CATMAID data model

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CATMAID, the Collaborative Annotation Toolkit for Massive Amounts of Image Data, is a web-based platform for the annotation of very large data sets, such as those produced by serial section transmission electron microscopy. The aim of the CATMAID data model is to describe annotations of image data in terms of Subject-Relation-Object triples suitable for mapping to semantic web technologies and incorporation into controlled vocabularies such as those defined by the Open Biomedical Ontologies group. Our other aim is to be able to store these logical relationships efficiently in PostgreSQL. The following sections describe our current model and the plans for possible changes.

Current Entities and Relationships

The highest level distinction between the entities we are modelling is between **location**s and **class_instance**s. The distinction between these is that a **location** is a physically located point in space, whereas a **class_instance** is an abstract concept, only linked to a point in space via relationships with other entities.

There are two subclasses of **location**: **treenode**, which represents a node as part of a tree structure, and **connector**, which is used to link locations in a many-to-many fashion, typically used for annotating synapses.

Examples of **class_instance**s include neurons, groups of neurons, groups of groups, skeletons, synapses and preand postsynaptic terminals.

The current set of relationships which are allowed between these entities are:

subject	relation	object
treenode	labeled_as	class_instance
treenode	element_of	class_instance
treenode	model_of	class_instance
connector	model_of	class_instance
class_instance	model_of	class_instance
class_instance	part_of	class_instance
class_instance	is_a	class_instance
class_instance	postsynaptic_to	class_instance
class_instance	presynaptic_to	class_instance

These relationships are all asymmetric. Every **class_instance** is an instance of a particular **class** - at the moment, the defined **class**es are **skeleton**, **neuron**, **group**, **label**, **root**, **soma**, **synapse**, **presynaptic terminal**, **postsynaptic terminal**.

Examples of the uses of each of these relationships are given below:

labeled_as

During annotation, it is frequently useful to annotate a point in the tree with a text-based tag, which might be "TODO", "possible synapse here", etc. This relationship links a treenode to a label of this kind.

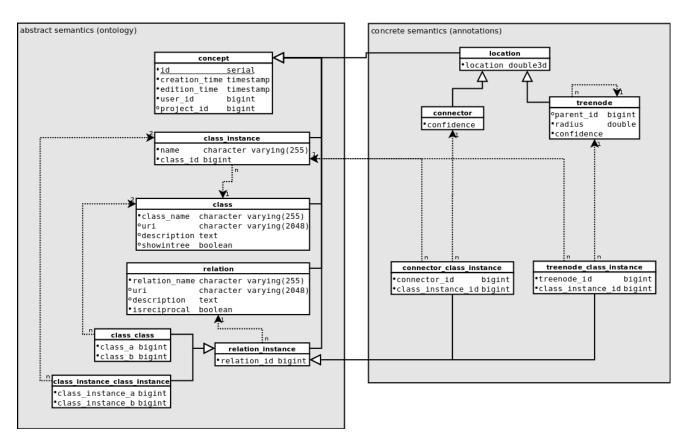


Figure 1: Database schema

postsynaptic_to, presynaptic_to

We are representing the polyadic synapses found in our data with a connector placed over the synaptic cleft. A connector is a model_of a synapse. Treenodes that are pre-synaptic and post-synaptic partners are in a model_of relationship with presynapticterminal and postsynapticterminal. The postynaptic_to and presynaptic_to relationships are then used to link the synaptic terminals to their synapse. These relationships could describe many-to-many relationships across synapses, although there is typically only a single presynaptic terminal in the relationship.

element_of

The element_of relationship is currently used to indicate that a treenode is part of a particular skeleton. This is distinct from the standard part_of relationship to indicate that the subject is a treenode rather than a class_instance.

model of

The concrete annotations used in CATMAID should be tissue-type agnostic; in other words, they should not be specific to annotating neuronal structures. The model_of relationship links a basic annotation to the particular biological structure that it is representing. One example use of this is to create a link between a skeleton and the neuron it represents the midline of. The same neuron can be modelled by skeletons traced by different users. As another example, a connector is a model of a synapse.

part_of

This relationship indicates that one class_instance is part of another; for example, a soma is part_of a neuron or a presynaptic terminal is part_of a skeleton. This should be the same as the part_of relationship used in the OBO foundry¹ ontologies.

is_a

This relationship indicates that one class_instance is a subtype of another; for example, a neuron is_a cell. This should be the same as the part_of relationship used in the OBO foundry.

