



# Future -omics developments

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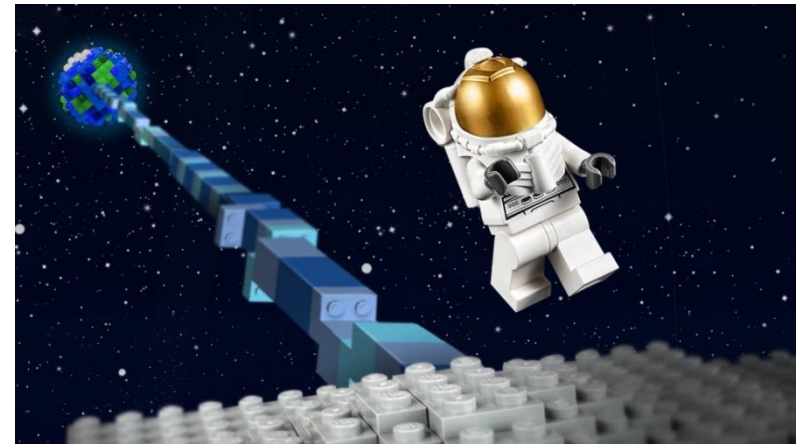
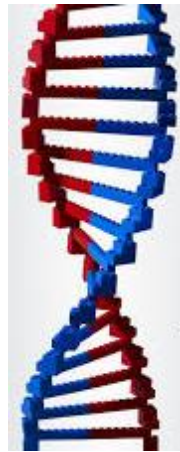
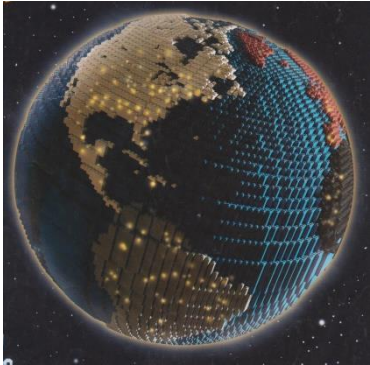
Department of Pharmaceutical Chemistry

Faculty of Pharmacy



HELLENIC REPUBLIC  
**National and Kapodistrian  
University of Athens**

## Where do we stand?



### Points to work

- An **aging population** mainly in Europe/USA - **HEALTH**
- Consumption continuously increasing/**waste production** - **ENVIRONMENTAL DEGRADATION**
- **Exhaustive use of natural sources** - **RESOURCES**
- Huge differences in **food access/poverty rates** –**LIMITED ACCESS**



## Where do we want to be?



Exploration of the marine environment

Support biomass production and processing

Contribute to product innovation and differentiation

Enabling technologies and infrastructure

Providing policy support and stimulation

- ✓ Described land species ~ 1.5 million (85%)
- ✓ Described marine species ~ 250,000
- ✓ There are more phyla and classes in the oceans (of 33 animal phyla, 21 exclusively marine, only 1 terrestrial)

## Biodiversity

<u>TYPE OF ECOSYSTEM</u>	<u>TYPE OF ORGANISM</u>	<u>TYPE OF ZONATION</u>	<u>TYPE OF TROPHIC RELATIONSHIP</u>
* <i>Deep sea</i>	* <i>Plankton</i>	* <i>Neritic</i>	* <i>Autotrophs</i>
* <i>Cold coral reefs</i>	* <i>Nekton</i>	* <i>Epipelagic</i>	* <i>Heterotrophs</i>
* <i>Seamounts</i>	* <i>Benthos</i>	* <i>Photic</i>	
* <i>Pelagic ecosystems</i>		* <i>Mesopelagic</i>	
* <i>Coastal waters</i>		* <i>Bathypelagic</i>	
* <i>Intertidal ecosystems</i>		* <i>Aphotic</i>	
* <i>Estuaries</i>		* <i>Abyssopelagic</i>	
		* <i>Hadal</i>	

