

Data life cycle for Life Science

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SZN Seminar, Naples, 2023-02-09

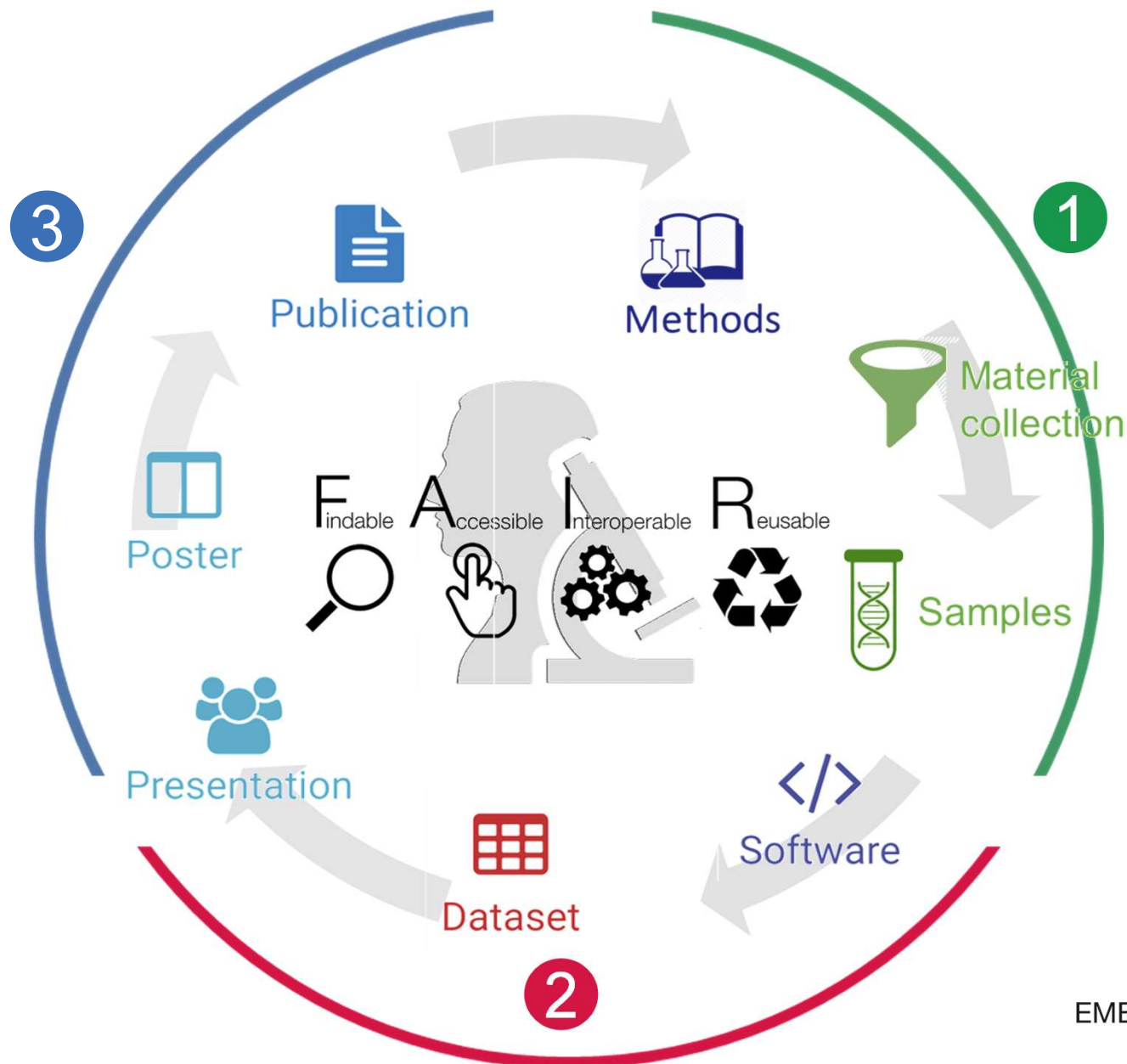


Data life cycle

Sampling
Best Practices

Data Sharing
Best Practices

Publication
Best Practices



Sampling best practices

Exemplar initiatives - developing and experimenting with best practices

- Tara Oceans
- MicroB3
- EMO-BON
- AtlantECO



System of protocols

International System of Units (SI)

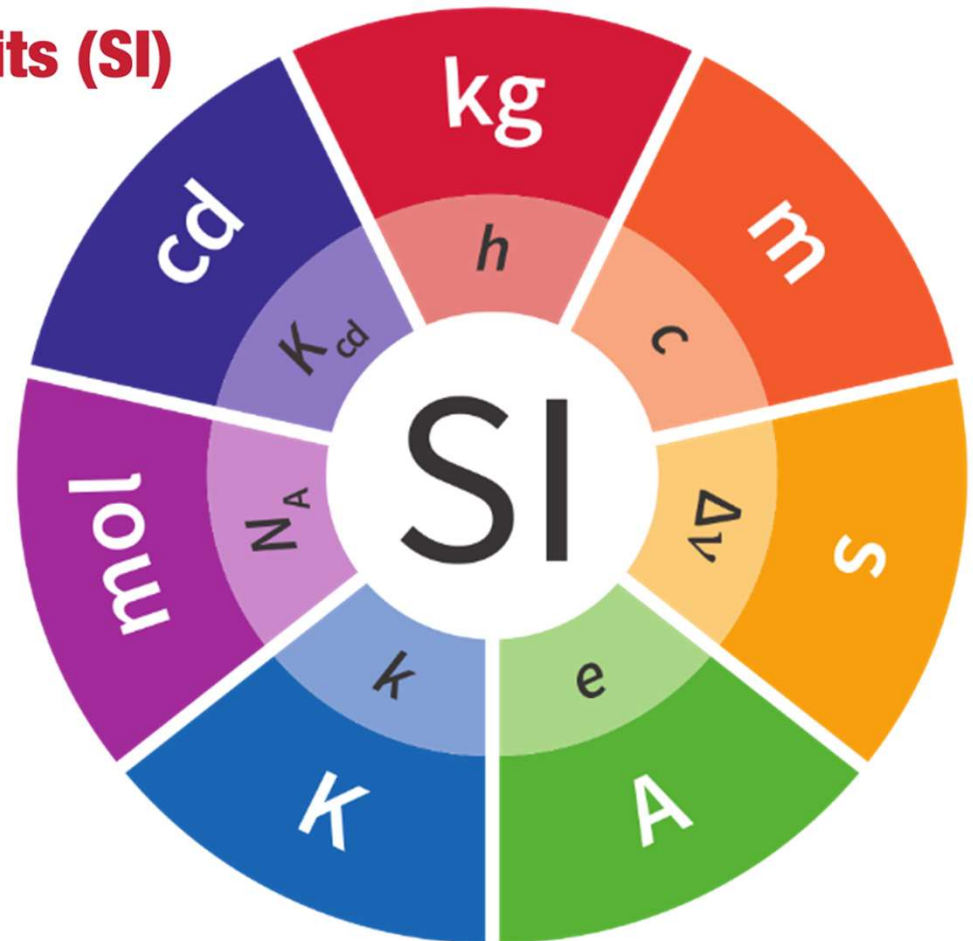
SI Base Units

| Base Quantity | Name | Symbol |
|---------------------------|----------|--------|
| Length | meter | m |
| Mass | kilogram | kg |
| Time | second | s |
| Electric current | ampere | A |
| Thermodynamic temperature | kelvin | K |
| Amount of substance | mole | mol |
| Luminous intensity | candela | cd |

SI Derived Units

| Derived Quantity | Name | Symbol | Equivalent SI units |
|---------------------|----------------|-------------|---------------------------|
| Frequency | hertz | Hz | s^{-1} |
| Force | newton | N | $m \cdot kg \cdot s^{-2}$ |
| Pressure | pascal | Pa | N/m^2 |
| Energy | joule | J | $N \cdot m$ |
| Power | watt | W | J/s |
| Electric charge | coulomb | C | $s \cdot A$ |
| Electric potential | volt | V | W/A |
| Electric resistance | ohm | Ω | V/A |
| Celsius temperature | degree Celsius | $^{\circ}C$ | K^* |

