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# **pycropml Documentation**

***Release 1.1.0***

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# CHAPTER 1

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## PyCrop2ML documentation

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### 1.1 Module description

#### 1.1.1 What is PyCrop2ML?

**PyCrop2ML** is a free, open-source library for defining and exchanging CropML models. It is used to generate components of modeling and simulation platforms from the CropML specification and allow component exchange between different platform.

It allows to parse the models described in CropML format and automatically generate the equivalent executable Python, java, C#, C++ components and packages usable from existing crop simulation platform.

#### 1.1.2 What is Crop2ML ?

**CropML** is a XML-(JSON-)based language used to represent different biological processes involved in the crop models.

CropML project aims to provide common framework for defining and exchanging descriptions of crop growth models between crop simulation frameworks.

### Objectives

Our main objectives are:

- define a **declarative language** to describe either an atomic model or a composition of models
- add semantic dimension to CropML language by annotation of the models to allow the composition of components of different platforms by using the standards of the semantic web
- develop a library to allow the transformation and the exchange of CropML model between different Crop modelling and simulation platform
- provide a **web repository** enabling registration, search and discovery of CropML Models

- facilitate Agricultural Model Exchange Initiative

## Context

Nowadays, we observe the emergence of plant growth models which are built in different platforms. Although standard platform development initiatives are emerged, there is a lack of transparency, reusability, and exchange code between platforms due to the high diversity of modeling languages leading to a lack of benchmarking between the different platforms.

This project aims to gather developers and plant growth modelers to define a standard framework based on the development of declarative language and libraries to improve exchange model components between platforms.

## Motivation

### Our motivation is to:

- Strengthen the synergy between crop modelers, users and scientific researchers
- Facilitate model intercomparison (at the process level) and model improvement through the exchange of model components (algorithms) and code reuse between platforms/models.
- Bridge the gap between ecophysicists who develop models at the process level with crop modelers and model users and facilitate the integration in crop models of new knowledge in plant science (i.e. we are seeking the exchange of knowledge rather than black box models).
- Increase capabilities and responsiveness to stakeholder' needs.
- Propose a solution to the AgMIP community for NexGen crop modeling tools.

## Vision

- Facilitate the development of complex models
- Use modular modelling to share knowledge and rapidly develop operational tools.
- Reuse model parts to leverage the expertise of third parties;
- Renovate legacy code.
- Realize the benefit of sharing and complementing different expertise.
- Promote model sharing and reuse

### 1.1.3 Crop2ML Description

In Crop2ML, a model is either a model unit or a composition of models. ModelUnit represents an atomic unit of a crop model. A model composition is a model resulting from the composition of two or more atomic model or composite models.

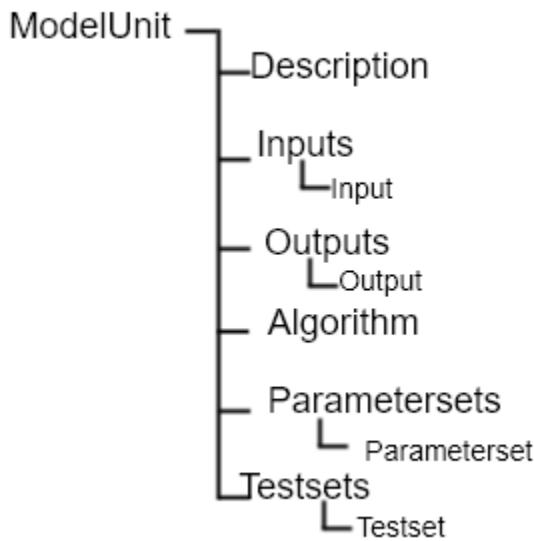
These models have a specific formal definition in Crop2ML.

#### Formal definition of a Model Unit in Crop2ML

The structure of a Model Unit in Crop2ML MUST be conform to a specific Document Type Definition named [ModelUnit.dtd](#).

So a Model Unit Crop2ML document is a XML document well-formed and also obeys the rules given in the ModelUnit structure.

This structure MAY be described by the below tree:



For more details, consult [Crop2ML model unit specification](#).

### Formal definition of a Composite Model in Crop2ML

A Composite Model Crop2ML is an assembly of processes which are described by a set of model units or a composition of models. Given a composite model is a model, this one has also inputs, outputs and internal state which describe the orchestration of different independent models composed.

The structure of a Composite Model in Crop2ML MUST conform to a specific Document Type Definition named [ModelComposition.dtd](#).

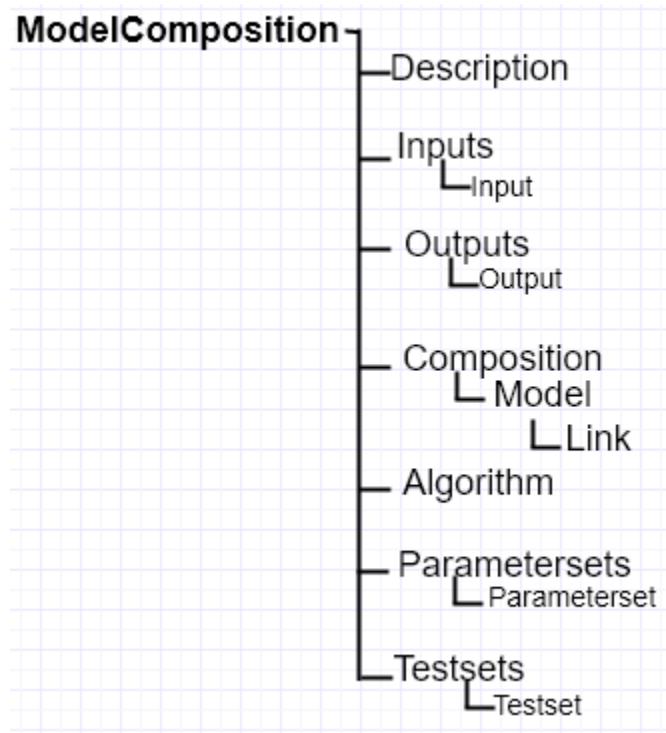
The composition is represented as a directed port graph of models:

Vertices are the different models that form the composition.

Ports are the inputs and outputs of each model.

Edges are directed and connect one output port to an input port of another model.

It contains in addition to all Elements of a model unit a Composition Element for the composition of models. This structure MAY be described by the below tree:



| For more details, consult [Crop2ML model composite specification](#) .

#### 1.1.4 CyML Language Specification

This document specifies the CyML language, an extended Cython subset supported.

#### 1.1.5 Cython file types

- The implementation files, carrying a `.pyx` suffix.

#### Basic Types

The following basic types are supported

- `bool`
- `int`
- `float`
- `double`
- `string`

#### Complex Types

The following complex types are supported

- array
- list
- datetime

## Conditional Statements

- IF
- IF/Else
- IF/ElseIf/Else

The ELIF and ELSE clauses are optional. An IF statement can appear anywhere that a normal statement or declaration can appear

## Integer For Loops

CyML recognises the usual Python for-in-range integer loop pattern:

```
for i in range(n):  
    ...
```

Like other Python looping statements, break and continue may be used in the body, without the loop that have an else clause.

## While Loop

- Like Python While loop

## Function

- Parameters with declaration
- Default value is possible

## Return Statement

- A function needs to return the same data type.

The following code is valid:

```
def fibonacci(int n):  
    if n <= 2:  
        return 1  
    else:  
        return fibonacci(n-1)+fibonacci(n-2)
```

## Call functions

- Call to CyML functions are supported if the function
- is accessible on the module
- is accessible from import statement

## Operators

### Assignment

Assign	<code>b = a</code>
--------	--------------------

### Unary operators

UAdd	<code>+a</code>
USub	<code>-a</code>

### Binary operators

Add	<code>a + b</code>
Sub	<code>a - b</code>
Mult	<code>a * b</code>
Div	<code>a / b</code>
FloorDiv	<code>a // b</code>
Pow	<code>a ** b</code>
Mod	<code>a % b</code>
LShift	<code>a &lt;&lt; b</code>
RShift	<code>a &gt;&gt; b</code>
BitOr	<code>a   b</code>
BitXor	<code>a ^ b</code>
BitAnd	<code>a &amp; b</code>

### Augmented assign statements

AugAdd	<code>a += b</code>
AugSub	<code>a -= b</code>
AugMult	<code>a *= b</code>
AugDiv	<code>a /= b</code>

### Comparison Operators

Eq	<code>a == b</code>
NotEq	<code>a != b</code>
Lt	<code>a &lt; b</code>
LtE	<code>a &lt;= b</code>
Gt	<code>a &gt; b</code>
GtE	<code>a &gt;= b</code>

### Bool Operators

<code>&amp;&amp;</code>	a and b
<code>  </code>	a or b

## Array creation routines

<code>“ “</code>	Return a new array of given shape and type, without initializing entries.
<code>“ “</code>	Return a new array of given shape and type, filled with ones.
<code>“ “</code>	Return a new array of given shape and type, filled with zeros.

## Mathematical functions

### Trigonometric functions

<code>sin(x)</code>	Trigonometric sine, element-wise.
<code>cos(x)</code>	Cosine elementwise.
<code>tan(x)</code>	Compute tangent element-wise.
<code>arcsin(x)</code>	Inverse sine, element-wise.
<code>arccos(x)</code>	Trigonometric inverse cosine, element-wise.
<code>arctan(x)</code>	Trigonometric inverse tangent, element-wise.

### Hyperbolic functions

<code>sinh(x)</code>	Hyperbolic sine, element-wise.
<code>cosh(x)</code>	Hyperbolic cosine, element-wise.
<code>tanh(x)</code>	Compute hyperbolic tangent element-wise.

