

Generation and Exploitation of Marine Natural Product Extract Collections as Source of New Bioactive Molecules

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November 2018

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10YEARS

FUNDACIÓN CENTRO DE EXCELENCIA EN INVESTIGACIÓN DE MEDICAMENTOS INNOVADORES EN ANDALUCÍA





 Introduction of Fundación MEDINA and our collection

Creation of Natural Product Extracts
 Collections

• Exploitation of collections in search of new bioactive molecules



ABOUT US



Independent, Non-profit Research Organization established as a Private-public partnership between:

- Government of Andalucía (Spain)
- University of Granada (Spain)
- Merck Sharp and Dohme de España S.A.







Our Mission

Discovery of new bioactive compounds and innovative therapies for unmet medical and industrial needs.

HEALTH SCIENCES TECHNOLOGY PARK, GRANADA, SPAIN









Faculty of Health Sciences

Faculty of Medicine

CSIC- I. Lopez Neyra

UGR-I.Biomedicine

Bio Incubator

General Hospital

Genyo







- Multidisciplinary team of scientists (28 permanent staff, up to 40 employees)
- 2300 m² fully equipped facilities
- One of the largest microbial collections and libraries for natural products drug discovery
- High capacity center for high throughput screening
- Drug discovery research programs in infectious diseases, cancer and neurodegeneration



MEDINA at a glance





Contribution of CIBE-MRL Spain to Merck Natural Product Drug Discovery





MICROBIAL NATURAL PRODUCTS

Maximizing chemical diversity by maximizing biodiversity







100IN.

Microbial Strain Collections 190,000 strains*

Fungi, Actinomycetes & unicellular Bacteria

Natural Products Libraries
 >200,000 extracts

from a large diversity of well-identified microbial strains

Proven genetic and chemical diversity

Geographical, Ecological & Taxonomical Diversity

* Recent donation of new 74,000 strains from former Cubist-Merck & Co. Inc Collection



LARGEST MICROBIAL COLLECTION WORLDWIDE WITH 190.000 STRAINS







Fungal plant endophytes: 32% Soil & plant rhizosphere actinomycetes: 86% Marine actinomycetes: < 10%







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Types of HTS Libraries

Extracts/Fractions NP Libraries

 Unknown Composition/

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- Potentially new chemistry
- Unknown
 concentration [WBE]
- vehicle DMSO/AQ
- Dereplication required LC-MS/NMR
- Hit follow-up required for identification

Purified Natural Products Libraries

 Known/unknown Composition

- Know/unknown concentration [1 - 5 mg/mL]
- vehicle DMSO
- Three freeze-thaw cycles maximum
- Degradation analyses required LC-HRMS

Synthetics/Combin atorial Libraries

- Known Composition
- Known concentration [10 mM]
- Vehicle DMSO
- Three freeze-thaw
 cycles maximum
- Recurrent Degradation analyses required LC-HRMS (QC)



Collections of extracts-General Rules

- Cover a wide and representative range of biological species
- Created using robust and reproducible protocols

- Must represent the most significant biodiversity (ideally all) included in the microbial strains or macroorganism sample collection
- Ideally a back-up sample or a protocol to generate it must be available for scaling-up purposes
- Extracts must be **stored in appropriate conditions** and in a format suitable for the rapid generation of multiple aliquots
- Associated information managed through a Laboratory Information Management System (LIMS)
- Samples included in the collection **must have been legally collected** in compliance with Nagoya protocol and other national and international legislation



Expeditions, Samples and Extract Collections Pharma Mar 200.000 extracts!!!

CLASIFICATION

COLLECTION

EXPEDITIONS

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Harvesting and Logistics





Taxonomy & Labelling



GPS, Images

DATA MANAGEMENT





Storage, Safety



Preparation of extracts

EXTRACTION Water/MeOH:DCM

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1/



MARINE BIODISCOVERY IN IRELAND



FUNCTIONING OF THE NMBLI



Voucher

SOP MB-104

Marine

Biodiscovery











Microbiological Approaches to Maximize Chemical Diversity



- Culture conditions (OSMAC)
- Stress factors

- Addition chemical inducers
- Interspecies crosstalk: Co-culturing
- Epigenetic modifiers

- Genes encoding biosynthetic enzymes in bacteria and fungi outnumber known secondary metabolites
- We are missing large part of NP biosynthetic capacity
- Only a subset of pathways is expressed in laboratory conditions
- Access to this reservoir of molecules needs activation of cryptic genes









"The research leading to these results has received funding from the European Union's Seventh Framework Programme (FP7/2007-2013 under grant agreement n $^{\circ}$ 312184)"



Geographical Origin of MEDINA's Marine Strains



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Taxonomic Diversity of the Marine Collection



Flavobacteriaceae Pseudomonadaceae Alteromonadaceae Pseudoalteramanadaceae Vibrionaceae Erythrobacteraceae Sphingomonadaceae Caulobacteraceae Phyllobacteriaceae Rhadobacteraceae Streptomycetoceae Nocardiaceae Mycobacteriaceae Dietziaceae Nocardioidaceae Micromonosporaceae Streptosporangiaceae Nocardiopsaceae Micrococcacege Promicromonosporaceae Intrasporangiaceae Microbacteriaceae

lacillaceae

Bacilli Flavobacteria Gammaproteobacteria Alphaproteobacteria Actinobacteria High taxonomic diversity within the collection, which includes obligate marine bacteria such as *Salinispora* spp.

over **5,500 marine bacteria**, mainly actinobacteria (~75%), non-actinobacteria taxa (~25%).

