

YAML	Turtle
<pre> # Study identifier, specifically created for the website. # The identifier is the last part of the URL directing to the particular study study_id: 'Covid19Icarius' # Analytics Use Case of the Study, choose 0, 1, 2 or 3: # 0: Characterization # 1: Population-Level Estimation # 2: Patient-Level Prediction # 3: Characterization and Population-Level Estimation study_usecase: [1] # Conditions studied; if multiple conditions are being studied, # duplicate all keys under "conditions" conditions: - concept_name: "Disease caused by 2019-nCoV" concept_id: "37311061" code: concept_code: "840539006" vocabulary_id: "SNOMED CT" concept_code_url: "http://snomed.info/id/840539006" # Drug studied; if multiple subjects are being studied, # duplicate all keys under "study_subject" study_subject: - concept_name: "quinaprilat" concept_id: "45775375" code: concept_code: "1546359" vocabulary_id: "RxNorm" concept_code_url: "http://purl.bioontology.org/ontology/RXNORM/1546359" </pre>	<pre> @prefix ohdsi: <http://data.ohdsi.org/> . @prefix concept: <http://data.ohdsi.org/concept/> . @prefix study: <https://covid19.ohdsi.app/study/> . @prefix schema: <http://schema.org/> . study:Covid19Icarius a schema:MedicalObservationalStudy ; ohdsi:analyticsUseCase ohdsi:PopulationLevelEstimation ; schema:healthCondition concept:37311061 ; schema:studySubject concept:45775375 . concept:37311061 a schema:MedicalCondition ; schema:name "Disease caused by 2019-nCoV" ; schema:identifier "37311061" ; schema:code [a schema:MedicalCode ; schema:codeValue "840539006" ; schema:codingSystem "SNOMED CT" ; schema:sameAs "http://snomed.info/id/840539006"] . concept:45775375 a schema:Drug ; schema:name "quinaprilat" ; schema:identifier "1777087" ; schema:code [a schema:MedicalCode ; schema:codeValue "1546359" ; schema:codingSystem "RxNorm" ; schema:sameAs "http://purl.bioontology.org/ontology/RXNORM/"1546359"] . </pre>