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# **pycropml Documentation**

***Release 0.1.1***

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## Contents

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<b>1 CropML documentation</b>	<b>3</b>
<b>Python Module Index</b>	<b>57</b>
<b>Index</b>	<b>59</b>



**Contents .. \_pycropml:**



# CHAPTER 1

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## CropML 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 CropML Description

In CropML, a model is either a model unit or a composition of models. A ModelUnit represents the atomic unit of a crop model define by the modelers. A model composition is a model resulting from the composition of two or more atomic model or composite models.

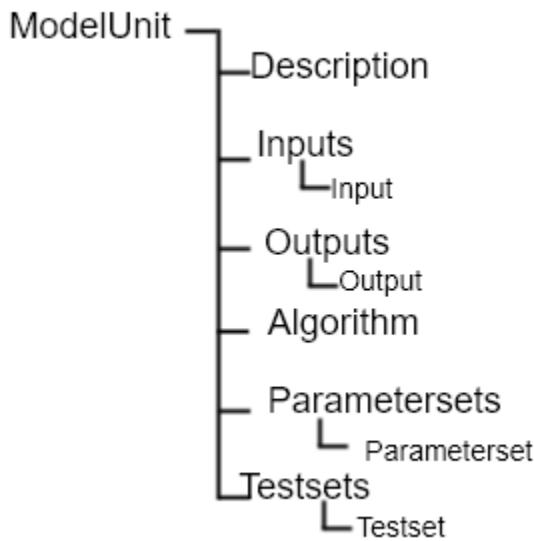
These models have a specific formal definition in CropML.

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

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

So a Model Unit CropML 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:



Element	Description
ModelUnit	The root of an atomic model in CropML which make the difference from a composite model.
Description	some basic information related to the name of the model, its authors and others elements used to reference it.
Inputs	A list of inputs characterized by their names, initial states, the range of values and others. Its input variables are related to climate, soil and cropping system
Outputs	A list of outputs defining the processes involved, the variables whose dynamics we want to observe.
Algorithm	The description of the behaviour of the model made by the mathematical relationship between the inputs and the outputs with some control structure.
Parametersets	Some sets of parameters which are invariant and used for the simulation of the models.
Testsets	Set of model configuration used to compare estimated and desired outputs .

In the next, we define the major elements of a CropML model unit.

## ModelUnit element

An atomic model in CropML is declared with `<>ModelUnit><` element, the usual root of CropML ModelUnit document.

```
<?xml version="1.0" encoding="utf-8"?>
<!DOCTYPE ModelUnit PUBLIC "-//SIMPLACE/DTD SOL 1.0//EN" "https://raw.githubusercontent.com/AgriculturalModelExchangeInitiative/xml_representation/master/ModelUnit.dtd">
<ModelUnit modelid=" " timestep=" " name=" " version="">
    ...
</ModelUnit>
```

This element MUST contain a Description, an Algorithm, Parametersets and Testsets elements and MAY optionally have Inputs and Outputs elements. The restriction of the length of different lists is not imposed.

ModelUnit element MUST have an modelid and name attributes which are used to reference an atomic model. It MUST also contain a timestep attribute to define the temporality of the model and a version attribute for each version of the model.

## Description element

This element gives the general information on the model and is composed by a set of character elements. It MUST contain Title, Authors, Institution and abstract elements and MAY optionally contain URI and Reference elements.

```
<ModelUnit modelid=" " timestep=" " name=" " version =" ">
    <Description>
        <Title>title</Title>
        <Authors>authors</Authors>
        <Institution>institution</Institution>
        <URI>uri</URI>
        <Abstract><![CDATA[abstract]]></Abstract>
    </Description>
    ...
</ModelUnit>
```

## Inputs elements

The inputs of Model are listed inside an XML element called Inputs within a [dictionary structure](#) composed by their attributes which declarations are optional(default, max, min, parametercategory, variablecategory and uri) or required(name, datatype, description, inputtype, unit ) and their corresponding value. *Inputs* element MUST contain one or more *Input* elements.

```
<ModelUnit modelid=" " timestep=" " name=" " version =" ">
    ...
    <Inputs>
        <Input name=" " description=" " parametercategory=" " datatype=" " min=" " max=" "
            default=" " unit=" " uri="" inputtype=" "/>
```

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```

<Input name="" description="" parametercategory="" datatype="" min="" max=
↪ " " default="" unit="" uri="" inputtype="" />
...
</Inputs>
...
</ModelUnit>
```

- The required *datatype* attribute is the type of input value specified in *default* (the default value in the input), *min* (the minimum value in the input) and *max* (the maximum value in the input). It MAY be one type of the set of types used in the existing crop modeling platform.
- The *inputtype* attribute makes it possible to distinguish the variables and the parameters of the model. So it MUST take one of two possible values: *parameter* and *variable*.
- The *parametercategory* attribute defines the category of parameter which is specified by one of the following values: *constant*, *species*, *soil* and *genotypic*.
- The *variablecategory* defines the category of variable depending on whether it is a *state*, a *rate* or an “auxiliary” variable. State variable characterize the behavior of the model and rate variable characterizes the changes in state variables.

## Outputs element

The outputs of Model are listed inside an XML element called Outputs within a dictionary structure composed by their attributes which declarations are:

- optional(variabletype and URI)
- required(name, datatype, description, unit, max and min )
- and their corresponding value

*Outputs* MUST contain zero or more output elements.

```

<ModelUnit modelid="" timestep="" name="" version =" ">
...
<Outputs>
    <Output name="" description="" datatype="" min="" max="" unit="" uri="" />
    <Output name="" description="" datatype="" min="" max="" unit="" uri="" />
    ...
</Outputs>
...
</ModelUnit>
```

The definition of different attributes is same as Input’s attributes.

## Algorithm element

The *Algorithm* element defines the building block of CropML model unit and shows the computational method to determine the outputs from the inputs.

It consists of a set of mathematical equations (relation between inputs), loops and conditional instructions which are well structured in a specific *language*, the algorithm’s attribute.

```
<ModelUnit modelid=" " timestep=" " name=" " version =" ">
...
<Algorithm language =""><! [CDATA[
...
]]>
</Algorithm>
...
</ModelUnit>
```

## Parametersets element

*Parametersets* element contains one or more *Parameterset* elements that define the different ways of setting the model. Each *Parameterset* element MUST have *name* and *description* attributes that respectively represents the name and the description of each setting.

The different parameterset MUST contain a list of Param elements that show in attribute the name of the parameter (an input which inputtype equals *parameter*) and the fixed value of this one.

```
<ModelUnit modelid=" " timestep=" " name=" " version =" ">
...
<Parametersets>
  <Parameterset name="" description="" uri = ""/>
  <Parameterset name="" description="" >
    <Param name="">value</Param>
    <Param name="">value</Param>
    ...
  </Parameterset>
  ...
  ...
</ModelUnit>
```

## Testsets element

*Testsets* element contains one or more *Testset* elements that define the different run for evaluating the outputs of the model.

Each *Testset* element MUST have *name*, *description* and *parameterset* attributes that respectively represents the name, the description of each run and the name of the parameterset related to the Testset. This one allow to retrieve the name and the value of different parameters includes in this parameterset.

The different Testset MUST contain a list of InputValue and OutputValue elements corresponding respectively to the values of inputs used in the run and the values of Outputs that will be asserted.

```
<ModelUnit modelid=" " timestep=" " name=" " version =" ">
...
<Testsets>
```

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```

<Testset name="" parameterset = "" description="" uri = ""/>
<Testset name="" parameterset = "" description="" >
  <Test name="">
    <InputValue name="">value</InputValue>
    ...
    <OutputValue name="" precision ="">value</OutputValue>
    ...
  </Test>
  ...
</Testset>
...
</Testsets>
...
</ModelUnit>

```

## Formal definition of a Composite Model in CropML

A Composite Model CropML 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 CropML MUST conform to a specific Document Type Definition named [ModelComposition.dtd](#).

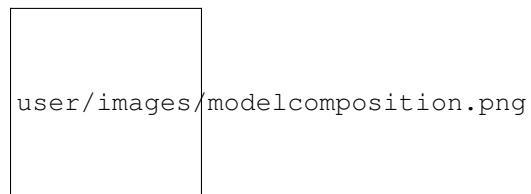
The composition is represented as a directed port graph of models:

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:



In the next, we define the major elements of a CropML model unit.

## Inputs element

It MUST contain one or more *input* element which provide a set of independent models entries. If two or more input variables of independent models are the same (same unit, interval, description) a link should be made to one input variable of the composite model.

## Outputs element

It MUST contain one or more *output* element which provide a set of independent models outputs or a result of a combination of models .

## Composition element

It's a list of *models* elements which contains a list of *links* elements. Link provides the mechanism for mapping inputs declared within one modelUnit to output in another modelUnit, allowing information to be exchanged between the various atomic models in the composite model.

## Algorithm element

The implementation differs from the platform:

- Discrete Events Models and Formalisms (RECORD)
- Actor model framework (OpenAlea)
- A sequence of algorithmic instructions which implement the control flow (BIOMA)

## 1.1.4 PyCropML User Guide

**Version** 0.1.1

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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

### 1.1.5 src

#### pycropml package

##### Subpackages

##### pycropml.interface package

###### Submodules

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

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

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

```
pycropml.interface.version.MAJOR = 0  
(int) Version major component.
```

```
pycropml.interface.version.MINOR = 0  
(int) Version minor component.
```

```
pycropml.interface.version.POST = 2  
(int) Version post or bugfix component.
```

##### 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)
```

## pycroml.transpiler.generators.csharpGenerator module

```
class pycroml.transpiler.generators.csharpGenerator.CsharpCompo(tree=None,
                                                               model=None,
                                                               name=None)
Bases: pycroml.transpiler.generators.csharpGenerator.CsharpTrans, pycroml.
transpiler.generators.csharpGenerator.CsharpGenerator
```

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

```
assignParam()
constrWrap()
copyconstrWrap()
copyconstructor(node)
estimateWrap()
format()
get_mo(varname)
initCompo()
initWrap()
instanceModels()
loadParamWrap()
outputWrap()
privateWrap()
```

```
    setCompo(p)
    tranAssignParam()
    visit_assignment(node)
    visit_declaration(node)
    visit_function_definition(node)
    visit_local(node)
    visit_module(node)
    visit_return(node)
    wrapper()

class pycropml.transpiler.generators.csharpGenerator.CsharpGenerator(tree,
                                                               model=None,
                                                               name=None)
Bases: pycropml.transpiler.codeGenerator.CodeGenerator, pycropml.transpiler.rules.csharpRules.CsharpRules
```

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

```
    add_features(node)
    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_continuestatnode(node)
    visit_custom_call(node)
        TODO
    visit_datetime(node)
    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)
    TODO
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.csharpGenerator.CsharpTrans (models)
Bases: pycropml.transpiler.codeGenerator.CodeGenerator, pycropml.transpiler.rules.csharpRules.CsharpRules

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

DATATYPE = {'BOOLEAN': 'bool', 'DATE': 'DateTime', 'DATEARRAY': ['array', 'DateTime'],
copyconstructor (node)
generate (nodes, typ)
getset (node, wrap=False)
model2Node ()
private (node)
visit_DateTime (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.csharpGenerator.to_struct_cs (models, rep, name)
pycropml.transpiler.generators.csharpGenerator.to_wrapper_cs (models, rep, name)
```

## 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 - descrrition of the code - Inputs details - Outputs details

comment (doc)
doc (x, name)
generate_desc (model)
generate_header (model)
```

## pycroml.transpiler.generators.fortranGenerator module

```
class pycroml.transpiler.generators.fortranGenerator.FortranCompo (tree=None,  

model=None,  

name=None)  
Bases: pycroml.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 pycroml.transpiler.generators.fortranGenerator.FortranGenerator (tree,  

model=None,  

name=None)  
Bases: pycroml.transpiler.codeGenerator.CodeGenerator, pycroml.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)  
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_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_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_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_tab(node)
visit_unary_op(node)
visit_while_statement(node)

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

## pycroml.transpiler.generators.javaGenerator module

```
class pycroml.transpiler.generators.javaGenerator.JavaCompo(tree, model=None,
                                                               name=None)
Bases: pycroml.transpiler.generators.javaGenerator.JavaTrans, pycroml.
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 pycroml.transpiler.generators.javaGenerator.JavaGenerator(tree,
                                                               model=None,
                                                               name=None)
Bases: pycroml.transpiler.codeGenerator.CodeGenerator, pycroml.transpiler.
rules.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_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.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)
```

## 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

```
class pycropml.transpiler.generators.siriusGenerator.SiriusCompo(tree=None,
    model=None,
    name=None)
Bases: pycropml.transpiler.generators.csharpGenerator.CsharpCompo

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

constrWrap()
copyconstrWrap()
visit_module(node)
wrapper()
```

```
class pycropml.transpiler.generators.siriusGenerator.SiriusGenerator(tree=None,
model=None,
name=None)
Bases: pycropml.transpiler.generators.csharpGenerator.CsharpGenerator
```

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

