







# **BabelFSH**

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### **About us**

- "Module 2B Project" in the Medical Informatics Initiative: advancing the harmonization of health data and IT solutions at the university hospital sites in cooperation with the NUM
- Project funded 2023-2026 with three partner sites
  - University of Luebeck, Josef Ingenerf
  - University of Cologne, Oya Beyan & Andreas Beyer
  - Hanover Medical School, Michael Marschollek
- Goal: provide and support a central terminology server to DICs and the MII/NUM in general, to support semantic interoperability
  - Support for CDS designers in the MII for the development of the CDS
  - Support for DICs to provide a reference server for terminology used in MII/NUM



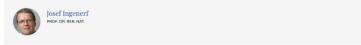
















### Köln



Andreas Beyer PROF. DR.



Oya Deniz Beyan











### Hannover



Michael Marschollek







### Alumni









### The Problem

- Using FHIR Terminology server operations requires access to FHIR CodeSystem resources
  - Except for SNOMED CT and LOINC, but that's a different story
- Many terminological resources aren't natively FHIR
  - [BfArM resources, i.e. ICD-10-GM, ICD-10-WHO, ICD-0-3, ICF, [Orphanet], Alpha-ID-SE,
     OPS]: ClaML and CSV
  - ATC: Excel sheets
  - EDQM Standard Terms: API
  - OncoTree: API or RDF
- Some CodeSystems also "define" ValueSets and/or ConceptMaps
- Conversion to FHIR requires the same thing being done over and over again
  - o Define Metadata of the resource (-s), sometimes cleanly, sometimes hard-coded
  - Add content (concepts for CS, includes for VS, elements for CM) to the resource (-s),
     domain-specific fashion
  - Write out a FHIR JSON resource and/or upload to FHIR server.









# FSH and SUSHI, briefly

- FSH = FHIR Shorthand; SUSHI Unshortens Short Hand Inputs
- FSH used for definition of profiles and IGs (mainly)
  - o In-development specification for a domain-specific language
  - o R2 of FSH released in Feb 2022; ongoing development for FSH 3.x
  - Specification defines syntax and semantics of the language
- SUSHI is the go-to method for converting FSH into FHIR
  - reference implementation and de-facto standard FSH "compiler"
  - Usable both "standalone" and together with the IG publisher
  - o also defines an ANTLR4 grammar for parsing the formal language specification







## **BabelFSH**

- Use of FSH for definition of metadata in a language users are familiar with
  - Same language for definition of super-simple CS and especially profiles
- Use of "plugins" for domain-specific concept generation
  - o If possible, general-purpose implementation
  - o If not, that's fine 😁
  - Very simple API for the definition of plugins: only produce a stream of concepts/includes/elements
- Parametrization of the plugins using "command line switches"
- Optionally take in input data from files or other mechanisms
- Write out FHIR JSON to an output folder
- Use of the SUSHI grammar and ANTLR4 parser with obvious restrictions and one extension
  - Generation of FSH items other than CS/VS/CM and RuleSet not intended
  - Comments give the parameters for the invoked plugins, delimited with recognition tokens









