NOMAD structures data into \*\*sections\*\*, where each section can contain data and more sections.

This allows to browse complex data like you would browse files and directories on your computer.

Each section follows a \*\*definition\*\* and all the contained data and subsection have a specific name, description, possible type, shape, and unit. This means that all data follows a \*\*schema\*\*.

This not only helps the human exploration, but also makes it machine interpretable, increases consistency and interoperability, enables search, APIs, visualization, and analysis.

!processed data screenshot (images/screenshot.png)

Browsing structured data in the NOMAD UI (link)

Schema language

The bases for structured data are schemas written in a \*\*schema language\*\*. Our schema language is called the \*\*NOMAD Metainfo\*\* language. The name is evocative of the rich \*\*metadata information\*\* that should be associated with the research data and made available in a machine-readable format.

It defines the tools to define sections, organize definitions into \*\*packages\*\*, and define section properties (\*\*subsections\*\* and \*\*quantities\*\*).

!schema language (images/schema\_language.png)

The NOMAD Metainfo schema language for structured data definitions

Packages contain section definitions, section definitions contain definitions for

subsections and quantities. Sections can inherit the properties of other sections. While

subsections allow to define containment hierarchies for sections, quantities can

use section definitions (or other quantity definitions) as a type to define references.

If you are familiar with other schema languages and means to defined structured data

(json schema, XML schema, pydantic, database schemas, ORM, etc.), you might recognize

these concept under different names. Sections are similar to \*classes\*, \*concepts\*, \*entities\*, or \*tables\*.

Quantities are related to \*properties\*, \*attributes\*, \*slots\*, \*columns\*.

subsections might be called \*containment\* or \*composition\*. subsections and quantities

with a section type also define \*relationships\*, \*links\*, or \*references\*.

Our guide on how to write a schema (../howto/customization/basics.md) explains these concepts with an example.

#### Schema

NOMAD represents many different types of data. Therefore, we cannot speak of just \*the one\*

schema. The entirety of NOMAD schemas is called the \*\*NOMAD Metainfo\*\*.

Definitions used in the NOMAD Metainfo fall into three different categories. First, we have sections that define a \*\*shared entry structure\*\*. Those are independent of the

type of data (and processed file type). They allow to find all generic parts without any deeper understanding of the specific data. Second, we have definitions of \*\*re-usable base sections\*\* for shared common concepts and their properties.

Specific schemas can use and extend these base sections. Base sections define a fixed interface or contract that can be used to build tools (e.g. search, visualizations, analysis)

around them. Lastly, there are \*\*specific schemas\*\*. Those re-use base sections and complement the shared entry structure. They define specific data structures to represent

specific types of data.

!schema language (images/schema.png)

The three different categories of NOMAD schema definitions

### Base sections

Base section is a very loose category. In principle, every section definition can be inherited from or can be re-used in different contexts. There are some dedicated (or even abstract)

base section definitions (mostly defined in the `nomad.datamodel.metainfo` package and sub-packages),

but schema authors should not strictly limit themselves to these definitions.

The goal is to re-use as much as possible and to not re-invent the same sections over and over again. Tools build around certain base section, provide an incentive to use them.

# !!! warning "Attention"

There is no detailed how-to or reference documentation on the existing base sections and how to use them yet.

One example for re-usable base section is the workflow package (../howto/customization/workflows.md).

These allow to define workflows in a common way. They allow to place workflows in the shared entry structure, and the UI provides a card with workflow visualization and navigation for all entries that have a workflow inside.

## !!! warning "Attention"

Currently there are two version of the workflow schema. They are stored in two top-level `EntryArchive` subsections (`workflow` and `workflow2`). This

will change soon to something that supports multiple workflows used in specific schemas and results.

Specific schemas

Specific schemas allow users and plugin developers to describe their data in all detail. However, users (and machines) not familiar with the specifics, will struggle to interpret these kinda of data. Therefore, it is important to also translate (at least some of) the data

into a more generic and standardized form.

!schema language (images/data.png)

From specific data to more general interoperable data.

The \*\*results\*\* section provides a shared structure designed around base section definitions.

This allows you to put (at least some of) your data where it is easy to find, and in a form that is easy to interpret. Your non-interoperable, but highly detailed data needs to be transformed into an interoperable (but potentially limited) form.

Typically, a parser will be responsible to populate the specific schema, and the interoperable schema parts (e.g. section results) are populated during normalization.