```
class pycropml.transpiler.generators.siriusGenerator.SiriusTrans(models)
Bases: pycropml.transpiler.generators.csharpGenerator.CsharpTrans
```

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

```
pycropml.transpiler.generators.siriusGenerator.to_struct_sirius(models, rep,
name)
```

```
pycropml.transpiler.generators.siriusGenerator.to_wrapper_sirius(models, rep,
name)
```

## Module contents

### pycropml.transpiler.lib package

#### Module contents

### pycropml.transpiler.rules package

#### Submodules

##### pycropml.transpiler.rules.csharpRules module

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

binary_op = {'!=': ' !=', '*': '*', '+': '+', '-': '-', '/': '/', '<': '<', '<=':
constructor = '\n public %s()\n {\n }\n '
copy_constr = '\n public %s(%s toCopy, bool copyAll) // copy constructor\n {\n if (copyAll)\n _%s = toCopy.%s;\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 = '\n public %s(%s toCopy): this() // copy constructor\n {\n _%s = toCopy.%s;\n }'
functions = {'datetime': {'datetime': ' new DateTime'}, 'io': {'print': 'Display.WriteLine'}}
methods = {'array': {'append': '.Add', 'len': '<function translateLenArray>'}, 'dict': {'clear': '.Clear'}, 'list': {'append': '.Add', 'len': '<function translateLenList>'}, 'set': {'add': '.Add', 'len': '<function translateLenSet>'}, 'tuple': {'len': '<function translateLenTuple>'}}
public_properties = '\n {\n get\n {\n return this._%s;\n }\n set\n {\n this._%s= value;\n }'
public_properties_compo = '\n {\n get\n {\n return _%s.%s;\n }\n set\n {\n %s.%s;\n }'
public_properties_wrap = '{ get { return %s.%s; } } \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.translateLenArray(node)
```

```
pycropml.transpiler.rules.csharpRules.translateLenDict(node)
pycropml.transpiler.rules.csharpRules.translateLenList(node)
pycropml.transpiler.rules.csharpRules.translateNotContains(node)
pycropml.transpiler.rules.csharpRules.translateSum(node)
pycropml.transpiler.rules.csharpRules.translateGet(node)
pycropml.transpiler.rules.csharpRules.translateKeyDict(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 FortranRules.<lambda>>}, 'math': {}}
    method()

    methods = {'array': {'append': <function FortranRules.<lambda>>, 'len': 'SIZE'}, '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.translateNotContains(node)
pycropml.transpiler.rules.fortranRules.translatePop(node)
pycropml.transpiler.rules.fortranRules.translatePow(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 = {'!=': '!=', '*': '*', '+': '+', '-': '-', '/': '/', '<': '<', '<=':
    constructor = '\n public %s()\n {\n \n }'

    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 {\n
    functions = {'datetime': {'datetime': 'format.parse'}, 'math': {'acos': 'Math.acos'}
    get_properties = '\n {\n return %s;\n }'
    get_properties_compo = '\n {\n return _%s.get%s();\n }'
    methods = {'array': {'append': '.add', 'len': <function translateLenArray>}, 'dict':
    set_properties = '\n {\n this.%s= _%s;\n } \n '
    set_properties_compo = '\n {\n %s\n } '
    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.trans_format_parse(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.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'}, 'math': {'acos': 'acos', 'asin':
    methods = {'array': {'len': 'len'}, 'datetime': {'datetime': 'datetime', 'day': 'day',
    types = {'bool': 'bool', 'datetime': 'datetime', 'dict': 'dict', 'float': 'float',
    unary_op = {'+': '+', '-': '-', 'not': 'not ', '~': '~'}

    pycropml.transpiler.rules.pythonRules.translateDictkeys(node)

```

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

## pycropml.transpiler.rules.sqRules module

```
class pycropml.transpiler.rules.sqRules.SqRules
    Bases: pycropml.transpiler.rules.generalRule.GeneralRule

    field_decl(node)
    header = '-----'
    method()
    methods = {'dict': {'len': 'SIZE'}, 'float': {'int': 'INT'}, 'int': {'float': 'R'}}
    namespace = {'headNamespace': '\nnamespace SiriusQualityEnergyBalance\n{\n    using System;
```

### Module contents

#### Submodules

## pycropml.transpiler.Parser module

```
class pycropml.transpiler.Parser.opt(**kwds)
    Bases: object
```

```
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
    Bases: object
```

Standard classes should respond to expand and to return valid nodes on expand

```
class pycropml.transpiler.api_transform.StandardCall(namespace, function, expand=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*, *default=None*, *expander=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.**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*)

Bases: *object*

**assert\_translatable**(*node*, *\*\*pairs*)

**newtype**()

**notdeclared**(*name*, *line*)

**retrieve\_library**(*func*)

**transformer**()

**visit\_addnode**(*node*, *operand1*, *operand2*, *location*)

**visit\_attributemode**(*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_condexprnode (node, test, true_val, false_val, location)
visit_continuestatnode (node, location)
visit_csimplebasetypenode (node, 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_unicodenode (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)
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
    Bases: object
```

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=(","))
    emit_string(node, prefix="")
    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_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)
    Bases: object

    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,
                                                               suggestions=None,
                                                               right=None,
                                                               wrong=None)
Bases: pycropml.transpiler.errors.PseudoError

exception pycropml.transpiler.errors.PseudoError(message,           suggestions=None,
                                                right=None, wrong=None)
Bases: 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)
Bases: object
    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

```
class pycropml.transpiler.main.Main(file, language, models=None, name=None)
Bases: object
    parse()
    to_ast(source)
```

```
    to_source()
    translate()
pycropml.transpiler.main.formater(code)
pycropml.transpiler.main.formaterNext(line)
```

## **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)
    Bases: object
```

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

License, Header  
Use pkglets  
Generate notebook from code source

```
class pycropml.code2nbk.Model2Nb(model, code, name, dir=None)
    Bases: object

Generate a Jupyter Notebook from a set of models.

generate_nb(language, tg_rep, namep)

nb = {'cells': [], 'metadata': {}, 'nbformat': 4, 'nbformat_minor': 4}
```

## 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**

Created on Tue Mar 19 22:59:23 2019

@author: midingoy

```
pycropml.cyml.ext(language)
pycropml.cyml.prefix(model)
pycropml.cyml.transpile_file(source, language)
pycropml.cyml.transpile_package(package, language)
```

## **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: 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

Created on Tue Mar 19 22:59:23 2019

@author: pradal

`pycropml.main.main()`

## 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 dictionary 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 wrlea.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 (key, default=None)
        Return the value for key if key is in the dictionary, else default.

    has_key (key)

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

    nb_public_values()
        Return the number of unprotected values

class pycropml.package.PackageManager
    Bases: object

    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 recursivly 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 pycropml.package.PseudoGroup (name)
    Bases: pycropml.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 pycroml.package.PyPackageReader(filename)
    Bases: pycroml.package.AbstractPackageReader
    Build packages from wralea file Use 'register_package' function

    build_package(wralemodule, 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 pycroml.package.PyPackageReaderModel(filename)
    Bases: pycroml.package.PyPackageReader
    Build a package from a __wralea__.py Use module variable

    build_package(wralemodule, pkgmanager)
        Build package and update pkgmanager

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

class pycroml.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 = '# This file has been generated at $TIME\n$PKG_DECLARATION\n'
    write_wralea(full_filename)
        Write the wralea.py in the specified filename

exception pycroml.package.UnknownNodeError(name)
    Bases: Exception

class pycroml.package.UserPackage(name, metainfo, path=None)
    Bases: pycroml.package.Package
    Package user editable and persistent

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

    pycroml.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>Ouputs (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 pkglets

Problems: - name of a model unit?

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

TODO

```
DATATYPE = {'BOOLEAN': <class 'bool'>, 'CHARLIST': <class 'list'>, 'DATE': <class 'str'>}
```

```
generate_algorithm(model_unit)
```

```
generate_component(model_unit)
```

Todo

```
generate_func_test(model_unit)
```

```
generate_function_doc(model_unit)
```

```
generate_function_signature(model_unit)
```

```
generate_package()
```

Generate a R 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)
```

```
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\_csharp module**

Add License, Header.

Use pkglets

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, model_unit)
```

```
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)', 'DATELIST': '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', 'DATELIST': 'ArrayList', 'DOUBLE':
    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 pkglets

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 ( )**

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

## pycrocml.render\_notebook\_java module

License, Header

Use pkglets

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.

```
create_notebook()
```

Generate a java package equivalent to the xml definition

Args: - models : a list of

#### **References** Nonparametric

**create\_test (model\_name)**

**TODO**

20m]

T. Fend

Use `pkglts`

## Problems:

## **class** pycropml.render pw

Bases: `object`  
TODO

DATA

**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.test_generator.generate_test_check (model, dir=None)
pycropml.test_generator.generate_test_cs (model, dir)
pycropml.test_generator.generate_test_f90 (model, dir)
pycropml.test_generator.generate_test_java (model, directory=None)
pycropml.test_generator.generate_test_openalea (model, dir=None)
pycropml.test_generator.generate_test_py (model, dir=None)
pycropml.test_generator.generate_test_simplace (model, dir=None)
pycropml.test_generator.generate_test_sirius (model, dir=None)
```

## **pycropml.topology module**

```
class pycropml.topology.Package (name, metainfo, path=None)
    Bases: object

class pycropml.topology.PackageManager (proj)
    Bases: object
```

```
check_exist()
contain_pkg()
get_path(pkg)

class pycrompi.topology.Topology(name, pkg=None)
Bases: object

algo2cyml()
algorithm()
check_compo(mc, m)
compotranslate(language)
createGraph()
create_edgeInOut()
decl(defa=True)
display_wf()
generate_function_signature(model)
get_mu_inp(pkgname, varname)
get_mu_out(pkgname, varname)
info_inputs_mu(ppkg, mu, varname)
info_minout()
info_outputs_mu(ppkg, mu, varname)
isPackage(name)
load_pkge(name)
meta_ext(pkgname)
meta_inp(pkgname)
meta_out(pkgname)
minout()
pkg_m(mc, m)
pkgs = {}
retrieve(pkgname)
topologicalSort()
translate()
translate_all(model)
val_init(model)
write_png()
write_xml()
```

## **pycropml.version module**

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

`pycropml.version.MAJOR = 0`

(int) Version major component.

`pycropml.version.MINOR = 1`

(int) Version minor component.