## 1.1.6 PyCrop2ML User Guide

**Version** 1.1.0

**Release** 1.1.0

**Date** Oct 07, 2020

This reference manual details functions, modules, and objects included in OpenAlea.Core, describing what they are and what they do. For a complete reference guide, see core\_reference.

**Warning:** This “Reference Guide” is still very much in progress. Many aspects of OpenAlea.Core are not covered.

## Manual

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**Note:** The following examples assume you have installed the packages and setup your python path correctly.

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## Installation

```
conda install -c openalea pycropml
```

or

```
python setup.py install
```

## Overview of the different classes

### 1.1.7 src

#### pycropml package

##### Subpackages

##### pycropml.interface package

###### Submodules

###### pycropml.interface.design module

###### pycropml.interface.version module

##### Module contents

#### pycropml.transpiler package

##### Subpackages

##### pycropml.transpiler.generators package

###### Submodules

###### pycropml.transpiler.generators.checkGenerator module

```
class pycropml.transpiler.generators.checkGenerator.CheckCompo(tree,
                                                               model=None,
                                                               name=None)
Bases: pycropml.transpiler.generators.checkGenerator.CheckGenerator
```

This class used to generates states, rates and auxiliary classes for C# languages.

```
class pycropml.transpiler.generators.checkGenerator.CheckGenerator(tree,
                                                               model=None,
                                                               name=None)
Bases: pycropml.transpiler.codeGenerator.CodeGenerator, pycropml.transpiler.
rules.pythonRules.PythonRules
```

This class contains the specific properties of python language and use the NodeVisitor to generate a python code source from a well formed syntax tree.

```
visit_ExprStatNode (node)
visit_array (node)
visit_assignment (node)
visit_binary_op (node)
visit_bool (node)
visit_breakstatnode (node)
visit_call (node)
visit_comparison (node)
visit_cond_expr_node (node)
visit_continuestatnode (node)
visit_custom_call (node)
visit_datetime (node)
visit_declaration (node)
visit_dict (node)
visit_else_statement (node)
visit_elseif_statement (node)
visit_float (node)
visit_for_iterator (node)
visit_for_iterator_with_index (node)
visit_for_range_statement (node)
visit_for_sequence (node)
visit_for_sequence_with_index (node)
visit_for_statement (node)
visit_function_definition (node)
visit_if_statement (node)
visit_implicit_return (node)
visit_import (node)
visit_importfrom (node)
visit_index (node)
visit_list (node)
visit_local (node)
visit_method_call (node)
visit_module (node)
visit_notAnumber (node)
```

```
visit_pair(node)
visit_sliceindex(node)
visit_standard_call(node)
visit_standard_method_call(node)
visit_str(node)
visit_tuple(node)
visit_unary_op(node)
visit_while_statement(node)
```

## **pycropml.transpiler.generators.cppGenerator module**

### **pycropml.transpiler.generators.csharpGenerator module**

### **pycropml.transpiler.generators.docGenerator module**

```
class pycropml.transpiler.generators.docGenerator.DocGenerator(model=None,
                                                               tag='#')
Bases: pycropml.transpiler.codeGenerator.CodeGenerator
Generate doc in different target language - descrichtion of the code - Inputs details - Outputs details
comment(doc)
doc(x, name)
generate_desc(model)
generate_header(model)
```

## **pycropml.transpiler.generators.fortranGenerator module**

```
class pycropml.transpiler.generators.fortranGenerator.FortranCompo(tree=None,
                                                               model=None,
                                                               name=None)
Bases: pycropml.transpiler.generators.fortranGenerator.FortranGenerator
This class used to generates states, rates and auxiliary classes for Fortran90 language.
visit_importfrom(node)
    self.newline(node) self.write('Use ') for idx, item in enumerate(node.name):
        if idx: self.write(',')
        self.write("%smod"%item.split("model_")[1].capitalize())
class pycropml.transpiler.generators.fortranGenerator.FortranGenerator(tree,
                                                               model=None,
                                                               name=None)
Bases: pycropml.transpiler.codeGenerator.CodeGenerator, pycropml.transpiler.rules.fortranRules.FortranRules
```

This class contains the specific properties of fortran language and use the NodeVisitor to generate a fortran code source from a well formed syntax tree.

```
add_features(node)
```

```

binop_precedence = {'!=': 4, '%': 11, '&': 7, '*': 11, '**': 12, '+': 9, '-': 9
body (statements)
checkIndex (node)
doc = None
    # get constant parameters in models if inp.inputtype=="parameter":
        #print(inp.name, model.name) if inp.parametercategory=="constant":
            self.mod_parameters.append(inp.name)

    Type for inp in self.model.inputs

internal_declaration (node)
part_declaration (node)
retrieve_params (node)
subOrFun (node)
transform_return (node)
unop_precedence = {'!': 3, '+': 10, '-': 10, 'not': 3, '~': 10}
visit_ExprStatNode (node)
visit_array_decl (node)
visit_assignment (node)
visit_binary_op (node)
visit_bool (node)
visit_bool_decl (node)
visit_breakstatnode (node)
visit_call (node)
visit_combine (node)
visit_comparison (node)
visit_cond_expr_node (node)
visit_continuestatnode (node)
visit_custom_call (node)
visit_datetime (node)
visit_datetime_decl (node)
visit_decl (nodeT)
visit_declaration (node)
visit_else_statement (node)
visit_elseif_statement (node)
visit_float (node)
visit_float_decl (node)
visit_for_iterator (node)

```

```
visit_for_iterator_with_index(node)
visit_for_range_statement(node)
visit_for_sequence(node)
visit_for_statement(node)
visit_function(node)
visit_function_definition(node)
visit_if_statement(node)
visit_implicit_return(node)
visit_import(node)
visit_importfrom(node)
    self.newline(node) self.write('from %s import ' % (node.namespace)) for idx, item in enumerate(node.name):
        if idx: self.write(',')
            self.write(item)
visit_index(node)
visit_int(node)
visit_int_decl(node)
visit_list(node)
visit_list_decl(node)
visit_local(node)
visit_method_call(node)
visit_module(node)
visit_notAnumber(node)
visit_pair(node)
visit_sliceindex(node)
visit_standard_call(node)
visit_standard_method_call(node)
visit_str(node)
visit_str_decl(node)
visit_subroutine(node)
visit_subroutine_def(node)
visit_tab(node)
visit_unary_op(node)
visit_while_statement(node)

pycrocml.transpiler.generators.fortranGenerator.checkList(list1, list2)
pycrocml.transpiler.generators.fortranGenerator.valParam(model, name)
```

## pycrocml.transpiler.generators.javaGenerator module

```
class pycrocml.transpiler.generators.javaGenerator.JavaCompo (tree, model=None,  

                                                 name=None)  

Bases:   pycrocml.transpiler.generators.javaGenerator.JavaTrans, pycrocml.  

         transpiler.generators.javaGenerator.JavaGenerator

This class used to generates states, rates and auxiliary classes for java language.

copyconstructor (node)  

get_mo (varname)  

initCompo ()  

instanceModels ()  

setCompo (p)  

visit_assignment (node)  

visit_declaration (node)  

visit_function_definition (node)  

visit_implicit_return (node)  

visit_module (node)  

visit_return (node)  

  
class pycrocml.transpiler.generators.javaGenerator.JavaGenerator (tree,  

                                                 model=None,  

                                                 name=None)  

Bases: pycrocml.transpiler.codeGenerator.CodeGenerator, pycrocml.transpiler.  

        rules.javaRules.JavaRules
```

This class contains the specific properties of Java language and use the NodeVisitor to generate a java code source from a well formed syntax tree.

```
add_features (node)  

gettype (arg)  

internal_declaration (node)  

retrieve_params (node)  

transform_return (node)  

visit_array (node)  

visit_array_decl (node)  

visit_assignment (node)  

visit_binary_op (node)  

visit_bool (node)  

visit_bool_decl (node)  

visit_breakstatnode (node)  

visit_call (node)  

visit_comparison (node)  

visit_cond_expr_node (node)
```

```
visit_constant (node)
visit_continuestatnode (node)
visit_custom_call (node)
    TODO
visit_datetime_decl (node)
visit_decl (node)
visit_declaration (node)
visit_dict (node)
visit_dict_decl (node)
visit_else_statement (node)
visit_elseif_statement (node)
visit_float (node)
visit_float_decl (node)
visit_for_iterator (node)
visit_for_iterator_with_index (node)
visit_for_range_statement (node)
visit_for_sequence_with_index (node)
visit_for_statement (node)
visit_function_definition (node)
visit_if_statement (node)
visit_implicit_return (node)
visit_import (node)
visit_importfrom (node)
visit_index (node)
visit_int_decl (node)
visit_list (node)
visit_list_decl (node)
visit_method_call (node)
visit_module (node)
visit_notAnumber (node)
visit_pair (node)
visit_print ()
visit_return (node)
visit_sliceindex (node)
visit_standard_call (node)
visit_standard_method_call (node)
```

```
visit_str (node)
visit_str_decl (node)
visit_tuple (node)
visit_tuple_decl (node)
visit_unary_op (node)
visit_while_statement (node)

class pycropml.transpiler.generators.javaGenerator.JavaTrans (models)
    Bases: pycropml.transpiler.codeGenerator.CodeGenerator, pycropml.transpiler.rules.javaRules.JavaRules

This class used to generates states, rates and auxiliary classes for java language.

DATATYPE = {'BOOLEAN': 'bool', 'DATE': 'datetime', 'DATEARRAY': ['array', 'datetime'],
access (node)
copyconstructor (node)
generate (nodes, typ)
getset (node)
gettype (arg)
model2Node ()
private (node)
visit_array_decl (node)
visit_bool_decl (node)
visit_datetime_decl (node)
visit_decl (node)
visit_dict_decl (node)
visit_float_decl (node)
visit_int_decl (node)
visit_list_decl (node)
visit_str_decl (node)
visit_tuple_decl (node)

pycropml.transpiler.generators.javaGenerator.argsToStr (args)
pycropml.transpiler.generators.javaGenerator.to_struct_java (models, rep, name)
```

## pycropml.transpiler.generators.openaleaGenerator module

```
class pycropml.transpiler.generators.openaleaGenerator.OpenaleaCompo (tree,
    model=None,
    name=None)
    Bases: pycropml.transpiler.generators.pythonGenerator.PythonCompo
```

This class used to generates states, rates and auxiliary classes for C# languages.