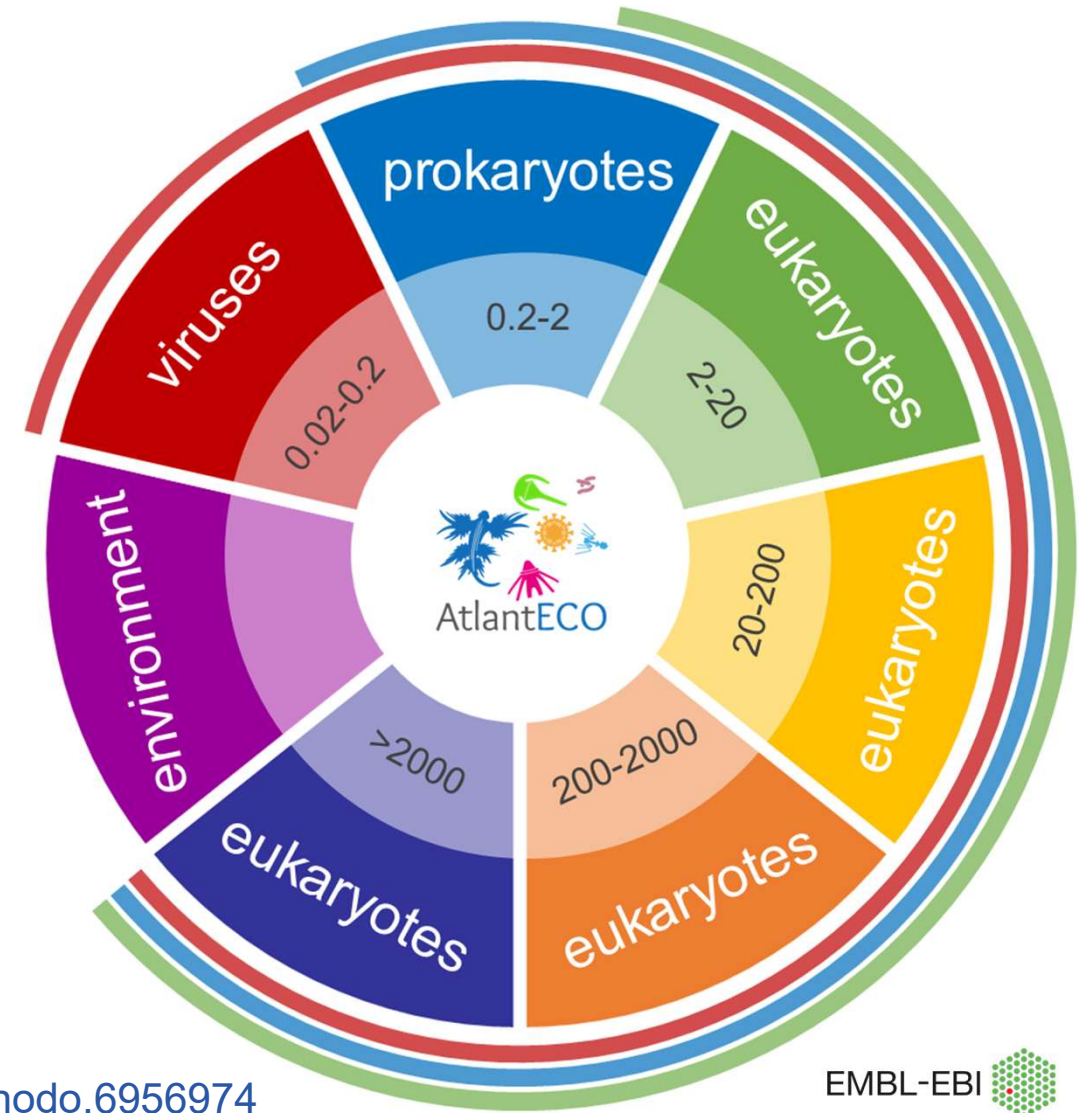
*Unit degree Celsius is equal in magnitude to unit kelvin.



System of protocols

- Size
- Taxonomy
- Across size-fractions

- Genomics
- Transcriptomics
- Proteomics
- Metabolomics
- Phenomics



<https://doi.org/10.5281/zenodo.6956974>

System of protocols

Base phenomics protocols

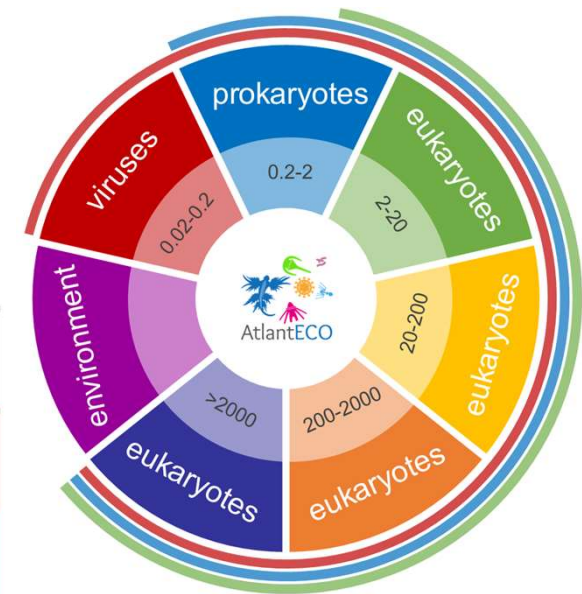
| Base Protocol | Target Size fraction | Target Analysis | Target Volume (L) | Target Concentration | Preservation |
|---------------|----------------------|-------------------------|-------------------|----------------------|--------------------------|
| I002 | 0.02-0.2 | Flow cytometry | 10^{-3} | | LN ₂ or -80°C |
| I02 | 0.2-2 | Flow cytometry | 10^{-3} | | LN ₂ or -80°C |
| I2 | 2-20 | Fluorescence microscopy | 10^2 | | +4°C |
| I20 | 20-200 | Flow imaging microscopy | 10^2-10^4 | | live |
| I200 | 200-2000 | Flatbed scan imaging | 10^3-10^5 | | formaldehyde |
| I2000 | >2000 | Flatbed scan imaging | 10^3-10^5 | | formaldehyde |
| environment | multiple | multiple | multiple | | |