Diversity tree: partial 16S rRNA sequences of 1,600 strains



PharmaSea: Selection of Strains

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Strains were selected in order to maximize the diversity in geographic origin, taxonomy and isolation source



Fermentation of each strain in 4 different media

Components (in g/L)	DEF15S-M	APM9-M R358		FPY12-M
Amicase				5
Bacto Peptone			2	5
Bacto Yeast Extract			4	
Fructose				20
Glucose		50		10
Maltose				10
Saccharose	5			
Soluble Starch (potato)	20	12	10	
Soybean Flour		30		
CoCl2.6H2O		2 mg		
FeSO4.7H2O			5 mL	
			(stock 8 g/L)	
KBr			5 mL	
			(stock 8 g/L)	
КН2РО4	1			
MgCl2.6H2O	1			
Na2SO4	2			
NaCl	1			
NH4Cl	2			
CaCO ₃	2	7		
Sigma Sea Salts	40	40	40	40
Trace Elements	1 mL			1 mL
	7.0 (before adding	7.0 (before adding	7	7
рп	CaCO ₃)	CaCO ₃)		





harma Sea Collection generated at MEDINA (6K)

Module	Type of Sample	# Samples	
PCA-000001 to PCA-000022	Crudes + blanks	1696	
PFA-000001 to PFA-000054	Fractions + blanks	4320	





Storage - Labeling



And the second second

LIMS Tracking of Samples: NAPIS





Medina_Carambolo - [Hierarchy - CA-253965-a01-MR016-EC01-FS004-FH175-FH001]

View Window Help



CA-253963	Pro	ompt Notes Report	Formulation	
E CA-253964 Strains CA-253965	Th	e following fields were	defined by the user during login.	
G CA-253965-a01		Field	Value	
🚋 📜 CA-253965-a01-MO001		Platanama		
🕀 <u>]</u> CA-253965-a01-MO002		Flatename	FL-HFF-000655	
CA-253965-a01-MO003 Small scale		PI Order	24	
⊕] CA-253965-a01-M0004		PIBow	8	
CA-253965-a01-MO005 Fermentations				
		Crow	H	
H JI CA-253965-a01-MO007		PI Column	3	
H I CA-253965-a01-M0008		Sampleid	PL-HPE-000659-H3	
Large scale				
I CA-253965-a01-MR011 Eormontations		Identifyer	(None)	
CA-253965-a01-M5012		Parent Name	CA-253965-a01-MR016-EC01-FS004-FH	
🕀 <u>]</u> CA-253965-a01-MR013		External Id	(None)	
⊡ 📃 CA-253965-a01-MR014		Excinding	(None)	
		Clients	(None)	
		Requesters	(None)	
GA-253965-a01-MR016-EC01-F5001 ■ CA-253965-a01-MR016-EC01-F5001		Category	SAMPLE	
CA-253965-a01-MR016-EC01-F5002				
🕀 🔟 CA-253965-a01-MR016-EC01-F5003		IDENTIFICATION	CA-20006-801-MR016-EC01-F0004-FH	
🖨 📜 CA-253965-a01-MR016-EC01-F5004		medAliquot Type	aF	
CA-253965-a01-MR016-EC01-F5004-a01 Assay Aliquots		Extraction Type	FBACTION	
CA-253965-a01-MR016-EC01-F5004-a03		Extraction Solvent	ACN/AQ	
		CTS_ID	(None)	
			(None)	
CA-253965-a01-MR016-EC01-E5004-EH004				
			ME	



Outline

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Antitumor Screening - Cytotoxicity



Human tumour



Cell culture



Inmortalized cell lines



In vitro cell culture





Inactive

Cytotoxic Control Sample preparation



Cell seeding



....

Quantification & Evaluation

Incubation



Cell treatment





ET-743, Trabectedin, Yondelis



Ecteinascidia turbinata



Pharma Mar

Approved for the treatment of Soft Tissue Sarcoma and Ovarian Cancer PM0184







Undergoing Phase II clinical trials against colorectal cancer as single agent

Lithoplocamia lithistoides

harma Sea Collection generated at MEDINA (6K)

Module	Type of Sample	# Samples	
PCA-000001 to PCA-000022	Crudes + blanks	1696	
PFA-000001 to PFA-000054	Fractions + blanks	4320	





MEDINA - PharmaSea Antimicrobial screening

Antibacterial

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- ✓ Staphylococcus aureus MRSA
- ✓ Pseudomonas aeruginosa
- ✓ Escherichia coli
- β-lactam synergy vs. multi-resistant Pseudomonas aeruginosa

