

# AlgebraicJulia: Applied Category Theory in Julia

Micah Halter, *James Fairbanks*  
Georgia Tech Research Institute (GTRI)

Evan Patterson  
MIT



# Spectrum of Scientific Computing Technology

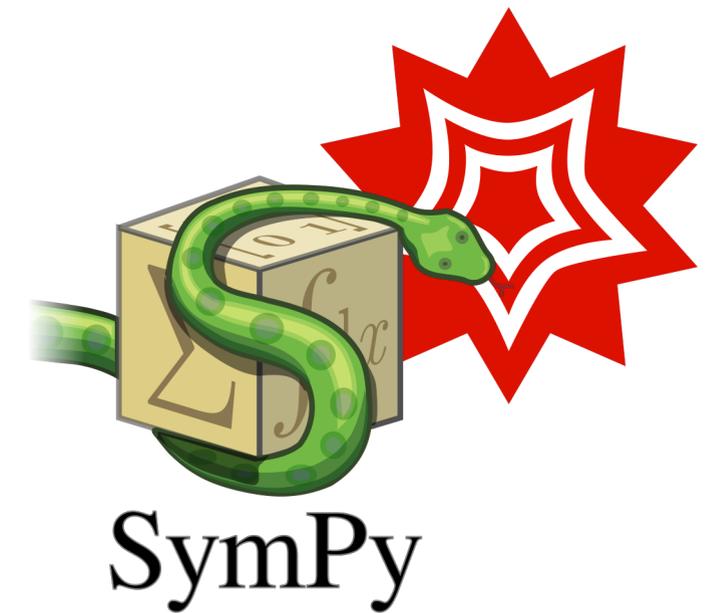
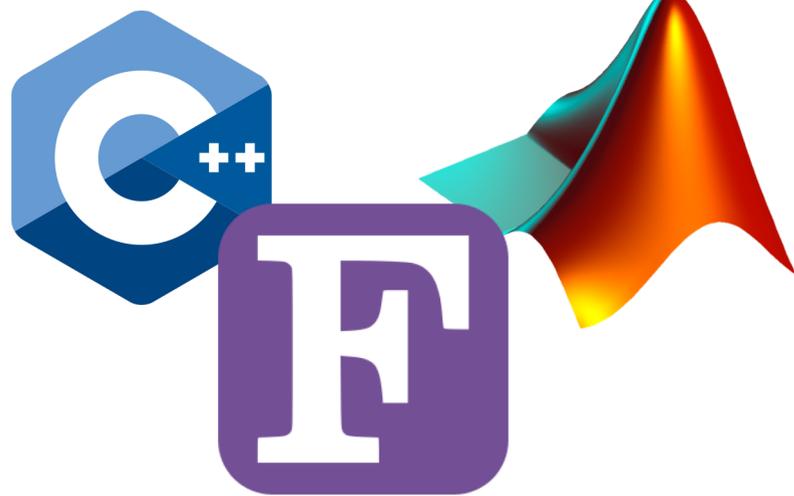


Arbitrary Code

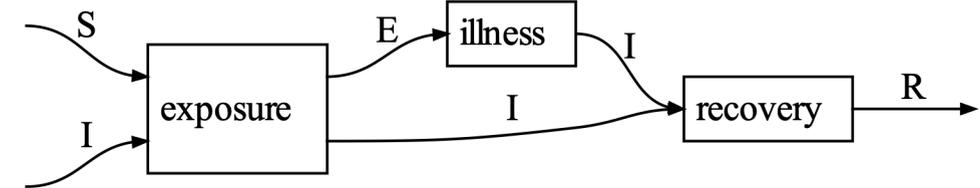
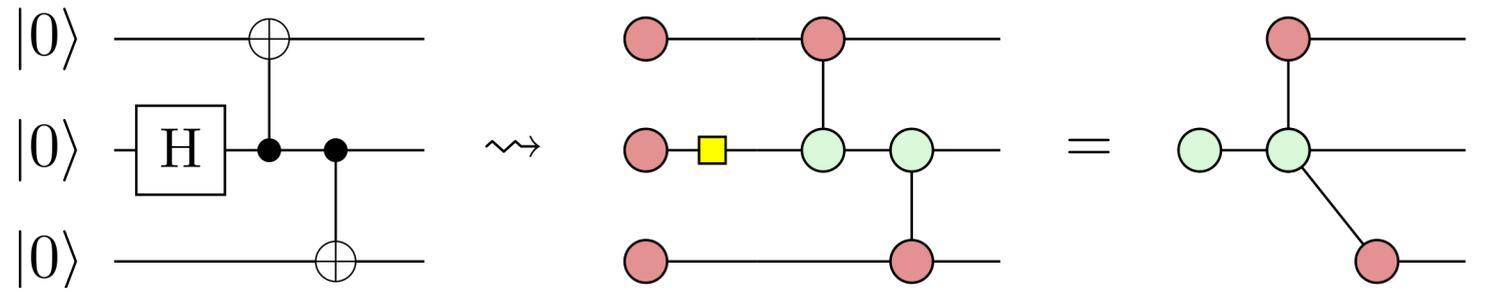
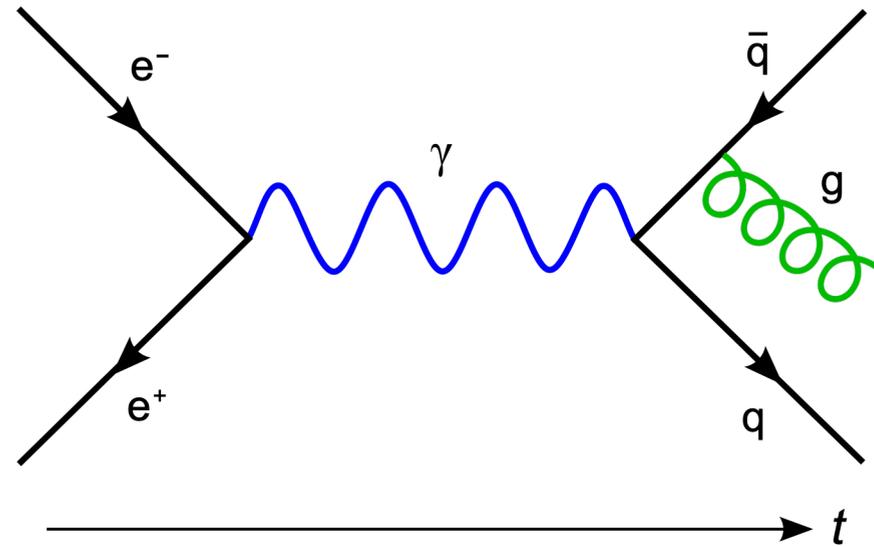
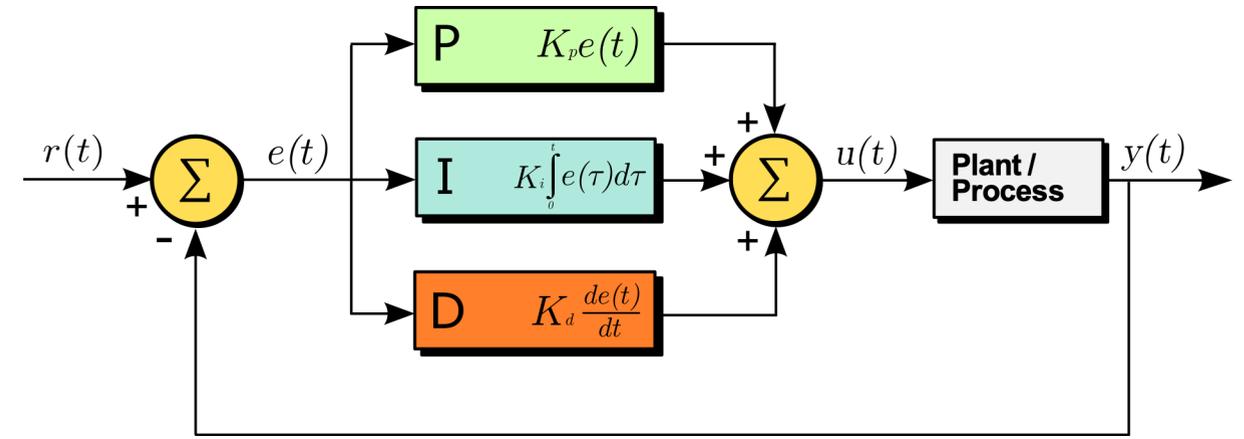
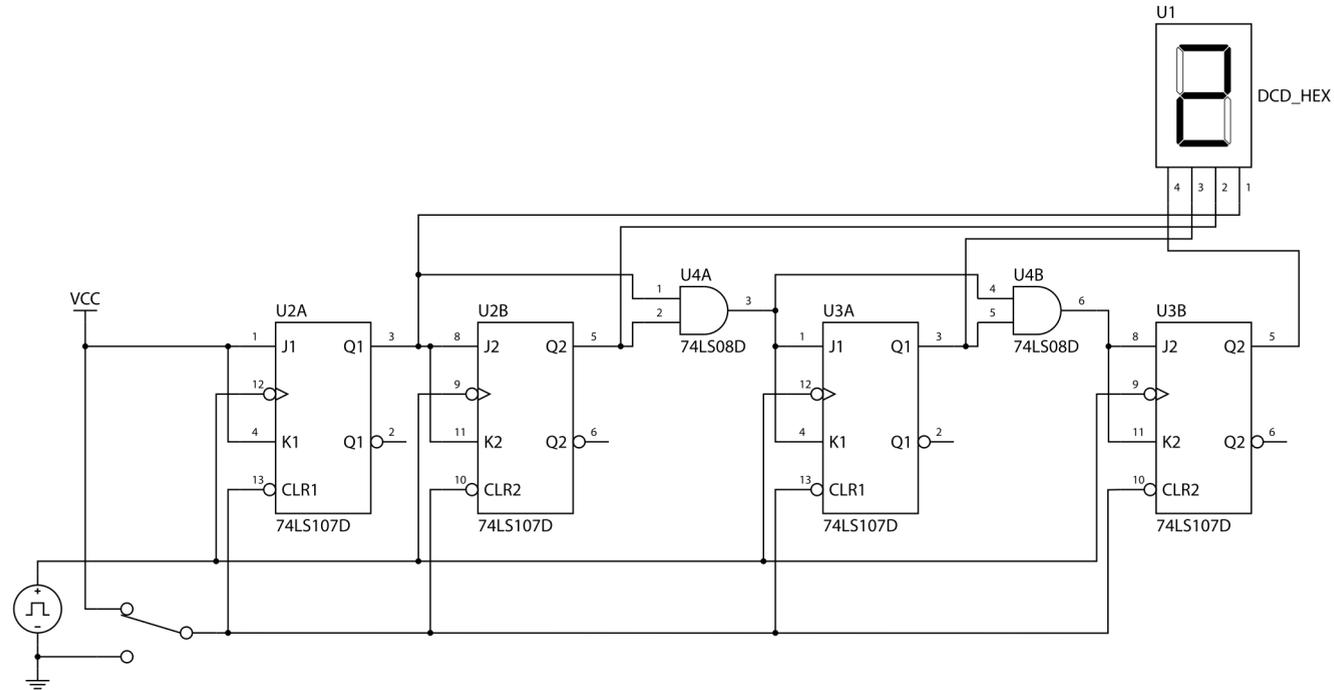
Domain Specific Languages

Modeling Frameworks

Computer Algebra Systems



# Formal Scientific Diagrams



# Three Layers of GAT Based Modeling

## Theory

$A, B, C$

$a : A, b : B, c : C$

$a \otimes b \cdot c$

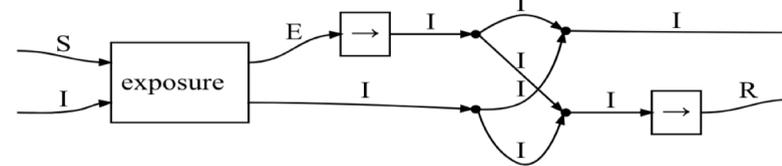
$(a \otimes b) \cdot (c \otimes d)$   
 $= (a \cdot c) \otimes (b \cdot d)$

## Syntax

### Formula Notation

$seir = expo \cdot (f_{E,I} \otimes id_I) \cdot \nabla_I \cdot \Delta_I \cdot (id_I \otimes f_{I,R})$

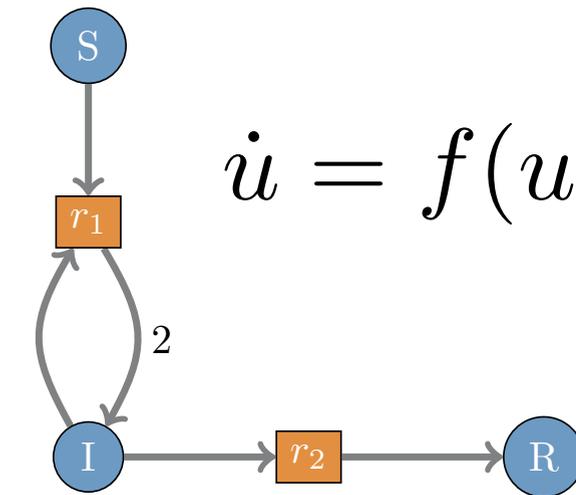
### Wiring Diagram



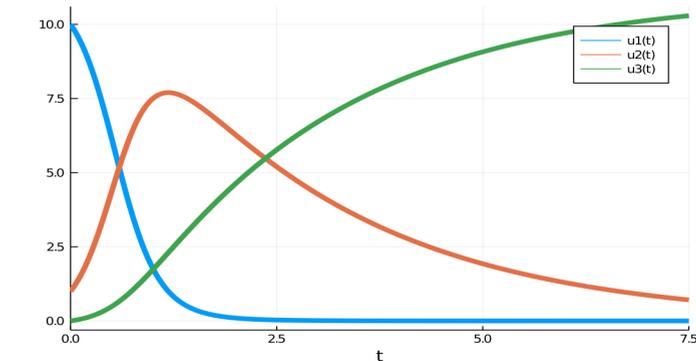
### Embedded Domain Specific Language

```
d = @program Disease (s::S, e::E, i::I) begin
  e1, i1 = exposure{S,I,E}(s,i)
  i2 = spontaneous{E,I}(e1)
  e = [e, e1]
  e_out = spontaneous{E,E}(e)
  i1 = [i1, i2]
  r = spontaneous{I,R}(i1)
  s_out = spontaneous{S,S}(s)
  return s_out, e_out, spontaneous{I,I}(i1)
end
```

## Instance



$$\dot{u} = f(u, t)$$



# DSLs implemented in Catlab

@theory

- Algebraic Structure that defines possible expressions (the theory of Groups)

@syntax

- **GATExpr**: Expr tied to a specific Theory

@presentation

- A specific example of the theory: (the Group of Integers mod 7)

@instance

- A Julia implementation (types and functions) implementing the theory

@program

- Lets you express formulas using program notation

# FinOrd: the Category of Natural Numbers



```
@theory Category (Ob, Hom) begin
  Ob :: TYPE
  Hom (dom :: Ob, codom :: Ob) :: TYPE

  id (A :: Ob) :: (A → A)
  compose (f :: (A → B), g :: (B → C)) :: (A → C)
    ⊢ (A :: Ob, B :: Ob, C :: Ob)
end
```

```
@instance Category (FinOrd, FinOrdMap) begin
  dom (f :: FinOrdMap) = FinOrd (f.dom)
  codom (f :: FinOrdMap) = FinOrd (f.codom)

  id (A :: FinOrd) = FinOrdFunction (identity, A, A)

  function compose (f :: FinOrdMap, g :: FinOrdMap)
    @assert codom (f) == dom (g)
    FinOrdFunction (compose (f.func, g.func), dom (f), codom (g))
  end
end
```

```
struct FinOrd n :: Int end

struct FinOrdFunc <: FinOrdMap
  func :: Function
  dom :: Int
  codom :: Int
end

(f :: FinOrdFunc) (x) = f.func (x)

struct FinOrdVec <: FinOrdMap
  func :: Vector {Int}
  codom :: Int
end

(f :: FinOrdVec) (x) = f.func [x]
```

# Monoidal Categories Support Programming



```
@present P(FreeSymmetricMonoidalCategory) begin
```

```
X::Ob
```

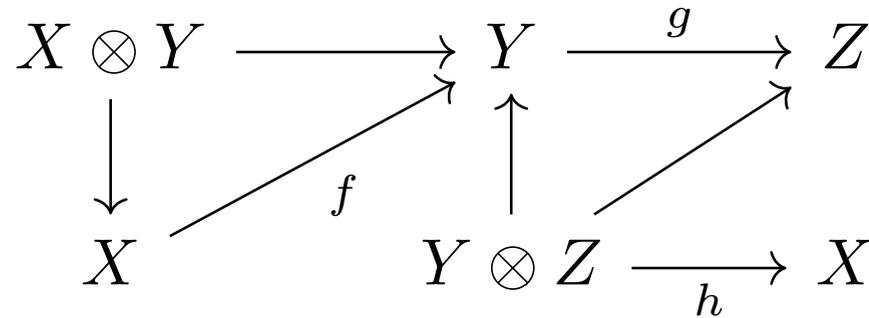
```
Y::Ob
```

```
Z::Ob
```

```
f::Hom(X, Y)
```

