

# Ontology for Biomedical Investigations

The OBI Consortium\*, <http://purl.obolibrary.org/obo/obi>

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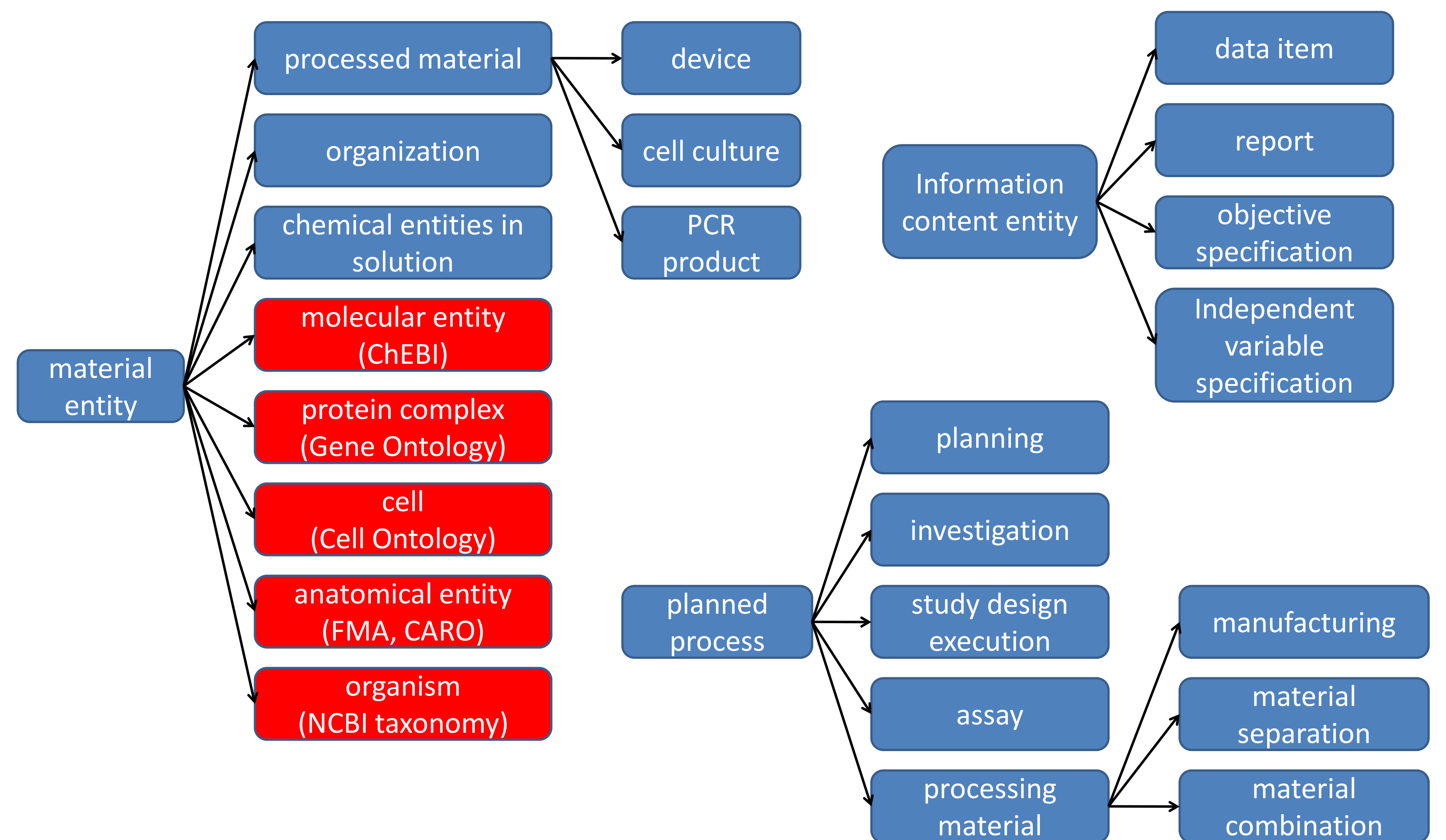
The goal of OBI is to enable a formal representation of biomedical investigations that captures the experimental evidence on which their findings are based. The scope of OBI includes: materials made in and produced for investigations, research objectives, experimental protocols, roles of people in investigations and processing and publication of data gathered in investigations. Use of OBI will allow comparison of experimental data from the wide array of scientific disciplines represented by domain experts in the OBI consortium. OBI follows the principles laid out by the OBO foundry, and integrates tightly with other foundry candidate ontologies, such as GO ([www.geneontology.org](http://www.geneontology.org)) and ChEBI ([www.ebi.ac.uk/chebi/](http://www.ebi.ac.uk/chebi/)) whose terms are used to describe biological reality. The use of OBI by the scientific community to represent or annotate their investigations within electronic data resources will facilitate interdisciplinary data synthesis, enable access to their data on the semantic web and improve third-party understanding of information related to life-science and clinical investigations.