`pycropml.version.POST = 1`

(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 pkglts

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()
    compositeNodeOutputs()
    connectInputs()
    connectInternal()
    connectOutputs()
    createNodes()
    retrievePackage(name)
    run()
```

## Module contents

### 1.1.6 Usecases

### 1.1.7 Licence

PyCropML is released under a MIT License.

## 1.1.8 Usecases

## 1.1.9 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

PyCropML is released under a MIT License.

## 1.7 Welcome to CropML'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|WrappingCorCpp| 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/dagss|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-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|Extension Types]]
- [[Sharing Declarations Between Pyrex Modules|http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/Manual/sharing\_declarations.html|Sharing Declarations Between Pyrex Modules]]
- [[FAQ|http://www.cosc.canterbury.ac.nz/greg.ewing/python/Pyrex/version/Doc/FAQ.html|FAQ]]

- [[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|differences between Pyrex and SageX]] (the predecessor of Cython).
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CategoryHomepage

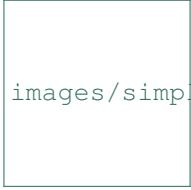
## 1.8 Indices and tables

### 1.9 Supported by:





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images/simpleplace.png

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## Python Module Index

---

### p

pycropml, 1  
pycropml.algorithm, 34  
pycropml.checking, 35  
pycropml.code2nbk, 35  
pycropml.composition, 35  
pycropml.cyml, 36  
pycropml.description, 36  
pycropml.error, 37  
pycropml.formater\_f90, 37  
pycropml.function, 37  
pycropml.initialization, 37  
pycropml.inout, 37  
pycropml.interface.version, 11  
pycropml.main, 38  
pycropml.package, 38  
pycropml.parameterset, 42  
pycropml.pparse, 42  
pycropml.render\_csharp, 44  
pycropml.render\_cyml, 44  
pycropml.render\_fortran, 45  
pycropml.render\_java, 45  
pycropml.render\_notebook, 46  
pycropml.render\_notebook\_csharp, 46  
pycropml.render\_notebook\_java, 47  
pycropml.render\_python, 47  
pycropml.render\_R, 43  
pycropml.test\_generator, 48  
pycropml.topology, 48  
pycropml.transpiler, 34  
pycropml.transpiler.api\_transform, 28  
pycropml.transpiler.ast\_transform, 29  
pycropml.transpiler.builtin\_typed\_api, 31  
pycropml.transpiler.checkingModel, 31  
pycropml.transpiler.codeGenerator, 32  
pycropml.transpiler.env, 32  
pycropml.transpiler.errors, 32  
pycropml.transpiler.generators, 25

pycropml.transpiler.generators.checkGenerator, 12  
pycropml.transpiler.generators.csharpGenerator, 13  
pycropml.transpiler.generators.docGenerator, 16  
pycropml.transpiler.generators.fortranGenerator, 17  
pycropml.transpiler.generators.javaGenerator, 19  
pycropml.transpiler.generators.openaleaGenerator, 22  
pycropml.transpiler.generators.pythonGenerator, 22  
pycropml.transpiler.generators.recordGenerator, 24  
pycropml.transpiler.generators.simplaceGenerator, 24  
pycropml.transpiler.generators.siriusGenerator, 24  
pycropml.transpiler.helpers, 33  
pycropml.transpiler.interface, 33  
pycropml.transpiler.lib, 25  
pycropml.transpiler.main, 33  
pycropml.transpiler.nodeVisitor, 34  
pycropml.transpiler.Parser, 28  
pycropml.transpiler.pseudo\_tree, 34  
pycropml.transpiler.rules, 28  
pycropml.transpiler.rules.csharpRules, 25  
pycropml.transpiler.rules.fortranRules, 26  
pycropml.transpiler.rules.generalRule, 26  
pycropml.transpiler.rules.javaRules, 27  
pycropml.transpiler.rules.pythonRules, 27  
pycropml.transpiler.rules.sqRules, 28  
pycropml.transpiler.version, 34  
pycropml.version, 50

`pycropml.writeTest`, 50  
`pycropml.writeTest_f90`, 50  
`pycropml.xml2wf`, 51

---

## Index

---

### A

abs\_expander() (in module `cropml.transpiler.api_transform`), 29  
AbstractPackageReader (class in `cropml.package`), 38  
access() (`pycropml.transpiler.generators.javaGenerator`.*JavaTrans* method), 21  
add() (in module `cropml.transpiler.builtin_typed_api`), 31  
add\_crop2ml\_path() (`cropml.package.PackageManager` method), 39  
add\_description() (`cropml.composition.ModelComposition` method), 35  
add\_description() (`cropml.modelunit.ModelUnit` method), 38  
add\_features() (`cropml.transpiler.generators.csharpGenerator`.*CsharpGenerator* method), 14  
add\_features() (`cropml.transpiler.generators.fortranGenerator`.*FortranGenerator* method), 17  
add\_features() (`cropml.transpiler.generators.javaGenerator`.*JavaGenerator* method), 19  
add\_modelunit() (`pycropml.package.Package` method), 38  
add\_name() (`pycropml.package.PseudoGroup` method), 40  
add\_package() (`pycropml.package.PackageManager` method), 39  
algo2cyml() (`pycropml.topology.Topology` method), 49  
Algorithm (class in `pycropml.algorithm`), 34  
Algorithm() (`pycropml.pparse.ModelParser` method), 42  
algorithm() (`pycropml.topology.Topology` method), 49

and\_() (in module `cropml.transpiler.builtin_typed_api`), 31  
api\_translate() (`cropml.transpiler.interface.middleware` method), 33  
APICheck() (in module `cropml.transpiler.builtin_typed_api`), 31  
argsToStr() (in module `cropml.transpiler.rules.fortranRules`), 26  
argsToStr() (in module `cropml.transpiler.rules.javaRules`), 27  
assert\_translatable() (`cropml.transpiler.ast_transform.AstTransformer` method), 29  
assignParam() (`cropml.transpiler.generators.csharpGenerator`.*CsharpCompo* method), 13  
AstTransformer (class in `cropml.transpiler.ast_transform`), 29

**B**

beautiful\_error() (in module `cropml.transpiler.errors`), 33  
binary\_and() (in module `cropml.transpiler.builtin_typed_api`), 31  
binary\_op (`pycropml.transpiler.rules.csharpRules`.*CsharpRules* attribute), 25  
binary\_op (`pycropml.transpiler.rules.fortranRules`.*FortranRules* attribute), 26  
binary\_op (`pycropml.transpiler.rules.javaRules`.*JavaRules* attribute), 27  
binary\_op (`pycropml.transpiler.rules.pythonRules`.*PythonRules* attribute), 27  
binary\_or() (in module `cropml.transpiler.builtin_typed_api`), 31  
binop\_precedence (`cropml.transpiler.codeGenerator`.*CodeGenerator* attribute), 32  
binop\_precedence (`cropml.transpiler.generators.fortranGenerator`.*FortranGenerator*

```

    attribute), 17
body () (pycropml.transpiler.codeGenerator.CodeGenerator      compareInterface () (pycropml.xml2wf.XmlToWf
    method), 32                                     method), 51
body () (pycropml.transpiler.generators.fortranGenerator.FortranGeneratorInputs () (py-
    method), 17                                         cropml.xml2wf.XmlToWf method), 51
body_or_else () (py-                                         compositeNodeOutputs () (py-
    cropml.transpiler.codeGenerator.CodeGenerator           cropml.xml2wf.XmlToWf method), 51
    method), 32
build_package () (py-                                         Composition () (pycropml.composition.ModelParser
    cropml.package.PyPackageReader     method),       method), 36
    41
build_package () (py-                                         compotranslate () (pycropml.topology.Topology
    cropml.package.PyPackageReaderModel   method),       method), 49
builtin_type_check () (in module py-                                         connectInputs () (pycropml.xml2wf.XmlToWf
    cropml.transpiler.builtin_typed_api), 31           method), 51
                                                        connectInternal () (pycropml.xml2wf.XmlToWf
                                                        method), 51
                                                        connectOutputs () (pycropml.xml2wf.XmlToWf
                                                        method), 51
constructor (pycropml.transpiler.rules.csharpRules.CsharpRules
    attribute), 25
constructor (pycropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
constrWrap () (pycropml.transpiler.generators.csharpGenerator.Csharp
    method), 13
constrWrap () (pycropml.transpiler.generators.siriusGenerator.SiriusCo
    method), 24
contain_pkg () (py-                                         copy_constr (pycropml.transpiler.rules.csharpRules.CsharpRules
    cropml.package.PyPackageReaderModel           attribute), 25
    method), 41
contain_pkg () (pycropml.topology.PackageManager
    method), 49
copy_constr (pycropml.transpiler.rules.csharpRules.CsharpRules
    attribute), 25
copy_constr (pycropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
copy_constr_compo (py-                                         copy_constr_compo (py-
    cropml.transpiler.rules.csharpRules.CsharpRules
    attribute), 25
copy_constr_compo (py-                                         copy_constr_compo (py-
    cropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
copy_constrArray (py-                                         copy_constrArray (py-
    cropml.transpiler.rules.csharpRules.CsharpRules
    attribute), 25
copy_constrArray (py-                                         copy_constrArray (py-
    cropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
copy_constrList (py-                                         copy_constrList (py-
    cropml.transpiler.rules.csharpRules.CsharpRules
    attribute), 25
copy_constrList (py-                                         copy_constrList (py-
    cropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
copy_compoFactor () (py-                                         copy_compoFactor () (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator
    method), 22                                         cropml.transpiler.generators.csharpGenerator.CsharpCompo
                                                        attribute), 27

```

*method), 13*

*copyconstructor () (py-  
cropml.transpiler.generators.csharpGenerator.CsharpTrans(pycroml.topology.Topology method), 49  
method), 16*

*copyconstructor () (py-  
cropml.transpiler.generators.javaGenerator.JavaComposition() (pycroml.composition.ModelParser  
method), 19*

*copyconstructor () (py-  
cropml.transpiler.generators.javaGenerator.JavaTrans  
method), 21*

*copyconstrWrap () (py-  
cropml.transpiler.generators.csharpGenerator.CsharpCompo  
method), 13*

*copyconstrWrap () (py-  
cropml.transpiler.generators.siriusGenerator.SiriusCompo  
method), 24*

*create\_edgeInOut () (pycroml.topology.Topology  
method), 49*

*create\_readers () (py-  
cropml.package.PackageManager  
method), 39*

*createGraph () (pycroml.topology.Topology  
method), 49*

*createNodes () (pycroml.xml2wf.XmlToWf method),  
51*

*CsharpCompo (class in py-  
cropml.transpiler.generators.csharpGenerator),  
13*

*CsharpGenerator (class in py-  
cropml.transpiler.generators.csharpGenerator),  
14*

*CsharpRules (class in py-  
cropml.transpiler.rules.csharpRules), 25*

*CsharpTrans (class in py-  
cropml.transpiler.generators.csharpGenerator),  
15*

**D**

*DATATYPE (pycroml.render\_csharp.Model2Package  
attribute), 44*

*DATATYPE (pycroml.render\_fortran.Model2Package  
attribute), 45*

*DATATYPE (pycroml.render\_java.Model2Package at-  
tribute), 45*

*DATATYPE (pycroml.render\_python.Model2Package  
attribute), 47*

*DATATYPE (pycroml.render\_R.Model2Package at-  
tribute), 43*

*DATATYPE (pycroml.transpiler.generators.csharpGenerator.CsharpTransmodule pycroml.cyml), 36  
attribute), 16*

*DATATYPE (pycroml.transpiler.generators.javaGenerator.JavaTrans  
attribute), 21*

*DATATYPE (pycroml.writeTest\_f90.Model2Package at-  
tribute), 50*

*datetime\_expander () (in module py-  
cropml.transpiler.api\_transform), 29*

*deepTrans(pycroml.topology.Topology method), 49*

*Description (class in pycroml.composition), 35*

*Description (class in pycroml.description), 36*

*Composition () (pycroml.composition.ModelParser  
method), 36*

*Description () (pycroml.pparse.ModelParser  
method), 42*

*dispatch () (pycroml.composition.ModelParser  
method), 36*

*dispatch () (pycroml.pparse.ModelParser method),  
42*

*dispatch () (pycroml.pparse.Parser method), 43*

*display\_wf () (pycroml.topology.Topology method),  
49*

*div () (in module py-  
cropml.transpiler.builtin\_typed\_api), 31*

*doc (pycroml.transpiler.generators.fortranGenerator.FortranGenerator  
attribute), 17*

*doc () (pycroml.transpiler.generators.docGenerator.DocGenerator  
method), 16*

*DocGenerator (class in py-  
cropml.transpiler.generators.docGenerator),  
16*

**E**

*emit\_sequence () (py-  
cropml.transpiler.codeGenerator.CodeGenerator  
method), 32*

*emit\_string () (py-  
cropml.transpiler.codeGenerator.CodeGenerator  
method), 32*

*Env (class in pycroml.transpiler.env), 32*

*Error, 37*

*estimateWrap () (py-  
cropml.transpiler.generators.csharpGenerator.CsharpCompo  
method), 13*

*expand () (pycroml.transpiler.api\_transform.StandardCall  
method), 29*

*expand () (pycroml.transpiler.api\_transform.StandardCallAttrib  
method), 29*

*expand () (pycroml.transpiler.api\_transform.StandardMethodCall  
method), 29*

*expand () (pycroml.transpiler.api\_transform.StandardSwapper  
method), 29*

*field\_decl () (pycroml.transpiler.rules.sqRules.SqRules  
method), 28*

**F**

```

filename_to_module()           (py-          method), 47
    cropml.package.PyPackageReader   method), generate_algorithm()           (py-
        41                           cropml.render_R.Model2Package   method),
find_and_register_packages()   (py-          43
    cropml.package.PackageManager   method), generate_algorithm()           (py-
        39                           cropml.writeTest_f90.Model2Package method),
find_crop2ml_dir()            (py-          50
    cropml.package.PackageManager   method), generate_component()           (py-
        40                           cropml.render_cyml.Model2Package method),
float_expander()              (in      module     py-          44
    cropml.transpiler.api_transform), 29 generate_component()           (py-
format() (pycropml.transpiler.generators.csharpGenerator.CsharpCroppd.render_python.Model2Package
    method), 13                         method), 47
formatDate() (in module pycropml.render_java), 46 generate_component()           (py-
formatDateList() (in module     py-          43
    cropml.render_java), 46                           cropml.render_R.Model2Package   method),
formater() (in module pycropml.formater_f90), 37 generate_component()           (py-
formater() (in module pycropml.transpiler.main), 34 cropml.writeTest_f90.Model2Package method),
formaterNext() (in module pycropml.formater_f90),
    37                                     50
generate_desc()                 (py-          generate_desc()           (py-
    cropml.transpiler.generators.docGenerator.DocGenerator
method), 16                           method), 16
generate_doc() (in module pycropml.render_cyml),
    45                                     generate_doc()           (in      module     py-
    cropml.render_fortran), 45
generate_doc() (in module     py-          48
    cropml.render_python), 48
generate_doc() (in module pycropml.render_R), 43 generate_doc()           (py-
generate_estimation()           (py-          50
    cropml.writeTest_f90.Model2Package method),
generate_factory()             (py-          generate_factory()           (py-
    cropml.render_python.Model2Package
method), 48                           method), 48
generate_func_factory()         (py-          generate_func_factory()           (py-
    cropml.transpiler.rules.fortranRules.FortranRules
attribute), 26                           cropml.transpiler.generators.openaleaGenerator.OpenaleaCompo
method), 22
functions(pycropml.transpiler.rules.javaRules.JavaRules
    attribute), 27                                     generate_func_test()           (py-
functions(pycropml.transpiler.rules.pythonRules.PythonRules
    attribute), 27                           cropml.render_cyml.Model2Package   method),
generate() (pycropml.transpiler.generators.csharpGenerator.CsharpCroppd.render_R.Model2Package
    method), 16                                     44
generate() (pycropml.transpiler.generators.javaGenerator.JavaTrans
    method), 21                                     generate_func_test()           (py-
generate_algorithm()           (py-          44
    cropml.render_cyml.Model2Package   method),
generate_algorithm()           (py-          44
    cropml.render_python.Model2Package
method), 44                                     generate_function_doc()           (py-
generate_function_doc()           (py-          48
    cropml.render_python.Model2Package
method), 48                                     generate_function_doc()           (py-
generate_function_doc()           (py-          48
    cropml.render_python.Model2Package
method), 48

```

## G