Representation in PostgreSQL

The representation of these entities and relationships in PostgreSQL makes use of the database's inheritance feature. If one table inherits from another, then the descendant has all the columns of the former, and various queries enacted on the parent table will also include rows from all descendant tables. At the top level of this hierarchy in CATMAID is the **concept** table, which provides columns for the *user_id*, *project_id*, *creation_time*, *modification_time* and a unique *id*. All the other tables described below similarly inherit from **concept**.

Each possible relation is described at the top level by a row in the **relation** table. For example (with some columns elided):

¹http://www.obofoundry.org/

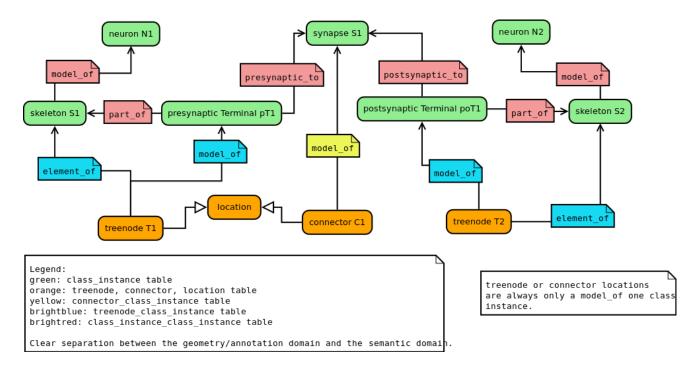


Figure 2: Treenode-connector relationships with the semantic annotation

relation_name	isreciprocal
labeled_as	f
postsynaptic_to	f
presynaptic_to	f
element_of	f
model_of	f
part_of	f
is_a	f

An instance of one of these relations is represented by a row in a table which inherits from the **relation_instance** table. The relation_instance table adds a *relation_id* column to specify which relation this is an instance of. The three tables that inherit from relation_instance are **class_class_instance_class_instance** and **treenode_class_instance**. The **class_class** table describes the possible classes that can be linked by that particular relation. Which table is used for the concrete relationships depends on whether the subject of the relationship is a **class** or a **treenode**. The treenode_class_instance table has rows that link a treenode to a class instance, whereas the **class_instance_class_instance** table links two **class_instance_class_instance**.

Examples of classes from the **class** table are:

showintree
t
t
t
f
f
t
t
t
t

The *showintree* column indicates whether this class should be shown in the tree display of the annotations in CATMAID's interface. Particular instances of each of these classes are represented by rows in the **class_instance** table, which provide a name and id for each class instance.

Instances of the Subject-Relation-Object relationship between instances of two classes are represented by rows in the class_instance_class_instance table:

relation_id	class_instance_a	class_instance_b
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Relations can also apply between a **class_instance** and a **treenode**.

The **treenode** and **connector** tables inherit from the **location** table which expresses the location in space.

The distinction between a treenode and a location is that a treenode additionally has these features:

- parent_id: A parent treenode, which may be null in the case of the root node in a skeleton.
- radius: a radius of the skeleton at that point.
- *confidence*: on a scale of 1 to 5, the confidence that the annotator has that the link to the parent node is correct. (5 is the most confident.)

These relationships between treenodes and class_instances are stored as rows in the **treenode_class_instance** table:

relation_id	treenode_id	class_instance_id
11 [element_of]	240	561
11 [element_of]	241	561
11 [element_of]	242	561
	•••	•••
23 [model_of]	268	2500
35 [labeled_as]	268	1562

('labeled_as' gives a label to a particular treenode, 'model_of' to mark the treenode as presynaptic terminal and 'element_of' links a treenode to the skeleton it is a part of.)

Similarily, relations can apply between a **connector** and a **class_instance**. Connector are points in space used to model the location of a synapse. Connector additionally has these features:

• *confidence*: on a scale of 1 to 5, the confidence that the annotator has that this connector actually is a synapse and links correctly. (5 is the most confident.)

These relationships between connectors and class_instances are stored as rows in the **connector_class_instance** table:

relation_id	location_id	class_instance_id
23 [model_of]	300	1800
23 [model_of]	303	4800
35 [labeled_as]	268	15623

File Format

For this web-based, database-backed system a concrete file format for import and export is not a critical as provision of an API for interrogating the database. However, we will intially use TrakEM2's XML-based format in order to import data.