**generate\_factory** (*model*)

Create a Node Factory from CropML model unit.

**generate\_wralea** (*mc*)

Generate wralea factories from the meta-information of the the model units.

**class** pycropml.transpiler.generators.openaleaGenerator.**OpenaleaGenerator** (*tree=None*,  
*model=None*,  
*name=None*)

Bases: *pycropml.transpiler.generators.pythonGenerator.PythonGenerator*

This class contains the specific properties of OpenAlea and use the NodeVisitor to generate a csharp code source from a well formed syntax tree.

pycropml.transpiler.generators.openaleaGenerator.**openalea\_interface** (*inout*)

pycropml.transpiler.generators.openaleaGenerator.**signature** (*model*)

## pycropml.transpiler.generators.pythonGenerator module

**class** pycropml.transpiler.generators.pythonGenerator.**PythonCompo** (*tree*,  
*model=None*,  
*name=None*)

Bases: *pycropml.transpiler.generators.pythonGenerator.PythonGenerator*

This class used to generates states, rates and auxiliary classes for C# languages.

**class** pycropml.transpiler.generators.pythonGenerator.**PythonGenerator** (*tree*,  
*model=None*,  
*name=None*)

Bases: *pycropml.transpiler.codeGenerator.CodeGenerator*, *pycropml.transpiler.rules.pythonRules.PythonRules*

This class contains the specific properties of python language and use the NodeVisitor to generate a python code source from a well formed syntax tree.

**comment** (*doc*)

**visit\_ExprStatNode** (*node*)

**visit\_array** (*node*)

**visit\_assignment** (*node*)

**visit\_binary\_op** (*node*)

**visit\_bool** (*node*)

**visit\_breakstatnode** (*node*)

**visit\_call** (*node*)

**visit\_comparison** (*node*)

**visit\_cond\_expr\_node** (*node*)

**visit\_continuestatnode** (*node*)

**visit\_custom\_call** (*node*)

**visit\_datetime** (*node*)

**visit\_declaration** (*node*)

**visit\_dict** (*node*)

---

```

visit_else_statement (node)
visit_elseif_statement (node)
visit_float (node)
visit_for_iterator (node)
visit_for_iterator_with_index (node)
visit_for_range_statement (node)
visit_for_sequence (node)
visit_for_sequence_with_index (node)
visit_for_statement (node)
visit_function_definition (node)
visit_if_statement (node)
visit_implicit_return (node)
visit_import (node)
visit_importfrom (node)
visit_index (node)
visit_list (node)
visit_method_call (node)
visit_module (node)
visit_notANumber (node)
visit_pair (node)
visit_sliceindex (node)
visit_standard_call (node)
visit_standard_method_call (node)
visit_str (node)
visit_tuple (node)
visit_unary_op (node)
visit_while_statement (node)

```

## pycroml.transpiler.generators.rGenerator module

```

class pycroml.transpiler.generators.rGenerator.RCompo (tree,           model=None,
                                                 name=None)
Bases: pycroml.transpiler.generators.rGenerator.RGenerator

This class used to generates states, rates and auxiliary classes for C# languages.

visit_tuple (node)

class pycroml.transpiler.generators.rGenerator.RGenerator (tree,      model=None,
                                                 name=None)
Bases: pycroml.transpiler.codeGenerator.CodeGenerator, pycroml.transpiler.rules.rRules.RRules

```

This class contains the specific properties of R language and use the NodeVisitor to generate a R code source from a well formed syntax tree.

```
comment (doc)
multValreturn (node)
visit_ExprStatNode (node)
visit_array (node)
visit_assignment (node)
visit_binary_op (node)
visit_bool (node)
visit_breakstatnode (node)
visit_comparison (node)
visit_cond_expr_node (node)
visit_continuestatnode (node)
visit_datetime (node)
visit_declaration (node)
visit_else_statement (node)
visit_elseif_statement (node)
visit_float (node)
visit_for_iterator (node)
visit_for_iterator_with_index (node)
visit_for_range_statement (node)
visit_for_sequence (node)
visit_for_sequence_with_index (node)
visit_for_statement (node)
visit_function_definition (node)
visit_if_statement (node)
visit_implicit_return (node)
visit_import (node)
visit_importfrom (node)
visit_index (node)
visit_list (node)
visit_method_call (node)
visit_module (node)
visit_notAnumber (node)
visit_pair (node)
visit_sliceindex (node)
```

---

```
visit_standard_call(node)
visit_standard_method_call(node)
visit_str(node)
visit_tuple(node)
visit_unary_op(node)
visit_while_statement(node)
```

**pycropml.transpiler.generators.recordGenerator module**

```
<vle_project version="1.1.x" date="2012-Oct-03 13:01:13" author="Eric Casellas">
    <structures>
        <model width="2184" height="1280" name="2CV_parcelle" type="coupled">
            <submodels> <model observables="vueDecision" conditions="condDecFSA" dynamics="dynDecFSA" debug="false" width="100" height="75" x="132" y="26" name="Decision" type="atomic"></model> <model width="100" height="165" x="316" y="127" name="2CV" type="coupled"></model>
        </submodels> <connections> </connections>
    </model>
</structures> <dynamics> </dynamics> <experiment name="2CV_parcelle"> </experiment>
</vle_project>
```

**pycropml.transpiler.generators.simplaceGenerator module**

```
class pycropml.transpiler.generators.simplaceGenerator.SimplaceCompo(tree,
    model=None,
    name=None)
Bases: pycropml.transpiler.generators.javaGenerator.JavaGenerator

class pycropml.transpiler.generators.simplaceGenerator.SimplaceGenerator(tree,
    model=None,
    name=None)
Bases: pycropml.transpiler.generators.javaGenerator.JavaGenerator

visit_declaration(node)
visit_function_definition(node)
visit_import(node)
visit_local(node)
visit_module(node)
visit_return(node)
```

## pycropml.transpiler.generators.siriusGenerator module

### Module contents

#### pycropml.transpiler.lib package

### Module contents

#### pycropml.transpiler.rules package

### Submodules

#### pycropml.transpiler.rules.cppRules module

```
class pycropml.transpiler.rules.cppRules.CppRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    binary_op = {'!=': '!=', '*': '*', '+': '+', '-': '-', '/': '/', '<': '<', '<=':
    constant = {'math': {'pi': 'M_PI'}}
    constructor = '\n %s::%s()\n {\n \n }\n '
    copy_constr = '\n public %s(%s toCopy, bool copyAll) // copy constructor\n {\n if (copyAll)\n     copyConstr(toCopy);\n }'
    copy_constrArray = '\n for (int i = 0; i < %s; i++)\n {\n     _%s[i] = toCopy._%s[i];\n }'
    copy_constrList = '\n for (int i = 0; i < toCopy.%s.Count; i++)\n {\n     %s.Add(toCopy.%s[i]);\n }'
    copy_constr_compo = '\nvoid %s(%s& toCopy): this() // copy constructor\n{\n    copyConstr(toCopy);\n}'
    functions = {'datetime': {'datetime': 'new DateTime'}, 'io': {'print': 'cout<<'}, 'methods': {'array': {'append': '.Add', 'len': <function translateLenArray>}, 'dict': {'public_properties': '\n    get\n    {\n        return this._%s;\n    }\n    set\n    {\n        this._%s = value;\n    }', 'public_properties_compo': '\n    get\n    {\n        return _%s.%s;\n    }\n    set\n    {\n        %s.%s = value;\n    }', 'public_properties_wrap': '{\n        get\n        {\n            return %s.%s;\n        }\n        set\n        {\n            %s.%s = value;\n        }\n    }', 'types': {'array': 'array<%s, %s>', 'bool': 'bool', 'datetime': 'string', 'dict': '{}', 'unary_op': {'+': '+', '-': '-', 'not': '!', '~': '~'}}}, 'translateContains': (node), 'translateIndex': (node), 'translateInsert': (node), 'translateLenArray': (node), 'translateLenDict': (node), 'translateLenList': (node), 'translateLenStr': (node), 'translateMIN': (node), 'translateNotContains': (node), 'translatePop': (node)}
```

---

```
pycropml.transpiler.rules.cppRules.translateSum(node)
pycropml.transpiler.rules.cppRules.translateget(node)
pycropml.transpiler.rules.cppRules.translatekeyDict(node)
```