System of protocols

Base genomics protocols

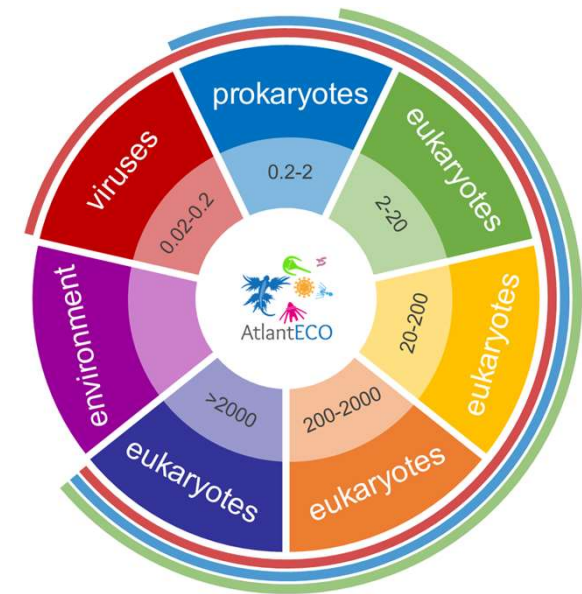
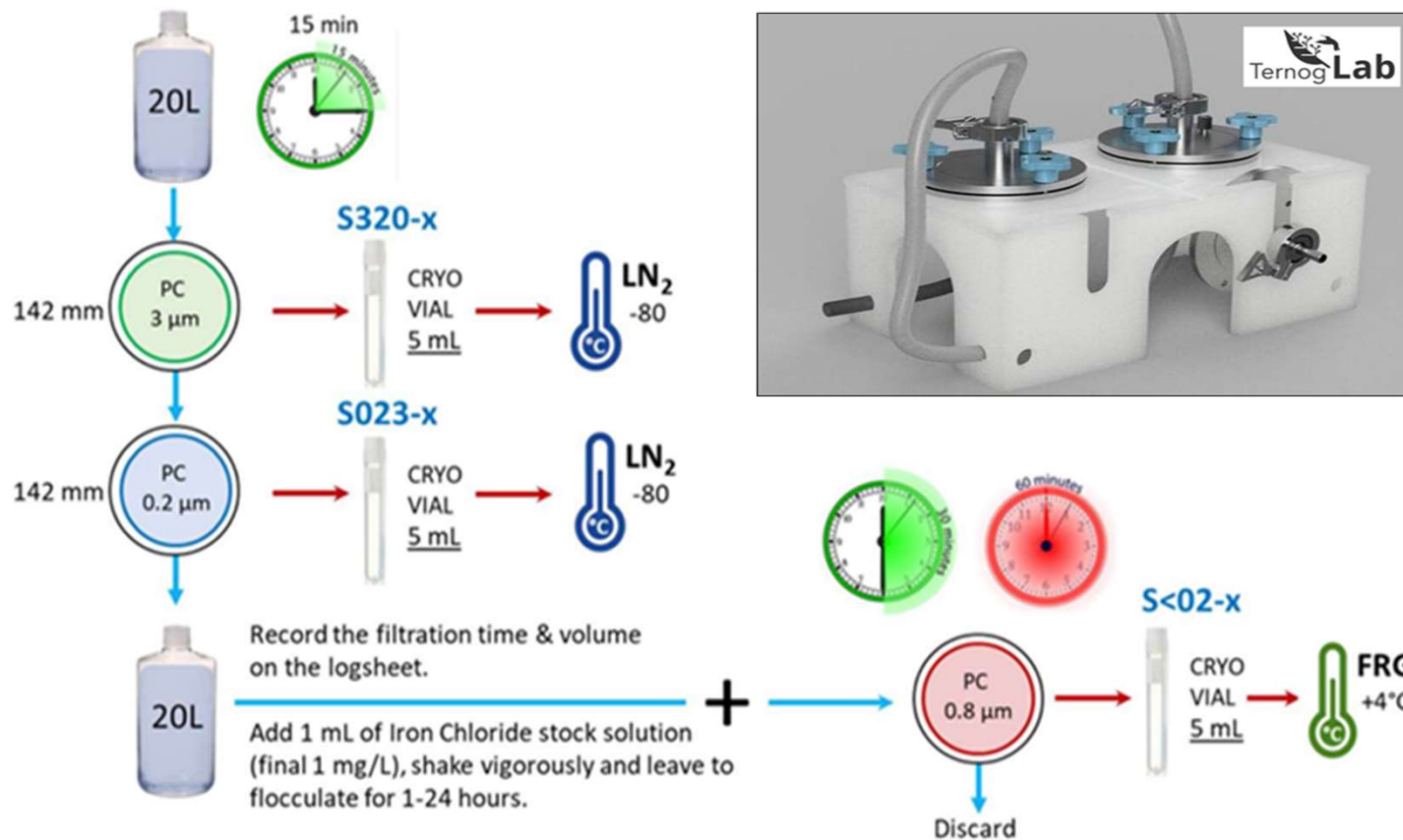
| Base Protocol | Target Size fraction | Target Analysis | Target Volume (L) | Target Time (min) | Preservation |
|---------------|----------------------|------------------------|----------------------------------|----------------------|--------------------------|
| S002 | 0.02-0.2 | MetaG, MetaT | 20 | Flocculated 4-24h | +4°C |
| S02 | 0.2-2 | MetaB, MetaG, MetaT | 20 | <15 | LN ₂ or -80°C |
| S2 | 2-20 | MetaB, MetaG, MetaT | 20 | <15 | LN ₂ or -80°C |
| S20 | 20-200 | MetaB, MetaG, MetaT | 10 ² -10 ⁴ | <15 | LN ₂ or -80°C |
| S200 | 200-2000 | MetaB, MetaG, MetaT | 10 ³ -10 ⁵ | <15 | LN ₂ or -80°C |
| S2000 | >2000 | MetaB, MetaG, MetaT | 10 ³ -10 ⁵ | <15 | LN ₂ or -80°C |
| eDNA | >0.2 | MetaB | 2 | | LN ₂ or -80°C |



