

# Taller GBIF.ES: Estandarización, documentación y publicación de datos de seguimientos de biodiversidad

28 – 30 de octubre de 2025, Universidad de Sevilla

## Estándares para la información de la biodiversidad

Francisco Pando  
GBIF España



# Guión

- Datos y ciencia
- Ciencia abierta y estándares
- Tipos de estándares
- TDWG
- Estándares en GBIF
  - Darwin Core
  - Humboldt Core
  - Plinian Core
- Darwin Core Archive
- GBIF new data model
- Darwin Core – DP





# Elementos que hacen la ciencia abierta

- Metadatos
- **Estándares**
- Identificadores
- Licencias
- Citación
- El cómo se sirve el dato

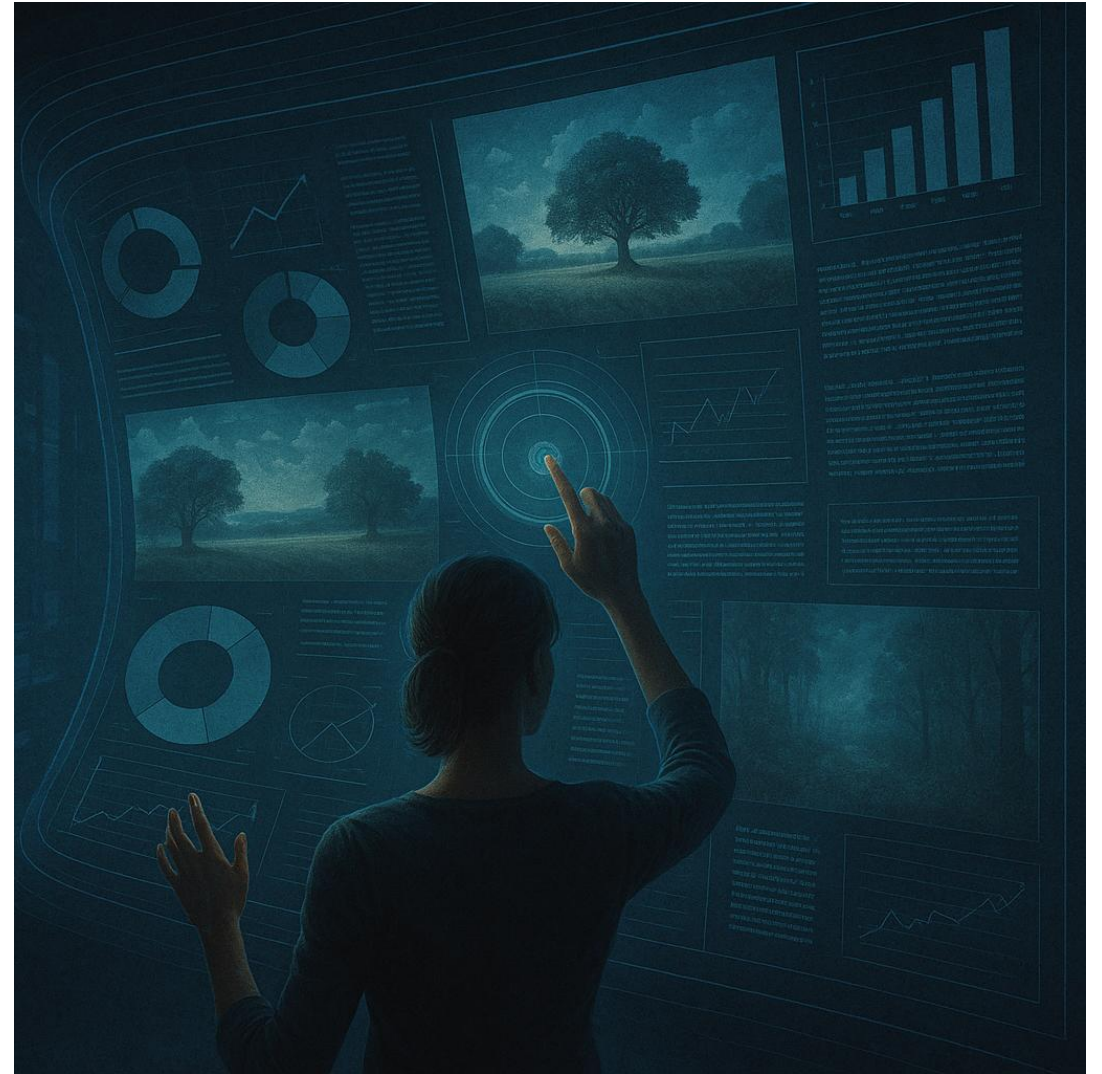
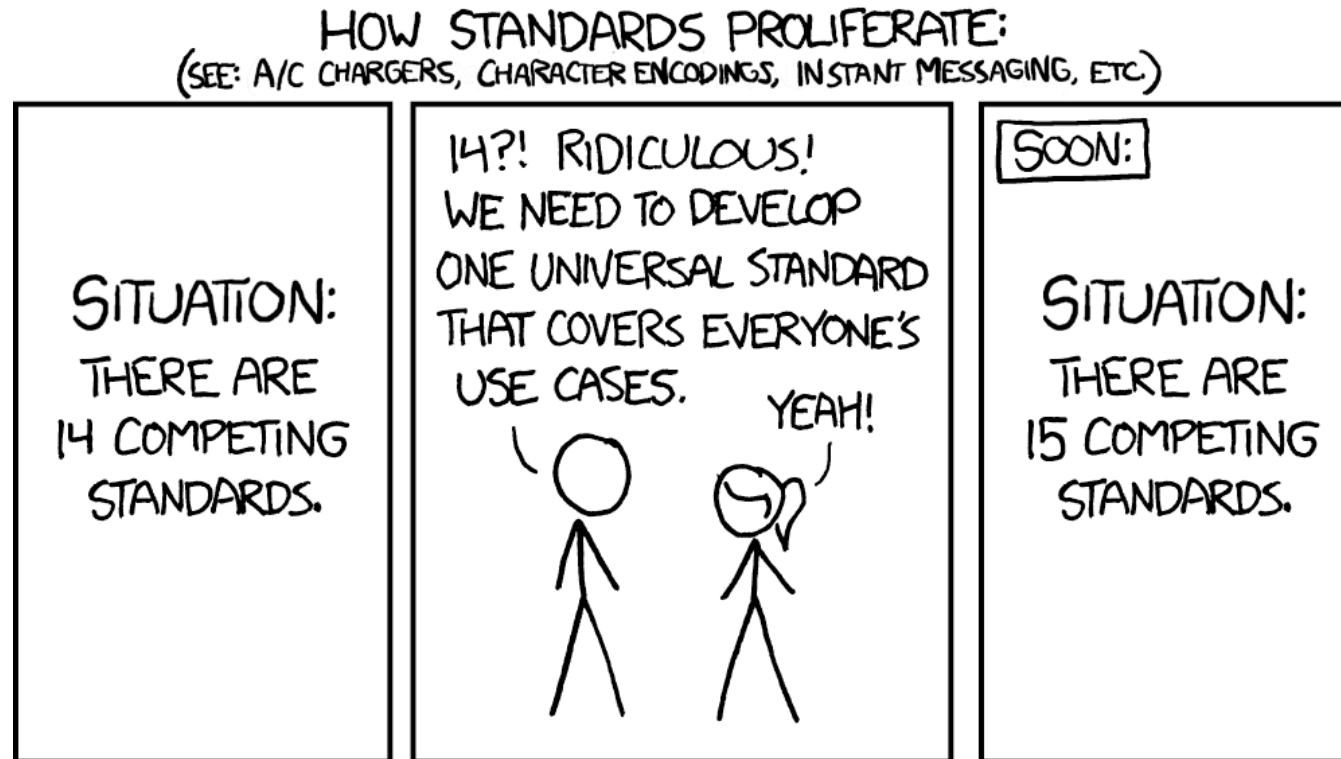