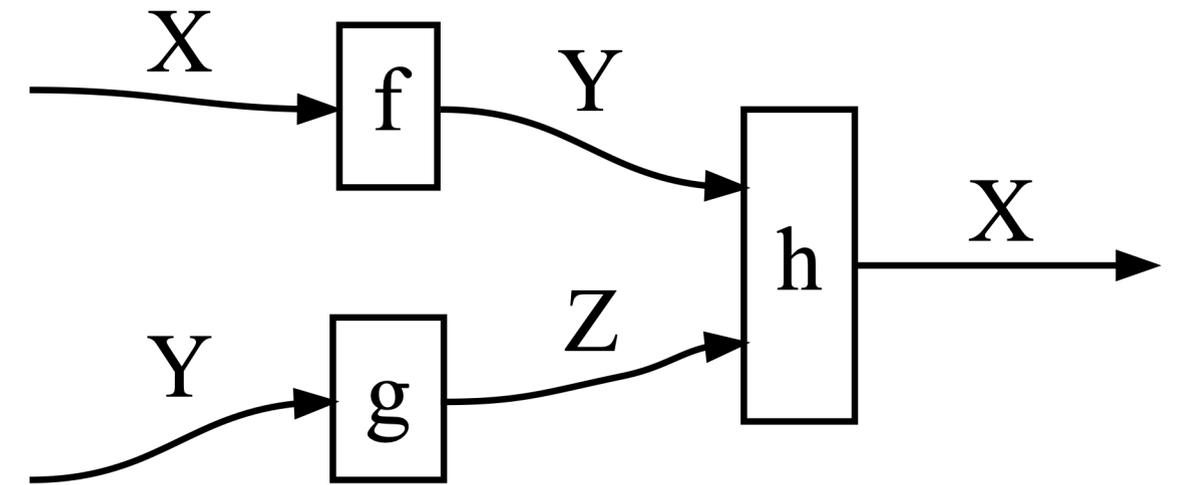
```
g::Hom(Y, Z)
```

```
h::Hom(Y⊗Z, X)
```



```
end
```

$$(f \otimes g) \cdot h: X \otimes Y \rightarrow X$$



```
@signature Category(Ob, Hom) => SymmetricMonoidalCategory(Ob, Hom)
```

```
otimes(A::Ob, B::Ob)::Ob
```

```
otimes(f::(A → B), g::(C → D))::((A ⊗ C) → (B ⊗ D)) ⊣
```

```
(A::Ob, B::Ob, C::Ob, D::Ob)
```

```
@op (⊗) := otimes
```

```
munit()::Ob
```

```
braid(A::Ob, B::Ob)::((A ⊗ B) → (B ⊗ A))
```

```
@op (σ) := braid
```

```
end
```

```
d = @program P (x::X, y::Y) begin
```

```
a = f(x)
```

```
b = g(y)
```

```
z = h(a, b)
```

```
return z
```

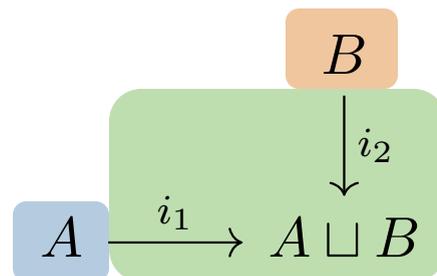
```
end
```

# Programming with Pictures

Initial Object

$$0 \longrightarrow A$$

Coproducts

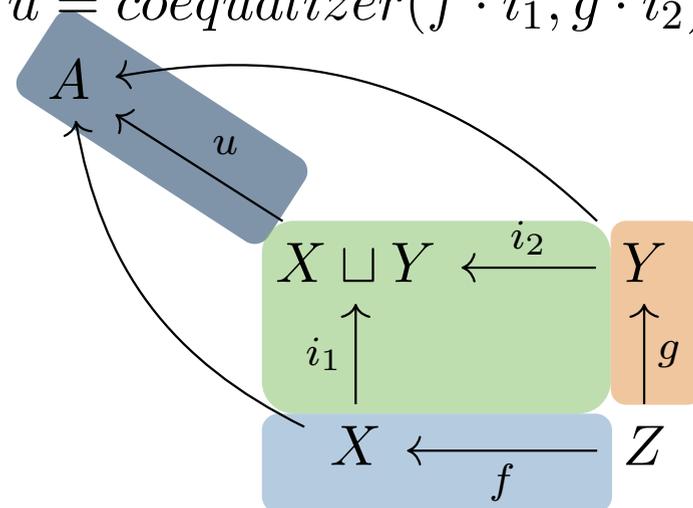


Coequalizers

$$E \xleftarrow{u} A \rightrightarrows B$$

Pushout of  $f, g$ ,

$$u = \text{coequalizer}(f \cdot i_1, g \cdot i_2)$$



```
initial(::Type{FinOrd}) = FinOrd(0)
```

```
function coproduct(A::FinOrd, B::FinOrd)
    m, n = A.n, B.n
    ι1 = FinOrdMap(1:m, m, m+n)
    ι2 = FinOrdMap(m+1:m+n, n, m+n)
    Cospan(ι1, ι2)
end
```

```
function pushout(span::Span{<:FinOrdMap, <:FinOrdMap})
    f, g = left(span), right(span)
    coprod = coproduct(codom(f), codom(g))
    ι1, ι2 = left(coprod), right(coprod)
    coeq = coequalizer(f·ι1, g·ι2)
    Cospan(ι1·coeq, ι2·coeq)
end
```

# Application: Simulating a Petri Net as ODEs

## Novel coronavirus 2019-nCoV: early estimation of epidemiological parameters and epidemic predictions

Version 2. Updated 27 Jan 2020

Jonathan M. Read<sup>1</sup>, Jessica R.E. Bridgen<sup>1</sup>, Derek A.T. Cummings<sup>2</sup>, Antonia Ho<sup>3</sup>, Chris P. Jewell<sup>1</sup>

### Affiliations:

1. Centre for Health Informatics, Computing and Statistics, Lancaster Medical School, Lancaster University, Lancaster, United Kingdom.

2. Department of Biology and Emerging Pathogens Institute, University of Florida, Gainesville, United States of America.

3. Medical Research Council-University of Glasgow Centre for Virus Research, Glasgow, United Kingdom.

Correspondence: [jonathan.read@lancaster.ac.uk](mailto:jonathan.read@lancaster.ac.uk)

## Impact of international travel and border control measures on the global spread of the novel 2019 coronavirus outbreak

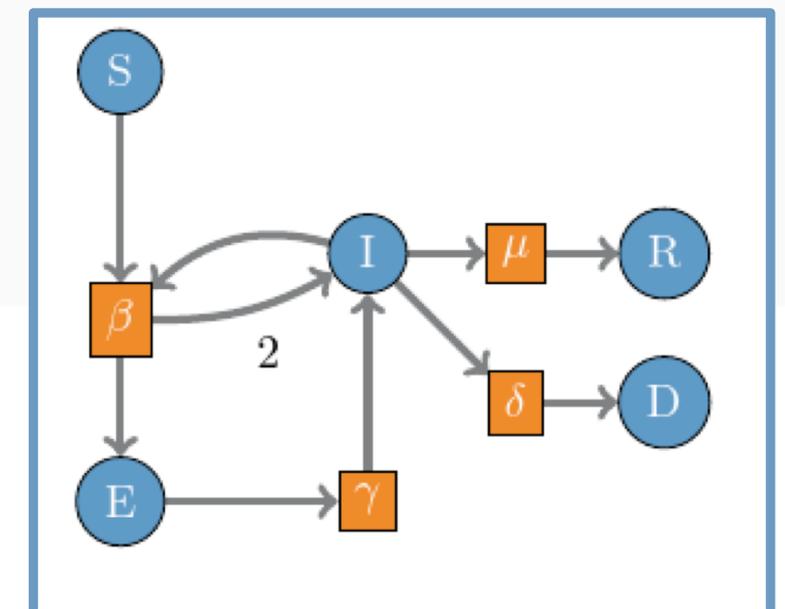
Chad R. Wells<sup>a</sup>, Pratha Sah<sup>a</sup>, Seyed M. Moghadas<sup>b</sup>, Abhishek Pandey<sup>a</sup>, Affan Shoukat<sup>a</sup>, Yaning Wang<sup>c</sup>, Zheng Wang<sup>d</sup>, Lauren A. Meyers<sup>e,f</sup>, Burton H. Singer<sup>g,1</sup>, and Alison P. Galvani<sup>a</sup>

<sup>a</sup>Center for Infectious Disease Modeling and Analysis, Yale School of Public Health, New Haven, CT 06520; <sup>b</sup>Agent-Based Modelling Laboratory, York University, Toronto, ON M3J 1P3, Canada; <sup>c</sup>State Key Laboratory of Mycology, Institute of Microbiology, Chinese Academy of Sciences, 100101 Beijing, China; <sup>d</sup>Department of Biostatistics, Yale School of Public Health, New Haven, CT 06510; <sup>e</sup>Department of Integrative Biology, The University of Texas at Austin, Austin, TX 78712; <sup>f</sup>Santa Fe Institute, Santa Fe, NM 87501; and <sup>g</sup>Emerging Pathogens Institute, University of Florida, Gainesville, FL 32610

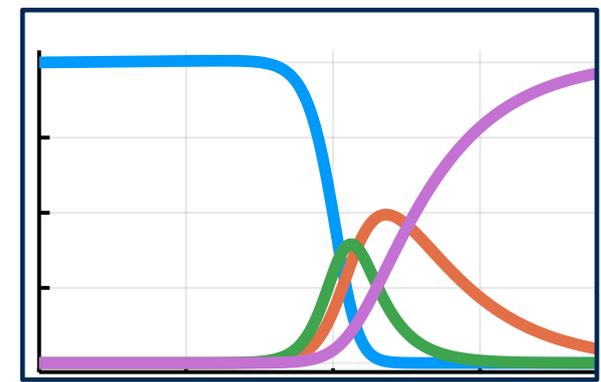
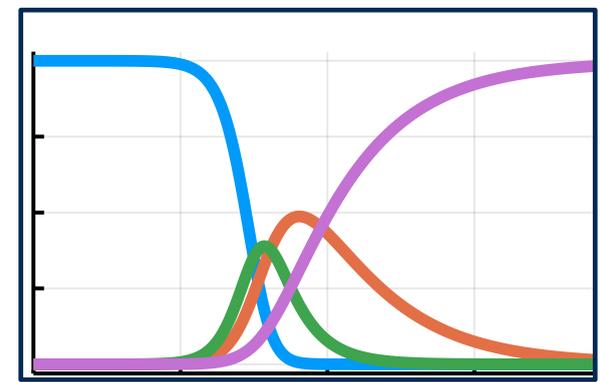
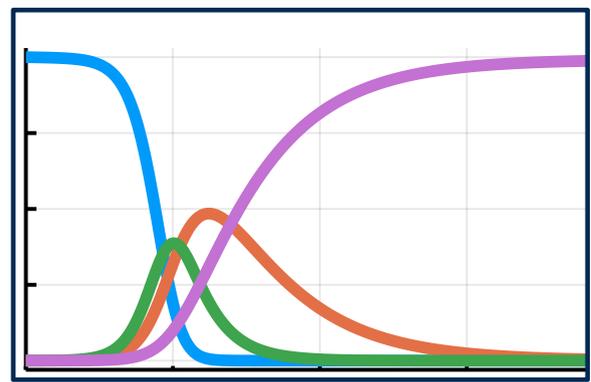
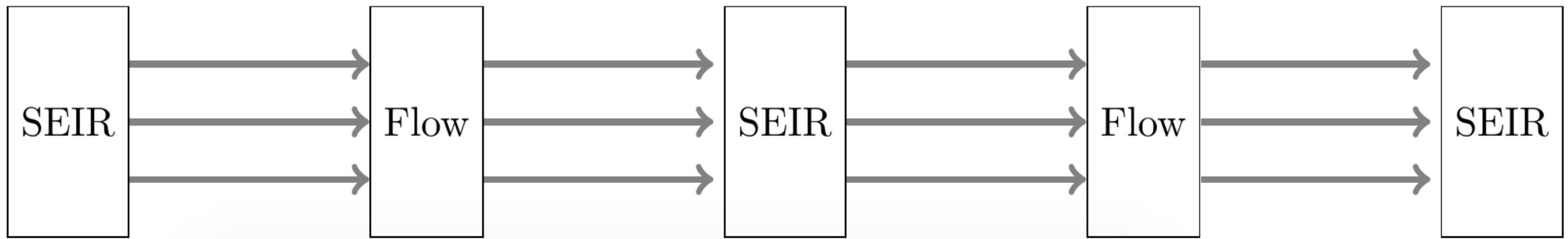
Contributed by Burton H. Singer, February 27, 2020 (sent for review February 12, 2020; reviewed by Yoav Keynan and Heman Shakeri)

Solve the initial value problem for  $\dot{u} = f(u, t)$

```
f(u, p) = begin
    for (i, t) in enumerate(reactions)
        ← reagents = t[1]
        → products = t[2]
        φ = p[i]*prod(u[reagents])
        du[reagents] .-= φ
        du[products] .+= φ
    end
    return du
end
```