Derived Protocols

- Size fractions
- Filtration volume
- Filtration time
- Preservation method

System of protocols

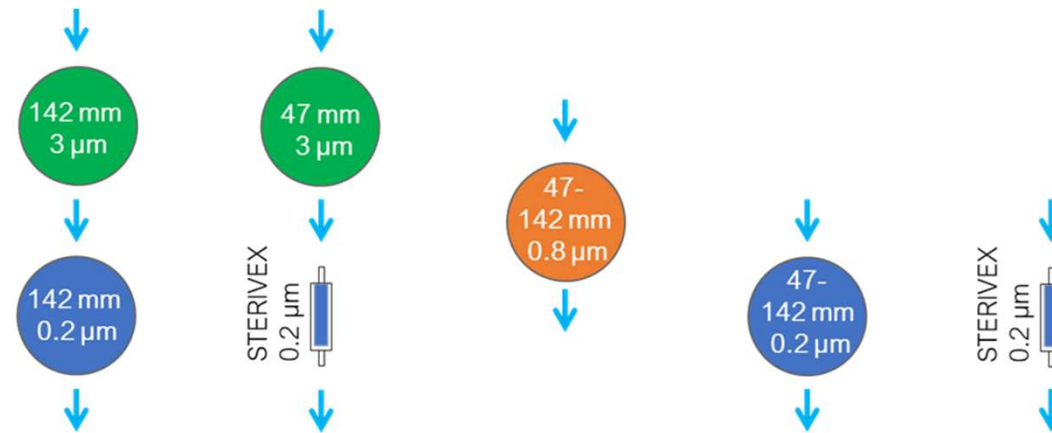


Derived Protocols

- Size fractions
- Filtration volume
- Filtration time
- Preservation method

Community Survey

All Atlantic Ocean
Microbiome Sampling



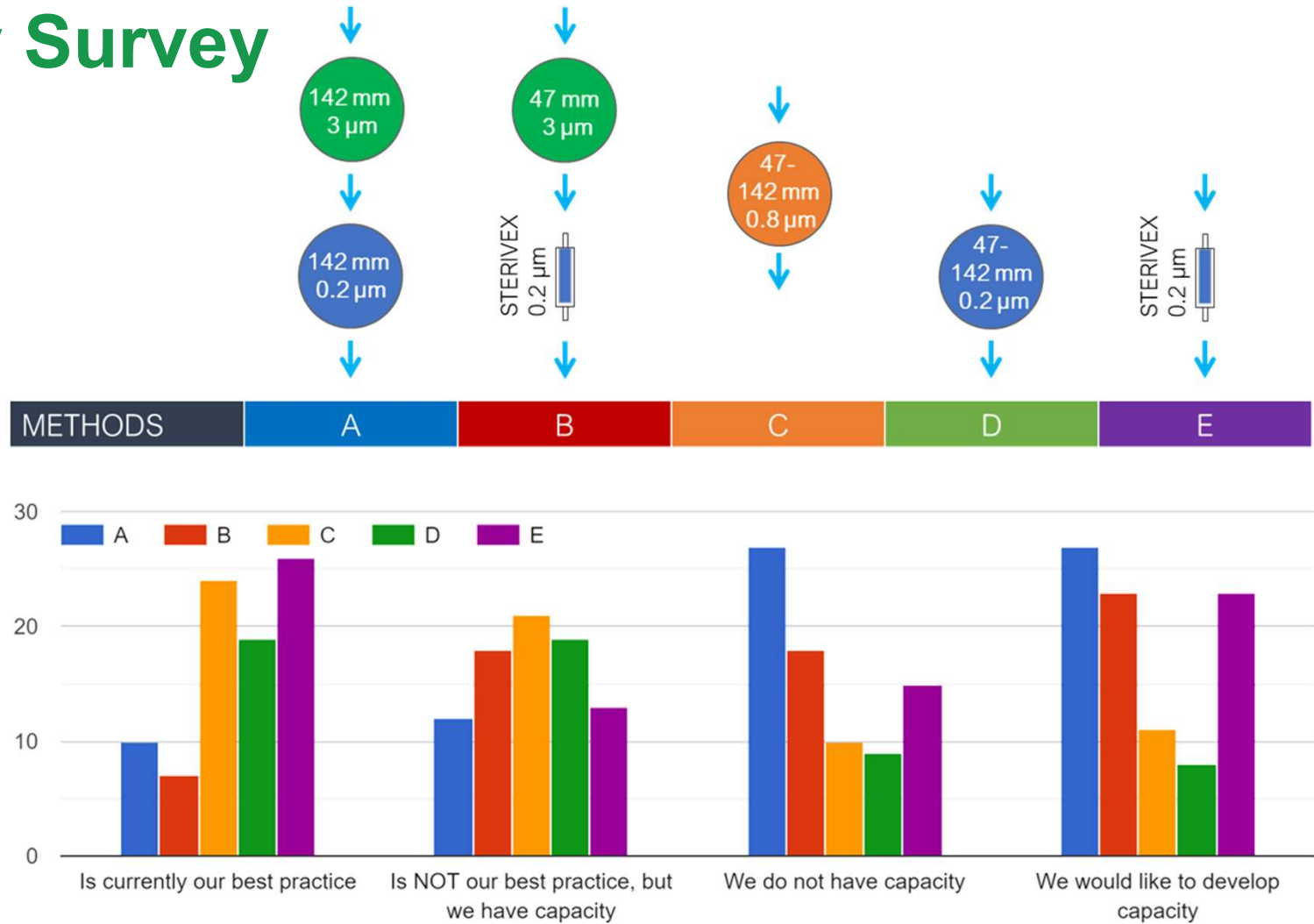
| METHODS | A | B | C | D | E |
|--------------------|---------------------|-------------------------|-----------------------|-----------------------|----------|
| Filters | 2x 142 mm membranes | 47 mm membr. + sterivex | 47 or 142 mm membrane | 47 or 142 mm membrane | sterivex |
| Vol. in 15 min. | 10-20 L | 2-5 L | 1-10L | 1-10L | 1-10 L |
| Pump system | large peristaltic | small peristaltic | various | various | various |
| ADOPTED BY | A | B | C | D | E |
| Ocean Sampling Day | | | (✓) | | ✓ |
| Bio-GO-SHIP* | | | | | ✓ |
| Tara Oceans | ✓ | | (✓) | (✓) | |
| EMO-BON** | ✓ | | | | |

*Bio-GO-SHIP Linking marine biodiversity and biogeochemistry (<https://biogoship.org/>)

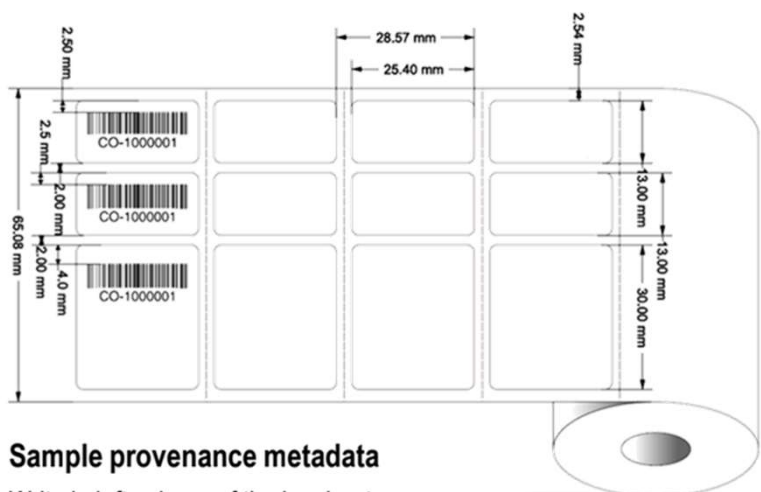
**European Marine Omics Biodiversity Observatory Network (<https://www.embrc.eu/emo-bon>)

Community Survey

All Atlantic Ocean
Microbiome Sampling



Collecting Metadata – unique identifiers



Sample provenance metadata

Write in left column of the logsheet:

1. Sampling depth (m)
2. Replicate # or Control #

Write on the large barcode sticker:

1. Station ID (e.g. **021**)
2. Sampling depth (e.g. **200** m)
3. Protocol label (e.g. **S320**)
4. Replicate # or Control # (e.g. **R1**)

Fix the large barcode sticker on the sample container

Fix the corresponding small barcode sticker on the logsheet, in the appropriate protocol column




Collecting Metadata – provenance & context

Fondation **taraocéan** explore and share **LOG_W-LAB-142** Mission **Microbiomes** C1

LOG_SAMPLES: 2022 04 09 STATION: 102 W-LAB-142
 OPERATOR(S): Cora

| Depth | Replicate | S320 Cryo-5mL LN2 #1 | S023 Cryo-5mL LN2 #1 | Filtration Volume (Litres) | Filtration Duration (minutes) | S<02 Cryo-5mL FRG +4°C | Filtration Volume (Litres) |
|-------|-----------|----------------------|--------------------------------------|----------------------------|-------------------------------|------------------------|----------------------------|
| 200 | R01 | 7908620 | 7908621 | 18.5 L | 15' min. | 7908634 | 10L 20L |
| 200 | R02 | ###-Z00 S320-2 | ###-Z00 S023-2 | 20L 50L | 15' min. | ###-Z00 S<02-2 | 10L 20L |
| 200 | R03 | ###-Z00 S320-3 | ###-Z00 S023-3 | 20L 50L | 15' min. | ###-Z00 S<02-3 | 10L 20L |
| 200 | R04 | ###-Z00 S320-4 | Collect filtrate for <3 µm protocols | 20L 50L | 15' min. | | 10L 20L |
| 202 | m | ###-Z02 S320 | ###-Z02 S023 | 20L 50L | 15' min. | ###-Z02 S<02 | 10L 20L |
| 204 | 60 m | 7918723 | 7918724 | 20L 50L | 14 min. | 7918725 | 10L 20L |
| 206 | m | ###-Z06 S320 | ###-Z06 S023 | 20L 50L | 15' min. | ###-Z06 S<02 | 10L 20L |
| Depth | Replicate | P320 Cryo-5mL LN2 #1 | P023 Cryo-5mL LN2 #1 | Filtration Volume (Litres) | Filtration Duration (minutes) | | |
| 200 | 0 m | 7908635 | 7908636 | 37 L | 15' min. | | |
| 202 | m | ###-Z02 P320 | ###-Z02 P023 | 20L 50L | 15' min. | | |
| 204 | 60 m | 7918726 | 7918727 | 18 L | 15' min. | | |
| 206 | m | ###-Z06 P320 | ###-Z06 P023 | 20L 50L | 15' min. | | |

AtlantECO  LOG_W-LAB-142_recto_V1




Collecting Metadata – structured checklists

Fondation **taraocéan** explore and share **LOG_W-LAB-142** Mission **Microbiomes** C1

LOG_SAMPLES: 2022 04 09 STATION: 102 W-LAB-142

OPERATOR(S): Gora

| Depth Replicate | S320 Cryo-5mL LN2 #1 | S023 Cryo-5mL LN2 #1 | Filtration Volume (Litres) | Filtration Duration (minutes) | S<02 Cryo-5mL FRG +4°C | Filtration Volume (Litres) |
|-----------------|----------------------|--------------------------------------|----------------------------|-------------------------------|------------------------|----------------------------|
| Z00 R01 0 m | 7908620 | 7908621 | 18.5 L | 15' min. | 7908634 | |
| Z00 R02 m | ###-Z00 S320-2 | ###-Z00 S023-2 | | | ###-Z00 S<02-2 | |
| Z00 R03 m | ###-Z00 S320-3 | ###-Z00 S023-3 | | | ###-Z00 S<02-3 | |
| Z00 R04 m | ###-Z00 S320-4 | Collect filtrate for <3 µm protocols | | | | |
| Z02 m | ###-Z02 S320 | ###-Z02 S023 | | | ###-Z02 S<02 | |
| Z04 60 m | 7918723 | 7918724 | 14 L | 14 min. | 7918725 | |
| Z06 m | ###-Z06 S320 | ###-Z06 S023 | | | ###-Z06 S<02 | |
| Depth Replicate | P320 Cryo-5mL LN2 #1 | P023 Cryo-5mL LN2 #1 | Filtration Volume (Litres) | Filtration Duration (minutes) | | |
| Z00 0 m | 7908635 | 7908636 | 37 L | 15' min. | | |
| Z02 m | ###-Z02 P320 | ###-Z02 P023 | | | | |
| Z04 60 m | 7918726 | 7918727 | 18 L | 15' min. | | |
| Z06 m | ###-Z06 P320 | ###-Z06 P023 | | | | |

