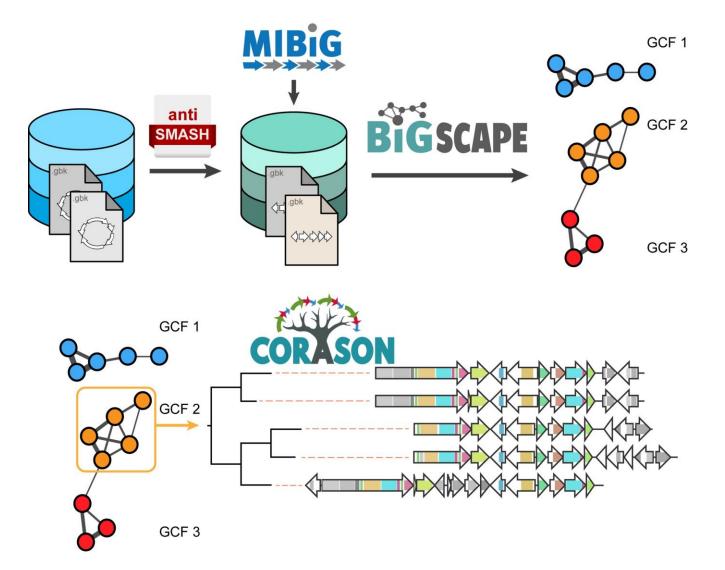




A computational framework for systematic exploration of biosynthetic diversity from large-scale genomic data



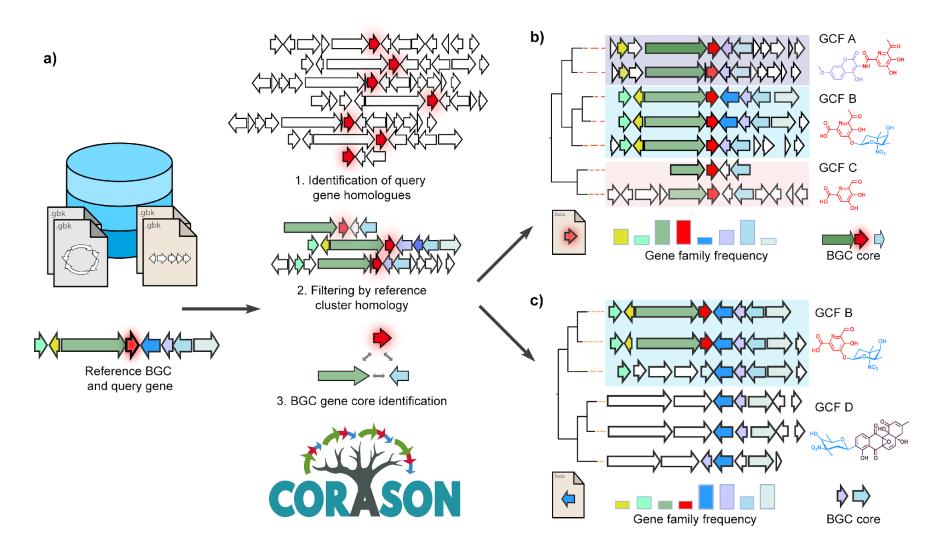
Jorge C. Navarro-Muñoz^{1*}, **Nelly Selem-Mojica^{2*}**, Michael Mullowney^{3*}, (many other co-authors), Neil L. Kelleher^{3#}, **Francisco Barona-Gomez^{2#}**, Marnix H. Medema^{1#} (2020) *Nature Chemical Biology*



