

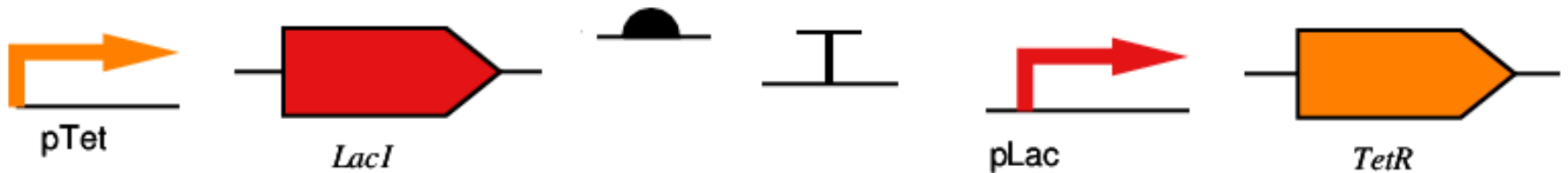
SBOL 2.0: Vision and Structure

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University of Utah

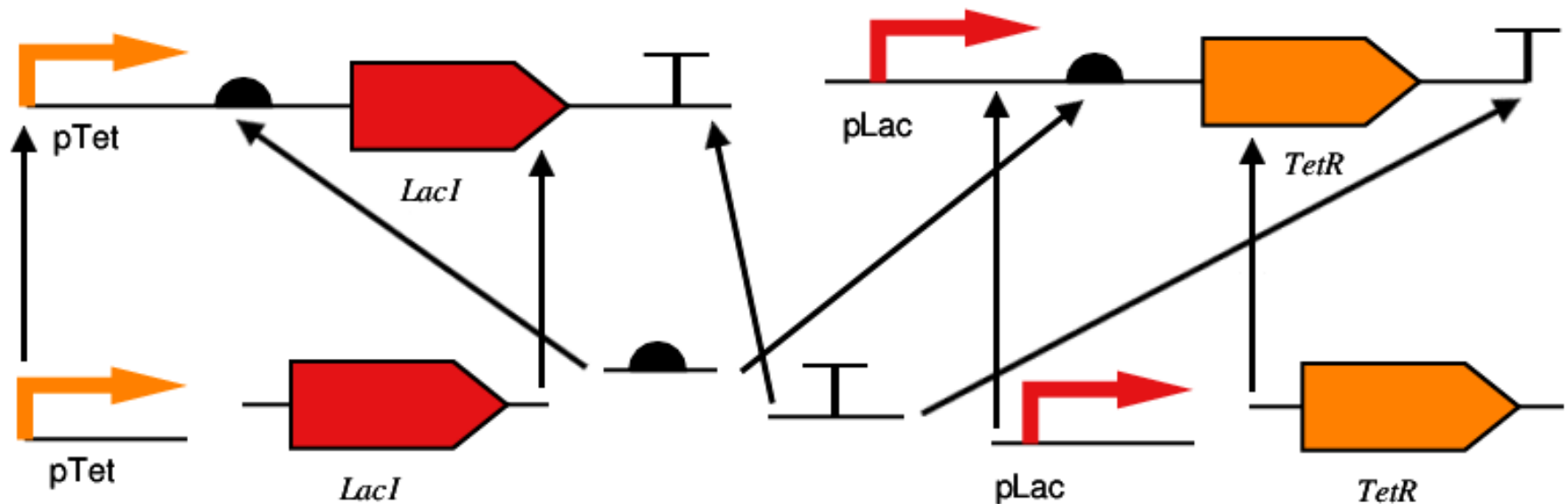
SBOL Workshop 10: UC Berkeley

Current Capabilities of SBOL 1.0



- Specification of DNA components

Current Capabilities of SBOL 1.0

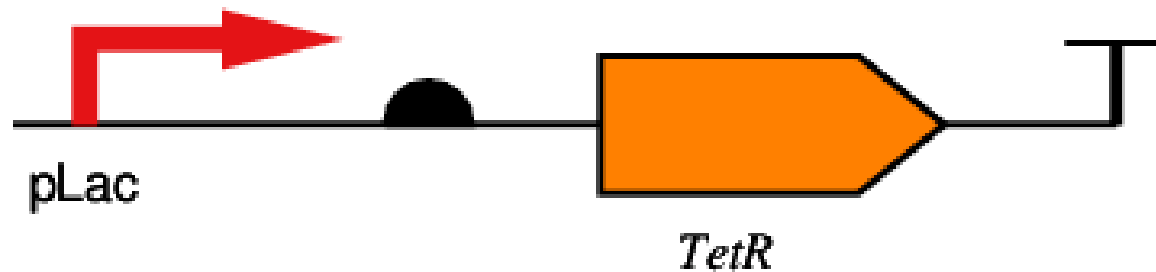


- Specification of DNA components
- Hierarchical composition of DNA components

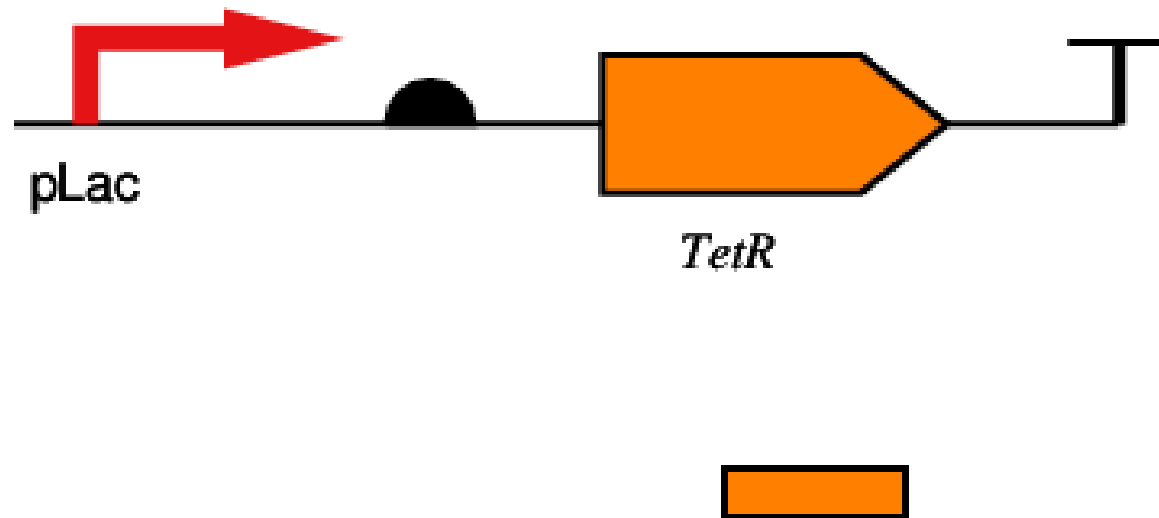
Goals for SBOL 2.0

- Increase the range of biological structure and function that we may specify.
- Provide an extensible basis for composition of functional modules with structural components and other functional modules.

Increasing Structural Range

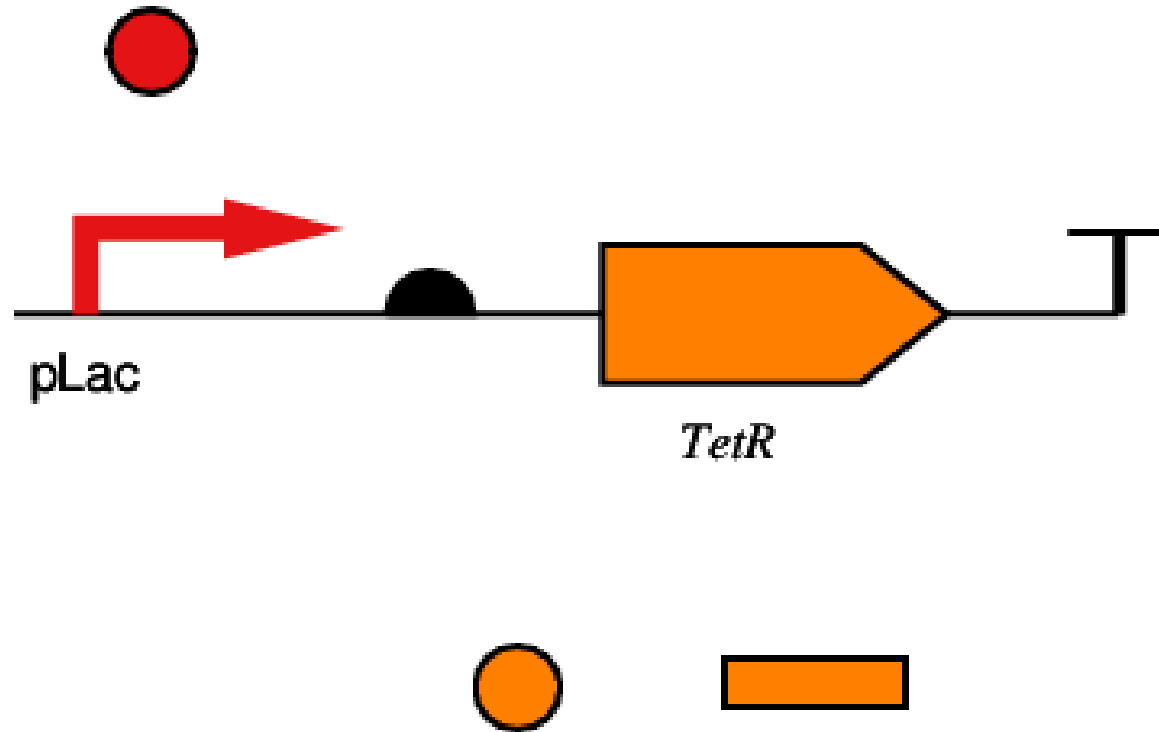


Increasing Structural Range



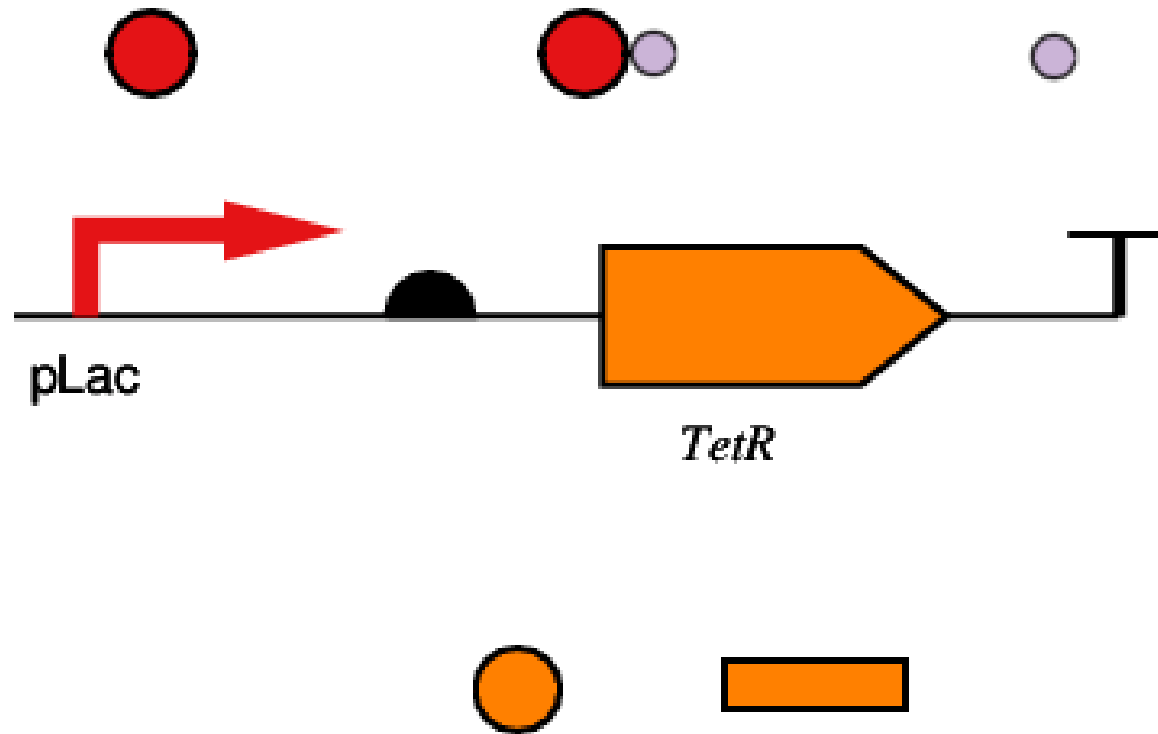
- RNA components (mRNA, tRNA, siRNA)

Increasing Structural Range



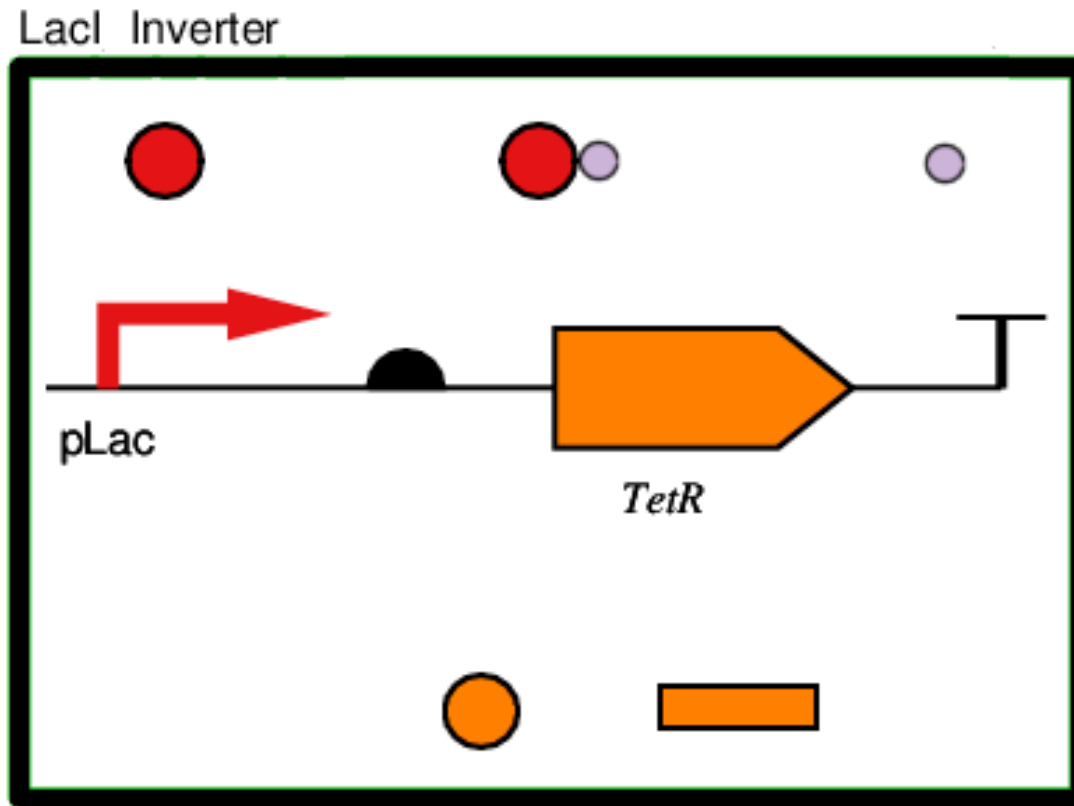
- RNA components
- Protein components (TFs, enzymes)