Imagen generada con  
IA

# Estándares



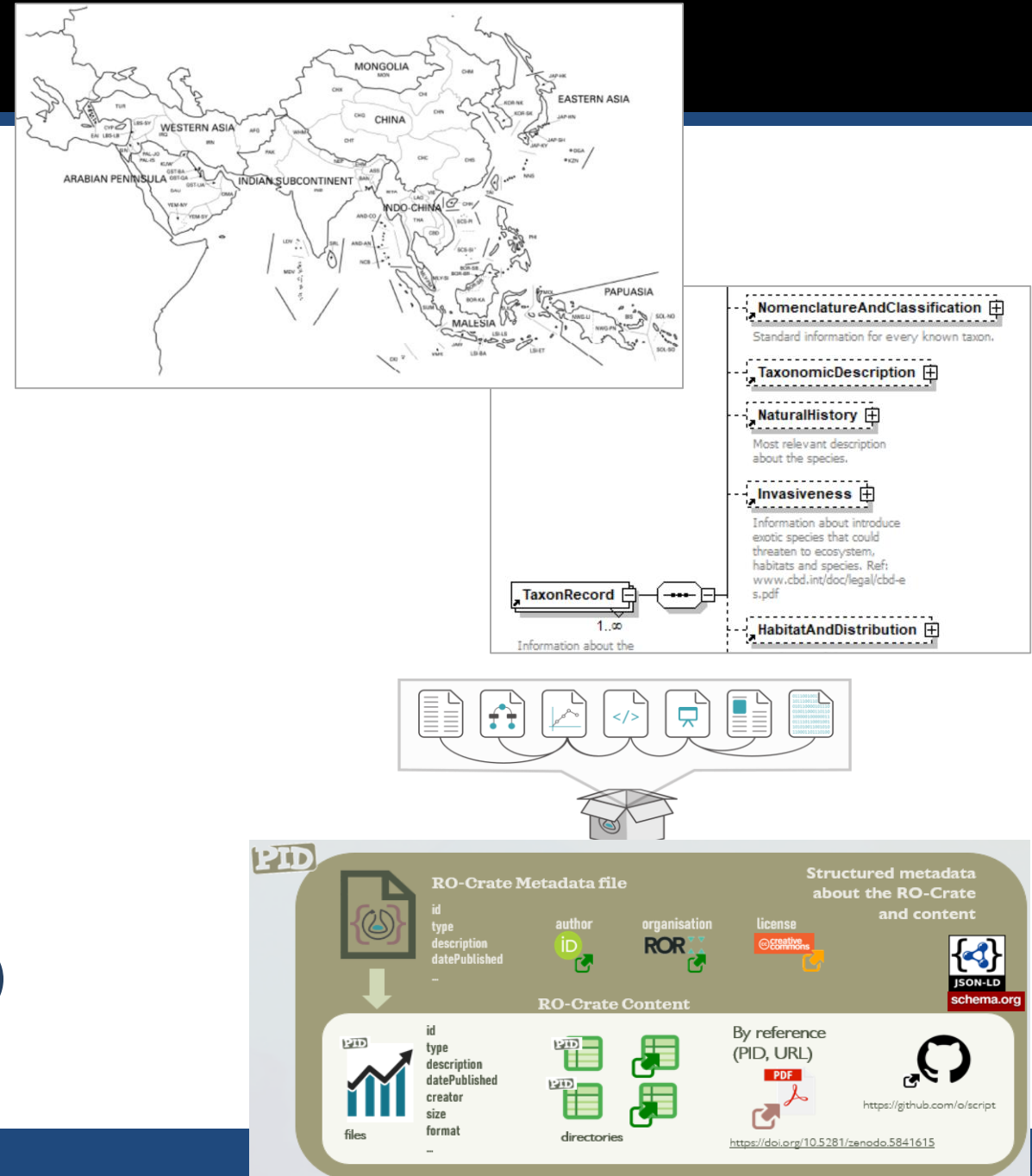
- Si lo cambias, ya no es un estándar
- Si solo lo usas tu, no es un estándar
- Los estándares son dinámicos y están, en constante evolución: revisiones y ediciones

<https://xkcd.com/927/>

“pensaba que después del COVID no tendría que contar estas cosas...”

# Estándares

- Vocabularios controlados (léxicos)
  - Que valores son válidos
  - (listas de géneros, de países...)
- Definiciones (semánticos)
  - para humanos, para máquinas
  - (v.gr.: Darwin Core [terms])
- Estructurales (sintácticos)
  - Como se codifica la información
  - (v.gr.: Frictionless Data Packages)



# “Estándares orquestados”

- CC es un vocabulario controlado
- Dublin Core es un estándar de términos
- XML es un estándar sintáctico

```
Xml                                                                    Cop
<metadata>
  <dc:title>Example Document</dc:title>
  <dc:creator>John Doe</dc:creator>
  <dc:subject>Metadata Example</dc:subject>
  <dc:description>This is an example of a metadata section in XML.</dc:des
  <dc:publisher>Example Publisher</dc:publisher>
  <dc:date>2025-01-23</dc:date>
  <dc:type>Text</dc:type>
  <dc:format>application/pdf</dc:format>
  <dc:identifier>http://example.com/document</dc:identifier>
  <dc:language>en</dc:language>
  <dc:rights>Creative Commons Attribution-NonCommercial 4.0 International
</metadata>
```

# Biodiversity Information Standards (TDWG)

We are a non-profit organization and a community dedicated to  
developing biodiversity information standards.

<https://www.tdwg.org>

## ABCD

Access to Biological Collection Data (ABCD) Schema

## Audiovisual Core

Audiovisual Core Multimedia Resources Metadata Schema

## Darwin Core

Darwin Core

## GUID Applicability Statements

GUID and Life Sciences Identifiers Applicability Statements

## Latimer Core

Latimer Core, a standard for describing collections

## SDS

TDWG Standards Documentation Standard (SDS)

## TAPIR

TDWG Access Protocol for Information Retrieval (TAPIR)

## TCS

Taxon Concept Schema (TCS)

## VMS

Vocabulary Maintenance Standard (VMS)

## WGSRPD

World Geographical Scheme for Recording Plant Distributions (WGSRPD)

## Interest and Task Groups

- MG** Access to Biological Collections Data
- IG** Annotations
- IG** Attribution
  - TG** People in Biodiversity Data
- MG** Audiovisual Core
  - TG** 3D Imagery and Data
  - TG** Views Controlled Vocabularies
- IG** Biodiversity Data Quality
  - TG** Framework on data quality
  - TG** Data quality tests and assertions
  - TG** Data quality use cases
  - TG** Best practices for development of vocabularies of values ("Vocabularies")
- IG** Biodiversity Informatics Curriculum
- IG** Biodiversity Services and Clients
- IG** Biological Interactions Data
- IG** Citizen Science
- IG** Collection Descriptions
  - MG** Latimer Core
  - TG** Minimum Information about a Digital Specimen (MIDS)
  - TG** Modelling Research Expeditions
- MG** Darwin Core
  - TG** Invasive Species
  - TG** Plant Phenology
  - TG** Realm and Biome
  - TG** Sensitive Species Extension
- IG** Earth Sciences and Paleobiology
  - TG** Darwin Core Chronometric Age Extension
  - TG** Extension for Geosciences (EFG)
  - TG** Mineralogy Extension
- IG** Genomic Biodiversity
  - TG** Environmental Samples and eDNA
  - TG** Sustainable DarwinCore MixS Interoperability
- IG** Geographical Schemes
  - TG** Marine Geo Schemes Task Group
  - TG** Terrestrial GeoSchemes TG
- IG** Machine Observations
- IG** Observations & Specimens
  - TG** How Did It Die?
  - TG** Humboldt Extension
  - TG** Material Sample
- IG** Process
- IG** Species Information
  - TG** Invasive Organism Information
  - TG** Plinian Core
- MG** Taxon Concept Schema (TCS) Maintenance Group
  - TG** Taxonomic Concept Schema (TCS2) Task Group
- IG** Technical Architecture Group
  - TG** Standards Mapping Task Group



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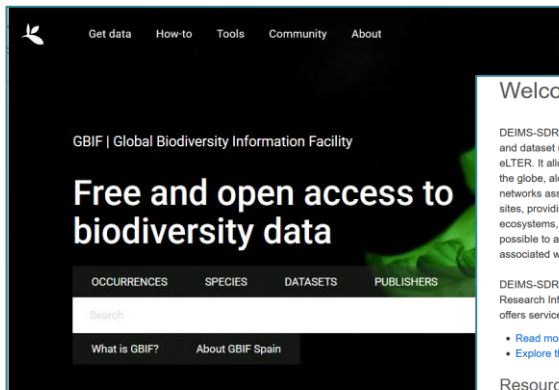
- MG Access to Biological Collections Data
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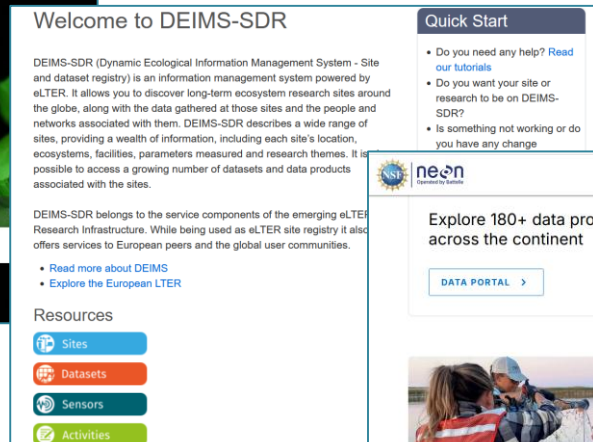
# EML (Ecological Metadata Language)

>> estándar de código abierto y basado en [XML](#), diseñado para documentar datos de investigación en ciencias ambientales y ecológicas. Permite describir el contenido, alcance espacial y temporal, métodos y protocolos de los datos

