How to write a YAML schema package

This guide explains how to write and upload NOMAD schema packages in the YAML format that can be uploaded as part of your data. This is a good way to start out experimenting with custom data structures in NOMAD, but for more advanced use cases you may need to use Python schema packages (../plugins/schema_packages.md). For more information on how an archive file is composed, visit Explanation > Data structure (../../explanation/data.md).

Example data

Let's assume we want to describe chemical compositions using the elements they contain.

The following structured data (in this example as a `.yaml` document) could describe the composition of water.

```
```yaml
```

{{ yaml\_snippet('examples/docs/basic\_schema/data.archive.yaml:data', '', 'm\_def') }}

In structured data formats (such as `.yaml` or `.json`), data is put into combinations of \*primitive values\* (e.g. `'H2O'`, `1.141`), \*objects\* (a set of \*keys\* and \*value\* pairs, where \*values\* can be \*objects\*, \*lists\*, or \*primitive values\*), and \*lists\* of \*values\*.

## Sections

In a schema package, we want to describe the structure of data, i.e. what are the allowed combinations of \*objects\*, \*lists\*, and \*primitive values\*.

The crucial task here is to define what \*keys\* certain \*types of objects\* can have and what possible \*values\* might exist for each of these keys.

In NOMAD, we call \*objects\* \*\*sections\*\* and we define \*types of objects\* with \*\*section

definitions\*\*. Since \*objects\* can be nested, \*\*sections\*\* become like the sections and subsections of a book or paper. Sections are a representation of data and they are the building blocks for \*\*archives\*\* (../../reference/glossary.md#archive). Section definitions form a schema package and they are

the building blocks for the \*\*metainfo\*\* (../../reference/glossary.md#metainfo).

In the above example, we have two \*types\* of \*objects\*: an overaching object for the entire structure

(with \*keys\* for `composition` and `elements`), and an additional object which describes the internal structure of

`elements` (with \*keys\* for `label`, `density`, and `isotopes`). Let's start with the \*definition\* for elements. This is what the \*section definition\* looks like in NOMAD's yaml-based schema package format:

```
```yaml
```

Element:

{{

yaml_snippet('examples/docs/basic_schema/package.archive.yaml:definitions/sections/ Element', ' ') }}

• • •

A *section definition* provides all the available *keys* for a *section* that instantiates this *definition*. For each *key*, e.g. `label`, `density`, `isotopes`, it provides more information on the possible values.

Let's have a look at the overall definition for our chemical composition:

```
```yaml
```

Composition:

{{

yaml\_snippet('examples/docs/basic\_schema/package.archive.yaml:definitions/sections/

```
Composition', ' ') }}
Again, all possible *keys* (`composition` and `elements`) are defined. But now we see
that there are two different types of *keys*, **quantities** and **subsections**. We
say that *section definitions* can have **properties** (e.g. the *keys* they define) and
there are two distinct types of *properties*.
Quantities
Quantities define possible *primitive values*. The basic properties that go into
a *quantity definition* are:
- **type**: what kind of *primitive value* can be used, e.g. `str` or `np.float64`
- **shape**: what is the shape of the value, e.g. scalar or list (`['*']`)
- **unit**: what is the physical meaning of the value
The *names* of *quantity definitions* serve as the *key*, used in respective *section
objects*.
Type
This is a list of supported quantity types.
|type|description|
|-|-|
|`string`||
|`str`||
|`float`||
|`integer`||
|`int`||
|`boolean`||
|`bool`||
```

|`np.int32`|Numpy based integer with 32 bits.|