Increasing Structural Range



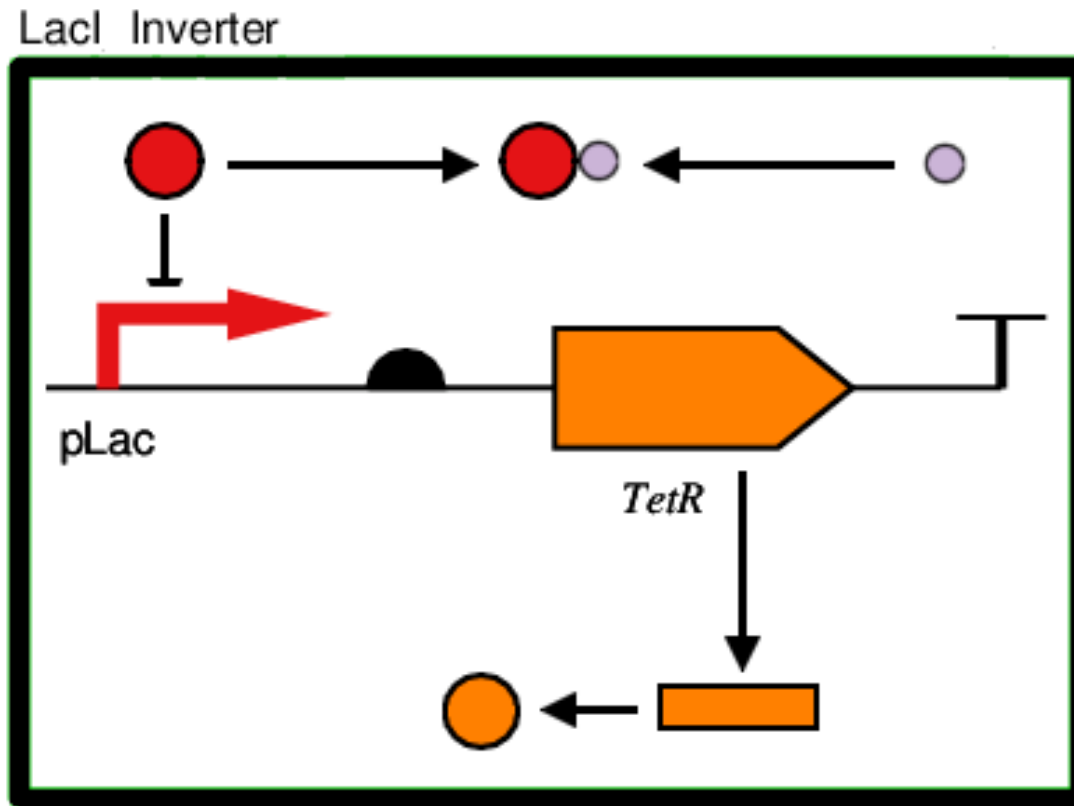
- RNA components
- Protein components
- Other Components (small molecules, complexes, light, pressure, pH, temp)

Increasing Functional Range



- Modules (logic gates, latches, oscillators, sensors, transducers, pathways, cascades)

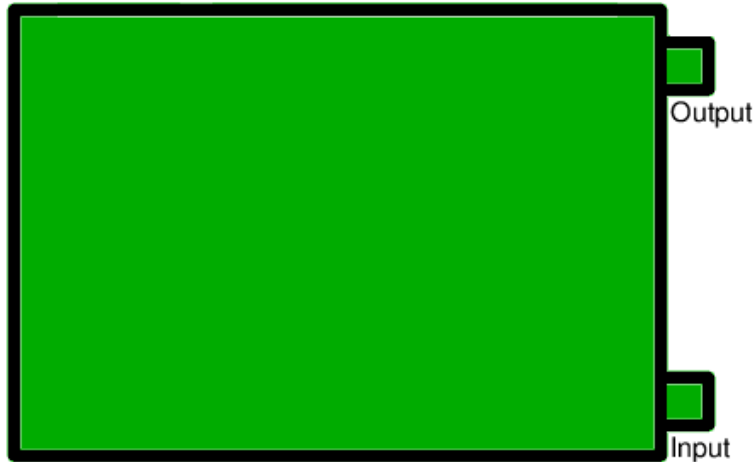
Increasing Functional Range



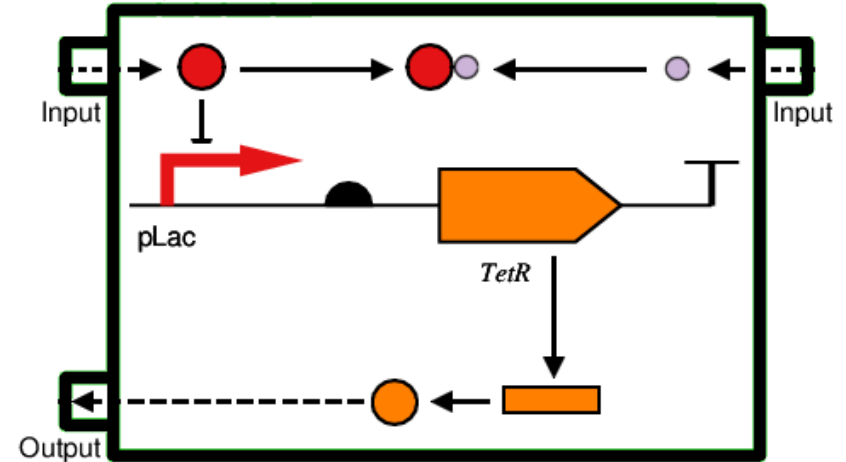
- Modules
- Interactions (activation, repression, complexation, transcription, translation, phosphorylation)

