

The IUPAC FAIRSpec Specification, Version 0.0.1

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[draft version 2021.12.07]

public PDF version found in

<https://github.com/IUPAC/IUPAC-FAIRSpec/tree/main/documents/specifications>

history:

BH 2021.12.07 updating guiding principles to PAC article post-submission rev. 1

BH 2021.08.05 updating IFS.property and IFS.representation tags

BH 2021.07.28 reorganizing content, giving subsections to basic principles section

BH 2021.07.04 minor regex description fix

BH 2021.07.01 added 5. Data and Metadata Extraction and Serialization

BH 2021.06.30 added Figure 2, IFSProperty and IFSReference; makes IFSRepresentation abstract

BH 2021.06.29 editing...

BH 2021.06.28 editing...

BH 2021.06.27 editing...

BH 2021.06.26 created

Abstract

This document describes a specification for data management that allows for a seamless process from generation of experimental data through analysis to publication and public archiving. Throughout the process, the key FAIR aspects of findability, accessibility, interoperability, and reuse are maintained and emphasized.

The GitHub repository for this project is at <https://github.com/BobHanson/IUPAC-FAIRSpec>, and a set of sample datasets with associated IUPAC FAIRSpec Finding Aids can be found in <https://chemapps.stolaf.edu/iupac/ifs> with an interactive demo at <https://chemapps.stolaf.edu/iupac/demo/demo.htm>.

Introduction

It is becoming more and more common for funding agencies to require that data involved in a study be managed using FAIR principles and ultimately made available in association with publication. To date, these requirements have been fulfilled with only minimal consideration for how data might be found and used by the broader science practitioner and science education community. It has been considered enough to produce what is effectively a "data dump" -- a monolithic, sometimes enormous, ZIP file containing hundreds or thousands (or worse, just a single) file that somehow is to be used by others.

The goal of the [IUPAC Project 2019-031-1-024](#) is to enable a standardized way of managing spectroscopic data digital collections. This document sets forth a set of standards for the description and cataloging of data and its associated metadata in ways that are practical, relatively simple to implement, modular, intuitive, easily extended, and flexible in terms of requirements for the data itself as well as format the metadata can be expressed in.

The scope of the project is *spectroscopic* data within a *chemical* context, but what is described here is fully extensible to any sort of data, within any context, and is already being proposed in the area of materials science. [Simon Cole]

The proposed standards involve **aspects**:

- a set of principles underlying what we mean by "FAIR" in relation to spectroscopic data.
- a recommendation for the **organization** of digital objects within a collection,
- a standard for **describing properties** of digital objects within the metadata records of the finding aid,
- a standard for the **serialization of the finding aid** for a collection,
- a proposal for methods of **data and metadata extraction** and the generation of FAIRSpec Finding Aids.
- a detailed **data model** for describing the contents of a "spectroscopic data collection" in terms of objects and relationships of objects,

1. Basic IUPAC FAIRSpec Principles

Spectroscopic data in the area of chemistry are intimately connected to chemical structure. The whole purpose of carrying out spectroscopic measurements is generally to discover the identity of pure compounds, to identify and quantitate the relative amounts of chemical impurities in a mixture, and to determine the structure of new compounds.

In addition, there are contexts where it is more appropriate to refer to spectroscopic data in relation to "samples." We want to know the identity of a sample, but we don't yet know its chemical structure. Or, perhaps we are working with a material that is not characterizable *per se* as a chemical compound. These standards cover such cases as well.

Spectroscopy datasets can be complex. We make the distinction here between *Digital Entities* (sequences of bytes, for example "files"), which, through metadata association, become *Digital Objects*. [def reference] A key concept in the IUPAC FAIRSpec data model is the **IUPAC FAIRSpec Digital Collection**, which uses metadata to make useful connections between digital objects and allows for a variety of "representations" of those objects (Figure 1.1).

One to One and One to Many FAIR Relationships

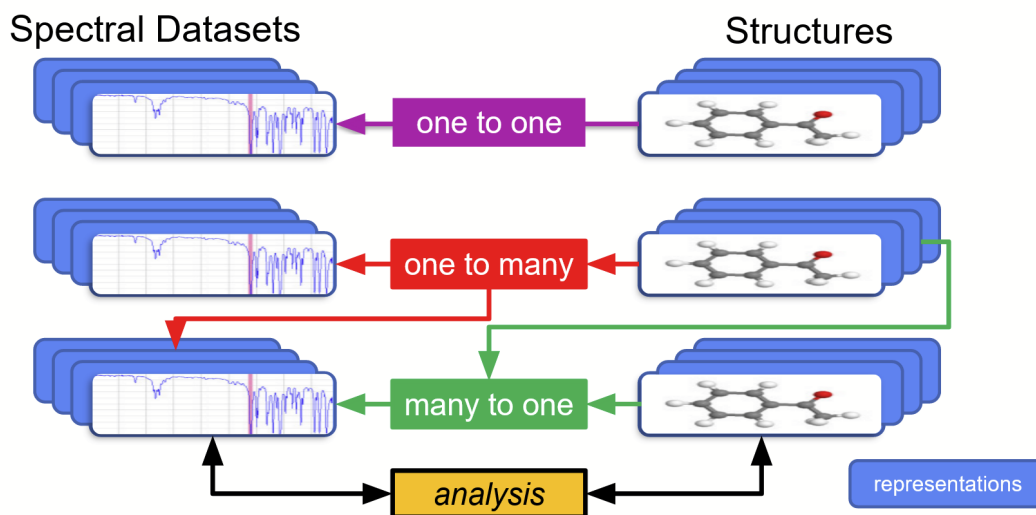


Figure 1.1. Key relationships among FAIRSpec Digital Objects.

Thus, the IUPAC FAIRSpec Standard data model is not about standardizing instrument data file formats (though it benefits from that) or requiring specific representations (though it might make specific recommendations in this regard). Rather, it is about providing a baseline common ground for the storage, transmission, and description of the contents of a digital collection derived from spectroscopic measurements or calculations. The ultimate goal is to be able to provide concise descriptions of complex data sets associated with manuscripts, ongoing laboratory work, and teaching, that can be findable, accessible, interoperable, and reusable by machines and humans alike.