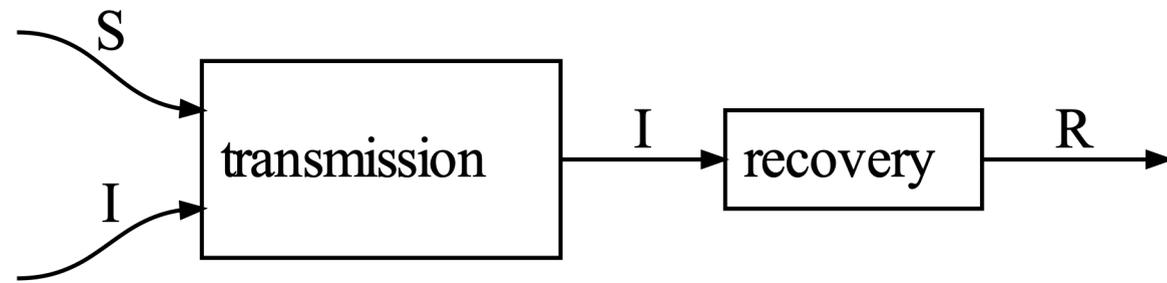


# Epidemiology Modeling Framework



# Basic SIR model

```
sir = transmission · recovery
```



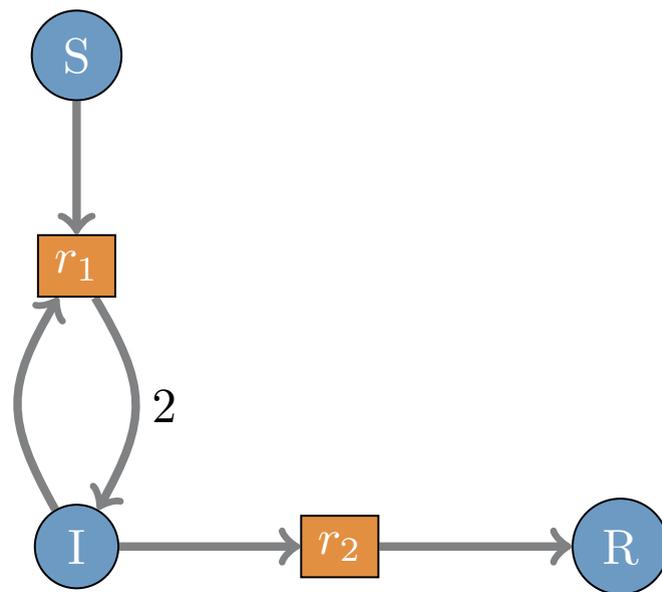
```
f = vectorfield(sird)
```

$$\dot{u}_1 = -r_1 u_1 u_2$$

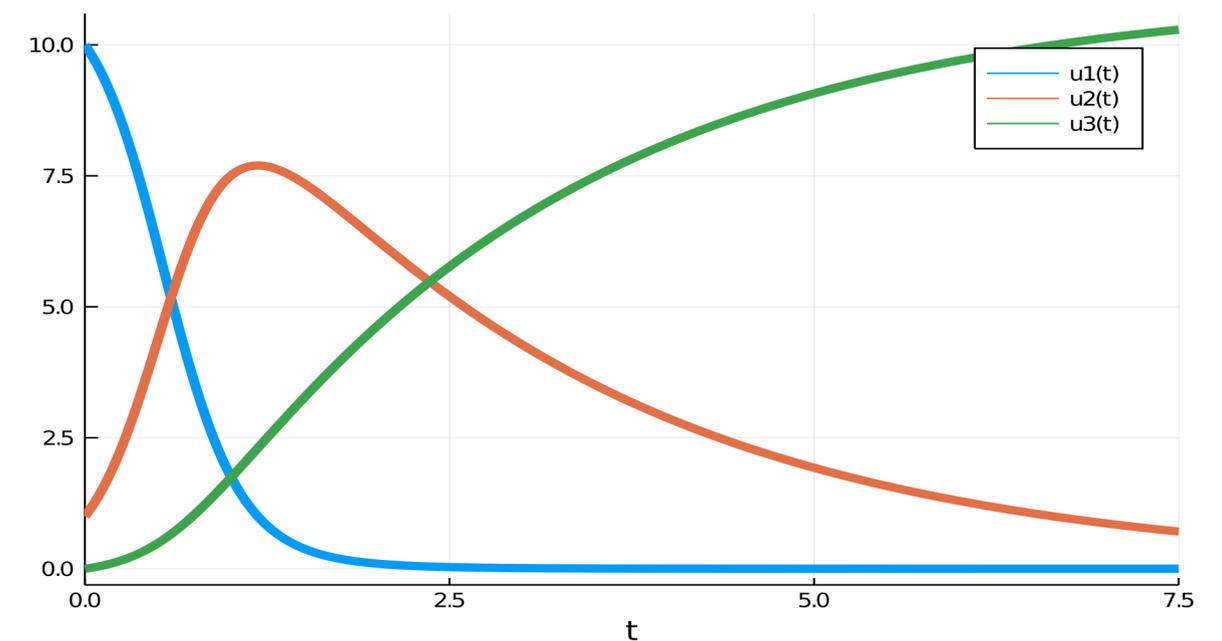
$$\dot{u}_2 = r_1 u_1 u_2 - r_3 u_2$$

$$\dot{u}_3 = r_3 u_2$$

```
sird = decoration(F(sir))
```

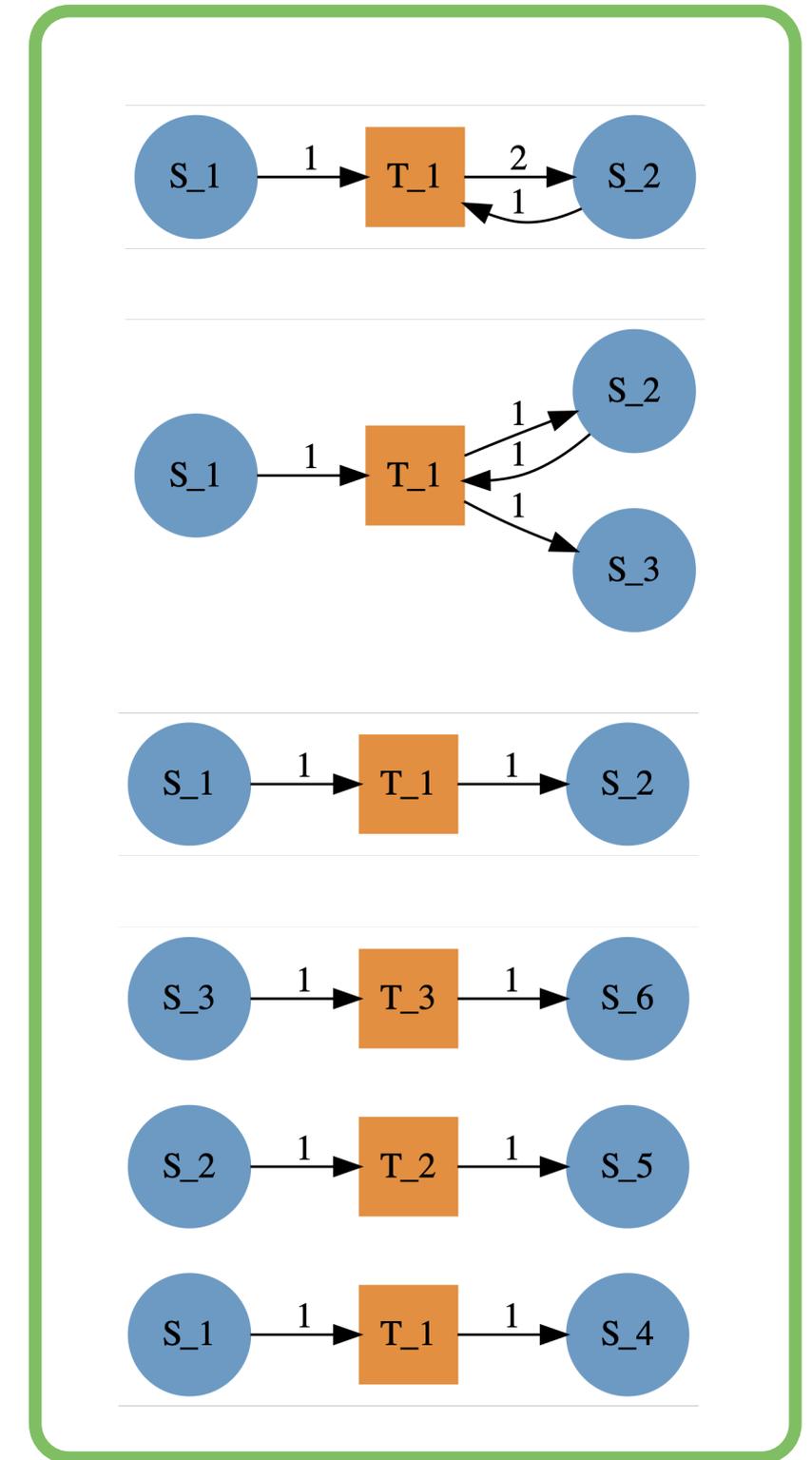
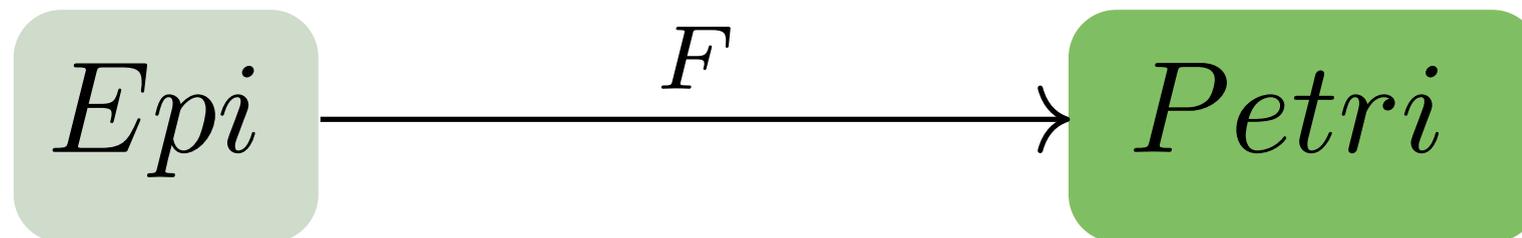


```
sol = solve(f, u0, r, (t0, t1))
```



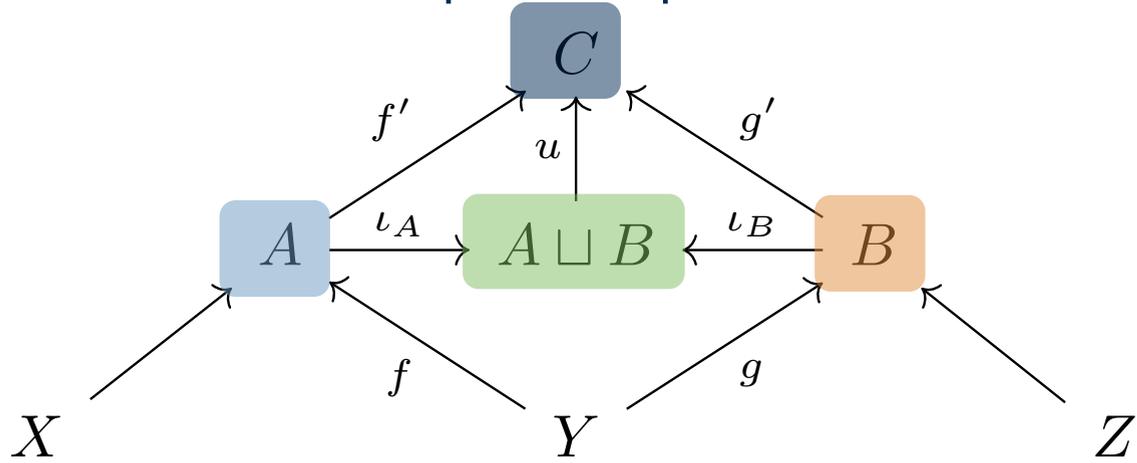
# Epidemiology Modeling

```
@present Epidemiology(FreeBiproductCategory) begin
  S::Ob
  E::Ob
  I::Ob
  R::Ob
  D::Ob
  transmission::Hom(S⊗I, I)
  exposure::Hom(S⊗I, E⊗I)
  illness::Hom(E, I)
  recovery::Hom(I, R)
  death::Hom(I, D)
  travel::Hom(S⊗E⊗I, S⊗E⊗I)
end
```



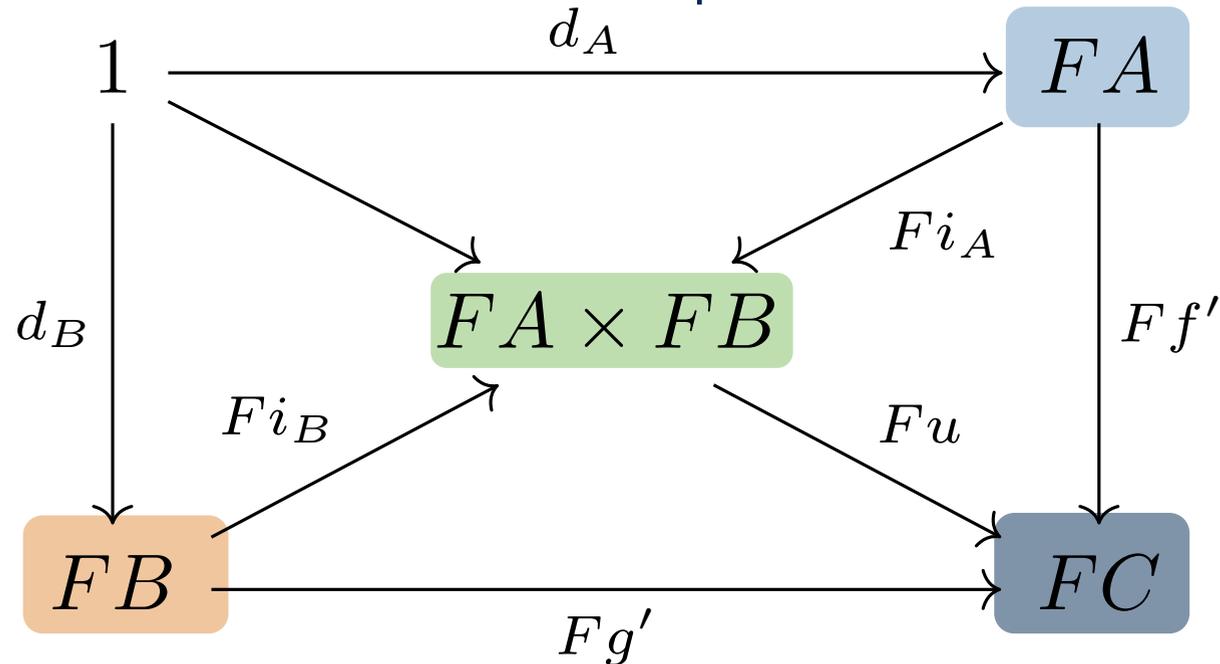
# Computing Decorated Cospans

Cospan Composition



$$FA \sqcup FB \xrightarrow{L} F(A \sqcup B)$$

Decorator Composition



```
compose(p::PetriCospan, q::PetriCospan) = begin
```

```
    u, f', g' = pushout(f, g)
```

```
    F = functor(decorator(p))
```

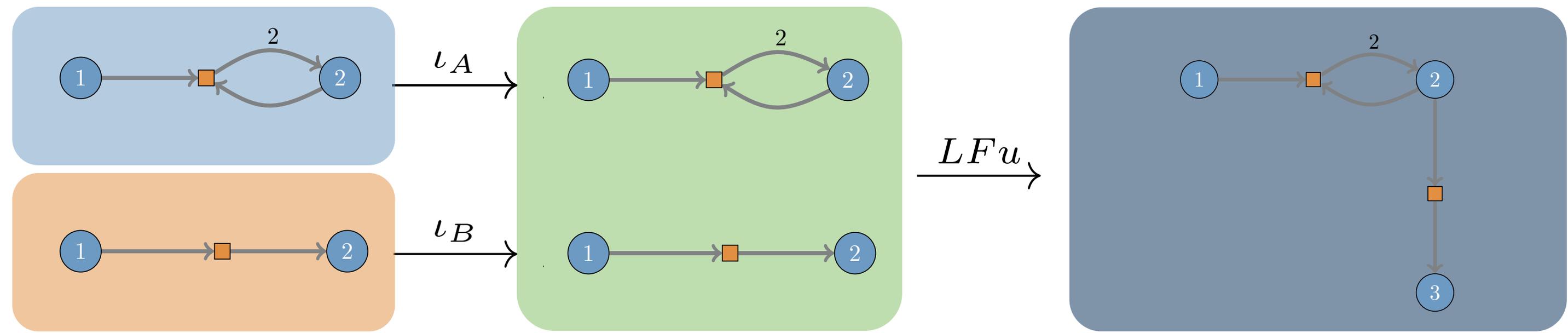
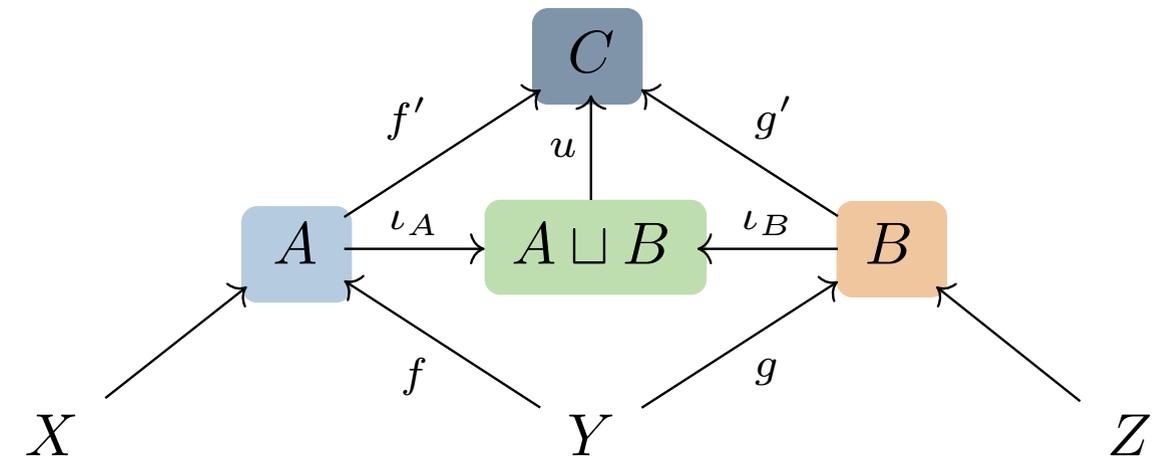
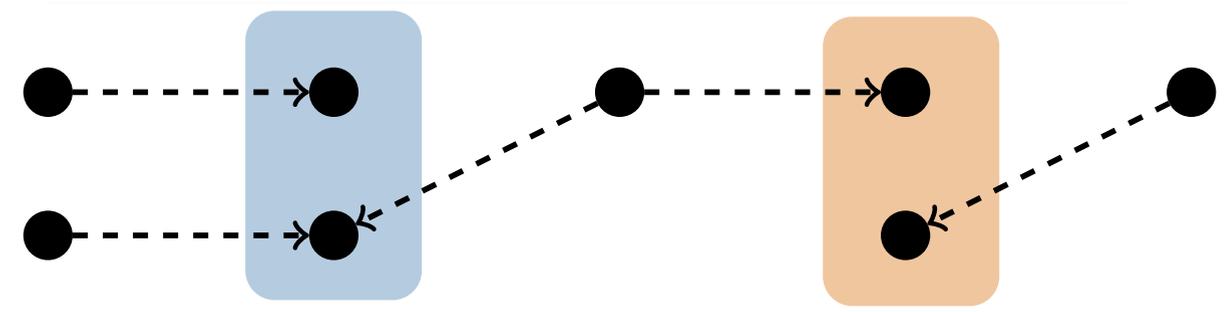
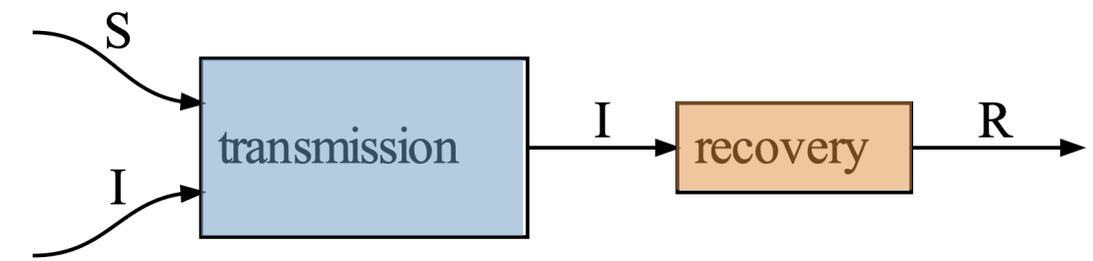
```
    L = laxator(decorator(p))
```

```
    dc = F(u) (L(decoration(p), decoration(q)))
```

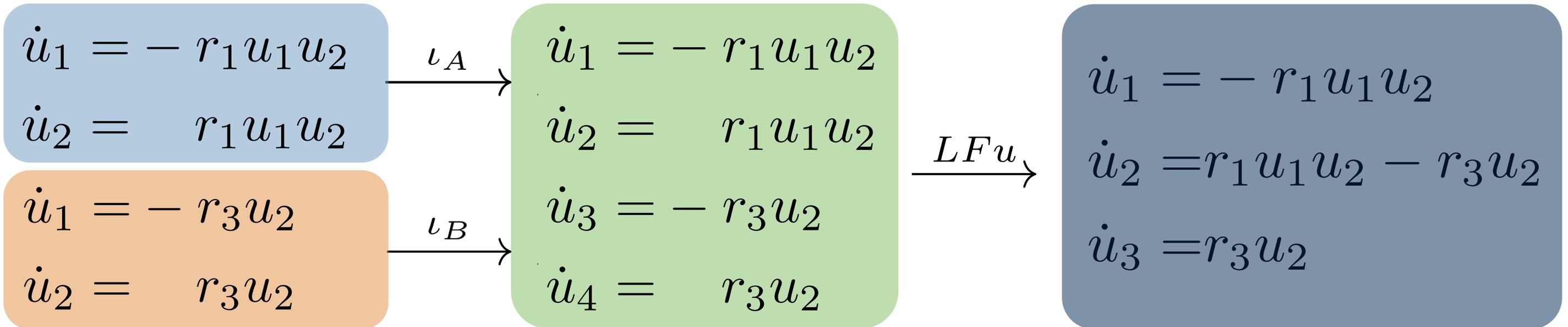
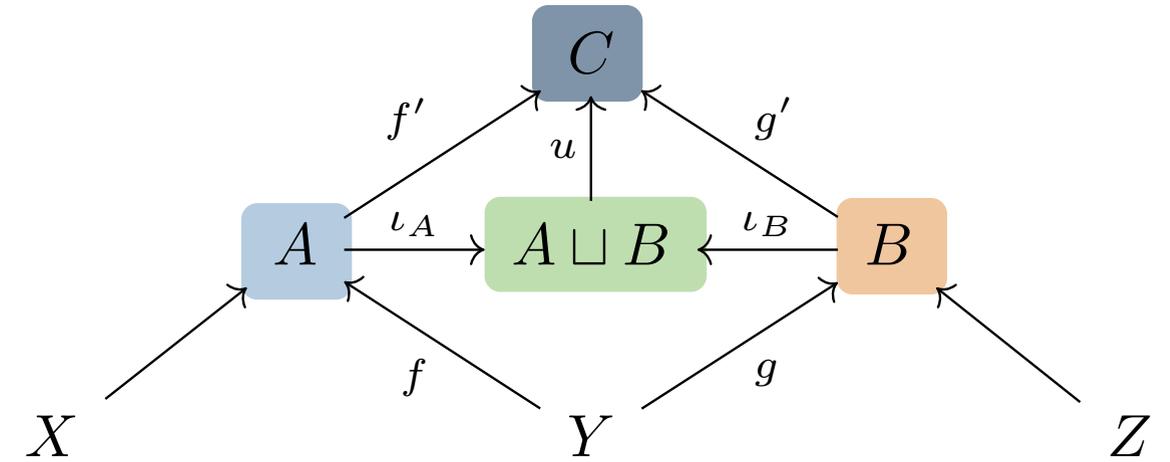
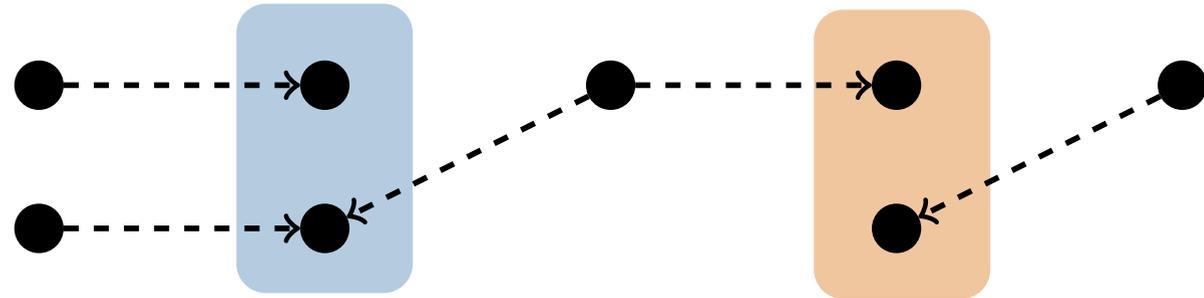
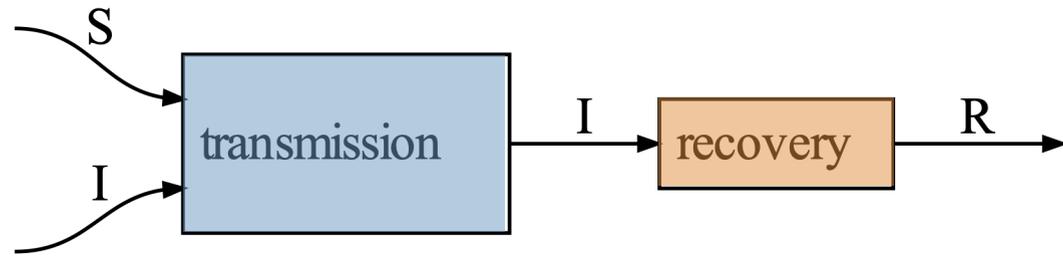
```
    return PetriCospan(Cospan(f', g'), decorator(p), dc)
```

```
end
```

