



PAN-EUROPEAN INFRASTRUCTURE  
FOR OCEAN & MARINE DATA  
MANAGEMENT

## ***Making SeaDataNet more fit for handling biological data***

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<sup>2</sup> International Council for the Exploration of the Seas (ICES)

**IMDIS, 23-25 September 2013**

# Objective

- **SDN II extension to marine biological data**

SeaDataNet II will undertake actions to make SeaDataNet better fit for handling marine biological data sets and establishing interoperability with biology infrastructure developments.

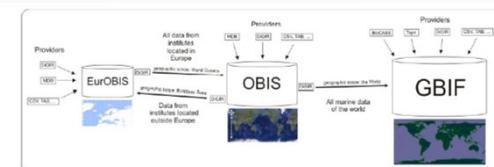
EMODNET biology

EUROBIS & (OBIS and GBIF)

EUROMARINE

WORMS (& PESI, 4D4Life, EOL)

LIFEWATCH



# *Objective*

Original DOW task description :

WP8.3: Defining extended metadata format for CDI to support handling of marine biological data

- D8.4 Analysis report with required adaptions for marine biological data
  - ✓ required adaptions of the CDI format
  - ✓ required extensions of the vocabularies
  - ✓ possible need for support of extra data formats
  - ✓ proposed exchange protocol for handling marine biological data

# *Approach*

- Look at the types of data concerned and corresponding minimal requirements
- Look at data use and applications and corresponding minimal requirements
- Look at existing standards and practices in the biological community
- Compare to available standards within SeaDataNet approach