Basis for Composition

TetR Inverter



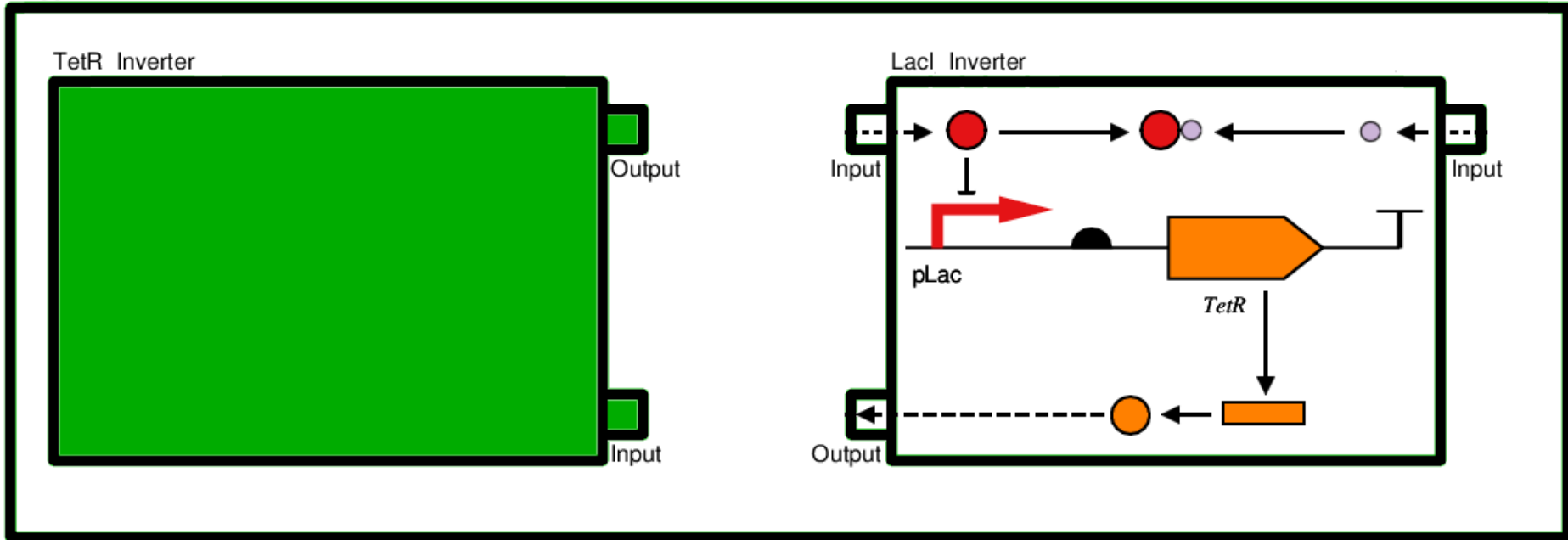
LacI Inverter



- Ports

Basis for Composition

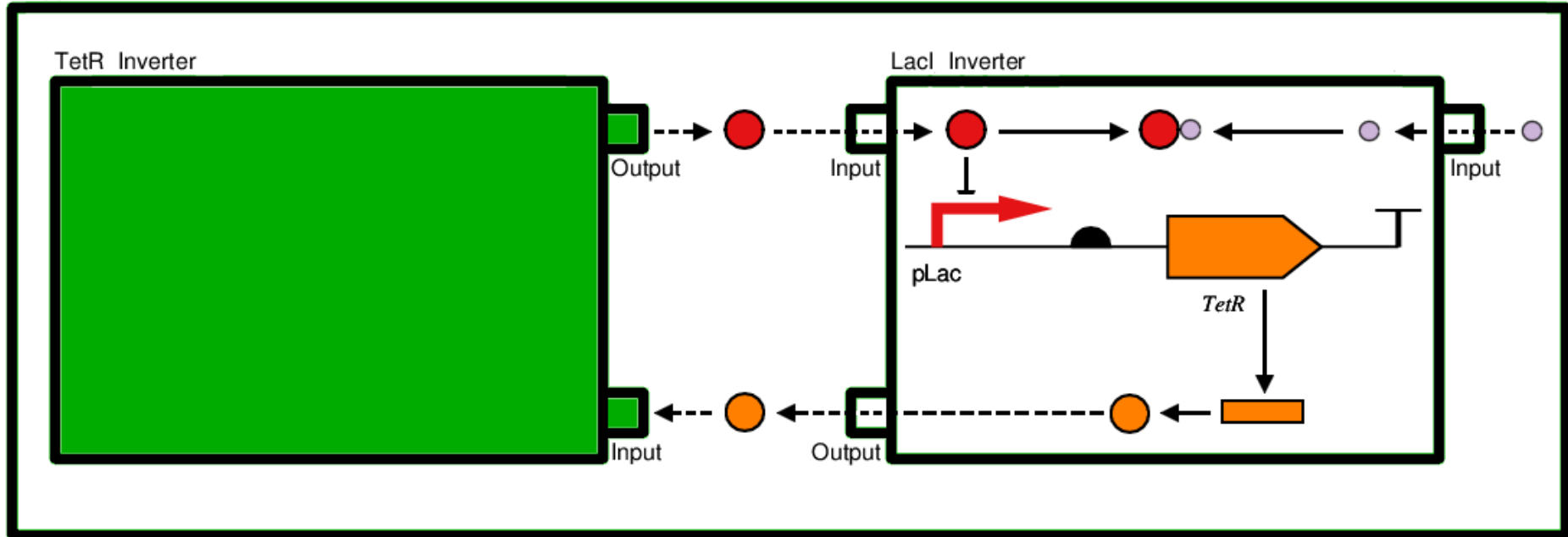
Toggle Switch



- Ports
- Instantiation

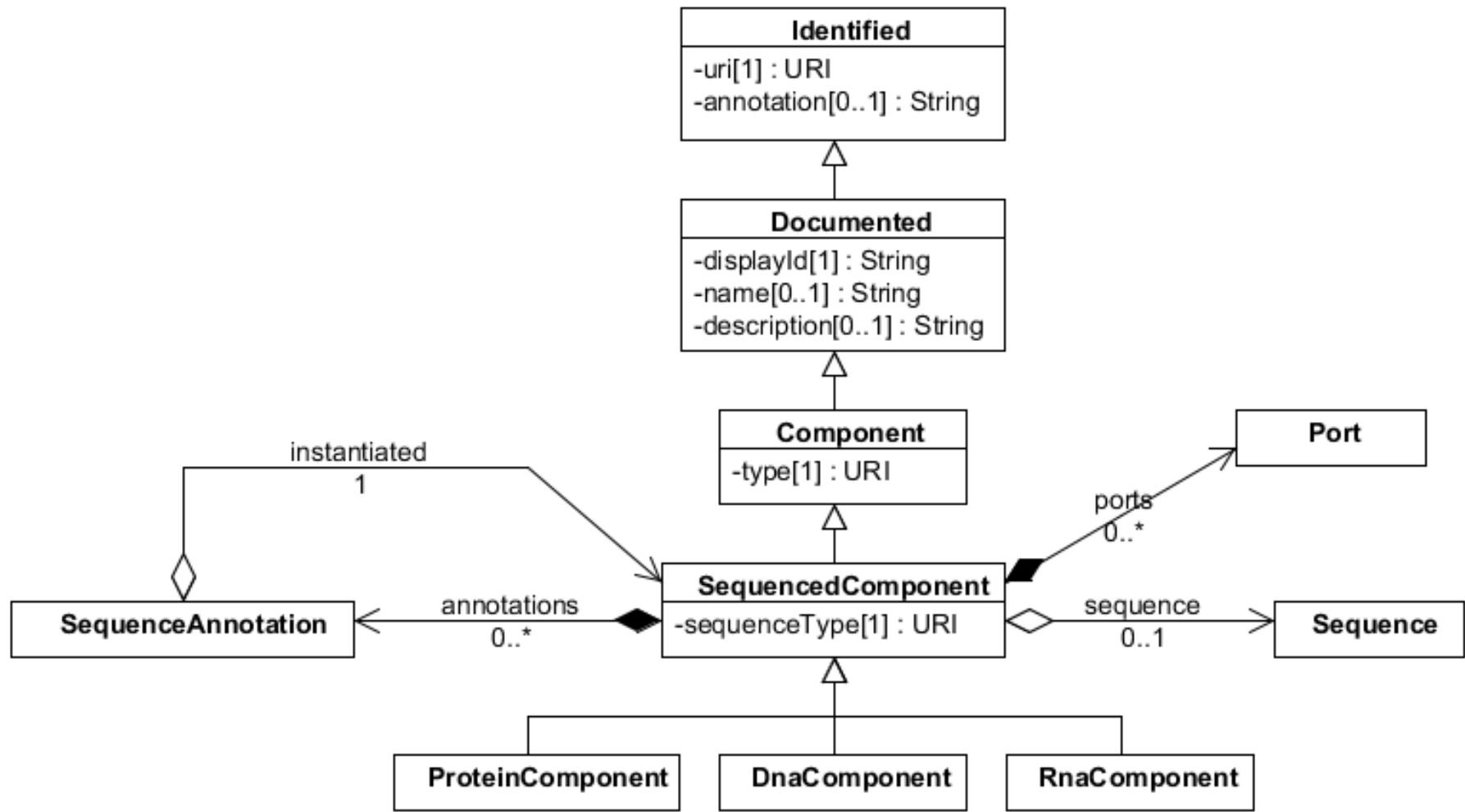
Basis for Functional Composition

Toggle Switch

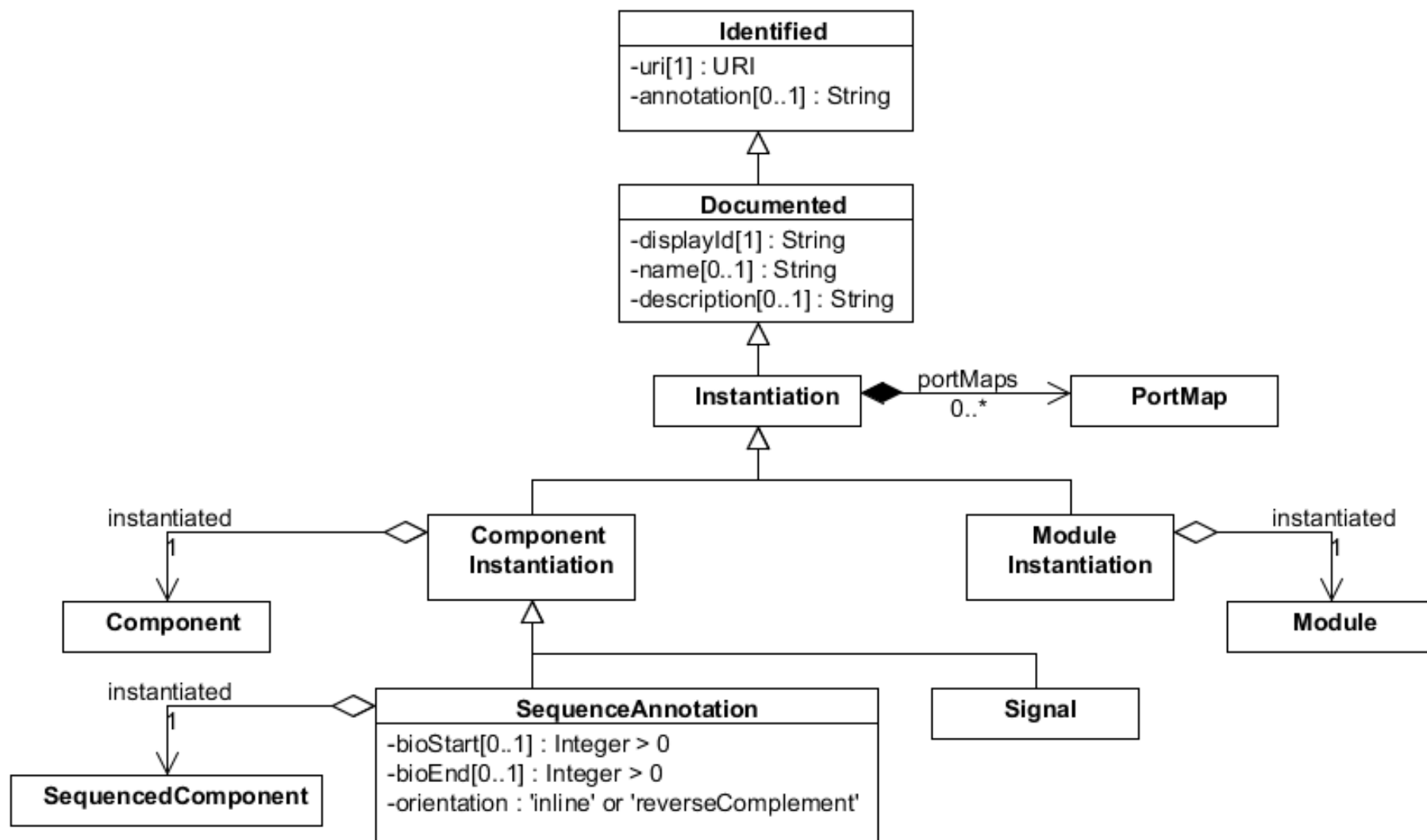


- Ports
- Instantiation
- Port Mapping and Mixed Hierarchy

Data Model: Sequenced Component



Data Model: Instantiation



Example: Components

DnaComponent
-displayId : "BBa_R0010"
-sequenceType : promoter

DnaComponent
-displayId : "BBa_J61120"
-sequenceType : RBS

DnaComponent
-displayId : "BBa_C0040"
-sequenceType : CDS

DnaComponent
-displayId : "ECK120033736"
-sequenceType : terminator

DnaComponent
-displayId : "LacI_Inverter"
-sequenceType : gene

ProteinComponent
-displayId : "LacI"
-sequenceType : transcription factor

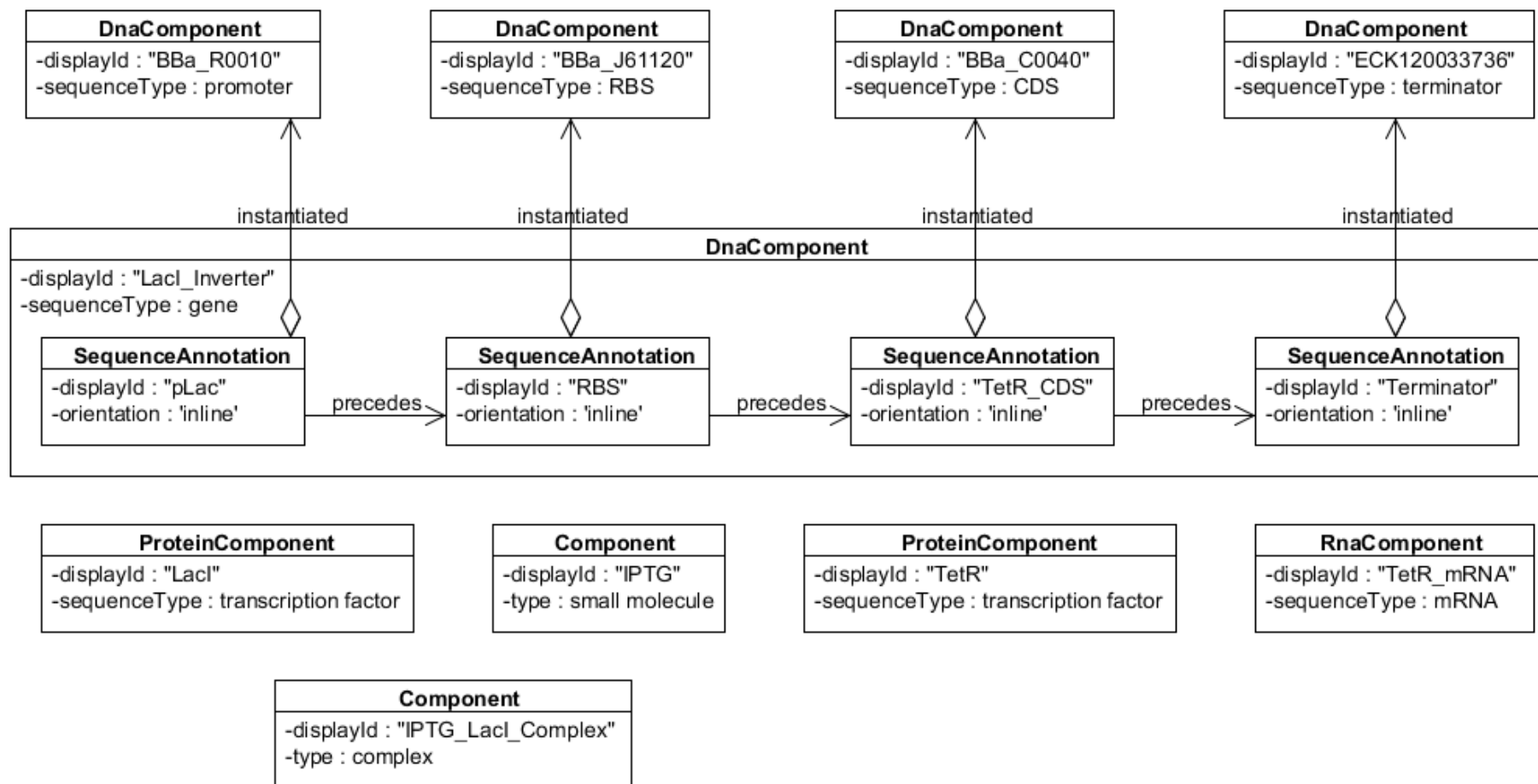
Component
-displayId : "IPTG"
-type : small molecule

ProteinComponent
-displayId : "TetR"
-sequenceType : transcription factor

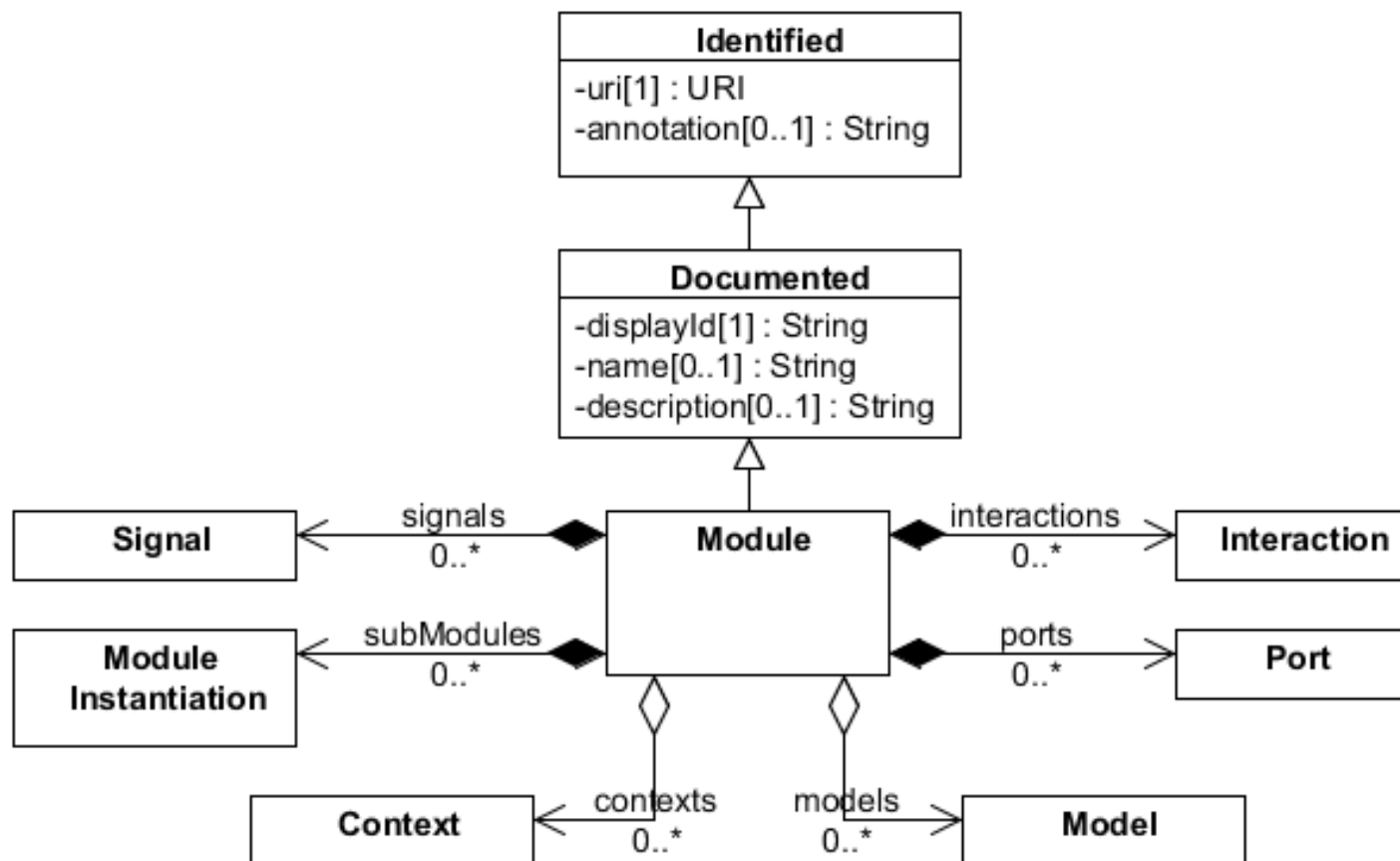
RnaComponent
-displayId : "TetR_mRNA"
-sequenceType : mRNA

Component
-displayId : "IPTG_LacI_Complex"
-type : complex

Example: Sequence Instantiation



Data Model: Module



Example: Module

Component
-displayId : "IPTG_LacI_Complex"
-type : complex

ProteinComponent
-displayId : "TetR"
-sequenceType : transcription factor

RnaComponent
-displayId : "TetR_mRNA"
-sequenceType : mRNA

Module
-displayId : "LacI_Inverter"