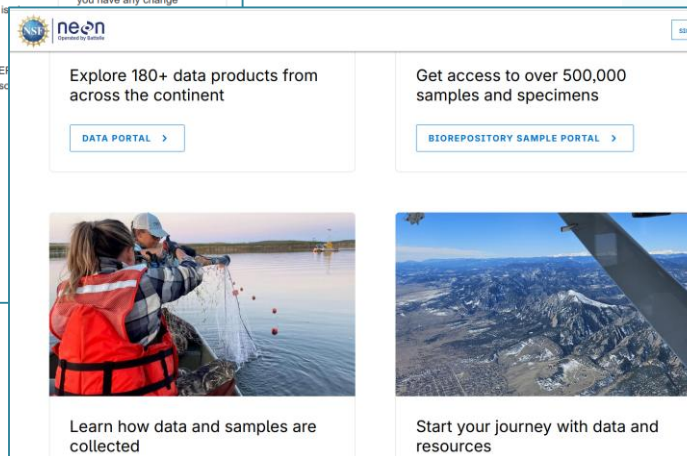
<https://www.gbif.org>



<https://deims.org>

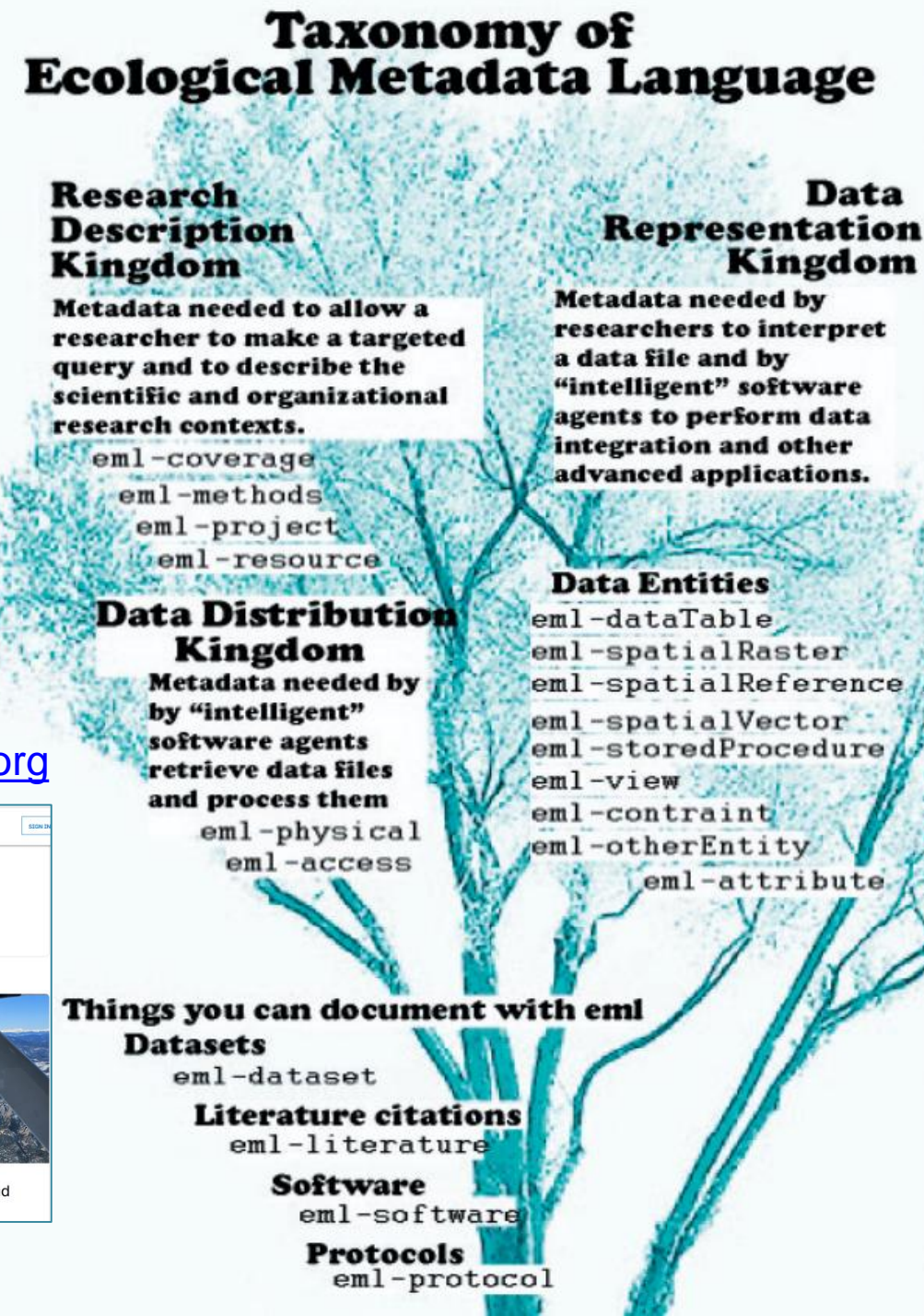


[www.neonscience.org](http://www.neonscience.org)



<https://eml.ecoinformatics.org/>

<https://lternet.edu/wp-content/uploads/2010/12/EMLHandbook-2.pdf>



# Darwin Core



TDWG

Darwin Core

## Darwin Core Quick Reference Guide

<https://dwc.tdwg.org/terms/>

Record-level  
Occurrence  
Organism  
MaterialEntity  
MaterialSample  
Event  
Location  
GeologicalContext  
Identification  
Taxon  
MeasurementOrFact  
ResourceRelationship  
UseWithIRI  
LivingSpecimen  
PreservedSpecimen  
FossilSpecimen  
MaterialCitation  
HumanObservation  
MachineObservation

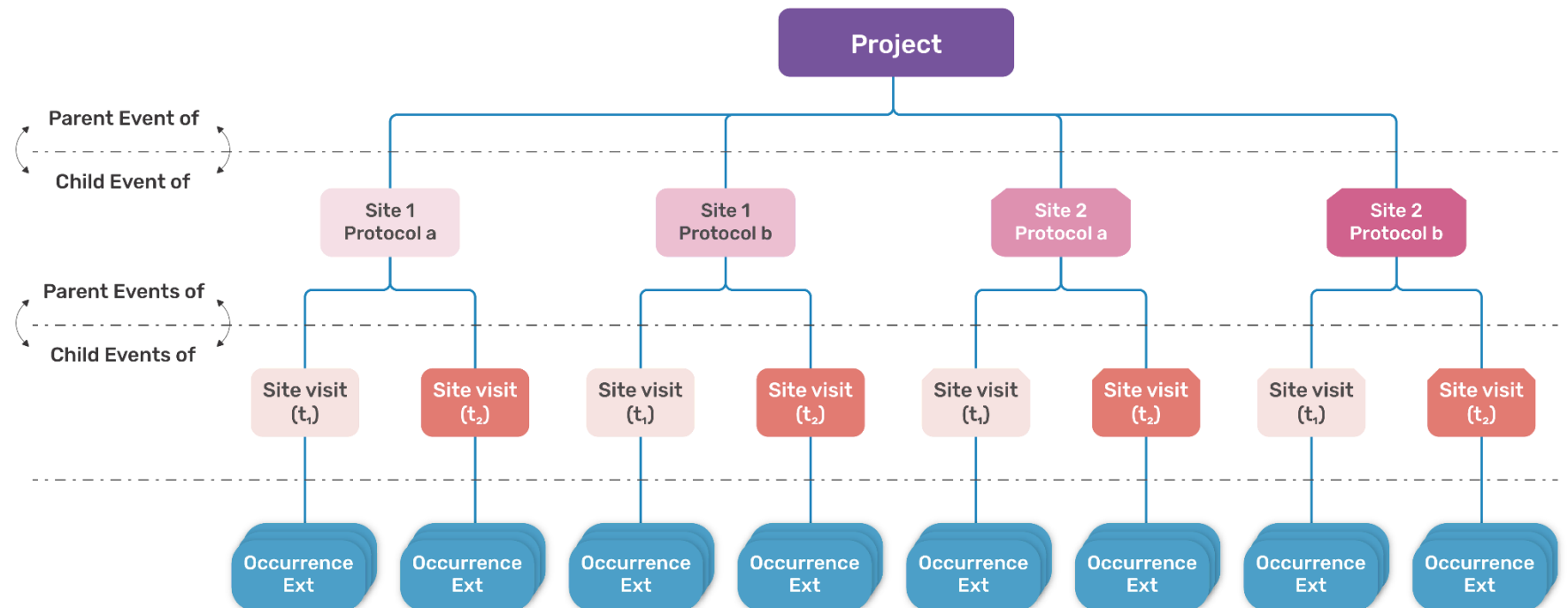
Record-level Terms	Dublin Core terms, institutions, collections, nature of data record	Simple Darwin Core (flat)
Occurrence	evidence of species in nature, observers, behavior, associated media, references.	
Event	sampling protocols and methods, date, time, field notes	
Location	geography, locality descriptions, spatial data	
Identification	linkage between Taxon and Occurrence	
Taxon	scientific names, vernacular names, names usages, taxon concepts, and the relationships between them	
GeologicalContext	geologic time, chrono-stratigraphy, biostratigraphy, lithostratigraphy	
ResourceRelationship	explicit relationships between identified resources (e.g., one organism to another, taxon to location, etc.)	Generic Darwin Core (relational)
MeasurementOrFact	measurements, facts, characteristics, assertions, references	

[https://www.researchgate.net/figure/Darwin-Core-Categories-Simple-Darwin-Core-is-comprised-of-seven-categories-of-terms\\_fig3\\_221741592](https://www.researchgate.net/figure/Darwin-Core-Categories-Simple-Darwin-Core-is-comprised-of-seven-categories-of-terms_fig3_221741592)