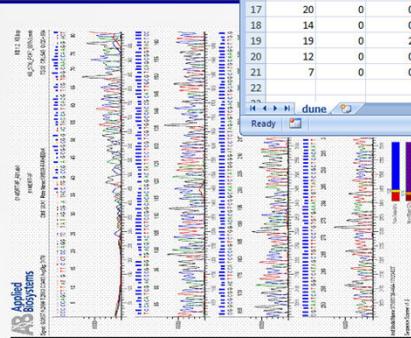


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## Example data

WGPDMOexample.txt - Kladblok									
Bestand Bewerken Opmak Beeld Help									
00, BLUK, 74, 2004, 3, 2, ,,,									
20, BLUK, 1, GRT, ,,, 4									
23, 1, BLUK, , TI9, NA, NON, NA, ,,, GRS									
90, 74E9, CEND7/04, DEFRA, ,,,									
91, CEND7/04, 77, +55 19. 234, -001 15.123, HLS, 20040705									
03, CEND7/04, 77, CF, 1, 1, , LIMANDA LIMANDA, ITLN, RS, ,,,									
04, CEND7/04, 77, 1, 1, OW, M, NS, K, 3, ,,,									
10, CEND7/04, 77, 1, 1, WO, , LNMEA, CM, 1, ,,, 17									
10, CEND7/04, 77, 1, 1, EP, , LYMP CYS, AFNR, 1, 1, ,,,									
10, CEND7/04, 77, 1, 1, EP, , SKIN ULC, AFNR, 1, 0, ,,,									
10, CEND7/04, 77, 1, 1, EP, , EPID PAP, AFNR, 1, 0, ,,,									
04, CEND7/04, 77, 1, 2, 1, EP, , LYMP CYS, AFNR, 1, 1, ,,,									
10, CEND7/04, 77, 1, 2, WO, , LNMEA, CM, 1, ,,, 18									
10, CEND7/04, 77, 1, 2, EP, , LYMP CYS, AFNR, 1, 1, ,,,									
10, CEND7/04, 77, 1, 2, EP, , SKIN ULC, AFNR, 1, 0, ,,,									
10, CEND7/04, 77, 1, 2, EP, , EPID PAP, AFNR, 1, 1, ,,,									
04, CEND7/04, 77, 1, 3, 4, OW, F, NS, K, 2, ,,,									
10, CEND7/04, 77, 1, 3, WO, , LNMEA, CM, 1, ,,, 21									
10, CEND7/04, 77, 1, 3, EP, , SKIN ULC, AFNR, 1, 0, ,,,									
10, CEND7/04, 77, 1, 3, EP, , LYMP CYS, AFNR, 1, 4, ,,,									
04, CEND7/04, 77, 1, 4, 1, OW, F, NS, K, 3, ,,,									
10, CEND7/04, 77, 1, 4, WO, , LNMEA, CM, 1, ,,, 23									
10, CEND7/04, 77, 1, 4, EP, , SKIN ULC, GRADE, 1, ,,, 3									
10, CEND7/04, 77, 1, 4, EP, , EPID PAP, GRADE, 1, ,,, 1									
10, CEND7/04, 77, 1, 4, EP, , LYMP CYS, GRADE, 1, ,,, 0									



```
<?xml version="1.0" encoding="UTF-8"?>
<SimpleDarwinRecords>
  xmlns="http://rs.tdwg.org/dwc/simpledarwincore/"
  xmlns:dwc="http://purl.org/dc/terms/"
  xmlns:dc="http://www.w3.org/2001/XMLSchema-instance"
  xsi:schemaLocation="http://rs.tdwg.org/dwc/simpledarwincore/ http://rs.tdwg.org/dwc/xsd
<SimpleDarwinRecord>
  <dcm:modified>2006-05-04T18:13:51.0Z</dcm:modified>
  <dcl:language>en</dcl:language>
  <dwc:basisOfRecord>Taxon</dwc:basisOfRecord>
  <dwc:scientificNameID>http://research.calacademy.org/research/ichthyology/catalog/fishatg...
  <dwc:acceptedNameUsage>http://research.calacademy.org/research/ichthyology/catalog/fishatg...
  <dwc:nameAccordingToID>http://research.calacademy.org/research/ichthyology/catalog/getref.asp?id=2764</dwc:nameAccordingToID>
  <dwc:scientificName>Centropyge flavicauda Fraser-Brunner 1933</dwc:scientificName>
  <dwc:acceptedNameUsage>Centropyge fisheri (Snyder 1904)</dwc:acceptedNameUsage>
  <dwc:parentNameUsage>Centropyge Kaup, 1860</dwc:parentNameUsage>
  <dwc:originalNameUsage>Centropyge flavicauda Fraser-Brunner 1933</dwc:originalNameUsage>
  <dwc:nameAccordingToAllen>Allen, G.R. 1980. Butterfly and angelfishes of the world. Volume II. Mergus Publishers. Pp. 149-352.</dwc:namePublishedIn>
  <dwc:higherClassification>Meetings for Scientific Business of the Zoological Society of London 1933 (pt 3, no.30): 543-599, Pl. 1.</dwc:higherClassification>
  <dwc:kingdom>Animalia</dwc:kingdom>
  <dwc:phylum>Chordata</dwc:phylum>
  <dwc:class>Osteichthyes</dwc:class>
  <dwc:order>Perciformes</dwc:order>
  <dwc:family>Centriscidae</dwc:family>
  <dwc:genus>Centropyge</dwc:genus>
  <dwc:specificEpithet>flavicauda</dwc:specificEpithet>
  <dwc:scientificNameAuthorship>Fraser-Brunner 1933</dwc:scientificNameAuthorship>
  <dwc:taxonRank>Species</dwc:taxonRank>
  <dwc:taxonomicStatus>Accepted</dwc:taxonomicStatus>
</SimpleDarwinRecord>
</SimpleDarwinRecords>
```



Vissoort :	Schol	Datum :	04/07/2008
Vaartuig :	O.89	PK :	Vissenij : bokken
Zone :	N	Visuren:	Totale vangst : 11060
Sortering	1	2	3
Tot. vangst	1880	3491	4289
Monster	25.460	36.450	48.100
24			
25			
26		3	
27		7	
28		20	
29		23	
30		27	1
31		14	12
32		5	29
33		1	5
34		21	16
35		6	20
36		1	27
37		15	8
38		7	21
39			28
40		2	2
41		1	16
42			4
43			3
44			10
45			22
46			1
47			5
48			10
49			4
50			2