**The exploration of marine ecosystems has only began in the mid 1970**



# OMICS TECHNOLOGIES

GENOMICS

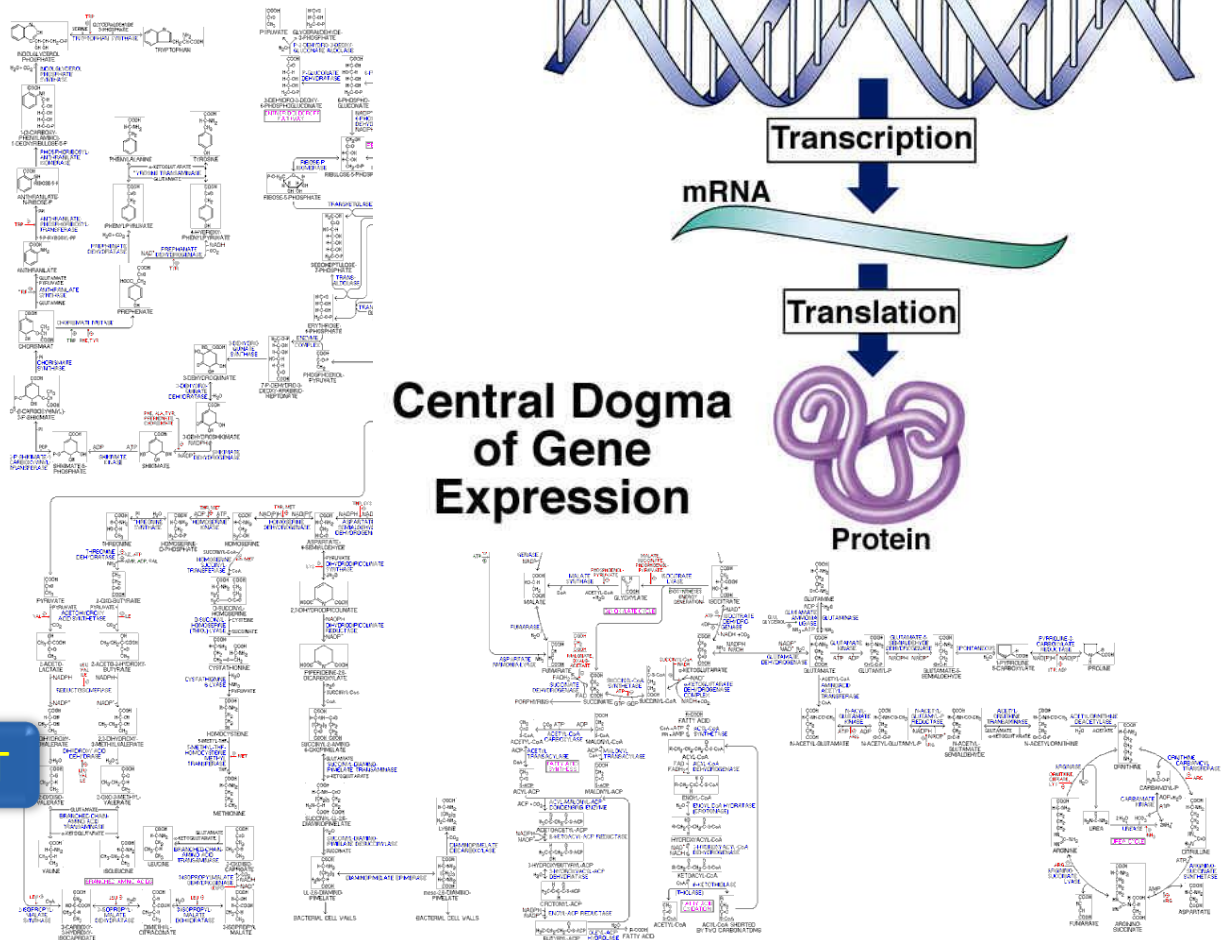
TRANSCRIPTOMICS

PROTEOMICS

METABOLOMICS

PHENOTYPE

ENVIRONMENT



## ➤ GENOMICS

The life aquatic: advances in marine vertebrate genomics

Joanna L. Kelley, Anthony P. Brown, Nina Overgaard Therkildsen & Andrew D. Foote

Affiliations | Corresponding author

*Nature Reviews Genetics* 17, 523–534 (2016) | doi:10.1038/nrg.2016.66

## METAGENOMICS

Marine genomics goes viral

Linda Koch

*Nature Reviews Genetics* (2016) | doi:10.1038/nrg.2016.130

Published online 03 October 2016

- **HABs** are present in nearly all aquatic environments as naturally occurring phenomena



### ARTICLE

Received 5 Sep 2013 | Accepted 14 Mar 2014 | Published 16 Apr 2014

DOI: 10.1038/ncomms4652

## Forensic genomics as a novel tool for identifying the causes of mass mortality events

Pierre De Wit<sup>1,2</sup>, Laura Rogers-Bennett<sup>3</sup>, Raphael M. Kudela<sup>4</sup> & Stephen R. Palumbi<sup>1</sup>

- More than 1/3 of Reef building corals threatened with extinction



## Genomic basis for coral resilience to climate change

Daniel J. Barshis<sup>1,2</sup>, Jason T. Ladner, Thomas A. Oliver, François O. Seneca, Nikki Traylor-Knowles, and Stephen R. Palumbi

Department of Biology, Hopkins Marine Station, Stanford University, Pacific Grove, CA 93950

*PNAS* | January 22, 2013 | vol. 110 | no. 4 | 1387–1392

## ➤ PROTEOMICS

- Existing methodological limitations
- Non-model organisms
- Lack of the proteomes of species commonly used in aquaculture

*Proteomics for the authentication of fish species*  
*Journal of Proteomics, Volume 147, 2016, 119–124*



*Mar. Drugs* **2014**, *12*(6), 3416-3448; doi:10.3390/md12063416

Open Access

Article

## Metabolomic Tools for Secondary Metabolite Discovery from Marine Microbial Symbionts

Lynsey Macintyre <sup>1,\*</sup>, Tong Zhang <sup>1</sup>, Christina Viegelmann <sup>1</sup>, Ignacio Juarez Martinez <sup>1</sup>, Cheng Cheng <sup>1,2</sup>, Catherine Dowdells <sup>1</sup>, Usama Ramadan Abdelmohsen <sup>2</sup>, Christine Gernert <sup>2</sup>, Ute Hentschel <sup>2</sup> and RuAngelie Edrada-Ebel <sup>1,\*</sup>

*Mar. Drugs* **2014**, *12*(6), 3681-3705; doi:10.3390/md12063681

Open Access

Article

## Accurate Dereplication of Bioactive Secondary Metabolites from Marine-Derived Fungi by UHPLC-DAD-QTOFMS and a MS/HRMS Library

Sara Kildgaard, Maria Mansson, Ina Dosen, Andreas Klitgaard, Jens C. Frisvad, Thomas O. Larsen and Kristian F. Nielsen \*

## Microbial Strain Prioritization Using Metabolomics Tools for the Discovery of Natural Products

Yanpeng Hou†, Doug R. Braunt†, Cole R. Michelt†, Jonathan L. Klassen†, Navid Adnani†, Thomas P. Wyche†, and Tim S. Bugni††

† Pharmaceutical Sciences Division, School of Pharmacy, University of Wisconsin, Madison, Wisconsin 53705, United States

‡ Department of Bacteriology, University of Wisconsin, Madison, Wisconsin 53705, United States

