

Technical Presentations Report

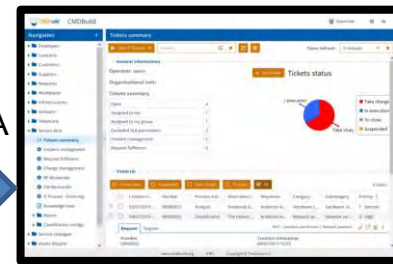
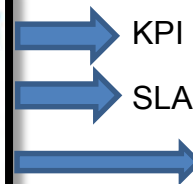


Antonio José Sáenz-Albanés

ICT Core e-Infrastructure Operations Coordinator

Infrastructure offered by LifeWatch ERIC

- Distributed e-Infrastructure
- Data processing and authoring tools
- VREs, Workflows, LifeBlock
- Availability
 - Redundancy, monitoring, SOC
 - KPIs -> SLAs
- FitSM

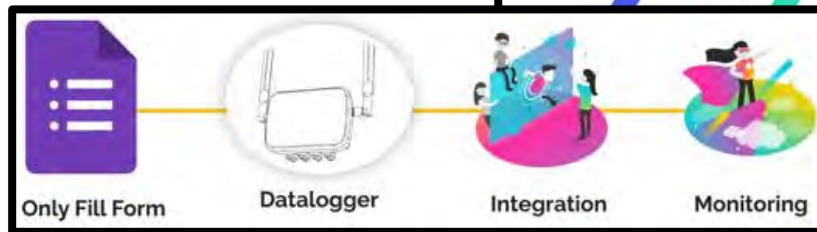
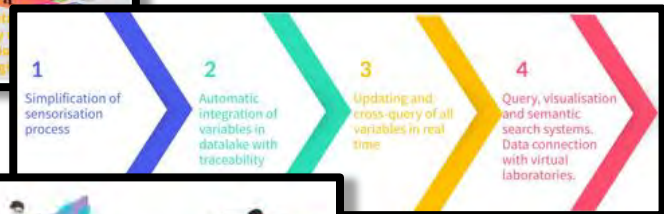
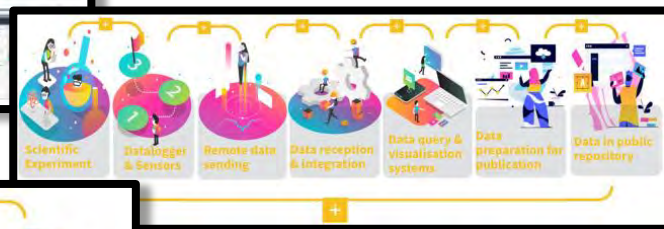
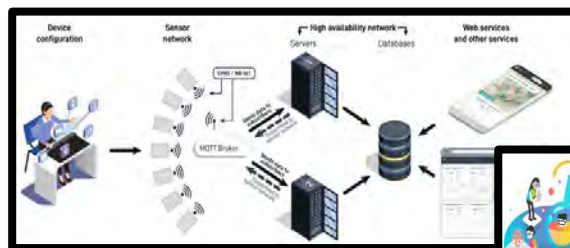


Improving the environmental monitoring cycle, remote sensing & space technologies

Pablo David Guerrero-Alonso

Data capture process

- Traditional IoT environmental monitoring cycle
- Improving the environmental monitoring cycle
 - Simplifying sensorisation process
 - Automatic integration in DB & traceability
 - Real-time updating & queries
 - Exploration through vLabs
- Enhanced Dataloggers

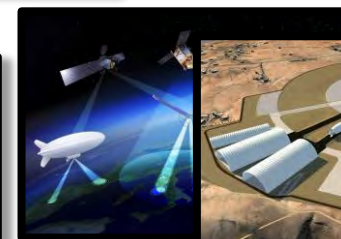
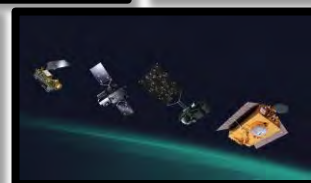
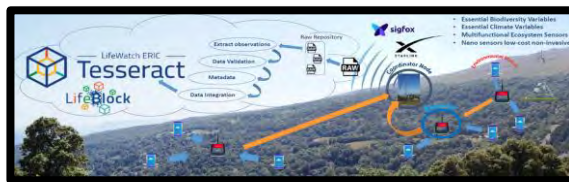