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## ***Types of data***

- Observation and results
    - Occurrence
    - Density
    - Biomass
    - Body morphology
    - Condition
    - Substance concentrations or ratios
    - Sequencing material
  - Biological components (=non-taxonomic groups)
    - benthos, plankton, fish, birds, mammals, ...
- FORMAT & FIELD REQUIREMENTS
- FORMAT & FIELD REQUIREMENTS



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## ***Types of data***

- Geometry and sampling protocol
  - Point
    - soft-bottom grabs & cores (depth layer separation possible)
    - vertical net and water samples (multiple depths possible)
    - static net samples
    - hard-bottom sampling (scraping or visual)
    - static observations/underwater photography
  - Curve
    - net trawl, dredge or sledge
    - transect observations/underwater video
  - Surface
    - surface observations



**FORMAT & FIELD  
REQUIREMENTS**



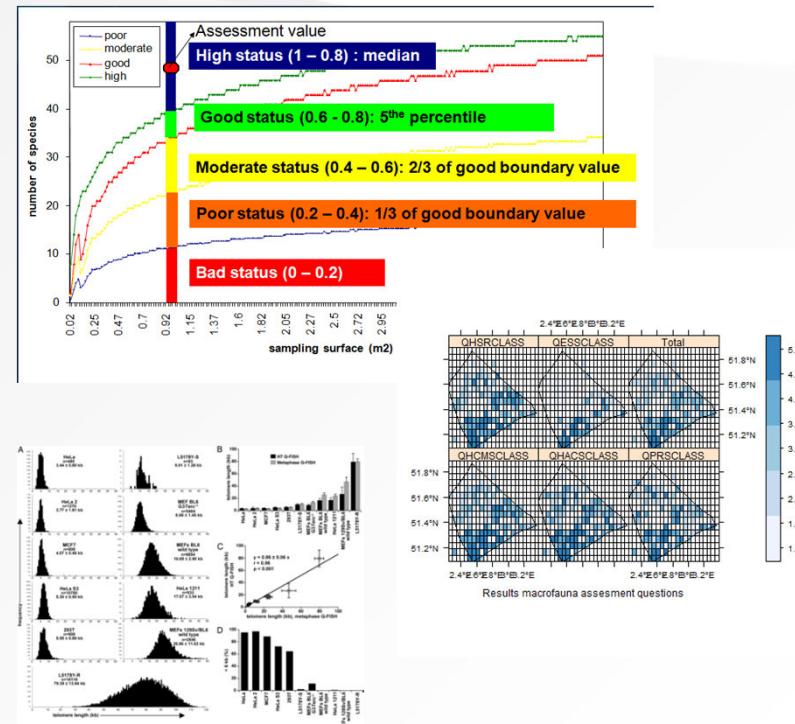
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# Data applications

- Species lists
- Species distribution
- Species richness
- Index calculations  
*BEQI, AMBI, BQI, ...*
- Population studies
- MSFD?, OSPAR?, ICES?...

→ FORMAT & FIELD  
REQUIREMENTS



## ***Standards and practices in the biological community***

### **Biodiversity Information Standards (TDWG)**

- non-profit scientific & educational association affiliated with International Union of Biological Sciences.
- development of standards for the exchange of biological/biodiversity data.

### **ABCD (Access to Biological Collections Data)**

- schema for biological collection records with numerous specific extensions (1200 concepts in ABCD 2.06)

### **Darwin Core (DwC)**

- versioned body of standards to facilitate the sharing of information about biological diversity

### **OBIS schema**

- Darwin core V2 specifically aimed at ocean biogeographic information

### **LSID: Life Science Identifier**

persistent, globally unique identifiers for biological objects including taxonomic names, specimen records, images and DNA sequences...