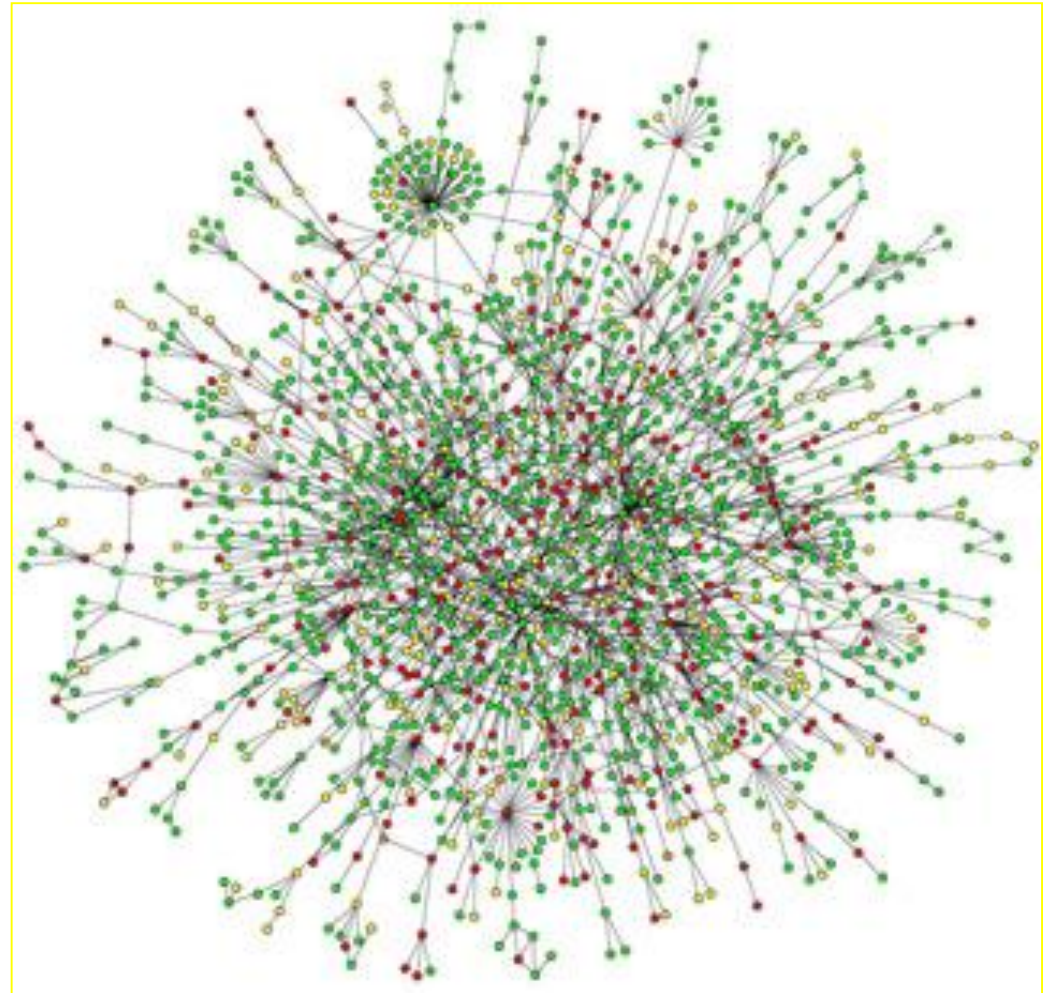
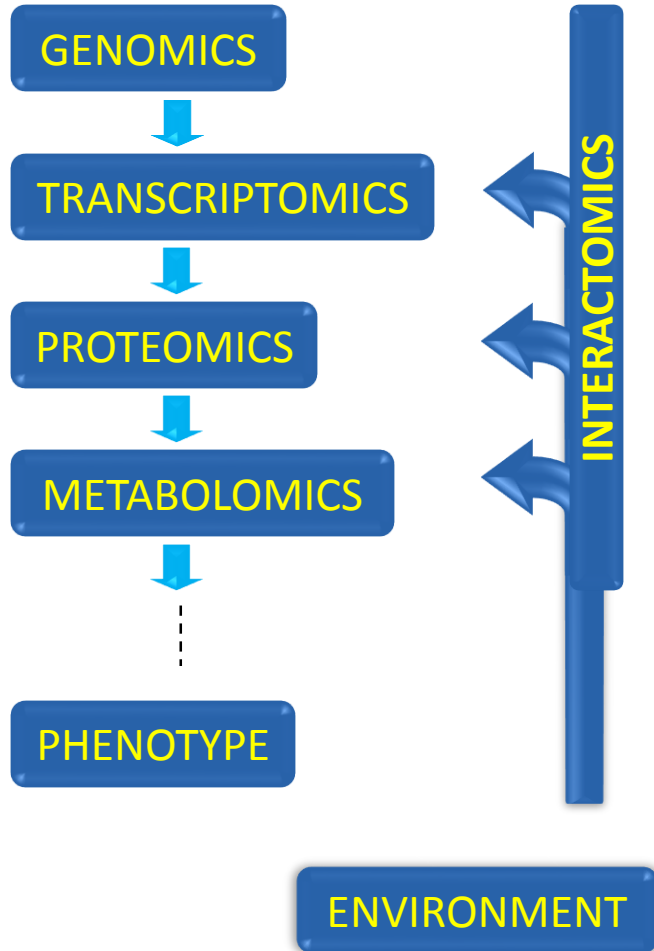
*Anal. Chem.*, **2012**, *84* (10), pp 4277–4283

DOI: 10.1021/ac202623g

Challenging issue for metabolomics analysis of marine organisms is the **high number of compounds** with unknown structures