The following principles underlie our understanding of the IUPAC FAIRSpec standards:

1. FAIR Management of data should be an ongoing concern.

- A. FAIR management of data must be an explicit part of research culture. FAIR data management should be a part of the design of all ongoing scientific endeavors from the very beginning and throughout its course, not something that is applied just at the time of publication. The key word here is *management*. All too often data are collected and organized in an arbitrary and idiosyncratic way. Management of the data is an afterthought, if at all. The IUPAC FAIRSpec standard is designed to be simple, practical, and, for the most part, automated, in order for it to be adopted and utilized throughout the data workflow, from collection to publication and beyond.
- B. FAIR management of data should be of intrinsic value. The value of spectroscopic data management has to have real-time value for the originating research group(s), providing value-added benefits within and between those groups that go far beyond the need to satisfy granting agencies.
- C. Good data management requires distributed curation. By **curation** we mean the association of meaning to a digital aggregation, thus turning its **digital entities** into *digital objects* and making them findable. Some of this curation can be done automatically; some must be done by hand. Simply designing an electronic filing system for data or associating data with an electronic laboratory notebook (ELN) are examples of curation. While much of what we describe here can be discovered programmatically, ultimately it is the task of researchers themselves to properly prepare and maintain their data and to make the connections between spectroscopic data and chemical structure or sample identity that make for intrinsic value. The FAIRSpec standard is designed to allow for this sort of management from the very beginning of data collection, allowing for data representations that are vendor specific, for example.
- D. Experimental work is by nature iterative. FAIR management of data associated with a research endeavor must allow for cycles of data *generation*, (re)*processing*, and (re)*analysis*. The IUPAC FAIRSpec standard is designed to allow a progression of associations from sample to structure.

2. Context is important.

- A. Digital objects are generally part of a collection. Spectra are taken for a reason, generally associated with a grant or project. They are maintained in relation to a student, researcher, research group, or institution. They are presented as part of a publication or presentation. Taking a cue from the field of digital archiving, FAIR management of data standards should emphasize the value of a *collection*, however that may be defined. A FAIR data management standard should describe the relationship among the different components of a collection, making sure that the finding of one component can lead to the finding of the whole.
- B. Chemical properties are related to chemical structure. This includes spectroscopic properties as discerned from the collection and analysis of spectroscopic data. This is why interpretation of spectra from structure and *vice versa* is an integral part of undergraduate chemistry courses. As such, spectroscopic data without reference to chemical identifiers of some kind (a compound name, a drawing, an InChI or SMILES), is unlikely to be useful to anyone, including its creator. The IUPAC FAIRSpec standard optimizes the connection between structure and spectra.
- C. Data relationships are diverse and develop over time. The association of spectroscopic data with chemical structure is not something that happens automatically or immediately. Indeed, initially there is a specific *sample* - generally the result of experimentation. The process involves the collection of spectroscopic data associated with that sample in a "one to many" relationship. That is, for any given sample, there may be several associated spectroscopic

data items. FAIR management of data should allow for contexts in which it may not yet be possible (or not ever possible) to connect a specific chemical structure to a spectrum and, by association, structure to sample. In addition, a "many to one" relationship must also be possible, either because the sample is a mixture or because the material involved does not lend itself to such a single-structure description in the first place.

- D. FAIR management of data should allow for validation. When data is made available in machine-readable form, it allows for the possibility of automated processes to be developed for use in validation of the data before, during, and after publication. Simply publishing an image of a spectrum in a PDF document does not allow any such validation. One of the most important potential impacts of the IUPAC FAIRSpec standard is that it will allow not just the routine validation of spectral data in relation to proposed structure, but also a potentially enormous data resource for the development of better validation tools, particularly in the area of NMR spectroscopy.

3. FAIR management of data requires curation.

- A. Data reuse relies upon practical findability. A 180 MB zip file containing 1200 files may be "findable," but unless there is a set of key registered metadata describing the data within that zip file, just having the file in hand does not constitute "found". Again, drawing from the area of digital archiving, a FAIR data management standard should describe a digital finding aid that exposes key metadata and allows a **reuser** to quickly ascertain whether additional scrutiny of the data collection is warranted, and/or the ability to conduct a rich fielded search of the database store where the metadata describing the collection have been registered.
- B. Data has to be organized to be accessible. It is probably obvious that data cannot be reused unless it is accessible. This accessibility can be seen at two levels. The IUPAC FAIRSpec standard does not assume that data are **open**. Owners of data have the right of restricting access to their data under whatever contract it was collected. But even for authorized reusers, data can be effectively inaccessible. For example, the reuser should not have to download and unpack a zip file containing a spectrum that is within a zip file for a compound that itself is within a zip file associated with a publication just to check to see if that spectrum is of immediate value. Thus, the IUPAC FAIRSpec standard allows for the automated or semi-automated repackaging of data and metadata extraction from an original dataset in order to provide a better reuse experience. Accessibility also means having sufficient information in the metadata elements to allow unsupervised machine-based retrieval of the data from a complex collection.
- C. Data interoperability requires well-designed metadata. The IUPAC FAIRSpec standard must be clearly defined and, as much as possible, mappable onto other metadata standards (e.g., bibliographic standards or instrument identifiers) that are in use or will be in future use in the area of machine-based knowledge discovery. That is, truly "FAIR" as in "Fully Artificial Intelligence Ready".
- D. Value is in the eye of the reuser. The IUPAC FAIRSpec standard respects the fact that data can have multiple *representations*. One might argue that only the "raw" data from an instrument – a free induction decay in the case of NMR spectroscopy – is the essential representation of the data that would allow for those data to be described as FAIR. However, that is not necessarily the case. The *reuse* of data relies upon data being in a form that is meaningful *for the reuser*. For a practicing spectroscopist, it may be that retrieving the raw spectroscopic dataset is critical, but for others - reviewers, readers, students - the "real" 1D or 2D spectrum may be the only representation they are able to utilize. Sometimes an image is more appropriate, and sometimes just a simple peak list is desired. Key to this principle is that the full nature of reuse cannot be predicted by the data originator.

4. Metadata must be registered and standardized.

- A. Register key metadata. A certain amount of "globally significant" metadata must be registered with a **metadata registration agency**. While the finding aid for a collection provides metadata that serve the purpose of describing the contents of the collection once the reuser

arrives at the site, full findability requires that the most salient aspects of the collection's metadata be separately registered with a registration agency, giving it an existence that is independent from the finding aid and, in fact, independent of the data itself, and allowing the metadata to be aggregated on a global scale. This registration requires dialog between registration agencies and standards developers to ensure that the sort of metadata being produced under the standards can be integrated into the schemas developed by those agencies.