Improving the environmental monitoring cycle, remote sensing & space technologies

Jaime Lobo Domínguez-Roqueta

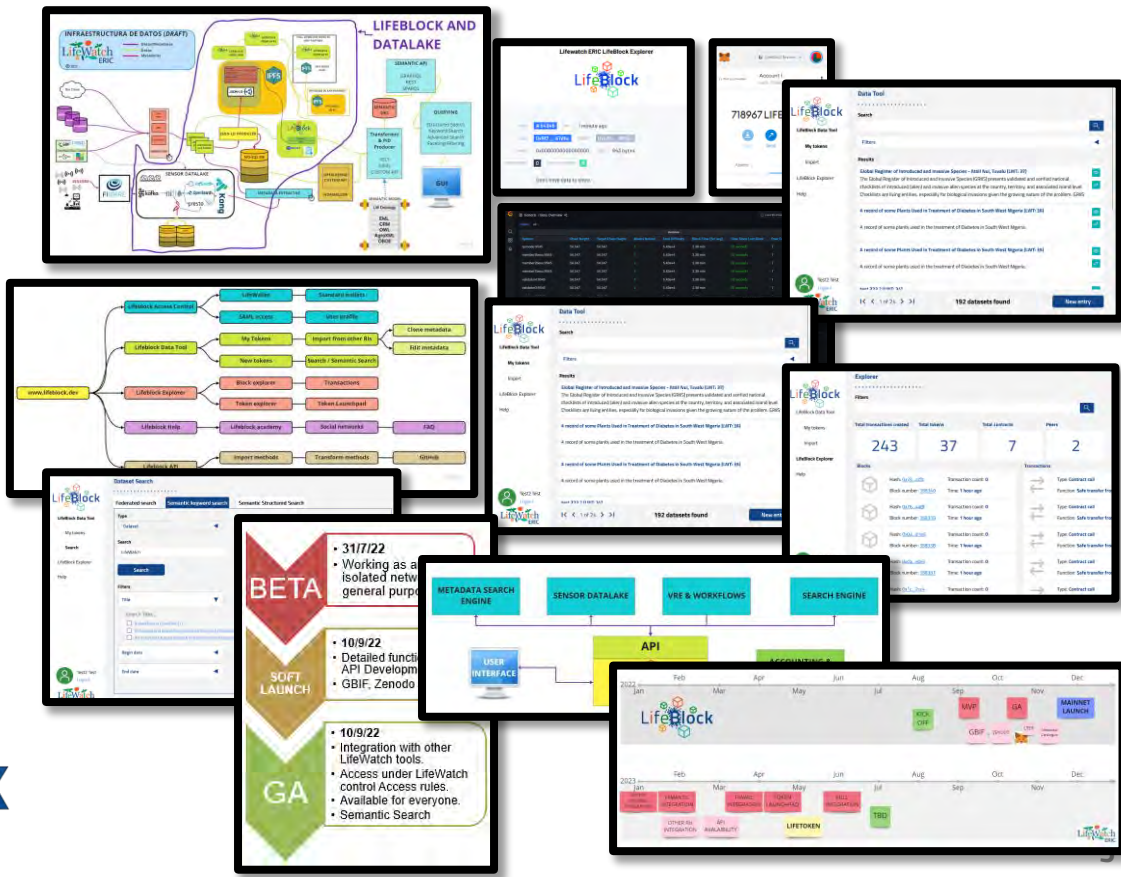
Biodiversity, Remote Sensing & Space Technologies

- FAIR DATA & e-Services
- Satellites
 - UNOOSA, Copernicus, Galileo
 - Space 4 SDGS
- High Altitude Balloons
- HAPS
 - High Altitude Pseudo Satellites
- First Andalusian Earth Observation Nanosatellite