```

GeneralRule      (class      in      py-          generate_func_test()           (py-
    cropml.transpiler.rules.generalRule), 26                           cropml.render_python.Model2Package
method), 48
generate() (pycropml.transpiler.generators.csharpGenerator.CsharpCroppd.render_R.Model2Package
    method), 16                                     generate_func_test()           (py-
generate() (pycropml.transpiler.generators.javaGenerator.JavaTrans
    method), 21                           cropml.render_cyml.Model2Package   method),
generate_algorithm()           (py-          44
    cropml.render_cyml.Model2Package   method),
generate_algorithm()           (py-          44
    cropml.render_python.Model2Package
method), 44                                     generate_function_doc()           (py-
generate_function_doc()           (py-          48
    cropml.render_python.Model2Package
method), 48                                     generate_function_doc()           (py-
generate_function_doc()           (py-          48
    cropml.render_python.Model2Package
method), 48

```

```

    cropml.render_R.Model2Package      method), generate_test()          (py-
43                                         (py-  cropml.render_java.Model2Package  method),
generate_function_doc()           (py- 45                                         (py-
    cropml.writeTest_f90.Model2Package method), generate_test()          (py-
50                                         (py-  cropml.render_notebook.Model2Nb   method),
generate_function_signature()     (py- 46                                         (py-
    cropml.render_cyml.Model2Package method), generate_test()          (py-
44                                         (py-  cropml.render_notebook_csharp.Model2Nb
generate_function_signature()     (py-  method), 47                                         (py-
    cropml.render_python.Model2Package method), generate_test()          (py-
48                                         (py-  cropml.render_notebook_java.Model2Nb
generate_function_signature()     (py-  method), 47                                         (py-
    cropml.render_R.Model2Package   method), generate_test()          (py-
43                                         (py-  cropml.render_python.Model2Package
generate_function_signature()     (py-  method), 48                                         (py-
    cropml.topology.Topology method), 49 generate_test()          (py-
generate_header()               (py-  cropml.render_R.Model2Package   method),
method), 16                                         (py-  generate_test()          (py-
generate_nb()                   (py-  cropml.writeTest_f90.Model2Package method),
method), 35                                         (py- 51                                         (py-
generate_notebook()             (py-  generate_test_check()  (in  module  py-
    cropml.render_notebook.Model2Nb  method), 46                                         (py-
generate_notebook()             (py-  generate_test_cs()    (in  module  py-
    cropml.render_notebook_csharp.Model2Nb
method), 46                                         (py-  generate_test_f90()   (in  module  py-
generate_notebook()             (py-  method), 47                                         (py-
    cropml.render_notebook_java.Model2Nb
method), 47                                         (py-  generate_test_java()  (in  module  py-
generate_package()              (py-  method), 48                                         (py-
    cropml.render_cyml.Model2Package method), 44                                         (py-
generate_package()              (py-  generate_test_openalea() (in  module  py-
    cropml.render_python.Model2Package method), 48                                         (py-
generate_package()              (py-  generate_test_py()    (in  module  py-
    cropml.render_R.Model2Package   method), 43                                         (py-
generate_package()              (py-  generate_test_simplace() (in  module  py-
    cropml.writeTest_f90.Model2Package method), 50                                         (py-
generate_public_class()         (py-  generate_test_sirius()  (in  module  py-
    cropml.writeTest_f90.Model2Package method), 50                                         (py-
generate_test()                 (py-  generate_wralea()    (py-
    cropml.render_csharp.Model2Package method), 44                                         (py-
generate_test()                 (py-  cropml.render_python.Model2Package
    cropml.render_cyml.Model2Package method), 44                                         (py-
generate_test()                 (py-  method), 48                                         (py-
    cropml.render_fortran.Model2Package method), 45                                         (py-
get() (pycrocml.package.PackageDict method), 39
get() (pycrocml.package.PackageManager method), 40
get_crop2ml_path() (pycrocml.package.Package method), 38
get_default_home_dir() (in  module  py-
    cropml.package), 41
get_id() (pycrocml.package.Package method), 38
get_id() (pycrocml.package.PseudoGroup method),
40
get_metainfo() (pycrocml.package.Package method), 38

```

```

get_mo() (pycrocml.transpiler.generators.csharpGenerator.CsharpCompo
    method), 13
get_mo() (pycrocml.transpiler.generators.javaGenerator.JavaCompo
    method), 19
get_modelunit() (pycrocml.package.Package
    method), 38
get_mu_inp() (pycrocml.topology.Topology method),
    49
get_mu_out() (pycrocml.topology.Topology method),
    49
get_names() (pycrocml.package.Package method), 38
get_openalea_home_dir() (in module pycrocml.package), 41
get_path() (pycrocml.package.PyPackageReaderModel
    method), 41
get_path() (pycrocml.topology.PackageManager
    method), 49
get_pkg_files() (pycrocml.package.Package
    method), 39
get_pkg_name() (pycrocml.package.PyPackageReader
    method), 41
get_pkgrader() (pycrocml.package.PackageManager
    method), 40
get_properties() (pycrocml.transpiler.rules.javaRules.JavaRules
    attribute), 27
get_properties_compo (pycrocml.transpiler.rules.javaRules.JavaRules
    attribute), 27
get_str() (pycrocml.package.PyPackageWriter
    method), 41
get_tip() (pycrocml.package.Package method), 39
get_tip() (pycrocml.package.PseudoGroup method),
    40
get_userpkg_dir() (in module pycrocml.package),
    41
getset() (pycrocml.transpiler.generators.csharpGenerator.CsharpGenerator
    method), 16
getset() (pycrocml.transpiler.generators.javaGenerator.JavaTrans
    method), 21
gettype() (pycrocml.transpiler.generators.javaGenerator.JavaGenerator
    method), 19
gettype() (pycrocml.transpiler.generators.javaGenerator.JavaTrans
    method), 21

H
has_key() (pycrocml.package.PackageDict method),
    39
has_key() (pycrocml.package.PackageManager
    method), 40
header (pycrocml.transpiler.rules.sqRules.SqRules
    attribute), 28

    info_inputs_mu() (pycrocml.topology.Topology
        method), 49
    info_minout() (pycrocml.topology.Topology
        method), 49
    info_outputs_mu() (pycrocml.topology.Topology
        method), 49
    init() (pycrocml.package.PackageManager method),
        40
    initCompo() (pycrocml.transpiler.generators.csharpGenerator.CsharpCompo
        method), 13
    initCompo() (pycrocml.transpiler.generators.javaGenerator.JavaCompo
        method), 19
Initialization (class in pycrocml.initialization), 37
Initialization() (pycrocml.composition.ModelParser
    method), 36
Initialization() (pycrocml.pparse.ModelParser
    method), 42
initialization() (pycrocml.render_cyml.Model2Package
    method), 44
initWrap() (pycrocml.transpiler.generators.csharpGenerator.CsharpCompo
    method), 13
Input (class in pycrocml.inout), 37
Input() (pycrocml.pparse.ModelParser method), 42
InputOutput (class in pycrocml.inout), 37
Inputs() (pycrocml.pparse.ModelParser method), 42
instanceModels() (pycrocml.transpiler.generators.csharpGenerator.CsharpCompo
    method), 13
instanceModels() (pycrocml.transpiler.generators.javaGenerator.JavaCompo
    method), 19
int_expander() (in module pycrocml.transpiler.api_transform), 29
internal_declaration() (pycrocml.transpiler.generators.csharpGenerator.CsharpGenerator
    method), 14
internal_declaration() (pycrocml.transpiler.generators.fortranGenerator.FortranGenerator
    method), 17
internal_Declaration() (pycrocml.transpiler.generators.javaGenerator.JavaGenerator
    method), 19
is_directory() (pycrocml.package.Package
    method), 39
is_editable() (pycrocml.package.Package method),
    39
is_protected() (in module pycrocml.package), 42
isPackage() (pycrocml.topology.Topology method),
    49
items() (pycrocml.package.PackageManager method),
    40

```

```

iter_public_values() (py- method() (pycroml.transpiler.rules.sqRules.SqRules
    cropml.package.PackageDict method), 39 method), 28
iteritems() (pycroml.package.PackageManager methods (pycroml.transpiler.rules.csharpRules.CsharpRules
    method), 40 attribute), 25
iterkeys() (pycroml.package.PackageManager methods (pycroml.transpiler.rules.fortranRules.FortranRules
    method), 40 attribute), 26
itervalues() (pycroml.package.PackageManager methods (pycroml.transpiler.rules.javaRules.JavaRules
    method), 40 attribute), 27
methods (pycroml.transpiler.rules.pythonRules.PythonRules
attribute), 27
methods (pycroml.transpiler.rules.sqRules.SqRules
attribute), 28
middleware (class in pycroml.transpiler.interface), 33
mimetype (pycroml.package.Package attribute), 39
mimetype (pycroml.package.PseudoGroup attribute), 40
min_expander() (in module pycroml.transpiler.api_transform), 29
MINOR (in module pycroml.interface.version), 11
MINOR (in module pycroml.transpiler.version), 34
MINOR (in module pycroml.version), 50
minout() (pycroml.topology.Topology method), 49
mod() (in module pycroml.transpiler.builtin_typed_api), 31
Model, 52
Model() (pycroml.composition.ModelParser method), 36
Model1Nb (class in pycroml.code2nbk), 35
Model1Nb (class in pycroml.render_notebook), 46
Model1Nb (class in pycroml.render_notebook_csharp), 46
Model1Nb (class in pycroml.render_notebook_java), 47
model2Node() (pycroml.transpiler.generators.csharpGenerator.Csharp
method), 16
model2Node() (pycroml.transpiler.generators.javaGenerator.JavaTrans
method), 21
Model2Package (class in pycroml.render_csharp), 44
Model2Package (class in pycroml.render_cyml), 44
Model2Package (class in pycroml.render_fortran), 45
Model2Package (class in pycroml.render_java), 45
Model2Package (class in pycroml.render_python), 47
Model2Package (class in pycroml.render_R), 43
Model2Package (class in pycroml.writeTest_f90), 50
model_parser() (in module pycroml.composition), 36
model_parser() (in module pycroml.ppparse), 43
ModelComposition (class in pycroml.composition), 35
ModelComposition() (py-

```

**J**

```

JavaCompo (class in pycroml.transpiler.generators.javaGenerator), 19
JavaGenerator (class in pycroml.transpiler.generators.javaGenerator), 19
JavaRules (class in pycroml.transpiler.rules.javaRules), 27
JavaTrans (class in pycroml.transpiler.generators.javaGenerator), 21

```

**K**

```

keys() (pycroml.package.PackageManager method), 40

```

**L**

```

len_expander() (in module pycroml.transpiler.api_transform), 29
Links() (pycroml.composition.ModelParser method), 36
load_directory() (pycroml.package.PackageManager method), 40
load_pkge() (pycroml.topology.Topology method), 49
loadParamWrap() (pycroml.transpiler.generators.csharpGenerator.CsharpCom
method), 13
lower() (in module pycroml.package), 42

```

**M**

```

Main (class in pycroml.transpiler.main), 33
main() (in module pycroml.main), 38
MAJOR (in module pycroml.interface.version), 11
MAJOR (in module pycroml.transpiler.version), 34
MAJOR (in module pycroml.version), 50
max_expander() (in module pycroml.transpiler.api_transform), 29
meta_ext() (pycroml.topology.Topology method), 49
meta_inp() (pycroml.topology.Topology method), 49
meta_out() (pycroml.topology.Topology method), 49
method() (pycroml.transpiler.rules.fortranRules.FortranRules
method), 26

```