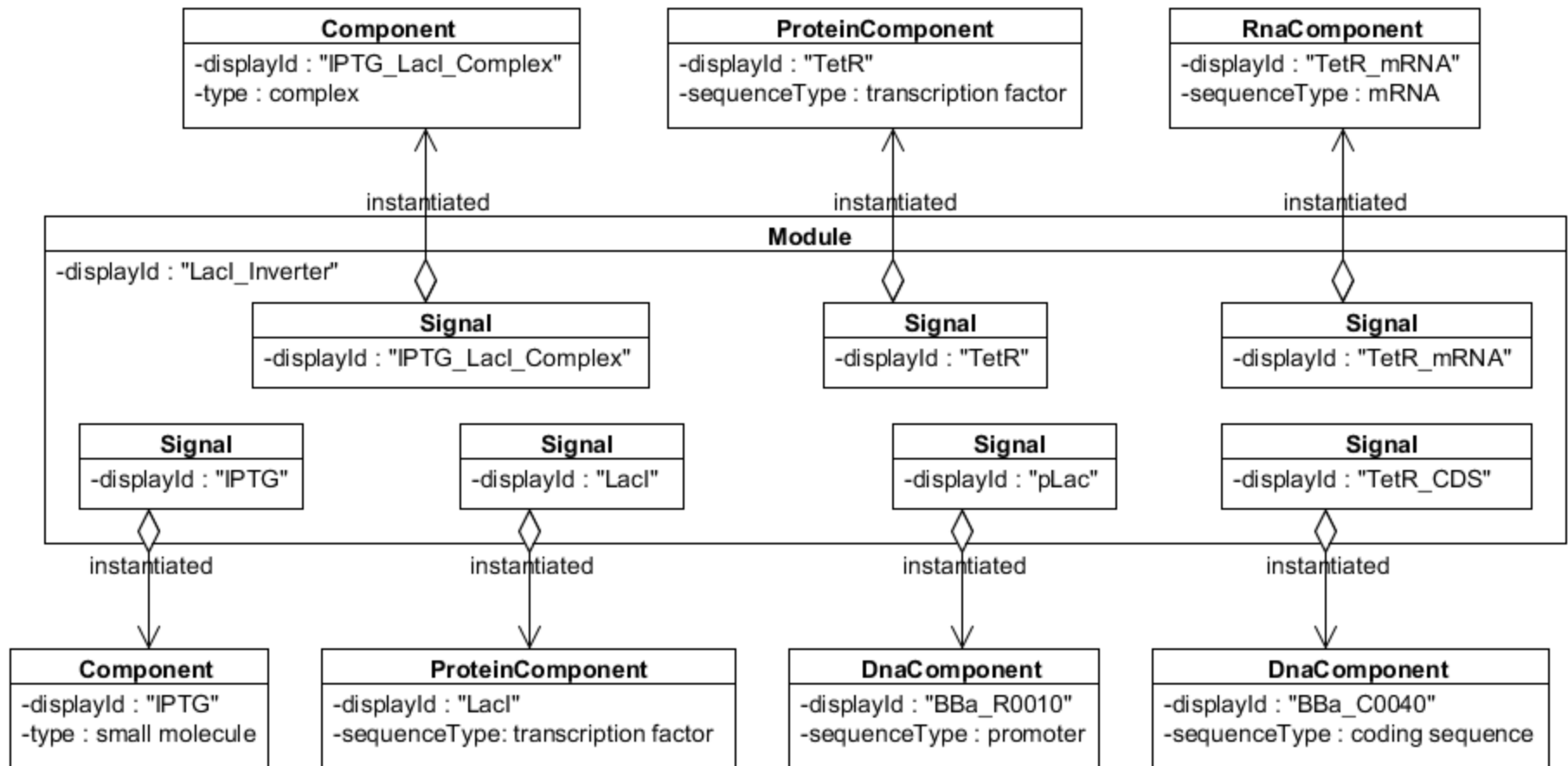
Component
-displayId : "IPTG"
-type : small molecule

ProteinComponent
-displayId : "LacI"
-sequenceType: transcription factor

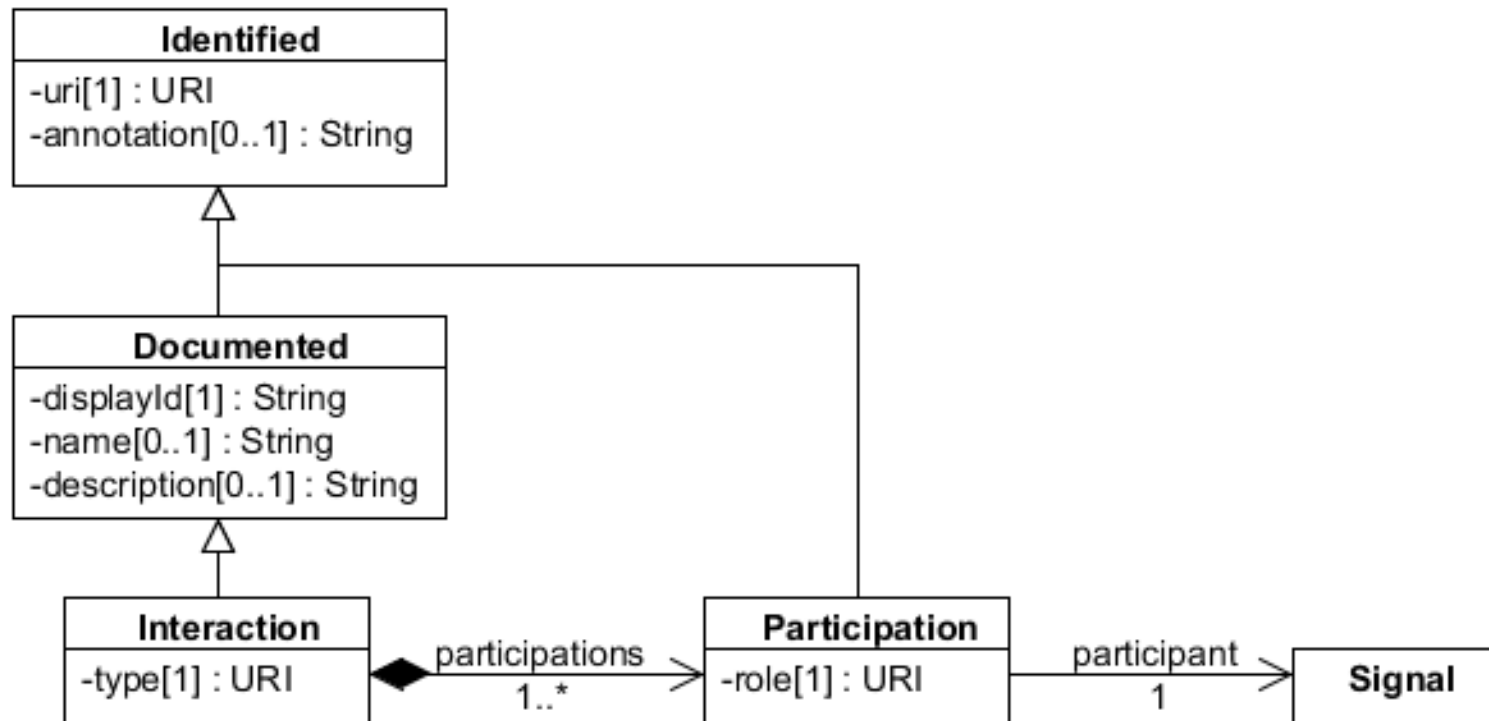
DnaComponent
-displayId : "BBa_R0010"
-sequenceType : promoter

