

# Bioschemas Community Call Rolling Minutes - 2026

Every third Monday of the month at 17:00 CET/CEST

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This document -

<https://docs.google.com/document/d/1kd5F97ogdiPNhLTnkei-RVR8TC8Ohpc5QSPX3KsfDrk/edit#>

This is a rolling minute for the [Bioschemas Community](#) meeting. Expected participants are everyone involved in any Bioschemas activities worldwide, which include the Bioschemas leadership team, [Bioschemas - ELIXIR Strategic Implementation Study](#) members, Bioschemas collaborators (schema.org, OpenAIRE, Biothings-METADATAPLUS, etc.).

## Call-in instructions

**Current ZOOM for 2026 (from March 2026)**

<https://us02web.zoom.us/j/84701740969?pwd=BEMswQXAoiEShf8RJNLPivvHfPy87s.1>

## ELIXIR Zoom:

<https://elixir-europe-org.zoom.us/j/83477517512?pwd=LI5wZ41D4lib4TKGnl30zoLt1zsl7r.1> (until Feb.2026)

From March.2026 <https://us02web.zoom.us/j/84701740969?pwd=BEMswQXAoiEShf8RJNLPivvHfPy87s.1>

## Bioschemas resources

- ❖ [Useful information](#) - training resource, tooling, ELIXIR Strategic Implementation Study use cases
- ❖ Planning for Bioschemas at conferences/events - [index file](#)
- ❖ [Bioschemas Dataset cross-walk](#) (v0.2)
- ❖ [Crosswalks among commonly used metadata schemes](#) - publication includes Bioschemas (Zenodo)
- ❖ [Recordings folder](#)

## Communication Channels

- ❖ Slack - [invite link](#) for bioschemas.slack.com
- ❖ [Mailing list](#)
- ❖ [Bioschemas Github](#)
- ❖ [Bioschemas Google Drive](#)

## Archive of previous meetings

[Archive from 2020.06 to 2022.11](#)

[Archive for 2023](#)

[Archive for 2024](#)

[Archive for 2025](#)

**This year 2026**

## Meeting Notes Template

2026.mm.dd Next Meeting template - do not modify (Empty template for next meeting)		
Chair		Minuting:
Attendees		

<b>Apologies</b>	
<p>Community agenda (every odd month)</p> <ul style="list-style-type: none"> <li>❖ Newcomers</li> <li>❖ Presentation/Demo <ul style="list-style-type: none"> <li>➢ [Who][What]</li> <li>➢ [Who][What]</li> <li>➢ [Who][What]</li> <li>➢ [Who][What]</li> <li>➢ [Who][What]</li> </ul> </li> <li>❖ Round-table update/issue of discussion <p>Please indicate here (resource: person name) if you want to add an update</p> <ul style="list-style-type: none"> <li>➢</li> </ul> </li> <li>❖ Recent Changes/News (for information, not be read): <ul style="list-style-type: none"> <li>➢</li> </ul> </li> </ul>	
<p>Technical issues/Issue tracker (every even month)</p> <ul style="list-style-type: none"> <li>❖ Newcomers</li> <li>❖ Actions from previous call (please fill updated in before the meeting)</li> <li>❖ New issues and prioritisation</li> </ul>	
<p>Next meetings planning</p> <ul style="list-style-type: none"> <li>❖ Candidates for future presentations</li> </ul> <p>AOB/Suggested topics for next meeting</p> <ul style="list-style-type: none"> <li>❖ ...</li> </ul> <p><b>Next meeting</b></p>	

## Meetings 2026

<p>2026.03.16 - Community meeting</p> <p><a href="#">Zoom link for today</a> (automatic recording so we can later share the presentation)</p>		
<b>Chair</b>		<i>Minuting: everybody</i>
<b>Attendees</b>		

## Apologies

### Community agenda (every odd month - this month)

- ❖ Newcomers
  -
- ❖ **Presentation/Demo Invited talk - Egon Willighagen will discuss search engines and Bioschemas in his presentation/talk "Search engines and finding the long tail of science: is Bioschemas the solution?"**
  - [Who][What]
  - [Who][What]
  - [Who][What]
- ❖ Next community meetings
  - **18.May.2026 - Confirmed - NFDI4Biodiversity - Marcus Ernst**
  - **Nick & Alban Biodiversity (kickoff, sample/biosample)**
  - **Alban (around summer?)- tools metadata harvesting - research software ecosystem**
  - LabProtocol and LabProcess Germany/NFDI?
  - Microbiota: tentative, pending confirmation from Maja
    - Leyla to contact
  - RO-crates and models representation - Eli Chadwick
    - Nick to check with Eli
  - DALIA metadata schema
    - Nick to check dates
  - DCAT application profiles? Chemical community - Steffan? Used with RO-Crates. ELI? Similarities to Bioschemas profiles. Investigate. Data Plant (Sebastian Baier)
    - Leyla explore
- ❖ Round-table update/issue of discussion