```

    cropml.composition.ModelParser      method), openalea_interface()      (in module py-
    36                                         cropml.transpiler.generators.openaleaGenerator),
ModelDefinition (class in pycropml.composition), 22
    35                                         OpenaleaCompo (class      in py-
ModelDefinition (class in pycropml.modelunit), 38  cropml.transpiler.generators.openaleaGenerator),
    38                                         22
ModelParser (class in pycropml.composition), 35  OpenaleaGenerator (class      in py-
    35                                         cropml.transpiler.generators.openaleaGenerator),
ModelParser (class in pycropml.pparse), 42  22
Models (class in pycropml.composition), 36
    36                                         operator_enter()          (py-
ModelUnit (class in pycropml.modelunit), 38  cropml.transpiler.codeGenerator.CodeGenerator
    38                                         method), 32
ModelUnit () (pycropml.pparse.ModelParser method),  operator_exit()          (py-
    42                                         cropml.transpiler.codeGenerator.CodeGenerator
modulo_expander () (in module py-  method), 32
    cropml.transpiler.api_transform), 29
mul () (in module py-  opt (class in pycropml.transpiler.Parser), 28
    cropml.transpiler.builtin_typed_api), 31
my_input () (in module pycropml.render_cyml), 45
my_input () (pycropml.render_fortran.Model2Package  or_()          (in module py-
    method), 45                                         cropml.transpiler.builtin_typed_api), 31
                                         Output (class in pycropml.inout), 37
                                         Output () (pycropml.pparse.ModelParser method), 42
                                         Outputs () (pycropml.pparse.ModelParser method), 42
                                         outputWrap () (pycropml.transpiler.generators.csharpGenerator.Csharp
                                         method), 13

N
namespace (pycropml.transpiler.rules.sqRules.SqRules
    attribute), 28
nb (pycropml.code2nbk.Model2Nb attribute), 35
nb_public_values () (py-
    cropml.package.PackageDict method), 39
new () (pycropml.package.PseudoGroup method), 40
newline () (pycropml.transpiler.codeGenerator.CodeGenerator
    method), 32
newtype () (pycropml.transpiler.ast_transform.AstTransformer
    method), 29
Node (class in pycropml.transpiler.pseudo_tree), 34
NodeVisitor (class      in py-
    cropml.transpiler.nodeVisitor), 34
notdeclared () (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 29
num (pycropml.render_csharp.Model2Package attribute), 44
num (pycropml.render_cyml.Model2Package attribute), 44
num (pycropml.render_fortran.Model2Package attribute), 45
num (pycropml.render_java.Model2Package attribute), 45
num (pycropml.render_python.Model2Package attribute), 48
num (pycropml.render_R.Model2Package attribute), 43
num (pycropml.writeTest_f90.Model2Package attribute), 51

O
openalea_interface() (in module py-
    cropml.render_python), 48
                                         P
                                         Package (class in pycropml.package), 38
                                         PackageManager (class in pycropml.topology), 48
                                         PackageDict (class in pycropml.package), 39
                                         PackageManager (class in pycropml.topology), 48
                                         param () (pycropml.pparse.ModelParser method), 42
                                         Parameterset (class in pycropml.parameterset), 42
                                         parameterset () (in module pycropml.parameterset), 42
                                         Parameterset () (pycropml.pparse.ModelParser
                                         method), 42
                                         Parametersets () (pycropml.pparse.ModelParser
                                         method), 42
                                         parse () (pycropml.composition.ModelParser method), 36
                                         parse () (pycropml.composition.Parser method), 36
                                         parse () (pycropml.pparse.ModelParser method), 42
                                         parse () (pycropml.pparse.Parser method), 43
                                         parse () (pycropml.transpiler.main.Main method), 33
                                         Parser (class in pycropml.composition), 36
                                         Parser (class in pycropml.pparse), 42
                                         parser () (in module pycropml.transpiler.Parser), 28
                                         part_declaraction () (py-
                                         cropml.transpiler.generators.fortranGenerator.FortranGenerator
                                         method), 17
                                         pkg_m () (pycropml.topology.Topology method), 49
                                         pkg_template (pycropml.package.PyPackageWriter
                                         attribute), 41
                                         pkgs (pycropml.topology.Topology attribute), 49

```

POST (*in module pycrocml.interface.version*), 11  
 POST (*in module pycrocml.transpiler.version*), 34  
 POST (*in module pycrocml.version*), 50  
 pow\_() (*in module pycrocml.transpiler.builtin\_typed\_api*), 31  
 pow\_expander() (*in module pycrocml.transpiler.api\_transform*), 29  
 prefix() (*in module pycrocml.cyml*), 36  
 prepare\_table() (*in module pycrocml.transpiler.helpers*), 33  
 private() (*pycrocml.transpiler.generators.csharpGenerator*), 16  
 private() (*pycrocml.transpiler.generators.javaGenerator*), 21  
 privateWrap() (*pycrocml.transpiler.generators.csharpGenerator.CsharpCompiler*), 13  
 protected() (*in module pycrocml.package*), 42  
 PseudoCythonNotTranslatableError, 32  
 PseudoCythonTypeCheckError, 32  
 PseudoError, 33  
 PseudoGroup (*class in pycrocml.package*), 40  
 public\_properties (*pycrocml.transpiler.rules.csharpRules.CsharpRules attribute*), 25  
 public\_properties\_compo (*pycrocml.transpiler.rules.csharpRules.CsharpRules attribute*), 25  
 public\_properties\_wrap (*pycrocml.transpiler.rules.csharpRules.CsharpRules attribute*), 25  
 pycrocml (*module*), 1, 51  
 pycrocml.algorithm (*module*), 34  
 pycrocml.checking (*module*), 35  
 pycrocml.code2nbk (*module*), 35  
 pycrocml.composition (*module*), 35  
 pycrocml.cyml (*module*), 36  
 pycrocml.description (*module*), 36  
 pycrocml.error (*module*), 37  
 pycrocml.formater\_f90 (*module*), 37  
 pycrocml.function (*module*), 37  
 pycrocml.initialization (*module*), 37  
 pycrocml.inout (*module*), 37  
 pycrocml.interface.version (*module*), 11  
 pycrocml.main (*module*), 38  
 pycrocml.modelunit (*module*), 38  
 pycrocml.package (*module*), 38  
 pycrocml.parameterset (*module*), 42  
 pycrocml.pparse (*module*), 42  
 pycrocml.render\_csharp (*module*), 44  
 pycrocml.render\_cyml (*module*), 44  
 pycrocml.render\_fortran (*module*), 45  
 pycrocml.render\_java (*module*), 45  
 pycrocml.render\_notebook (*module*), 46  
 pycrocml.render\_notebook\_csharp (*module*), 46  
 pycrocml.render\_notebook\_java (*module*), 47  
 pycrocml.render\_python (*module*), 47  
 pycrocml.render\_R (*module*), 43  
 pycrocml.test\_generator (*module*), 48  
 pycrocml.topology (*module*), 48  
 pycrocml.transpiler (*module*), 34  
 pycrocml.transpiler.api\_transform (*module*), 28  
 pycrocml.Transpiler.ast\_transform (*module*), 29  
 pycrocml.Transpiler.builtin\_typed\_api (*module*), 31  
 pycrocml.Transpiler.checkingModel (*module*), 31  
 pycrocml.Transpiler.codeGenerator (*module*), 32  
 pycrocml.Transpiler.env (*module*), 32  
 pycrocml.Transpiler.errors (*module*), 32  
 pycrocml.Transpiler.generators (*module*), 25  
 pycrocml.Transpiler.generators.checkGenerator (*module*), 12  
 pycrocml.Transpiler.generators.csharpGenerator (*module*), 13  
 pycrocml.Transpiler.generators.docGenerator (*module*), 16  
 pycrocml.Transpiler.generators.fortranGenerator (*module*), 17  
 pycrocml.Transpiler.generators.javaGenerator (*module*), 19  
 pycrocml.Transpiler.generators.openaleaGenerator (*module*), 22  
 pycrocml.Transpiler.generators.pythonGenerator (*module*), 22  
 pycrocml.Transpiler.generators.recordGenerator (*module*), 24  
 pycrocml.Transpiler.generators.simplaceGenerator (*module*), 24  
 pycrocml.Transpiler.generators.siriusGenerator (*module*), 24  
 pycrocml.Transpiler.helpers (*module*), 33  
 pycrocml.Transpiler.interface (*module*), 33  
 pycrocml.Transpiler.lib (*module*), 25  
 pycrocml.Transpiler.main (*module*), 33  
 pycrocml.Transpiler.nodeVisitor (*module*), 34  
 pycrocml.Transpiler.Parser (*module*), 28  
 pycrocml.Transpiler.pseudo\_tree (*module*), 34  
 pycrocml.Transpiler.rules (*module*), 28  
 pycrocml.Transpiler.rules.csharpRules (*module*), 25

```

pycropml.transpiler.rules.fortranRules
    (module), 26
pycropml.transpiler.rules.generalRule
    (module), 26
pycropml.transpiler.rules.javaRules
    (module), 27
pycropml.transpiler.rules.pythonRules
    (module), 27
pycropml.transpiler.rules.sqRules (mod-
    ule), 28
pycropml.transpiler.version (module), 34
pycropml.version (module), 50
pycropml.writeTest (module), 50
pycropml.writeTest_f90 (module), 50
pycropml.xml2wf (module), 51
PyPackageReader (class in pycropml.package), 41
PyPackageReaderModel (class in py-
    cropml.package), 41
PyPackageWriter (class in pycropml.package), 41
PythonCompo (class in py-
    cropml.transpiler.generators.pythonGenerator),
    22
PythonGenerator (class in py-
    cropml.transpiler.generators.pythonGenerator),
    22
PythonRules (class in py-
    cropml.transpiler.rules.pythonRules), 27

R
rebuild_category () (py-
    cropml.package.PackageManager method), 40
register_packages () (py-
    cropml.package.AbstractPackageReader method), 38
register_packages () (py-
    cropml.package.PyPackageReader method), 41
reload() (pycropml.package.Package method), 39
reload() (pycropml.package.PackageManager method), 40
remove_files() (pycropml.package.Package method), 39
retrieve_library () (py-
    cropml.transpiler.ast_transform.AstTransformer method), 29
retrieve_params () (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        (in module pycropml.render_csharp), 47
retrieve_params () (py-
    cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        (in module pycropml.render_notebook_fortran), 47
retrieve_params () (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator)
        (in module pycropml.render_java), 46
method), 19
retrieve_path () (in module pycropml.composition), 36
retrievePackage () (pycropml.xml2wf.XmlToWf
    method), 51
retrieve() (pycropml.topology.Topology method), 49
run() (pycropml.render_cyml.Model2Package method),
    44
run() (pycropml.render_notebook.Model2Nb method),
    46
run() (pycropml.render_notebook_csharp.Model2Nb
    method), 47
run() (pycropml.render_notebook_java.Model2Nb
    method), 47
run() (pycropml.render_python.Model2Package
    method), 48
run() (pycropml.render_R.Model2Package method), 43
run() (pycropml.writeTest_f90.Model2Package
    method), 51
run() (pycropml.xml2wf.XmlToWf method), 51