# Humboldt Core

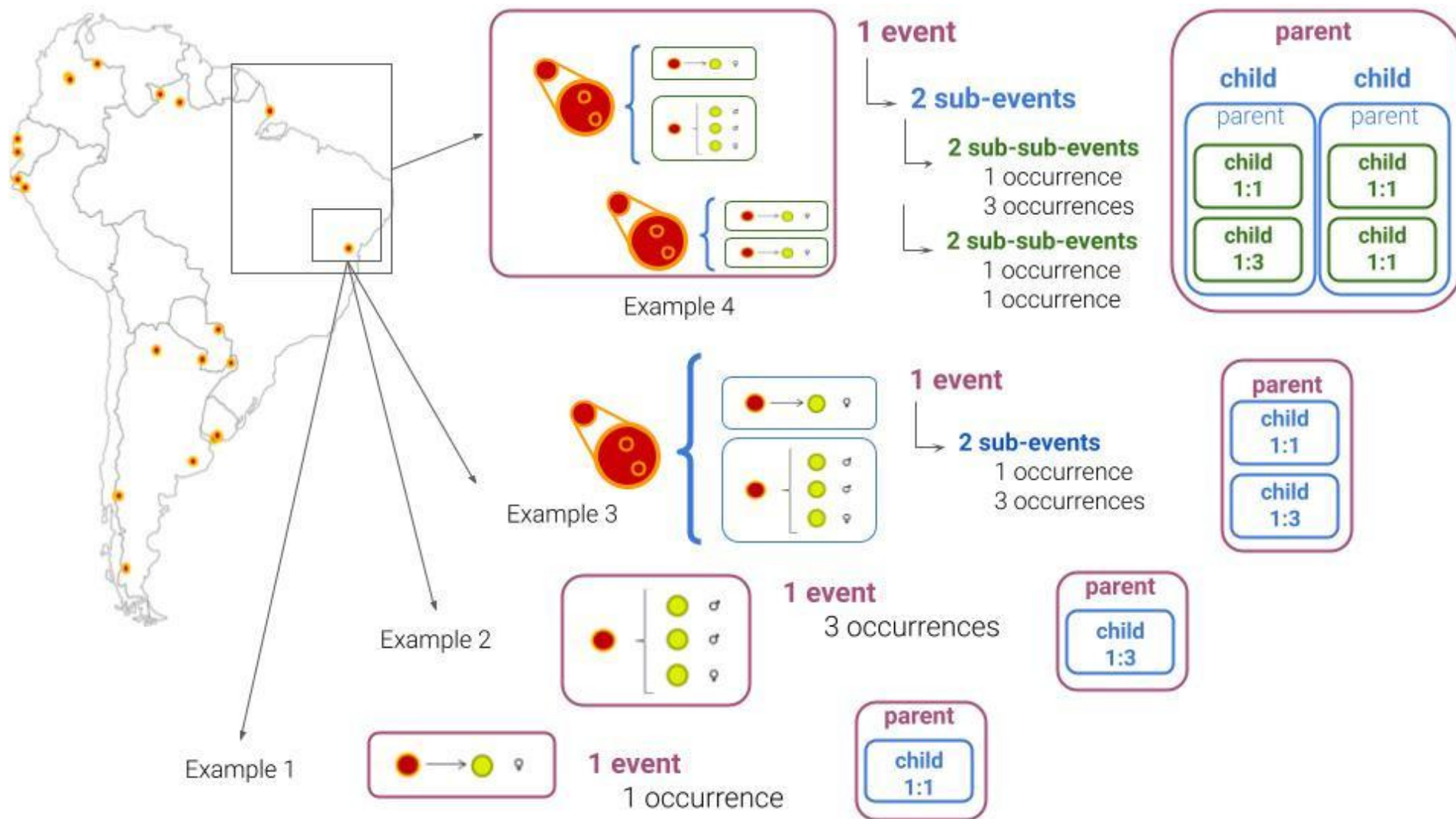
Es una extensión del estándar Darwin Core (DwC) es un vocabulario estructurado (términos y definiciones) desarrollada para mejorar la descripción de inventarios y muestreos estructurados de biodiversidad. Sirve para describir y documentar datos tanto procedentes de muestreos tradicionales como los recogidos por redes de sensores-

- 55 elementos
- Jerárquico
- Ratificado por TDWG en 2024
- <https://eco.tdwg.org/HumboldtExtensionforEcologicalInventories:UserGuide>
- <https://docs.gbif.org/survey-monitoring-quick-start/en>





# Humboldt Core



Visual representation of a biodiversity inventory illustrating the four examples of inventory data ranging from simple, individual occurrences (Example 1) to more complex, hierarchically structured data



# Humboldt Core

- [Site](#)
- [Habitat Scope](#)
- [Temporal Scope](#)
- [Taxonomic Scope](#)
- [Organismal Scope](#)
- [Methodology Description](#)
- [Material Collected](#)
- [Sampling Effort](#)
- [UseWithIRI](#)

## Site

siteCount siteNestingDescription verbatimSite

geospatialScopeAreaValue geospatialScopeAreaUnit

totalAreaSampledUnit reportedWeather reportedWeatherUnit

## Organismal Scope

targetLifeStageScope excludedLifeStageScope isLifeStageScopeFullyReported

targetDegreeOfEstablishmentScope excludedDegreeOfEstablishmentScope

isDegreeOfEstablishmentScopeFullyReported targetGrowthFormScope

excludedGrowthFormScope isGrowthFormScopeFullyReported

hasNonTargetOrganisms verbatimTargetScope

## Habitat Scope

targetHabitatScope excludedHabitatScope

## Temporal Scope

eventDurationValue eventDurationUnit

## Methodology Description

compilationTypes compilationSourceTypes inventoryTypes protocolNames

protocolDescriptions protocolReferences isAbundanceReported

isAbundanceCapReported

isLeastSpecificTargetCategory

## Material Collected

hasVouchers voucherInstitutions hasMaterialSamples materialSampleTypes

## Taxonomic Scope

targetTaxonomicScope excludedTaxonomicScope taxonCompletenessReported

taxonCompletenessProtocols isTaxonomicScopeFullyReported isAbsenceReported

absentTaxa hasNonTargetTaxa nonTargetTaxa areNonTargetTaxaFullyReported

## Sampling Effort

samplingPerformedBy isSamplingEffortReported samplingEffortProtocol

samplingEffortValue samplingEffortUnit

# Plinian Core

- Compartir información a nivel de especies (un estándar de intercambio).
- Interoperabilidad: expresado en xml, se apoya en otros estándares
- Información = propiedades y traits relativos al taxón: descripciones, nomenclatura, estado de conservación, manejo, historia natural..
- Este proyecto ha sido liderado por:
  - El Instituto Nacional de Biodiversidad de Costa Rica (INBio, Costa Rica)
  - Nodo Español de GBIF (GBIF, España)
  - La Universidad de Granada (UG, España)
  - El Instituto Alexander von Humboldt (IAvH, Colombia)
  - La Comisión Nacional del Conocimiento y el Uso de Biodiversidad de México (CONABIO)
  - La Universidad de Sao Paulo y SiBBR Brazil)
- Auto-contenido y con capacidad de dar soporte a la integración de datos en múltiples bases de datos, con diferentes niveles de granularidad.
- Clases: 13 Genéricas, 27 Complejas y 22 Simples.
- Adoptado por el Ministerio de Agricultura..... Y Medio Ambiente de España como estándar
- Parte de los procesos de la organización “Biodiversity Information Standards” TDWG:  
<http://www.tdwg.org/>



tdwg / PlinianCore



<> Code



Issues

5



Pull requests



Discussions



Actions



Projects



Wiki



PlinianCore

Public



Edit Pins



Unwatch

32



Fork

5

# Plinian Core

Pages 236

Plinian Core is a standard oriented to share species level information. Its hierarchical schema allows --for instance-- to develop species pages or data sheets in websites.

[About Plinian Core](#)

## Abstract Model [Current v3.2.2.7]

Available at: [xsd abstract-model stable-version](#)

In order to validate the abstract model schema, some additional schemas are needed. These are also available at the same location.

## Application profiles already defined:

It's important to know how we can deploy an Application Profile. If we want to check an AP, we have to download, in the same folder where we have our AP, all the files that are located in this link: [xsd abstract-model stable-version](#). Then, when we open the AP in the editor, e.g: Altova, this one will validate the AP

### Plinian Core terms documentation



- [Home](#)
- [Plinian Core Terms Index](#)
- [Plinian Core Terms arranged hierarchically](#)
  - [Metadata](#)
  - [TaxonRecord](#)
    - [BaseElements](#)
    - [RecordMetadata](#)
    - [NomenclatureAndClassification](#)
    - [TaxonomicDescription](#)
    - [NaturalHistory](#)
    - [Invasiveness](#)
    - [HabitatAndDistribution](#)
    - [DemographyAndThreat](#)

<https://github.com/tdwg/PlinianCore>

# Términos

## Term index

This quick guide provides a list of all current elements of the Plinian core. elements are organized by categories (in bold) in the index. The categories correspond to Plinian Core elements that are classes (elements that have elements to describe them). The elements that describe a given class (the properties) appear in the list immediately below the name of the category. The index provides links to the element descriptions in the table below the

### Metadata

[Dataset\\_ID](#) | [dateStamp](#) | [citation](#) | [eml-dataset](#) | [resourceLogoUrl](#) | [Refer](#)

### Base Elements

[TaxonRecordID](#) | [taxonConceptID](#) | [GlobalUniqueIdentifier](#) | [Abstract](#)

### Record Metadata

[Language](#)  
([TargetAudiences](#)) > [Audience](#) | [AudiencesUnstructured](#)  
([Version](#)) > [Major](#) | [Minor](#) | [Modifier](#) | [DateIssued](#) | [PreferredFlag](#)  
([Revision](#)) > [associatedParty](#) | [pubDate](#) | [created](#)