- B. Assign a variety of persistent identifiers. All registered metadata should be assigned a persistent identifier. The persistent identifier serves to allow a persistent link to the **landing page** for data at some future point in time, regardless of whether or not the data are in their original location. The landing page is typically an HTML-based webpage designed for human access, whereas a digital finding aid in the form of XML or JSON may be more appropriate for machine-based or artificial intelligence-based applications. The fact that XML-based finding aids (with associated style sheets) can serve for both purposes has made them popular in the digital archival community. Thus, it could be that just the finding aid itself needs to be registered, with multiple entry points into a digital collection for subcollections of data objects. The most common form of a persistent identifier is the DOI, or **Digital Object Identifier**.
- C. Enable metadata crosswalks. Metadata must be standardized and allow for **metadata crosswalks** that allow the interoperability and exchange of metadata among different systems. The registration of metadata ensures that it will conform to a predictable structure specified by one or more **metadata schemas**. Currently popular schemas for scientific data include those established and curated by the DataCite organization, Schema.org, and the DDI Alliance. Such schemas may overlap to a considerable extent, but each has nuances that make it especially suited to particular needs or interests. For example, Schema.org was launched by several large commercial search engine providers to help improve the quality of their own data indexing, and the focus of the DDI Alliance is social, behavioral, economic, and health sciences. The process of mapping the metadata from one schema to another, referred to as "crosswalking", allows metadata standardized for one schema to be also representable by another. The IUPAC FAIRSpec standard must be mappable to one or more of these schemas, at least in part.
- D. Allow for value-added benefits. Once one or more digital finding aids are registered with persistent identifiers (e.g., DOI's or handles), they become part of a **metadata store**. Registering the finding aid with a DOI will enable it to become part of a queryable system of metadata, which can undergo **metadata harvesting** to provide a range of value-added services. For example, a service might index the metadata in ways that make it searchable on a much larger scale. Citation and access statistics can be provided, and **PID graphs** can be constructed that provide information about how one digital collection is related to another. A PID graph can be used to discover connections between diverse entities in the research landscape, connecting foundations and instrument manufacturers with information about their market and impact. But for that to be possible at the granularity that we require, harvesting methods must allow for standardized discipline-specific metadata such as we are proposing in this project to be represented in discipline-aware ways.

5. FAIR data management standards should be *modular, extensible, and flexible*

- A. Modularity allows specialization. By this, we mean that it should allow for core standards to be developed in different subdisciplines in parallel and sequentially, with different emphases and nuances that are unique to those subdisciplines. The goal of the IUPAC FAIRSpec project is to provide the templates for current and future subdisciplines to expand upon in whatever way is appropriate to their area.
- B. Design to adapt to future needs. FAIR data management standards should be extensible. That is, they should express clear versioning and allow for the inclusion of metadata descriptions that will meet future needs.
- C. Respect digital diversity. Taking another cue from the digital archivist community, FAIR data management standards should respect variety and should not require (though IUPAC might

recommend) one data format over another. Our motto is: *The collection is what it is. Work with it.* The IUPAC FAIRSpec standard must allow for workflows that start with data aggregations that contain a wide variety of possible data formats and organizational structure. Thus, while standardized, open, nonproprietary data formats are ideal in many contexts, the IUPAC FAIRSpec standard does not assume knowledge of a reuse context and does not in any way require any specific data format, open or otherwise. What is required is that the format of the data be identified. In addition, the exact format of the rendering of IUPAC FAIRSpec finding aids is not defined, allowing for different modes of expression, such as JSON or XML. This perspective is an 'added value' from IUPAC that facilitates interoperability across sectors, users, and disciplines.

- D. *All data formats should be valued.* To whatever extent is possible, preserving the original native instrument format of data is to be valued. This value derives from the need to have a clear pathway of ***data provenance***, minimizing the loss of information and context, maximizing the ability of others to verify the data and analyses, and allowing modified analysis that can be used in contexts not considered by the original data creators. Unlike specifications that transform raw data into other formats such as JCAMP-DX, nmrML, or NMR-STAR, the IUPAC FAIRSpec standard accepts data in native instrument formats (as well as these derived formats), opting instead for automated metadata and data extraction in order to provide more generally reusable data representations of that raw data.

2. Preferred Collection Organization

2.1 Structured data collections

The first step to good data management is having a well-defined and clearly structured data collection. This is the first step of curation. The **IFS Finding Aid** is designed to be able to handle collections that are organized in the form of *file systems*. For example, these could be directories on a local or host computer, remote repositories, ZIP files, or entirely abstract references into a relational database. Whether these are arranged hierarchically, in directories of directories, or as a "flat" list of entries (as is the underlying case in all ZIP files, actually), is completely up to the implementer. What is important, however, is that there be clear connections between structure and spectra. In the simplest form, this can be simply a hierarchical directory structure that can be interpreted readily by human and machine.

For example, Figure 2.1 shows the layout of a well-structured starting-point collection that could easily be transformed into an IUPAC FAIRSpec data collection and described by an [IFS Finding Aid](#). While not perfect, the context can reasonably inform the (next) curator that these are the compound numbers associated with the [acs.orglett.0c0051](#) supporting information [dataset](#) collection.

```
graph TD
    Root[acs.orglett.0c0051] --> FID[FID for Publication]
    FID --> 1c[1c]
    FID --> 1d[1d]
    1c --> 13C_1c[13C-NMR]
    1c --> 1H_1c[1H-NMR]
    1c --> HRMS_1c[HRMS]
    13C_1c --> 81[81]
    1H_1c --> 80[80]
    HRMS_1c --> PDF_1c[68075_mari0099_maxis_pos.pdf]
    HRMS_1c --> Mol_1c[1c.mol]
    1d --> 13C_1d[13C-NMR]
    1d --> 1H_1d[1H-NMR]
    1d --> HRMS_1d[HRMS]
    HRMS_1d --> Mol_1d[1d.mol]
    Root --> 3a[3a]
    Root --> 3b[3b]
```

Figure 2.1. A well-structured collection of digital entities ready for repackaging as an IUPAC FAIRSpec data collection.

2.2 Describing a collection - The IFS Finding Aid

Given a collection such as shown in Figure 2.1, the job of the curator (human or otherwise) is to produce a guiding metadata object that describes the contents of the collection to whatever extent is possible -- or at least *reasonable*. The IFS Finding Aid specification starts by cataloging the contents of a data collection based on file names. **IFS Finding Aid** notation allows for a combination of zip files and directories. The "pipe" symbol "|" indicates "entering a zip file", whereas the "forward slash" symbol "/" indicates moving into a subdirectory. Together, any sort of combination of nested directories and zip files can be described. For example, this string:

FID for Publication/1c.zip|1c/13C-NMR.zip|13C-NMR/81/

(from the <https://doi.org/10.1021/acs.orglett.0c00571> supporting information FAIR Data collection) describes a resource that is a directory of files (13C-NMR/81/) within a zip file (13C-NMR.zip) that itself is in a directory (1c/) contained in a zip file (1c.zip) that is in a subdirectory (FID for Publication/), which, by the way, is actually the top directory in the zip file submitted by the authors.

It is the job of the curator of the collection to determine whether this is appropriate or not. While it looks well organized, from a reusability perspective, is it optimal? Is this the best we can do? Should the reuser have to open zip file upon zip file upon zip file to retrieve a given spectrum? Probably not. Then again, it is one single download for a whole publication. So it certainly has value.

Additionally, this is a fine way for data to be *ingested* into a digital collection. It is well organized, and it has natural chemist-friendly descriptors. I can guess that if I drill down to this resource, I will probably get a 13C NMR spectrum for compound 1c as described in this publication. Whether or not that is a Bruker dataset or a Varian dataset, I don't know, since both of those instrument formats involved directories. The objective, of course, is to not have to guess about anything. Still, it's not bad, at least for ingestion.