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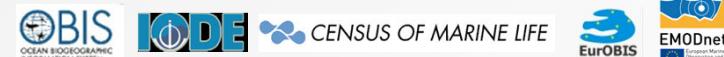
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## ***Standards and practices in the biological community***

- Some large marine integrative databases
  - Macroben [MarBEF (FP6)]
    - **Macrobenthos**
  - Manuela [MarBEF (FP6)]
    - **Meiobenthos**
  - LargeNet [MarBEF (FP6)]
    - **Benthos and plankton**
  - Hermione DB [Hermione (FP7)]
    - **Deep sea observations**
  - IMERS [regional survey database]
    - **Benthos, Plankton, Seabirds, ...**

	OBIS required	MarBEF MacroBEN	MarBEF LargeNet	MarBEf Manuela	Hermione DB	IMERS
Survey		/	/	/	cruise	trip
LocationID		statcode	stat_code	stat_code	seapname	namedplace
Longitude	X	lat	lat	lat	lat	start latitude
Latitude	X	long	long	long	long	start longitude
WKTFootPrint		/	/	/	/	start latitude
WKTFootPrint		/	/	/	/	start longitude
WKTFootPrint		/	/	/	/	end latitude
WKTFootPrint		/	/	/	/	end longitude
/						
MinimumDepth		depth	bottom depth	depth_min	sample_depth	sampling depth
MaximumDepth		depth	bottom depth	depth_max	sample_depth	sampling depth
EventID		/	/	/	eventID, eventlabel	eventid
InstitutionCode	X	institute	institute	institute	institute	Institute
SamplingProtocol		(mesh, equipment)	sample_method, subsample protocol,	sample_method	samplingequipment	instrument
EventDate		day, month, year	sample_year, sample_month, sample_day	sample_year, sample_month, sample_day	date_time, sample_year, sample_month, sample_day	startdatetime, enddatetime
FieldNumber		replicate	sample_id, replicate	sample, replicate	replica_id	Sample ID
SampleSize		Area	sample_area	sample_area	sample_surface_value	distancetrawled, volumefiltered
ScientificName	X	/	cleanname	cleanname	species	Taxon
ScientificNameID		aphia_id	aphia_id	aphia_id	aphia_ID	aphia_ID
Sex	X	/	/	/	/	gender
LifeStage		stage	stage	stage	lifestage	lifestage
ObservedIndividualCount		count	count	abundance	count	count

## ***Compare to available standards within SeaDataNet***

- Current Data transport format useful for biological data?

⇒ Not as it is;  
in order to adequately describe the biological observations we need:

- Additional information elements needed
  - In format (or in CDI)
- Adapted parameter vocab
- More generic semantic header

A	B	C	D	E	F	G	H	I	J	K	L	M
1	//Data documentation at <a href="http://www.bodc.ac.uk/data/documents/series/49486/">http://www.bodc.ac.uk/data/documents/series/49486/</a>											
2	//SON_parameter_mapping											
3	//subject>SON:LOCAL:Chronological Julian Date</subject><object>SON:P011::CJ(Y1101)</object><units>SON:P061::UTAA</units>											
4	//subject>SON:LOCAL:CurDir</subject><object>SON:P011::COAE01</object><units>SON:P061::UBB</units>											
5	//subject>SON:LOCAL:CurrSpd</subject><object>SON:P011::CSAE01</object><units>SON:P061::UVB</units>											
6	//subject>SON:LOCAL:Temp</subject><object>SON:P011::TEMP001</object><units>SON:P061::UPAA</units>											
7	//											
8												
9	Cruise	Station	Type	yyyy-mm-Latitude	LOCAL_CCEDMO_co-Bot.Depth	Chronolog.QV	SEADeCurDir [d QV:SEADeCurDir]					
10	RCM.C	1185/C/B/C *		1983-08-2	1.4153	54.5855	49486	43	27	2445568	1	322
11										2445568	1	321
12										2445568	1	321
13										2445568	1	322
14										2445568	1	323
15										2445568	1	324
16										2445568	1	322
17										2445568	1	323
18										2445568	1	324
19										2445568	1	327
20										2445568	1	328
21										2445569	1	329