```
|`np.int64`|Numpy based integer with 64 bits.|
|`np.float32`|Numpy based float with 32 bits.|
|`np.float64`|Numpy based float with 64 bits.|
|`Datetime`||
|`User`|A type for NOMAD users as values.|
|`Author`|A complex type for author information.|
|`{type_kind: Enum, type_data: []}`|Use `type_data` to specify enum values as list of strings.|
|**|To define a quantity that is a reference to a specific section.|
```

The shape of a quantity is a list of \*dimensions\*, where each \*dimension\* defines the possible size of that \*dimension\*. The empty list (or no shape) describes a scalar value, a list with one \*dimension\* a list or vector, a list with two \*dimensions\* a matrix, etc.

Dimensions can be given as:

- an integer number to define a fixed size, e.g. a 3x3 matrix would have shape `[3, 3]`.
- the string `'\*'` to denote am arbitrary sized dimension, e.g. a list quantity would have shape `['\*']`.
- A string that describes the name of a sibling quantity with an integer type, e.g. `['number\_of\_atoms', 3]`

Unit

Shape

NOMAD manages units and data with units via the Pint (https://pint.readthedocs.io/en/stable/) Python package. A unit is given as a string that is parsed by pint. These strings can

be simple units (or their aliases) or complex expressions. Here are a few examples: 'm', 'meter', 'mm', 'millimeter', 'm/s', 'm/s\*\*2'.

While you can use all kinds of units in your uploaded schema packages, the built-in

NOMAD schema (Metainfo) uses only SI units.

Subsections

\*Subsections\* define a \*part-of-relationship\* between two \*sections\*. \*Subsection definitions\* are \*properties\* of the parent \*section definition\* and name a child \*section definition\*. In the data, we can now contain instances of the target (e.g. `Element`) in instances of the source (e.g. `Composition`). A \*subsection\* can be defined as \*repeating\* to allow many child \*sections\* of the same \*type\*. In our example,

one 'Composition' can contain many 'Elements'.

The \*names\* of \*subsection definitions\* serve as the \*key\*, used in respective \*section objects\*.

Uploading schema packages

NOMAD archive files allow you to upload data in NOMAD's native file format. An archive file can be a .yaml or .json file. It ends with `.archive.json` or `.archive.yaml`.

Archive files are mainly used to convey data. Since YAML schema packages are also "just" data, archive

files can also be used to convey a schema package.

You can upload schema packages and data in separate files.

```
`schema_package.archive.yaml`

'`yaml

--8<-- "examples/docs/basic_schema/package.archive.yaml"

'``

and `data.archive.yaml`

```yaml

--8<-- "examples/docs/basic schema/data.archive.yaml"
```

```
Or, you can upload the schema package and data in the same file:
```yaml
--8<-- "examples/docs/basic schema/package.archive.yaml"
data:
 m def: Composition
{{ yaml snippet('examples/docs/basic schema/data.archive.yaml:data', ' ', 'm def') }}
References
Reference quantities
We already saw that we can define a *part-of* relationship between sections. When we
want to represent highly inter-linked data, this is often insufficient. *References*
allow us to create a more lose relationship between sections.
A reference is a uni-directional link between a *source* section and a *target* section.
References can be defined in a schema package as a quantity in the *source* section
definition
that uses the *target* section definition as a type.
Instead of connecting the elements in a composition with subsections, we can also
connect a composition section to elements with a quantity:
```yaml
Composition:
{{
yaml snippet('examples/docs/references/single.archive.yaml:/definitions/sections/Comp
osition', ' ') }}
. . .
Here, `type: Element` refers to the section definition `Element`, very similar to
`section: Element` in a subsection definition.
```

We saw above that subsections are represented as nested *objects* in data (forcing a *part-of* relationship). References are represented as string-typed *primitive values* in serialized data. Here is an example `Composition` with references to elements: ```yaml {{ yaml_snippet('examples/docs/references/single.archive.yaml:/data/compositions/0', '') }}

These string-references determine the *target* section's place in the same archive.

Each `/`-separated segment represents a *key*. A reference starts from the

root *object* and following the sequence of *keys* to a specific *object* (i.e. section).