2.3 Data and metadata extraction

The key to proper curation is that it enables additional curation by others. It is quite likely that additional representations might be added some time after, for example, an author has delivered a dataset to a publisher. We call this **data and metadata extraction**. For example, if a MOL file is found, it certainly has intrinsic value of its own at the level of a finding aid. And using available software tools and services, such as Jmol and NCI/CADD, that MOL structure can lead to an InChI, and InChIKey, several flavors of SMILES representations, conversion from 2D to 3D or the reverse, and the generation of images of the chemical drawing for the structure. NMR data sets in proprietary or open vendor formats might be able to be standardized as JCAMP-DX, nmrML, or NMR-STAR formats, *in addition* to being made available in their original format.

Our recommendation is as follows:

1. The originally ingested (creator-produced) format should be preferably one but not more than a few ZIP files or repository references (that can be downloaded as ZIP files).

2. This original collection should be preserved and always made available when a subset from that collection is delivered.
3. The collection should be "extracted" for data and metadata that is recognizable and of perceived "high-level" searchability value. That decision, of course, is contextual. The IUPAC FAIRSpec data management standard allows for a wide range of detail, customizable to the implementer of the standard and the overall context of the data service involved.
4. Data sets should be deliverable as ZIP files or their equivalent (TAR, for example). This again, though, is totally up to the implementer of the standard.
5. Representations of data (images, PDF files, etc.) should be extracted and catalogued as **IFS Representations** or **IFS Property** values associated with **IFS Objects**. The **IFS Finding Aid** associated with this collection should report as many of these properties and representations as deemed appropriate to the implementer.
6. It is preferable to make all of these Digital Objects retrievable individually, whether the original hierarchy is maintained or not.
7. It is **highly recommended** that any data ingest include SMILES and/or a 2D or 3D MOL file for each structure entry. "Structures" generated by drawing programs, unless done properly, are not sufficient. (For example, even standard protecting group names such as "Ac" and "TMS" cause problems with automated systems.) InChI and InChIKey are also desirable, but they are not in and of themselves sufficient, unless they can be "[resolved](#)" to an actual molecular structure.

3. The IUPAC FAIRSpec Metadata Model

Metadata "records" associated with IUPAC FAIRSpec datasets involve standardized IFS property names and associated (controlled-vocabulary or free-text) property values. Names start with "IFS.property" and continue hierarchically with categories and subcategories. Similarly, IFS representations have names start with "IFS.representation." Except for a very few parameters that are associated with the collection as a whole, these parameters are all described within spectroscopy-specific categories as "IFS.property.spec.*" or "IFS.representation.spec.*".

To date, the list is rather short, but it will grow.

3.1 IFS Properties

3.1.1 Collection properties

IFS.property.collection.id
IFS.property.collection.len
IFS.property.collection.object
IFS.property.collection.ref
IFS.property.collection.source.data.uri
IFS.property.collection.source.publication.uri

3.1.2 Spectroscopic Data Properties

IFS.property.spec.hrms.expt.label
IFS.property.spec.ir.expt.label
IFS.property.spec.ms.expt.label
IFS.property.spec.nmr.expt.dim
IFS.property.spec.nmr.expt.freq.1
IFS.property.spec.nmr.expt.freq.2
IFS.property.spec.nmr.expt.freq.3
IFS.property.spec.nmr.expt.label
IFS.property.spec.nmr.expt.nucl.1
IFS.property.spec.nmr.expt.nucl.2
IFS.property.spec.nmr.expt.nucl.3
IFS.property.spec.nmr.expt.pulse.prog
IFS.property.spec.nmr.expt.solvent
IFS.property.spec.nmr.expt.temperature.K
IFS.property.spec.nmr.instr.freq.nominal
IFS.property.spec.nmr.instr.manufacturer.name
IFS.property.spec.nmr.instr.probe.type
IFS.property.spec.raman.expt.label
IFS.property.spec.uvvis.expt.label

3.1.3 Chemical Structure Properties

3.1.4 Chemical Sample Properties

3.2 IFS Representations

3.2.1 Spectroscopic Data Representations

IFS.representation.spec.hrms.spectrum.description
IFS.representation.spec.hrms.spectrum.image
IFS.representation.spec.hrms.spectrum.pdf
IFS.representation.spec.hrms.vendor.dataset

IFS.representation.spec.ir.jcamp
IFS.representation.spec.ir.peaklist
IFS.representation.spec.ir.spectrum.description
IFS.representation.spec.ir.spectrum.image
IFS.representation.spec.ir.spectrum.pdf
IFS.representation.spec.ir.vendor.dataset

IFS.representation.spec.ms.jcamp
IFS.representation.spec.ms.peaklist
IFS.representation.spec.ms.spectrum.description
IFS.representation.spec.ms.spectrum.image
IFS.representation.spec.ms.spectrum.pdf
IFS.representation.spec.ms.vendor.dataset

IFS.representation.spec.nmr.jcamp.fid.1d
IFS.representation.spec.nmr.jcamp.fid.2d
IFS.representation.spec.nmr.jcamp.spec.1i1r.1d
IFS.representation.spec.nmr.jcamp.spec.1r.1d
IFS.representation.spec.nmr.jcamp.spec.2d
IFS.representation.spec.nmr.peaklist
IFS.representation.spec.nmr.spectrum.description
IFS.representation.spec.nmr.spectrum.image
IFS.representation.spec.nmr.spectrum.pdf
IFS.representation.spec.nmr.vendor.dataset

IFS.representation.spec.raman.jcamp
IFS.representation.spec.raman.peaklist
IFS.representation.spec.raman.spectrum.description
IFS.representation.spec.raman.spectrum.image
IFS.representation.spec.raman.spectrum.pdf
IFS.representation.spec.raman.vendor.dataset

3.2.2 Chemical Structure Representations

IFS.representation.struc.cdx
IFS.representation.struc.cdxml
IFS.representation.struc.mol
IFS.representation.struc.mol.2d
IFS.representation.struc.mol.3d
IFS.representation.struc.png
IFS.representation.struc.sdf
IFS.representation.struc.sdf.2d
IFS.representation.struc.sdf.3d
IFS.representation.struc.unknown

3.2.3 Chemical Sample Representations

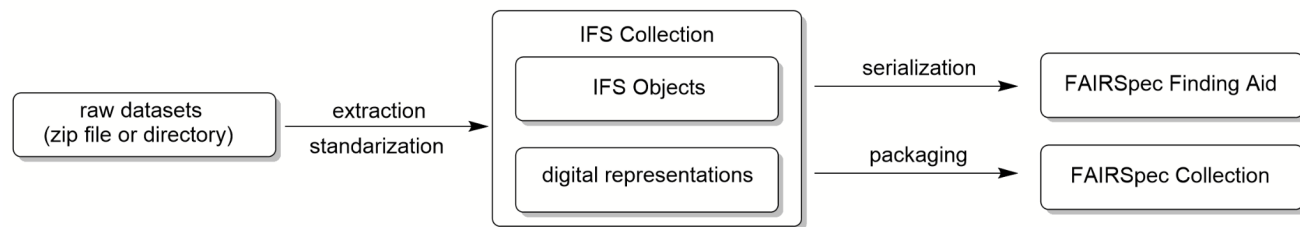
(todo)

