### SBGN-ML

Milestone 1

### Generalities

- SBGN-ML is XML based
  - cf. schema SBGN.xsd

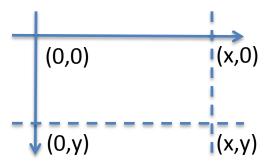
- SBGN-ML files represent SBGN maps
  - PD (Process Description) only in milestone 1

Basic geometry and labels

#### **COMMON TYPES AND ELEMENTS**

### **Point**

- PointAttributes attribute group
  - absolute 2D Cartesian coordinates
    - x (float): horizontal, from left to right
    - y (float): vertical, from top to bottom
  - origin in top-left corner of map
  - no unit (scale free)



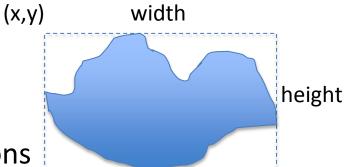
- point element: has PointAttributes
  - <point x="25.4" y="12.3" />

# Bounding box

- bbox element
  - describes a rectangle
    - **PointAttributes** = top left corner
    - width & height (float) = dimensions



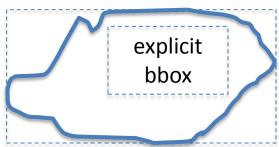
- outer limit of a shape (can be irregular)
- applies to glyph and label elements
- for process nodes = central shape only (not input/output ports)



### Label

- label element = text of a glyph element
  - text attribute (string, mandatory)
    - can be multiline: use line break
  - bbox child (optional) = position
    - the text is centered in the bbox
    - use parent glyph's bbox when omitted
    - represents text size hint when explicit
  - <label text="inherited bbox">





General structure of a SBGN-ML file

#### **HIGH LEVEL ELEMENTS**

### Structure of a SBGN-ML file

- **sbgn** root element
  - single map element: SBGN PD map
    - list of glyph elements (can be empty)
    - list of arc elements (can be empty)

Stand-alone high level SBGN nodes, and sub-nodes

#### **GLYPH ELEMENT**

# class attribute (optional)

- Semantic of the glyph:
  - Shape visual appearance
  - Map syntactic validity
- Entity Pool Nodes (EPN):
  - unspecified entity [DEFAULT]
  - simple chemical
  - macromolecule
  - nucleic acid feature
  - complex
  - simple chemical multimer
  - macromolecule multimer
  - nucleic acid feature multimer
  - complex multimer
  - source and sink
  - perturbing agent

- Process Nodes (PN):
  - process
  - omitted process
  - uncertain process
  - association
  - dissociation
  - phenotype
- Logic Operator Nodes (LON): and, or, not
- Sub-glyphs on Nodes: state variable, unit of information
- Sub-glyphs on Arcs: stoichiometry
- Other glyphs: compartment, submap, tag

### Other attributes

- id attribute (mandatory)
  - alphanumeric identifier, starting with a letter
  - usually meaningless: <glyph id="glyph1234" ...> ... </glyph>
- orientation attribute (optional)
  - Process node: horizontal, vertical
  - Tag: left, right, up, down
  - <glyph class="tag" orientation="right" id="g1">

... </glyph>

Label text here



## Text inside a glyph: label or state

- state (optional): state variables only
  - value attribute
  - variable attribute
- label (optional): all others, except PN and LON
  - <glyph ...> <label text="my label" /> ... </glyph>

#### clone element

- Optional
- Means the glyph has a clone marker
- Can contain a label
  - <glyph class="complex" id="g1">
    ... <clone> <label text="clone label" /> </clone> ...
    </glyph>

# Geometry of a glyph: bbox

- Mandatory
- <glyph id="g1">
   <bbox x="2" y="5" width="67" height="24" />
   ...
   </glyph>
- Cf. definition of bbox

## Children elements: glyph list

- Optional, any size
- <glyph id="g1">

```
...

<glyph id="g1.1"> ... </glyph>

<glyph id="g1.2"> ... </glyph>

...

</glyph>
```

- Examples:
  - Unit of information of a compartment
  - State variable of an EPN
  - Inner EPN-like component of a complex

## Children elements: port list

- Optional, any size
  - 2 port elements required for PN (in & out ends)
- Attributes:
  - PointAttributes absolute 2D Cartesian coordinates
  - id: alphanumeric identifier starting with a letter
- <glyph class="process node" id="g1" orientation="horizontal">

Visible link between two nodes

#### **ARC ELEMENT**

#### class attribute

- Semantic of the arc:
  - > Line visual appearance (usually, decorators at the end)
  - ➤ Map syntactic validity (cf. class of source and target)
- production and consumption arcs
- all types of modification arcs:
  - modulation
  - stimulation
  - catalysis
  - inhibition
  - necessary stimulation
- logic arc
- equivalence arc

### source and target attributes

- Mandatory
- Can refer to the id of a glyph or a port
- <arc class="consumption" source="epn1" target="pn1.1">

• • •

</arc>

## stoichiometry of an arc

- Optional child element
- glyph of class stoichiometry

```
    <arc class="production" source="pn1.2" target="epn2"></arc>
    <glyph class="stoichiometry" id="s1"></arc>
    <label text="3"/></box x="23" y="45" width="10" height="10"></glyph>
    </arc>
```

## Geometric path of an arc

- **start** element
  - PointAttributes: start point of the path
- optional list of next elements
  - PointAttributes: next point in the path
  - point elements (between 0 and 2)
    - control points of the Bezier curve (1: quadratic, 2: cubic)
- end element
  - PointAttributes: end point of the path
  - point elements (between 0 and 2)
    - control points of the curve (1: quadratic, 2: cubic)

# Example of paths (straight lines)

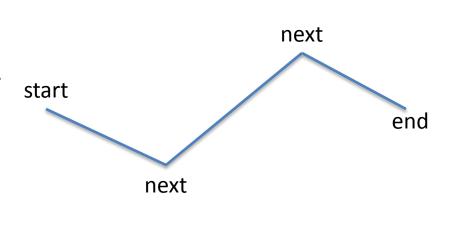
Single straight line

```
<arc ...>
  <start x="10" y="20"/>
  <end x="60" y="20"/>
  </arc>
```



Polyline

```
<arc ...>
  <start x="10" y="20"/>
  <next x="30" y="30"/>
   <next x="50" y="10"/>
   <end x="60" y="20"/>
</arc>
```



# Example of paths (curves)

Quadratic Bezier curve

```
<arc ...>
  <start x="10" y="20"/>
    <end x="60" y="20">
      <point x="34" y="23" />
      </end>
</arc>
```

Cubic Bezier curve

```
<arc ...>
  <start x="10" y="20"/>
    <end x="60" y="20">
      <point x="12" y="23" />
      <point x="34" y="21" />
      </end>
```

# Example of paths (mixed)

```
• <arc ...>
   <start x="10" y="20"/>
   <next x="30" y="30"/>
   <next x="50" y="10">
    <point x="14" y="53">
   </next>
   <end x="60" y="20">
    <point x="12" y="23" />
    <point x="34" y="21" />
   </end>
  </arc>
```