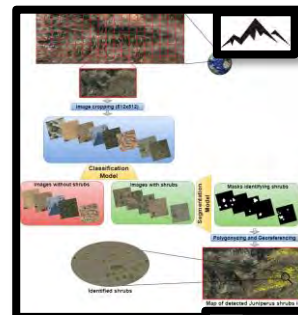
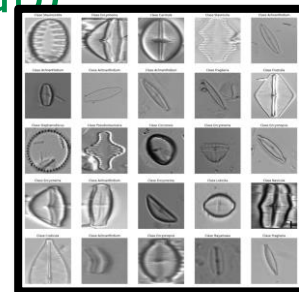
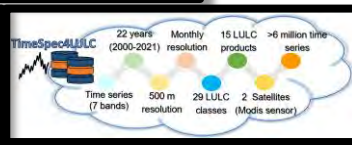
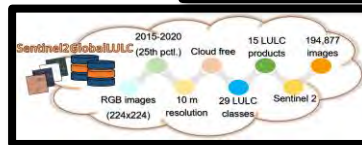


Dr. Joaquín López-Lérida

- LifeBlock and datalake architecture
- LifeBlock objectives
- LifeBlock proof of concept (2021)
- Development status
 - Sept. '22: Testnet
- LifeBlock functional map
- LifeBlock data tool: Sept '22
- LifeBlock explorer: Sept '22
- Semantic search and RI: Oct '22
- Short term roadmap
- LifeBlock API (REST, GRAPHQL, SPARQL)



- Projects:
 - Land use/cover mapping
 - High mountain shrubs detection.
 - Diatoms recognition
 - Photosynthetic pigments concentration estimation
 - Herbarium species identification



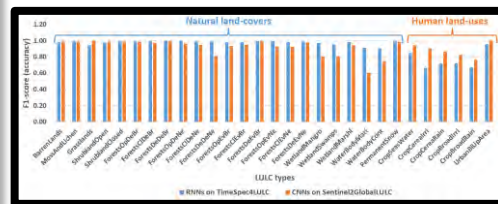
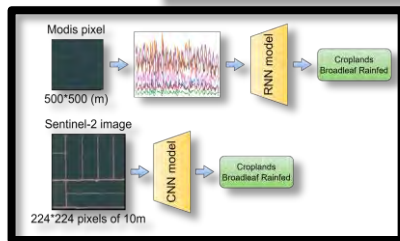
Clasificador de diatomeas automático

Predicir Nombre

La clase predicha por el modelo es **Amphora**

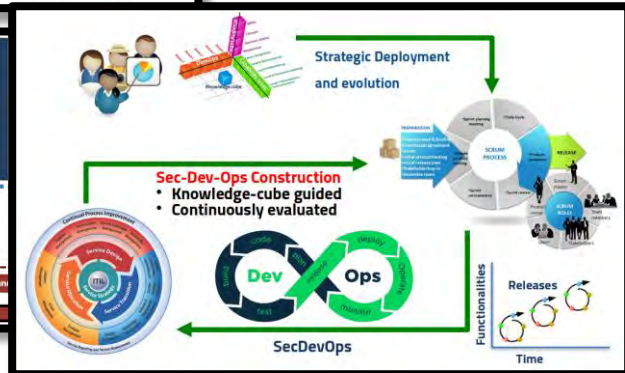
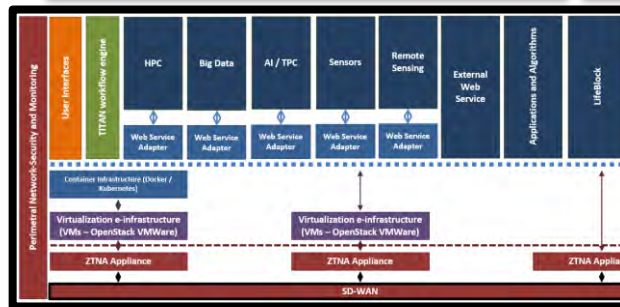
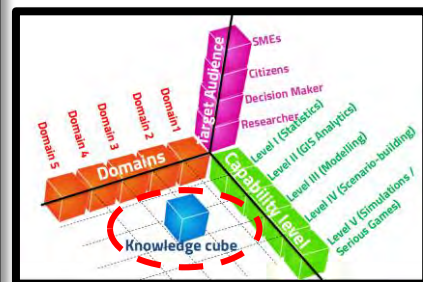
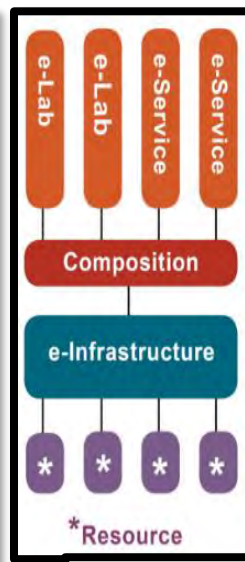
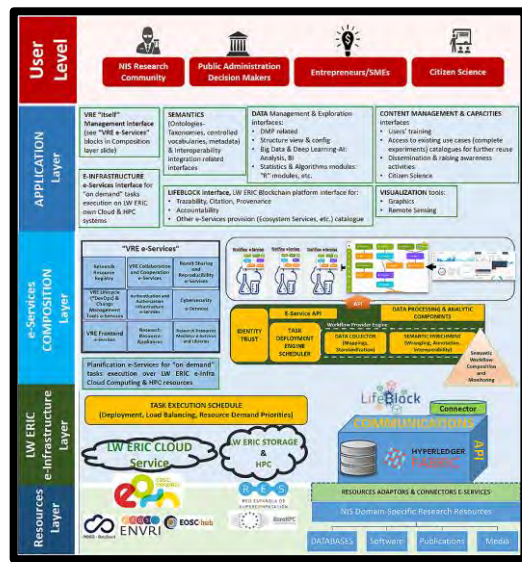
Predicciones:

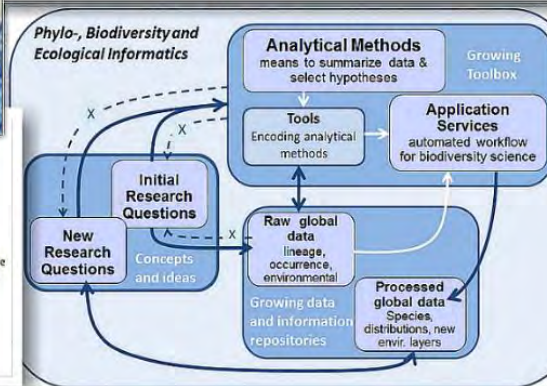
- Amphora - probabilidad: 100.0 %
- Encyonema - probabilidad: 0.0 %
- Gomphonema - probabilidad: 0.0 %
- Luticola - probabilidad: 0.0 %
- Psammodium - probabilidad: 0.0 %



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- The image illustrates a metabarcoding workflow, likely for environmental DNA (eDNA) analysis. It is divided into three main sections: a workflow diagram, a software interface, and a data table.
- ### Metabarcoding Validation Case - Workflow
- The workflow diagram shows the process from eDNA sample to OTU table, involving steps 1 through 8. The steps are:
- STEP 1 (MBC): eDNA Sample
 - STEP 2: Reference Database
 - STEP 3: Distribution of OTUs
 - STEP 4: OTU Table
 - STEP 5: OTU Table
 - STEP 6: OTU Table
 - STEP 7: OTU Table
 - STEP 8: OTU Table
- ### Tesseract Workflow Environment
- The Tesseract software interface shows the workflow environment. The main window displays the workflow overview, which includes the following steps:
- Workflow overview
 - Workflow description
 - Workflow execution
 - Workflow results
- The workflow overview shows the process flow from eDNA sample to OTU table, with intermediate steps for reference database, distribution of OTUs, and OTU table.
- ### OTU Table
- The OTU table displays the results of the metabarcoding analysis. The columns are:
- Accession
 - Scientific Name
 - Kingdom
 - Phylum
 - Class
 - Order
 - Family
 - Genus
 - Species
 - Abundance
 - Date
 - Location
 - Identifier
- The table contains 10 rows of data, representing different species and their abundance across different samples.

- Reference architecture
- ORCHESTRA model-based service oriented architecture
- User layer
- Application layer
- E-Service composition layer
- LW-ERIC e-Infrastructure layer
- Resources layer
- Design of functional layers on deployment
- Development methodology
 - SecDevOps
 - Knowledge cubes





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Thanks!