# INTEGRATIVE OMICS

DNA- protein, protein-protein and protein-ligand interactions



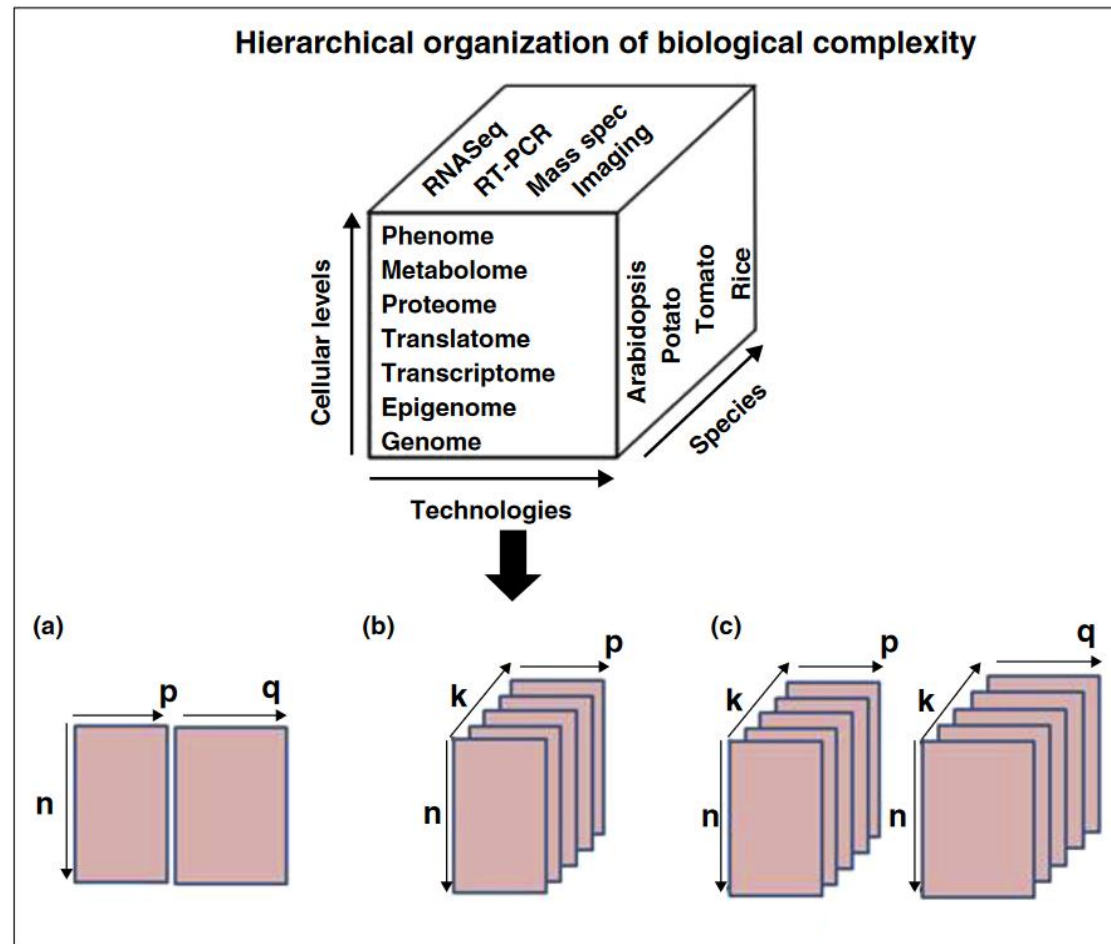
# 1+1>2

Systems biology is based on the understanding that the whole is greater than the sum of the parts

## Statistical Correlation of Heterogeneous Data

- Detection of hidden features
- Explore new dimension of data space

*PROPER DESIGN OF EXPERIMENTS*





**A success story of the correlation of heterogeneous data**



# DISCOVERY OF NOVEL NATURAL PRODUCTS

## Drug discovery pipeline:

### Drug R&D

### Clinical Trials

1-2 years

Target ID & Validation

**6 YEARS**

Optimization

1 year

Pre Clinical Animal Studies

20-100 people  
1-2 years

Phase 1: Safety

100-300 people  
1-2 years

Phase 2: Efficacy Safety

1,000-3,000 people  
2-3 years

Phase 3: Efficacy Safety

1-2 years

FDA Review & Approval

*Steve Blank, 2013, University of California*

Potency Studies  
Selectivity Studies  
PK/ADME-Tox properties  
SAR pharmacophore modeling

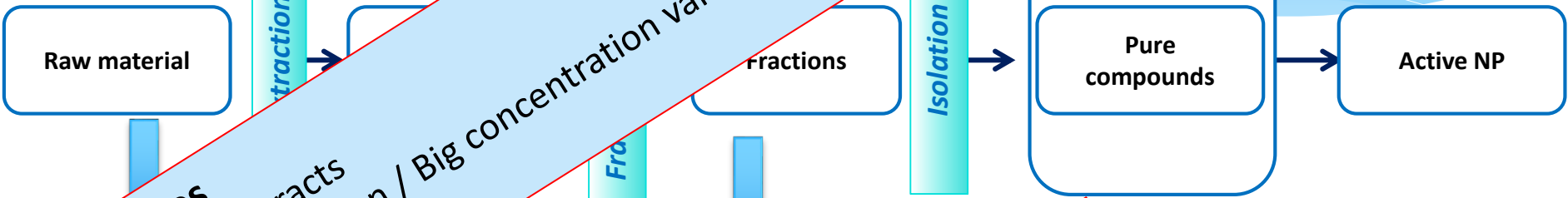
Pharmacological profile  
Administration route  
Drug interactions

t=0

- Pure Compounds
- Compound Libraries
- High Throughput Computer Automated Screening

Biological evaluation

Biological evaluation



**Challenges**

- Complex Extracts
- Low concentration / Big concentration variance
- Diverse Structures

**Advance Analytical Techniques**

- Isolation - Chromatographic
- Identification - Spectroscopic

**Classical pipeline pitfalls**

Laborious      Expensive

**KNOWN COMPOUNDS**



Extraction  
ASE, MVE, SFE

Extracts library

1<sup>st</sup> LEVEL  
Screening

Screening

HPLC - DAD / ELS  
& LC-MS  
Profiling

2<sup>nd</sup> LEVEL

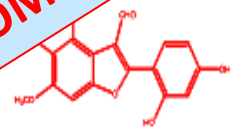
Separation  
Isolation  
LPLC, MPLC  
HPLC, FCPC  
ART

3<sup>rd</sup> LEVEL

UV and MS/MS  
libraries  
ChromQuest & MassFrontier

Identification  
(UV, IR)  
1 & 2D NMR  
400 & 600MHz

Microprobe 1 mm



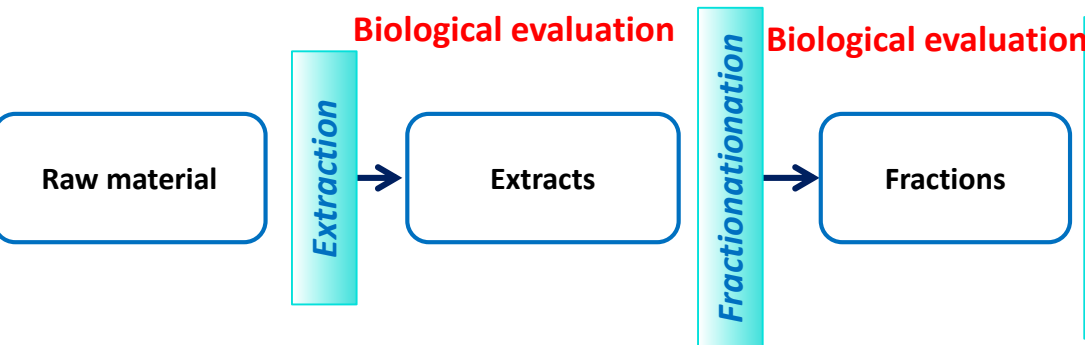
Dereplication methods have been emerged aiming to rationalize the entire discovery process

How **-omics** tools can accelerate our efforts from research bench to new products of natural origin?