S
safe_double () (py-
    cropml.transpiler.codeGenerator.CodeGenerator
    method), 32
safe_serialize_type () (in module py-
    cropml.transpiler.helpers), 33
sep (pycropml.package.PseudoGroup attribute), 41
serialize_type () (in module py-
    cropml.transpiler.helpers), 33
set_properties (py-
    cropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
set_properties_compo (py-
    cropml.transpiler.rules.javaRules.JavaRules
    attribute), 27
set_sys_crop2ml_path () (py-
    cropml.package.PackageManager method), 40
setCompo () (pycropml.transpiler.generators.csharpGenerator.CsharpCom-
    position method), 13
setCompo () (pycropml.transpiler.generators.javaGenerator.JavaCompo-
    nent method), 19
signature () (in module pycropml.render_csharp), 44
signature () (in module pycropml.render_cyml), 45
signature () (in module pycropml.render_fortran), 45
signature () (in module pycropml.render_java), 46
signature () (in module pycropml.render_notebook_csharp), 47
signature () (in module pycropml.render_notebook_fortran), 47
signature () (in module pycropml.render_notebook_java), 47
signature () (in module pycropml.render_python), 48
signature () (in module pycropml.render_R), 43

```

signature() (in module `pycroml.transpiler.generators.openaleaGenerator`), 22

signature() (in module `pycroml.writeTest_f90`), 51

`SimplaceCompo` (class in `pycroml.transpiler.generators.simplaceGenerator`), 24

`SimplaceGenerator` (class in `pycroml.transpiler.generators.simplaceGenerator`), 24

`simplify()` (in module `pycroml.transpiler.builtin_typed_api`), 31

`SiriusCompo` (class in `pycroml.transpiler.generators.siriusGenerator`), 24

`SiriusGenerator` (class in `pycroml.transpiler.generators.siriusGenerator`), 24

`SiriusTrans` (class in `pycroml.transpiler.generators.siriusGenerator`), 25

`SqRules` (class in `pycroml.transpiler.rules.sqRules`), 28

`Standard` (class in `pycroml.transpiler.api_transform`), 28

`StandardCall` (class in `pycroml.transpiler.api_transform`), 28

`StandardCallAttrib` (class in `pycroml.transpiler.api_transform`), 29

`StandardMethodCall` (class in `pycroml.transpiler.api_transform`), 29

`StandardSwapper` (class in `pycroml.transpiler.api_transform`), 29

`sub()` (in module `pycroml.transpiler.builtin_typed_api`), 31

**T**

`tab_aware()` (in module `pycroml.transpiler.errors`), 33

`Test` (class in `pycroml.checking`), 35

`Testset` (class in `pycroml.checking`), 35

`testset()` (in module `pycroml.checking`), 35

`Testset()` (`pycroml.pparse.ModelParser` method), 42

`Testsets()` (`pycroml.pparse.ModelParser` method), 42

`to_ast()` (`pycroml.transpiler.main.Main` method), 33

`to_source()` (`pycroml.transpiler.main.Main` method), 33

`to_struct_cs()` (in module `pycroml.transpiler.generators.csharpGenerator`), 16

`to_struct_java()` (in module `pycroml.transpiler.generators.javaGenerator`), 21

`to_struct_sirius()` (in module `pycroml.transpiler.generators.siriusGenerator`), 25

`to_wrapper_cs()` (in module `pycroml.transpiler.generators.csharpGenerator`), 16

`to_wrapper_sirius()` (in module `pycroml.transpiler.generators.siriusGenerator`), 25

`topologicalSort()` (`pycroml.topology.Topology` method), 49

`Topology` (class in `pycroml.topology`), 49

`tranAssignParam()` (in module `pycroml.transpiler.generators.csharpGenerator.CsharpCompo` method), 14

`trans_format_parse()` (in module `pycroml.transpiler.rules.javaRules`), 27

`transBool()` (in module `pycroml.render_cyml`), 45

`transf()` (in module `pycroml.render_csharp`), 44

`transf()` (in module `pycroml.render_cyml`), 45

`transf()` (in module `pycroml.render_java`), 46

`transf()` (in module `pycroml.render_notebook_csharp`), 47

`transf()` (in module `pycroml.writeTest_f90`), 51

`transfDate()` (in module `pycroml.render_csharp`), 44

`transfDate()` (in module `pycroml.render_cyml`), 45

`transfDate()` (in module `pycroml.render_java`), 46

`transfDate()` (in module `pycroml.render_notebook_csharp`), 47

`transfDate()` (in module `pycroml.writeTest_f90`), 51

`transfDateList()` (in module `pycroml.render_csharp`), 44

`transfDateList()` (in module `pycroml.render_cyml`), 45

`transfDateList()` (in module `pycroml.render_java`), 46

`transfDateList()` (in module `pycroml.render_notebook_csharp`), 47

`transfDateList()` (in module `pycroml.writeTest_f90`), 51

`transfDouble()` (in module `pycroml.render_csharp`), 44

`transfDouble()` (in module `pycroml.render_java`), 46

`transfDouble()` (in module `pycroml.render_notebook_csharp`), 47

`transfDouble()` (in module `pycroml.writeTest_f90`), 51

`transfList()` (in module `pycroml.render_csharp`), 44

`transfList()` (in module `pycroml.render_java`), 46

`transform()` (`pycroml.transpiler.interface.TreeInterface`

*method), 33*  
*transform\_block ()* (in module *cropml.transpiler.interface.TreeInterface*)  
*method), 33*  
*transform\_default ()* (in module *cropml.transpiler.interface.TreeInterface*)  
*method), 33*  
*transform\_return ()* (in module *cropml.transpiler.generators.csharpGenerator*)  
*method), 14*  
*transform\_return ()* (in module *cropml.transpiler.generators.fortranGenerator*)  
*method), 17*  
*transform\_return ()* (in module *cropml.transpiler.generators.javaGenerator*)  
*method), 19*  
*transform\_to\_syntax\_tree ()* (in module *cropml.transpiler.ast\_transform*, 31)  
*transformer ()* (in module *cropml.transpiler.ast\_transform.AstTransformer*)  
*method), 29*  
*transfSDIList ()* (in module *cropml.render\_notebook\_csharp*, 47)  
*transfSDIList ()* (in module *cropml.writeTest\_f90*, 51)  
*transfString ()* (in module *cropml.render\_csharp*, 44)  
*transfString ()* (in module *pycropml.render\_java*, 46)  
*transfString ()* (in module *cropml.render\_notebook\_csharp*, 47)  
*transfString ()* (in module *cropml.writeTest\_f90*, 51)  
*translate ()* (in module *pycropml.topology.Topology*)  
*method), 49*  
*translate ()* (in module *pycropml.transpiler.main.Main*)  
*method), 34*  
*translate\_all ()* (in module *pycropml.topology.Topology*)  
*method), 49*  
*translateAppend ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translateCeil ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translateContains ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translateDictkeys ()* (in module *cropml.transpiler.rules.javaRules*, 27)  
*translateDictkeys ()* (in module *cropml.transpiler.rules.pythonRules*, 27)  
*translateFind ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translateGet ()* (in module *cropml.transpiler.rules.csharpRules*, 26)  
*translateIndex ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translatekeyDict ()* (in module *cropml.transpiler.rules.csharpRules*, 26)  
*translateLenArray ()* (in module *cropml.transpiler.rules.csharpRules*, 25)  
*translateLenArray ()* (in module *cropml.transpiler.rules.javaRules*, 27)  
*translateLenDict ()* (in module *cropml.transpiler.rules.javaRules*, 27)  
*translateLenList ()* (in module *cropml.transpiler.rules.csharpRules*, 26)  
*translateNotContains ()* (in module *cropml.transpiler.rules.csharpRules*, 26)  
*translateNotContains ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translateNotContains ()* (in module *cropml.transpiler.rules.javaRules*, 27)  
*translateNotContains ()* (in module *cropml.transpiler.rules.pythonRules*, 27)  
*translatePop ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translatePow ()* (in module *cropml.transpiler.rules.fortranRules*, 26)  
*translateSum ()* (in module *cropml.transpiler.rules.csharpRules*, 26)  
*translateSum ()* (in module *cropml.transpiler.rules.javaRules*, 27)  
*translation\_error ()* (in module *pycropml.errors*, 33)  
*transpile\_file ()* (in module *pycropml.cyml*, 36)  
*transpile\_package ()* (in module *pycropml.cyml*, 36)  
*TreeInterface* (class in *cropml.transpiler.interface*, 33)  
*type\_check\_error ()* (in module *pycropml.transpiler.errors*, 33)  
*types* (*pycropml.transpiler.rules.csharpRules.CsharpRules* attribute), 25  
*types* (*pycropml.transpiler.rules.fortranRules.FortranRules* attribute), 26  
*types* (*pycropml.transpiler.rules.javaRules.JavaRules* attribute), 27  
*types* (*pycropml.transpiler.rules.pythonRules.PythonRules* attribute), 27  
*types2* (*pycropml.transpiler.rules.javaRules.JavaRules* attribute), 27  
*U*  
*unary\_op* (*pycropml.transpiler.rules.csharpRules.CsharpRules* attribute), 25

unary\_op (*pycropml.transpiler.rules.fortranRules.FortranRules*.*attribute*), 26  
 unary\_op (*pycropml.transpiler.rules.javaRules.JavaRules*.*attribute*), 27  
 unary\_op (*pycropml.transpiler.rules.pythonRules.PythonRules*.*attribute*), 27  
 UnknownNodeError, 41  
 unop\_precedence (*pycropml.transpiler.codeGenerator.CodeGenerator*.*attribute*), 32  
 unop\_precedence (*pycropml.transpiler.generators.fortranGenerator.FortranGenerator*.*attribute*), 17  
 update\_category () (*pycropml.package.PackageManager*.*method*), 40  
 update\_modelunit () (*pycropml.package.Package*.*method*), 39  
 UserPackage (*class in pycropml.package*), 41

**V**

val\_init () (*pycropml.topology.Topology*.*method*), 49  
 valParam () (*in module* *pycropml.transpiler.generators.fortranGenerator*), 19  
 values () (*pycropml.package.PackageManager*.*method*), 40  
 visit () (*pycropml.transpiler.nodeVisitor.NodeVisitor*.*method*), 34  
 visit\_addnode () (*pycropml.transpiler.ast\_transform.AstTransformer*.*method*), 29  
 visit\_array () (*pycropml.transpiler.codeGenerator.CodeGenerator*.*method*), 32  
 visit\_array () (*pycropml.transpiler.generators.checkGenerator.CheckGenerator*.*method*), 12  
 visit\_array () (*pycropml.transpiler.generators.csharpGenerator.CsharpGenerator*.*method*), 14  
 visit\_array () (*pycropml.transpiler.generators.javaGenerator.JavaGenerator*.*method*), 19  
 visit\_array () (*pycropml.transpiler.generators.pythonGenerator.PythonGenerator*.*method*), 22  
 visit\_array\_decl () (*pycropml.transpiler.generators.csharpGenerator.CsharpGenerator*.*method*), 14  
 visit\_array\_decl () (*pycropml.transpiler.generators.csharpGenerator.CsharpGenerator*.*method*), 16

visit\_array\_decl () (*pycropml.transpiler.generators.fortranGenerator.FortranGenerator*.*method*), 17  
 visit\_assignment () (*pycropml.transpiler.generators.javaGenerator.JavaTrans*.*method*), 21  
 visit\_assignment () (*pycropml.transpiler.generators.checkGenerator.CheckGenerator*.*method*), 12  
 visit\_assignment () (*pycropml.transpiler.generators.csharpGenerator.CsharpCompo*.*method*), 14  
 visit\_assignment () (*pycropml.transpiler.generators.csharpGenerator.CsharpGenerator*.*method*), 14  
 visit\_assignment () (*pycropml.transpiler.generators.fortranGenerator.FortranGenerator*.*method*), 17  
 visit\_assignment () (*pycropml.transpiler.generators.javaGenerator.JavaCompo*.*method*), 19  
 visit\_assignment () (*pycropml.transpiler.generators.javaGenerator.JavaGenerator*.*method*), 20  
 visit\_assignment () (*pycropml.transpiler.generators.pythonGenerator.PythonGenerator*.*method*), 22  
 visit\_attributemode () (*pycropml.transpiler.ast\_transform.AstTransformer*.*method*), 29  
 visit\_binary\_op () (*pycropml.transpiler.generators.checkGenerator.CheckGenerator*.*method*), 12  
 visit\_binary\_op () (*pycropml.transpiler.generators.csharpGenerator.CsharpGenerator*.*method*), 14  
 visit\_binary\_op () (*pycropml.transpiler.generators.fortranGenerator.FortranGenerator*.*method*), 17  
 visit\_binary\_op () (*pycropml.transpiler.generators.javaGenerator.JavaGenerator*.*method*), 20  
 visit\_binary\_op () (*pycropml.transpiler.generators.pythonGenerator.PythonGenerator*.*method*), 22  
 visit\_binopnode () (*pycropml.transpiler.ast\_transform.AstTransformer*.*method*), 29  
 visit\_bool () (*pycropml.transpiler.generators.checkGenerator.CheckGenerator*.*method*), 12  
 visit\_bool () (*pycropml.transpiler.generators.csharpGenerator.CsharpGenerator*.*method*)

```

        method), 14
visit_bool() (pycropml.transpiler.generators.fortranGenerator.FortranGenerator
        method), 17
visit_bool() (pycropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_bool() (pycropml.transpiler.generators.pythonGenerator.PythonGenerator
        method), 22
visit_bool_decl() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 14
visit_bool_decl() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 16
visit_bool_decl() (py-
        cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        method), 17
visit_bool_decl() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_bool_decl() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_bool_decl() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 21
visit_boolbinopnode() (py-
        cropml.transpiler.ast_transform.AstTransformer)
        method), 29
visit_boolnode() (py-
        cropml.transpiler.ast_transform.AstTransformer)
        method), 29
visit_breakstatnode() (py-
        cropml.transpiler.ast_transform.AstTransformer)
        method), 30
visit_breakstatnode() (py-
        cropml.transpiler.generators.checkGenerator.CheckGenerator)
        method), 12
visit_breakstatnode() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 14
visit_breakstatnode() (py-
        cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        method), 17
visit_breakstatnode() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_breakstatnode() (py-
        cropml.transpiler.generators.pythonGenerator.PythonGenerator)
        method), 22
visit_call() (pycropml.transpiler.generators.checkGenerator.CheckGenerator
        method), 12
visit_call() (pycropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 14
visit_call() (pycropml.transpiler.generators.fortranGenerator.FortranGenerator
        method), 17
visit_call() (pycropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_call() (pycropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_call() (pycropml.transpiler.generators.pythonGenerator.PythonGenerator
        method), 20
visit_cargdeclnode() (py-
        cropml.transpiler.generators.checkGenerator.CheckGenerator)
        method), 12
visit_cargdeclnode() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 14
visit_cargdeclnode() (py-
        cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        method), 18
visit_cargdeclnode() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_cargdeclnode() (py-
        cropml.transpiler.generators.pythonGenerator.PythonGenerator)
        method), 22
visit_cond_expr_node() (py-
        cropml.transpiler.generators.checkGenerator.CheckGenerator)
        method), 12
visit_cond_expr_node() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 14
visit_cond_expr_node() (py-
        cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        method), 14
visit_cond_expr_node() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 18
visit_cond_expr_node() (py-
        cropml.transpiler.generators.pythonGenerator.PythonGenerator)
        method), 20
visit_continuestatnode() (py-
        cropml.transpiler.ast_transform.AstTransformer)
        method), 30
visit_continuestatnode() (py-
        cropml.transpiler.ast_transform.AstTransformer)
        method), 30
visit_continuestatnode() (py-
        cropml.transpiler.generators.checkGenerator.CheckGenerator)
        method), 12
visit_continuestatnode() (py-
        cropml.transpiler.generators.pythonGenerator.PythonGenerator)
        method), 22
visit_continuestatnode() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 12
visit_continuestatnode() (py-
        cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        method), 18
visit_continuestatnode() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_continuestatnode() (py-
        cropml.transpiler.generators.pythonGenerator.PythonGenerator)
        method), 20
visit_continuestatnode() (py-
        cropml.transpiler.generators.csharpGenerator.CsharpGenerator)
        method), 18
visit_continuestatnode() (py-
        cropml.transpiler.generators.fortranGenerator.FortranGenerator)
        method), 20
visit_continuestatnode() (py-
        cropml.transpiler.generators.javaGenerator.JavaGenerator)
        method), 20
visit_continuestatnode() (py-
        cropml.transpiler.generators.pythonGenerator.PythonGenerator)
        method), 20

```