AtlantECO  LOG_W-LAB-142_recto_V1

| attribute | Format / units | comment |
|---|--|---|
| sample id | SAMEA0000000 | |
| sample label | Alpha-numeric, e.g. "project_date-time_station_environment_size-fraction_method" | human readable and meaningful label |
| sampling design, label(s) | alpha-numeric | campaign, station, site, transect, etc. |
| sampling device | alpha-numeric | device name & specifications |
| operator | alpha-numeric | initials or full name |
| sampling date and time | yyyy-mm-dd T hh:mm | in UTC |
| latitude | N/S dd.dxxxxx or N/S dd mm.mmm | ultimately in decimal degree N |
| longitude | E/W ddd.dxxxxx or E/S ddd mm.mmm | ultimately in decimal degree E |
| elevation, depth below soil surface | cm | ultimately in metre |
| elevation, depth below sediment surface | cm | ultimately in metre |
| elevation, depth below water surface | m | |
| elevation, altitude above sea level | m | |
| methodological details | alpha-numeric | e.g. processing time & volume |

Data sharing best practices



Permanent archives selected for the different data types:

- BioSamples for metadata (<https://www.ebi.ac.uk/biosamples/>)
- ENA for genomics data (<http://www.ebi.ac.uk/ena>)
- MGnify for metagenomic data (<https://www.ebi.ac.uk/metagenomics/>)
- PRIDE for proteomics data (<https://www.ebi.ac.uk/pride/>)
- Metabolights for metabolomics data (<https://www.ebi.ac.uk/metabolights/>)
- BioImage Archive for imaging data (<https://www.ebi.ac.uk/bioimage-archive/>)

Data workflow

Environmental

EMODnet, Copernicus, etc.



Provenance & Envir. Context

BioSamples



Genomics

ENA, Mgnify & Ensembl



Proteomics

PRIDE



Metabolomics

Metabolights



Imaging

BiImage archive & EcoTaxa



primary data
curation & archiving

annotated data
analysis & archiving

Data workflow

Environmental

EMODnet, Copernicus, etc.

Provenance & Envir. Context

BioSamples

Genomics

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EMODnet



Harmonised context

- Georeferences
- Taxonomy
- Marine domain

- Longhurst Biogeo. Provinces
- IHO Sea Areas
- Oceanographic Products
- Astronomical Products

- Countries & Territories
- Exclusive Economic Zones
- Nagoya requirements
- ABS regulations



Data workflow

Environmental
EMODnet, Copernicus, etc.