DnaComponent
-displayId : "BBa_C0040"
-sequenceType : coding sequence

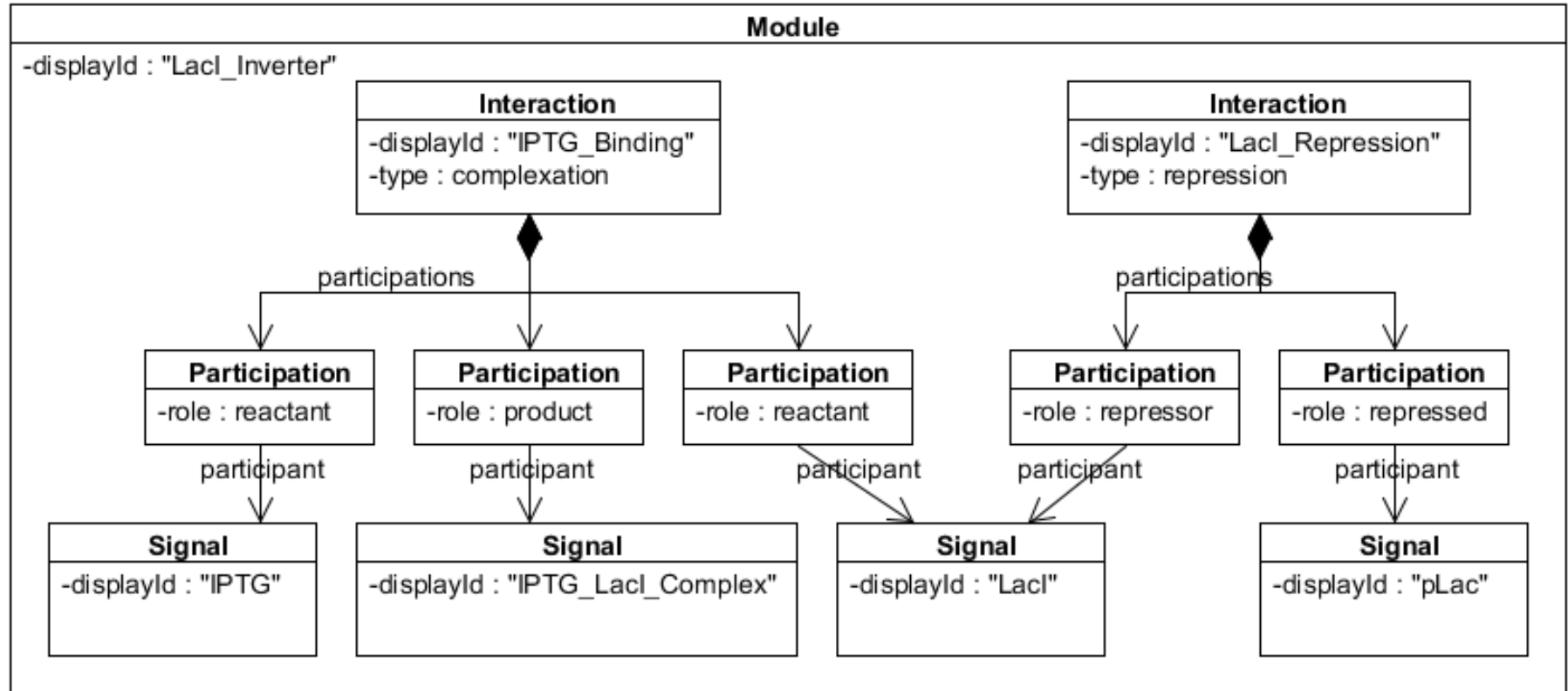
Example: Signal Instantiation



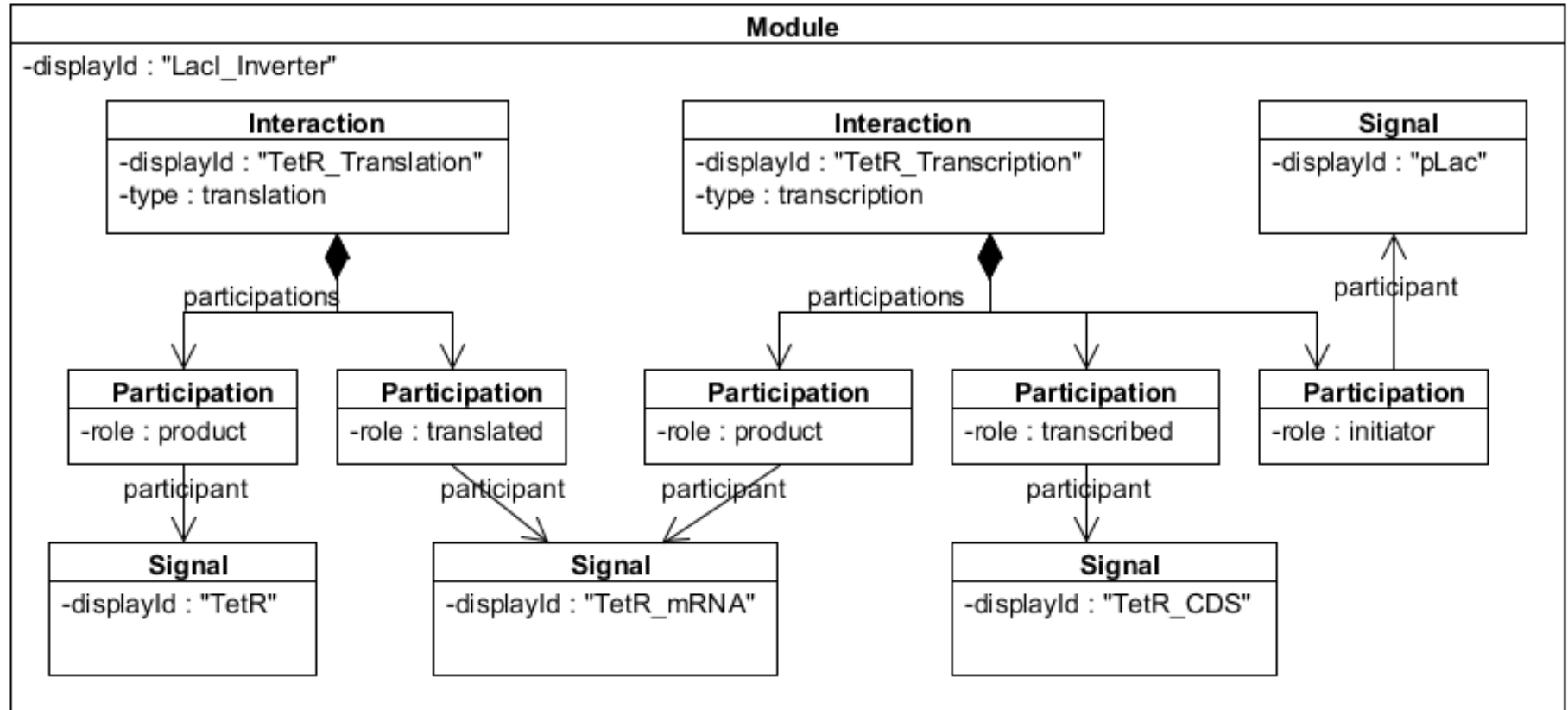
Data Model: Interactions



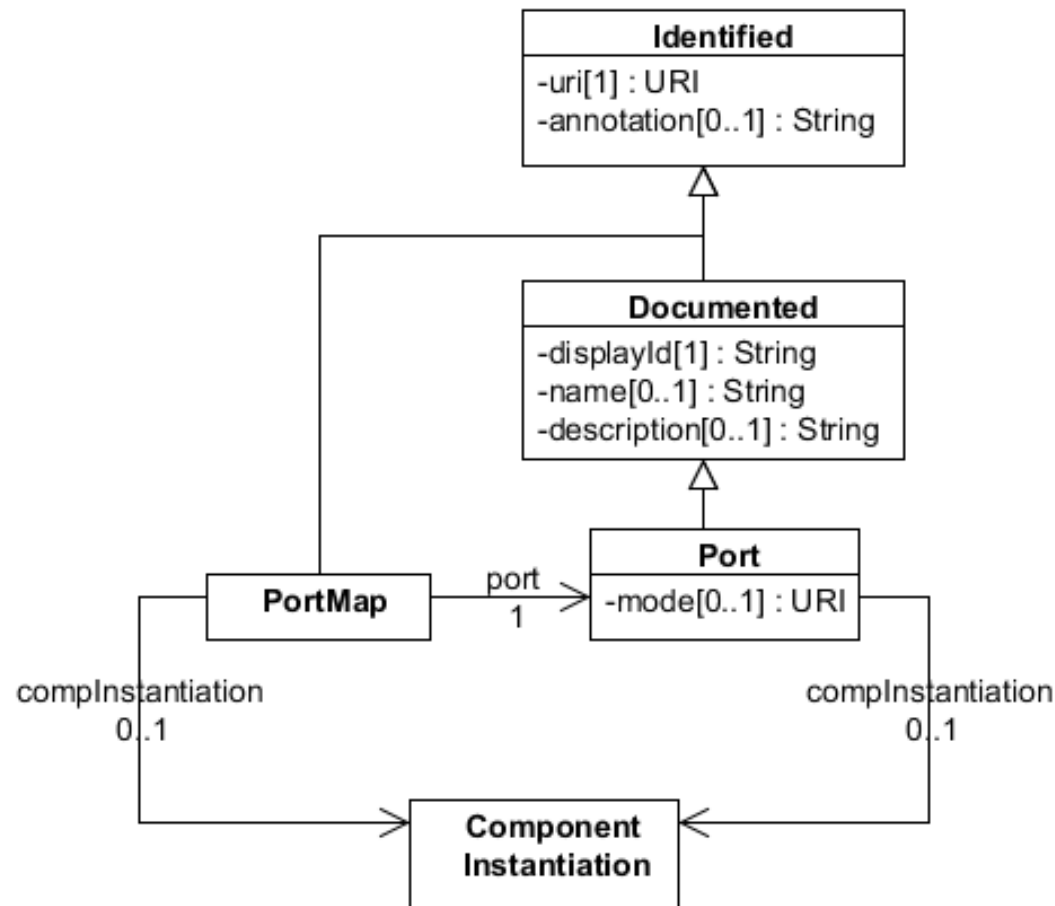
Example: Interactions



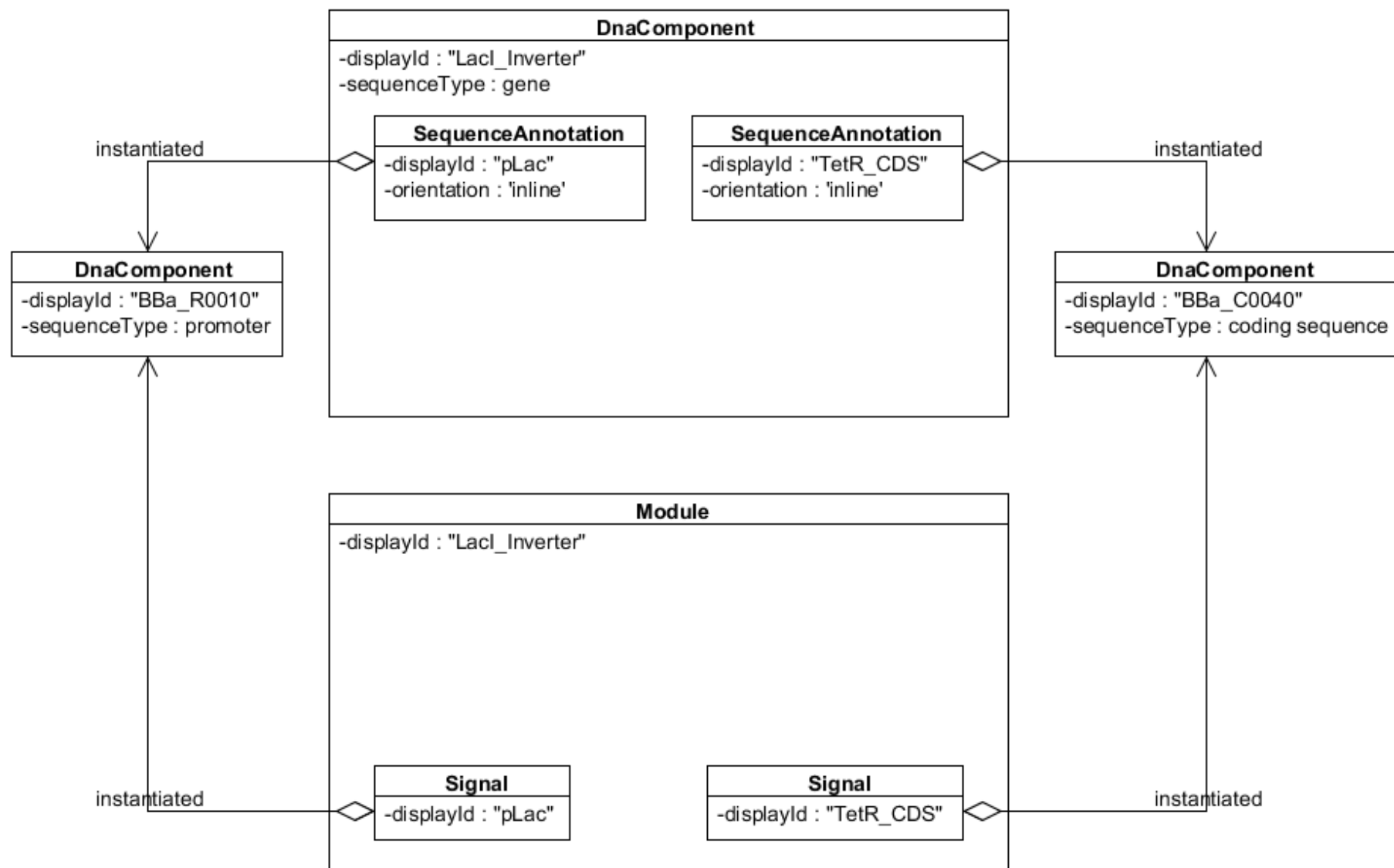
Example: Interactions



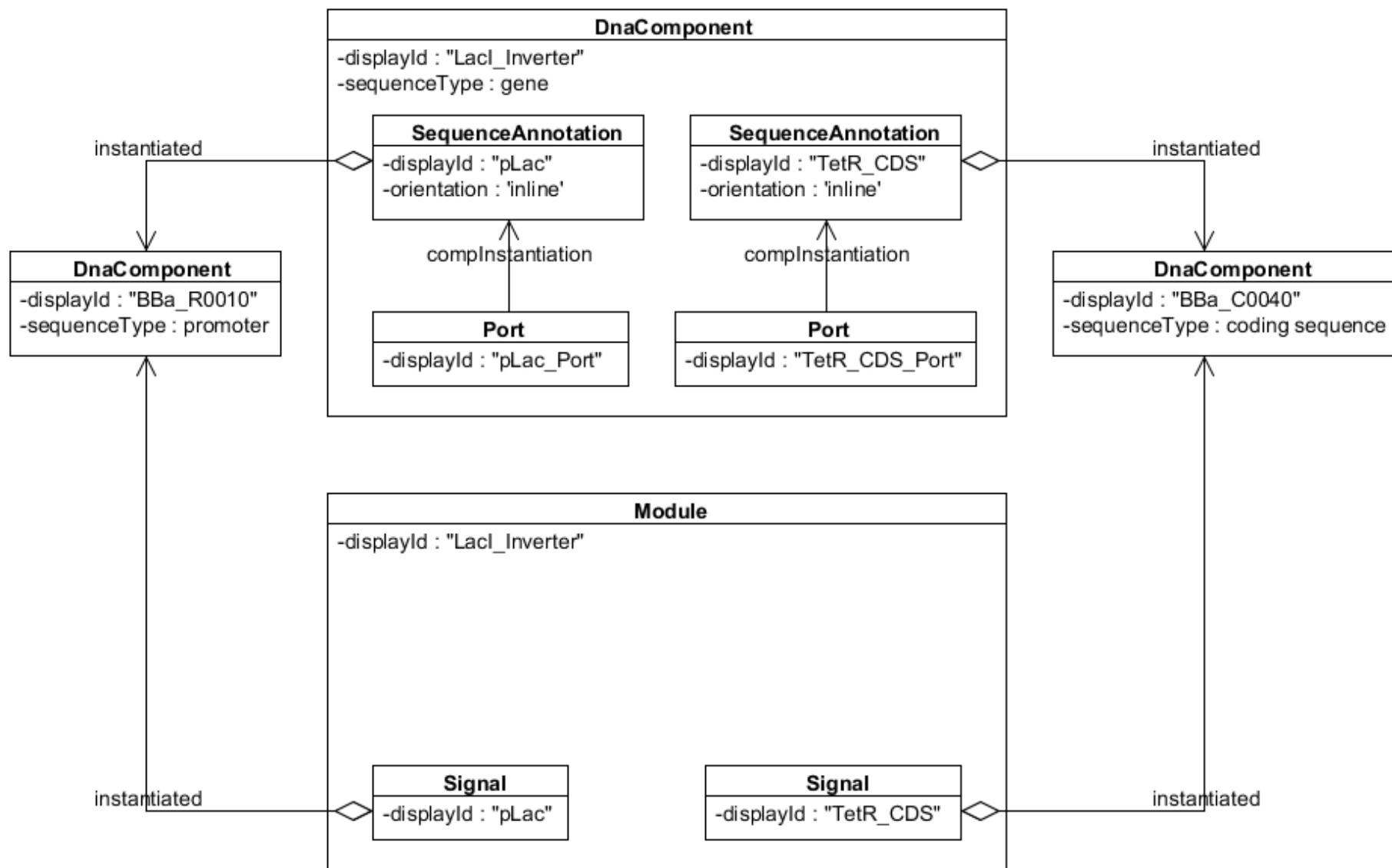
Data Model: Ports and PortMaps



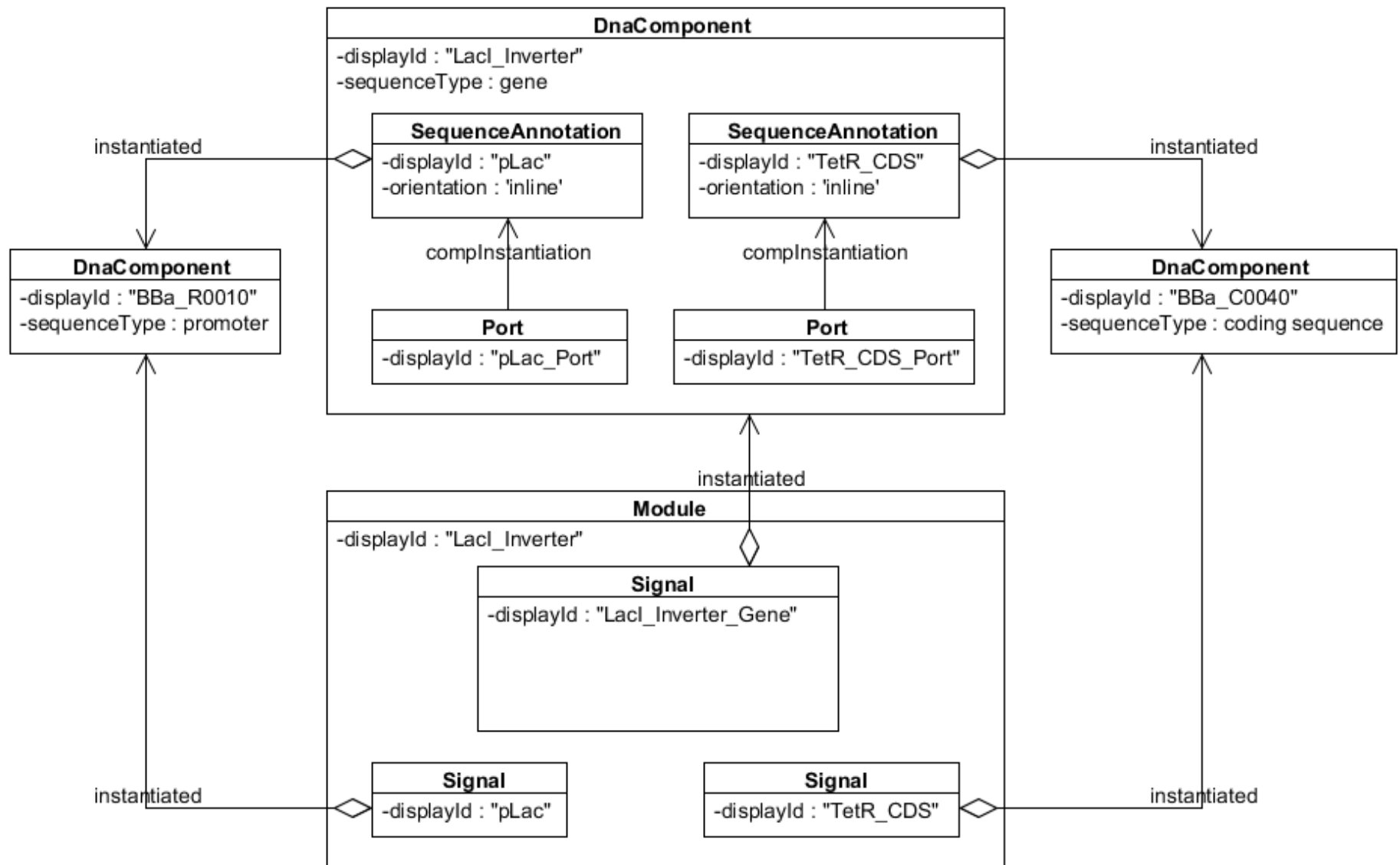
Composing Inverter Component with Module