**Please indicate here (resource: person name) if you want to add an update**

  - Topics for this meeting
    -
  - Topics from previous meetings
    - New types/profiles or guidance
      - (Sara, Nick) Biosamples type and profile
        - ◆ Leyla gave Nick more comments
      - (Leyla) LabProtocols type and profile
        - ◆ Mail sent
      - (Steffan) DefinedTerm
        - ◆ Ping Ginger (Nick)
      - (Carole) Models?
        - ◆ FAIR for AI; ELIXIR AI ecosystems focus group
        - ◆ Talk to [AIRBDS](#) as part of AIBIO-UK (Melanie)
          - Charlie Harrison
          - Presentation with FAIR4ML RDA
        - ◆ Biocroissant (Nick & Leyla joined community)
          - Leyla present Nick's slides at meeting next Tues
      - (requested by the FAIRAgro) Provenance
        - ◆ AI generated, how to represent AI-generated metadata (no progress - Gabriel) (of interest to Tess

- (Phil) as well
  - (Phil) LearningPath, LearningPathTopic
  - [learning paths new schemas proposal](#)
    - ◆ Gbh eu bh - Learning Path schema/profile
    - ◆ Sequential training materials
    - ◆ ELIXIR group happy with this work
- Schemas.science update
  - See AOB
- Events/workshops
  - ☐ ISCB-UK abstract Bioschemas (21-22 April, Cambridge)
  - ☐ ELIXIR platforms; ELIXIR AH (Jun, Lyon)
  - ☐ Biodiv-FAIR-checker k/o this week
  - ☐ ECCB - tentative
  - ☐ AlinBiology (Heidelberg) (March 10-13)
  - ☐ SWAT4LS
  - ☐ 20th Feb - AI readiness workshop in Hinxton
- Projects
  - (Alban) Bioschemas profile recommender
  - Datacite - completeness score for their profiles
- ❖ Recent Changes/News (for information, not be read):
  -
- ❖ AoB
  - The Dec 2025 ELIXIR TeSS Scientific Advisory Board meeting is recommending TeSS to move towards schemas.science over/as well as Bioschemas as the TeSS Platform moves into new domains.

### Technical issues/Issue tracker (every even month, not this time)

- ❖ Newcomers
  -
- ❖ Discussion points for this meeting (which should be discussed this meeting and cannot wait in the long list from previous actions)
  -
- ❖ Actions from previous call (please fill updated in before the meeting)
  -

### AOB/Suggested topics for next meeting

- ❖ Coming events
  - Next meeting
  -

## 2026.02.16 - technical call

[Zoom link for this meeting](#)

**Chair**

*Minuting: all*

**Attendees**

Nick, Alban, Phil, Leyla

**Apologies**

### Community agenda (every odd month, not this month)

- ❖ Newcomers
- ❖ Presentation/Demo
  - [Who][What]
- ❖ Round-table update/issue of discussion
  - Please indicate here (resource: person name) if you want to add an update
- ❖ Recent Changes/News (for information, not be read):
  -

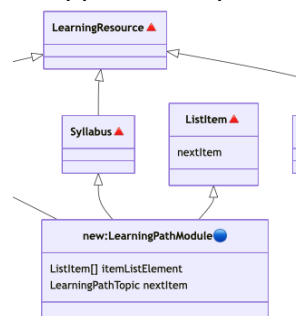
### Technical issues/Issue tracker (every even month, this month)

- ❖ Newcomers
- ❖
- ❖ Round table
  - (Alban, LJ) Mapping Bioschemas with CodeMeta
    - To be mapped to [Codemeta 3.1](#)
    - Crosswalk :  
<https://docs.google.com/spreadsheets/d/1vG25kGUyKoHACs8tusFJRI2H1AWUrfjaM9klCuSbbtM/edit?usp=sharing>
  - Workflows
    - Missing examples using bioschemas properties over [schema.org](#) in URIs
    - New request to join but it was spam email
  - LearningPath
    - Following instructions in specifications and governance docs, raised new issues ([spec#717](#), [gov#7](#))
    - What are the next steps? (Have [properties](#), [sandbox](#))
    - DDE does not have Syllabus (is it out of date? Can it do multiple inheritance like schema.org does now?)
  - RSec
    - ...
  - Biodiv
    - Links with FAIRsharing ?
      - Dataset
      - DataCatalog
      - Nuanced URLs in Fairsharing eg. resolving to their metadata record rather than a resource itself
    - FAIRassist → many ≠ tools / approaches

