



Medizinische Hochschule Hannover



SU-TermServ

BabelFSH

Interoperabilitätsforum, June 2024, Berlin @ DIN Joshua Wiedekopf IT Center for Clinical Research & Institute of Medical Informatics, University of Lübeck

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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



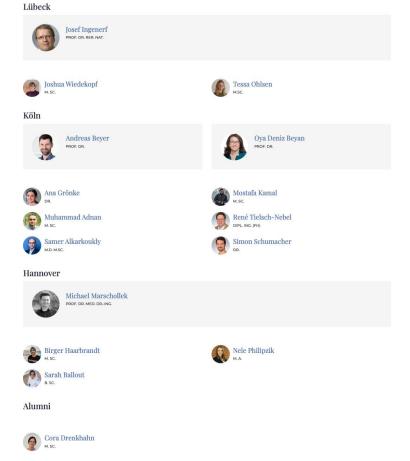


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Our team







The Problem

- Using FHIR Terminology server operations requires access to FHIR CodeSystem resources
 - \circ ~ 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
 - 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)
 - In-development specification for a domain-specific language
 - 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"
 - \circ ~ Usable both "standalone" and together with the IG publisher
 - also defines an ANTLR4 grammar for parsing the formal language specification







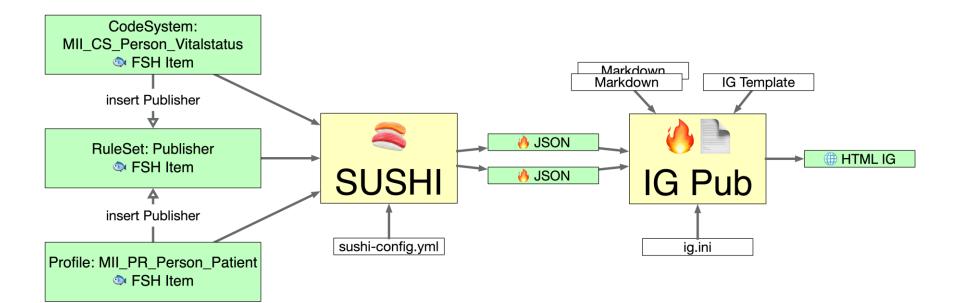
BabelFSH

- 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
 - If possible, general-purpose implementation
 - 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

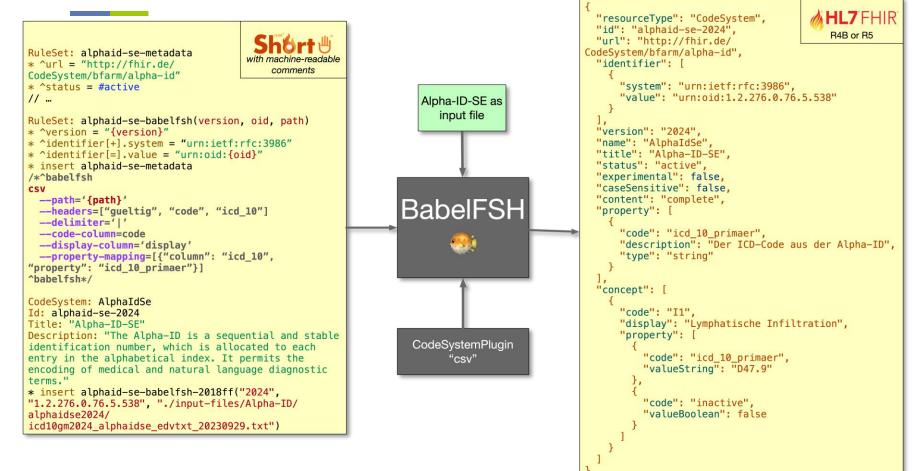






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Development perspective

- Implemented as of 2024-05-23: only CodeSystem plugins
 - o 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

- <u>https://gitlab.com/mii-termserv/babelfsh</u>
 - Also provided: example **.babelfsh.fsh** files
- Generation of FHIR resources used within the MII CDS and other projects
 - 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



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