### Nomenclature and Classification

([TaxonRecordName](#)) > [ScientificName](#)  
([Synonyms](#)) > [SynonymName](#) | [SynonymStatus](#) | [SynonymsUnstructured](#)  
([CommonNames](#)) > [Name](#) | [Language](#) | [UsedIn](#) | [UsedBy](#) | [CommonNamesUnstructured](#)  
([Hierarchy](#)) > [kingdom](#) | [phylum](#) | [class](#) | [order](#) | [family](#) | [genus](#) | [subgenus](#) | [specificEpithet](#) | [infraspecificEpithet](#) | [higherClassification](#) | [ParentTaxon](#)  
([MiscDetails](#)) > [MeasurementOrFact](#) | [DetailUnstructured](#)  
[NomenclatureAndClassificationUnstructured](#)

### Taxonomic Description

[BriefDescription](#)  
([FullDescription](#)) > [MeasurementOrFact](#) | [FullDescriptionUnstructured](#)  
([IdentificationKeys](#)) > [Keys](#)  
[TaxonomicDescriptionUnstructured](#)

### Natural History

([LifeForm](#)) > [MeasurementOrFact](#) | [LifeFormUnstructured](#)  
([LifeCycle](#)) > [MeasurementOrFact](#) | [LifeCycleUnstructured](#)  
([Reproduction](#)) > [MeasurementOrFact](#) | [ReproductionUnstructured](#)  
([AnnualCycles](#)) > [Event](#) | [starTimeInterval](#) | [endTimeInterval](#) | [AnnualCyclesUnstructured](#)  
([Feeding](#)) > [Strategy](#) | [StrategyRemarks](#) | [FeedingUnstructured](#)  
([Dispersal](#)) > [Purpose](#) | [Type](#) | [StructureDispersed](#) | [Distance](#) | [DispersalUnstructured](#)  
([Behavior](#)) > [MeasurementOrFact](#) | [BehaviorUnstructured](#)  
([Interactions](#)) > [InteractionSpecies](#) | [InteractionSpeciesType](#) | [InteractionUnstructured](#)  
([MolecularData](#)) > [MeasurementOrFact](#) | [MolecularDataUnstructured](#)  
([Migratory](#)) > [Causes](#) | [Patterns](#) | [Routes](#) | [Season](#) | [MigratoryUnstructured](#)  
([EcologicalSignificance](#)) > [MeasurementOrFact](#) | [EcologicalSignificanceUnstructured](#)  
([MiscDetails](#))  
([EnvironmentalEnvelope](#)) > [MeasurementOrFact](#) | [EnvironmentalEnvelopeUnstructured](#)  
[NaturalHistoryUnstructured](#)

### Invasiveness

[origin](#) | [presence](#) | [persistence](#) | [distribution](#) | [harmful](#) | [modified](#) | [startValidDate](#) | [endValidDate](#) | [countryCode](#) | [stateProvince](#) | [county](#) | [localityName](#) | [language](#) | [citation](#) | [abundance](#) | [trend](#) | [rateOfSpread](#) | [regulatoryListing](#) | [memo](#) | [publicationDate](#) | [localityType](#) | [WhatImpact](#) | [vector](#) | [Route](#) | [impactTarget](#) | [impactMechanism](#)  
[InvasivenessUnstructured](#)

### Habitat and Distribution

([Habitats](#)) > [MeasurementOrFact](#) | [HabitatUnstructured](#)  
([Distribution](#)) > [DistributionScope](#) | [temporalCoverage](#) | [GeographicEntity](#) | [country](#) | [stateProvince](#) | [county](#) | [municipality](#) | [locality](#) | [DistributionUnstructured](#)  
([Endemic](#)) > [EndemicTo](#) | [EndemicIn](#) | [EndemicUnstructured](#)  
[HabitatAndDistributionUnstructured](#)

### Demography and Threat

([Territory](#)) > [ExtentOfOccurrence](#) | [AreaOfOccupancy](#) | [TerritoryUnstructured](#)  
([PopulationBiology](#)) > [AbundanceData](#) | [DensityData](#) | [PatternDistribution](#) | [Size](#) | [SexRatio](#) | [Fecundity](#) | [MortalityRate](#) | [BirthRate](#) | [NumberIndividualsPerObservation](#) | [AverageDensity](#) | [PopulationTrend](#) | [Recruitment](#) | [PopulationGrowthRate](#) | [Emigration](#) | [Immigration](#) | [DescriptionLifeStages](#) | [ProportionIndividualsPerStageLife](#) | [CarryingCapacity](#) | [PopulationBiologyUnstructured](#)  
([ThreatStatus](#)) > [ThreatCategory](#) | [Authority](#) | [AppliesTo](#) | [ThreatStatusUnstructured](#)  
([DirectThreats](#)) > [MeasurementOrFact](#) | [DirectThreatUnstructured](#)  
([Legislation](#)) > [LegislationName](#) | [ProtectionLegalStatus](#) | [LegislationRead](#) | [Status](#) | [Type](#) | [Norm](#) | [AppliesTo](#) | [LegislationUnstructured](#)  
[DemographyAndThreatUnstructured](#)

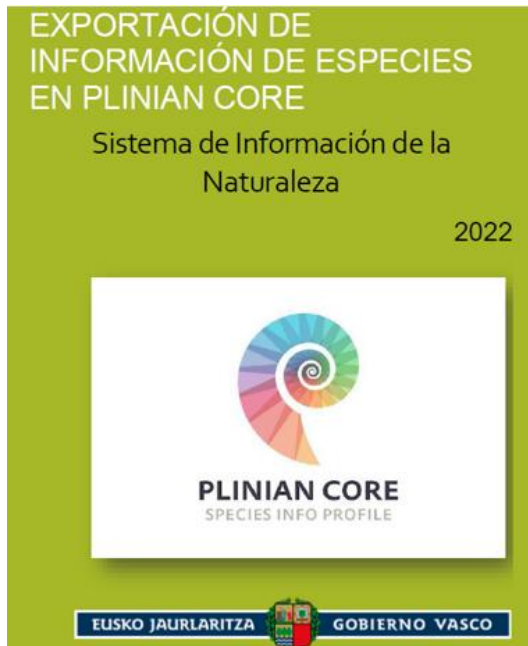
### Uses, Management and Conservation

([Uses](#)) > [SourceOfInformation](#) | [Use-Value](#) | [PartUsed](#) | [Users](#) | [Organisms](#) | [VernacularNameUseAnnotations](#) | [ProductionDetails](#) | [MeansOfApplication-Administration](#) | [SeasonOfAvailability-Use](#) | [Conservation-ExploitationData](#) | [UseTypeAtomized](#) | [Economics](#) | [RatingPopularity](#) | [Properties](#) | [Potential](#) | [UseNotes](#) | [UsesUnstructured](#)  
([ManagementAndConservation](#)) > [Objetives](#) | [ManagementPlan](#) | [Actions](#) | [HumanAndEnvironmentalrelevance](#) | [ManagementAndConservationUnstructured](#)  
[UsesManagementAndConservationUnstructured](#)

**[associatedParty](#), [MeasurementOrFact](#), [References](#), [AncillaryData](#)**



# Plinian Core is used



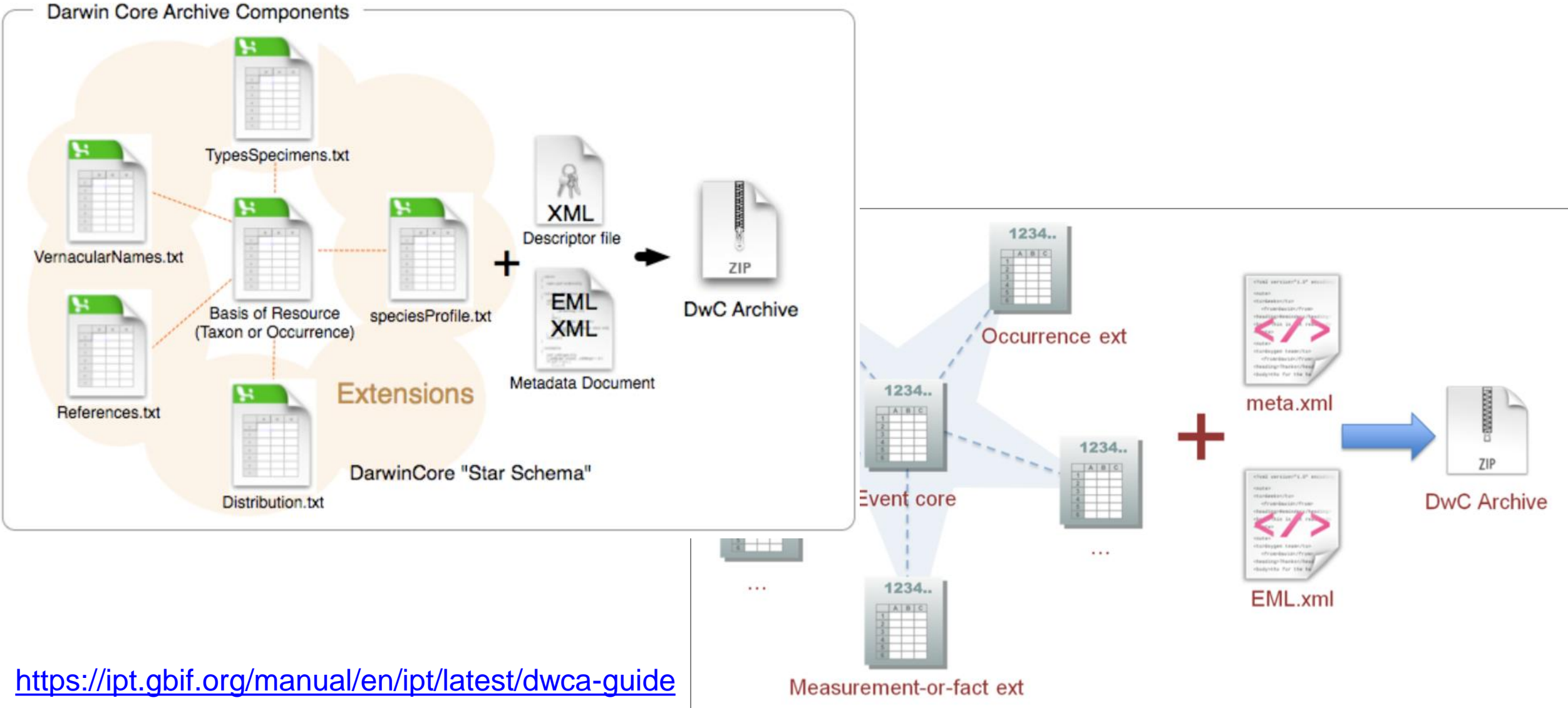
## Bienvenidos al inventario de especies silvestres

Esta página Web posee información tanto de historia natural como de gestión sobre especies silvestres nativas de Chile así como también algunas especies exóticas asilvestradas en nuestro territorio.