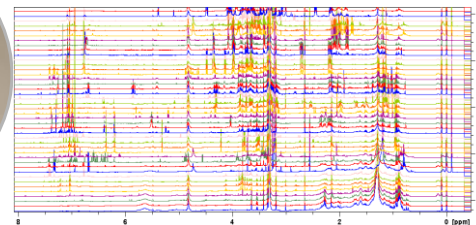
**Skip the classical pipeline**



*In vitro*  
evaluation



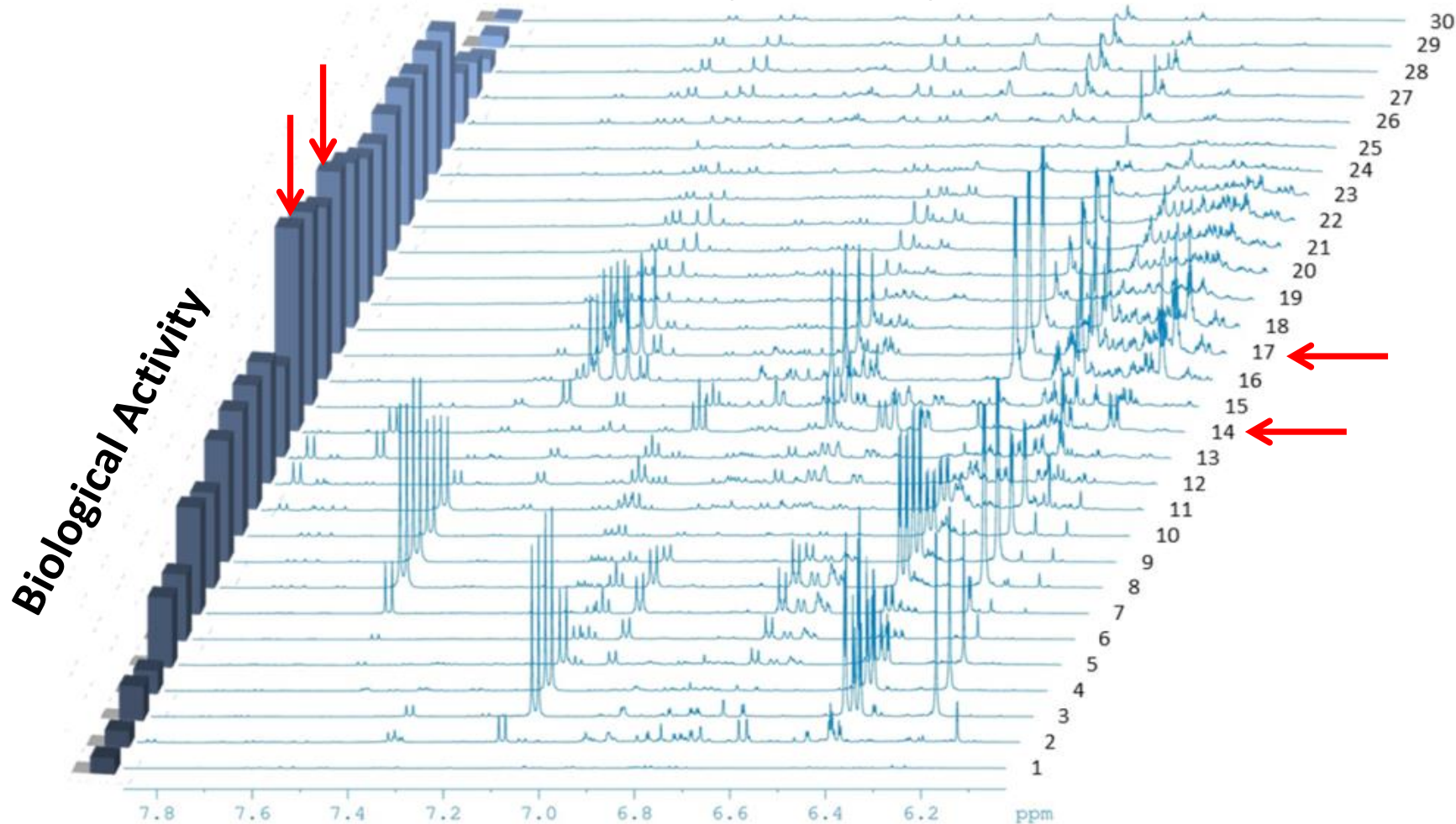
**NMR Profiling**



**HeteroCovariance  
Approach**

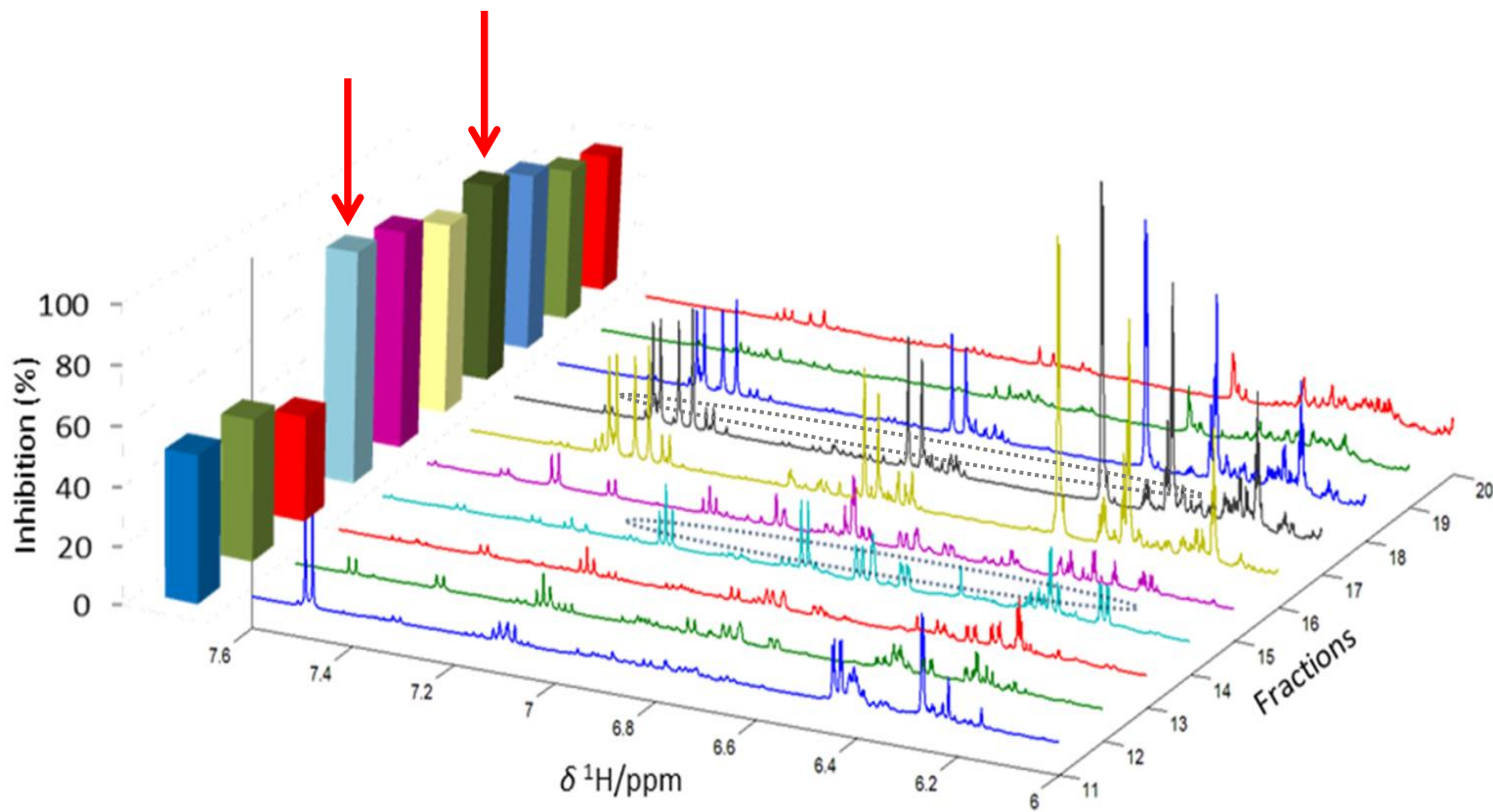


## Spectroscopic Features (NMR)

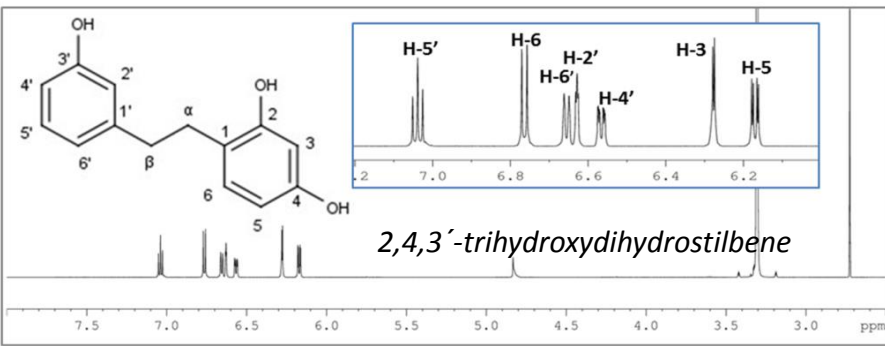