## pycropml.transpiler.rules.csharpRules module

```
class pycropml.transpiler.rules.csharpRules.CsharpRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    binary_op = {'!=': '!=', '*': '*', '+': '+', '-': '-', '/': '/', '<': '<', '<=':
    constant = {'math': {'pi': 'Math.PI'}}
    constructor = '\n    public %s() { }\n'
    copy_constr = '\n    public %s(%s toCopy, bool copyAll) // copy constructor\n    {\n        if (copyAll)\n            foreach (var item in toCopy)\n                Add(item);\n    }\n'
    copy_constrArray = '\n    for (int i = 0; i < %s; i++)\n        _%s[i] = toCopy._%s[i];\n'
    copy_constrList = '\n    for (int i = 0; i < toCopy.%s.Count; i++)\n        %s.Add(toCopy.%s[i]);\n'
    copy_constr_compo = '\n    public %s(%s toCopy): this() // copy constructor\n    {\n        foreach (var item in toCopy)\n            Add(item);\n    }\n'
    functions = {'datetime': {'datetime': 'new DateTime'}, 'io': {'print': <function print>}}
    methods = {'array': {'append': '.Add', 'len': <function translateLenArray>}, 'dict': {'clear': 'Clear', 'keys': 'Keys', 'values': 'Values'}}
    public_properties = '\n    get { return this._%s; }\n    set { this._%s= value; }\n'
    public_properties_compo = '\n    get { return _%s.%s; }\n    set\n        {\n            %s = value;\n            foreach (var item in %s)\n                item.%s = value;\n        }\n'
    public_properties_wrap = '{\n        get { return %s.%s; }\n        set { %s.%s= value; }\n    }\n'
    types = {'DateTime': 'DateTime', 'array': '%s[] %s= new %s', 'bool': 'bool', 'date': 'Date', 'double': 'double', 'float': 'float', 'int': 'int', 'long': 'long', 'string': 'string'}
    unary_op = {'+': '+', '-': '-', 'not': '!', '~': '~'}

    pycropml.transpiler.rules.csharpRules.linq(name, z=True, swap=False)
    pycropml.transpiler.rules.csharpRules.translateLenArray(node)
    pycropml.transpiler.rules.csharpRules.translateLenDict(node)
    pycropml.transpiler.rules.csharpRules.translateLenList(node)
    pycropml.transpiler.rules.csharpRules.translateNotContains(node)
    pycropml.transpiler.rules.csharpRules.translatePow(node)
    pycropml.transpiler.rules.csharpRules.translatePrint(node)
    pycropml.transpiler.rules.csharpRules.translateSum(node)
    pycropml.transpiler.rules.csharpRules.translateget(node)
    pycropml.transpiler.rules.csharpRules.translatekeyDict(node)
    pycropml.transpiler.rules.csharpRules.translatevalueDict(node)
```

## pycropml.transpiler.rules.fortranRules module

```
class pycropml.transpiler.rules.fortranRules.FortranRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    binary_op = {'!=': '.NE.', '*': '*', '**': '**', '+': '+', '-': '-', '/': '/',
    functions = {'datetime': {'datetime': <function <lambda>>}, 'io': {'print': <funct
    method()

    methods = {'array': {'append': <function <lambda>>, 'len': 'SIZE'}, 'dict': {'len'
        dependencies = {

            'list': { 'index': 'list_sub', 'append': 'list_sub'

            }

        }

    }

    types = {'array': '%s, DIMENSION(%s)', 'bool': 'LOGICAL', 'datetime': 'CHARACTER(65
    unary_op = {'+': '+', '-': '-', 'not': '.NOT.', '~': '~'}

pycropml.transpiler.rules.fortranRules.argsToStr(args)
pycropml.transpiler.rules.fortranRules.translateAppend(node)
pycropml.transpiler.rules.fortranRules.translateCeil(node)
pycropml.transpiler.rules.fortranRules.translateContains(node)
pycropml.transpiler.rules.fortranRules.translateFind(node)
pycropml.transpiler.rules.fortranRules.translateIndex(node)
pycropml.transpiler.rules.fortranRules.translateMIN(node)
pycropml.transpiler.rules.fortranRules.translateNotContains(node)
pycropml.transpiler.rules.fortranRules.translatePop(node)
pycropml.transpiler.rules.fortranRules.translatePow(node)
pycropml.transpiler.rules.fortranRules.translatePrint(node)
```

## pycropml.transpiler.rules.generalRule module

```
class pycropml.transpiler.rules.generalRule.GeneralRule
    Bases: object

    """ Abstract class of Rules
```

## pycropml.transpiler.rules.javaRules module

```
class pycropml.transpiler.rules.javaRules.JavaRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    binary_op = {'!=': '!=', '*': '*', '+': '+', '-': '-', '/': '/', '<': '<', '<='
    constant = {'math': {'pi': 'Math.PI'}}
    constructor = '\n public %s() { }'
```

```

copy_constr = '\n public %s(%s toCopy, boolean copyAll) // copy constructor \n {\n if
copy_constrArray = '\n for (int i = 0; i < %s; i++)\n {\n _%s[i] = toCopy._%s[i];\n }'
copy_constrList = '\n for (%s c : toCopy.%s)\n {\n _%s.add(c);\n }\n this.%s = _%s;'
copy_constr_compo = '\n public %s(%s toCopy) // copy constructor \n {'
functions = {'datetime': {'datetime': <function translateDateTime>}, 'io': {'print':
get_properties = '\n { return %s; }'
get_properties_compo = '\n { return _%s.get%s(); }'
methods = {'array': {'append': '.add', 'len': <function translateLenArray>}, 'dict':
set_properties = '\n { this.%s= _%s; } \n '
set_properties_compo = '\n { %s } '
types = {'array': '%s[] %s= new %s', 'bool': 'boolean', 'datetime': 'Date', 'dict':
types2 = {'Date': 'Date', 'bool': 'Boolean', 'datetime': 'Date', 'float': 'Double'
unary_op = {'+': '+', '-': '-', 'not': '!', '~': '~'}
pycropml.transpiler.rules.javaRules.argsToStr(args)
pycropml.transpiler.rules.javaRules.translateDateTime(node)
pycropml.transpiler.rules.javaRules.translateDictValues(node)
pycropml.transpiler.rules.javaRules.translateDictkeys(node)
pycropml.transpiler.rules.javaRules.translateLenArray(node)
pycropml.transpiler.rules.javaRules.translateLenDict(node)
pycropml.transpiler.rules.javaRules.translateLenList(node)
pycropml.transpiler.rules.javaRules.translateNotContains(node)
pycropml.transpiler.rules.javaRules.translatePow(node)
pycropml.transpiler.rules.javaRules.translatePrint(node)
pycropml.transpiler.rules.javaRules.translateSum(node)

```

## pycropml.transpiler.rules.pythonRules module

```

class pycropml.transpiler.rules.pythonRules.PythonRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    binary_op = {'!=': '!=', '%': '%', '*': '*', '+': '+', '-': '-', '/': '/',
    '<': '<'
    functions = {'datetime': {'datetime': 'datetime'}, 'io': {'print': <function trans
    methods = {'array': {'append': '.append', 'len': 'len'}, 'datetime': {'datetime':
    types = {'bool': 'bool', 'datetime': 'datetime', 'dict': 'dict', 'float': 'float',
    unary_op = {'+': '+', '-': '-', 'not': 'not ', '~': '~'}
pycropml.transpiler.rules.pythonRules.translateDictkeys(node)
pycropml.transpiler.rules.pythonRules.translateModulo(node)
pycropml.transpiler.rules.pythonRules.translateNotContains(node)

```

```
pycropml.transpiler.rules.pythonRules.translatePrint(node)
```

## pycropml.transpiler.rules.rRules module

```
class pycropml.transpiler.rules.rRules.RRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    binary_op = {'!=': '!=', '%': '%%', '*': '*', '+': '+', '-': '-', '/': '/', '<'}
    functions = {'datetime': {'datetime': <function <lambd>>}, 'io': {'print': <funct>
methods = {'array': {'append': '.append', 'len': 'len'}, 'datetime': {'datetime': <function <lambd>>}}
types = {'bool': 'bool', 'datetime': 'str', 'float': 'float', 'int': 'int', 'list': 'list'}
unary_op = {'+': '+', '-': '-', 'not': 'not ', '~': '~'}

pycropml.transpiler.rules.rRules.argsToStr(args)
pycropml.transpiler.rules.rRules.translateDictkeys(node)
pycropml.transpiler.rules.rRules.translateModulo(node)
pycropml.transpiler.rules.rRules.translateNotContains(node)
pycropml.transpiler.rules.rRules.translatePrint(node)
```

## pycropml.transpiler.rules.sqRules module

### Module contents

#### Submodules

### pycropml.transpiler.Parser module

```
class pycropml.transpiler.Parser.opt(**kwds)
```

```
pycropml.transpiler.Parser.parser(module)
```

Read, parse a Cython code and generate an abstract syntax tree.

Context: Compilation context: contains every pxd ever loaded, path information and the data related to the compilation. Class where it is implemented Cython parse method.

