



# QUANTIFYING LIFE WITH DIGITAL BIOLOGY

- Biotechnology company with the mission to embed DNA technologies in business operations.
- We provide end-to-end, personalised solutions to our customers using cutting-edge technologies.
- We use innovative technologies to sequence genetic material and we analyse complex data using Bioinformatics and Artificial Intelligence methods.
- We employ and collaborate with scientists world-renowned in their field. Thus, we have the possibility to provide consulting services, add to the theoretical background of the problem and co-shape with our clients their personalised solution.

#### **Areas of Expertise**

#### **Metagenomics of Built Environments**







- Detection and monitoring of all microorganisms found in a built environment.
- The sampling method is non-invasive and can be applied to air, liquids and surfaces.
- Huge range of applications such as hospitals, public buildings, means of transport, livestock units, water supply systems, sewage, archaeological excavation sites and works of art.

#### **Agrigenomics**





- Genetic identification of ingredients in organic samples such as food and soil samples, using state-of-the-art technologies (Next Generation Sequencing, microarrays).
- Applications in agriculture, animal husbandry and across the wide range of food industry.

#### The scientific team - MBE



**CSO (Chief Scientific Officer)** 

Prof. Christos Ouzounis is the CSO and cofounder of the company, and a Prof. in Bioinformatics at AUTH. He holds a B.Sc. in Biological Sciences (AUTH), an M.Sc. in Biological Computation and a DPhil in Computational Chemistry and Structural Biology, both from the University of York. He has lead core bioinformatic facilities at EMBL-EBI, King's College London and CERTH in Thessaloniki.



**CTO (Chief Technology Officer)** 

Christos Karapiperis is the CTO and co-founder of the company. He has a Bachelor's degree in Computer Science, an MBA degree from the University of Sheffield and is a PhD Candidate in Bioinformatics at AUTH. He has over 20 years of experience in IT and Bioinformatics projects.



**Bioinformatics Expert** 

Dr. Anastasia Chasapi holds a
Bachelor's degree in Biological
Sciences (AUTH), an M.Sc. in
Proteomics and Bioinformatics from
the University of Geneva and a PhD
in Computational Biology from the
University of Lausanne. She has
worked for years at the Swiss
Institute of Bioinformatics as a
collaborating scientist. Currently she
holds a Research position at CERTH.

#### The scientific team - Agrigenomics



Agrigenomics Team Leader
Dr. Zoe Chilioti is an agronomist, she holds an M.Sc. in Greenhouses / Protected Crops and a PhD in Molecular Plant Biology. She has many years of experience as a collaborating scientist at Johns Hopkins University and the NIH in USA. Currently she is leading a research group at CERTH in Thessaloniki.



Agrigenomics Expert

Dimitrios Valasiadis is an agronomist,
he holds an M.Sc. in Genetics & Plant
Breeding and is a PhD candidate in
Systems Biology & Post-Harvest
Physiology. Currently, he is an active
researcher in the field of Applied Life
Sciences.



Athanasia Alexandridou recently obtained a Bachelor's degree in Biological Sciences from the University of Oklahoma, USA. She is currently conducting her first research steps in Applied Life Sciences.

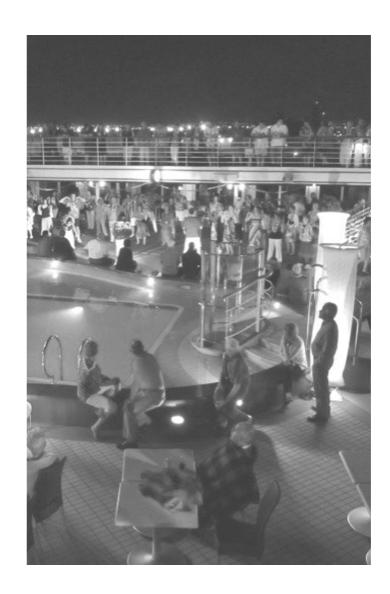
#### **Metagenomics in Biosafety**

- Detection and monitoring of infectious agents in key areas of human activity (eg hospitals, schools, water supply system) is essential for effective prevention and control of health risks.
- MBE technology allows the exhaustive identification of all microbes in a sample and constitutes a universal solution for the identification of pathogenic microbial communities in an environment, without the need for special tests for expected or well-characterised pathogens.
- Our advanced bioinformatics platform permits the identification of biological risk factors and helps us set priorities and provide guidelines for the prevention of public health issues.



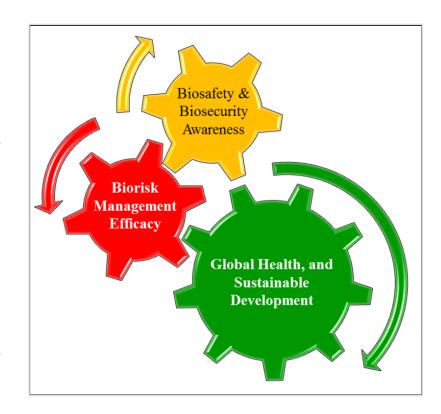
#### The problem

- Cruise ships are intense mobility hubs among people from all over the world, who can carry infectious agents.
- The problem is exacerbated as microbial diversity varies around the world. As a result, there is risk of exposing customers and staff to unknown pathogens with increased infectivity.
- The recent SARS-Cov2 pandemic has created a climate of insecurity towards the use of leisure services, especially when it involves using indoor public spaces.
- At the same time, citizens are now more aware and better informed about biosafety issues, and therefore more demanding from services provided to them.