4. Data and Metadata Extraction and Serialization

4.1 Workflow

The FAIRSpec Finding Aid is not expected to be created by hand. Rather, it is expected to be the end result of an automated or semiautomated *curation* process, much the way the finding aid in the area of digital archiving is created in that community. The general workflow is shown in Figure 4.1.

newer version (FAIRSpec-workflow.cdx)



more technical version:

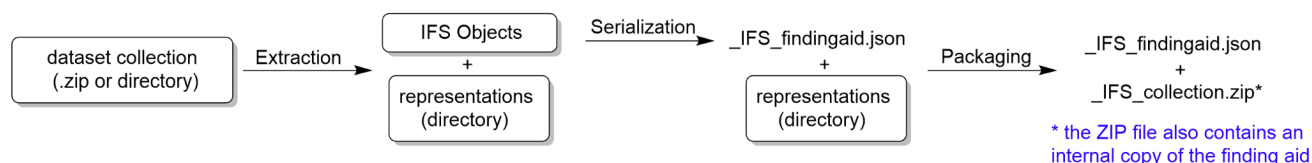


Figure 4.1. The expected workflow for creating IFS FAIRSpec data collections

The job of the Extractor software is to extract information into the IFS data model. During serialization, the Serializer is passed keys, attributes, and values along with serial arrays and key/value maps that need to be formatted according to the serializer's design. There is nothing particularly difficult about serialization -- the default JSON serializer is about 100 lines of code. Packaging simply involves adding a copy of the finding aid along with the representation directory and its contents to a ZIP file and possibly zipping that zip file up with the original finding aid (depending upon whether the intent is to deliver the directory or a zip file as the data associated with the finding aid).

Note that it is entirely possible that this workflow could be part of a standard operating procedure for a research lab. Spectra are taken in the course of sample analysis. If the structure is known (or even if it is not), an IFS Extractor is used to *insert* critical structural or sample information into the system, whether that simply be a directory on an NMR instrument's data drive, or some sort of cloud-based system. The IFS Extractor would maintain the finding aid for this private, local research collection. Basic web desktop tools could be used to display and work with the finding aid in the general course of experimentation, allowing the researcher to quickly find spectra, assign structures, and make structure-spectrum associations. Then, at the time of publication, perhaps, final compound numbers would be added, and the collection or (more probably) a subset of the collection could be extracted for release as supporting information. There would be no need for the manual construction of a "supporting information PDF" file. If that is required, software designed with the IUPAC FAIRSpec Specification in mind could produce that document automatically.

4.2 Data and Metadata Extraction

Extraction is the challenge. How do we identify and catalog the contents of a general dataset? Two general scenarios are envisioned:

- 1) An institution or portal has created datasets with IFS FAIRSpec in mind. In that case, extraction should be quite straightforward and is likely to require no human intervention.
- 2) A dataset collection in the form of a zip file with spectroscopic data and related structural or sample information associated with it is delivered by a research group to a publisher or repository. Depending upon how that was accomplished, this may take some human curation.

In either case, extraction involves some sort of data analysis. Here, we simply describe one possible solution, what we are calling the "reference IFS Extractor"

(com.integratedgraphics.ifs.Extractor.java). In this reference Extractor, the process starts with a small manually generated configuration document, *IFS-extract.json*, a sample of which is given below, and is referred to here as the *extractor script*:

```
{
  "IFS-extract-version": "0.1.0-alpha",
  "keys": [
    {
      "license": "{IFS.property.findingaid.data.license.uri::https://creativecommons.org/licenses/by-nc/4.0/},{IFS.property.findingaid.data.license.name::cc-by-nc-4.0}",
      "hash": "0c00770",
      "ifsid": "acs.joc.{hash}",
      "figshareid": "22567817",
      "puburi": "{IFS.property.findingaid.source.publication.uri::https://doi.org/10.1021/{ifsid}}",
      "data": "{IFS.property.findingaid.source.data.uri::https://ndownloader.figshare.com/files/{figshareid}}",
      "zip": "{data}{id=IFS.property.struc.compound.label::*.zip}",
      "objects": "{zip}{IFS.representation.spec.nmr.vendor.dataset::{IFS.property.spec.nmr.expt.label::*}}",
      "objects": "{zip}{IFS.representation.spec.nmr.vendor.dataset::{IFS.property.spec.nmr.expt.label::*.mnova}}"
    }
  ]
}
```

The Extractor's overall task is to carry out "regular expression" matches on the file names of the entries in the ZIP file or the files in a directory structure. The extractor script specifies how to process the zip file names. The values that the Extractor will need are for keys **license**, **ifsid**, **puburi**, and **objects** (of which there may be any number). It is only the **objects** that will be matched to the file names in the zip file. **license** and **puburi** are scanned for particular values, and **ifsid** is used to name the IFSFindingAid directly. All the rest (*hash*, *figshareid*, *data*, and *zip*) are just definitions chosen by the writer of the script to make it more generalizable or understandable to a human reader.

The extraction notation used is relatively straightforward: Braces { } start and end a definition. Two colons "::" separate the name of an IFS FAIRSpec standard parameter name and its value. In the license value, we are assigning the special IFSFindingAid properties

IFS.property.findingaid.data.license.uri to be <https://creativecommons.org/licenses/by-nc/4.0/> and **IFS.property.findingaid.data.license.name** to be **cc-by-nc-4.0**. The publication URI is assigned the value <https://doi.org/10.1021/acs.joc.0c00770>.

An additional option is to give a name to a matched part of the file name. The syntax for this involves adding "key=" before the parameter name, and then using <key> when that is needed. For example:


```
{ "path": "{data}|FID for Publication/{id=IFS.property.struc.compound.label::*}.zip|",  
  "objects": "{path}<id>/{IFS.representation.struc.mol.2d::<id>.mol}" },
```

defines the name of the zip file to be **id** and then looks for that in the name in two places. This particular example would match a file by the name of "3a/3a.mol" in a zip file (the pipe symbol) by the name of "3a.zip" in the "FID for Publication" directory of the zip file (another pipe symbol) indicated by {data}. In addition, it would identify the mol file itself as a 2d MOL structure representation, and will assign the value of IFS.property.struc.compound.label for that structure to be "3a". Note that "{key=::}.....{key}", without a property name, also works.

The asterisk here is one of the few special characters recognized by the default Extractor:

*	any characters up to but not including or /
**	and characters including and /
<char>	any number of *<char>*, so "*-*.pdf" would match "2-1H.pdf" or "2-b-1H.pdf"
{regex::....}	any standard Java regex pattern such as {regex::[a-z0-9]}.