**options:** To set Compilation Options as language\_level: The source language level to use, formal\_grammar: to define if it will be used to Parse the file evaluate\_tree\_assertions: To evaluate parse tree show\_version : To display version number use\_listing\_file: Generate a .lis file errors\_to\_stderr: Echo errors to stderr when using .lis include\_path: Directories to search for include files output\_file: Name of generated .c file generate\_pxi: Generate .pxi file for public declarations capi\_reexport\_cincludes: Add cincluded headers to any auto-generated header files. timestamps: Only compile changed source files verbose : Always print source names being compiled compiler\_directives: Overrides for pragma options (see Options.py) embedded\_metadata: Metadata to embed in the C file as json. evaluate\_tree\_assertions: Test support: evaluate parse tree assertions cplus : Compile as c++ code

Here default options were used except language level

Scanning.FileSourceDescriptor: Represents a code source. Only file sources for Cython code supported

**pycropml.transpiler.api\_transform module**

```
class pycropml.transpiler.api_transform.Standard
    Standard classes should respond to expand and to return valid nodes on expand

class pycropml.transpiler.api_transform.StandardCall(namespace, function, ex-
    pander=None)
    Bases: pycropml.transpiler.api_transform.Standard
    converts to a standard call of the given namespace and function

    expand(args)

class pycropml.transpiler.api_transform.StandardCallAttrib(namespace, function,
    expander=None)
    Bases: pycropml.transpiler.api_transform.Standard
    converts to a standard call of the given namespace and function

    expand(args=[])

class pycropml.transpiler.api_transform.StandardMethodCall(type, message, de-
    fault=None, ex-
    pander=None)
    Bases: pycropml.transpiler.api_transform.Standard
    converts to a method call of the same class

    expand(args)

class pycropml.transpiler.api_transform.StandardSwapper(type, message)
    Bases: pycropml.transpiler.api_transform.Standard

    expand(args)

pycropml.transpiler.api_transform.abs_expander(type, message, args)
pycropml.transpiler.api_transform.array_expander(type, message, args)
pycropml.transpiler.api_transform.datetime_expander(type, message, args)
pycropml.transpiler.api_transform.float_expander(type, message, args)
pycropml.transpiler.api_transform.int_expander(type, message, args)
pycropml.transpiler.api_transform.len_expander(type, message, args)
pycropml.transpiler.api_transform.max_expander(type, message, args)
pycropml.transpiler.api_transform.min_expander(type, message, args)
pycropml.transpiler.api_transform.modulo_expander(type, message, args)
pycropml.transpiler.api_transform.pow_expander(type, message, args)
```

**pycropml.transpiler.ast\_transform module**

```
class pycropml.transpiler.ast_transform.AstTransformer(tree, code, model=None)

    accessReturn(node)
    assert_translatable(node, **pairs)
    checktype(base)
```

```
newtype (name)
notdeclared (name, line)
retrieve_library (func)
transformer ()
translate_comprehensionnode (rhs, lhs, location)
visit_addnode (node, operand1, operand2, location)
visit_attributenode (node, obj, location)
visit_binopnode (node, operand1, operand2, location)
visit_boolbinopnode (node, operand1, operand2, location)
visit_boolnode (node, location)
visit_breakstatnode (node, location)
visit_cargdeclnode (node, base_type, declarator, default, annotation, location)
visit_comprehensionappendnode (node, expr, location)
visit_comprehensionnode (node, loop, location)
visit_condexprnode (node, test, true_val, false_val, location)
visit_continuestatnode (node, location)
visit_csimplebasetypenode (node, location)
visit_cstructoruniondefnode (node, attributes, location)
visit_cvardefnode (node, base_type, declarators, location)
visit_definitions ()
visit_defnode (node, args, star_arg, starstar_arg, decorators, body, return_type_annotation, location)
visit_dictnode (node, key_value_pairs, location)
visit_divnode (node, operand1, operand2, location)
visit_elements (elements, kind, homogeneous=True)
visit_exprstatnode (node, expr, location)
visit_floatnode (node, location)
    if float(node.value) < 0.0: return { 'type': 'unary_op',
        'operator': '-', 'value': str(-float(node.value)), 'pseudo_type': "float" }
visit_forinstatnode (node, target, iterator, item, body, else_clause, location)
visit_ifclausenode (node, body, condition, location)
visit_ifstatnode (node, if_clauses, else_clause, location)
visit_indexnode (node, base, index, location)
visit_inplaceassignmentnode (node, lhs, rhs, location)
visit_intnode (node, location)
visit_listnode (node, args, mult_factor, location)
visit_modnode (node, operand1, operand2, location)
```

```
visit_mulnode(node, operand1, operand2, location)
visit_namenode(node, location)
visit_node(node)
visit_notnode(node, operand, location)
visit_pownode(node, operand1, operand2, location)
visit_primarycmpnode(node, operand1, operand2, coerced_operand2, cascade, location)
visit_printstatnode(node, arg_tuple, stream, location)
visit_pyclassdefnode(node, location)
visit_returnstatnode(node, value, location)
visit_simplecallnode(node, function, coerced_self, args, arg_tuple, location)
visit_singleassignmentnode(node, lhs, rhs, location)
visit_sliceindexnode(node, start, stop, base, slice, location)
visit_statlistnode(node, stats, location)
visit_stringnode(node, location)
visit_subnode(node, operand1, operand2, location)
visit_top_level(nodes)
visit_tuplenode(node, args, mult_factor, location)
visit_unaryminusnode(node, operand, location)
visit_unaryplusnode(node, operand, location)
visit_unicodeinode(node, location)
visit_whilestatnode(node, condition, body, else_clause, location)

pycropml.transpiler.ast_transform.transform_to_syntax_tree(tree)
Generate a Node class from the tree in dict format
```

## pycropml.transpiler.builtin\_typed\_api module

```
pycropml.transpiler.builtin_typed_api.add(l, r)
pycropml.transpiler.builtin_typed_api.and_(l, r)
pycropml.transpiler.builtin_typed_api.arg_check(expected_type, args, a)
pycropml.transpiler.builtin_typed_api.binary_and(l, r)
pycropml.transpiler.builtin_typed_api.binary_or(l, r)
pycropml.transpiler.builtin_typed_api.builtin_type_check(namespace, function, receiver, args)
pycropml.transpiler.builtin_typed_api.div(l, r, lo)
pycropml.transpiler.builtin_typed_api.mod(l, r)
pycropml.transpiler.builtin_typed_api.mul(l, r)
pycropml.transpiler.builtin_typed_api.or_(l, r)
pycropml.transpiler.builtin_typed_api.pow_(l, r)
```

```
pycropml.transpiler.builtin_typed_api.simplify(kind, generics)
pycropml.transpiler.builtin_typed_api.sub(l, r)
```

## pycropml.transpiler.checkingModel module

```
class pycropml.transpiler.checkingModel.Checking
```

Module used to check units validity in model equation based on model xml files. This checking can also use for python code with metadata

## pycropml.transpiler.codeGenerator module

```
class pycropml.transpiler.codeGenerator.CodeGenerator(add_line_information=False)
    Bases: pycropml.transpiler.nodeVisitor.NodeVisitor

    binop_precedence = {'!=': 4, '%': 10, '&': 7, '*': 10, '**': 12, '+': 9, '-': 9}
    body(statements)
    body_or_else(node)
    comma_separated_list(items)
    emit_sequence(node, parens=(u" ", u"))
    emit_string(node, prefix=u")
    newline(node=None, extra=0)
    operator_enter(new_prec)
    operator_exit()
    safe_double(node)
    unop_precedence = {'!': 3, '+': 11, '-': 11, 'not': 3, '~': 11}
    visit_ExprStatNode(node)
    visit_array(node)
    visit_call(node)
    visit_custom_call(node)
    visit_for_sequence(node)
    visit_int(node)
    visit_local(node)
    visit_simpleCall(node)
    write(x)
```

## pycropml.transpiler.env module

```
class pycropml.transpiler.env.Env(values=None, parent=None)
```

```
    child_env(values=None)
```

## pycropml.transpiler.errors module

```
exception pycropml.transpiler.errors.PseudoCythonNotTranslatableError(message,  

    sug-  

    ges-  

    tions=None,  

    right=None,  

    wrong=None)  

    Bases: pycropml.transpiler.errors.PseudoError  

exception pycropml.transpiler.errors.PseudoCythonTypeCheckError(message,  

    sug-  

    ges-  

    tions=None,  

    right=None,  

    wrong=None)  

    Bases: pycropml.transpiler.errors.PseudoError  

exception pycropml.transpiler.errors.PseudoError(message, suggestions=None,  

    right=None, wrong=None)  

    Bases: exceptions.Exception  

pycropml.transpiler.errors.beautiful_error(exception)  

pycropml.transpiler.errors.cant_infer_error(name, line)  

pycropml.transpiler.errors.tab_aware(location, code)  

    if tabs in beginning of code, add tabs for them, otherwise spaces  

pycropml.transpiler.errors.translation_error(data, location=None, code=None,  

    wrong_type=None, **options)  

pycropml.transpiler.errors.type_check_error(data, location=None, code=None,  

    wrong_type=None, **options)
```

## pycropml.transpiler.helpers module

```
pycropml.transpiler.helpers.prepare_table(types, original_methods=None)  

pycropml.transpiler.helpers.safe_serialize_type(l)  

    serialize only with letters, numbers and _  

pycropml.transpiler.helpers.serialize_type(l)
```

## pycropml.transpiler.interface module

```
class pycropml.transpiler.interface.TreeInterface(tree)  

    visits recursively nodes of the tree with defined transform_<node_type> methods and transforms in place  

        transform(tree, in_block=False)  

        transform_block(tree)  

        transform_default(tree)  

class pycropml.transpiler.interface.middleware(tree)  

    Bases: pycropml.transpiler.interface.TreeInterface  

        api_translate()
```

## pycropml.transpiler.main module

### pycropml.transpiler.nodeVisitor module

```
class pycropml.transpiler.nodeVisitor.NodeVisitor
Bases: object
```

Define a method which browse the graph and call a methode constructed from the type of each node of the graph

```
visit(node)
```

## pycropml.transpiler.pseudo\_tree module

```
class pycropml.transpiler.pseudo_tree.Node(type, **fields)
```

The new Node generated with specific properties. These properties are automatically set”

Example: Node(type='local', name='l', pseudo\_type="int") to represent a int variable declaration

```
y
```

## pycropml.transpiler.version module

Maintain version for this package. Do not edit this file, use ‘version’ section of config.

```
pycropml.transpiler.version.MAJOR = 0
```

(int) Version major component.

```
pycropml.transpiler.version.MINOR = 0
```

(int) Version minor component.

```
pycropml.transpiler.version.POST = 2
```

(int) Version post or bugfix component.