Provenance & Envir. Context
BioSamples

Genomics
ENA, Mgnify & Ensembl

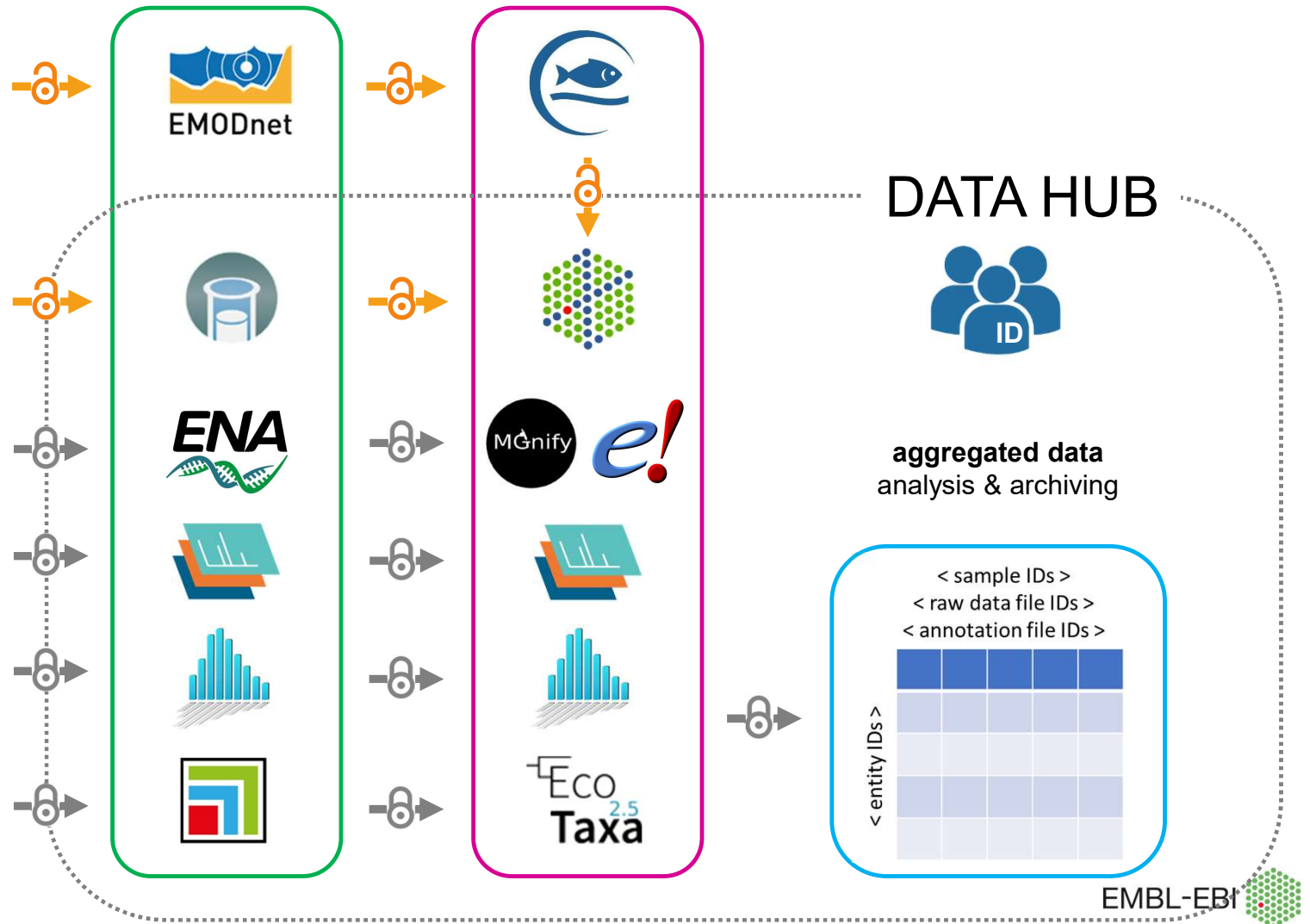
Proteomics
PRIDE

Metabolomics
Metabolights

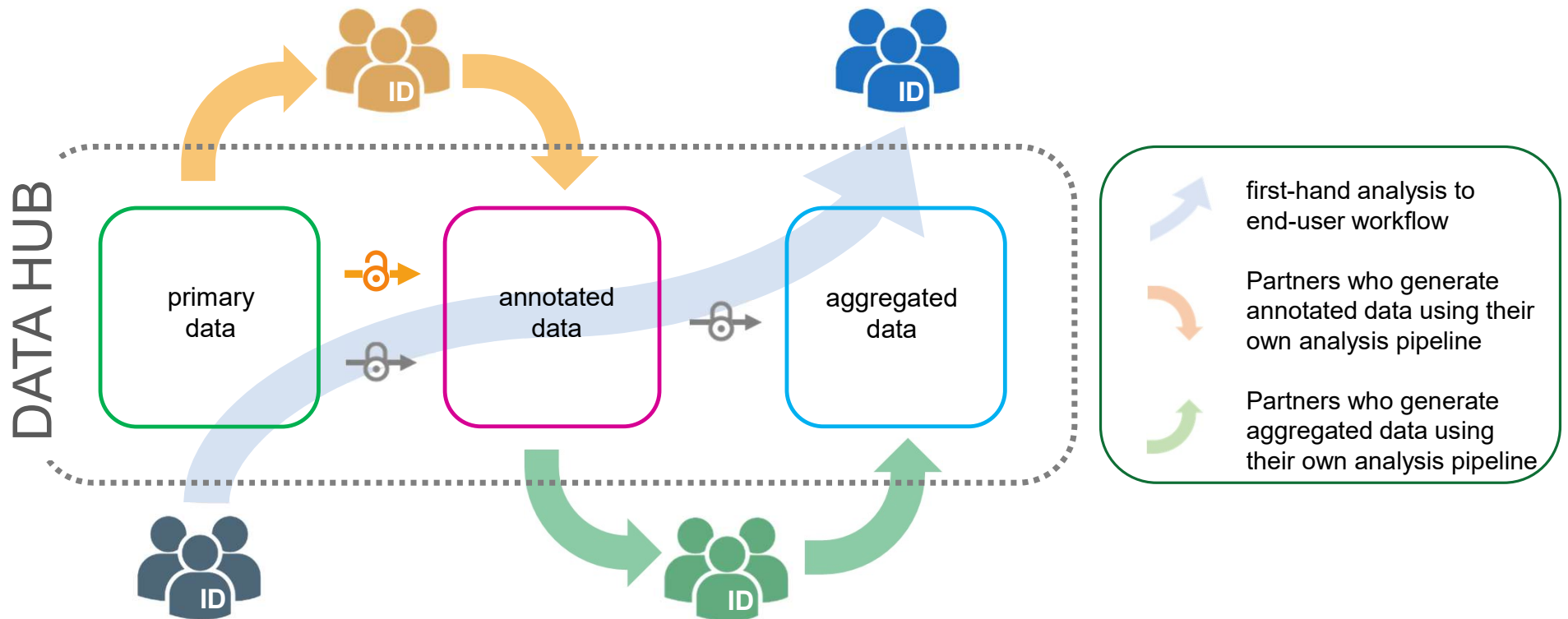
Imaging
BioImage archive & EcoTaxa

primary data
curation & archiving

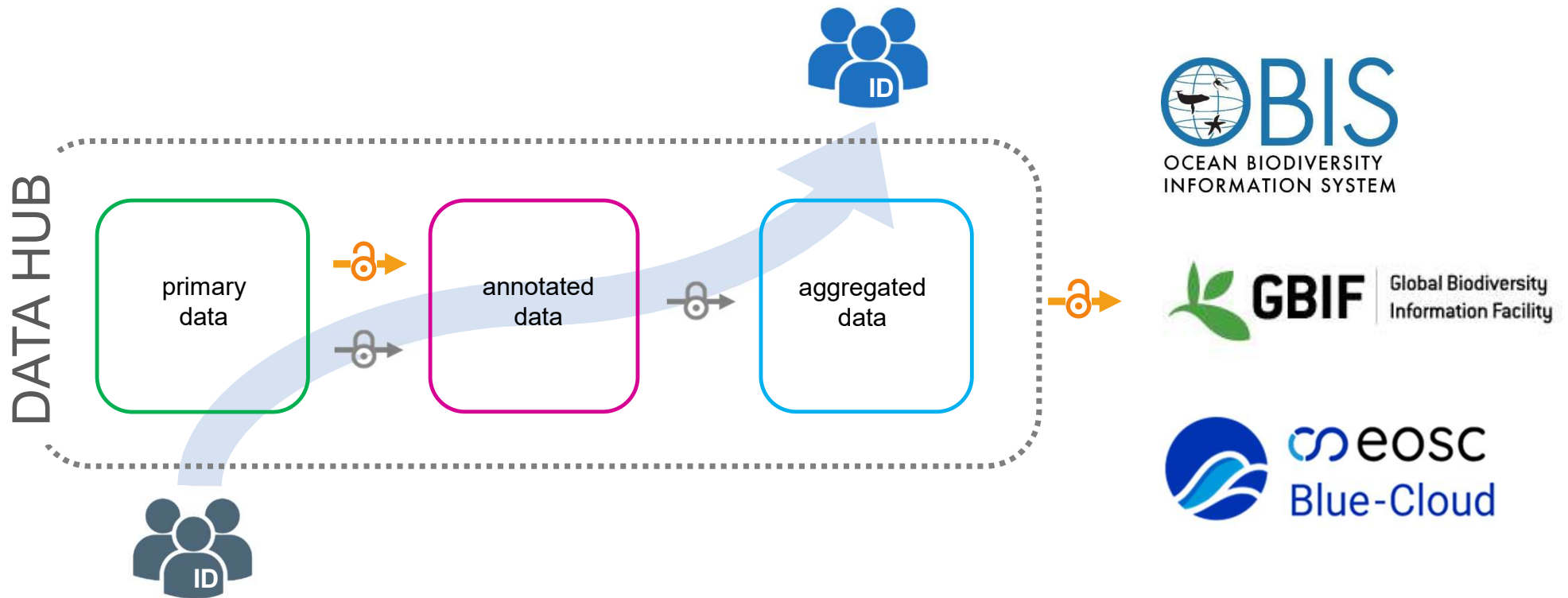
annotated data