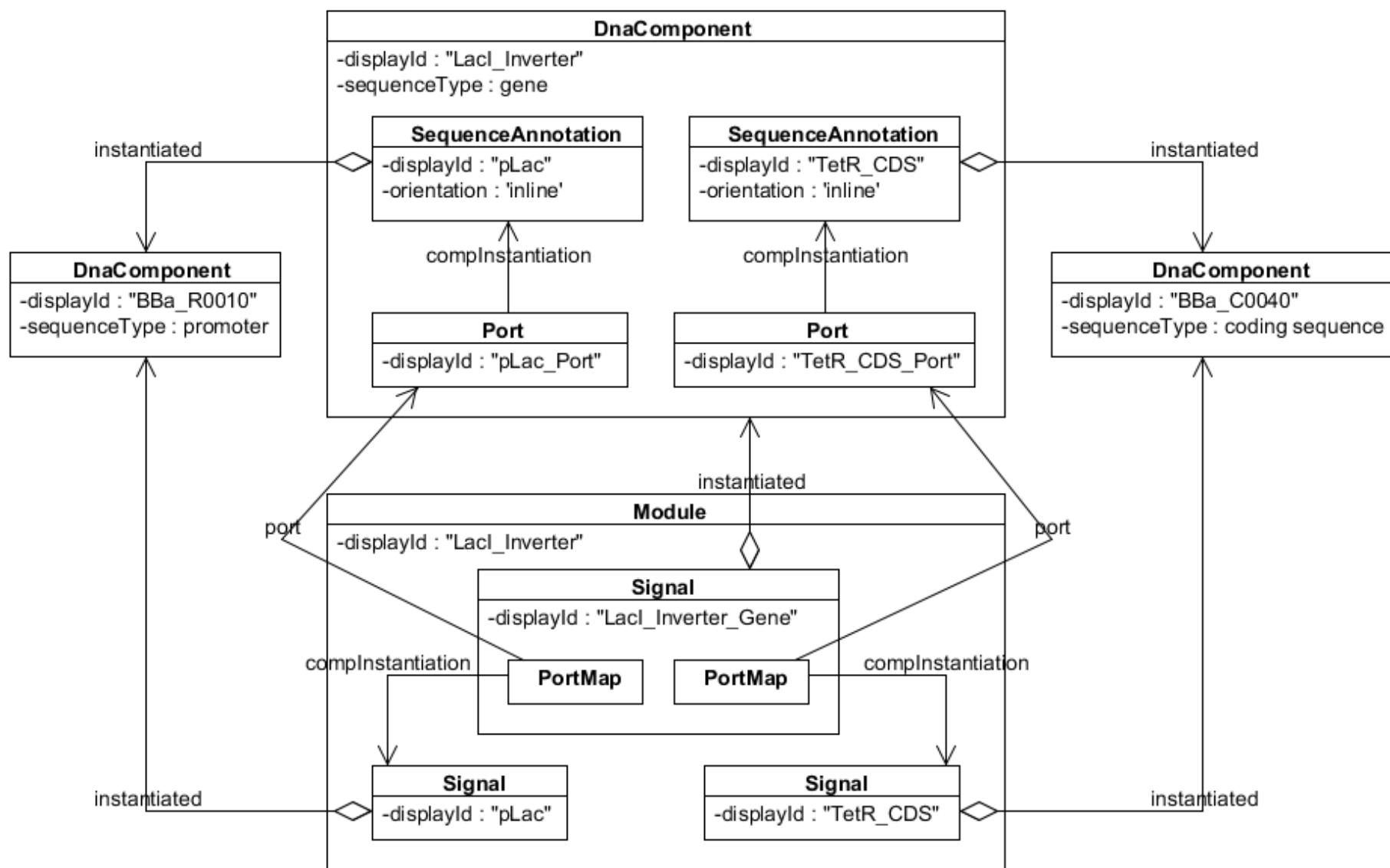
Example: Component Ports



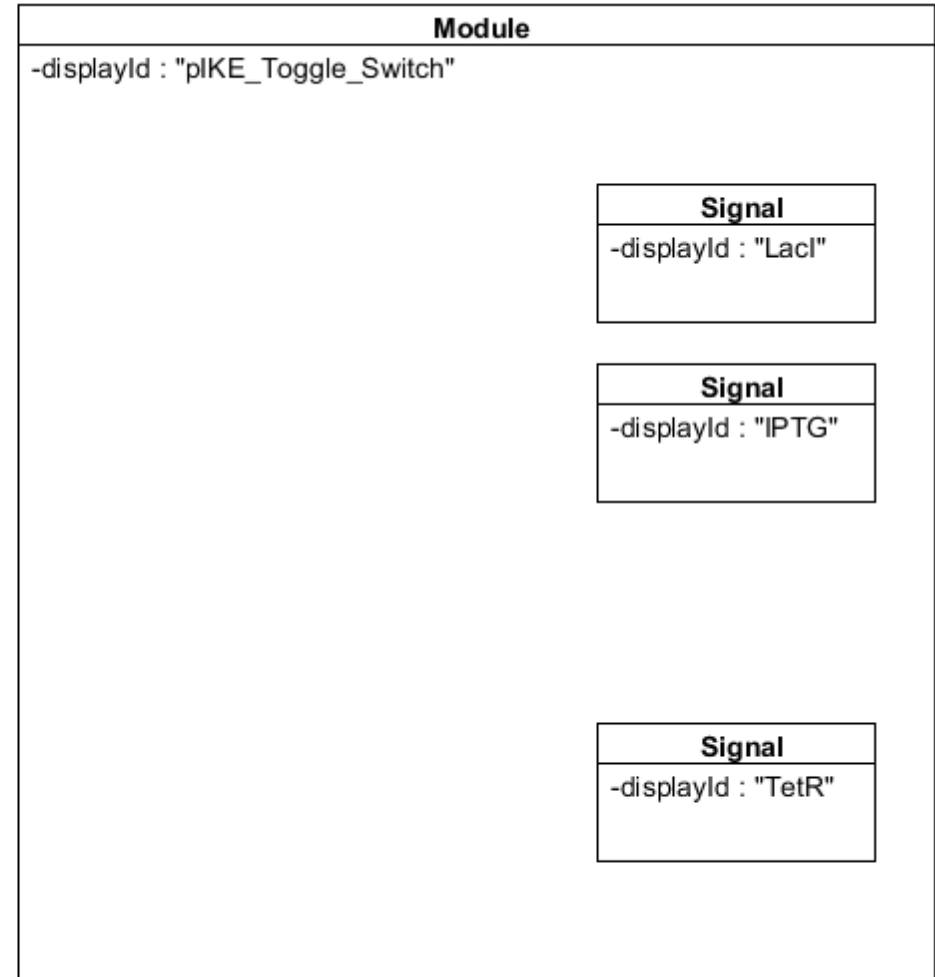
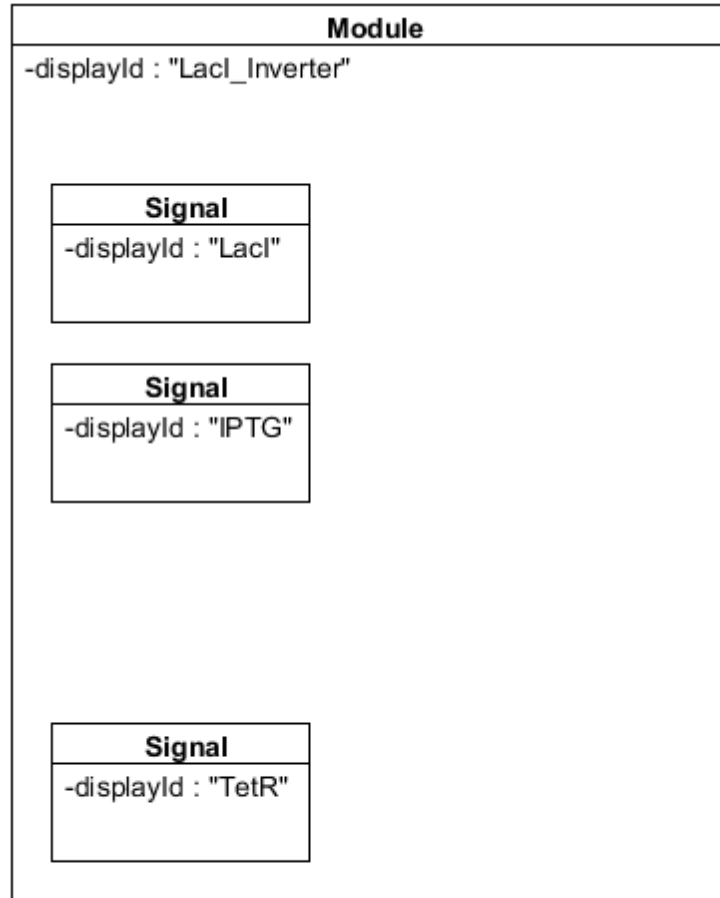
Example: Signal Instantiation



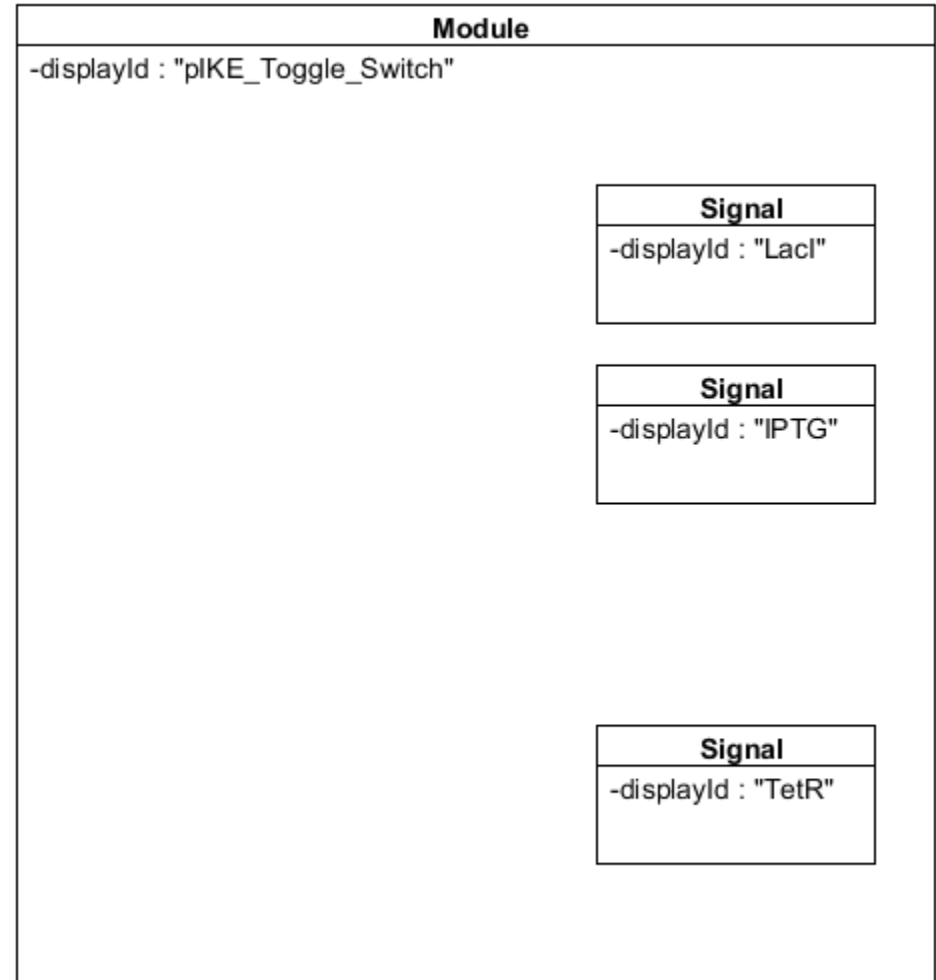
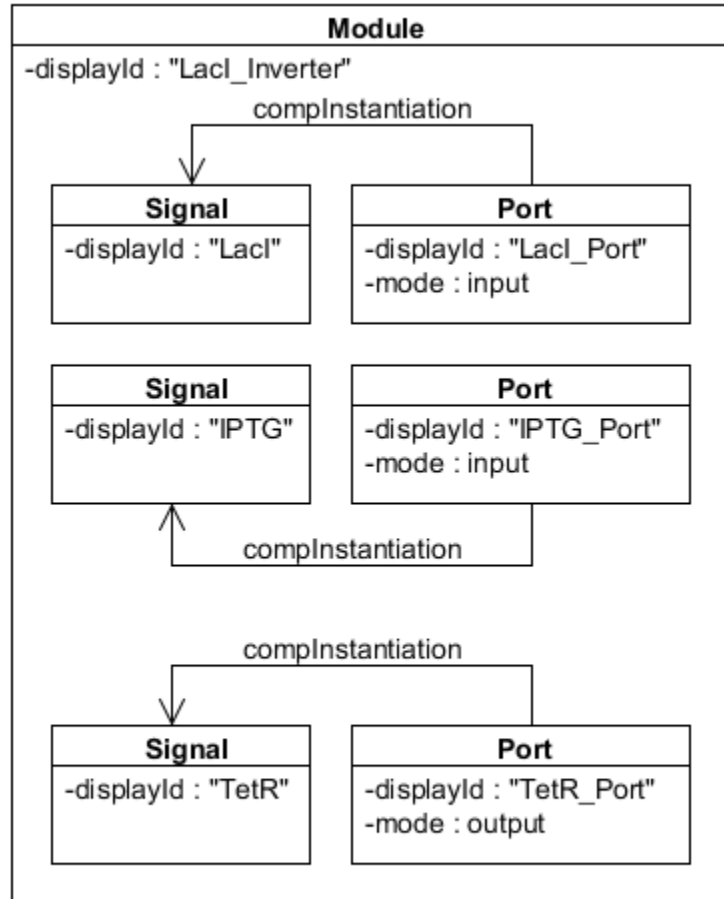
Example: Port Mapping