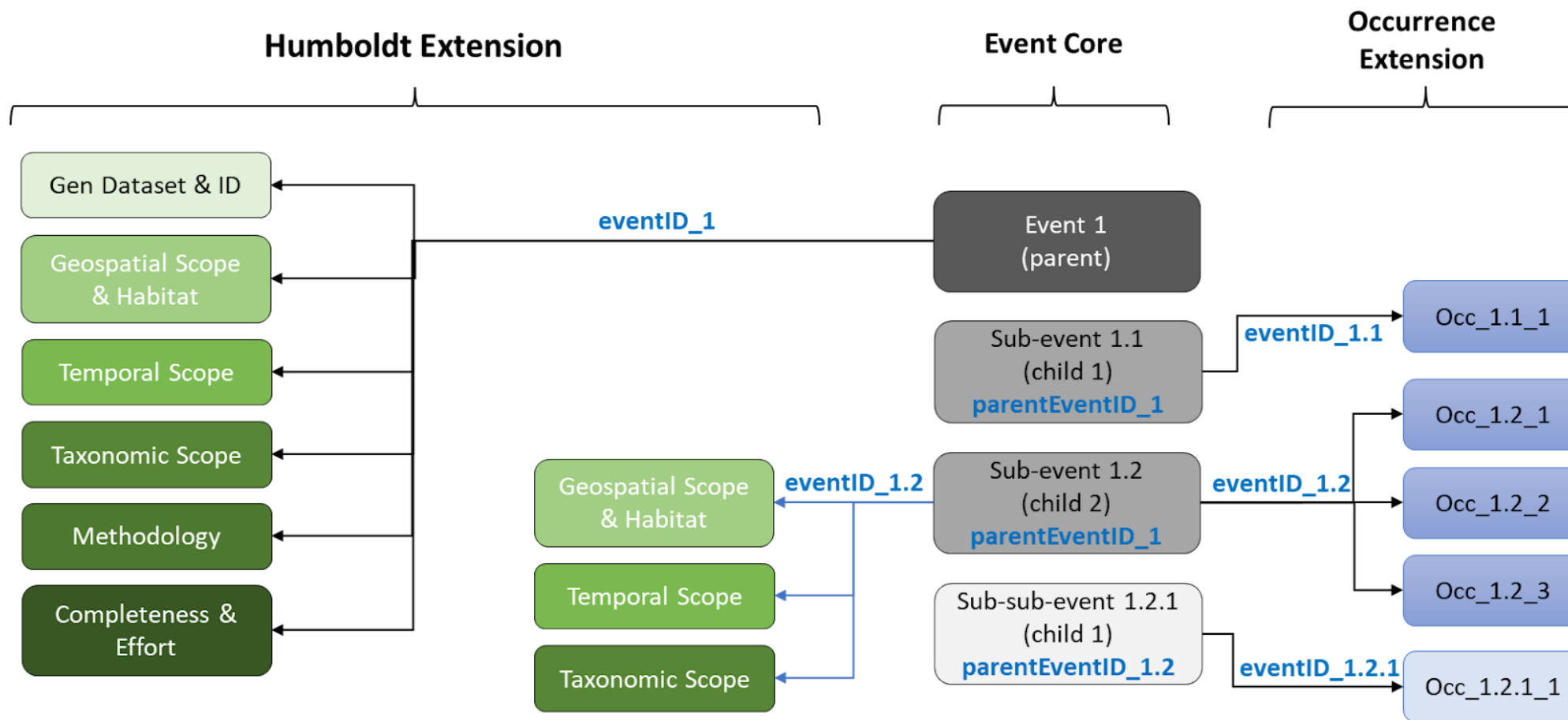
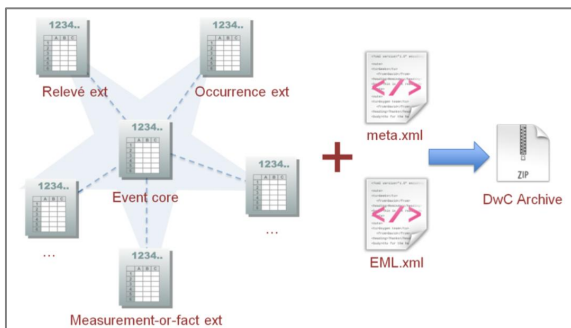
El Inventario Nacional de Especies cuenta solamente con un subconjunto del total de especies nativas descritas para nuestro país. Lo mismo para especies exóticas. Esperamos que al mediano plazo contemos con la totalidad de la flora y fauna nativa incorporada en este sitio.



# Darwin Core. Implementación: Darwin Core Archive



# Darwin Core Archive



Visual representation of a structured biodiversity inventory using the Darwin Core Event class and the Occurrence and Humboldt Extensions. Information on different hierarchies is linked via each unique dwc:eventID



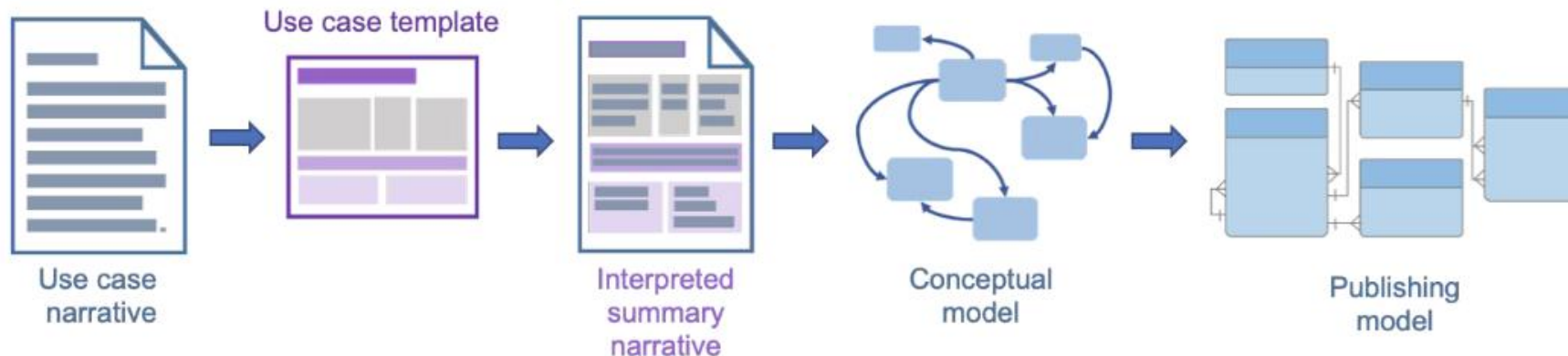
# Nuevo modelo conceptual para Darwin Core

>> define la estructura lógica y las relaciones entre los distintos elementos del estándar Darwin Core (DwC). Sobre el conjunto de términos o vocabulario controlado que es Darwin Core, el modelo conceptual describe cómo se conectan entre sí esos términos —es decir, la arquitectura conceptual que sustenta el intercambio de datos de biodiversidad.

“...biodiversity data is more complicated than ‘just’ the occurrence of species in time and space; organisms interact, co-occur, move and evolve.

This implies a need for richer and more varied types of data than GBIF has thus far supported.”