- OSTRails - focused on tracking provenance now, rather than cross-compare between FAIR assessment results directly
- BioCroissant: Bioschemas presented on 10.Feb.2026, waiting for this community to decide the way to go
  - Some interest : “Bioschemas maybe useful on top of Croissant to specify and assess profile completeness”
  - Need some follow-up
- Events
  - (LJ) will present something related to training materials (possible update to TrainingMaterial profile) to the FAIR training materials group in ELIXIR, 04.March.2026
    - Analysing similarities with the MODALIA ontologies and the TrainingMaterial Bioschemas profile →
    - All updated crosswalks at <https://drive.google.com/drive/folders/1LLyba9zCug8UZKyVCw2BX4H5eXcMJceO?usp=sharing>
  - (Phil) very interested in the mapping for an EOSC workshop
  - German Bioinformatics Conference, do we want to [submit a workshop](#) together with BioCroissant? Deadline 31.March.2026
- (LJ) Batch modifications about includedInDataset
  - hasPart for CreativeWork branch: Tutorial
  - Add includedInDataset to BioChemEntity

❖ TODO for the next meeting

- (Nick/Alban?) REVISE the governance “update/creation” of profiles section
- (Phil) Ask Ginger about <https://schema.org/Syllabus> in DDE
  - How DDE supports multiple inheritance (Syllabus + ListItem +



Course)

- Invitation to first [mTeSS-X Metadata Model Workshops](#) 24 February
- (Phil/Alban) issues with website/GitHub - Latest action: organize call with Alban

❖ PENDING Actions from previous call (please fill updated in before the meeting)

- [From previous technical call](#)

- (Phil/Alban) issues with website/GitHub - Latest action: organize call with Alban
- Phil: **Learning Path** new profiles queries 2026-02-16
  - See above
- [MolecularEntity](#) - One of many for minimum properties
  - Done: DDE,
  - Ongoing: FAIRChecker
  - Pending: Review by Steffen
  - Latest action: doing during hackathon at ZB MED but nothing scheduled for this and not done in previous hackathons
- Add includedInDataset to all profiles [#617](#)
  - Get the list of profiles that use isPartOf as minimum, recommended or optional → [DONE](#)
  - For CreativeWork branch, use isPartOf
  - **Action needed: what to do for BioChemEntity? → add includedInDataset to type BioChemEntity and corresponding profiles**
  - **Action needed: write tutorials**
- DefinedTerm / DefinedTermSet [#618](#)
  - **Action needed: ping Ginger**
- New generic types in Bioschemas, e.g., CreativeWork so we can include generic properties such as intendedUse (for datasets, software, workflows, etc.)
  - Pending: Check how it would work with [schem.org](https://www.schem.org) validators
- [Issue 629](#). Description of the (nature of) content of a Dataset (or any other CreativeWork in Bioschemas) → RO-crates use **about**
  - **Pending Action: get [PR status](#) from Ginger**
- ❖ New subjects for this meeting:
  -
- ❖ New issues and prioritisation

## Next meetings planning

❖

## AOB/Suggested topics for next meeting

❖ ...

## Next meeting



## 2026.01.19 - Community meeting

[Zoom link for today](#) (automatic recording so we can later share the presentation)

Chair		Minuting: everybody
Attendees	Gabriel Schneider, Victor Lu, Leyla Castro, Alban Gaignard, Nick Juty, Melanie Vollmar, Phil Reed	
Apologies		

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- ❖ Newcomers
  -
- ❖ **Presentation/Demo Invited talk - Victor Lu presenting licenses with SPDX (and more you can do with it)**
  - Presentation available at [https://docs.google.com/presentation/d/1\\_Az1r3zQdr17iwke5140PvXjDA2ja7zwrtcy\\_zCLHys/edit?usp=sharing](https://docs.google.com/presentation/d/1_Az1r3zQdr17iwke5140PvXjDA2ja7zwrtcy_zCLHys/edit?usp=sharing)
  - SPDX is an ontology, see for instance Dataset at SPDX Dataset : <https://spdx.github.io/spdx-spec/v3.0.1/model/Dataset/Classes/DatasetPackage/>
  - Paper about AI/dataset profile at <https://arxiv.org/abs/2504.16743>
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  - **Open slot 16 March**
    - **Nick & Alban Biodiversity (kickoff, sample/biosample)**
  - **Alban (around summer?)- tools metadata harvesting - research software ecosystem**
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      - Gbh - need biosample and lab protocol. They will contact us
      - Lukas Wiel - maybe could present too..
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