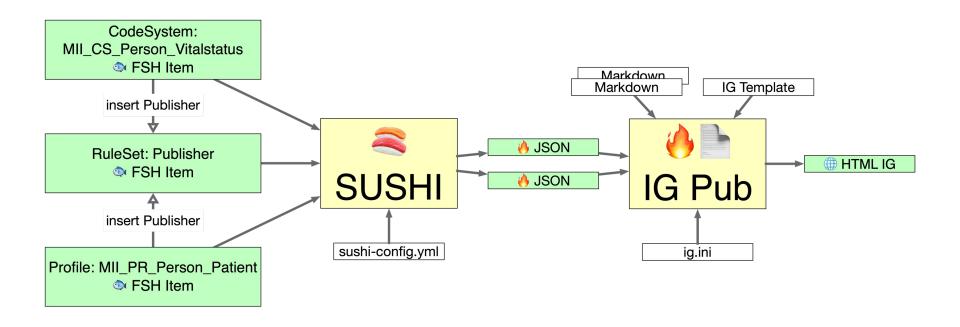








# SUSHI, 🍄 FSH and the 🤚 🖿 IG Publisher











```
RuleSet: alphaid-se-metadata
                                      with machine-readable
* ^url = "http://fhir.de/
                                          comments
CodeSvstem/bfarm/alpha-id"
* ^status = #active
// ...
RuleSet: alphaid-se-babelfsh(version, oid, path)
* ^version = "{version}"
* ^identifier[+].system = "urn:ietf:rfc:3986"
* ^identifier[=].value = "urn:oid:{oid}"
* insert alphaid-se-metadata
/*^babelfsh
CSV
 --path='{path}'
 --headers=["gueltig", "code", "icd_10"]
 --delimiter='|'
 --code-column=code
 --display-column='display'
 --property-mapping=[{"column": "icd_10",
"property": "icd 10 primaer"}]
^babelfsh*/
CodeSystem: AlphaIdSe
Id: alphaid-se-2024
Title: "Alpha-ID-SE"
Description: "The Alpha-ID is a sequential and stable
identification number, which is allocated to each
entry in the alphabetical index. It permits the
encoding of medical and natural language diagnostic
terms."
* insert alphaid-se-babelfsh-2018ff("2024",
"1.2.276.0.76.5.538", "./input-files/Alpha-ID/
alphaidse2024/
icd10gm2024 alphaidse edvtxt 20230929.txt")
```

```
Alpha-ID-SE as
     input file
BabelFSH
 CodeSystemPlugin
       "csv"
```

```
HL7 FHIR
 "resourceType": "CodeSystem",
 "id": "alphaid-se-2024",
                                          R4B or R5
 "url": "http://fhir.de/
CodeSystem/bfarm/alpha-id",
  "identifier": [
      "system": "urn:ietf:rfc:3986",
      "value": "urn:oid:1.2.276.0.76.5.538"
  "version": "2024".
  "name": "AlphaIdSe".
  "title": "Alpha-ID-SE",
  "status": "active",
  "experimental": false.
  "caseSensitive": false,
  "content": "complete",
  "property": [
      "code": "icd 10 primaer",
      "description": "Der ICD-Code aus der Alpha-ID",
      "type": "string"
  "concept": [
      "code": "I1",
      "display": "Lymphatische Infiltration",
      "property": [
          "code": "icd 10 primaer".
          "valueString": "D47.9"
          "code": "inactive".
          "valueBoolean": false
```







# **Development perspective**

- Implemented as of 2024-05-23: only CodeSystem plugins
  - CSV
  - Excel (started development for ATC)
  - ClaML: general using <u>fhir-claml</u> and BfArM-specific flavours
- Planned:
  - OncoTree (API)
  - EDQM Standard Terms (API); including several ValueSets → API for VS/CM generation still NYI
  - ... as required by the MII CDS and other partners
  - CM from OHDSI Athena
- Fancy ideas:
  - SQL using JDBC for "catalogs" in primary systems
  - Provision of FHIR API for nomenclatures (UCUM)
  - Web-based interface and/or language server integration in VS Code for simpler creation and validation of BabelFSH source files









### BabelFSH in use and in the future

- https://gitlab.com/mii-termserv/babelfsh
  - o Also provided: example **.babelfsh.fsh** files
- Generation of FHIR resources used within the MII CDS and other projects
  - o ultimately for upload to the SU-TermServ server; c.f. our (WIP) package registry
- Tool can help with difficult-to-license terminologies: share BabelFSH files instead of FHIR resources
- Representation of terminological artefacts subject to variation
  - Naming Conventions
  - Property definitions
  - Establishment of BabelFSH and sharing of definition files as a commonly-used tool
     can help alignment to conventions by making conversion process more transparent



# MII Service Unit Terminological Services (SU-TermServ)

https://mii-termserv.de team@mail.mii-termserv.de

A partnership between the University of Luebeck, the University of Cologne, and the Hannover Medical School