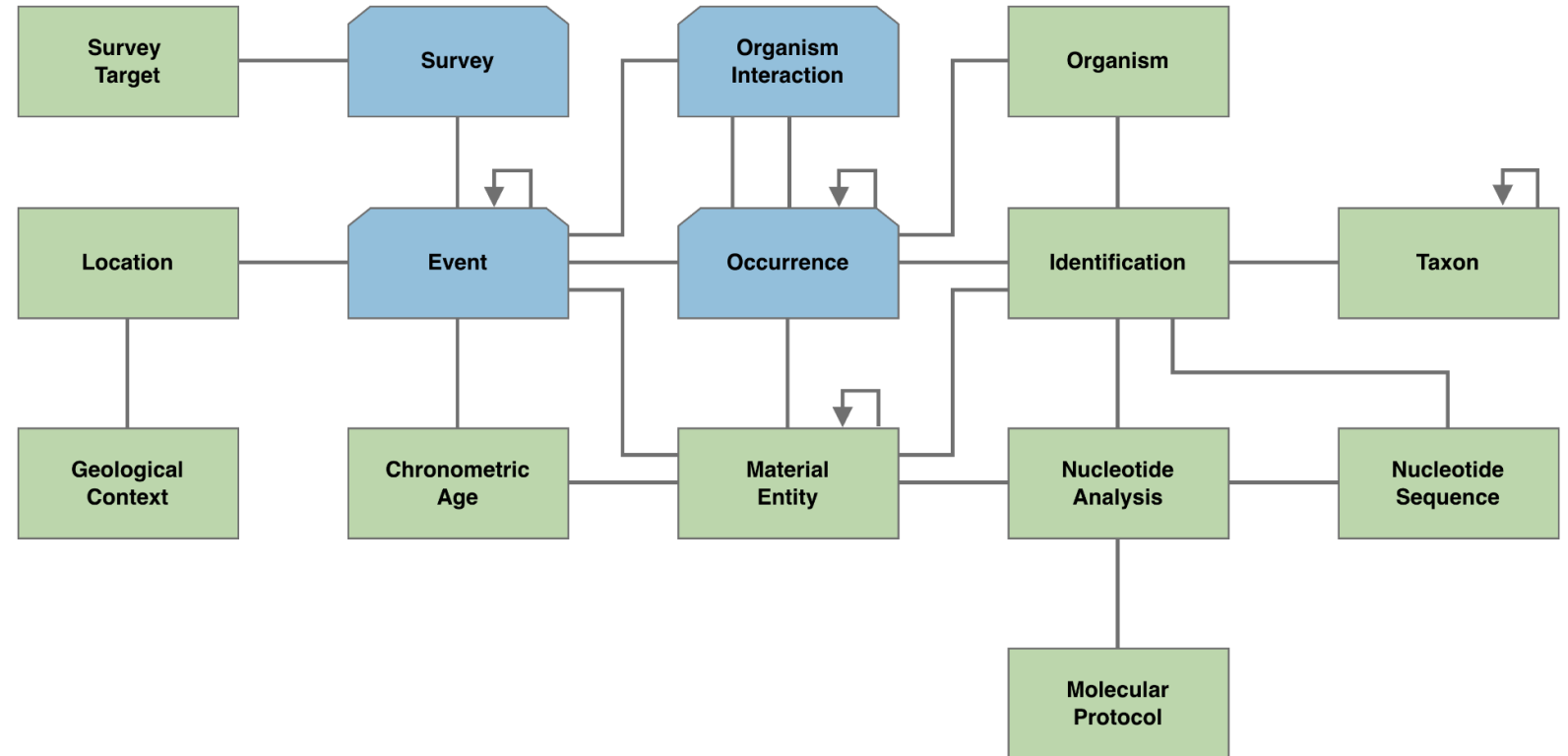
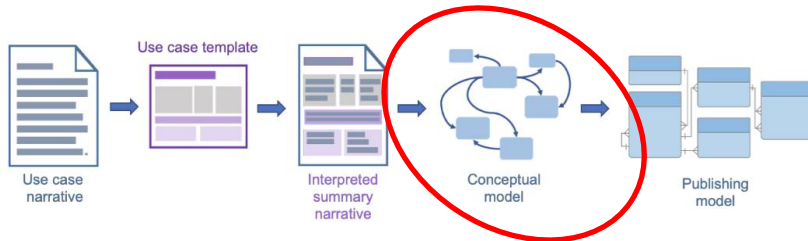
Heberling et al. (2021) Data integration enables global biodiversity synthesis <https://doi.org/10.1073/pnas.2018093118>



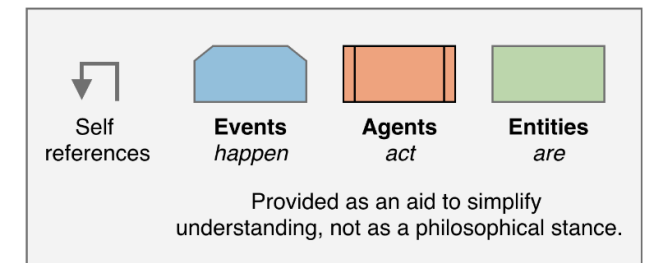
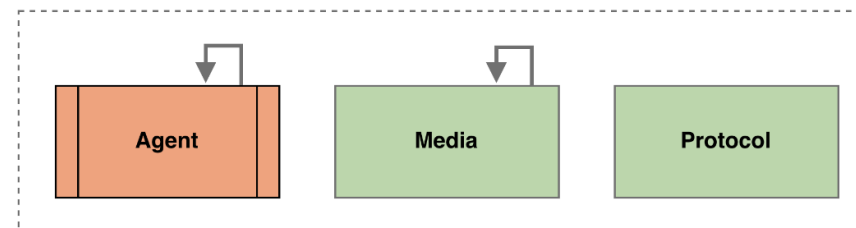


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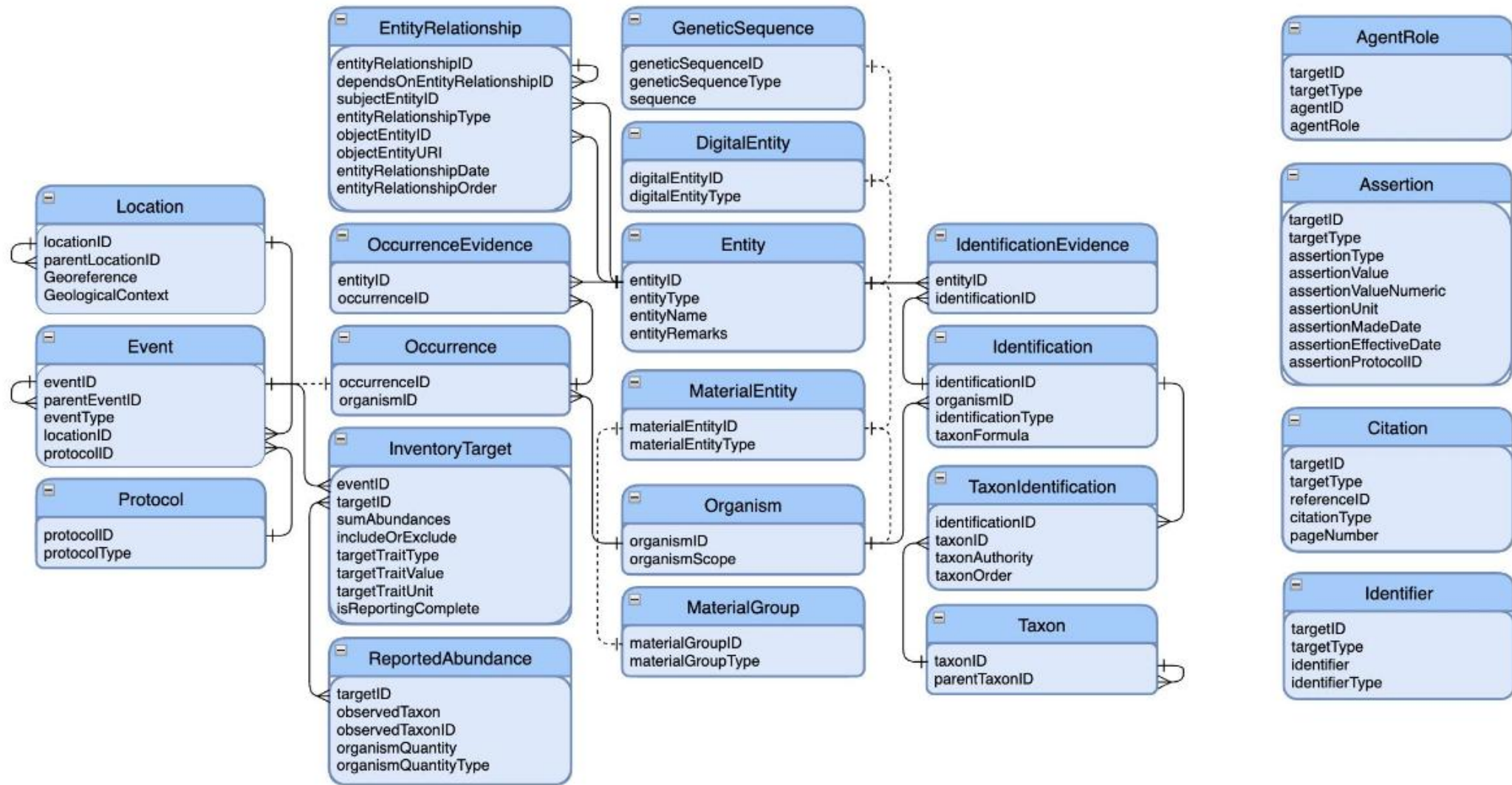
<https://gbif.github.io/dwc-dp/cm/>