Antifungal

✓ Candida albicans
 ✓ Aspergillus fumigatus

Anti-TB

✓ Mycobacterium tuberculosis H37Ra
 ✓ Mycobacterium bovis BCG







Zebrafish-based identification of neuroactive and antiepileptic hits (KU Leuven)



- Photomotor response assay: neuroactive hits
- Epilepsy seizure model: anticonvulsant hits
- Maximum Tolerated Concentration (MTC) analysis: general toxicity analysed for hits

3777 screened samples
727 neuroactive hits
43 anticonvulsant hits
5 out of 11 hits validated
→ To be continued

PHARMASEA

Screening Campaigns Results

Count of Hits	Qualifier			Extracts	
per Assay	Α	Q	I	Tested	
Absorbance_MB2884 (E.coli)	68	137	5761	5966	
Absorbance_MB5393 (MRSA)	114	10	5842	5966	
Resazurine_A.fum_ATCC46645	196	30	5740	5966	
Absorbance_C.alb_MY1055	142	62	3316	3520	
Resazurine_C.alb_MY1055	73	35	2338	2446	
Resazurine_P.aer_PAO1 + Imipenem	0	0	2480	2480	
Resazurine_A.bau_MB5973	0	0	1840	1840	
Resazurine_A.bau_MB5973 + Imipenem	25	28	3673	3726	
Fluorescence_M.tub_H37Ra	201	44	5721	5966	
Grand Total	843	374	41025	42242	
	12	17	_		

PharmaSea Drug Discovery Campaign

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6016 samples (blanks+extracts+fractions) from 405 strains

1217 Hits (843 Active, 374 Questionable) selected for CherryPiking 416 confirmed Hits selected for LCMS Dereplication of knowns

163 samples with no matches in LCMS DB

71 strain-condition-assay selected for confirmation

71 ferment + extraction + assay confirmation

HPLC fractionation+assay+ LCMS and NMR dereplication

> 20 Strains selected for Big Scale Purification

52 Compounds





Streptomyces zhaozhouensis CA-185989



Biological activity	MIC [µg/mL]				
Compound ID	MRSA	C. albicans	A. fumigatus		
Isoikarugamycin	2-4	2-4	4-8		
Ikarugamycin	2-4	4	4-8		
28-N-methyllkarugamycin	1-2	4	4-8		
30-oxo-28-N-methyllkarugamycin	32-64	>64	>64		

R₁= H $R_2=H$ Ikarugamycin

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- $R_1 = CH_3 R_2 = H$ 28-N-methyllkarugamycin $R_1 = CH_3$ $R_2 = O$
- New
- 30-oxo-28-N-methyllkarugamycin New

Mar. Drugs, 2015, 13, 128-140.



New antifungal macrolides

Streptomyces caniferus CA-271066





R ₁ = -OH	R ₂ = -H	$R_3 = -H$	Compound 1
R ₁ = -OH	R ₂ = -OH	$R_3 = -H$	GT-35
R ₁ = -H	R ₂ = -OH	$R_3 = -SO_3H$	Compound 2
R ₁ = -OH	R ₂ = -OH	$R_2 = -SO_3H$	Compound 3

			MIC values (µg/mL)			
Fungal strains	Strain number	Compound 1	Compound 2	Compound 3	Compound 4	
C. albicans	ATCC64124	0.5-1	2	0.5-1	8	
C. tropicalis	ATCC750	0.5-1	2	0.5-1	8	
C. glabrata	ATCC90030	0.5-1	2	0.5-1	8	
C. krusei	ATCC6258	0.5-1	1	0.5-1	8	
C. parapsilosis	ATCC22019	0.5-1	2	0.5-1	16	
A. fumigatus	ATCC46645	4	4	4	8	



Full absolute configuration of macrolides



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Bioinformatic prediction



Bioinformatic + NMR prediction

Post-Tailoring NMR predicted

Bioinformatic prediction

DISCREPANCY

New spirotetronate derivatives. Phocoenamicins

Micromonospora chaiyaphumensis CA-214671



compounds	MIC (mg/mL)				ZOI* mm(mg)
	S. Aureus	M. tuberculosis	M. bovis	E. faecium	B. subtilis
	MB5393	ATCC 25177	ATCC 35734	MB5571	MB964
phocoenamicin B	8-16	>128	>128	>128	7 (2)
phocoenamicin C	32-64	32	>128	>128	7 (4)
phocoenamicin	4-8	16-32	>128	32-64	7 (4)
vancomycin	2-4			>128	
streptomycin		1.6-3.2	0.4-0.8		
gentamicin					8 (0.25)
penicillin G					19 (0.06)

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Mar. Drugs, 2018, 16, 95.





PRIORITY LIST AND BIOACTIVE COMPOUNDS



Antibacterial activity





Halidrys siliquosa

New bioactive meroterpenoid derivatives

New C-glycosilated angucycline derivatives

Streptomyces sp. CA-237351

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Neuropharmacology, 2018, 141, 283-295



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