## Module contents

### Submodules

#### pycropml.algorithm module

```
class pycropml.algorithm.Algorithm(language, development, platform, filename=None)
Bases: object
```

#### pycropml.checking module

```
class pycropml.checking.Test(name)
Bases: pycropml.checking.Testset
```

```
class pycropml.checking.Testset(name, parameterset, description, uri=None)
Bases: object
```

Test

```
pycropml.checking.testset(model, name, kwds)
```

**pycropml.code2nbk module****pycropml.composition module**

Read xml representation of a model composite

**class pycropml.composition.Description**  
Bases: `object`

Model Composition Description.

**A description is defined by:**

- Title
- Authors
- Institution
- Reference
- Abstract

**class pycropml.composition.ModelComposition(kwds)**  
Bases: `pycropml.composition.ModelDefinition`

Formal description of a Model Composite.

**add\_description(description)**  
TODO

**class pycropml.composition.ModelDefinition(kwds)**  
Bases: `object`

Model name, id, version and step

**class pycropml.composition.ModelParser**  
Bases: `pycropml.composition.Parser`

Read an XML file and transform it in our object model.

**Composition(elts)**

**Description(Title, Author, Institution, Reference, Abstract)**

**Initialization(elt)**

**Links(elt)**

Retrieve different types of links

**Model(elt)**

Models

**ModelComposition(elts)**

ModelComposition (Description, Models, Inputlink,Outputlink,externallink)

**dispatch(elt)**

**parse(fn)**

**class pycropml.composition.Models(name, modelid, file, package\_name=None)**

Bases: `pycropml.composition.ModelComposition`, `pycropml.modelunit.ModelUnit`

**class pycropml.composition.Parser**  
Bases: `object`

Read an XML file and transform it in our object model.

**dispatch** (*elt*)

**parse** (*fn*)

`pycropml.composition.model_parser (fn)`

Parse a composite model and return the model.

Returns ModelComposite object of the CropML Model.

`pycropml.composition.retrieve_path (fn)`

## **pycropml.cyml module**

### **pycropml.description module**

**class** `pycropml.description.Description`

Bases: `object`

Model Unit Description.

**A description is defined by:**

- Title
- Author
- Institution
- Reference
- Abstract

## **pycropml.error module**

Created on Wed Apr 10 17:01:34 2019

@author: midingoy

**exception** `pycropml.error.Error (message)`

Bases: `exceptions.Exception`

## **pycropml.formater\_f90 module**

`pycropml.formater_f90.formater (code)`

`pycropml.formater_f90.formaterNext (line)`

## **pycropml.function module**

**class** `pycropml.function.Function (name, language, filename, type, description)`

Bases: `object`

## pycropml.initialization module

```
class pycropml.initialization.Initialization(name, language, filename)
```

Bases: object

Function

## pycropml.inout module

```
class pycropml.inout.Input(kwds)
```

Bases: pycropml.inout.InputOutput

```
class pycropml.inout.InputOutput(kwds)
```

Bases: object

```
class pycropml.inout.Output(kwds)
```

Bases: pycropml.inout.InputOutput

## pycropml.main module

### pycropml.model module

#### pycropml.modelunit module

Model Description and Model Unit.

```
class pycropml.modelunit.ModelDefinition(kwds)
```

Bases: object

```
class pycropml.modelunit.ModelUnit(kwds)
```

Bases: pycropml.modelunit.ModelDefinition

Formal description of a Model Unit.

```
add_description(description)
```

TODO

## pycropml.package module

```
from pycropml import composition from pycropml.pparse import model_parser from path import Path import networkx as nx from collections import defaultdict from IPython.display import Image, display from networkx.drawing.nx_pydot import to_pydot from pycropml.render_cyml import signature
```

```
class pycropml.package.AbstractPackageReader(filename)
```

Bases: object

Abstract class to add a package in the package manager.

```
register_packages(pkgmanager)
```

Create and add a package in the package manager.

```
class pycropml.package.Package(name, metainfo, path=None)
```

Bases: pycropml.package.PackageDict

A Package is a dictionnary of node factory. Each node factory is able to generate node and their widgets. Meta informations are associated with a package.

```
add_modelunit (modelunit)
    Add to the package a factory ( node or subgraph )

get_crop2ml_path ()
    Return the full path of the wralea.py (if set)

get_id ()
    Return the package id

get_metainfo (key)
    Return a meta information. See the standard key in the __init__ function documentation. :param key: todo

get_modelunit (modelid)
    Return the factory associated with id

get_names ()
    Return all the factory names in a list

get_pkg_files ()
    Return the list of xml filename of the package. The filename are relative to self.path

get_tip ()
    Return the package description

is_directory ()
    New style package. A package is embeded in a unique directory. This directory can not contain more than one package. Thus, you can move, copy or delete a package by acting on the directory without ambiguity.
    Return True if the package is embeded in a directory.

is_editable ()
    A convention (for the GUI) to ensure that the user can modify the package.

mimetype = 'pycrop2ml/package'

reload ()
    Reload all xml file of the package

remove_files ()
    Remove pkg files

update_modelunit (old_name, modelunit)
    Update factory (change its name)

class pycropml.package.PackageDict (*args)
Bases: dict

    Dictionary with case insensitive key This object is able to handle protected entry begining with an '#'
    get (k[, d]) → D[k] if k in D, else d. d defaults to None.

    has_key (k) → True if D has a key k, else False

    iter_public_values ()
        Iterate through dictionary value (remove protected value)

    nb_public_values ()
        Return the number of unprotected values

class pycropml.package.PackageManager

add_crop2ml_path (path, container)
    Add a search path for wralea files
```

#### Parameters

- **path** – a path string
- **container** – set containing the path

**add\_package** (*package*)  
 Add a package to the pkg manager

**clear** ()  
 Remove all packages

**create\_readers** (*crop2ml\_files*)

**find\_and\_register\_packages** (*no\_cache=False*)  
 Find all composite model on the system and register them If *no\_cache* is True, ignore cache file

**find\_crop2ml\_dir** (*directory, recursive=True*)  
 Find in a directory wralea files, Search recursively is recursive is True  
 :return : a list of pkgreader instances

**get** (\**args*)

**get\_pkgreader** (*filename*)  
 Return the pkg reader corresponding to the filename

**has\_key** (\**args*)

**init** (*dirname=None, verbose=True*)  
 Initialize package manager  
 If *dirname* is None, find composition files on the system else load directory

**items** ()

**iteritems** ()

**iterkeys** ()

**itervalues** ()

**keys** ()

**load\_directory** (*dirname*)  
 Load a directory containing wraleas

**rebuild\_category** ()  
 Rebuild all the category

**reload** (*pkg=None*)  
 Reload one or all packages. If the package *pkg* is None reload all the packages. Else reload only *pkg*.

**set\_sys\_crop2ml\_path** ()  
 Define the default composition files search path  
 For that, we look for “composition” entry points and deprecated\_wralea entry point if a package is declared as deprecated\_wralea, the module is not load

**update\_category** (*package*)  
 Update the category dictionary with package contents

**values** ()

**class** pycroml.package.**PseudoGroup** (*name*)  
 Bases: *pycroml.package.PackageDict*  
 Data structure used to separate dotted naming (packages, category)

```
add_name (name, value)
    Add a value in the structure with the key name_tuple

get_id()
    todo

get_tip()
    todo

mimetype = 'pycrop2ml/package'

new(name)
    todo

sep = '.'

class pycrocml.package.PyPackageReader(filename)
    Bases: pycrocml.package.AbstractPackageReader
    Build packages from wralea file Use 'register_package' function

    build_package (wraleamodule, pkgmanager)
        Build package and update pkgmanager

    filename_to_module (filename)
        Transform the filename ending with .py to the module name

    get_pkg_name ()
        Return the OpenAlea (uniq) full package name

    register_packages (pkgmanager)
        Execute model.py

class pycrocml.package.PyPackageReaderModel(filename)
    Bases: pycrocml.package.PyPackageReader
    Build a package from a __wralea__.py Use module variable

    build_package (wraleamodule, pkgmanager)
        Build package and update pkgmanager

    check_exist ()
    contain_pkg (pkg)
    get_path (pkg, name)

class pycrocml.package.PyPackageWriter(package)
    Bases: object
    Write a wralea python file

    get_str ()
        Return string to write

    pkg_template = '\n$PKGNAME\n$METAINFO\n'

    wralea_template = '\n# This file has been generated at $TIME\n$PKG_DECLARATION\n'

    write_wralea (full_filename)
        Write the wralea.py in the specified filename

exception pycrocml.package.UnknownNodeError (name)
    Bases: exceptions.Exception
```

```
class pycropml.package.UserPackage(name, metainfo, path=None)
Bases: pycropml.package.Package

    Package user editable and persistent

pycropml.package.get_default_home_dir()
    Return the home directory (valid on linux and windows)

pycropml.package.get_openalea_home_dir(name='pycrop2ml')
    Return the crop2ml home directory If it doesn't exist, create it

pycropml.package.get_userpkg_dir(name='user_pkg')
    Get user package directory (the place where are the wralea.py files). If it doesn't exist, create it

pycropml.package.is_protected(item)
    Return true the item is protected

pycropml.package.lower(item)

pycropml.package.protected(item)
    Return corresponding protected name for item
```

## pycropml.parameterset module

```
class pycropml.parameterset.Parameterset(name, description, uri=None)
Bases: object

    Parameter set

pycropml.parameterset.parameterset(model, name, kwds)
```

## pycropml.pparse module

License, Header

```
class pycropml.pparse.ModelParser
Bases: pycropml.pparse.Parser

    Read an XML file and transform it in our object model.

    Algorithm(elt)
    Description>Title, Author, Institution, Reference, Abstract
    Function(elt)
    Initialization(elt)
    Input(elts)
    Inputs(Input)
    ModelUnit(elts)
        ModelUnit (Description,Inputs,Outputs,Algorithm,Parametersets, Testsets)
    Output(elts)
    Outputs(elts)
        Outputs (Output)
    Parameterset(elts)
    Parametersets(Parameterset)
```

```
Testset (Test)
Testsets (Testset)
dispatch (elt)
param (pset, elt)
    Param
parse (crop2ml_dir)

class pycropml.pparse.Parser
    Bases: object

    Read an XML file and transform it in our object model.

    dispatch (elt)
    parse (crop2ml_dir)

pycropml.pparse.model_parser (crop2ml_dir)

Parse a set of models as xml files contained in crop2ml directory and algorithm in src directory This function returns models as python object.

    Returns ModelUnit object of the Crop2ML Model.
```

## **pycropml.render\_R module**

Add License, Header.