It's no more complicated than that. Anything that is missing should be representable using {regex::....}. All the extractor is doing is transforming all this into standard Java regex patterns and then applying that to the file name.

Examples of extraction scripts can be found in the extract/ directory at the GitHub site.

4.3 IFSSpecDataFindingAid Serialization

An **IFSSpecDataFindingAid** object, in order to be useful, must be *serialized*. That is, it must be represented as a Digital Object that can be stored digitally, read by an appropriate computer program. The description here involves a JSON serialization, but that could just as well be XML or some other form of serialization. The reference implementation contains a default JSON serializer, but an implementer can relatively straightforwardly subclass that to produce another sort of serialization.

-- TODO --

5. The IUPAC FAIRSpec Data Model

5.1 Definitions

What follows is a detailed discussion of the data and metadata model underlying the IUPAC FAIRSpec specification. The description of the model given here borrows the language of Java. This allows the discussion to be framed within an existing object-oriented standard (Java 8). And, as it turns out, the reference implementation is written in Java (and, through co-transpiling, deliverable as both Java and JavaScript). The following terms are drawn from Java, but they are applicable to whatever context the data model is expressed:

- *object*

A construct containing *fields* (variables) and *methods* (functions that take inputs, modify the fields of one or more objects, create new objects, and return values).

- *class*

A description of an object's type, expressing its capabilities and limitations. Essentially a template. One says that an object is an "instance" of its class, or that "instantiation" of a class creates an object. In the reference implementation, all classes start with "IFS". In this document, classes are shown in **bold**.

- *subclass* and *superclass*

The "child" of a *superclass* is a *subclass*. We say that a subclass *extends* its superclass so that it can "inherit" the superclass's fields and methods. This efficient design allows general characteristics to be described in one class (a superclass) and propagated to a whole family of other classes (its subclasses). It allows for rational future expansion via subclassing. So, for example, we will see ***IFSCollection*** as a general concept, and ***IFSStructureCollection*** and ***IFSSpecDataCollection*** as unique specializations of that concept. These two types of collection share all the characteristics of an ***IFSCollection***, but differ in their details. And ***IFSCollection*** is itself a subclass of ***IFSObject***, so anything we can say about ***IFSObject*** also relates to ***IFSCollection***, ***IFSStructureCollection***, and ***IFSSpecDataCollection***. For example, they all have a *name*, a *type*, and a list of defined *properties*.

- *abstract class*

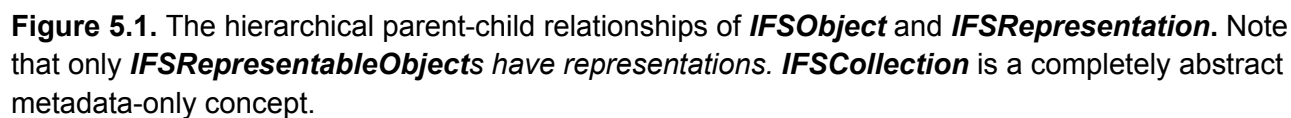
An abstract class is a class that can only be a superclass. It can never be instantiated itself. It is basically just a template that provides common fields and methods for its various subclasses. For example, ***IFSObject*** is an abstract class and cannot be instantiated, but it holds fields *name*, *type*, and *id* that all of its many subclasses have in common. Abstract classes in this document are given in ***bold italics***.

- *interface*

A description of the key methods of a class that are generally useful. Or just a tag that identifies a class as a certain kind of class. We say a class "implements" an interface. For example, ***IFSObject*** implements ***IFSSerializable***, meaning all ***IFSObject*** subclasses must have the two methods: *serialize* and *getSerializedType*. In the

IUPAC FAIRSpec reference implementation, all interfaces start with "IFS" and end with "I".

These classes are fundamental classes that have no specific reference to spectroscopy. They could easily be extended to non-spectroscopy FAIR data management applications. The two high-level superclasses in the IUPAC FAIRSpec Data Model are the abstract classes ***IFLObject*** and ***IFSRepresentation***. Almost all objects in the model subclass one of these two classes. The overall hierarchy of ***IFLObject*** and ***IFSRepresentation*** are shown in **Figure 5.1**. Two additional classes include **IFSPROPERTY** and **IFSReference**.



IFSOject

abstract class **IFSOject** extends **ArrayList** implements IFSOjectI, IFSSerializable

IFSOject is the primary abstract superclass for all IFS Data Model metadata objects. It holds the essential characteristics of what it means to be an **IFSOject**, specifically:

- *name* (some sort of name for this object)
- *id* (some sort of unique identifier)
- *path* (some sort of indicator where this object can be found in its collection)
- *properties* (values with standardized keys, with type and units)
- *params* (values without type or units that are outside the standardized system)
- *type* (one of a specific allowable types, such as *Structure*, *NMRSpecData*, or *Unknown*).
- *subtype* (some sort of qualifier to the type)

Note that **IFSOject.properties** and **IFSOject.params** both hold metadata. The *properties* field is intended for properties that are described by the standard, while *params* holds properties that are not (yet) part of the standard. This allows implementers to introduce properties that are important to their work without breaking the standard. Future versions of the standard may include those properties using standardized syntax, adding information about units and type. In addition, because **IFSOject** extends **ArrayList** (a Java class maintaining an array of objects that can be added to dynamically), all **IFSOjects** are inherently list-like. **IFSOject** and its subclasses come in two flavors: **IFSRepresentableObject** and **IFSCollection**, both of which are themselves abstract. **IFSRepresentableObjects** hold lists of **IFSRepresentations**; **IFSCollections** hold lists of other **IFSOjects**.

IFSRepresentation

abstract class **IFSRepresentation** implements IFSSerializableI

IFSRepresentation instances reference resources or contain byte sequence data themselves. These are the Digital Entities that are being described by metadata (the **IFSOjects**) in order to become Digital Objects. Note that **IFSRepresentations** are not **IFSOjects** and are not list-like. The essential characteristics of an **IFSRepresentation** include:

- *type* (one of a specific allowable types, such as *Structure*, *NMRSpecData*, or *UnknownI*)
- *subtype* (generally a media type, such as *image/png*)
- *ref* (the reference to this file in the form of an **IFSReference**)
- *data* (the bytes, if not referenced)
- *len* (the length of the data at this *ref* or in the *data* field.)

IFSRepresentation.data could be anything, but most likely will be of the type `String` or `byte[]` (byte array). These are the 2D or 3D MOL or SDF models, InChI or SMILES strings representing chemical structure, and the experimental, predicted, or simulated NMR or IR data associated with a chemical compound or mixture of compounds.

