# GreenPhyl RDF data model

Prefixes: @base <a href="http://www.southgreen.fr/agrold/">http://www.southgreen.fr/agrold/>... rdf:<http://www.w3.org/1999/02/22-rdf-syntax-ns#>. @prefix @prefix rdfs:<http://www.w3.org/2000/01/rdf-schema#> . @prefix xsd:<http://www.w3.org/2001/XMLSchema#> . @prefix agrold vocabulary:<a href="http://www.southgreen.fr/agrold/vocabulary/">http://www.southgreen.fr/agrold/vocabulary/</a>. obo:<http://purl.obolibrary.org/obo/>. @prefix uniprot:<http://purl.uniprot.org/uniprot/>. @prefix @prefix tigrlocus:<a href="http://www.southgreen.fr/agrold/tigr.locus/">http://www.southgreen.fr/agrold/tigr.locus/</a>>. @prefix tairlocus:<a href="http://identifiers.org/tair.locus/">http://identifiers.org/tair.locus/</a>>. pubmed:<http://identifiers.org/pubmed/>. @prefix interpro:<http://identifiers.org/interpro/>. @prefix @prefix ncbi taxon:<a href="http://purl.obolibrary.org/obo/NCBITaxon">http://purl.obolibrary.org/obo/NCBITaxon</a>>. greenphyl family:<a href="mailto:family/">http://www.southgreen.fr/agrold/greenphyl.family/">... @prefix @prefix greenphy\_annotation:<http://www.southgreen.fr/agrold/greenphyl.annotation/>. Predicates to be used:

rdf:type rdf:subject rdf:predicate rdf:object rdfs:subClassOf rdfs:label agrold vocab:description agrold vocab:taxon agrold\_vocab:contains agrold\_vocab:is\_orthologous to agrold vocab:is paralogous to agrold vocab:is member of agrold\_vocab:xRef agrold vocab:has annotation agrold vocab:has score agrold\_vocab:assigned\_by agrold vocab:has uniprot accession agrold vocab:number of sequences agrold vocab:has thresold agrold\_vocab:curation\_status agrold\_vocab:has\_uniprot\_accession agrold vocab:has go identifier obo:BFO 0000056 # participates in - Biological process

```
obo: BFO_0000085 # has_function - Molecular Function obo:BFO_0000082 # located_in - Cellular Component
```

### **Protein information:**

ID: Uniprot < <a href="http://purl.uniprot.org/uniprot/">http://purl.uniprot.org/uniprot/</a>>

Name: Literal Description: literal

Protien Ontology term: SO\_0000104 <obo:SO\_0000104> Taxon: NCBI <a href="http://purl.obolibrary.org/obo/NCBITaxon">http://purl.obolibrary.org/obo/NCBITaxon</a> >

Gene: MSU/TIGR Locus ID <a href="http://www.southgreen.fr/agrold/tigr.locus/">http://www.southgreen.fr/agrold/tigr.locus/</a>> (O.sativa);

TAIR ID < <a href="http://identifiers.org/tair.locus/">http://identifiers.org/tair.locus/</a>> (A.thaliana)

IPR: InterPro < <a href="http://identifiers.org/interpro/">http://identifiers.org/interpro/</a>
GO: ID <a href="http://purl.obolibrary.org/obo/">http://purl.obolibrary.org/obo/</a>
Homology info: Orthology: Uniport ID (URI)

Paralogy: Uniport ID (URI)

Score: literal

Annotation: Literal

DB xref: URI or literal, depends on the kind of xref e.g IDs Reference: Pubmed ID <a href="http://identifiers.org/pubmed/">http://identifiers.org/pubmed/</a>

#### **Protein RDF Model:**

E.g. Arabidopsis

**Note:** For the corresponding ortholog protein a prefix need to be based on the source organism.

#### tair:ATXXXX

```
rdf:type
             agro vocab:Protein;
rdfs:subClassOf
                   obo:SO_0000104
rdfs:label
             "name":
agrold vocab:description
                         "description";
agrold_vocab:taxon ncbi_taxon:XXXX;
obo:BFO 0000056 # participates in
                                      obo:GO ID;
obo: BFO 0000085 # has function obo: GO ID;
obo:BFO 0000082 # located in
                                obo:GO ID;
agrold vocab:contains
                                interpro:XXXXX;
                                greenphyl_family:XXXXX; # family ID
agrold vocab:is member of
agrold vocab:is orthologous to
                                uri:XXXXXX;
agrold vocab:has annotation
                                greenphy annotation:ID1XXXXX ID2XXXX;
agrold_vocab:is_paralogous_to
                                uri:XXXXXX ;
                                greenphy_annotation:ID1XXXXX_ID3XXXX;
agrold_vocab:has_annotation
agrold vocab:has uniprot accession
                                             uniprot:XXXXX;
agrold_vocab:xRef pubmed:XXXXXX;
```

```
# Reification
greenphy_annotation:ID1XXXXX_ID2XXXX
rdf:type rdf:Statement;
rdf:subject tair:XXXXX ; # ID1
rdf:predicate agro_vocab:is_orthologous_to;
rdf:object tair:XXXXX ; # ID2
agrold_vocab:has_score "score"; # Literal
agrold_vocab:assigned_by "GreenPhyl".
```

## Family information:

ID: Internal < http://www.southgreen.fr/agrold/greenphyl.family/>

cluster (family): OBI\_0000251

Name: Literal Level: Literal

Number\_of\_sequences: Literal

Evidence:Literal

GO: ID < <a href="http://purl.obolibrary.org/obo/">http://purl.obolibrary.org/obo/</a>>

DB xref: URI or literal, depends on the kind of xref e.g IDs

## Family RDF model:

```
greenphyl_family:GXXXX
      rdf:type
                    agro_vocab:Protein_Family;
      rdf:subClassOf
                          obo:OBI_0000251;
                    "name";
      rdfs:label
      agrold vocab:description
                                 "description";
      agrold vocab:has go identifier
                                        obo:GO ID;
      agrold_vocab:evidence
                                        "Evidence";
                                               "#"^^xsd:integer;
      agrold_vocab:number_of_sequences
      agrold_vocab:has_thresold "#"^^xsd:integer; # e.g. 1, 2,3
      agrold_vocab:curation_status
                                        "Status"; # Annotation in progress
      agrold_vocab:xRef pubmed:XXXXXX .
```