Use pkglts

Problems: - name of a model unit?

```
Model2Package (models, dir=None, pkg_name=None)
    Bases: object

    TODO
    generate_func_test (model_unit)
    generate_function_doc (model_unit)
    generate_test (model_unit)
    num = 0
    run ()
        TODO.

    write_tests ()
        TODO: Manage several models rather than just one.

pycropml.render_R.comment (line)
pycropml.render_R.generate_doc (model)
pycropml.render_R.signature (model)
pycropml.render_R.transf (type_v, elem)
```

## pycropml.render\_cpp module

Add License, Header.

Use pkglets

Problems: - name of a model unit?

```
class pycropml.render_cpp.Model2Package (models, dir=None)
    Bases: object
```

TODO

```
DATATYPE = {'BOOLEAN': 'bool', 'DATE': 'string', 'DATELIST': 'vector<string>', 'DOUBLE':
```

```
generate_test (model_unit)
```

```
num = 0
```

```
write_tests ()
```

TODO: Manage several models rather than just one.

```
pycropml.render_cpp.signature (model)
```

```
pycropml.render_cpp.transf (type_v, elem)
```

```
pycropml.render_cpp.transfDate (type, elem)
```

```
pycropml.render_cpp.transfDateList (type, elem)
```

```
pycropml.render_cpp.transfDouble (type_v, elem)
```

```
pycropml.render_cpp.transfList (type_v, elem)
```

```
pycropml.render_cpp.transfString (type_v, elem)
```

## pycropml.render\_csharp module

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Problems: - name of a model unit?

```
class pycropml.render_csharp.Model2Package (models, dir=None)
    Bases: object
```

TODO

```
DATATYPE = {'BOOLEAN': 'bool', 'DATE': 'string', 'DATELIST': 'List<string>', 'DOUBLE':
```

```
generate_test (model_unit)
```

```
num = 0
```

```
write_tests ()
```

TODO: Manage several models rather than just one.

```
pycropml.render_csharp.signature (model)
```

```
pycropml.render_csharp.transf (type_v, elem)
```

```
pycropml.render_csharp.transfDate (type, elem)
```

```
pycropml.render_csharp.transfDateList (type, elem)
```

```
pycropml.render_csharp.transfDouble (type_v, elem)
```

```
pycropml.render_csharp.transfList (type_v, elem)
pycropml.render_csharp.transfString (type_v, elem)
```

## **pycropml.render\_cyml module**

Add License, Header. Use pkglets Problems: - name of a model unit?

```
class pycropml.render_cyml.Model2Package (models, dir=None, pkg_name=None)
Bases: object
```

TODO

```
generate_algorithm (model_unit)
```

```
generate_component (model_unit)
```

Todo

```
generate_func_test (model_unit)
```

```
generate_function_doc (model_unit)
```

```
generate_function_signature (func_name, inputs)
```

```
generate_package ()
```

Generate a Cyml package equivalent to the xml definition. Args: - models : a list of model - dir: the directory where the code is generated. Returns: - None or status

```
generate_test (model_unit)
```

```
initialization (model_unit)
```

**num** = 0

```
run ()
```

TODO.

```
write_tests ()
```

TODO: Manage several models rather than just one.

```
pycropml.render_cyml.generate_doc (model)
```

```
pycropml.render_cyml.my_input (_input, defa=True)
```

```
pycropml.render_cyml.signature (model)
```

```
pycropml.render_cyml.transBool (type, elem)
```

```
pycropml.render_cyml.transf (type_, elem)
```

```
pycropml.render_cyml.transfDate (type, elem)
```

```
pycropml.render_cyml.transfDateList (type, elem)
```

## **pycropml.render\_fortran module**

Add License, Header.

Use pkglets

Problems: - name of a model unit?

---

```

class pycropml.render_fortran.Model2Package (models, directory=None, pkg_name=None)
    Bases: object

    TODO

    DATATYPE = {'BOOLEAN': 'LOGICAL', 'DATE': 'CHARACTER(65)', 'DATERLIST': 'CHARACTER(65),'
    generate_test (model_unit)
    my_input (_input)
    num = 0
    write_tests ()
        TODO: Manage several models rather than just one.

    pycropml.render_fortran.generate_doc (model)
    pycropml.render_fortran.signature (model)

```

## **pycropml.render\_java module**

Add License, Header.

Use pkglets

Problems: - name of a model unit?

```

class pycropml.render_java.Model2Package (models, dir=None)
    Bases: object

    DATATYPE = {'BOOLEAN': 'boolean', 'DATE': 'String', 'DATERLIST': 'Arrays.asList', 'DOUB'
    generate_test (model_unit)
    num = 0
    write_tests ()
        TODO: Manage several models rather than just one.

    pycropml.render_java.formatDate (elem)
    pycropml.render_java.formatDateList (elem)
    pycropml.render_java.signature (model)
    pycropml.render_java.transf (type_v, elem)
    pycropml.render_java.transfDate (categ, name, elem)
    pycropml.render_java.transfDateList (categ, name, elem)
    pycropml.render_java.transfDouble (type_v, elem)
    pycropml.render_java.transfList (type_v, elem)
    pycropml.render_java.transfString (type_v, elem)

```

## **pycropml.render\_notebook module**

License, Header

Use pkglets

Problems: - name of a model unit?

```
class pycropml.render_notebook.Model2Nb (models, dir=None)
Bases: pycropml.render_python.Model2Package

Generate a Jupyter Notebook from a set of models.

generate_notebook()
    Generate a Python package equivalent to the xml definition.

    Args: - models : a list of model - dir: the directory where the code is generated.

    Returns: - None or status

generate_test (model_unit)
run()
    TODO.
```

## **pycropml.render\_notebook\_csharp module**

License, Header

Use pkglts

Problems: - name of a model unit?

```
class pycropml.render_notebook_csharp.Model2Nb (models, dir=None)
Bases: pycropml.render_csharp.Model2Package

Generate a Jupyter Notebook from a set of models in Csharp.

generate_notebook()
    Generate a csharp package equivalent to the xml definition.

    Args: - models : a list of model - dir: the directory where the code is generated.

    Returns: - None or status

generate_test (model_unit)
run()
    TODO.
```

```
pycropml.render_notebook_csharp.signature (model)
pycropml.render_notebook_csharp.transf (type, elem)
pycropml.render_notebook_csharp.transfDate (type, elem)
pycropml.render_notebook_csharp.transfDateList (type, elem)
pycropml.render_notebook_csharp.transfDouble (type, elem)
pycropml.render_notebook_csharp.transfSDIList (type, elem)
pycropml.render_notebook_csharp.transfString (type, elem)
```

## **pycropml.render\_notebook\_java module**

License, Header

Use pkglts

Problems: - name of a model unit?

---

```
class pycropml.render_notebook_java.Model2Nb (models, dir=None)
Bases: pycropml.render\_java.Model2Package

Generate a Jupyter Notebook from a set of models in Java.

generate_notebook()
    Generate a java package equivalent to the xml definition.

    Args: - models : a list of model - dir: the directory where the code is generated.

    Returns: - None or status

generate_test (model_unit)
run()
    TODO.

pycropml.render_notebook_java.signature (model)
```

## pycropml.render\_python module

Add License, Header.

Use pkglts

Problems: - name of a model unit?

```
class pycropml.render_python.Model2Package (models, dir=None, pkg_name=None)
Bases: object

TODO

DATATYPE = {'BOOLEAN': <type 'bool'>, 'CHARLIST': <type 'list'>, 'DATE': <type 'str'>,
generate_algorithm (model_unit)
generate_component (model_unit)
    Todo

generate_factory (model)
    Create a Node Factory from CropML model unit.

generate_func_test (model_unit)
generate_function_doc (model_unit)
generate_function_signature (model_unit)
generate_package()
    Generate a Python package equivalent to the xml definition.

    Args: - models : a list of model - dir: the directory where the code is generated.

    Returns: - None or status

generate_test (model_unit)
generate_wralea()
    Generate wralea factories from the meta-information of the the model units.

num = 0

run()
    TODO.
```

```
    write_tests()  
        TODO: Manage several models rather than just one.  
  
pycropml.render_python.generate_doc(model)  
pycropml.render_python.openalea_interface(inout)  
pycropml.render_python.signature(model)
```

## **pycropml.test\_generator module**

### **pycropml.topology module**

### **pycropml.version module**

Maintain version for this package. Do not edit this file, use ‘version’ section of config.

```
pycropml.version.MAJOR = 1  
    (int) Version major component.  
  
pycropml.version.MINOR = 1  
    (int) Version minor component.  
  
pycropml.version.POST = 0  
    (int) Version post or bugfix component.
```

## **pycropml.wf2xml module**

### **pycropml.writeTest module**

Created on Mon Mar 18 15:46:31 2019

@author: midingoy

```
class pycropml.writeTest.WriteTest(models, language, dir)  
    Bases: object  
  
    write()  
        Populate and write the test files.
```

## **pycropml.writeTest\_f90 module**

Created on Thu Mar 28 15:39:28 2019

@author: midingoy

Add License, Header.