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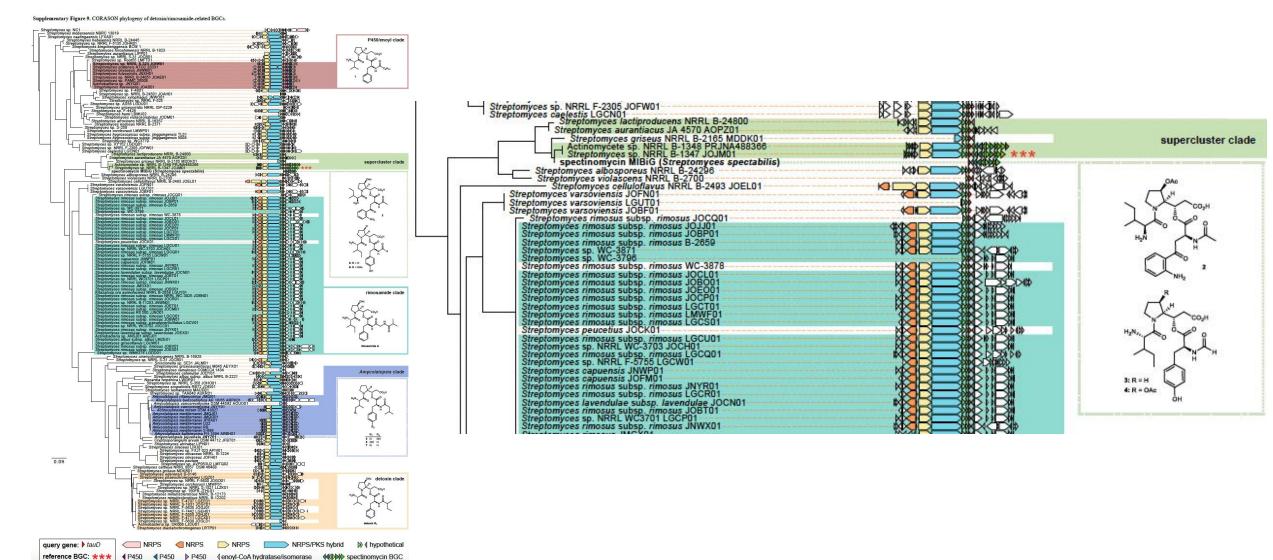
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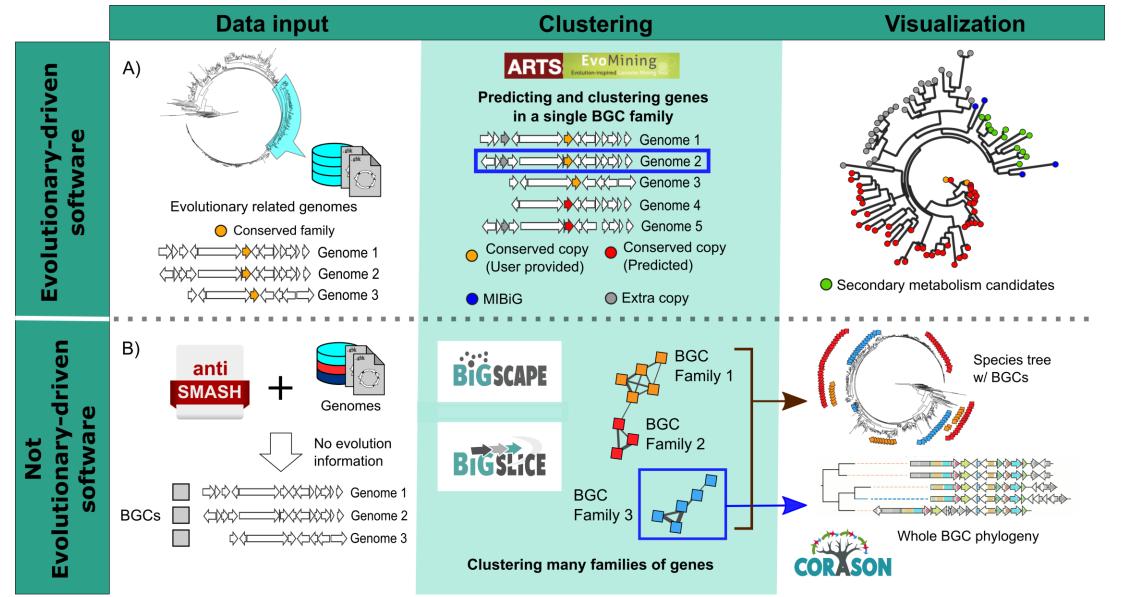


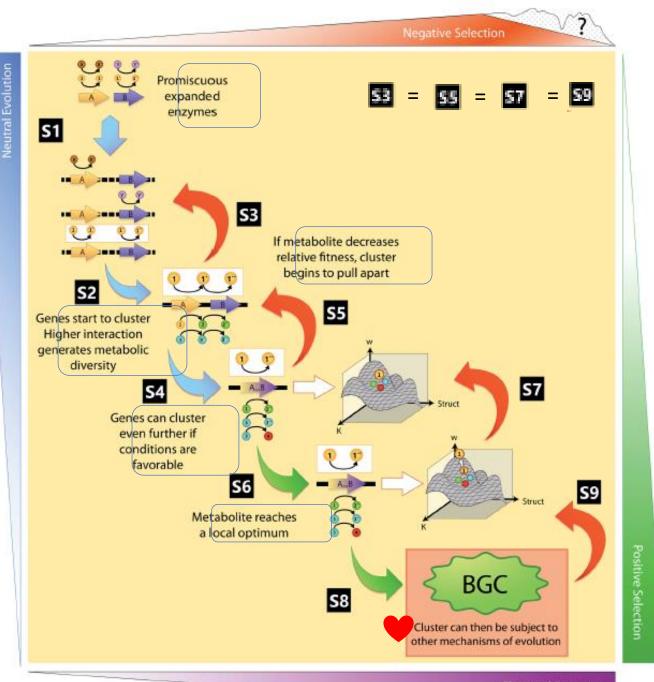
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Chevrette et al (2021) The confluence of Big Data and evolutionary genome mining for the dscoviery of natural products. *Natural Product Reports.* 2024-2040. Chevrette et al (2022) Evolutionary Genome Mining for the Discovery and Engineering of Natural Product Biosynthesis. In Engineering Natural Product Biosynthesis. 129-155







 Catechol-type

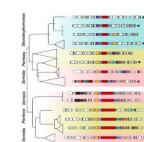
 Promouninetar

 Hydroxample-type

 Mixed Sgand

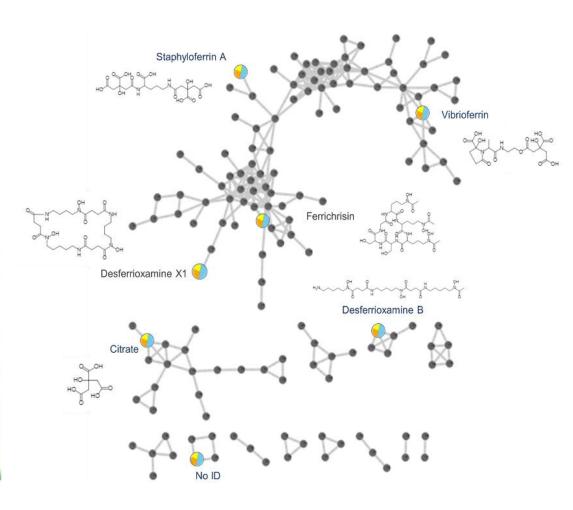
 Mixed Sgand

 Image Sgand





PopGenNP – computational?



Chemical Diversity