# Petri Net Decorated Cospan Composition

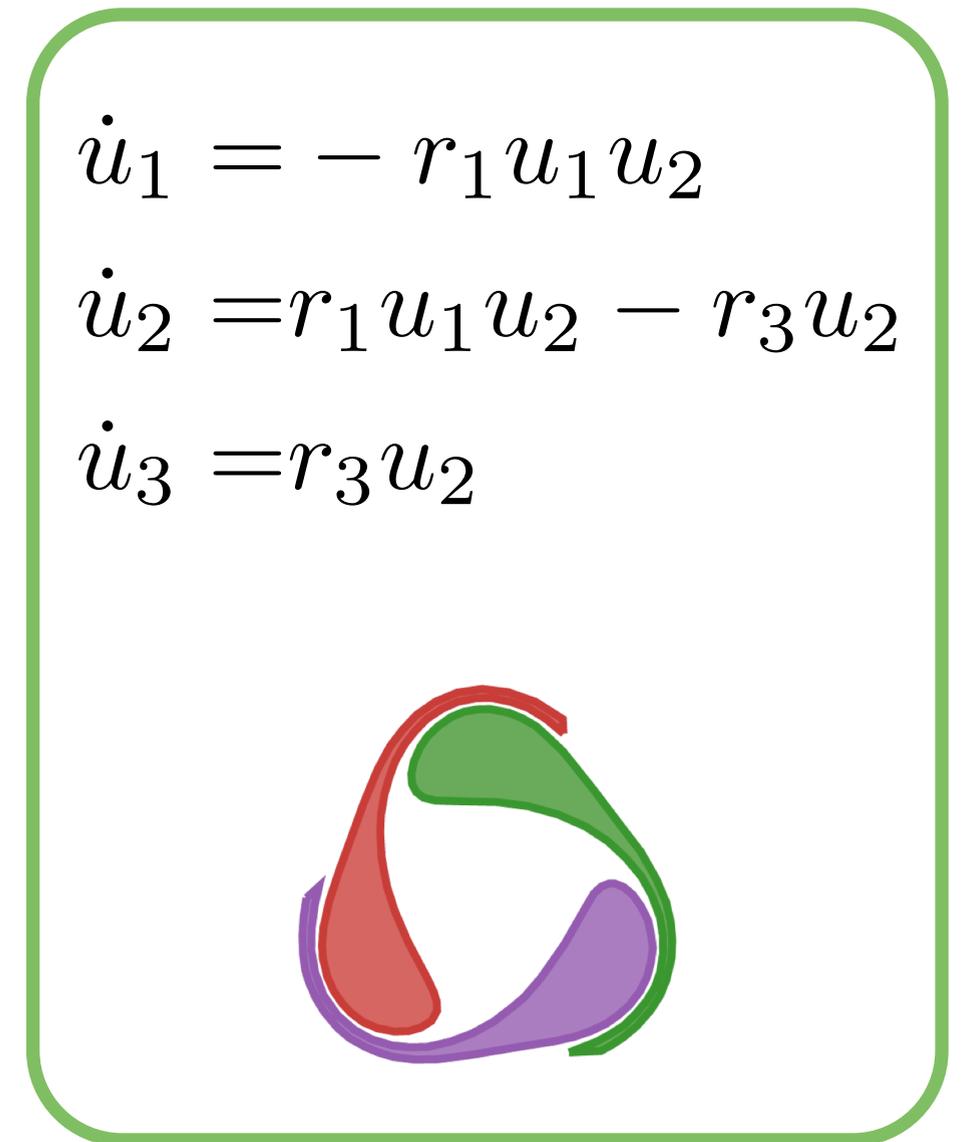
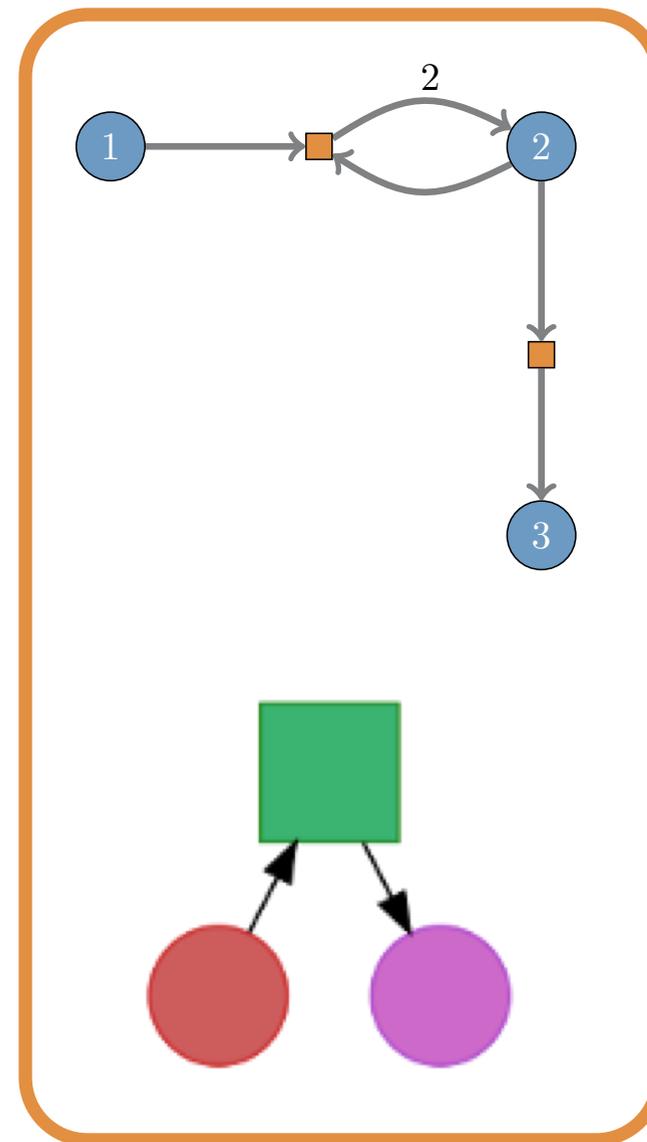
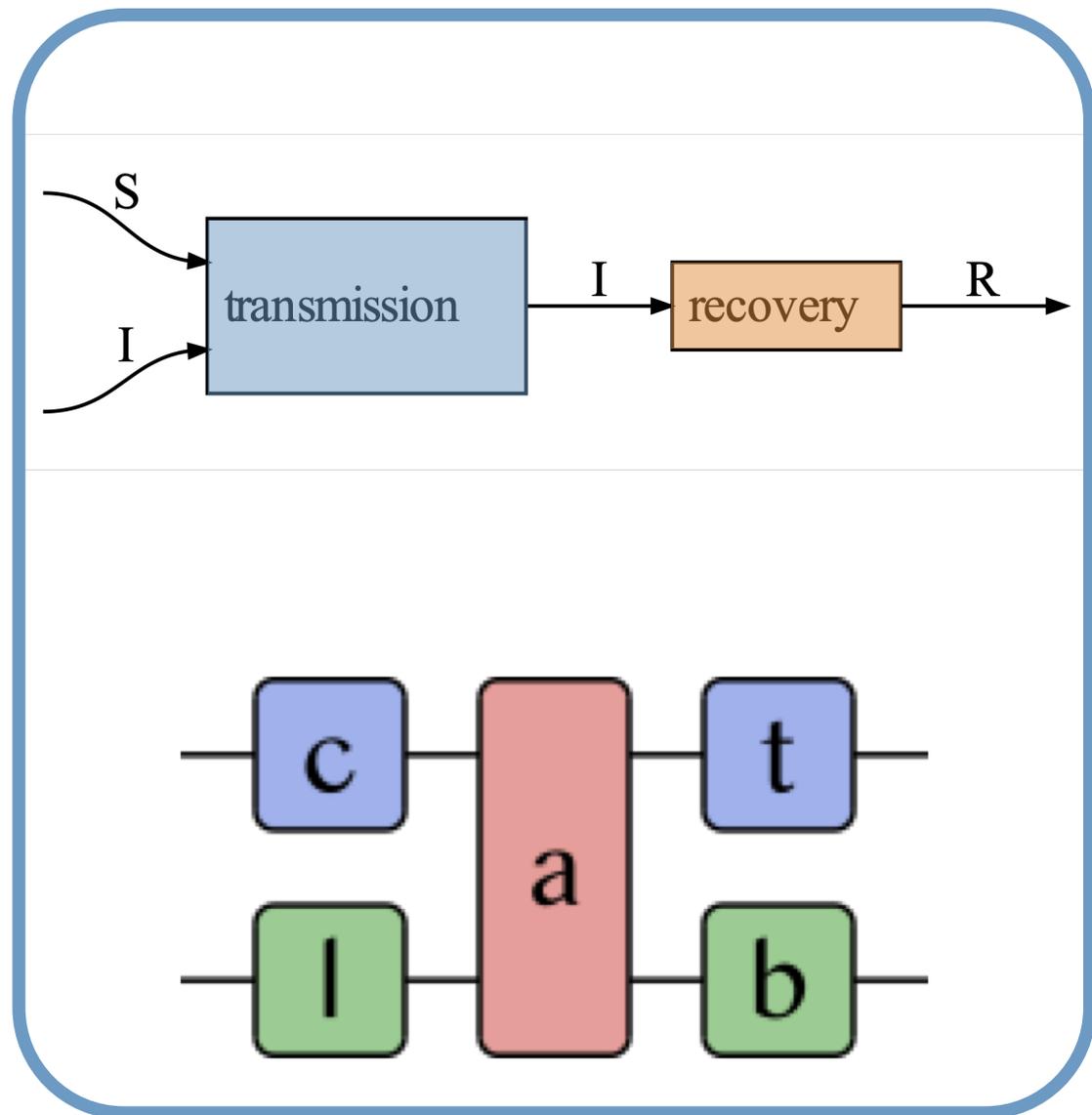


# Resource Sharing to Assign Kinetics



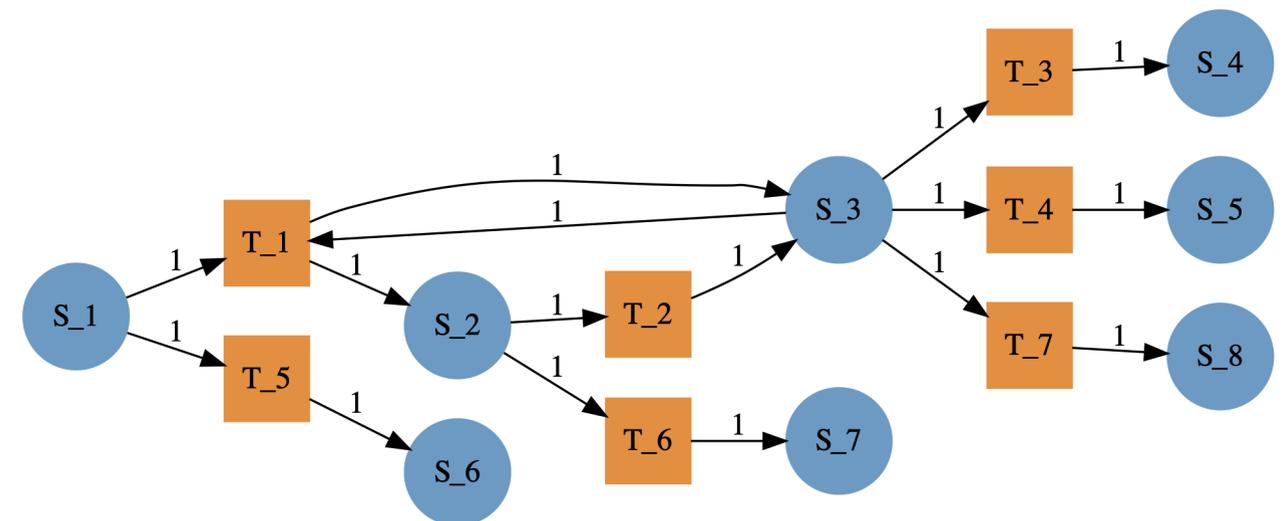
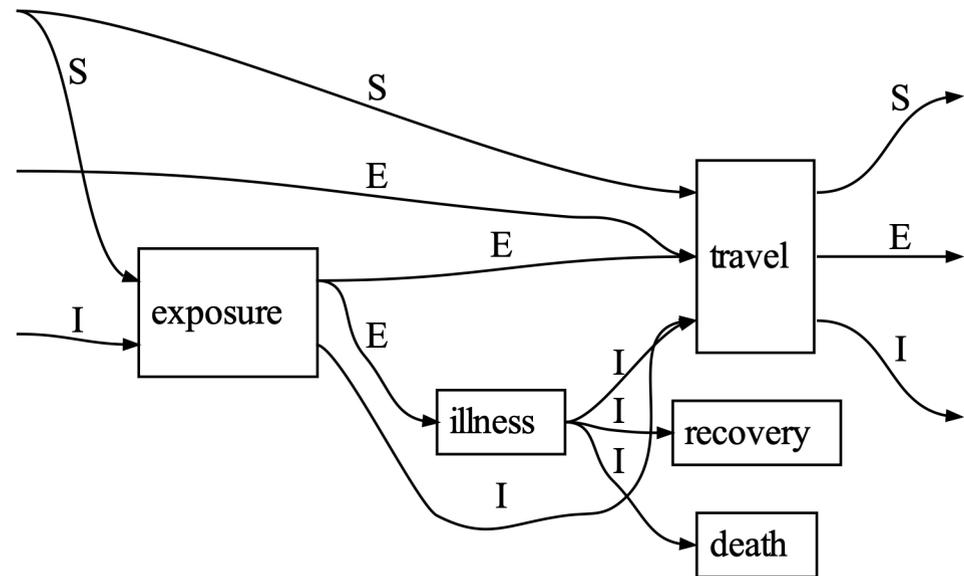
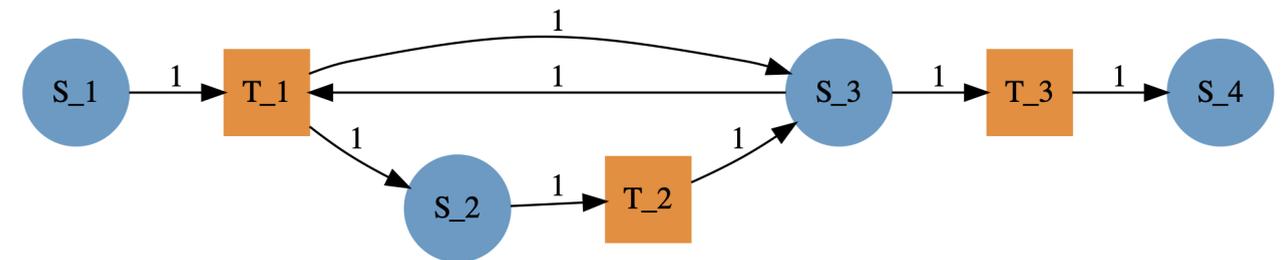
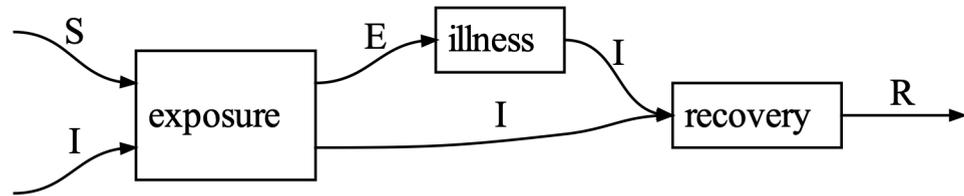
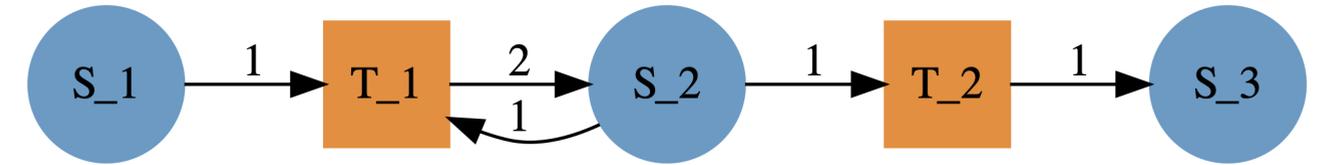
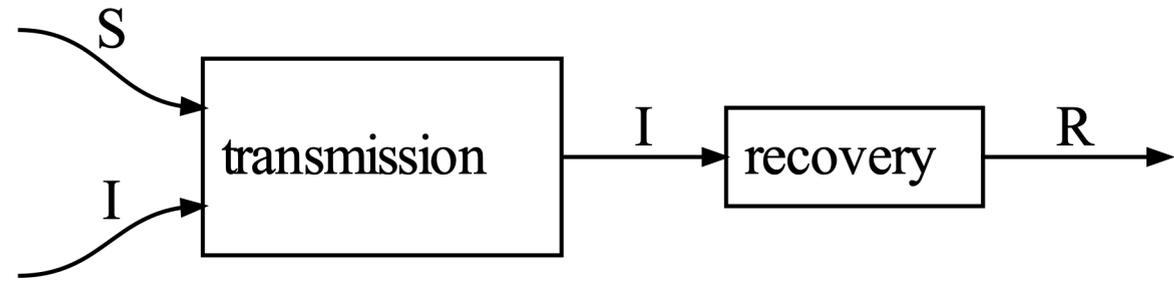
# Functorial Modeling Pipeline

