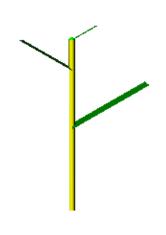
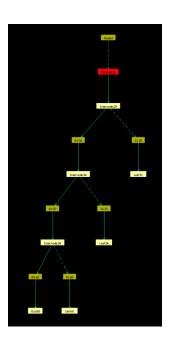
XL language: Queries & Co

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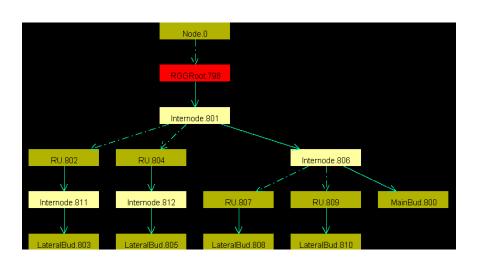




Everything is represented as a graph...

- Plant structure described by a graph composed of nodes (plant organs), connected via edges (relations between organs)
- We can search for a specific pattern in the graph and collect information







Graph operators

- Queriesto search for a specific pattern
- Aggregators to collect values
- Analyse relations between nodes

Find:

all leaves
total leaf area
stem length
first internode below bud
etc.



Graph query

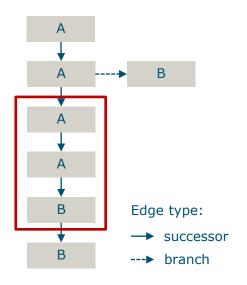
XL syntax: enclosed in asterisked parentheses (* *)

■ The elements are given in their expected order, e.g.:

(* A A B *)

searches for a subgraph that consists of a sequence of nodes of the types A A B, connected by *successor* edges

([blank] – successor edge notation)





Query: Examples

- Find all nodes B, connected to A with a branch edge (+>)
- Find all internodes
- ... and print them out

A query can contain a condition

- Find all newly created leaves (with age 0)
- Search for all leaves with length > 0.05
- Find all pairs of F-s with distance < 1</p>

```
(*A +> B *)
(* Internode *)
println((* Internode *));
(* 1:Leaf, (1[age] == 0) *)
(* 1:Leaf, (1[length] > 0.05) *)
(* f:F, g:F, ((f != g) && (distance(f, g) < 1)) *)
```



Aggregators

- Collect multiple values when traversing the graph and return one single value as a result
- Standard aggregate operations:

count, sum, empty, exist, forall, first, last, max, min, mean, selectRandomly, selectWhereMin, selectWhereMax, ...

Applied to the results of a query:

```
count( (* Leaf *) )
```



Aggregators: Examples

• Get stem length:

Search for all internodes and sum up their lengths

Get total leaf area:

Search for all leaves and sum up their surface areas

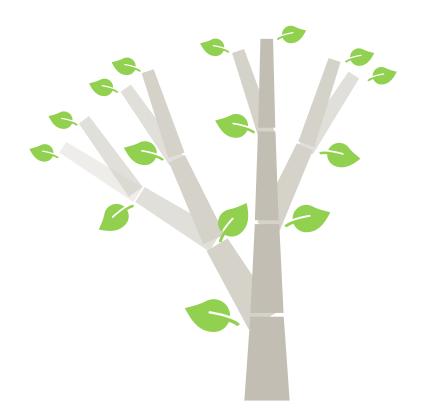
Get total absorbed light by a plant:

Search for all leaves and sum up the amount of light they absorbed

```
sum((* Internode *)[length])
sum((* Leaf *)[area]) I
sum((* Leaf *)[absLight]);
```



```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* 1:Leaf, (1[order] == 1 && 1[rank] == 1) *))
```





```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* l:Leaf, (l[order] == 1 && l[rank] == 1) *))
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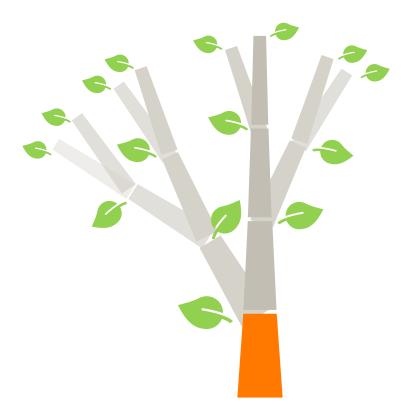


```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* l:Leaf, (l[order] == 1 && l[rank] == 1) *))
```





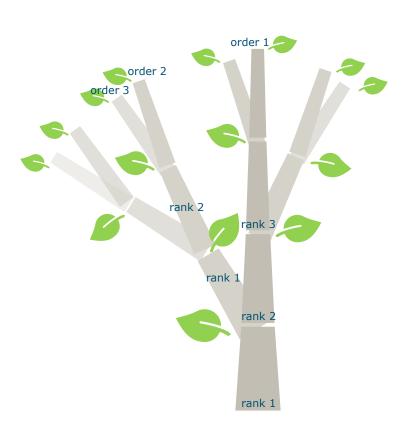
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```