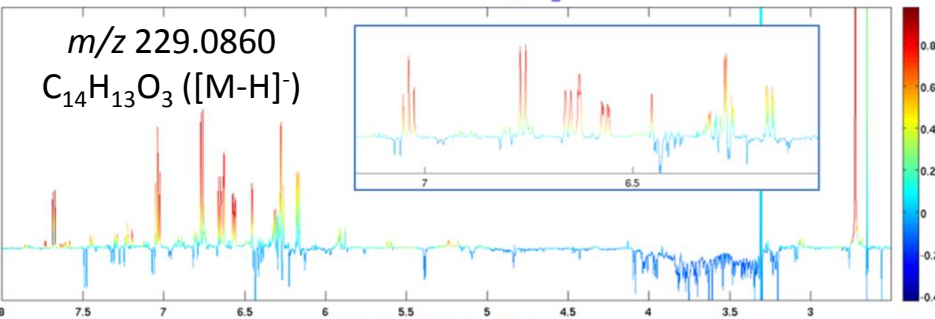
# HeteroCovariance Approach (HetCA)



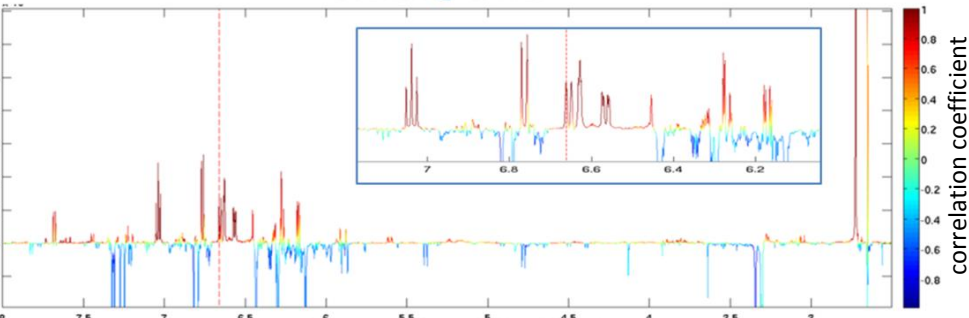
# HetCA TOOLBOX



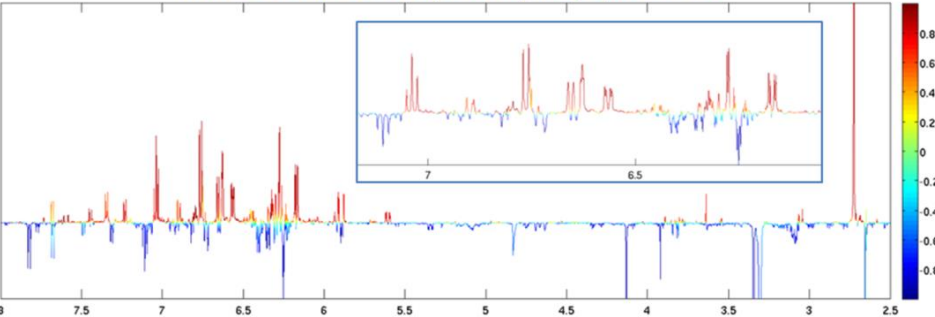
SHY @ 229.086\_7.91



STOCSY @ 6.6615



NMR covariance with activity fractions 12-16



- **Statistical Heterospectroscopy**  
(SHY, NMR – MS)