$$Epi \xrightarrow{F} Petri \xrightarrow{K} Dynam$$



# EpiModels: SIR, SEIR, SEIRD with Travel

$Epi \xrightarrow{F} Petri$

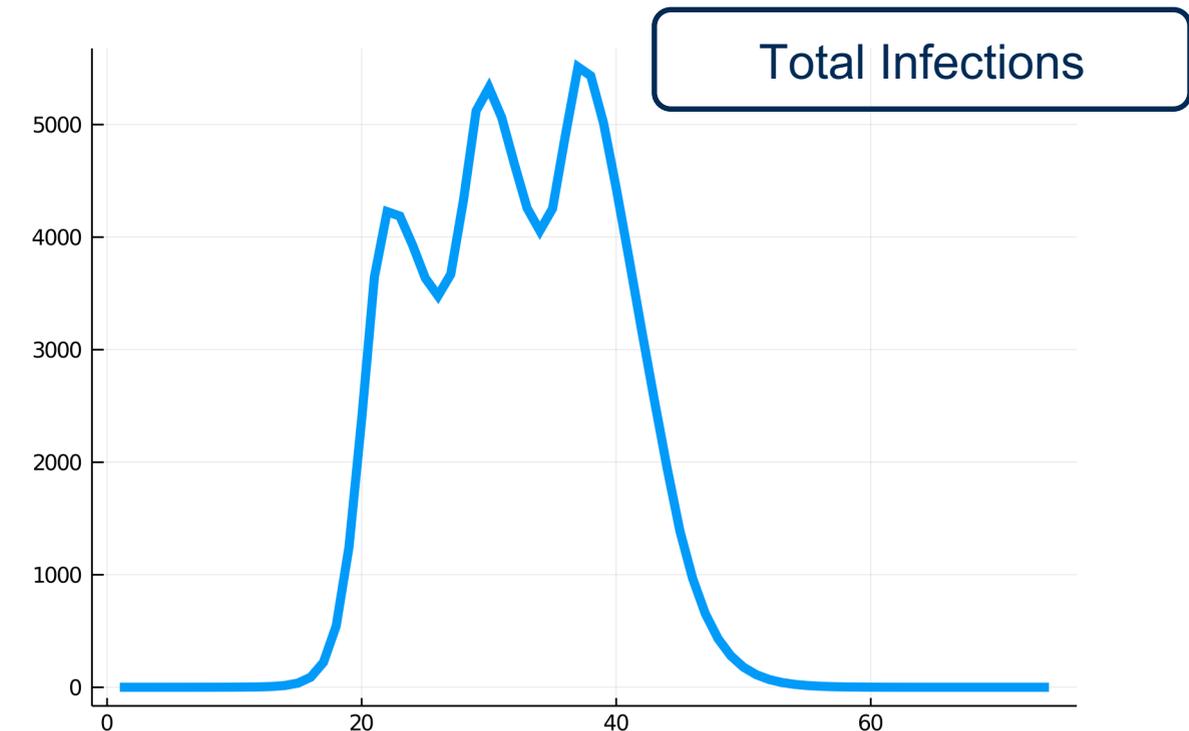
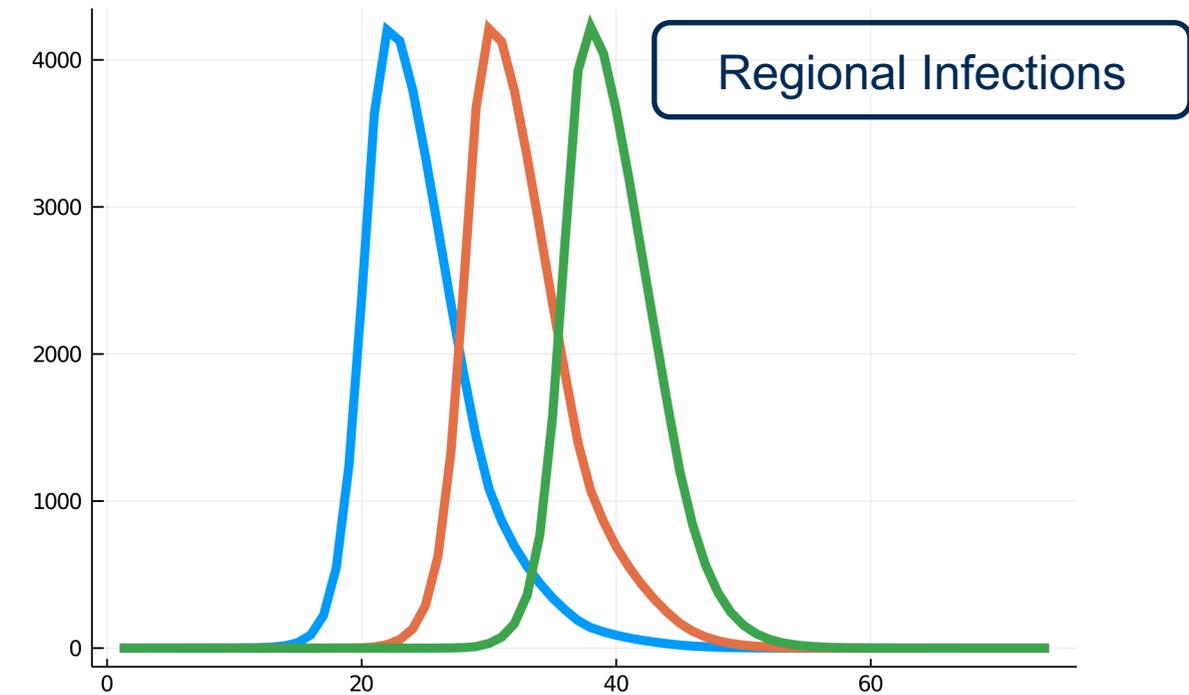
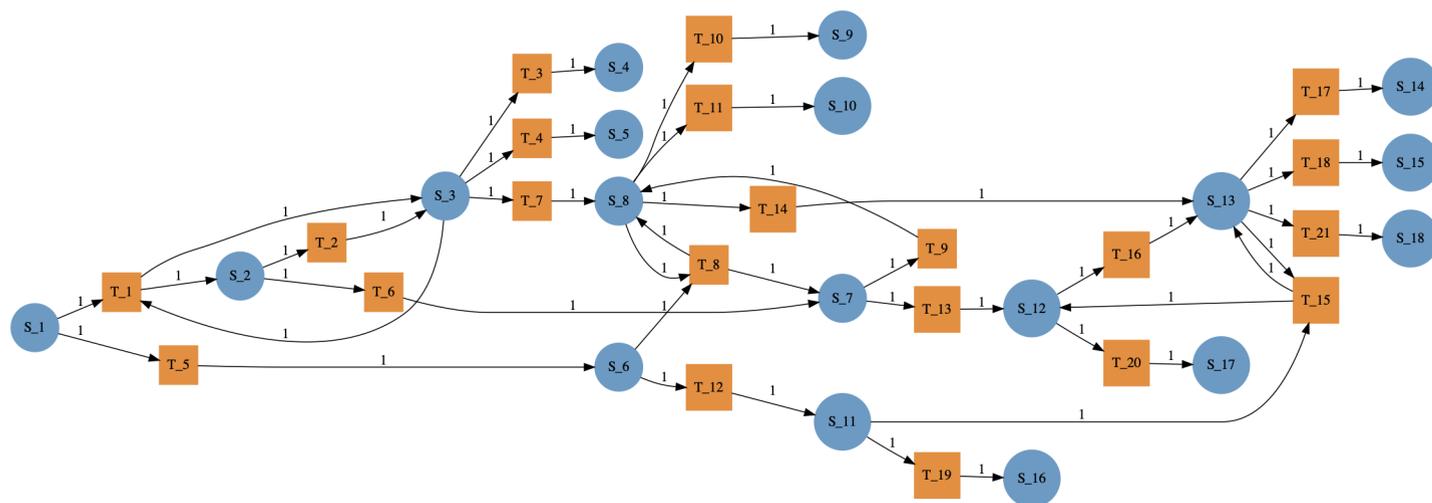
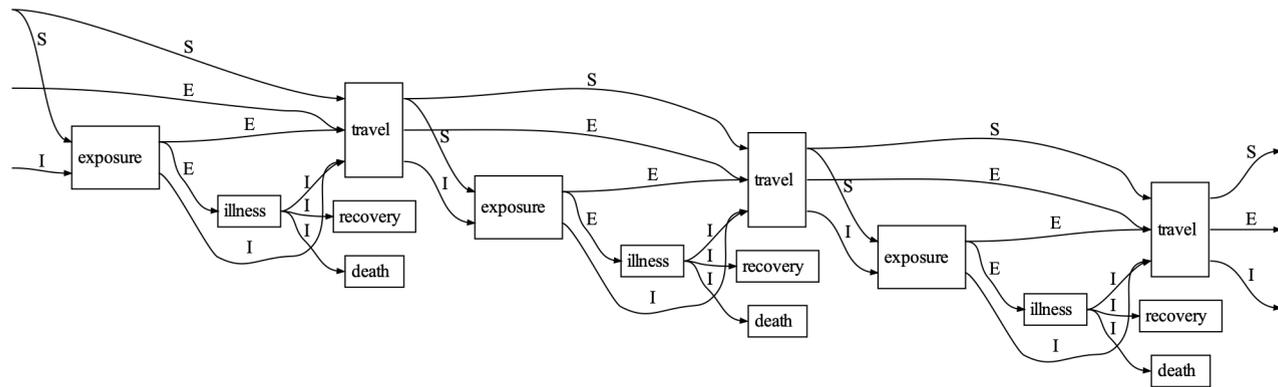


# Superposition of Regions Yield Multi-peaks



```
seird_city = @program Epidemiology (s::S, e::E, i::I) begin
  e2, i2 = exposure(s, i)
  i3 = illness(e2)
  d = death(i3)
  r = recovery(i3)
  return travel(s, [e, e2], [i2, i3])
end
```

```
^(f, n::Int) = fold(compose, repeat(f, n))
seird_3 = seird_city^3
```



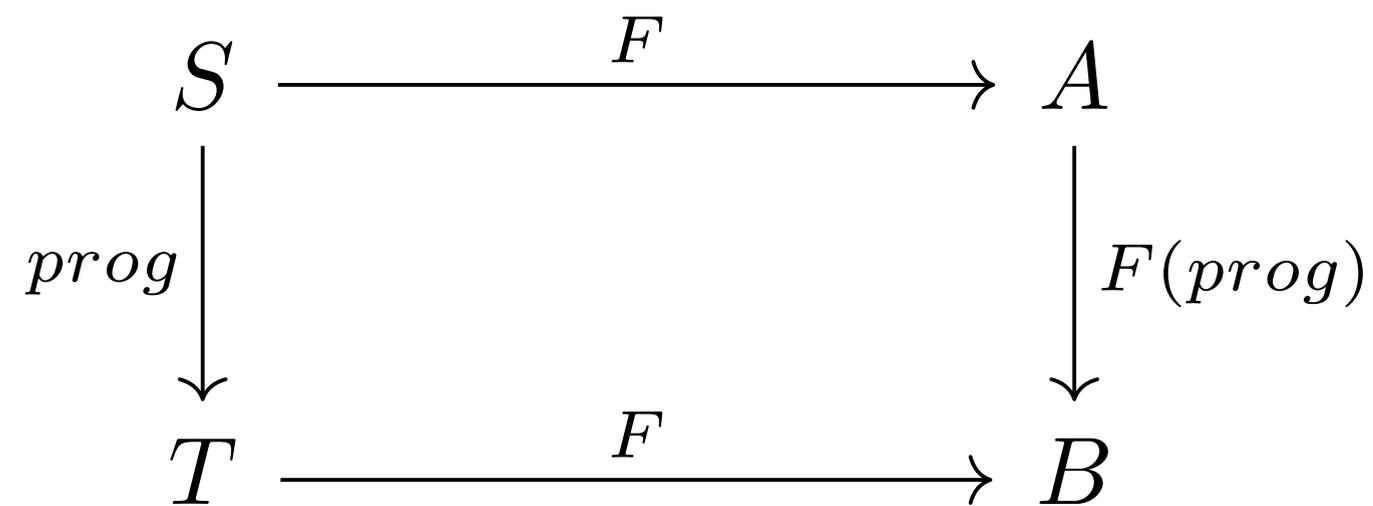
# Behavioral Approach to Modeling Systems



*Syntax*  $\longrightarrow$  *Semantics*

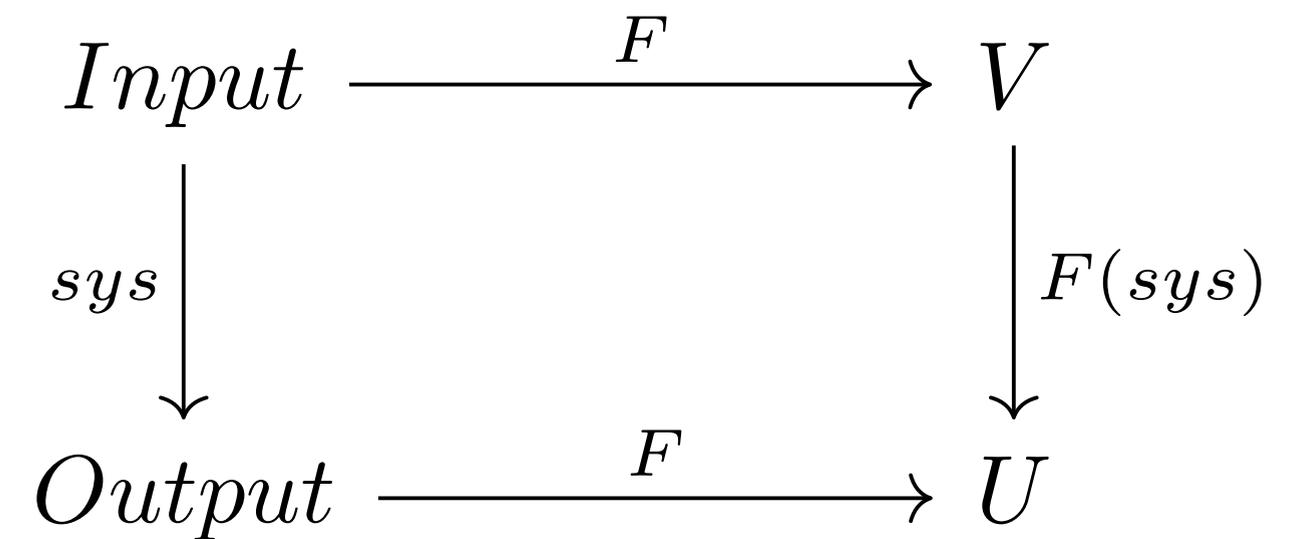
## Functional Programming

*Programs*  $\xrightarrow{F}$  *Functions*



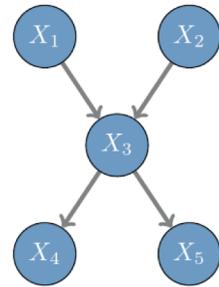
## Behavioral Semantics

*Designs*  $\xrightarrow{F}$  *Behaviors*



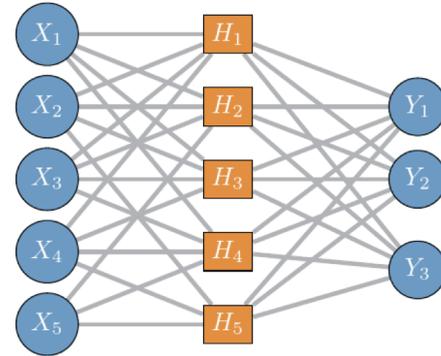
# Modeling Frameworks use Graphs + Math

Bayesian Network



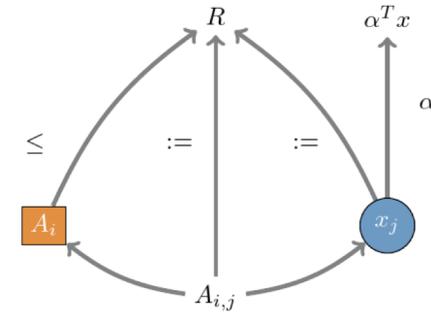
$$P(X_4, X_5 | X_3)P(X_3 | X_1, X_2)$$

Neural Network



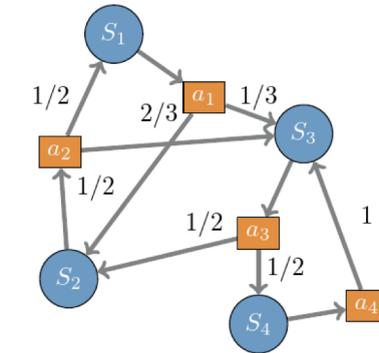
$$\min \sum_i \ell_\theta(y, x)$$

Optimization Problems



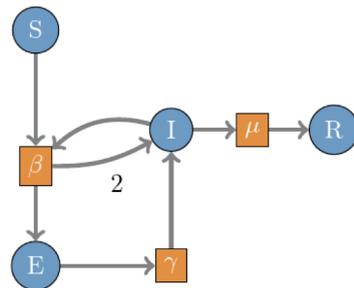
$$\min_x \alpha^T x \text{ s.t. } Ax < b$$

