FAIR Genomes semantic metadata schema

The FAIR Genomes semantic metadata schema to power reuse of NGS data in research and healthcare. Version 0.3-SNAPSHOT, 2021-02-25. This model consists of 9 modules that contain 110 metadata elements in total.

| Name | Ontology | Nr. of elements |
|--------------------------|---------------------|-----------------|
| Study | NCIT:C63536 | 9 |
| Personal | NCIT:C90492 | 12 |
| Leaflet and consent form | NCIT:C16468 | 8 |
| Individual consent | NCIT:C16735 | 13 |
| Clinical | NCIT:C25398 | 20 |
| Material | NCIT:C43376 | 16 |
| Sample preparation | OBI:0001902 | 9 |
| Sequencing | EDAM:topic_3168 | 12 |
| Analysis | EDAM:operation_2945 | 11 |

Table 1: FAIR Genomes v0.3-SNAPSHOT overview of all modules.

| Name | Ontology | Values |
|------------------------|-----------------|---------------------------------------|
| Identifier | OMIABIS:0000006 | UniqueID |
| Name | OMIABIS:0000037 | String |
| Description | OMIABIS:0000036 | Text |
| Inclusion criteria | OBI:0500027 | InclusionCriteria lookup (11 choices) |
| Principal investigator | OMIABIS:0000100 | String |
| Contact information | OMIABIS:0000035 | String |
| Study design | OBI:0500000 | Text |
| Start date | NCIT:C69208 | Date |
| Completion date | NCIT:C142702 | Date |

Table 2: Module: Study. A detailed examination, analysis, or critical inspection of a subject designed to discover facts about it. Ontology: NCIT:C63536.

| Name | Ontology | Values |
|-------------------------------|-----------------|------------------------------------|
| Personal identifier | NCIT:C164337 | UniqueID |
| Phenotypic sex | PATO:0001894 | PhenotypicSex lookup (4 choices) |
| Genotypic sex | PATO:0020000 | GenotypicSex lookup (11 choices) |
| Country of residence | NCIT:C171105 | Countries lookup (249 choices) |
| Ancestry | HANCESTRO:0004 | Ancestry lookup (305 choices) |
| Country of birth | GENEPIO:0001094 | Countries lookup (249 choices) |
| Year of birth | NCIT:C83164 | Integer |
| Inclusion status | NCIT:C166244 | InclusionStatus lookup (4 choices) |
| Age at death | NCIT:C135383 | Integer |
| Primary affiliated institute | SIO:000688 | Institutes lookup (218 choices) |
| Resources in other institutes | SIO:000688 | Institutes lookup (218 choices) |
| Participates in study | RO:0000056 | Reference to Study |

Table 3: Module: Personal. Data, facts or figures about an individual; the set of relevant items would depend on the use case. Ontology: NCIT:C90492.

| Name | Ontology | Values |
|----------------------------|-----------------|---------------------------------|
| Leaflet title | DC:title | String |
| Leaflet date | DC:date | Date |
| Leaflet version | DC:hasVersion | String |
| Consent form identifier | DC:identifier | UniqueID |
| Consent form accepted date | DC:dateAccepted | Date |
| Consent form valid until | DC:valid | Date |
| Consent form creator | SIO:000688 | Institutes lookup (218 choices) |
| Consent form version | DC:hasVersion | String |

Table 4: Module: Leaflet and consent form. A document explaining all the relevant information to assist an individual in understanding the expectations and risks in making a decision about a procedure. This document is presented to and signed by the individual or guardian. Ontology: NCIT:C16468.

| Name | Ontology | Values |
|-------------------------------|-------------|---------------------------------------|
| Individual consent identifier | ICO:0000044 | UniqueID |
| Person consenting | IAO:0000136 | Reference to Personal |
| Consent form used | IAO:0000136 | Reference to Leaflet and consent form |
| Collected by | NCIT:C45262 | Institutes lookup (218 choices) |
| Signing date | ICO:0000036 | Date |
| Valid from | DC:valid | Date |
| Valid until | DC:valid | Date |
| Represented by | NCIT:C51828 | Represented By lookup (2 choices) |
| Data use permissions | DUO:0000001 | DataUsePermissions lookup (5 choices) |
| Data use modifiers | DUO:0000017 | DataUseModifiers lookup (18 choices) |
| Modifiers specification | SIO:000090 | Text |
| Allow linkage | NCIT:C15424 | Boolean |
| Allow recontacting | NCIT:C25737 | Recontacting lookup (3 choices) |

Table 5: Module: Individual consent. Consent by a patient to a surgical or medical procedure or participation in a clinical study after achieving an understanding of the relevant medical facts and the risks involved. Ontology: NCIT:C16735.