- **Statistical Total Correlation Spectroscopy**  
(STOCSY, NMR-NMR)

- **Hetero Covariance Approach**  
(HetCA, Spectr. Features – Activity)

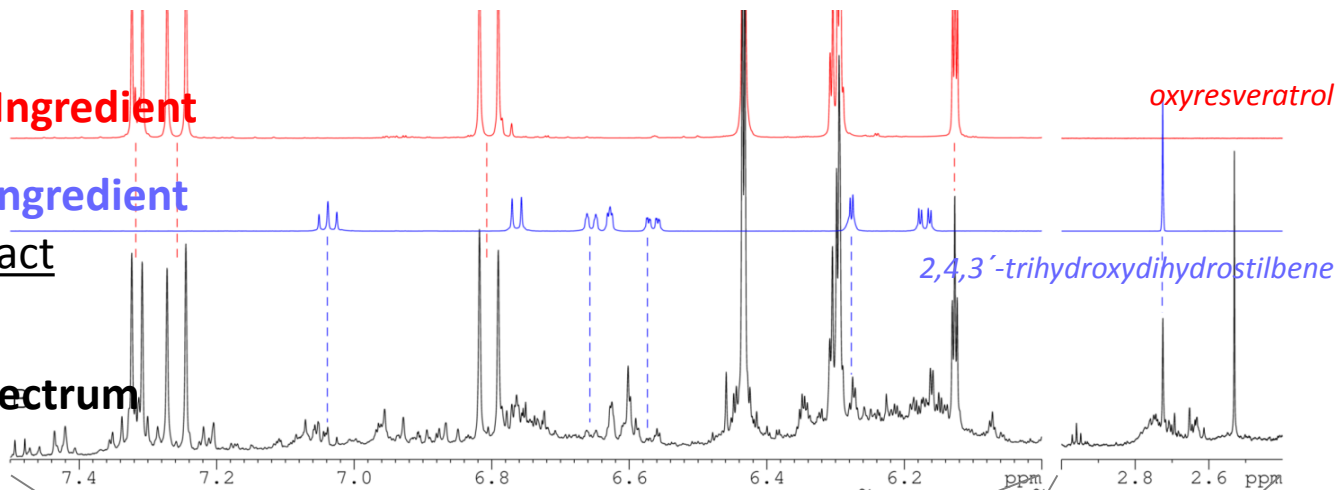
Increasing Correlation ↑

HetCA unmasks minor compounds with major activity from complex mixtures!

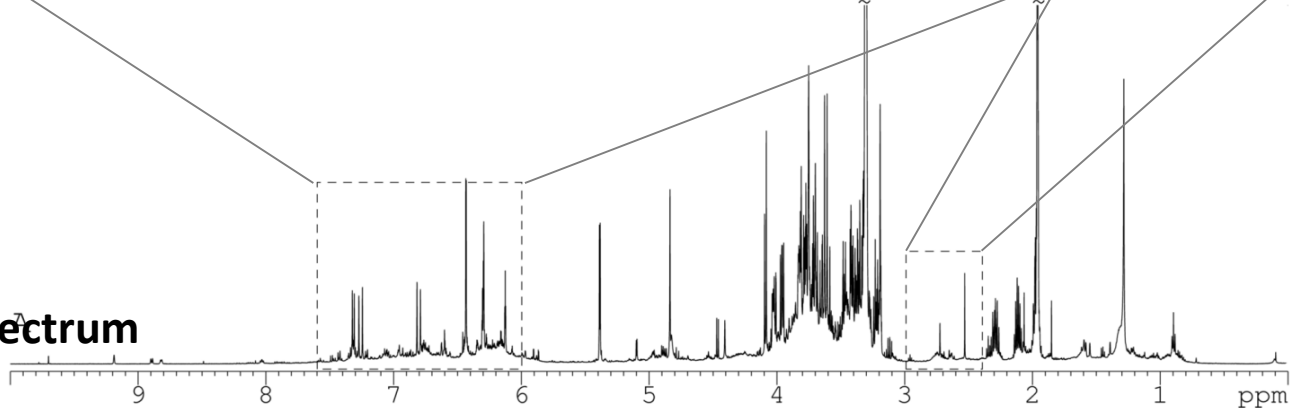
High Concentration **Active Ingredient**

Low Concentration **Active Ingredient**  
<1% of the total extract

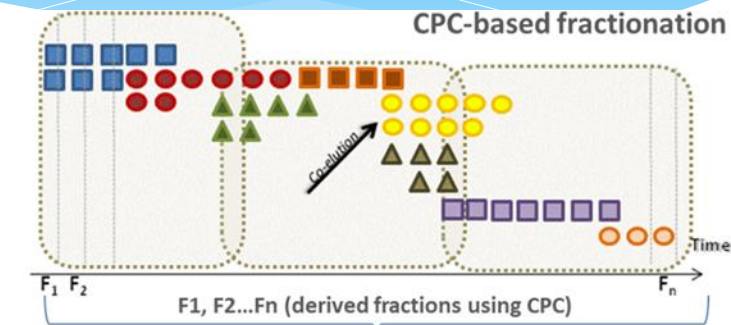
total extract 1D <sup>1</sup>H NMR spectrum