## Partial high-level is-a hierarchy of OBI classes.

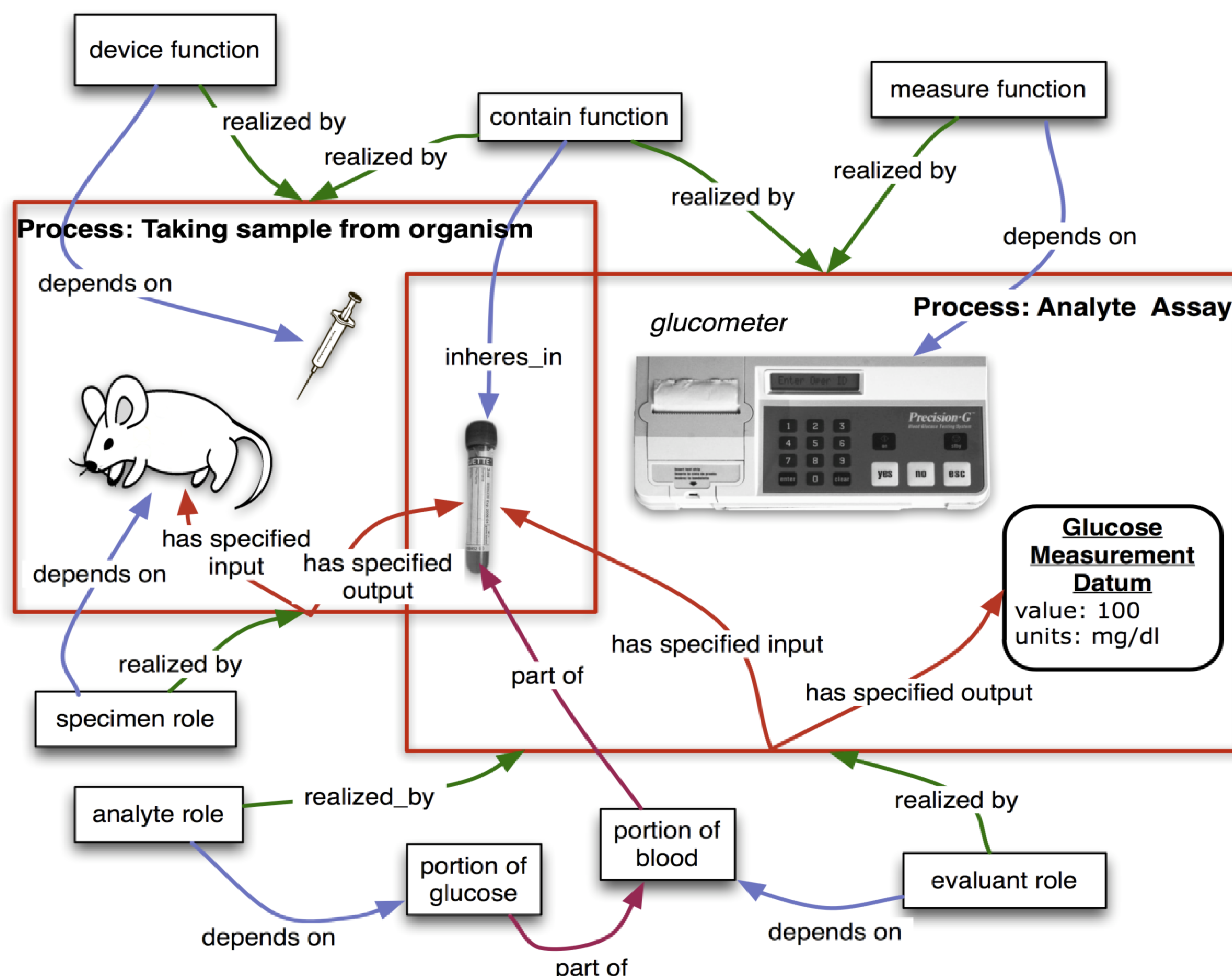
Under the Basic Formal Ontology (BFO) ([www.ifomis.org/bfo](http://www.ifomis.org/bfo)) *material entity*, several classes are imported from outside ontologies (indicated in red). The *processed material* class represents entities that were generated in a planned process, meaning that they were intentionally created.

*planned process* are initiated by an agent in order to achieve a specific objective. This includes all research activities, such as an *investigation* encompassing a planning, study design execution and documenting phase, or an individual *assay*.

*Information content entities* are about the material entities and processes in an investigation in the form of its design, outcomes, and reporting.



## A detailed example how to represent a blood glucose concentration measurement



**Measuring the glucose concentration in blood.** The large boxes represent processes and contain their participants. The *taking sample from organism* process takes place first. In this process, a syringe is used as a device to draw blood from the mouse which bears the specimen role. At the end of this process, a tube contains the blood specimen. In a second step, that blood will be used as the evaluant in an analyte assay in which the concentration of glucose in the blood is measured. A glucometer device is used to make this measurement. The analyte role inheres in the glucose molecules scattered throughout the blood specimen. The objective of this planned process is to analyze the analyte (glucose) concentration. This model of a specific use case is being used as the basis for generation of templates allowing us to automate subsequent additions of similar processes

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