Use pkglets

Problems: - name of a model unit?

```
class pycropml.writeTest_f90.Model2Package(models, dir=None)  
    Bases: object  
  
    TODO  
  
    DATATYPE = {'BOOLEAN': 'LOGICAL:::', 'DATE': 'CHARACTER(65)', 'DATELIST': 'CHARACTER(65)
```

---

```

generate_algorithm(model_unit)
generate_component(model_unit)
    Todo
generate_estimation(model_unit)
generate_function_doc(model_unit)
generate_package()
    Generate a csharp package equivalent to the xml definition.

    Args: - models : a list of model - dir: the directory where the code is generated.

    Returns: - None or status

generate_public_class(model_unit)
generate_test(model_unit)

num = 0

run()
TODO.

write_tests()
TODO: Manage several models rather than just one.

pycropml.writeTest_f90.signature(model)
pycropml.writeTest_f90.transf(type, elem)
pycropml.writeTest_f90.transfDate(type, elem)
pycropml.writeTest_f90.transfDateList(type, elem)
pycropml.writeTest_f90.transfDouble(type, elem)
pycropml.writeTest_f90.transfSDIList(type, elem)
pycropml.writeTest_f90.transfString(type, elem)

```

## pycropml.xml2wf module

```

class pycropml.xml2wf.XmlToWf(xmlwf, dir, pkg_name)
Bases: object

compareInterface(interfaces)
compoPack(name)
compositeNodeInputs()
compositeNodeOutputsconnectInputs()
connectInternal()
connectOutputs()
createNodes()
retrievePackage(name)
run()

```

## Module contents

### 1.1.8 Usecases

### 1.1.9 Licence

PyCropML is released under a MIT License.

### 1.1.10 Usecases

### 1.1.11 Glossary

Terminology

**Model** Simplified representation of the crop system within specific objectives.

#### Overview

## 1.2 Documentation

- A PDF version of **lcorel** documentation is available.

## 1.3 History

### 1.3.1 creation (2018-01-18)

- First release on PyPI.

## 1.4 Indices and tables

## 1.5 History

### 1.5.1 creation (2018-01-18)

- First release on PyPI.

## 1.6 License

**pycrop2ml** is released under a MIT License.

## 1.7 Welcome to PyCrop2ML's documentation!

Contents:

### 1.7.1 Contributing Guide

This is a wiki for anything related to the contributing on [[Crop2ML|https://github.com/AgriculturalModelExchangeInitiative/Crop2ML]] which is a project of the Agricultural Model Exchange Initiative. For more information about this project, please visit CropML documentation [[Crop2ML|https://cropmlformat.readthedocs.io/en/latest/?badge=latest/documentation]]:

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### 1.7.2 Quick Links

- [[Project Git Repository|https://github.com/cython/cython|Git Repository]] (and [[Change Log|https://github.com/cython/cython/blob/master/CHANGES.rst|Change Log]])
- [[ Differences between Cython and Pyrex|https://cython.readthedocs.io/en/latest/src/userguide/pyrex\_differences.html|Differences between Cython and Pyrex]]
- [[Unsupported Python features|https://cython.readthedocs.io/en/latest/src/userguide/limitations.html|Unsupported Python features]] (aka TODO list)
- [[ Hacker-Guide: How to work on the Cython compiler itself|HackerGuide|Hacker-Guide: How to work on the Cython compiler itself]]
- [[ Enhancement proposals|enhancements| Enhancement proposals]] (CEPs)
- [[ Projects using Cython|projects| Projects using Cython]]
- [[ Comparison with SWIG|SWIG| Comparison with SWIG]]
- [[Automatic .pxd/.pyx generation|AutoPxd|Automatic .pxd/.pyx generation]] from C or C++ header files.

### Cython Installers

- [[PyPi|http://pypi.python.org/pypi/Cython|PyPi]] via `easy_install` or `pip`
- [[Gentoo Ebuild|http://packages.gentoo.org/package/dev-python/cython|Gentoo Ebuild]]
- [[Debian package|http://packages.debian.org/sid/cython|Debian package]] (not always up to date)
- [[ Installing Cython on Windows|InstallingOnWindows| Installing Cython on Windows]]

### Tips and Tricks

- [[Getting started|http://docs.cython.org/src/quickstart/index.html|Getting started]]
- [[ Using early binding techniques to improve speed|http://docs.cython.org/en/latest/src/userguide/early\_binding\_for\_speed.html|Using early binding techniques to improve speed]]
- [[ Writing Cython programs in pure Python|http://docs.cython.org/src/tutorial/pure.html| Writing Cython programs in pure Python]]
- [[ Helpful notes for wrapping C++ APIs|http://docs.cython.org/en/latest/src/userguide/wrapping\_CPlusPlus.html| Helpful notes for wrapping C++ APIs]]

- [[ Discussion of all the options how to wrap C/C++ code to Python|WrappingCorCpl] Discussion of all the options how to wrap C/C++ code to Python]]
- [[WritingFastPyrexCode|http://www.sagemath.org:9001/WritingFastPyrexCode|WritingFastPyrexCode]]
- [[ Successful creation of a hierarchy of modules in a package|PackageHierarchy] Successful creation of a hierarchy of modules in a package]]
- [[ One method for source-level debugging|http://docs.cython.org/en/latest/src/userguide/debugging.html] One method for source-level debugging]]
- [[ Dynamic Memory Allocation (malloc, realloc, free)|http://docs.cython.org/en/latest/src/tutorial/memory\_allocation.html] Dynamic Memory Allocation (malloc, realloc, free)]]
- [[Profiling]]
- [[ Building a Windows Installer|BuildingWindowsInstaller] Building a Windows Installer]]
- [[Embedding Python|EmbeddingCython|Embedding Python]] to create standalone Cython programs.
- [[List Subclass Example|ListExample|List Subclass Example]] Adding mathematical operations to subclassed built-in list.
- Working with Numpy
  - [[Tutorial for NumPy users|http://docs.cython.org/en/latest/src/userguide(numpy\_tutorial.html|Tutorial for NumPy users]]]
  - [[Accessing a Numpy pointer for passing to C|http://docs.cython.org/en/latest/src/userguide/memoryviews.html#pass-data-from-a-c-function-via-pointer]]

### 1.7.3 People

[[Stefan Behnel|http://scoder.behnel.de/|Stefan Behnel]], [[Robert Bradshaw|http://www.math.washington.edu/~robertwb/|Robert Bradshaw]], [[Dag Seljebotn|http://heim.ifi.uio.no/dags/|Dag Seljebotn]], Lisandro Dalcin.

### 1.7.4 Mailing Lists

Our development mailing list is [[cython-dev|http://mail.python.org/mailman/listinfo/cython-devel|cython-dev]] and user mailing list at <http://groups.google.com/group/cython-users>.

In the past we also used a [[Google group|http://groups.google.com/group/cython|Google group]] and a list at [[BerliOS Developer|https://lists.berlios.de/mailman/listinfo/cython-dev|BerliOS Developer]]. You can still read [[the archives at Gmane|http://blog.gmane.org/gmane.comp.python.cython.devel|the archives at Gmane]].

### 1.7.5 Project Goals

- Fully supported easy-to-use test suite, including the normal CPython test suite.
- Easy installation and usage.
- Rich, accessible documentation. Make sure the examples are plenty and can be automatically tested.
- Make Cython part of the standard distribution of Python (like ctypes).
- Compile all Python code except for possibly some obvious exclusions, which will be worked out by developers.
- Very fast when the user explicitly declares types (but we're not going to make promises with type inference). Precise benchmarks.

- Mitigate or eliminate the need for users to invoke the Python/C API directly without sacrificing performance.

### 1.7.6 Documentation

- See <http://docs.cython.org/>.
- Official Pyrex [[Language Overview|<http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/LanguageOverview>]] (note the [[changes|<http://hg.cython.org/cython/changes>]] though).
- [[Extension Types|[http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/Manual/extension\\_types.html](http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/Manual/extension_types.html)]]
- [[Sharing Declarations Between Pyrex Modules|[http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/Manual/sharing\\_declarations.html](http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/Manual/sharing_declarations.html)]]
- [[FAQ|<http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/FAQ.html>]]
- [[Quick Guide to Pyrex|<http://ldots.org/pyrex-guide/>] Quick Guide to Pyrex]] from Michael Jason-Smith.
- CategoryCythonDoc lists pages that are related to Cython documentation.
- [[ Pure Python mode|pure Python mode]]
- SAGE Days 4 talk highlighting some of the [[differences between Pyrex and SageX|<http://cython.org/talks/SageX.pdf>]] (the predecessor of Cython).

[CategoryHomepage](#)

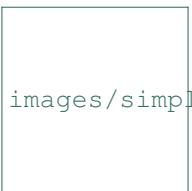
## 1.8 Indices and tables

## 1.9 Supported by:





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