IFSPProperty

class **IFSPProperty** implements IFSSerializableI

IFSPProperty is the storage class for all **IFSObjects**. It holds just a few pieces of information:

- *name* (a controlled-vocabulary name starting with "IFS.")
- *dataType* (one of the enumeration PROPERTY_TYPE)
- *units* (one of the enumeration PROPERTY_UNIT)
- *value* (the value of this property, which can be anything of type dataType)

The syntax of a **IFSPProperty.name** is as follows:

IFS.property.(key.)*key(.UNITS)

where * means zero or more, () means optional, *key* is a lower-case alphanumeric string, and *UNITS* is an upper-case string matching **IFSPProperty.units**. For example:

IFS.property.spec.nmr.expt.solvent
IFS.property.spec.nmr.expt.temperature.K

As of this writing, PROPERTY_TYPE values include { INT, FLOAT, STRING, NUCL, OBJ }, and PROPERTY_UNIT values include { NONE, HZ, MHZ, CELCIUS, KELVIN }. These options are expected to expand substantially as the standard is developed.

IFSReference

class **IFSReference** implements IFSSerializableI

IFSReference is the storage class connecting **IFSRepresentations** to their Digital Objects. This class has only three fields:

- *origin* (a reference to the originating Digital Entity)
- *path* (a relative path to the directory containing this resource)
- *localName* (the name of this resource in its extracted form)

When the representation is a ZIP file entry, the path will contain a "pipe" vertical line symbol "|"; directories are represented with the standard forward slash. Thus, *origin* might be something like

1.zip|1_13C/pdata/1/thumb.png

Note that this notation allows for resources that are from zip files that are themselves found within zip files.

Generally speaking, the *localName* will not be part of a zip file (other than perhaps the zip file used for the collection's delivery), and may even have no hierarchical directory structure, either. For example:

1.zip_1_13C_pdata_1_thumb.png

Note that an **IFSRepresentation** may or may not have an origin object, and it may or may not have a local path. The lack of an *origin* indicates that an extractor has created this representation during extraction. For example, using just a SMILES string, it has created a 2D MOL file, a 3D MOL file, InChI, InChIKey, etc. The lack of a *localName* indicates that a representation's byte data are being saved within the **IFSRepresentation** object itself, not separately.

The details of these fields and their serialization will depend upon the implementing IFSExtractorI design.

IFSRepresentableObject

abstract class **IFSRepresentableObject** extends **IFSObject**

Includes: **IFSSample**, **IFSStructure**, **IFSDataObject**, **IFSStructureAnalysis**, **IFSSampleAnalysis**

A class extending **IFSRepresentableObject** is expected to maintain a list of one or more distinctly different digital representations (byte sequences) that amount to more than just metadata.

Note that there is no mechanism for **IFSSample**, **IFSStructure**, or **IFSDataObject** objects to refer to each other. This is a key aspect of the IFS Data Model. The model is based on the idea of a collection of (relatively) independent objects, and it is the **IFSCollection** objects that do this referencing. This model allows for a relatively simple and flexible packaging of objects, with their relationships identified only in key metadata resources.

IFSSample

class **IFSSample** extends **IFSRepresentableObject**

IFSSample corresponds to a specific physical sample that may or may not (yet) have a chemical structure, spectroscopic data, or representations associated with it. Its properties include a unique id (such as a laboratory notebook reference), as well as melting point or boiling point. Its representations could include a photographic image of the solid, for example.

IFSStructure

class **IFSStructure** extends **IFSRepresentableObject**

An **IFSStructure** is a chemically-related structural object, which may have several representations in its list, such as a 2D or 3D MOL file, **PDB?**, an InChI or InChIKey, one or more chemical names, one or more SMILES strings, or even just a PNG image of a drawn structure. Each of these representations serves a purpose. Some are more "interoperable" than others, but each in its own

way may be more useful in a given context. An implementer would have the option to include these representations as data within the finding aid or as an *IFSRepresentation* reference to a resource within the collection.

IFSDataObject

abstract class *IFSDataObject* extends *IFSRepresentableObject*

The core *IFSDataObject* class references lists one or more Digital Object representations -- what a scientist would call their "data", such as a full vendor experiment dataset (a Bruker NMR experiment), a PNG image of a spectrum, or a peaklist. For the IUPAC FAIRSpec Project, we extend this class to *IFSSpecData* and its subclasses (*IFSNMRSpecData*, *IFSIRSpecData*, etc.). We recognize, however, that the system we are developing is not limited to spectroscopic data only, and future implementations of this data model may subclass *IFSDataObject* in completely different ways for completely different purposes.

IFSStructureAnalysis

abstract class *IFSStructureAnalysis* extends *IFSRepresentableObject*

A subclass of *IFSStructureAnalysis* is intended to represent a detailed correlation between chemical structure for a compound and its related experimental or theoretical spectroscopic data. For instance, it might correlate specific atoms or groups of atoms of a chemical structure with specific signals in a spectrum or other sort of data object.

IFSSampleAnalysis

abstract class *IFSSampleAnalysis* extends *IFSRepresentableObject*

A subclass of *IFSSampleAnalysis* is intended to represent a detailed correlation between a specific chemical *sample* and its molecular structure and spectroscopic data. The details of this analysis would be designed into the subclass extending this class.

IFSCollection

abstract class *IFSCollection* extends *IFSObject*

IFSCollection objects are "pure metadata" and thus have no digital representation outside of that role. They point to and associate *IFSObject* instances. *IFSCollection* objects may be collections of *IFSCollection* objects. This allows for a nesting of collections in meaningful ways. For example, the subclass *IFSStructureDataAssociation* maintains two specific collections: one of structures, and one of data objects.

To be sure, an **IFSCollection** could represent a digital object in the form of a zip file (and usually does). Nonetheless, this does not make it "representable" in the sense that it is likely to have multiple distinctly different representations that characterize an **IFSRepresentableObject**.

IFSStructureCollection, IFSDataObjectCollection, IFSSampleCollection

class **IFSSampleCollection** extends IFSCollection

class **IFSStructureCollection** extends IFSCollection

abstract class **IFSDataObjectCollection** extends IFSCollection

These classes each collect distinctly different metadata relating to structures, data, and samples, respectively, that are related in some way. For instance, all the compounds referred to in a publication, all the spectra for a publication, or all the spectra relating to a specific compound. (Or, perhaps, all the compounds in a mixture that are identified in an NMR spectrum.)

5.3 Common Associative Classes

One of the key aspects of the IUPAC FAIRSpec Data Model is that it allows subclasses of **IFSCollection** to do all the referencing between **IFSObjects**. Two abstract superclasses that make these connections include **IFSStructureDataAssociation** and **IFSSampleAssociation**. These classes associate specific structures, samples, and data to each other in some meaningful way. In addition, the analysis-related classes (discussed above) and the **IFSFindingAid** class fall into this category.

IFSStructureDataAssociation

abstract class **IFSStructureDataAssociation** extends **IFSCollection**

This class is the key class for connecting a chemical structure with its spectroscopic data. It can represent all possible relationships: one-to-one (the structure associated with a spectrum), one-to-many (a set of spectra associated with a give structure), many-to-one (a set of structures describing a mixture associated with a spectrum), and many-to-many (a set of spectra associated with a mixture of compounds).