analysis & archiving



Data Hub workflow



Data Hub workflow



eosc

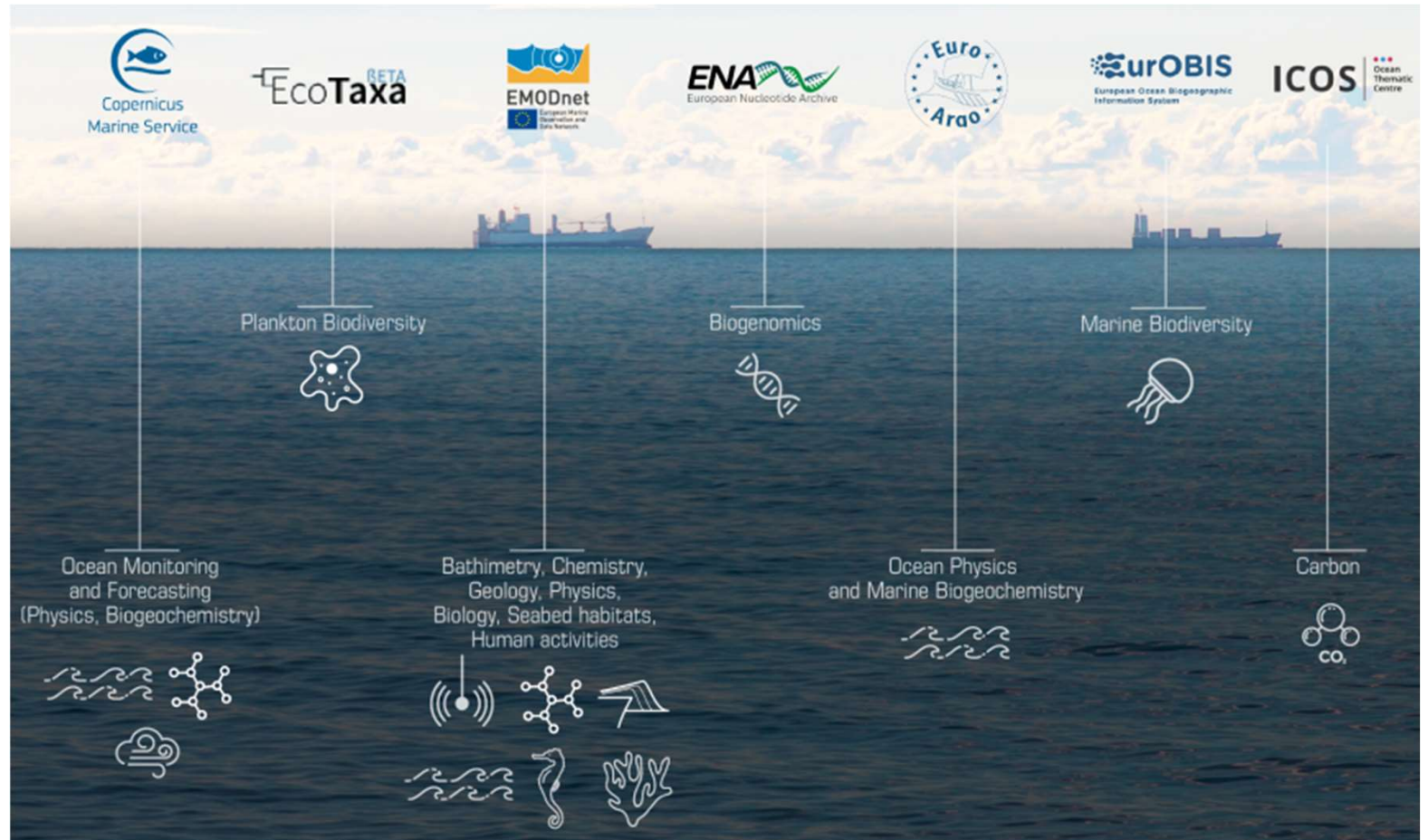
Blue-Cloud



<https://blue-cloud.org/>



Discovery & Access

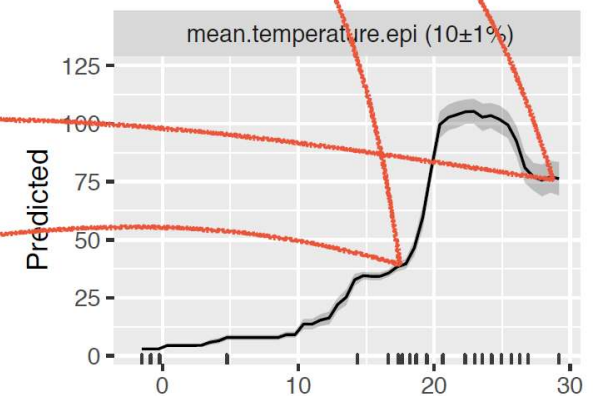
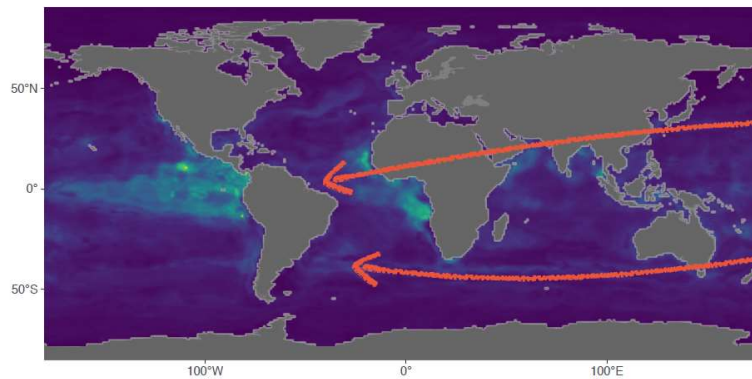
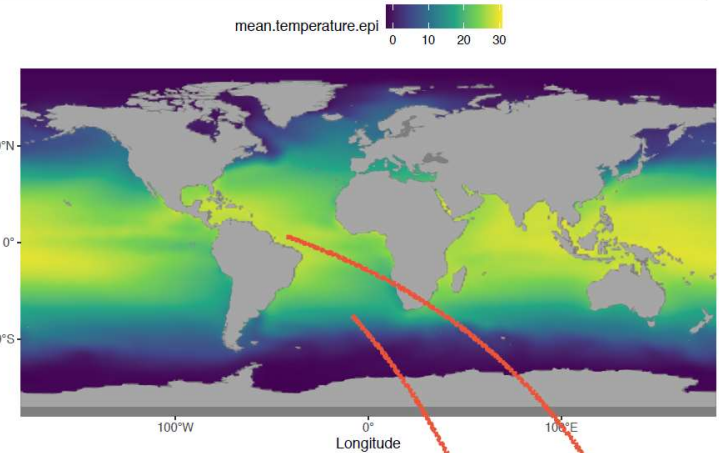
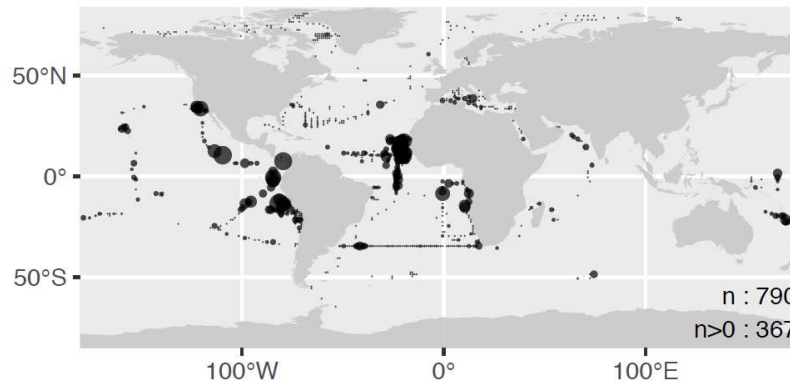