| Name | Ontology | Values |
|----------------------------|--------------------|--------------------------------------|
| Clinical identifier | NCIT:C87853 | UniqueID |
| Belongs to person | IAO:0000136 | Reference to Personal |
| Phenotype | NCIT:C16977 | Phenotypes lookup (15802 choices) |
| Unobserved phenotype | HL7:C0442737 | Phenotypes lookup (15802 choices) |
| Phenotypic data available | NCIT:C15783 | DCMITypes lookup (6 choices) |
| Clinical diagnosis | NCIT:C15607 | Diseases lookup (9700 choices) |
| Molecular diagnosis gene | NCIT:C20826 | Genes lookup (19202 choices) |
| Molecular diagnosis other | NCIT:C20826 | Text |
| Age at diagnosis | SNOMEDCT:423493009 | Integer |
| Age at last screening | NCIT:C81258 | Integer |
| Medication | NCIT:C459 | Drugs lookup (5632 choices) |
| Drug regimen | NCIT:C142516 | Text |
| Family members affected | HP:0032320 | FamilyMembers lookup (41 choices) |
| Family members sequenced | NCIT:C71384 | FamilyMembers lookup (41 choices) |
| Consanguinity | GSSO:007578 | String |
| Medical history | NCIT:C18772 | MedicalHistory lookup (1154 choices) |
| Age of onset | Orphanet:C023 | Integer |
| First contact | LOINC:MTHU048806 | Date |
| Functioning | NCIT:C21007 | Text |
| Material used in diagnosis | SIO:000641 | String |

Table 6: Module: Clinical. Relating to the examination and treatment of patients dependent on direct observation. Ontology: NCIT:C25398.

| Name | Ontology | Values |
|--|----------------|--|
| Material identifier | NCIT:C93400 | UniqueID |
| Collected from person | SIO:000244 | Reference to Personal |
| Belongs to diagnosis | SIO:000068 | Reference to Clinical |
| Sampling timestamp | EFO:0000689 | DateTime |
| Registration timestamp | NCIT:C25646 | DateTime |
| Sampling protocol | EFO:0005518 | Text |
| Sampling protocol deviation | NCIT:C50996 | String |
| Reason for sampling protocol deviation | NCIT:C93529 | String |
| Biospecimen type | NCIT:C70713 | MaterialTypes lookup (13 choices) |
| Anatomical source | UBERON:0001062 | AnatomicalSources lookup (13827 choices) |
| Pathological state | NCIT:C25687 | PathologicalState lookup (4 choices) |
| Storage conditions | NCIT:C96145 | StorageConditions lookup (26 choices) |
| Expiration date | NCIT:C164516 | Date |
| Percentage tumor cells | NCIT:C127771 | Decimal |
| Physical location | GAZ:00000448 | String |
| Derived from | NCIT:C28355 | String |

Table 7: Module: Material. Natural substances derived from living organisms such as cells, tissues, proteins, and DNA. Ontology: NCIT:C43376.

| Name | Ontology | Values |
|-------------------------|-----------------|------------------------------|
| Sampleprep identifier | NCIT:C132299 | UniqueID |
| Belongs to material | NCIT:C25683 | Reference to Material |
| Input amount | AFRL:0000010 | Integer |
| Library preparation kit | GENEPIO:0000085 | NGSKits lookup (615 choices) |
| PCR free | NCIT:C17003 | Boolean |
| Target enrichment kit | GENEPIO:0000081 | NGSKits lookup (615 choices) |
| UMIs present | EFO:0010199 | Boolean |
| Intended insert size | FG:0000001 | Integer |
| Intended read length | NCIT:C153362 | Integer |

Table 8: Module: Sample preparation. A sample preparation for assay that preparation of nucleic acids for a sequencing assay. Ontology: OBI:0001902.

| Name | Ontology | Values |
|-----------------------------|-----------------|--|
| Sequencing identifier | NCIT:C171337 | UniqueID |
| Belongs to sample | NCIT:C25683 | Reference to Sample preparation |
| Sequencing date | GENEPIO:0000069 | Date |
| Sequencing platform | GENEPIO:0000071 | SequencingPlatform lookup (7 choices) |
| Sequencing instrument model | GENEPIO:0001921 | SequencingInstrumentModels lookup (39 choices) |
| Sequencing type | NCIT:C25284 | Sequencing Types lookup (35 choices) |
| Average read depth | NCIT:C155320 | Integer |
| Observed read length | NCIT:C153362 | Integer |
| Observed insert size | FG:0000002 | Integer |
| Percentage Q30 | GENEPIO:0000089 | Decimal |
| Percentage TR20 | FG:0000003 | Decimal |
| Other quality metrics | EDAM:data_3914 | Text |

Table 9: Module: Sequencing. The determination of complete (typically nucleotide) sequences, including those of genomes (full genome sequencing, de novo sequencing and resequencing), amplicons and transcriptomes. Ontology: EDAM:topic_3168.

| Name | Ontology | Values |
|---|------------------|--------------------------------------|
| Analysis identifier | AFR:0001979 | UniqueID |
| Belongs to sequencing | NCIT:C25683 | Reference to Sequencing |
| Physical data location | GAZ:00000448 | String |
| Abstract data location | NCIT:C142494 | String |
| Data formats stored | EDAM:format_1915 | DataFormats lookup (582 choices) |
| Algorithms used | NCIT:C16275 | Text |
| Reference genome used | EDAM:data_2340 | GenomeAccessions lookup (29 choices) |
| Bioinformatic protocol used | EDAM:data_2531 | Text |
| Bioinformatic protocol deviation | NCIT:C50996 | String |
| Reason for bioinformatic protocol deviation | NCIT:C93529 | String |
| WGS guideline followed | NCIT:C17564 | String |

Table 10: Module: Analysis. Apply analytical methods to existing data of a specific type. Ontology: EDAM:operation_2945.