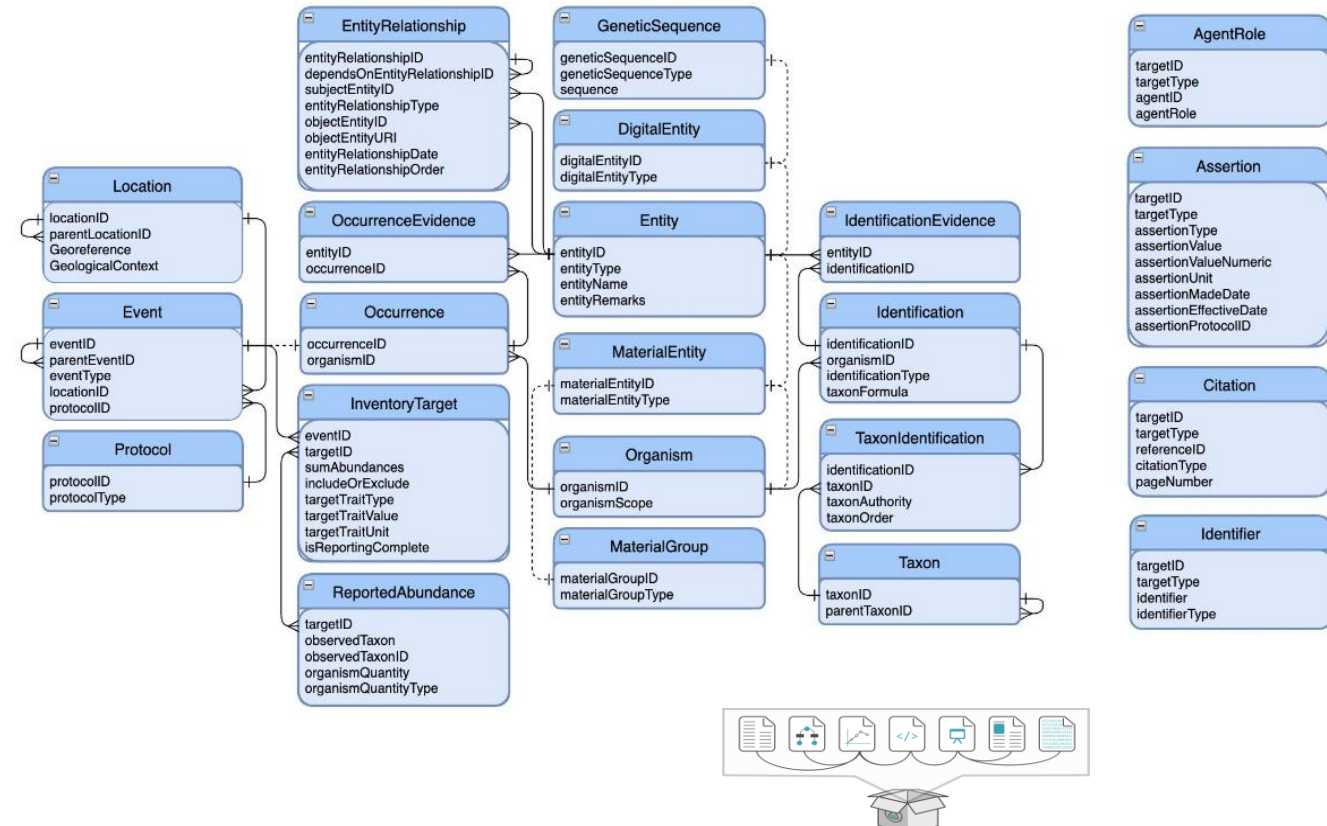
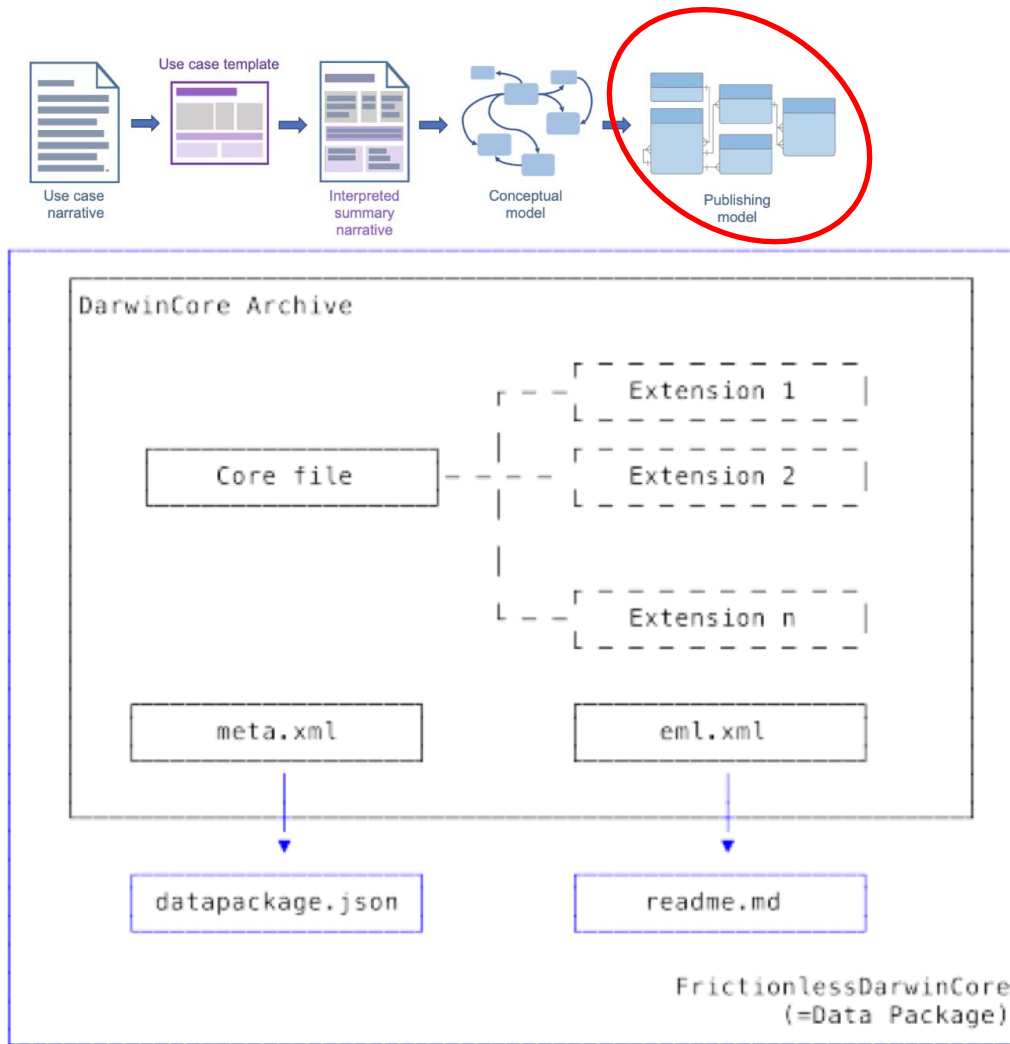
These connect in many-to-many relationships  
to targets above in context-specific ways



# Nuevo modelo conceptual para Darwin Core

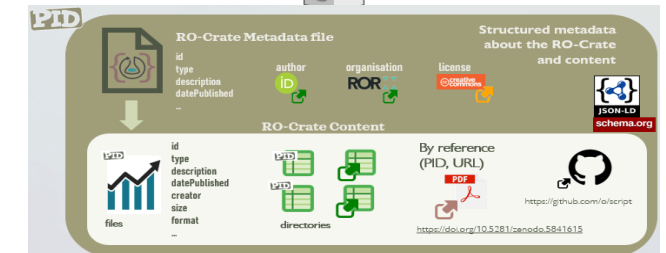


# Darwin Core – DP



Darwin Core – DP es una implementación de “Frictionless data Packages”

<https://gbif.github.io/dwc-dp/dp/>



# Hemos visto...

- Datos y ciencia
- Ciencia abierta y estándares
- Tipos de estándares
- TDWG
- Estándares en GBIF
- Darwin Core
- Humboldt Core
- Plinian Core
- Darwin Core Archive
- GBIF new data model
- Darwin Core – DP

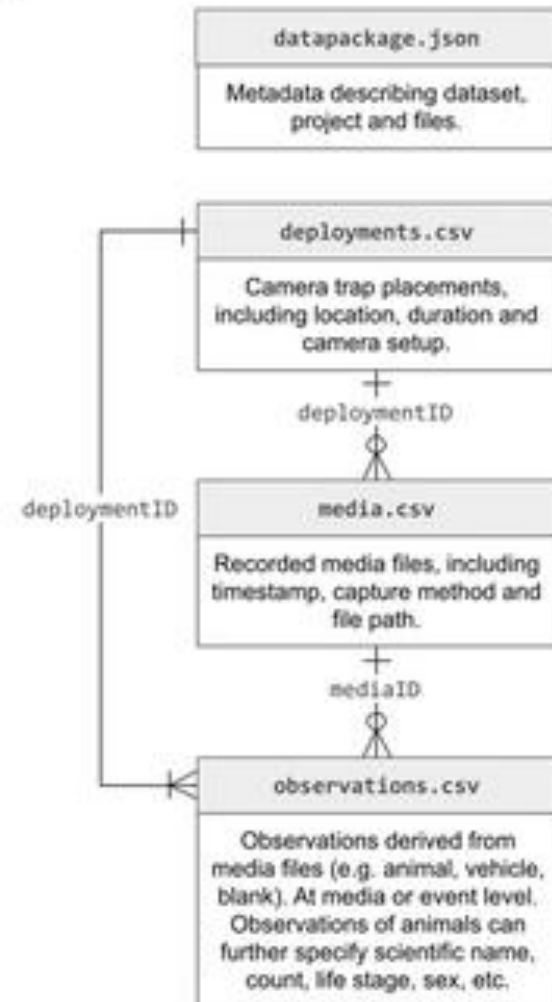


# Camtrap DP es una implementación de Darwin Core DP

<https://camtrap-dp.tdwg.org/>



(B)



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