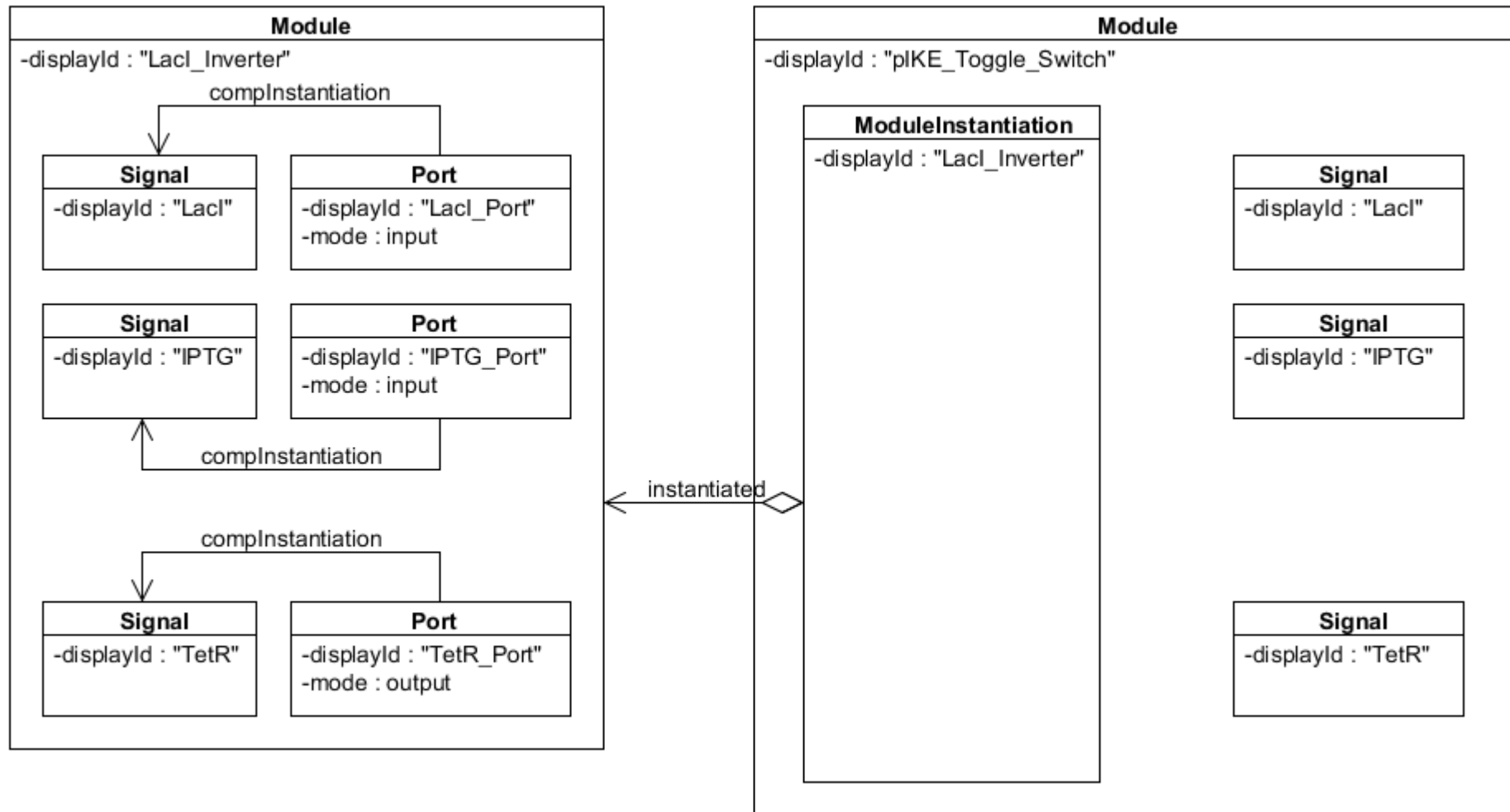
Composing Inverter Module with Toggle Module