Markov Decision Process



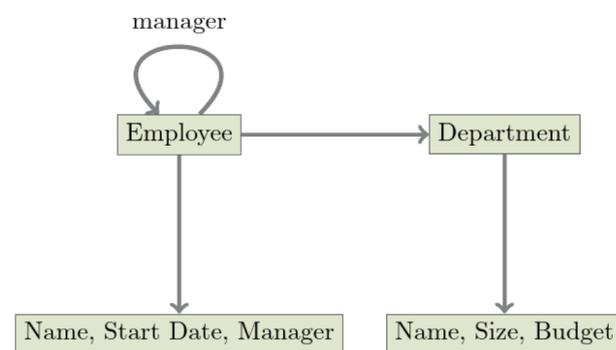
Construct  $\pi : State \rightarrow Action$  maximizing  $E[R]$

Petri Net



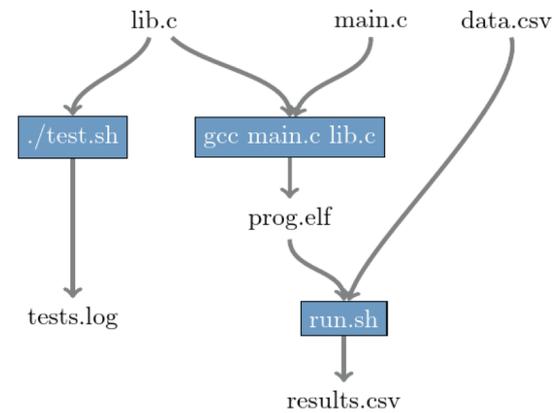
$$\text{Law of Mass Action: } \dot{u} = f(u, t)$$

Databases



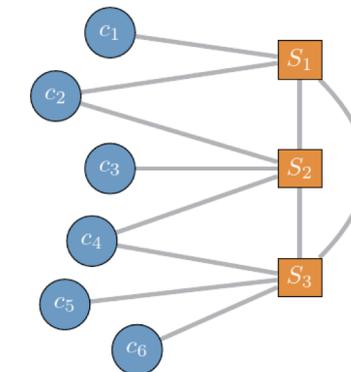
Relational Algebra

Dependency Graph



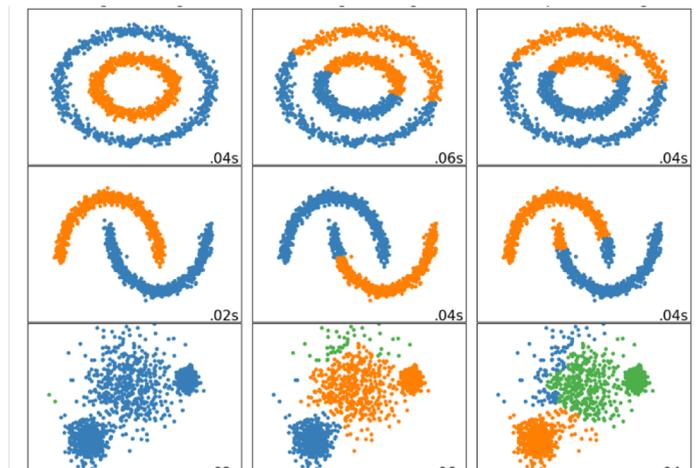
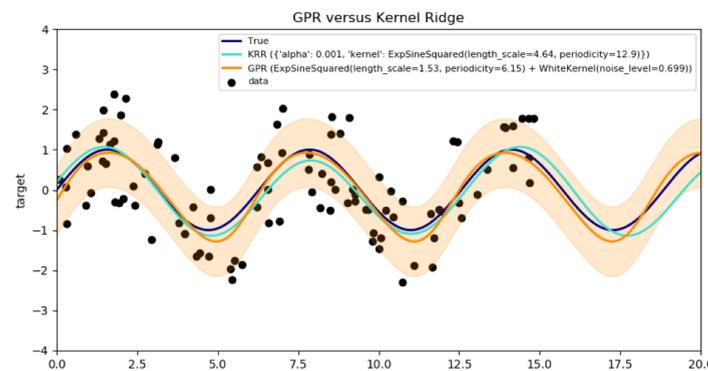
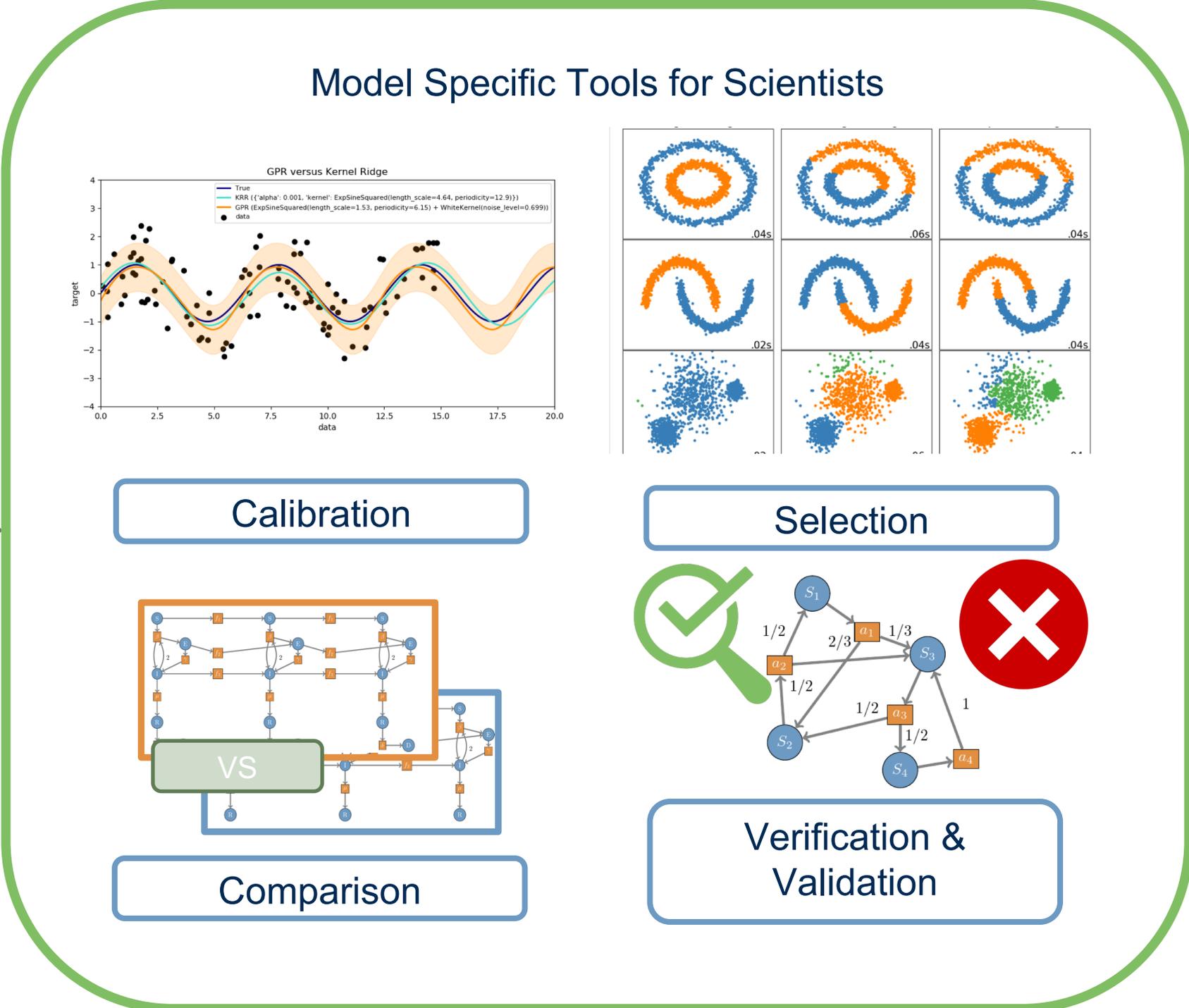
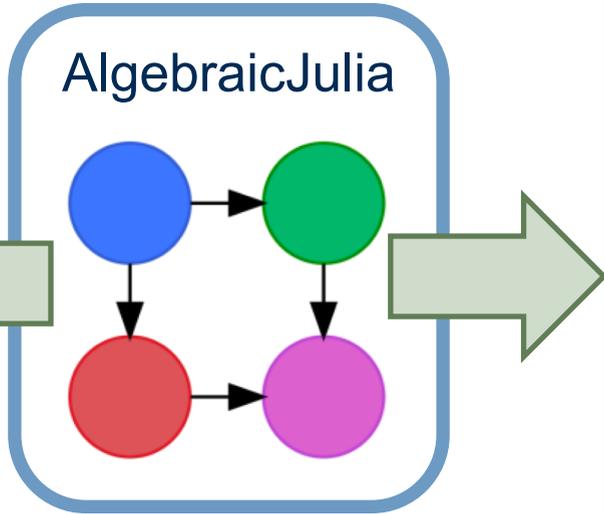
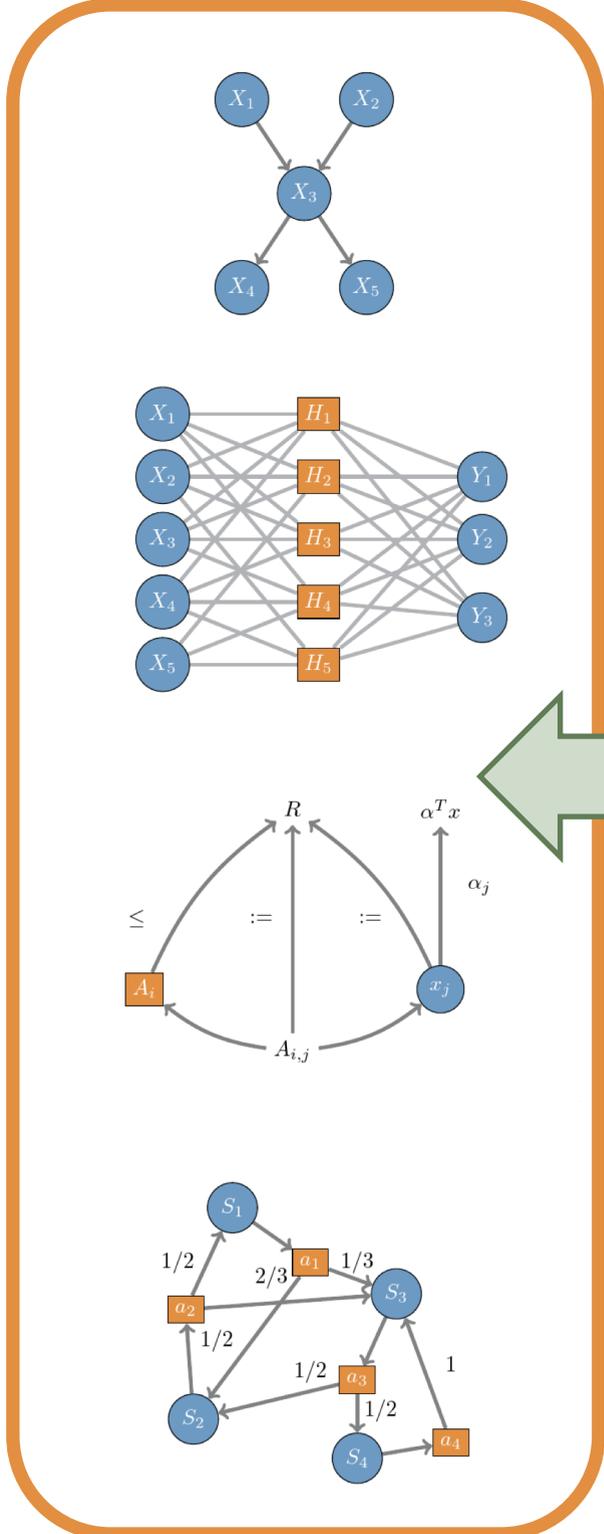
Scheduling and Compilation

Computer Network



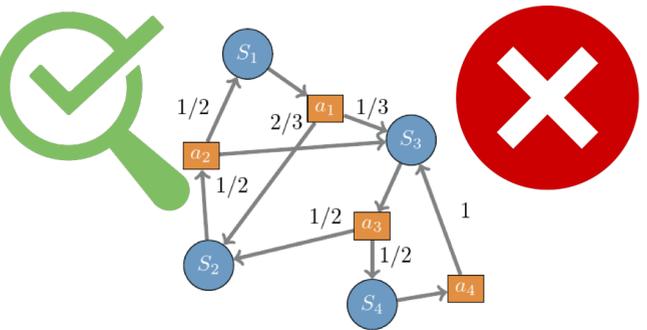
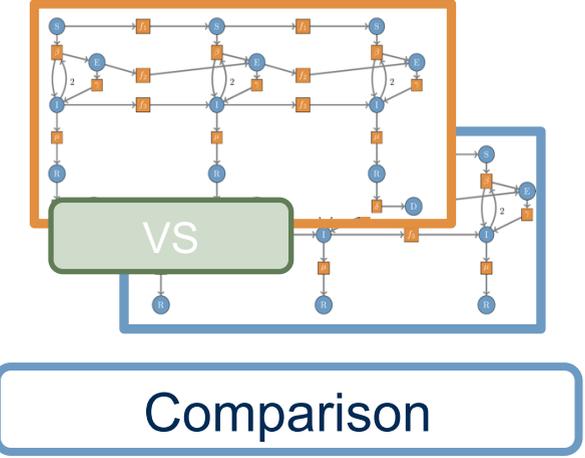
Bandwidth, Latency, Routing

# Model Aware Scientific Computing Vision



Calibration

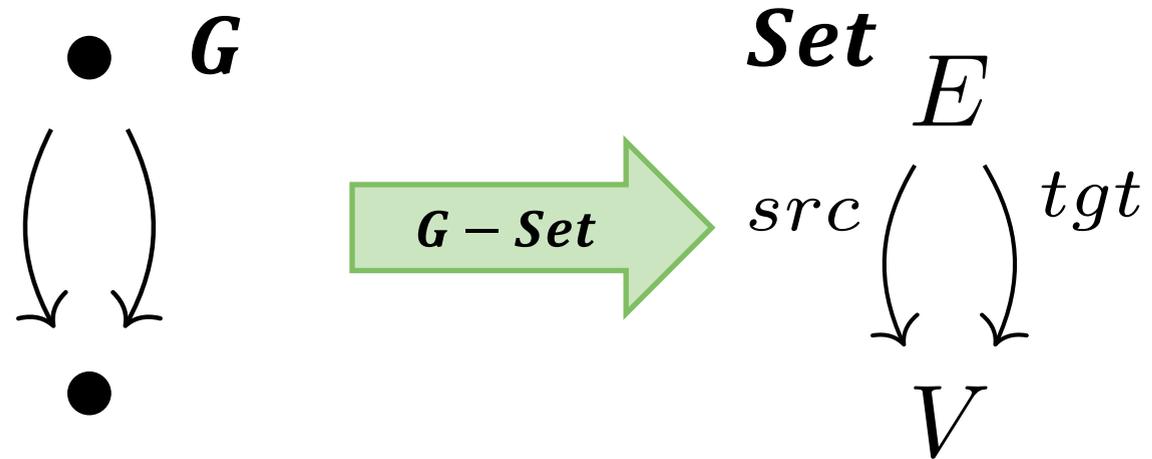
Selection



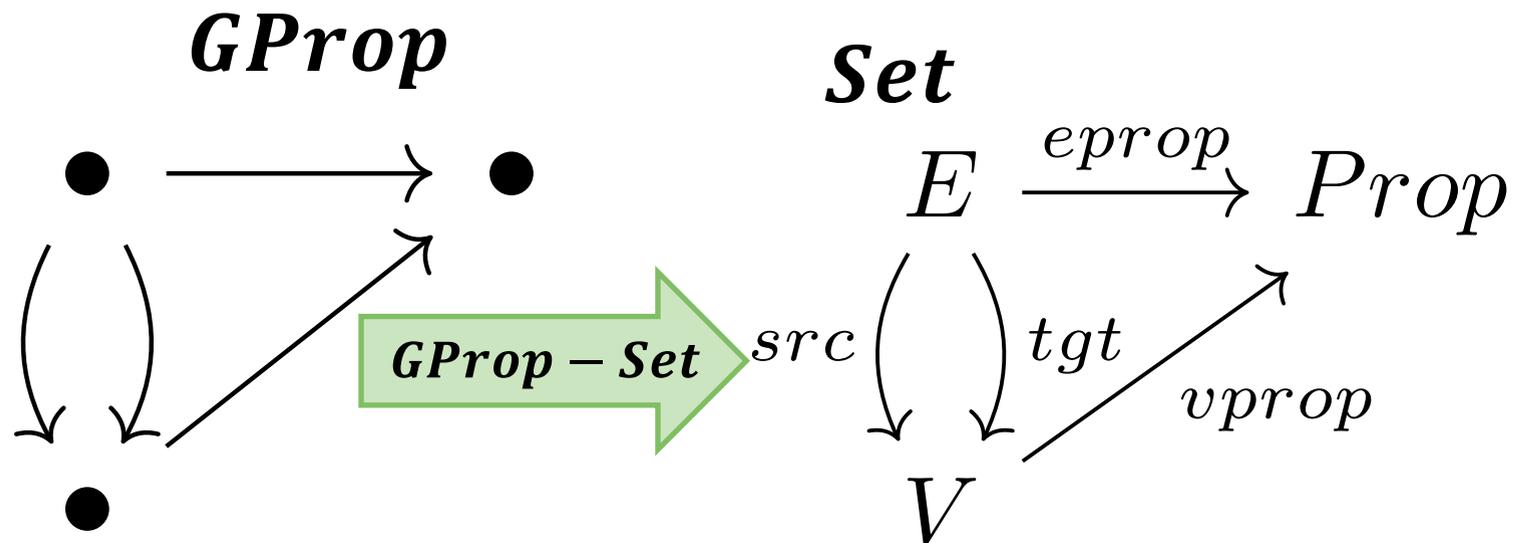
Verification & Validation

# C-Sets: Categorical Data Structures

## Logically extending a data structure with ACT principles



```
@present TheoryGraph (FreeCategory) begin
  V :: Ob
  E :: Ob
  src :: Hom (E, V)
  tgt :: Hom (E, V)
end
Graph = CSetType (TheoryGraph)
```