```
cropml.transpiler.generators.pythonGenerator.PythonGeneratorMethod), 14
method), 23 visit_decl() (py- visit_decl() (py- visit_decl() (py-
visit_csimplebasetypenode() (py- method), 16 method), 16 method), 18
cropml.transpiler.ast_transform.AstTransformer visit_decl() (py- visit_decl() (py-
method), 30 method), 18 method), 18
visit_custom_call() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.checkGenerator.CheckGeneratorMethod), 20 method), 21 method), 21
method), 12 visit_decl() (py- visit_decl() (py-
visit_custom_call() (py- method), 21 method), 21
cropml.transpiler.generators.csharpGenerator.CsharpGeneratorDeclaration() (py-
method), 14 method), 14
visit_custom_call() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.fortranGenerator.FortranGeneratorDeclaration() method), 12 method), 12
method), 18 method), 18
visit_custom_call() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.csharpGenerator.CsharpCompo method), 14 method), 14
method), 20 method), 14
visit_custom_call() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.javaGenerator.JavaGeneratorDeclaration() method), 14 method), 14
method), 23 method), 14
visit_cvardefnode() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.ast_transform.AstTransformer visit_decl() (py- method), 18 method), 18
method), 30 method), 18
visit_datetime() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.checkGenerator.CheckGeneratorDeclaration() method), 19 method), 19
method), 12 method), 12
visit_datetime() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.csharpGenerator.CsharpGeneratorDeclaration() method), 20 method), 20
method), 14 method), 14
visit_DateTime() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.csharpGenerator.CsharpTransDeclaration() method), 23 method), 23
method), 16 method), 16
visit_datetime() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.fortranGenerator.FortranGeneratorDefinitions() method), 24 method), 24
method), 18 method), 18
visit_datetime() (py- visit_decl() (py- visit_decl() (py-
cropml.transpiler.generators.pythonGenerator.PythonGeneratorDeclaration() method), 30 method), 30
method), 23 method), 23
visit_datetime_decl() (py- visit_dict() (py- visit_dict() (py-
cropml.transpiler.generators.csharpGenerator.CsharpGeneratorMethod) (py- visit_dict() (py-
method), 14 method), 14
method), 14 method), 12
visit_datetime_decl() (py- visit_dict() (py- visit_dict() (py-
cropml.transpiler.generators.csharpGenerator.CsharpTransMethod), 14 method), 14
method), 16 visit_dict() (py- visit_dict() (py-
visit_datetime_decl() (py- method), 20 method), 20
method), 18 visit_dict() (py- visit_dict() (py-
visit_datetime_decl() (py- method), 23 method), 23
method), 18 method), 18
visit_datetime_decl() (py- visit_dict_decl() (py- visit_dict_decl() (py-
cropml.transpiler.generators.javaGenerator.JavaGenerator cropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 20 method), 14 method), 14
visit_datetime_decl() (py- visit_dict_decl() (py- visit_dict_decl() (py-
cropml.transpiler.generators.javaGenerator.JavaTrans cropml.transpiler.generators.csharpGenerator.CsharpTrans
method), 21 method), 16 method), 16
visit_decl() (py- visit_decl() (py- visit_decl() (py-
```

```

    cropml.transpiler.generators.javaGenerator.JavaGenerator cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 20                                         method), 17                                         (py-
visit_dict_decl()                               (py- visit_ExprStatNode()                               (py-
    cropml.transpiler.generators.javaGenerator.JavaTrans   cropml.transpiler.generators.pythonGenerator.PythonGenerator
method), 21                                         method), 22
visit_dictnode()                                (py- visit_float()                                (py-
    cropml.transpiler.ast_transform.AstTransformer        cropml.transpiler.generators.checkGenerator.CheckGenerator
method), 30                                         method), 12
visit_divnode()                                 (py- visit_float()                                (py-
    cropml.transpiler.ast_transform.AstTransformer        cropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 30                                         method), 15
visit_elements()                                (py- visit_float()                                (py-
    cropml.transpiler.ast_transform.AstTransformer        cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 30                                         method), 18
visit_else_statement()                           (py- visit_float()                                (py-
    cropml.transpiler.generators.checkGenerator.CheckGenerator) opml.transpiler.generators.javaGenerator.JavaGenerator
method), 12                                         method), 20
visit_else_statement()                           (py- visit_float()                                (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator) opml.transpiler.generators.pythonGenerator.PythonGenerator
method), 15                                         method), 23
visit_else_statement()                           (py- visit_float_decl()                               (py-
    cropml.transpiler.generators.fortranGenerator.FortranGenerator) opml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 18                                         method), 15
visit_else_statement()                           (py- visit_float_decl()                               (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator) opml.transpiler.generators.csharpGenerator.CsharpTrans
method), 20                                         method), 16
visit_else_statement()                           (py- visit_float_decl()                               (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator) opml.transpiler.generators.fortranGenerator.FortranGenerator
method), 23                                         method), 18
visit_elseif_statement()                         (py- visit_float_decl()                               (py-
    cropml.transpiler.generators.checkGenerator.CheckGenerator) opml.transpiler.generators.javaGenerator.JavaGenerator
method), 12                                         method), 20
visit_elseif_statement()                         (py- visit_float_decl()                               (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator) opml.transpiler.generators.javaGenerator.JavaTrans
method), 15                                         method), 21
visit_elseif_statement()                         (py- visit_floatnode()                               (py-
    cropml.transpiler.generators.fortranGenerator.FortranGenerator) opml.transpiler.ast_transform.AstTransformer
method), 18                                         method), 30
visit_elseif_statement()                         (py- visit_for_iterator()                               (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator) opml.transpiler.generators.checkGenerator.CheckGenerator
method), 20                                         method), 12
visit_elseif_statement()                         (py- visit_for_iterator()                               (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator) opml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 23                                         method), 15
visit_exprstatnode()                            (py- visit_for_iterator()                               (py-
    cropml.transpiler.ast_transform.AstTransformer        cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 30                                         method), 18
visit_ExprStatNode()                            (py- visit_for_iterator()                               (py-
    cropml.transpiler.codeGenerator.CodeGenerator       cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 32                                         method), 20
visit_ExprStatNode()                            (py- visit_for_iterator()                               (py-
    cropml.transpiler.generators.checkGenerator.CheckGenerator) opml.transpiler.generators.pythonGenerator.PythonGenerator
method), 12                                         method), 23
visit_ExprStatNode()                            (py- visit_for_iterator_with_index()                (py-

```

```

    cropml.transpiler.generators.checkGenerator.CheckGenerator(cropml.transpiler.generators.checkGenerator.CheckGenerator
method), 12
visit_for_iterator_with_index()      (py- visit_for_statement())          (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator(cropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 15
visit_for_iterator_with_index()      (py- visit_for_statement())          (py-
    cropml.transpiler.generators.fortranGenerator.FortranGenerator(cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 18
visit_for_iterator_with_index()      (py- visit_for_statement())          (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator(cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 20
visit_for_iterator_with_index()      (py- visit_for_statement())          (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator(cropml.transpiler.generators.pythonGenerator.PythonGenerator
method), 23
visit_for_range_statement()         (py- visit_forinstatnode())          (py-
    cropml.transpiler.generators.checkGenerator.CheckGenerator(cropml.transpiler.ast_transform.AstTransformer
method), 12
visit_for_range_statement()         (py- visit_function_definition())      (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator(cropml.transpiler.generators.checkGenerator.CheckGenerator
method), 15
visit_for_range_statement()         (py- visit_function_definition())      (py-
    cropml.transpiler.generators.fortranGenerator.FortranGenerator(cropml.transpiler.generators.csharpGenerator.CsharpCompo
method), 18
visit_for_range_statement()         (py- visit_function_definition())      (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator(cropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 20
visit_for_range_statement()         (py- visit_function_definition())      (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator(cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 23
visit_for_sequence()               (py- visit_function_definition())      (py-
    cropml.transpiler.codeGenerator.CodeGenerator(cropml.transpiler.generators.javaGenerator.JavaCompo
method), 32
visit_for_sequence()               (py- visit_function_definition())      (py-
    cropml.transpiler.generators.checkGenerator.CheckGenerator(cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 12
visit_for_sequence()               (py- visit_function_definition())      (py-
    cropml.transpiler.generators.fortranGenerator.FortranGenerator(cropml.transpiler.generators.pythonGenerator.PythonGenerator
method), 18
visit_for_sequence()               (py- visit_function_definition())      (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator(cropml.transpiler.generators.simplaceGenerator.SimplaceGenera
method), 23
visit_for_sequence_with_index()    (py- visit_if_statement())          (py-
    cropml.transpiler.generators.checkGenerator.CheckGenerator(cropml.transpiler.generators.checkGenerator.CheckGenerator
method), 12
visit_for_sequence_with_index()    (py- visit_if_statement())          (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator(cropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 15
visit_for_sequence_with_index()    (py- visit_if_statement())          (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator(cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 20
visit_for_sequence_with_index()    (py- visit_if_statement())          (py-
    cropml.transpiler.generators.pythonGenerator.PythonGenerator(cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 23
visit_for_statement()              (py- visit_if_statement())          (py-

```

```

    cropml.transpiler.generators.pythonGenerator.PythonGeneratoropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 23                                         method), 18
visit_ifclausenode()                               (py- visit_importfrom()          (py-
    cropml.transpiler.ast_transform.AstTransformer   method), 20
method), 30
visit_ifstatnode()                                (py- visit_importfrom()          (py-
    cropml.transpiler.ast_transform.AstTransformer   method), 23
method), 30
visit_implicit_return()                           (py- visit_index()           (py-
    cropml.transpiler.generators.checkGenerator.CheckGeneratoropml.transpiler.generators.checkGenerator.CheckGenerator
method), 12                                         method), 13
visit_implicit_return()                           (py- visit_index()           (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGeneratoropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 15                                         method), 15
visit_implicit_return()                           (py- visit_index()           (py-
    cropml.transpiler.generators.fortranGenerator.FortranGeneratoropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 18                                         method), 18
visit_implicit_return()                           (py- visit_index()           (py-
    cropml.transpiler.generators.javaGenerator.JavaCompo   cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 19                                         method), 20
visit_implicit_return()                           (py- visit_index()           (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator cropml.transpiler.generators.pythonGenerator.PythonGenerator
method), 20                                         method), 23
visit_implicit_return()                           (py- visit_indexnode()          (py-
    cropml.transpiler.generators.pythonGeneratorPythonGeneratoropml.transpiler.ast_transform.AstTransformer
method), 23                                         method), 30
visit_import()                                 (py- visit_inplaceassignmentnode() (py-
    cropml.transpiler.generators.checkGenerator.CheckGeneratoropml.transpiler.ast_transform.AstTransformer
method), 13                                         method), 30
visit_import()                                 (py- visit_int() (pycropml.transpiler.codeGenerator.CodeGenerator
    cropml.transpiler.generators.csharpGenerator.CsharpGeneratormethod), 32
method), 15                                         visit_int() (pycropml.transpiler.generators.fortranGenerator.FortranC
method), 18
visit_import()                                 (py- method), 18
    cropml.transpiler.generators.fortranGenerator.FortranGeneratordecl() (py-
method), 18                                         cropml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 15
visit_import()                                 (py- method), 15
    cropml.transpiler.generators.javaGenerator.JavaGeneratorint_decl() (py-
method), 20                                         cropml.transpiler.generators.csharpGenerator.CsharpTrans
method), 16
visit_import()                                 (py- method), 16
    cropml.transpiler.generators.pythonGenerator.PythonGenerator_decl() (py-
method), 23                                         cropml.transpiler.generators.fortranGenerator.FortranGenerator
method), 18
visit_import()                                 (py- method), 18
    cropml.transpiler.generators.simpleGenerator.SimpleGeneratorcl() (py-
method), 24                                         cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 20
visit_importfrom()                            (py- method), 20
    cropml.transpiler.generators.checkGenerator.CheckGeneratorint_decl() (py-
method), 13                                         cropml.transpiler.generators.javaGenerator.JavaTrans
method), 13
visit_importfrom()                            (py- method), 21
    cropml.transpiler.generators.csharpGenerator.CsharpGeneratornode() (py-
method), 15                                         cropml.transpiler.ast_transform.AstTransformer
method), 30
visit_importfrom()                            (py- method), 30
    cropml.transpiler.generators.fortranGenerator.FortranCompost() (pycropml.transpiler.generators.checkGenerator.CheckG
method), 17                                         method), 13
visit_importfrom()                            (py- visit_list() (pycropml.transpiler.generators.csharpGenerator.Csharp
method), 17
method), 17

```