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## **Information elements required**

- Identified crucial elements for mentioned applications:
  - Survey *(ABCD)*
  - LocationID *(DwC)*
  - Longitude *(OBIS)*
  - Latitude *(OBIS)*
  - WKTFootprint *(OBIS)*
  - MinimumDepth *(OBIS)*
  - MaximumDepth *(OBIS)*
  - EventID *(DwC)*
  - SamplingProtocol *(DwC)*
  - EventDateTime *(DwC)*
  - FieldNumber *(OBIS)*
  - SampleSize *(OBIS)*
  - ScientificName *(OBIS, DwC)*
  - ScientificNameID *(DwC)*
  - LifeStage *(OBIS, DwC)*
  - Sex *(OBIS, DwC)*
  - ObservedIndividualCount *(OBIS, DwC)*

## **Adapted parameter vocabulary**

- P01 not very efficient for describing biological community observations
  - Generates huge amount of P01 entries
    - ex: MEIF0033 -Abundance of *Daptonema spp.* (*WoRMS 2455*) per unit area of the bed by sieving and picking under an optical microscope
    - 1 meiobenthos core => > 150 # taxa x lifestages x gender (x different taxonomic precision)
  - Link to WoRMS, but:
    - Taxonomy changes continuously (WoRMS has the most up-to-date names list)
    - No way of performing QC on taxon list
    - No way of validating new entries

## Adapted parameter vocabulary

Solution:

- Using more generic P011 terms
  - *SDBIOL01 Abundance of unspecified biological entity per unit volume of the water body*
  - *SDBIOL02 Abundance of unspecified biological entity per unit area of the bed*
  - *SDBIOL03 Ash-free dry weight biomass of unspecified biological entity per unit area of the bed*
  - *SDBIOL04 Wet weight biomass of unspecified biological entity per unit volume of the water body*
  - *SDBIOL05 Wet weight biomass of unspecified biological entity per unit area of the bed*
- Include separate taxon, lifestage and gender fields

<u>ScientificName</u>	ScientificNameID	Sex	LifeStage
Scophthalmus maximus	urn:lsid:marinespe	Unknown	Adult



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## ***More generic semantic header***

- Include references to external vocabularies
    - For example ICES vocab server: <http://vocab.ices.dk/matrix>
  - Clarify subject, object and unit
  - Use Darwin Core compliant terminology
- 
- Proposal:

	<u>example1</u>	<u>example2</u>
• Name	matrix	density[ind/m <sup>2</sup> ]
• Reference	<a href="http://vocab.ices.dk/matrix">http://vocab.ices.dk/matrix</a>	/
• MeasurementType	/	<a href="http://www.seadatanet.org/urnurl/SDN:P011::SDBIOL02">http://www.seadatanet.org/urnurl/ SDN:P011::SDBIOL02</a>
• MeasurementUnit	/	<a href="http://www.seadatanet.org/urnu/SDN:P061::UPMS">http://www.seadatanet.org/urnu/ SDN:P061::UPMS</a>
• MeasurementMethod	/	/



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## ***More generic semantic header***

- //<Field Index="21"  
Name="Density[m<sup>2</sup>]“  
Description="Abundance of unspecified biological entity per unit area of the bed (individuals/m<sup>2</sup>)“  
MeasurementTypeReference="http://www.seadatanet.org/urnurl/SDN:P01::SDBIOL02"  
MeasurementUnitReference="http://www.seadatanet.org/urnurl/SDN:P061::UPMS”  
MeasurementMethodReference=""  
/>>

## ***Suggested format***

**HEADER**  
Describing columns

//Field 1 is ....; Unit, Type of measurement; method of measurement...  
//Field 2 is.....; Unit, Type of measurement; method of measurement....  
//....