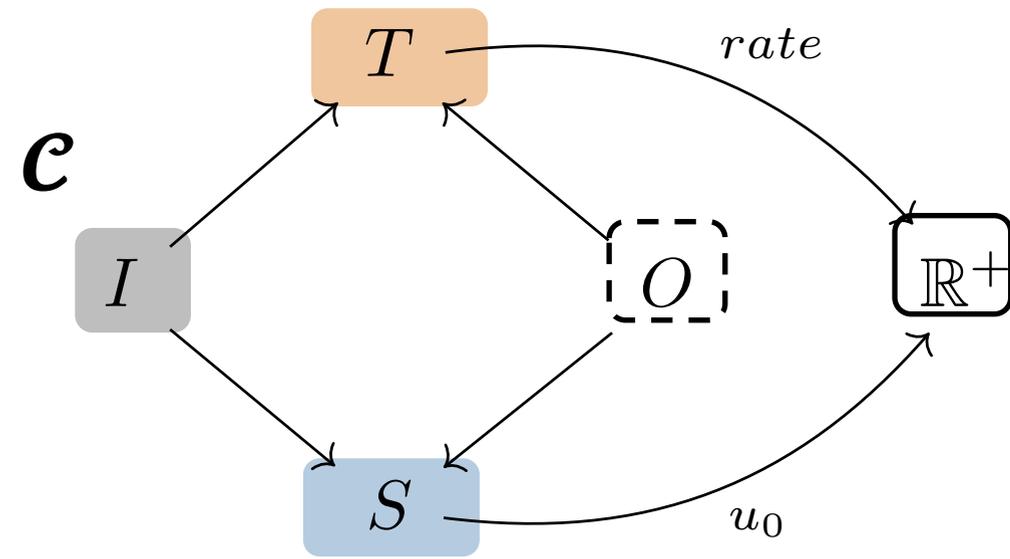


```
@present TheoryPropertyGraph <: TheoryGraph begin
  Prop :: Ob

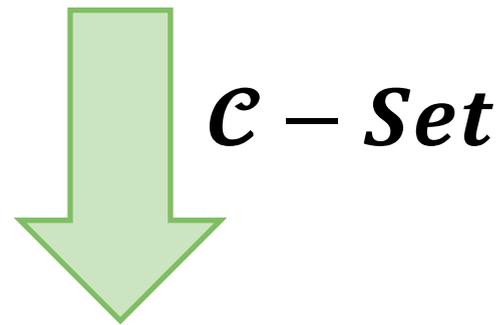
  vprops :: Hom (V, Prop)
  eprops :: Hom (E, Prop)
end

PropertyGraph = CSetType (TheoryPropertyGraph)
```

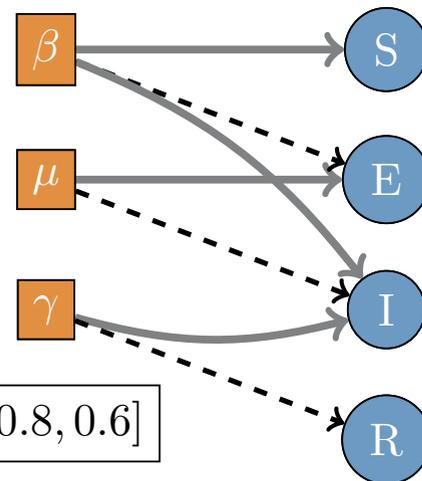
# $\mathcal{C}$ -Sets: Categorical Data Structures



- The data structure is a directed graph
- Instances are Sets with Functions
- Provides compositional framework for structured data
- Mathematically Elegant

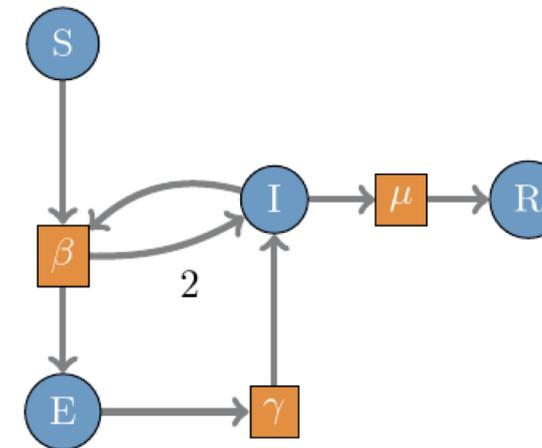
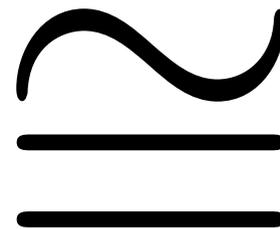


*Set*



$$\beta, \mu, \gamma = [1.2, 0.8, 0.6]$$

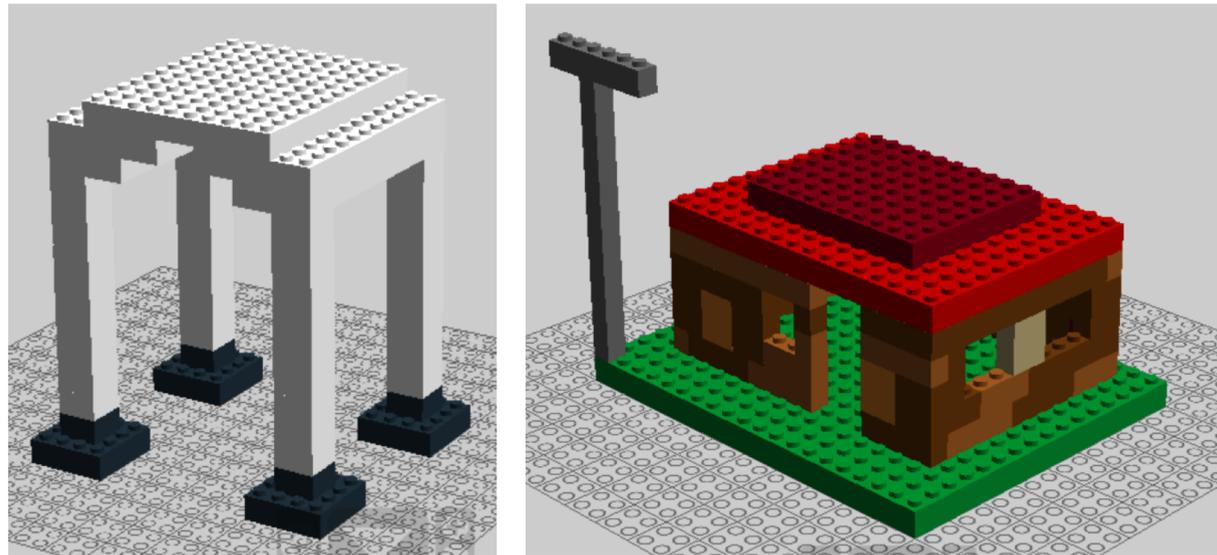
$$S, E, I, R = [0.99, 0.01, 0, 0]$$



$$\beta, \mu, \gamma = [1.2, 0.8, 0.6]$$

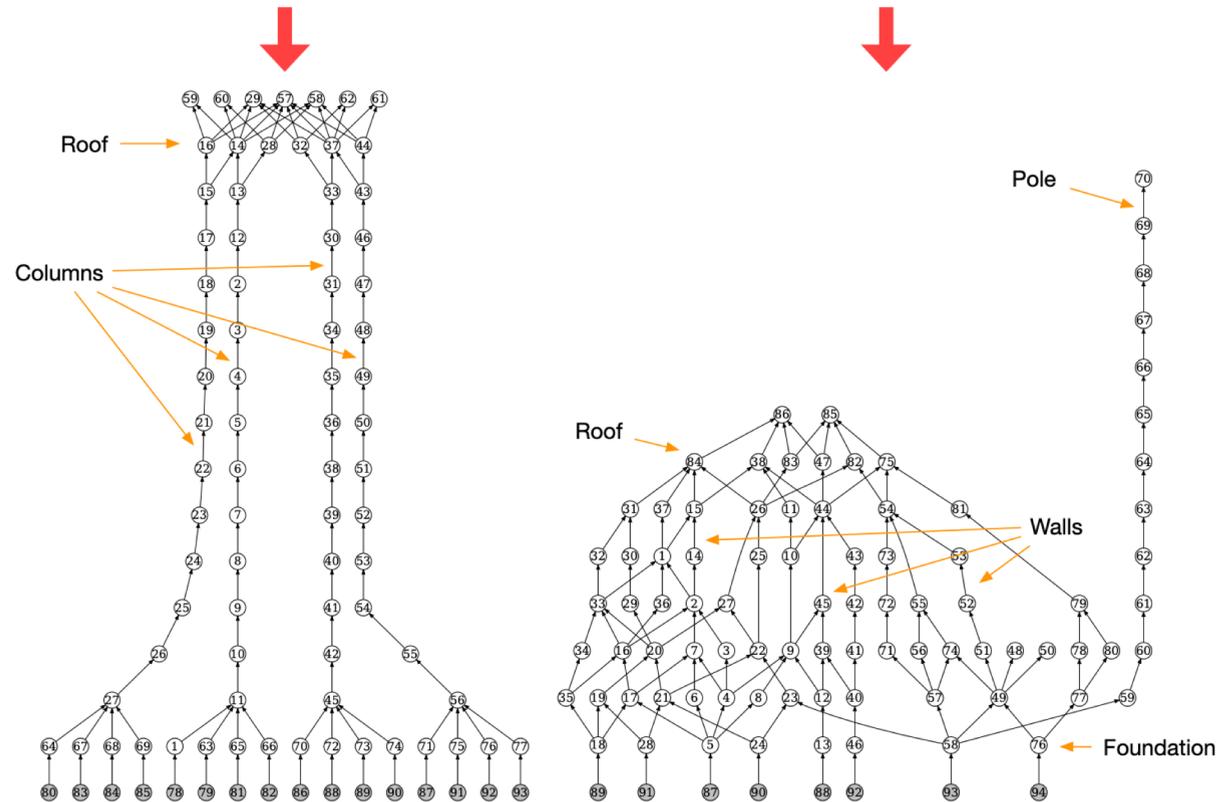
$$S, E, I, R = [0.99, 0.01, 0, 0]$$

# Automatic Parallelization for Construction



a) Columns - 77 bricks

b) House - 86 bricks



c) Columns - connectivity diagram

d) House - connectivity diagram

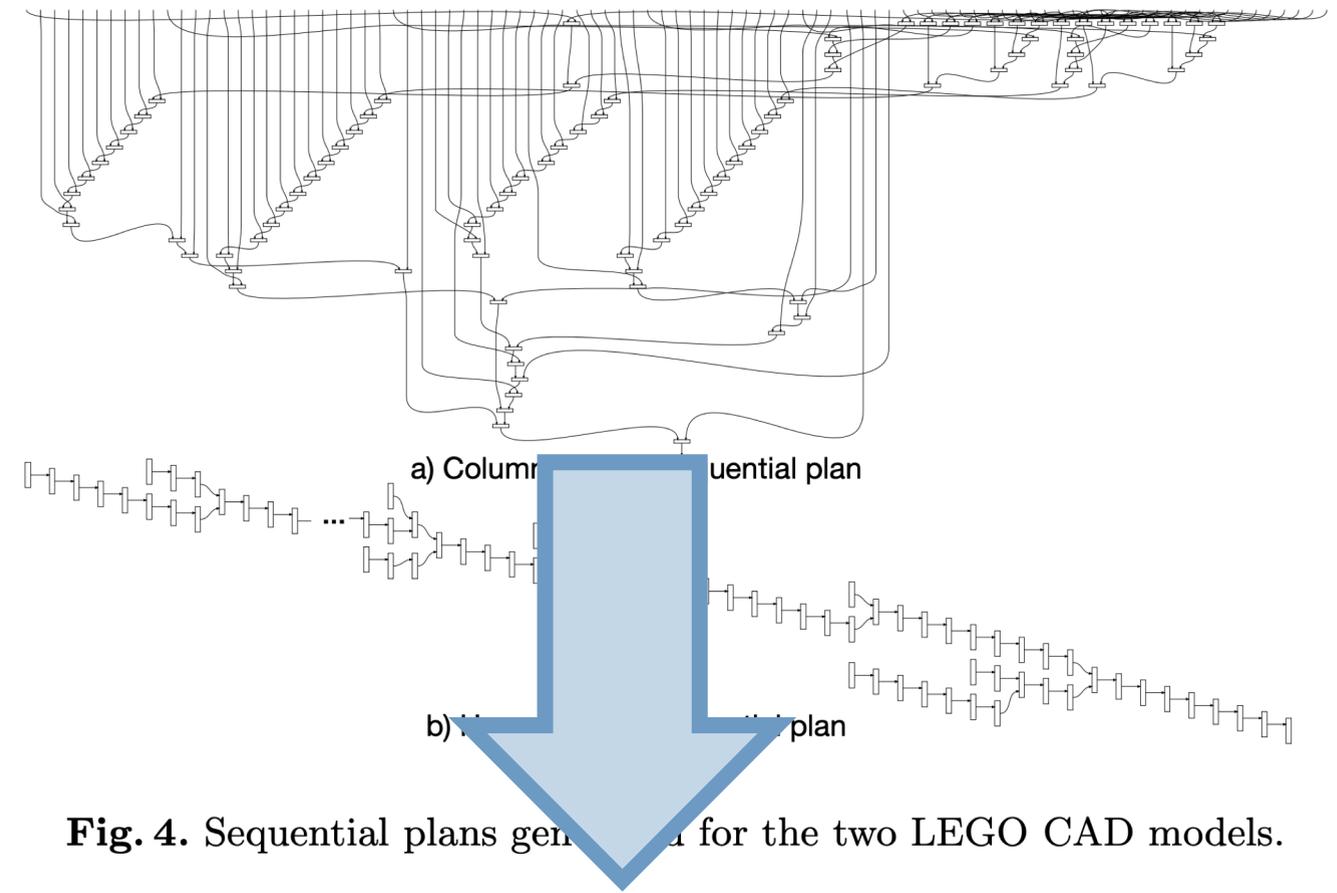
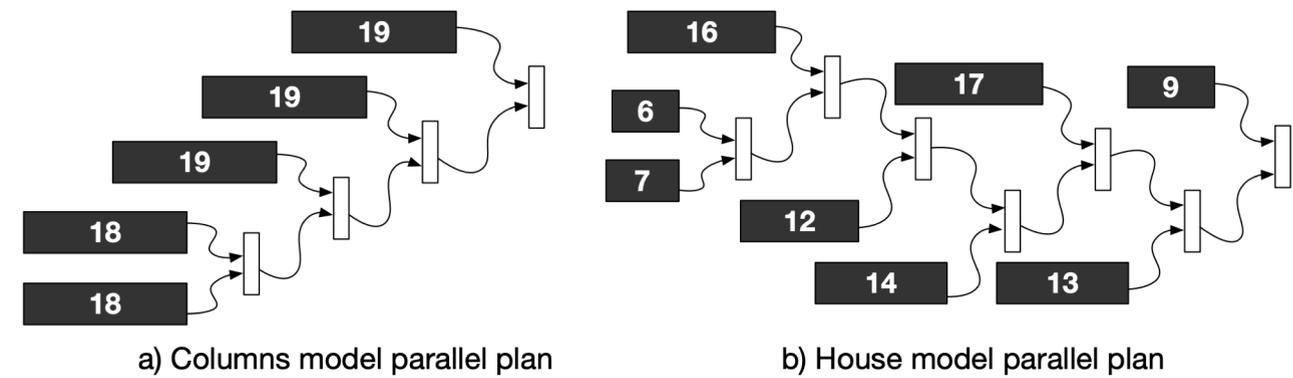


Fig. 4. Sequential plans generated for the two LEGO CAD models.



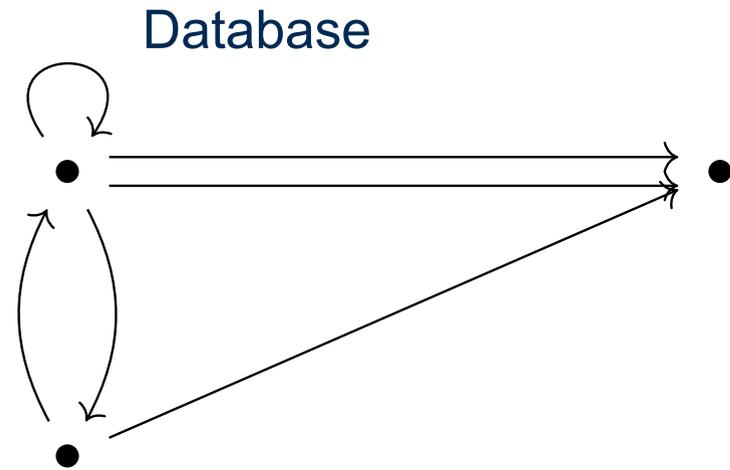
a) Columns model parallel plan

b) House model parallel plan

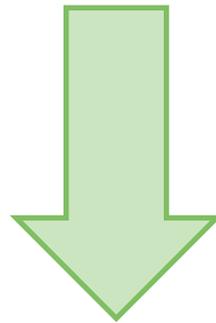
Fig. 5. Parallel plans generated for the two LEGO CAD models. The width of the black boxes represent a sub-plan (i.e., a stairs configuration).