```

        method), 15
visit_list () (pycrocml.transpiler.generators.fortranGenerator.FortranGenerator
        method), 18
(crocml.transpiler.generators.checkGenerator.CheckGenerator
visit_list () (pycrocml.transpiler.generators.javaGenerator.JavaGenerator
        method), 20
(crocml.transpiler.generators.javaGenerator.JavaGenerator
visit_list () (pycrocml.transpiler.generators.pythonGenerator.PythonGenerator
        method), 23
(crocml.transpiler.generators.csharpGenerator.CsharpCompo
method), 14
visit_list_decl () (py- visit_module ())
(crocml.transpiler.generators.csharpGenerator.CsharpGenerat(crocml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 15
visit_list_decl () (py- visit_module ())
(crocml.transpiler.generators.csharpGenerator.CsharpTrans(crocml.transpiler.generators.fortranGenerator.FortranGenerator
method), 16
visit_list_decl () (py- visit_module ())
(crocml.transpiler.generators.fortranGenerator.FortranGenerat(crocml.transpiler.generators.javaGenerator.JavaCompo
method), 18
visit_list_decl () (py- visit_module ())
(crocml.transpiler.generators.javaGenerator.JavaGenerator cropml.transpiler.generators.javaGenerator.JavaGenerator
method), 20
visit_list_decl () (py- visit_module ())
(crocml.transpiler.generators.javaGenerator.JavaTrans cropml.transpiler.generators.pythonGenerator.PythonGenerator
method), 21
visit_listnode () (py- visit_module ())
(crocml.transpiler.ast_transform.AstTransformer
method), 30
visit_local () (py- visit_module ())
(crocml.transpiler.codeGenerator.CodeGenerator
method), 32
visit_local () (py- visit_mulnode ())
(crocml.transpiler.generators.checkGenerator.CheckGenerator
method), 13
visit_local () (py- visit_namenode ())
(crocml.transpiler.generators.csharpGenerator.CsharpCompo(crocml.transpiler.ast_transform.AstTransformer
method), 14
visit_local () (py- visit_node ()) (pycrocml.transpiler.ast_transform.AstTransformer
(crocml.transpiler.generators.simplaceGenerator.SimplaceGener
method), 30
visit_local () (py- visit_notAnumber ())
(crocml.transpiler.ast_transform.AstTransformer
method), 24
visit_method_call () (py- visit_notAnumber ())
(crocml.transpiler.generators.checkGenerator.CheckGenerator
method), 13
visit_method_call () (py- visit_notAnumber ())
(crocml.transpiler.generators.csharpGenerator.CsharpGenerat(crocml.transpiler.generators.csharpGenerator.CsharpGenerator
method), 15
visit_method_call () (py- visit_notAnumber ())
(crocml.transpiler.generators.fortranGenerator.FortranGenerat(crocml.transpiler.generators.fortranGenerator.FortranGenerator
method), 18
visit_method_call () (py- visit_notAnumber ())
(crocml.transpiler.generators.javaGenerator.JavaGenerator
method), 21
visit_method_call () (py- visit_notAnumber ())
(crocml.transpiler.generators.javaGenerator.JavaGenerator
method), 20
visit_method_call () (py- visit_notAnumber ())
(crocml.transpiler.generators.pythonGenerator.PythonGenerat(crocml.transpiler.generators.pythonGenerator.PythonGenerator
method), 23
visit_modnode () (py- visit_notnode ())
(crocml.transpiler.ast_transform.AstTransformer
method), 30

```

```

visit_pair() (pycropml.transpiler.generators.checkGenerator.CheckGenerator
    method), 13                                         visit_sliceindex() (py-
                                                               cropml.transpiler.generators.csharpGenerator.CsharpGenerator
                                                               method), 13
visit_pair() (pycropml.transpiler.generators.fortranGenerator.FortranGenerator
    method), 18                                         (py-
                                                               cropml.transpiler.generators.csharpGenerator.CsharpGenerator
                                                               method), 15
visit_pair() (pycropml.transpiler.generators.javaGenerator.JavaGenerator
    method), 21                                         (py-
                                                               visit_sliceindex()
                                                               cropml.transpiler.generators.pythonGenerator.PythonGenerator
                                                               method), 15
visit_pair() (pycropml.transpiler.generators.pythonGenerator.PythonGenerator
    method), 23                                         (py-
                                                               visit_sliceindex()
                                                               cropml.transpiler.generators.fortranGenerator.FortranGenerator
                                                               method), 18
visit_pownode() (py- visit_sliceindex() (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 30                                         cropml.transpiler.generators.javaGenerator.JavaGenerator
                                                               method), 21
visit_primarycmpnode() (py- visit_sliceindex() (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 30                                         cropml.transpiler.generators.pythonGenerator.PythonGenerator
                                                               method), 23
visit_print() (py- visit_sliceindexnode() (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator
    method), 15                                         cropml.transpiler.ast_transform.AstTransformer
                                                               method), 30
visit_print() (py- visit_standard_call() (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator
    method), 21                                         cropml.transpiler.generators.checkGenerator.CheckGenerator
                                                               method), 13
visit_printstatnode() (py- visit_standard_call() (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 30                                         cropml.transpiler.generators.csharpGenerator.CsharpGenerator
                                                               method), 15
visit_pyclassdefnode() (py- visit_standard_call() (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 30                                         cropml.transpiler.generators.fortranGenerator.FortranGenerator
                                                               method), 18
visit_return() (py- visit_standard_call() (py-
    cropml.transpiler.generators.csharpGenerator.CsharpComporopml.transpiler.generators.javaGenerator.JavaGenerator
    method), 14                                         method), 21
visit_return() (py- visit_standard_call() (py-
    cropml.transpiler.generators.csharpGenerator.CsharpGeneratropml.transpiler.generators.pythonGenerator.PythonGenerator
    method), 15                                         method), 23
visit_return() (py- visit_standard_method_call() (py-
    cropml.transpiler.generators.javaGenerator.JavaCompo
    method), 19                                         cropml.transpiler.generators.checkGenerator.CheckGenerator
                                                               method), 13
visit_return() (py- visit_standard_method_call() (py-
    cropml.transpiler.generators.javaGenerator.JavaGenerator
    method), 21                                         cropml.transpiler.generators.csharpGenerator.CsharpGenerator
                                                               method), 15
visit_return() (py- visit_standard_method_call() (py-
    cropml.transpiler.generators.simplaceGenerator.SimplaceGene
    method), 24                                         cropml.transpiler.generators.fortranGenerator.FortranGenerator
                                                               method), 18
visit_returnstatnode() (py- visit_standard_method_call() (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 30                                         cropml.transpiler.generators.javaGenerator.JavaGenerator
                                                               method), 21
visit_simpleCall() (py- visit_standard_method_call() (py-
    cropml.transpiler.codeGenerator.CodeGenerator
    method), 32                                         cropml.transpiler.generators.pythonGenerator.PythonGenerator
                                                               method), 23
visit_simplecallnode() (py- visit_statlistnode() (py-
    cropml.transpiler.ast_transform.AstTransformer
    method), 30                                         cropml.transpiler.ast_transform.AstTransformer
                                                               method), 30
visit_singleassignmentnode() (py- visit_str() (pycropml.transpiler.generators.checkGenerator.CheckGen
    cropml.transpiler.ast_transform.AstTransformer
    method), 13                                         method), 13

```

```

visit_str() (pycrocml.transpiler.generators.csharpGenerator.CsharpCodeGenerator
    method), 15                                         visit_tuple_decl()                               (py-
visit_str() (pycrocml.transpiler.generators.fortranGenerator.FortranCodeGenerator
    method), 19                                         visit_tuple_decl()                               (py-
visit_str() (pycrocml.transpiler.generators.javaGenerator.JavaGenerator
    method), 21                                         visit_tuple_decl()                               (py-
visit_str() (pycrocml.transpiler.generators.pythonGenerator.PythonCodeGenerator
    method), 23                                         visit_tuplenode()                                (py-
visit_str_decl() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator
    method), 15                                         visit_unary_op()                                (py-
visit_str_decl() (py-                                         cropml.transpiler.generators.checkGenerator.CheckGenerator
    cropml.transpiler.generators.csharpGenerator.CsharpTrans
    method), 16                                         visit_unary_op()                                (py-
visit_str_decl() (py-                                         cropml.transpiler.generators.csharpGenerator.CsharpGenerator
    cropml.transpiler.generators.fortranGenerator.FortranGenerat
    method), 19                                         visit_unary_op()                                (py-
visit_str_decl() (py-                                         cropml.transpiler.generators.fortranGenerator.FortranGenerator
    cropml.transpiler.generators.javaGenerator.JavaGenerator
    method), 21                                         visit_unary_op()                                (py-
visit_str_decl() (py-                                         cropml.transpiler.generators.javaGenerator.JavaGenerator
    cropml.transpiler.generators.javaTrans
    method), 21                                         visit_unary_op()                                (py-
visit_stringnode() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.ast_transform.AstTransformer
    method), 31                                         visit_unaryminusnode()                            (py-
visit_subnode() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.ast_transform.AstTransformer
    method), 31                                         visit_unaryplusnode()                            (py-
visit_subroutine() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.generators.fortranGenerator.FortranGenerat
    method), 19                                         visit_unicodenode()                            (py-
visit_tab() (pycrocml.transpiler.generators.fortranGenerator.FortranCodeGenerator
    method), 19                                         visit_while_statnode()                           (py-
visit_top_level() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.ast_transform.AstTransformer
    method), 31                                         visit_while_statement()                          (py-
visit_tuple() (py-                                         cropml.transpiler.generators.checkGenerator.CheckGenerator
    cropml.transpiler.generators.checkGenerator.CheckGenerator
    method), 13                                         visit_while_statement()                          (py-
visit_tuple() (py-                                         cropml.transpiler.generators.csharpGenerator.CsharpGenerator
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator
    method), 15                                         visit_while_statement()                          (py-
visit_tuple() (py-                                         cropml.transpiler.generators.fortranGenerator.FortranGenerator
    cropml.transpiler.generators.javaGenerator.JavaGenerator
    method), 21                                         visit_while_statement()                          (py-
visit_tuple() (py-                                         cropml.transpiler.generators.javaGenerator.JavaGenerator
    cropml.transpiler.generators.javaTrans
    method), 23                                         visit_while_statement()                          (py-
visit_tuple_decl() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.generators.csharpGenerator.CsharpGenerator
    method), 15                                         visit_whilestatnode()                           (py-
visit_tuple_decl() (py-                                         cropml.transpiler.ast_transform.AstTransformer
    cropml.transpiler.generators.csharpGenerator.CsharpTrans
    method), 31

```

## W

wralea\_template (py-  
    *cropml.package.PyPackageWriter attribute*),  
    41  
wrapper () (pycropml.transpiler.generators.csharpGenerator.CsharpCompo  
    *nethod*), 14  
wrapper () (pycropml.transpiler.generators.siriusGenerator.SiriusCompo  
    *nethod*), 24  
write () (pycropml.transpiler.codeGenerator.CodeGenerator  
    *nethod*), 32  
write () (pycropml.writeTest.WriteTest method), 50  
write\_png () (pycropml.topology.Topology method),  
    49  
write\_tests () (py-  
    *cropml.render\_csharp.Model2Package method*), 44  
write\_tests () (py-  
    *cropml.render\_cyml.Model2Package method*),  
    45  
write\_tests () (py-  
    *cropml.render\_fortran.Model2Package method*), 45  
write\_tests () (py-  
    *cropml.render\_java.Model2Package method*),  
    45  
write\_tests () (py-  
    *cropml.render\_python.Model2Package method*), 48  
write\_tests () (pycropml.render\_R.Model2Package  
    *nethod*), 43  
write\_tests () (py-  
    *cropml.writeTest\_f90.Model2Package method*),  
    51  
write\_wralea () (py-  
    *cropml.package.PyPackageWriter method*),  
    41  
write\_xml () (pycropml.topology.Topology method),  
    49  
WriteTest (*class in pycropml.writeTest*), 50

## X

XmlToWF (*class in pycropml.xml2wf*), 51

## Y

y (pycropml.transpiler.pseudo\_tree.Node attribute), 34