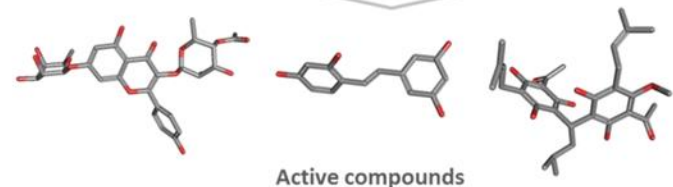
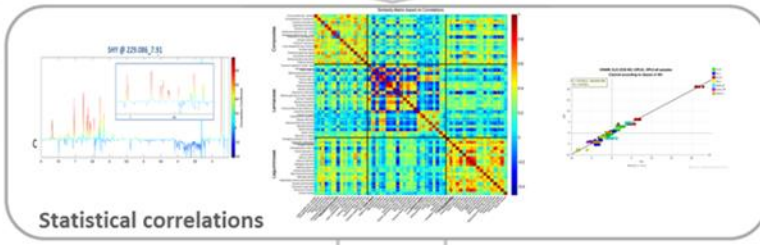
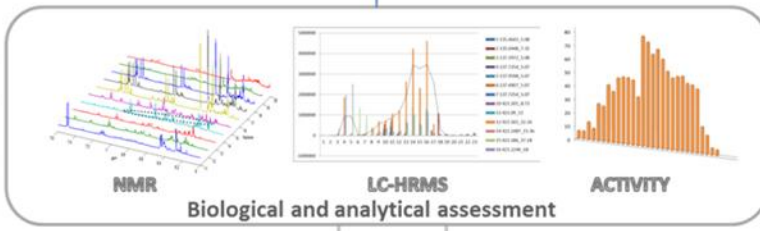
total extract 1D <sup>1</sup>H NMR spectrum



- Create an Artificial Variance
- Statistical correlation of heterogeneous data
- The **same spectral data** can be utilized for screening against any **possible pharmacological target** or any other **dose dependent property** on the basis of the HeteroCovariance Approach
- This highly innovative activity-based-metabolite-profiling approach can dramatically **accelerate the discovery of active natural products** challenging global biodiversity and chemodiversity



HETEROVARIANCE APPROACH





# Tascmar

## Tools And Strategies to access original bioactive compounds from Cultivation of **MAR**ine invertebrates and associated symbionts

Coordinator; CNRS

Budget: 6.7 Million euros

13 partners (100 researchers), among which 6 industrials

Project duration: 4 years starting the 1<sup>st</sup> April 2015

Participating countries: France, Greece, Israel, Germany, Italy, Spain, Sweden, Thailand



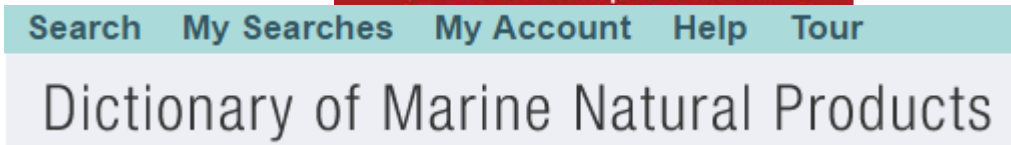
# AVAILABLE TOOLS

## ✓ We have the tools

- Advance analytical techniques (NMR, uHPLC-HRMS, bioautographic HPTLC, etc)

- **Compound Libraries**

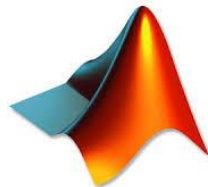
- Lack of NMR spectra
- MS annotations
- Non user-friendly



- Bioinformatics - “Big Data Era”



GitHub



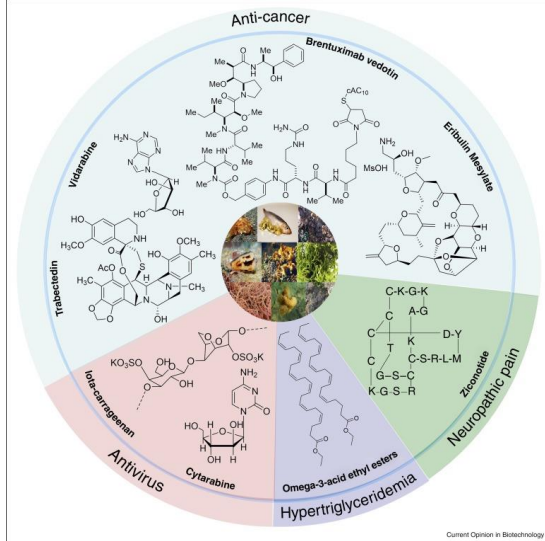
Exploration and  
Integration of  
Omics datasets





## FUTURE PROPOSAL

Current Opinion in Biotechnology, Volume 42, 2016, 13–23

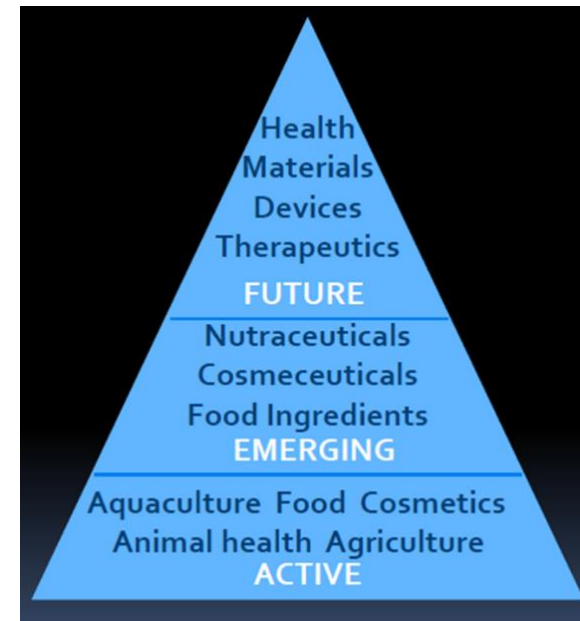


- Only **8 Drugs of marine origin** have been approved by FDA and EMA in **ca. 40 years**

✓ Smart connection of Heterogeneous Data

✓ Interdisciplinary Collaborations

✓ Fast Productive Solutions



MarineBiotech



**Thank you for your attention**