```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* l:Leaf, (l[order] == 1 && l[rank] == 1) *))
```

(order refers to branching order; main stem has order 1; each branch has one order higher; rank refers to organ number within a branch)





```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* 1:Leaf, (1[order] == 1 && 1[rank] == 1) *))
```





```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* 1:Leaf, (1[order] == 3 && 1[rank] == 2) *))
```





```
selectRandomly((* F *))
selectWhereMax((* f:F *), (f[diameter]))
first((* 1:Leaf, (1[order] == 3 && 1[rank] == 2) *))
```

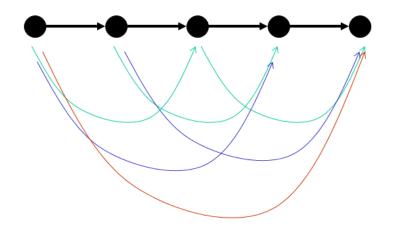




Derived relations

- Relation between nodes connected by several edges (one after the other) of the same type
- Example: find all descendants of some given node that are of type Internode

 "Transitive closure" of the original relation (iteration of edges)





XL syntax for the transitive closure

■ 1-to-n repetitions: +

0-to-n repetitions: *

Minimal elements (stop searching once a match has been found):

```
A (-edgetype->)+ :(B)
```

Examples:

Find all the internodes connected to the bud

```
(* Bud (<--)+ Internode *)
```

Find the first internode connected to the bud

```
(* Bud (<--)+ :(Internode) *)
```



Edge types

$$r = relation$$

Edge notation:

| ı | Arbitrary | > | |
|---|---------------|---------|--------|
| ı | Successor | (blank) | or > |
| ı | Branch | +> | or [(|
| | Decomposition | /> | |
| | User-defined | -r-> | |

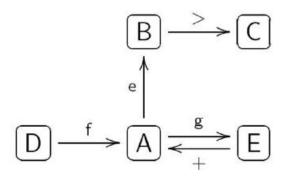
Edge direction:

| Forward | -r-> |
|----------------------|-------|
| Backward | <-r- |
| Forward or backward | -r- |
| Forward and backward | <-r-> |



Graphs in XL

successor: > or (blank)
branch: + or []



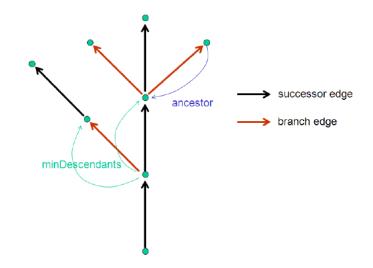
a:A
$$[-e-> B C] [<-f- D] -g-> E [a]$$

multiple representations possible!



Some predefined relations

- ancestor
 - nearest preceding node of a certain node type
- minDescendants
 - nearest successors of a certain node type (nodes of other types are skipped)
- descendants
 - all successors of a certain node type



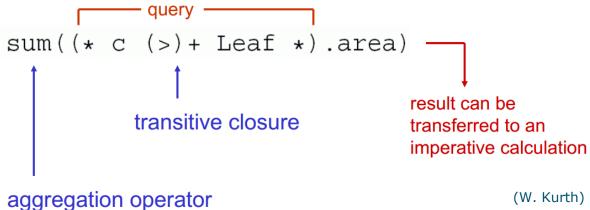
Syntax:

- -ancestor->
- -descendants->
- -minDescendants->



Summary – Graph operations

- Provide possibilities to connect structure and function
- Example: search for all leaves that are successors of node c and sum up their surface areas

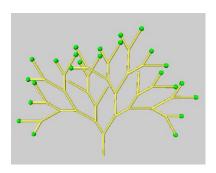


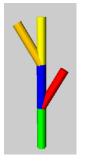


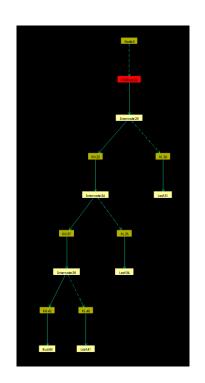
Examples

Tutorial: "tutorials:xl-queries-and-operators [GroIMP wiki]

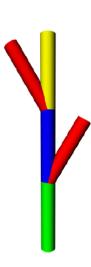
Further examples: binaryTree_distance.gsz nodeRelations.gsz











```
// all modules extend the module Internode
Axiom ==>
    InternodeFirst [RU(30) BranchFirst]
    InternodeSecond [RU(-30) BranchSecond]
    InternodeThird
// queries and their outputs:
(* InternodeFirst -minDescendants-> Internode *)
    BranchFirst
    InternodeSecond
(* InternodeFirst -descendants-> Internode *)
    InternodeSecond
    InternodeThird
    BranchSecond
    BranchFirst
(* InternodeThird -ancestor-> Internode *)
    InternodeSecond
(* InternodeThird (-ancestor->) + Internode *)
    InternodeSecond
    InternodeFirst
```





wiki.grogra.de