Here is the full archive data:

```yaml

data:

{{ yaml\_snippet('examples/docs/references/single.archive.yaml:/data', ' ', 'm\_def') }}
...

If you follow the \*keys\* `data`, `periodic\_table`, `elements`, `0`, you reach the section that represent hydrogen. Keep in mind that \*lists\* use index-numbers as \*keys\*.

Schema package references

References can look different depending on the context. Above we saw simple references

that point from one data section to another. But, you also already a saw a different type of reference. Schema packages themselves contain references: when we used `type: Element` or `section: Element` to refer to a \*section definition\*, we were writing down references that point to a \*section definition\*. Here we can use a convenience representation: `Element` simply replaces the otherwise cryptic

`#/definitions/sections/0`.

So far, we never discussed the use of `m\_def`. In the examples you might have seen this

as a special \*key\* in some objects. Whenever we cannot determine the \*section definition\*

for a \*section\* by its context (e.g. the \*key\*/\*subsection\* used to contain it in a \*parent section\*), we use `m\_def` to provide a reference to the \*section definition\*.

Different forms of references

Depending on where references are used, they might take a different serialized form.

Here are a few examples for different reference syntax:

|Example reference|Comments|

|---|

|`#/data/periodic\_table/elements/0`|Reference to a section within the subsection hierarchy of the same archive.|

|`Element`|Reference to a \*section definition\* in the same archive. Can only be used to target \*section definitions\*.|

|`nomad.datamodel.metainfo.workflow`|Reference to a \*section definition\* that was written in Python and is part of the NOMAD code. Can only be used to target \*section definitions\*.|

|`../upload/raw/data.archive.yaml#/data`|Reference to a section in a different

`.archive.yaml` file of the same upload.|

|`../upload/archive/mainfile/data.archive.yaml#/data`|Reference to a section in a processed archive given by entry \*mainfile\*.|

\`../upload/archive/zxhS43h2kqHsVDqMboiP9cULrS\_v#/data`|Reference to a section in a processed archive given by entry-id.|

|`../uploads/zxhS43h2kqHsVDqMboiP9cULrS\_v/raw/data.archive.yaml#/data`|Reference

to a section in an entry of a different upload.

|`https://mylab.eu/oasis/api/v1/uploads/zxhS43h2kqHsVDqMboiP9cULrS\_v/raw/data.arch ive.yaml#/data`|Reference to a section in an entry in a different NOMAD installation.|

References across entries

A references in the archive of one entry can point to a section in a different entry's archive. The following two example files, exemplify this use of reference between two NOMAD entries.

```
periodic_table.archive.yaml

```yaml

--8<-- "examples/docs/references/periodic_table.archive.yaml"

```

composition.archive.yaml

```yaml

--8<-- "examples/docs/references/composition.archive.yaml"
```

These inter-entry references have two parts: `#`, where *entry*

is a path or URL denoting the *target* entry and *section* a path within the *target* entry's subsection containment hierarchy.

Please note that also schema packages can be spread over multiple files. In the above example,

one file contained the schema package and data for a periodic table and another file contained

schema package and data for the composition of water (using the periodic table).

Base sections and inheritance

. . .

We add a relationship between *section definitions* that allows us to create more *specialized* definitions from more *abstract* definitions. Here the

properties of the *abstract* definition are inherited by the more *specialized definitions*

Base sections

Here is a simple schema package with two *specialization* of the same *abstract* section

definition:

```yaml

definitions:

```
{{ yaml_snippet('examples/docs/inheritance/basic.archive.yaml:/definitions', ' ') }}
```

The two \*specialized\* definitions `Annealing` and `Evaporation` define the \*abstract\* definition `Process` via the `base\_section` property. With this `Annealing` and `Evaporation`

inherit the quantity `time`. We do not need to repeat quantities from the base section, and we can add more properties. Here is an example `Evaporation` using both the inherited

and added quantity:

```yaml

```
{{ yaml_snippet('examples/docs/inheritance/basic.archive.yaml', '', 'definitions') }}
...
```

Polymorphy

What happens if we reference *abstract* definitions in subsections or reference quantities?

Here is an subsection example. In one schema, we define the relationship between 'Sample'

and `Process`. In another schema, we want to add more *specializations* to what a

```
process is.
**abstract.archive.yaml**
```yaml
--8<-- "examples/docs/inheritance/abstract.archive.yaml"
specialized.archive.yaml
```yaml
{{ yaml snippet('examples/docs/inheritance/specialized.archive.yaml', '', 'data') }}
The *section definition* use in the subsection `processes` defines what a contained
section has to be "at least". Meaning that any section based on a *specialization* of
`Process` would be a valid `processes` subsection.
**specialized.archive.yaml**
```yaml
definitions:
 # see above
data:
{{ yaml snippet('examples/docs/inheritance/specialized.archive.yaml:data', ' ') }}
```

The fact that a subsection or reference target can have different "forms" (i.e. based on different \*specializations\*) is called \*polymorphism\* in object-oriented data modelling.

Pre-defined sections

NOMAD provides a series of built-in \*section definitions\*. For example, there is `EntryArchive`, a definition for the top-level object in all NOMAD archives (e.g. `.archive.yaml` files). Here is a simplified except of the \*main\* NOMAD schema `nomad.datamodel`:

```
EntryArchive:

sub_sections:

metadata:

section: EntryMetadata

definitions:

section: nomad.metainfo.Package

data:

section: EntryData

... many more

EntryData:

empty
```

Compare this to the previous examples: we used the top-level \*keys\* `definitions` and `data` without really explaining why. Here you can see why. The `EntryArchive` \*property\* `definitions` allows us to put a \*schema package\* into our archives. And the `EntryArchive` \*property\* `data` allows us to put \*data\* into archives that is a \*specialization\* of `Schema`. The `Schema` definition is empty. It is merely an \*abstract\* placeholder that allows you to add \*specialized\* data sections to your archive. Therefore, all \*section definitions\* that define a top-level data section, should correctly use `nomad.datamodel.Schema` as a base section. This would be the first "correct" example:

```
```yaml
--8<-- "examples/docs/inheritance/hello.archive.yaml"
```

Here are a few other built-in section definitions and packages of definitions:

|Section definition or package|Purpose|

|---|

|nomad.datamodel.EntryArchive|Used for the root object of all NOMAD entries|
|nomad.datamodel.EntryMetadata|Used to add standard NOMAD metadata such as ids,
upload, processing, or author information to entries.|

|nomad.datamodel.EntryData|An abstract section definition for the `data` section.|
|nomad.datamodel.ArchiveSection|Allows to put `normalize` functions into your section
definitions.|

|nomad.datamodel.metainfo.eln.*|A package of section definitions to inherit commonly used quantities for ELNs. These quantities are indexed and allow specialization to utilize the NOMAD search.|

|nomad.parsing.tabular.TableData|Allows to inherit parsing of references .csv and .xls files.|

|nomad.datamodel.metainfo.workflow.*|A package of section definitions use by NOMAD to define workflows|

|nomad.metainfo.*|A package that contains all *definitions* of *definitions*, e.g. NOMAD's "schema language". Here you find *definitions* for what a sections, quantity, subsections, etc. is.|

Separating data and schema package

As we saw above, a NOMAD entry can contain schema package `definitions` and `data` at the

same time. To organize your schema package and data efficiently, it is often necessary to re-use

schema packages and certain data in other entries. You can use *references* to spread your

schema packages and data over multiple entries and connect the pieces via

references.

Here is a simple schema package, stored in a NOMAD entry with mainfile name `package.archive.yaml`:

```yaml

--8<-- "examples/docs/references/multiple files/package.archive.yaml"

. . .

Now, we can re-use this schema package in many entries via \*references\*. Here, we extend

a schema contained in the package and instantiate definitions is a separate mainfile `data-and-package.archive.yaml`:

```yaml

--8<-- "examples/docs/references/multiple_files/data-and-package.archive.yaml"

...

Here is a last example that re-uses the schema and references data from the two entries

above:

```yaml

--8<-- "examples/docs/references/multiple files/package.archive.yaml"

. . .

!!! warning "Attention"

You cannot create definitions that lead to circular loading of `\*.archive.yaml` files.

Each `definitions` section in an NOMAD entry represents a \*schema package\*. Each \*schema package\* needs to be fully loaded and analyzed before it can be used by other \*schema packages\* in other entries. Therefore, two \*schema packages\* in two entries cannot reference each other.

Conventions

## Conventions for labels

When assigning labels within your codebase, it's essential to follow consistent naming conventions for clarity and maintainability. The following guidelines outline the conventions for labeling different elements:

- \*\*Sections\*\*: Labels for sections should adhere to Python convention of CapitalizedCamelCase.

This means that each word in the label should begin with a capital letter, and there should be

no spaces between words. For example: `SectionLabelOne`, `SectionLabelTwo`.

- \*\*Quantities and Subsections\*\*: Labels for quantities and subsections should be in lower\_case. This convention involves writing all lowercase letters and separating words with whitespace. Abbreviations within these labels may be capitalized to enhance scientific readability. For example: `quantity label`, `subsection label`, `IV label`.