IFSSampleAssociation

abstract class **IFSSampleAssociation** extends **IFSCollection**

IFSSampleAssociation allows a single sample to be associated with any number of spectra and structure identifiers.

IFSFindingAid

abstract class ***IFSFindingAid*** extends IFSCollection

The key IUPAC FAIRSpec metadata object is the ***IFSFindingAid***, which constitutes the "master" metadata object for a collection. The ***IFSFindingAid*** is a special "collection of collections," providing:

1. metadata relating to the entire collection. For example, details about a publication or thesis.
2. high-level access to lower-level metadata. For example, a list of compound names or key spectroscopy data characteristics such as NMR nucleus (¹H, ¹³C, ³¹P) and spectrometer nominal frequency (300 MHz, 800 MHz) or type of IR analysis (such as ATR).
3. pointers to finding aids for subcollections, each of which may be a pointer to one or more additional finding aids.

It is the ***IFSFindingAid*** that ultimately distinguishes the IUPAC FAIRSpec data model from other models. It should contain all the information that forms the basis of what the user sees. It should reveal information about the collection that allows users to quickly determine whether data in this collection are relevant to their interests or not. The serialization of the ***IFSFindingAid*** object might be the target of a persistent identifier such as a DOI for the collection (if XML) or interpreted by that DOI's landing page (if JSON). Or, it could be dynamically created in response to a query, allowing standardized programmatic access to the collection.

The IUPAC FAIRSpec data model subclasses ***IFSFindingAid*** as ***IFSSpecDataFindingAid***. This class is discussed below.

5.4 Spectroscopy-Specific Classes

The above-mentioned classes are primarily abstract classes, meaning they can be used as the basis for other more specific classes, but they cannot themselves be instantiated. This instantiation is mostly carried out on subclasses in the data model's spectroscopy package. In this package we find specifics for the description of the various types of spectroscopy -- NMR, IR, UV-Visible, Raman, etc. This list may be extended to whatever extent necessary to cover the sorts of spectroscopy involved. The IUPAC FAIRSpec Standard specifies the identifiers used to extract information from "raw" data sets associated with these subclasses. In the reference implementation, these classes are all found in the *org.iupac.fairspec.spec* Java package:

Note that there are no references whatsoever to the *spec* classes from any class in any other package in the reference implementation. This is important. It means that any specialization or extension that involves spectroscopic details can be implemented within just this single *spec* package. This class reference isolation is a very important aspect of the overall design of the IUPAC FAIRSpec data model.

Included in the *spec* package are classes that subclass ***IFSSampleAnalysis***, ***IFSStructureAnalysis***, ***IFSSampleDataAssociation***, ***IFSStructureDataAssociation***, and ***IFSDataObject***, providing fields and methods specific to spectroscopy.

In addition, each spectroscopic type (nmr, ir, etc.) has its own specialized classes that subclass ***IFSSpecDataObject*** and ***IFSSpecDataRepresentation***. It is the various ***IFSSpecDataObject*** subclasses that specify the IFS FAIRSpec standards for a given type of spectroscopy. For example, in ***IFSNMRSpecDataObject*** we find specifications for syntax:

```
public static final String IFS_SPEC_NMR_INSTR_MANUFACTURER_NAME = "IFS.spec.nmr.instr.manufacturer.name";
public static final String IFS_SPEC_NMR_INSTR_FREQ_NOMINAL = "IFS.spec.nmr.instr.freq.nominal";
public static final String IFS_SPEC_NMR_INSTR_PROBEID = "IFS.spec.nmr.instr.probe.type";
public static final String IFS_SPEC_NMR_EXPT_DIM = "IFS.spec.nmr.expt.dim";
public static final String IFS_SPEC_NMR_EXPT_FREQ_1 = "IFS.spec.nmr.expt.freq.1";
public static final String IFS_SPEC_NMR_EXPT_FREQ_2 = "IFS.spec.nmr.expt.freq.2";
public static final String IFS_SPEC_NMR_EXPT_FREQ_3 = "IFS.spec.nmr.expt.freq.3";
public static final String IFS_SPEC_NMR_EXPT_NUCL_1 = "IFS.spec.nmr.expt.nucl.1";
public static final String IFS_SPEC_NMR_EXPT_NUCL_2 = "IFS.spec.nmr.expt.nucl.2";
public static final String IFS_SPEC_NMR_EXPT_NUCL_3 = "IFS.spec.nmr.expt.nucl.3";
public static final String IFS_SPEC_NMR_EXPT_PULSE_PROG = "IFS.spec.nmr.expt.pulse.prog";
public static final String IFS_SPEC_NMR_EXPT_SOLVENT = "IFS.spec.nmr.expt.solvent";
public static final String IFS_PROP_SPEC_NMR_EXPT_ID = "IFS.property.spec.nmr.expt.label";
```

and definitions that include units and data types:

```
super.setProperties(new IFSPROPERTY[] {
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_INSTR_MANUFACTURER_NAME),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_INSTR_FREQ_NOMINAL, IFSCONST.PROPERTY_TYPE.INT,
        IFSCONST.PROPERTY_UNITS.MHZ),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_INSTR_PROBEID),
    new IFSPROPERTY(IFSNMRSpecData.IFS_PROP_SPEC_NMR_EXPT_ID),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_DIM, IFSCONST.PROPERTY_TYPE.INT),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_PULSE_PROG),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_SOLVENT),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_FREQ_1, IFSCONST.PROPERTY_TYPE.INT,
        IFSCONST.PROPERTY_UNITS.MHZ),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_FREQ_2, IFSCONST.PROPERTY_TYPE.INT,
        IFSCONST.PROPERTY_UNITS.MHZ),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_FREQ_3, IFSCONST.PROPERTY_TYPE.INT,
        IFSCONST.PROPERTY_UNITS.MHZ),
    new IFSPROPERTY(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_NUCL_1, IFSCONST.PROPERTY_TYPE.NUCL,
        IFSCONST.PROPERTY_UNITS.NONE),
```

```

new IFSPProperty(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_NUCL_2, IFSCnst.PROPERTY_TYPE.NUCL,
    IFSCnst.PROPERTY_UNITS.NONE),
new IFSPProperty(IFSNMRSpecData.IFS_SPEC_NMR_EXPT_NUCL_3, IFSCnst.PROPERTY_TYPE.NUCL,
    IFSCnst.PROPERTY_UNITS.NONE)
});

```

The class system is completely modular. Using Java, if one were to introduce a new spectroscopic method, such as x-ray crystallography, one would simply:

1. create a new package `org.iupac.fairspec.spec.xray`, (*note capitalization*)
2. copy and adapt two new specialized classes, `IFSXRAYSSpecData.java` and `IFSXraySpecDataRepresentation.java`, adding properties as desired, and
3. fill in the template methods and fields as desired