Discovery & Access



Virtual Labs & Workbenches





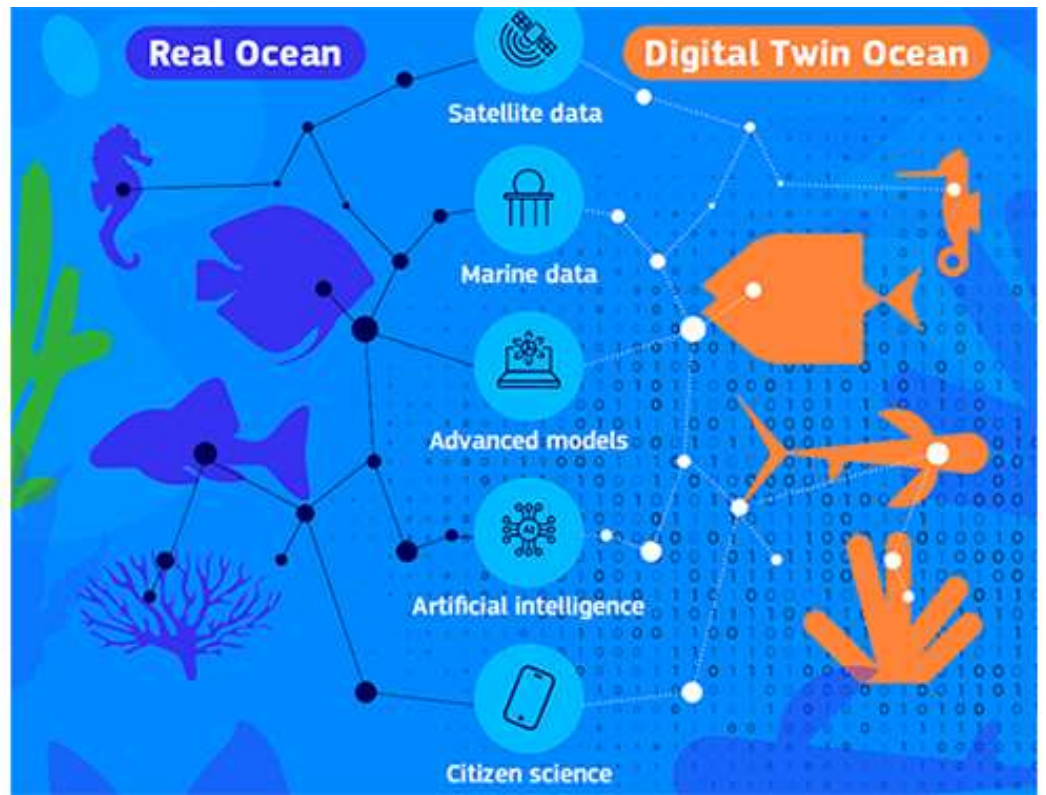
Discovery & Access



Virtual Environment



Publishing Services



Publication best practices

Exemplar initiatives

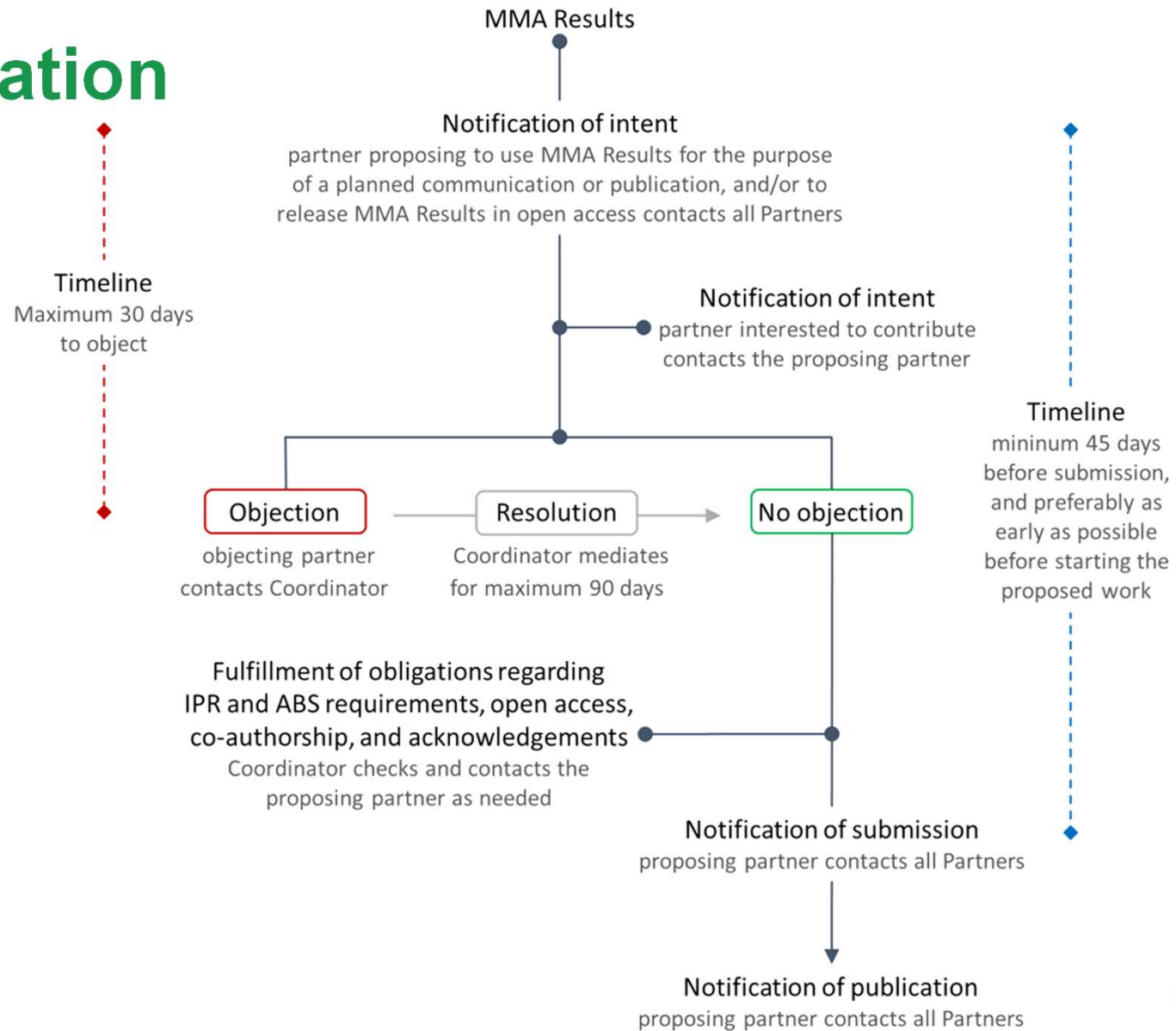
- Tara Oceans
- Tara Pacific
- AtlantECO

Best Practices

- Jointly owned results
- Early notification of intent to use “jointly owned results”
- Co-authorship
- Open Access publication



Early notification



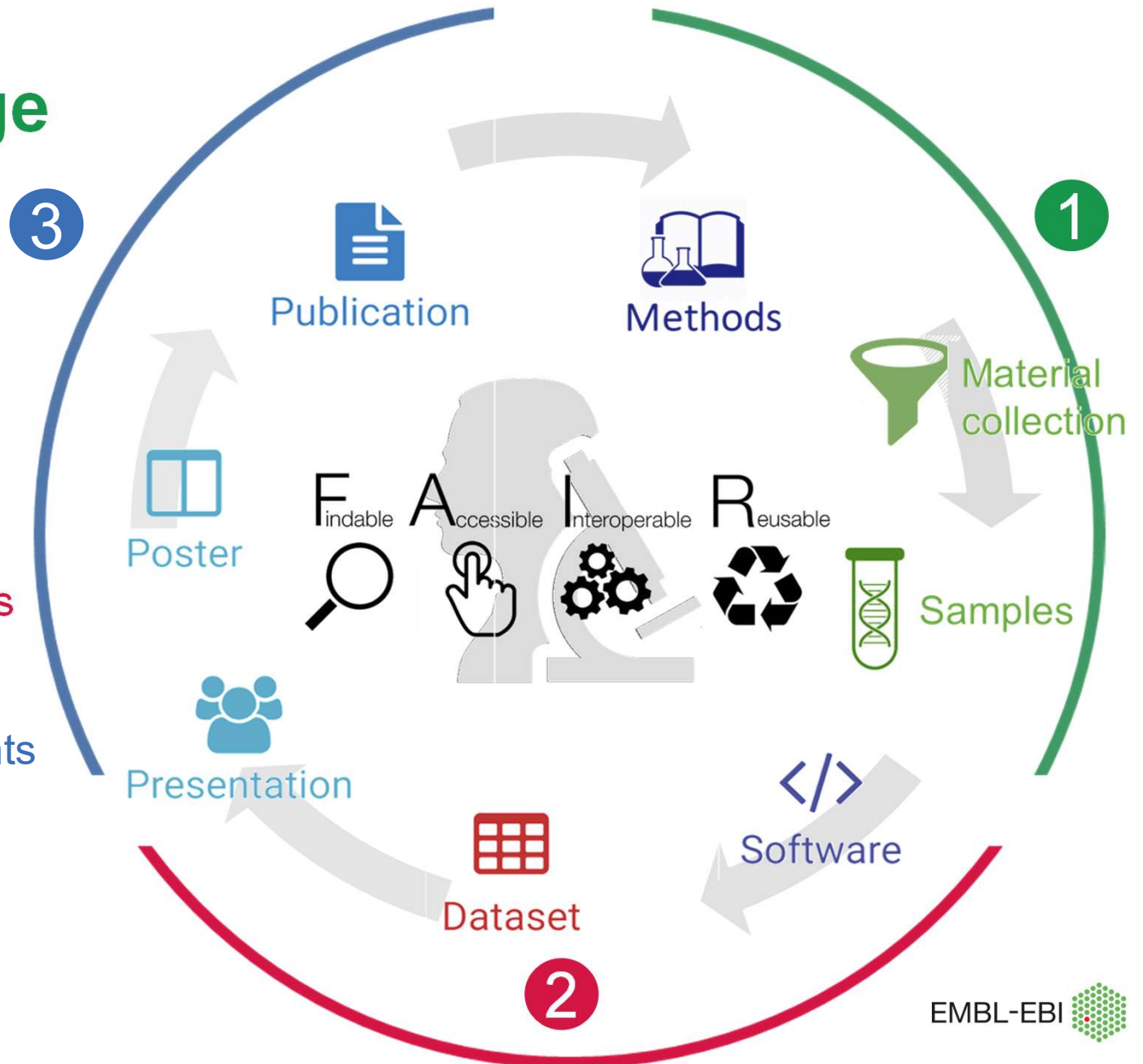
Take home message

Use standard methods
ADAPT
Collect structured metadata

Deposit metadata & data
EARLY
Progressive closed > open access

Communicate your research intents
EARLY
Seek collaboration

Publish in open access



**Thank you
for your kind attention**

Stéphane PESANT

Senior marine biocurator

pesant@ebi.ac.uk