**DATA TABLE**

Field 1	Field 2	Field 3	Field ...

## *Suggested format*

HEADER  
Describing columns

//Field 1 is ....; Unit, Type of measurement; method of measurement...  
//Field 2 is.....; Unit, Type of measurement; method of measurement....  
//....

Field 1	Field 2	Field 3	Field ...

DATA TABLE

17 core fields

# additional fields

From: SDN, OBIS, ICES,  
DwC, IOOSbiology, ...

Templates that include  
certain additional fields

# Core Fields

Field	BIODEF	SDN version	Origin
Core	Survey	Cruise	SDN ODV
Core	LocationID	Station	SDN ODV
Core	Longitude	Longitude[degrees_east]	SDN ODV
Core	Latitude	Latitude[degrees_north]	SDN ODV
Core	WKTFootPrint	Latitude1	CDI
Core	WKTFootPrint	Longitude1	CDI
Core	WKTFootPrint	Latitude2	CDI
Core	WKTFootPrint	Longitude2	CDI
Core	/	Bot. Depth (m)	SDN ODV
Core	MinimumDepth	MinimumObservationDepth (CDI)	CDI
Core	MaximumDepth	MaximumObservationDepth (CDI)	CDI
Core	EventID	localCDid	SDN ODV
Core	InstitutionCode	EDMOCode	SDN ODV
Core	SamplingProtocol	InstrumentType	CDI
Core	EventDate	vymmddThh:ss	ODV
Core	FieldNumber	FieldNumber	OBIS
Core	SampleSize	SampleSize	OBIS
Core	ScientificName	ScientificName	OBIS
Core	ScientificNameID	ScientificNameID	DwC
Core	Sex	Sex	OBIS
Core	LifeStage	LifeStage	OBIS
Core	ObservedIndividualCount	ObservedIndividualCount	OBIS

# Additional Fields



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## Examples/templates

- macrobenthos community data with density and biomass values.
- zooplankton community with samples from different depths
- demersal fish population data with densities for different size classes and individual fish measurements (examples of subsamples are included)
- pollutant concentrations in biota specimens

BIODEF1.1\_DemFish - Notepad

```
//<AdditionalFields>
//<Field Index="18" Name="SubSamplingCoefficient" Description="Proportion of the subsample compared to originally collected total sample in case of subsampling">
//<Field Index="19" Name="observedMinLengthInCm" Description="record minimum length in centimeters if measuring more than one individual for aggregate level">
//<Field Index="20" Name="observedMaxLengthInCm" Description="record maximum length in centimeters if measuring more than one individual for aggregate level">
//<Field Index="21" Name="observedIndividualLengthInCm" Description="record maximum length in centimeters if single individual is observed and measured (e.g. for fish)">
//<Field Index="22" Name="Density_per_m2" Description="Abundance of unspecified biological entity per unit area of the bed (individuals/m²)" Measurement="Count">
//</coreFields>
//
Survey LocationID Longitude Latitude WKTFootPrint EventID SamplingProtocol EventDate MinimumDepth MaximumDepth F
RV Pelagia/199402 Voordelta2 3,56706 51,80916 LINESTRING(3.56289 51.72211, 3.57123 51.89621) VD_1201 Beam Trawl; 5m width; 20cm high;
RV Pelagia/199402 Voordelta2 3,56706 51,80916 LINESTRING(3.56289 51.72211, 3.57123 51.89621) VD_1201 Beam Trawl; 5m width; 20cm high;
RV Pelagia/199402 Voordelta2 3,56706 51,80916 LINESTRING(3.56289 51.72211, 3.57123 51.89621) VD_1201 Beam Trawl; 5m width; 20cm high;
RV Pelagia/199402 voordelta2 3,56706 51,80916 LINESTRING(3.56289 51.72211, 3.57123 51.89621) VD_1201 Beam Trawl; 5m width; 20cm high;
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RV Pelagia/199402 Voordelta3 3,56732 51,80632 LINESTRING(3.56390 51.79867, 3.56988 51.80866) VD_1202 Beam Trawl; 5m width; 20cm high;
RV Pelagia/199402 Voordelta3 3,56732 51,80632 LINESTRING(3.56390 51.79867, 3.56988 51.80866) VD_1202 Beam Trawl; 5m width; 20cm high;
RV Pelagia/199402 Voordelta3 3,56732 51,80632 LINESTRING(3.56390 51.79867, 3.56988 51.80866) VD_1202 Beam Trawl; 5m width; 20cm high;
RV Pelagia/199402 Voordelta3 3,56732 51,80632 LINESTRING(3.56390 51.79867, 3.56988 51.80866) VD_1202 Beam Trawl; 5m width; 20cm high;
```