#### **Proposed solution**

- Our main goal is to detect and monitor microbial communities located in public and non-public areas and to record hotspot areas and transmission patterns.
- By constructing almost complete genomes from metagenomic samples, we are able to exhaustively identify all potential biological hazards as well as monitor their spread in cruise ships.
- Monitoring pathogens in the environment in almost real time allows targeted and effective infection control measures.
- The detection and monitoring of pathogens in indoor air, indoor and outdoor surfaces as well as in the water supply and produced wastewater provides a comprehensive record of the ship's epidemiological state.



### **Air - Water - Surface Monitoring**









**Cruiser rooms** 

**Indoor common areas** 

lounge, restaurants elevators, bathrooms casino, gym

Outdoor common areas

swimming pool water Swimming pool area decks

**Production facilities** 

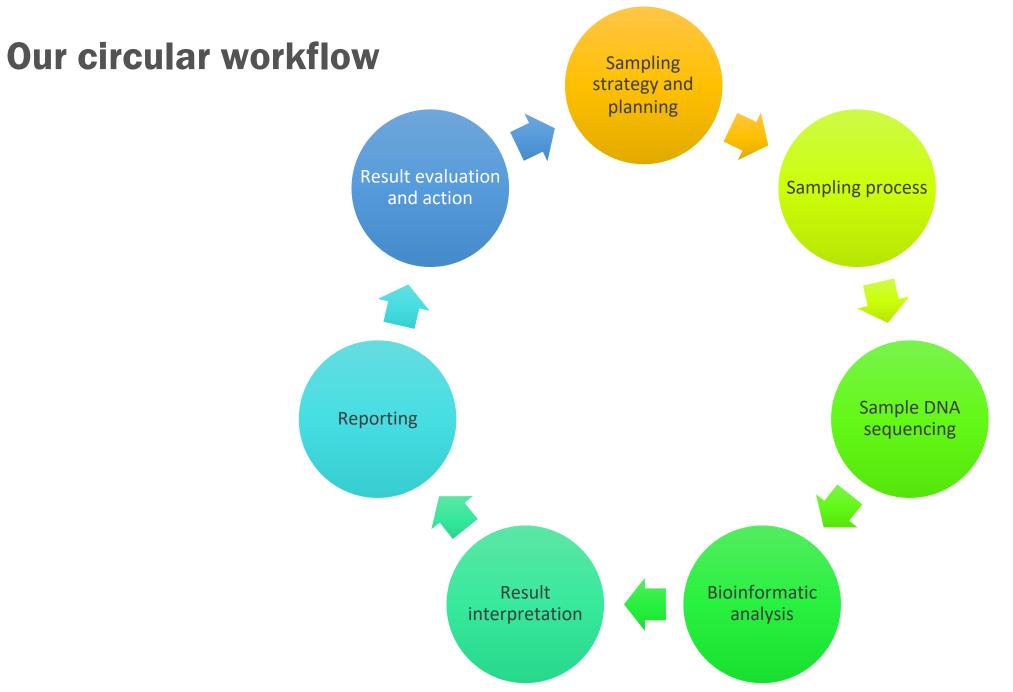
food preparation area washing rooms

Water supply and wastewater

### **Benefits for the Luxury Cruise Ship**

- Identifying and monitoring the cruiser's microbial footprint will create a safe and trustworthy environment both for the staff and customers.
- The cruiser administration will have imminent information of possible health threats and will be able to take the necessary measures.
- The interactive result reports generated by the DNA sequence platform can be used for controls and to inform interested third parties.





## **Comparison of Existing Approaches**

Criteria	Classic Microbiology	Molecular Analysis (PCR)	Metagenomic Analysis
Sample preparation difficulty	**	*	
Possibility for automated sample collection			***
Technical skill required for analysis	*	**	*
Time required for analysis	***	**	*
Result precision and accuracy		*	***
Amount of microbes that can be identified	*	*	***
Equipment & Hardware Cost	*	**	***

#### **Dedicated Biosafety Platform**

- Our Biosafety platform includes three main axes of analysis:
  - 1. Sample sequencing
  - 2. Bioinformatic analysis and result interpretation
  - 3. Statistical analysis and presentation of the results
- For the Biosafety platform hosting, we encourage the development of local infrastructures in the cruiser, which will give full control and immediate access of all produced data to the cruiser direction.
- The dedicated Biosafety Platform will consist of:
  - 1. A Data Center, equipped with sufficient computing power and storage space, holding all produced data
  - 2. Network and Data Security structures, such as back up devices
  - **3.** A Crisis Management Room, with teleconferencing capabilities.
- The above can be achieved through the collaboration with a contractor with expertise in the field.



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