This allows to separate certain aspects of conversions and potentially enables re-use for normalization routines. The necessary effort for normalization depends on how much

the specific schema deviates from base-sections. There are three levels:

- the parser (or uploaded archive file) populates section results directly
- the specific schema re-uses the base sections used for the results and normalization

can be automated

- the specific schema represents the same information differently and a translating normalization algorithm needs to be implemented.

Exploring the schema

All built-in definitions that come with NOMAD or one of the installed plugins can be explored with the Metainfo browser (https://nomad-lab.eu/prod/v1/gui/analyze/metainfo/nomad.datamodel.datamodel.Entry Archive){:target="\_blank"}. You can start with the root section `EntryArchive` and browse based on subsections, or explore the Metainfo through packages.

To see all user provided uploaded schemas, you can use a search for the subsection `definition`

 $\label{lab:euprod} $$ \left(\frac{1}{2} - \frac{1}{2} \right) = \frac{1}{2} . $$ (https://nomad-lab.eu/prod/v1/gui/search/entries?quantities=definitions) {:target="_blank"}.$ 

The subsection `definition` is a top-level `EntryArchive` subsection. See also our how-to on writing and uploading schemas (../howto/customization/basics.md#uploading-schemas).

Contributing to the Metainfo

The shared entry structure (including section results) is part of the NOMAD source-code. It interacts with core functionality and needs to be highly controlled.

Contributions here are only possible through merge requests.

Base sections can be contributed via plugins. Here they can be explored in the Metainfo browser, your plugin can provide more tools, and you can make use of normalize functions.

See also our how-to on writing schema packages (../howto/plugins/schema\_packages.md). You could

also provide base sections via uploaded schemas, but those are harder to explore and

distribute to other NOMAD installations.

Specific schemas can be provided via plugins or as uploaded schemas. When you upload

schemas, you most likely also upload data in archive files (or use ELNs to edit such files).

Here you can also provide schemas and data in the same file. In many case specific schemas will be small and only re-combine existing base sections.

See also our

how-to on writing YAML schemas (../howto/customization/basics.md).

Data

All processed data in NOMAD instantiates Metainfo schema definitions and the \*archive\* of

each entry is always an instance of `EntryArchive`. This provides an abstract structure for all data. However, it is independent of the actual representation of data in computer memory

or how it might be stored in a file or database.

The Metainfo has many serialized forms. You can write `.archive.json` or `.archive.yaml`

files yourself. NOMAD internally stores all processed data in message pack (https://msgpack.org/){:target="blank"}. Some

of the data is stored in mongodb or elasticsearch. When you request processed data via API, you receive it in JSON. When you use the ArchiveQuery (../howto/programmatic/archive\_query.md), all data is represented

as Python objects (see also example in schema package documentation (../howto/plugins/schema packages.md#schemapackage-class)).

No matter what the representation is, you can rely on the structure, names, types,

shapes, and units

defined in the schema to interpret the data.

Archive files: a shared entry structure

Broadening the discussion on the \*entry\* files that one can find in NOMAD, both schemas (#schema) or processed data (#data) are serialized as the same kind of \*archive file\*, either `.archive.json` or `.archive.yaml`.

The NOMAD archive file is indeed composed by several sections.

NOMAD archive file: `EntryArchive`

\* definitions: `Definitions`

\* metadata: `EntryMetadata`

\* data: `EntryData`

\* run: `Run`

\* nexus: `Nexus`

\* workflow: `LegacyWorkflow`

\* workflow2: `Workflow`

\* results: `Results`

They all instantiate the same root section `EntryArchive`. They all share common sections `metadata:Metadata`

and `results:Results`. They also all contain a \*data\* section, but the used section definition varies depending on the type of data of the specific entry. There is the literal `data:EntryData` subsection. Here `EntryData` is abstract and specific entries will use concrete definitions that inherit from `EntryData`. There are also specific \*data\*

sections, like `run` for simulation data and `nexus` for nexus data.

!!! note

As shown in Uploading schemas

(.../howto/customization/basics.md#uploading-schemas), one can, in principle, create an archive file with both `definitions` and one of the \*data\* sections filled, although this is not always desired because it will stick together a schema and a particular instance of that schema. They should be kept separate so that it is still possible to generate new data files from the same schema file.

# !!! warning "Attention"

The results, originally only designed for computational data, will soon be revised an replaced by a different section. However, the necessity and function of a section like this remains.

!schema language (images/super\_structure.png)

All entries instantiate the same section share the same structure.