

# Bioschemas Community Call Rolling Minutes - 2026

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

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[Community agenda \(every odd month\)](#)

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[Community agenda \(every odd month - this month\)](#)

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

[AOB/Suggested topics for next meeting](#)

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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=L15wZ41D4lib4TKGnI30zoLt1zsl7r.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		<i>Minuting:</i>
Attendees		

<b>Apologies</b>	
Community agenda (every odd month)	<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 <div style="background-color: #ffff00; border: 1px solid black; padding: 2px; display: inline-block;"> <b>Please indicate here (resource: person name) if you want to add an update</b> </div> <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>
Technical issues/Issue tracker (every even month)	<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>
Next meetings planning	<ul style="list-style-type: none"> <li>❖ Candidates for future presentations</li> </ul>
AOB/Suggested topics for next meeting	<ul style="list-style-type: none"> <li>❖ ...</li> </ul>
<b>Next meeting</b>	

## Meetings 2026

### 2026.03.16 - Community meeting

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

Chair		<i>Minuting: everybody</i>
Attendees		

Apologies	
Community agenda (every odd month - this month)	
<ul style="list-style-type: none"> <li>❖ Newcomers <ul style="list-style-type: none"> <li>➤</li> </ul> </li> <li>❖ <b>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?"</b> <ul style="list-style-type: none"> <li>➤ [Who][What]</li> <li>➤ [Who][What]</li> <li>➤ [Who][What]</li> </ul> </li> <li>❖ Next community meetings <ul style="list-style-type: none"> <li>➤ <b>18.May.2026 - Confirmed - NFDI4Biodiversity - Marcus Ernst</b></li> <li>➤ <b>Nick &amp; Alban Biodiversity (kickoff, sample/biosample)</b></li> <li>➤ <b>Alban (around summer?)- tools metadata harvesting - research software ecosystem</b></li> <li>➤ LabProtocol and LabProcess Germany/NFDI?</li> <li>➤ Microbiota: tentative, pending confirmation from Maja <ul style="list-style-type: none"> <li>■ Leyla to contact</li> </ul> </li> <li>➤ RO-crates and models representation - Eli Chadwick <ul style="list-style-type: none"> <li>■ Nick to check with Eli</li> </ul> </li> <li>➤ DALIA metadata schema <ul style="list-style-type: none"> <li>■ Nick to check dates</li> </ul> </li> <li>➤ DCAT application profiles? Chemical community - Steffan? Used with RO-Crates. ELI? Similarities to Bioschemas profiles. Investigate. Data Plant (Sebastian Baier) <ul style="list-style-type: none"> <li>■ Leyla explore</li> </ul> </li> </ul> </li> <li>❖ Round-table update/issue of discussion <ul style="list-style-type: none"> <li><b>Please indicate here (resource: person name) if you want to add an update</b> <ul style="list-style-type: none"> <li>➤ Topics for this meeting <ul style="list-style-type: none"> <li>■</li> </ul> </li> <li>➤ Topics from previous meetings <ul style="list-style-type: none"> <li>■ New types/profiles or guidance <ul style="list-style-type: none"> <li>• (Sara, Nick) Biosamples type and profile <ul style="list-style-type: none"> <li>◆ Leyala gave Nick more comments</li> </ul> </li> <li>• (Leyala) LabProtocols type and profile <ul style="list-style-type: none"> <li>◆ Mail sent</li> </ul> </li> <li>• (Steffan) DefinedTerm <ul style="list-style-type: none"> <li>◆ Ping Ginger (Nick)</li> </ul> </li> <li>• (Carole) Models? <ul style="list-style-type: none"> <li>◆ FAIR for AI; ELIXIR AI ecosystems focus group</li> <li>◆ Talk to <a href="#">AIRBDS</a> as part of AIBIO-UK (Melanie) <ul style="list-style-type: none"> <li>➤ Charlie Harrison</li> <li>➤ Presentation with FAIR4ML RDA</li> </ul> </li> <li>◆ Biocroissant (Nick &amp; Leyla joined community) <ul style="list-style-type: none"> <li>➤ Leyala present Nick's slides at meeting next Tues</li> </ul> </li> </ul> </li> <li>• (requested by the FAIRAgro) Provenance <ul style="list-style-type: none"> <li>◆ AI generated, how to represent AI-generated metadata (no progress - Gabriel) (of interest to Tess)</li> </ul> </li> </ul> </li> </ul> </li> </ul> </li> </ul> </li> </ul>	

- (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)

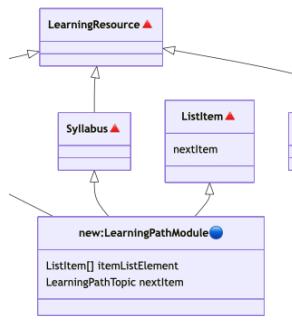
- ❖ Newcomers
- ❖ Presentation/Demo
  - [Who][What]
- ❖ Round-table update/issue of discussion
  - Please indicate here (resource: person name) if you want to add an update**
  -
- ❖ Recent Changes/News (for information, not be read):
  -

### 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/1vG25kGUyKoHACs8tusFJRI2H1AWUrfjaM9kICuSbbtM/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 +



- 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](#) 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

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

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

- ❖ 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/DatasetPac\\_kage/](https://spdx.github.io/spdx-spec/v3.0.1/model/Dataset/Classes/DatasetPac_kage/)
  - Paper about AI/dataset profile at <https://arxiv.org/abs/2504.16743>
  - [Who][What]
- ❖ Next community meetings
  - **18.May.2026 - Confirmed - NFDI4Biodiversity - Marcus Ernst**
  - **Open slot 16 March**
    - **Nick & Alban Biodiversity (kickoff, sample/biosample)**
  - **Alban (around summer?)- tools metadata harvesting - research software ecosystem**
  - Microbiota: tentative,pending confirmation from Maja
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      - (Sara, Nick) Biosamples type and profile
        - Leyala gave Nick more comments
      - (Leyla) LabProtocols type and profile
        - Gbh - need biosample and lab protocol. They will contact us
        - Lukas Wiel - maybe could present too..
    - ◆ Leyla to contact

- (Steffan) DefinedTerm
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