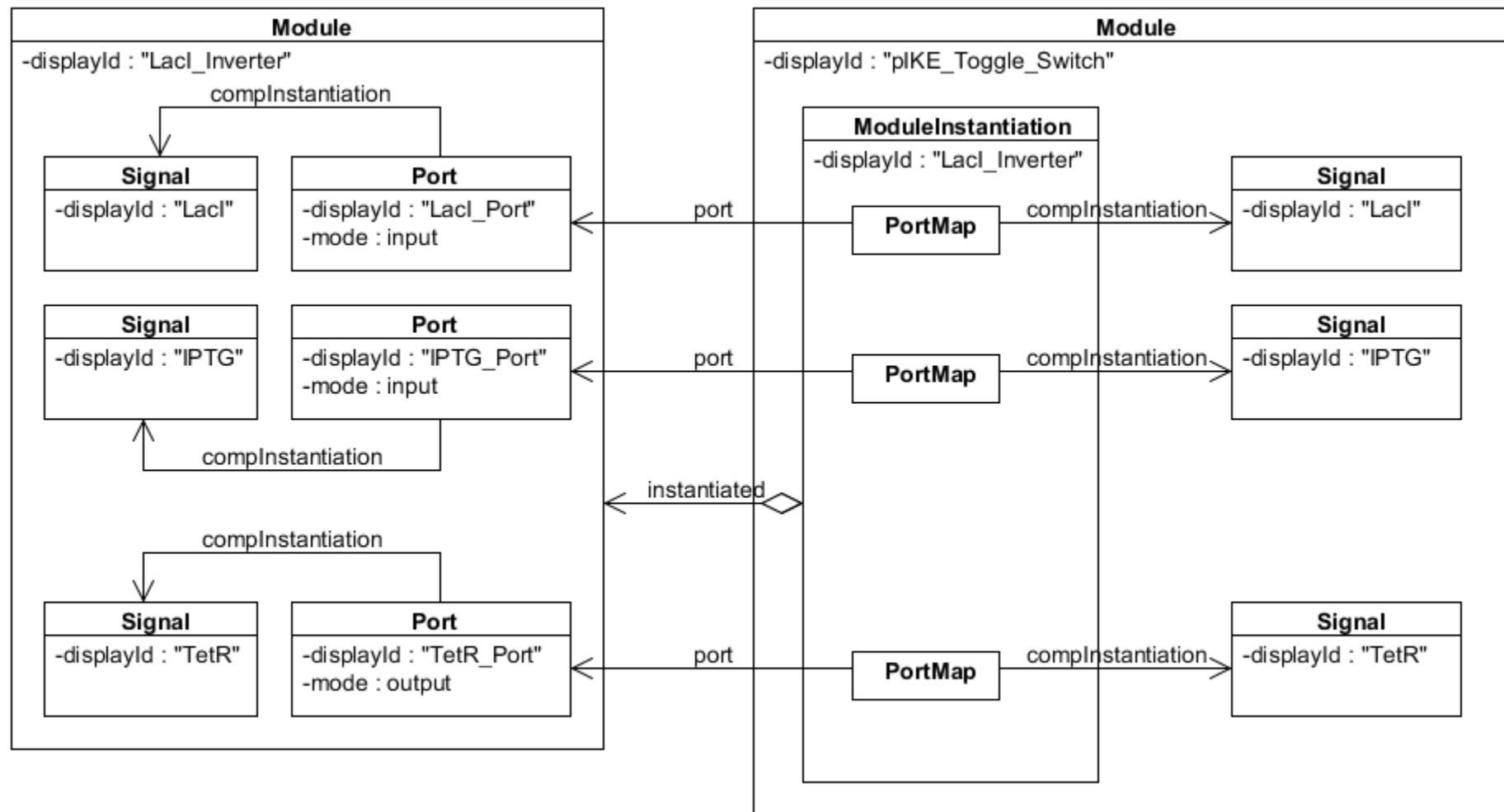
Example: Module Ports



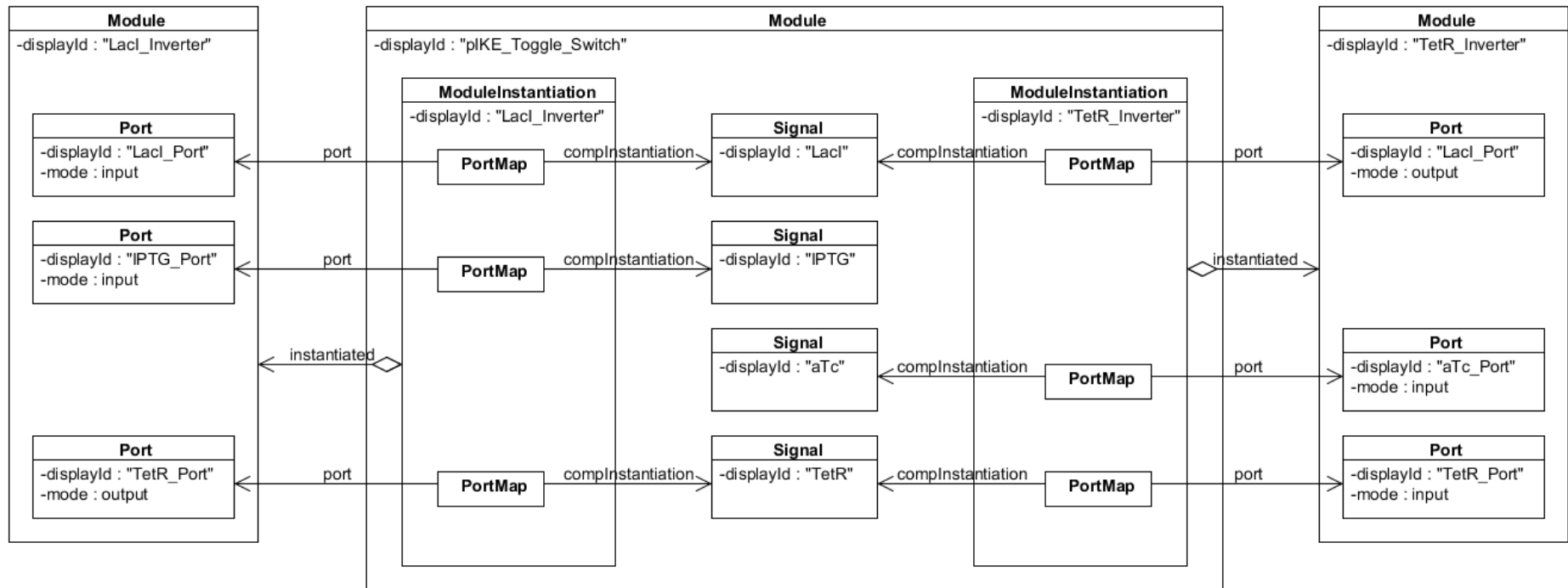
Example: Module Instantiation



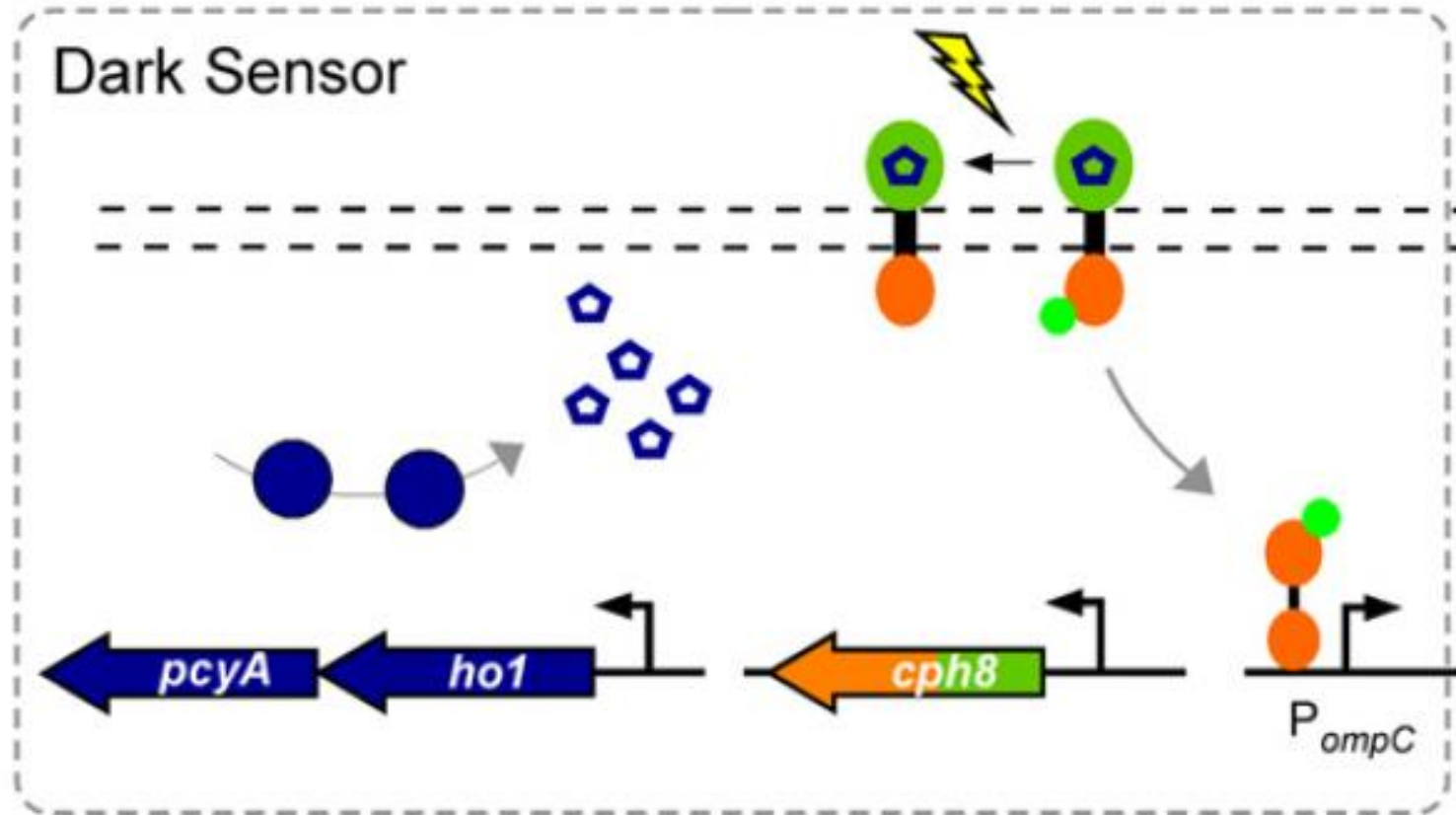
Example: Port Mapping



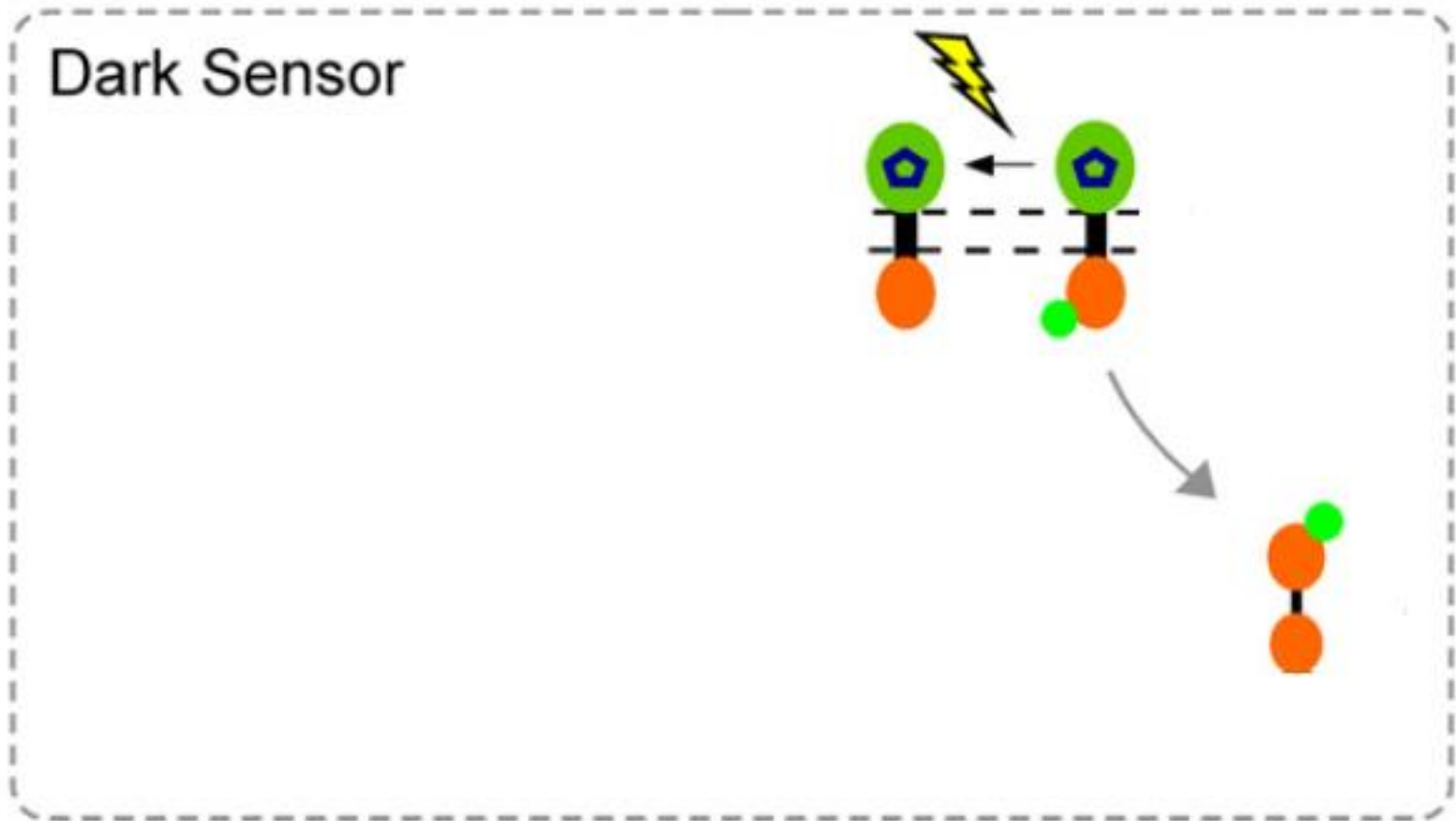
Toggle Module Composed



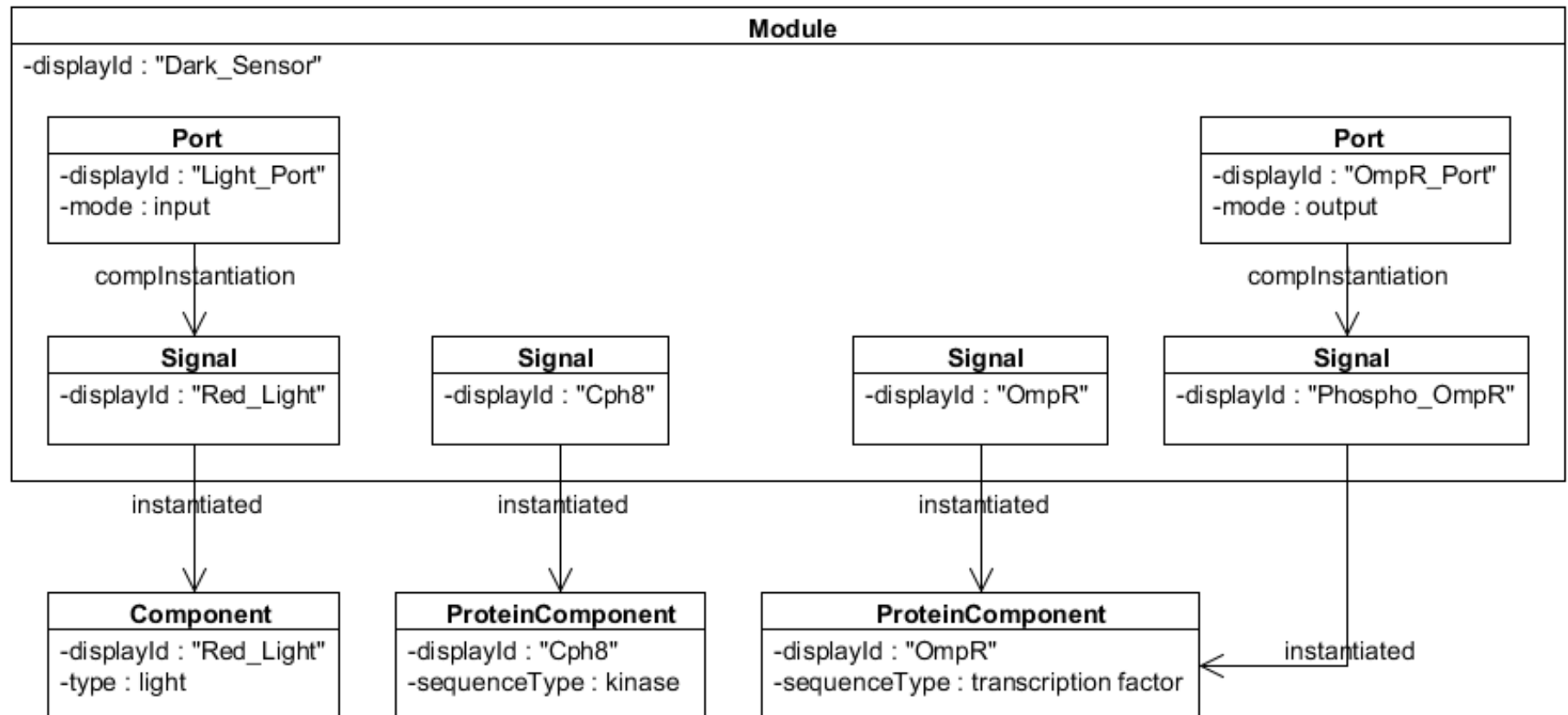
Dark Sensor (Tabor et al. 2009)



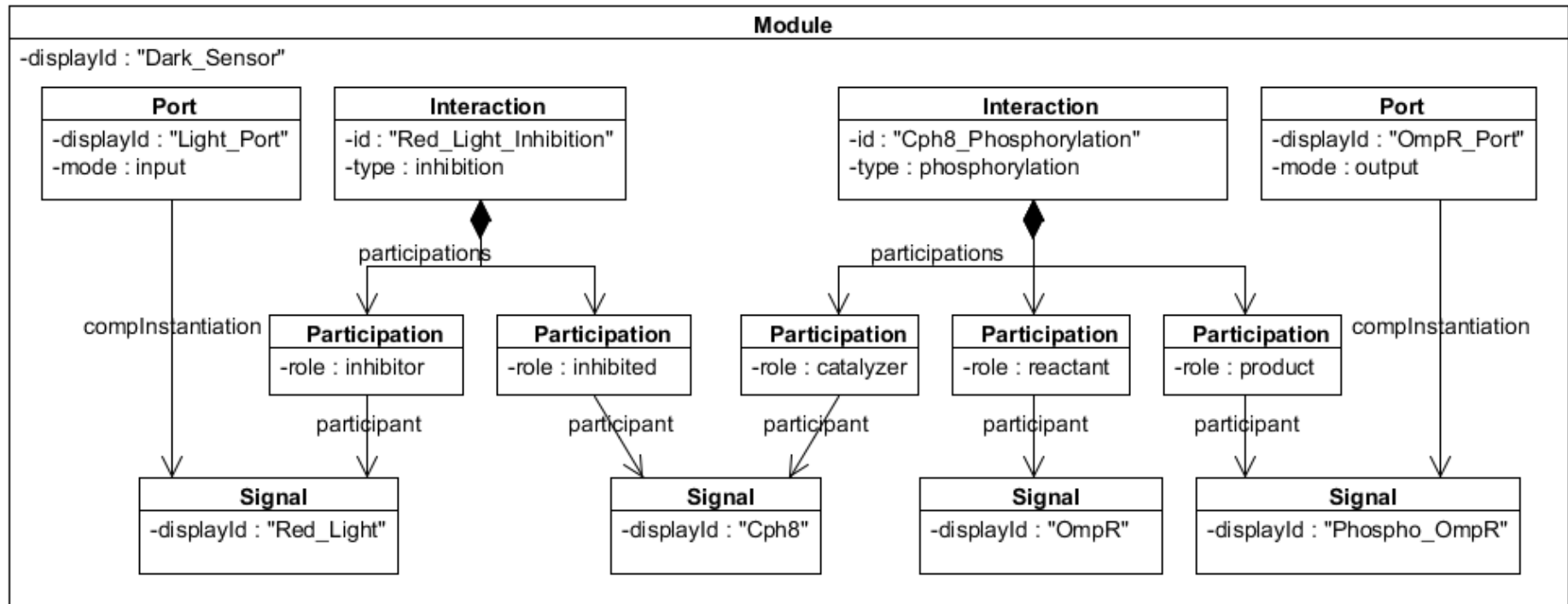
Dark Sensor (Tabor et al. 2009)



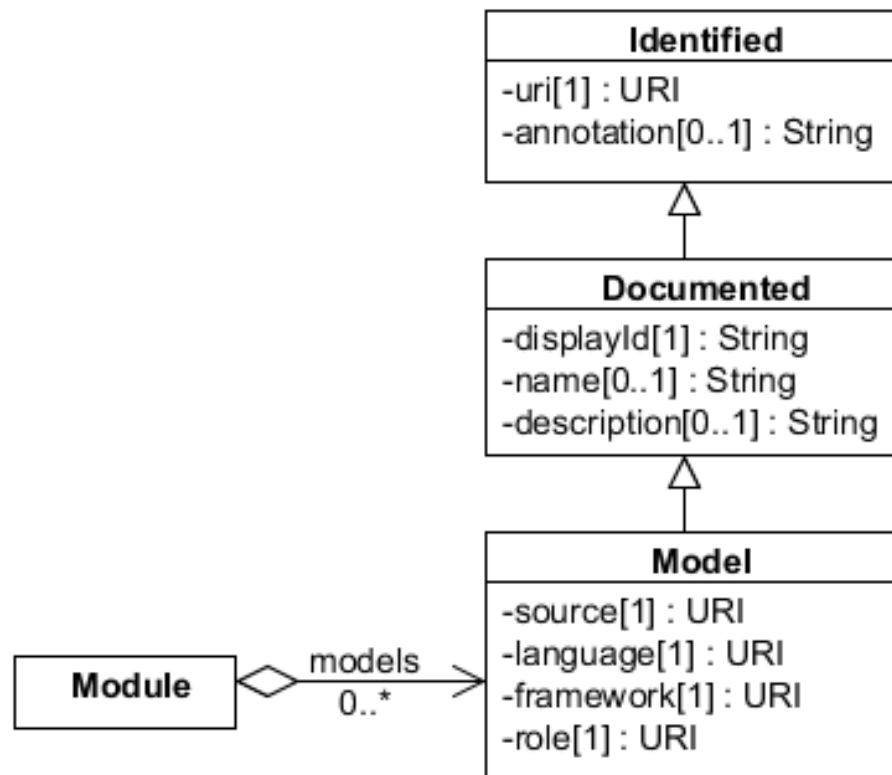
Dark Sensor: Instantiation



Dark Sensor: Interactions



Extension Status: Modeling



Extension Status: Context

Canonical EX: *Repressilator*



Measurement Device Zeiss Axiovert 135TV microscope

Environment The temperature of the samples was maintained at 30–32 °C by using Peltier devices (Melcor)

Container coverslip and microscope slide

Medium minimal media
1 ml of liquid 2% SeaPlaque low-melt agarose (FMC) in media
100 μ M IPTG inducer
antibiotic 20 g ml⁻¹ kanamycin or 20 g ml⁻¹ ampicillin
minimum initial cell density OD = 0.1

Host *E. coli* lac- strain MC4100

Composition Genome, Repressilator and Reporter plasmids

Summary UML