## ***Important differences with ODV***

- DwC & OBIS terminology for core fields
- High number of fixed fields
- Includes text fields
- Date time: iso 8601; extended possibilities
- Trajectory <> track of trawl
- Vertical profile <> vertical net haul plankton
- Header can include external references and vocabularies

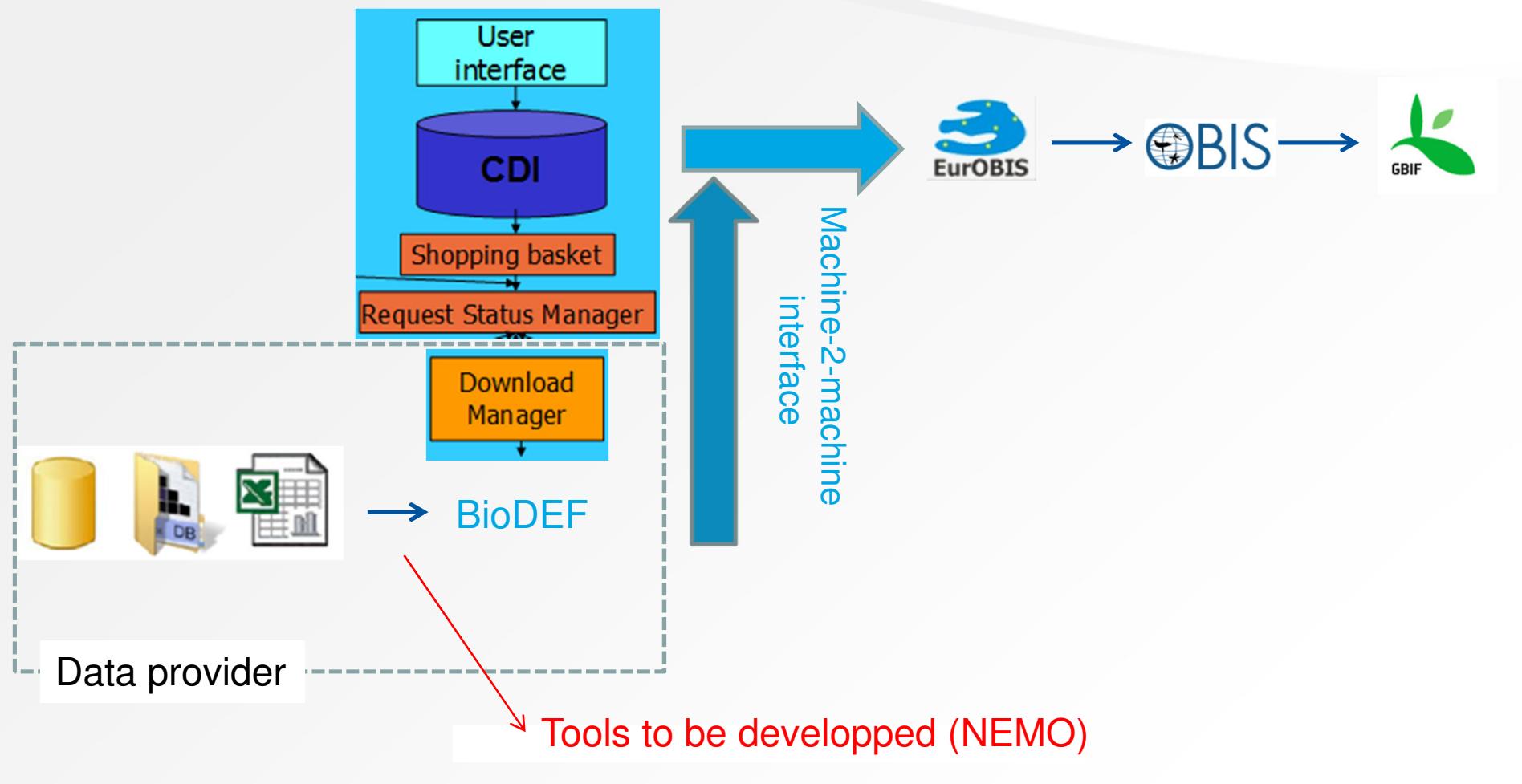
=> Increasing compatibility with SDN ODV



SeaDataNet

PAN-EUROPEAN INFRASTRUCTURE  
FOR OCEAN & MARINE DATA  
MANAGEMENT

## Data exchange protocols





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# Tools

- Ocean Data Viewer
- Taxonomic QC: WoRMS and LifeWatch taxon match tools
- Spatial QC: Geographic services
- Outlier occurrences (spatial and physico chemical)
- BioVEL data processing workflows (index calculations, species distr. maps,

Swedish LifeWatch - An analysis portal for biodiversity data

Overview Data sources Filter Calculation Format Result

Data sources: Species observation, Environmental data

Filter (Species observation search criteria): Occurrence, Taxa, Spatial

Calculation (data processing): Summary statistics, Grid statistics

Format (presentation settings): Table columns

Result: Maps, Tables, Diagrams

LIFEWATCH Register portal

New job Manual Changelog Results

Select one of the demo data files and choose from several web services, models and applications to work with other data files, please log in.

If you are new to this service, please read the manual.

File Browse... Use demo file: Marine View demo file Allowed filetypes: Plain text [TXT] Maximum rows in file: 10000

Row delimiter Return & linefeed (CRLF) First row contains column names

Column delimiter Tab

Decimal symbol Point(.)

Available services:

Servicetype	Name	Source	Description
Data validation and QC services			
Marineregions gazetteer services			
Taxon services			
Tidal services			
Geographical services - Administrative boundaries			
Geographical services - Bathymetry			
Geographical services - Biogeographical classification			
Geographical services - Features			
Geographical services - Protected areas			
Geographical services - Total biological valuation			

**BioVeL**  
Biodiversity Virtual e-Laboratory

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Workflows \ About Workflows

**About Workflows**

The quantity and heterogeneity of data in the biodiversity sciences have given rise to many distributed resources. Typically, researchers wish to combine these resources into multi-step computational tasks for a range of analytical purposes. Workflows, made of modularised units that can be repeated, shared, reused and repurposed, offer a practical solution for this task.

**RUN**

Workflows are executed through the BioVeL portal, a simple web interface that provides access to a pool of ready-made workflows and allows you to manage, share and save workflow results. You can monitor and interact with running workflows through the portal, changing parameters and directing your analyses.

**Data selection using BioStitch service through BioVeL Portal**

**Events**

**BioVeL's Meetings**

- European Biodiversity Informatics Conference, Italy, September 3-6, 2013
- BioVeL Third Annual Meeting and General Assembly, Budapest, Hungary, October 1-4, 2013

**BioVeL at Conferences**

- EGI Technical Forum, September 16-20, 2013, Madrid, Spain

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## **Outlook**

- Working towards more compatibility with SDN ODV
- Testing data submission in the framework of EMODNet Biology
- Tools to be developed/adapted
- Extensive training session in SDN training course in Ostend May 2014



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Thank You !

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