# Generating SQL Queries from *BiCatRel*

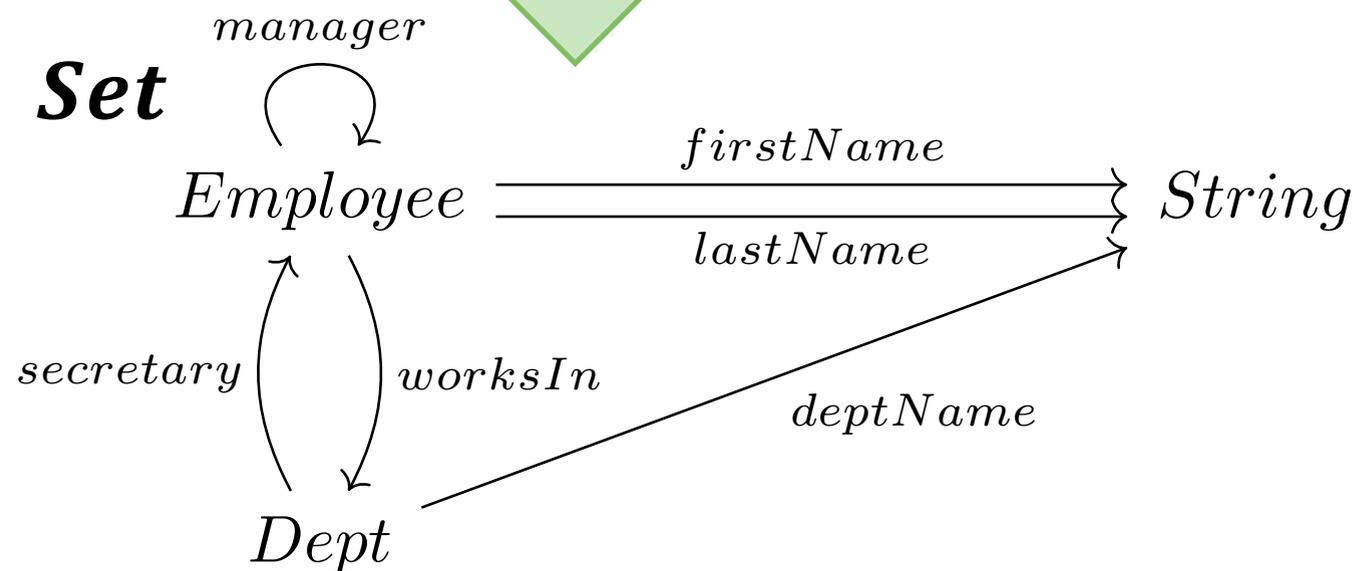
**Schema**



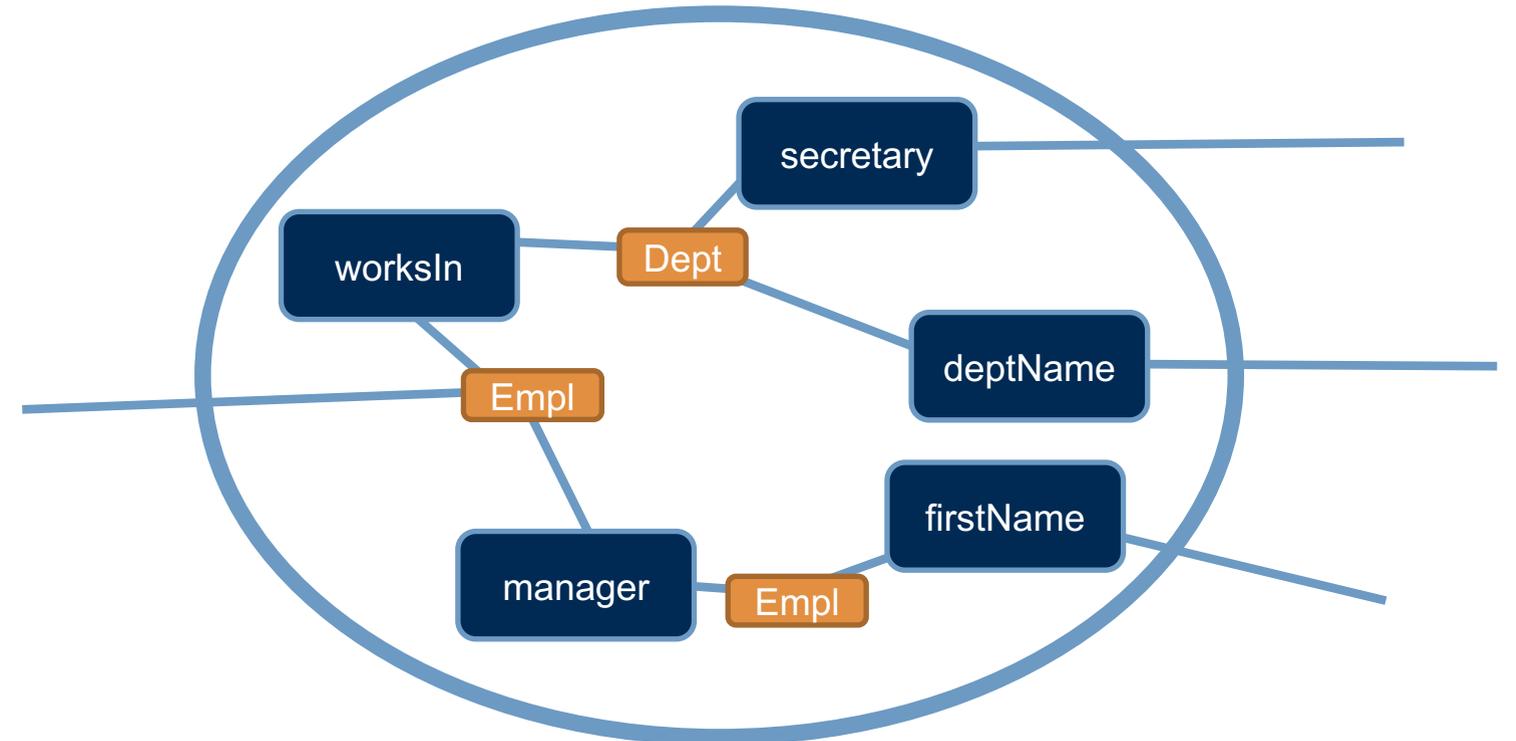
**Database**



**Set**

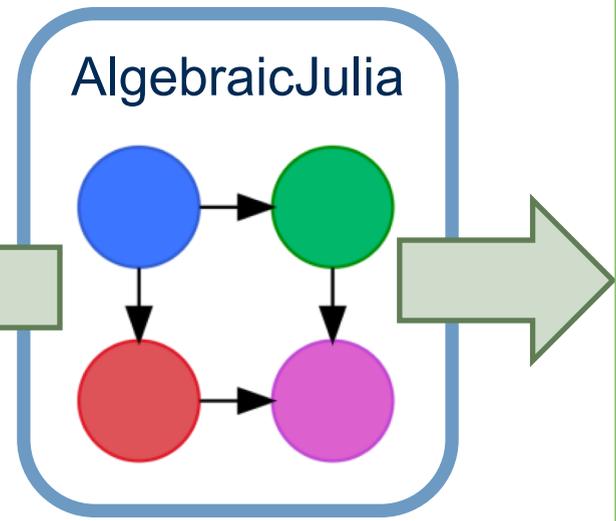
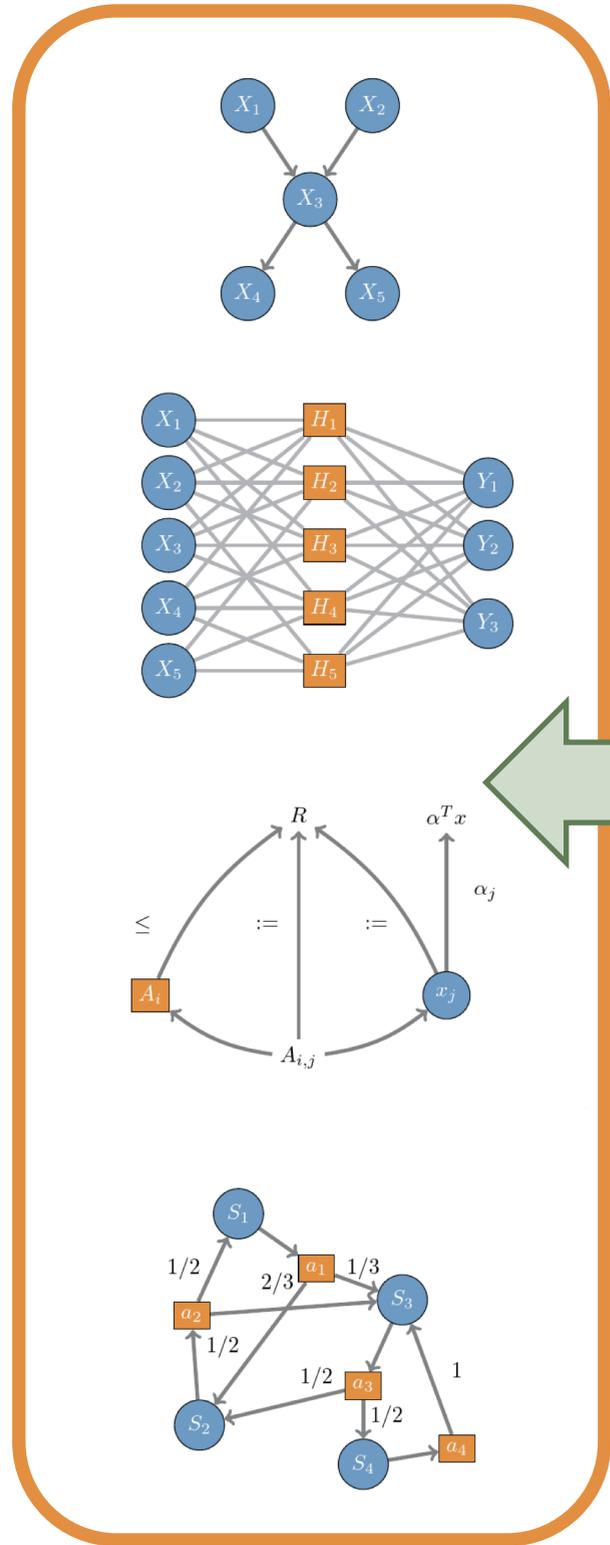


**Query**

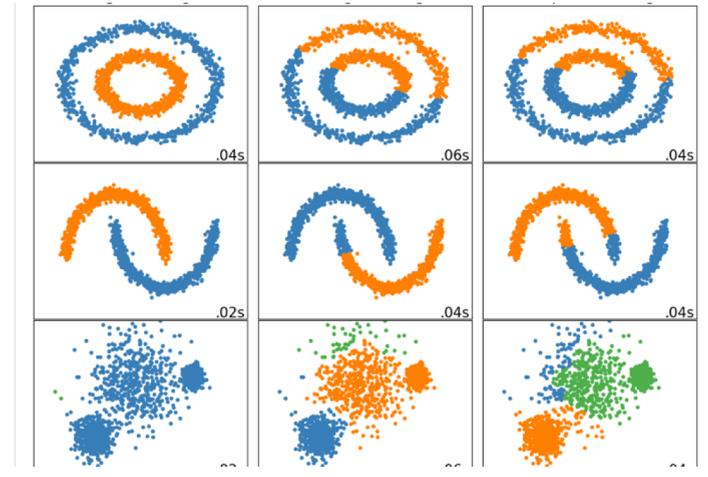
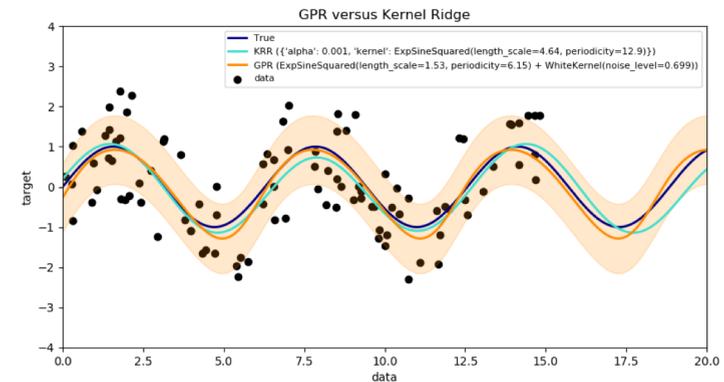


```
SELECT t4.employee AS employee,
       t6.secretary AS secretary, t7.name AS name,
       t9.first_name AS first_name
FROM worksIn AS t4, secretary AS t6, deptName AS t7,
     manager AS t8, firstName AS t9
WHERE t4.employee=t8.employee AND
      t4.department=t6.department AND
      t4.department=t7.department AND
      t8.employee=t9.employee;
```

# Model Aware Scientific Computing

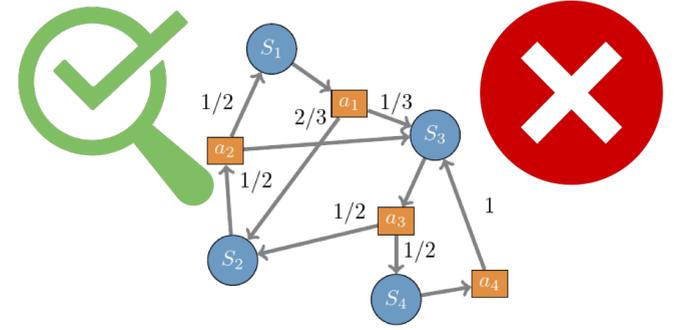
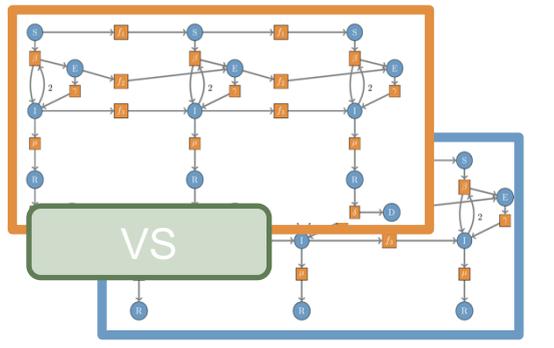


## Model Specific Tools for Scientists



Calibration

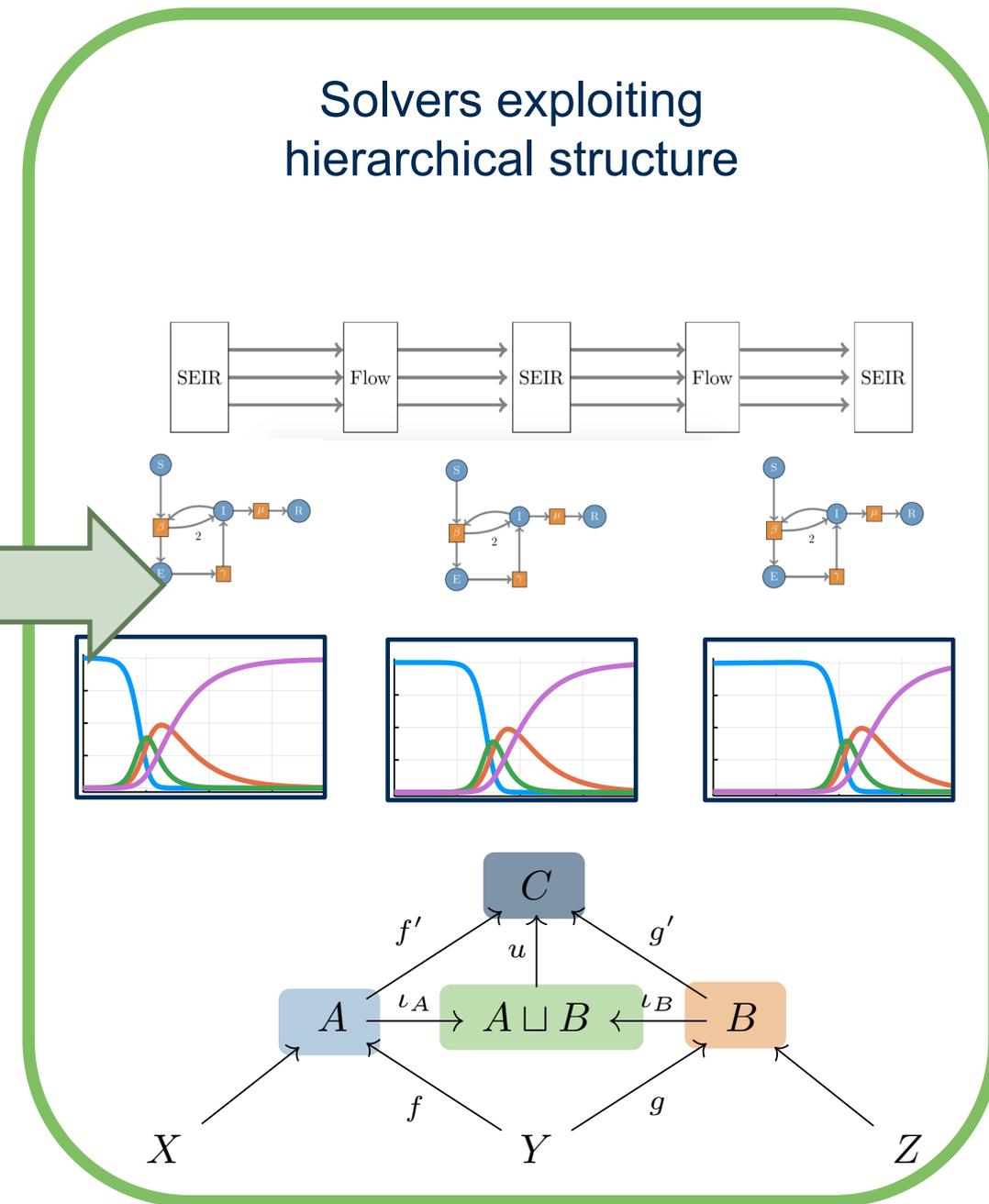
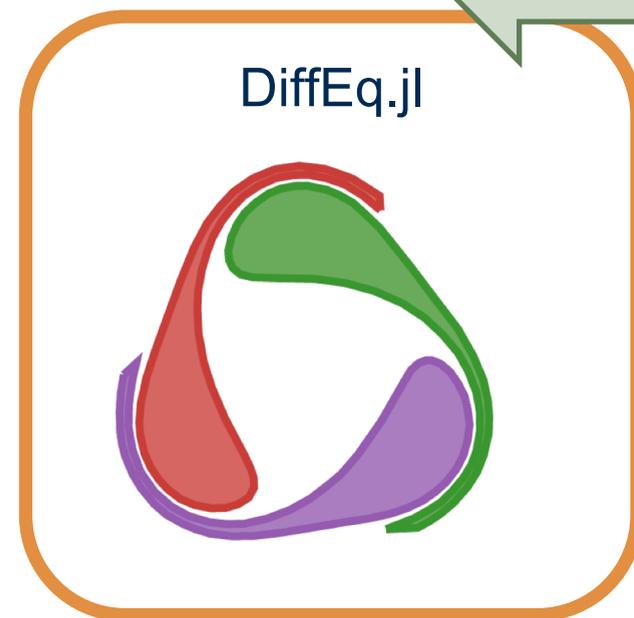
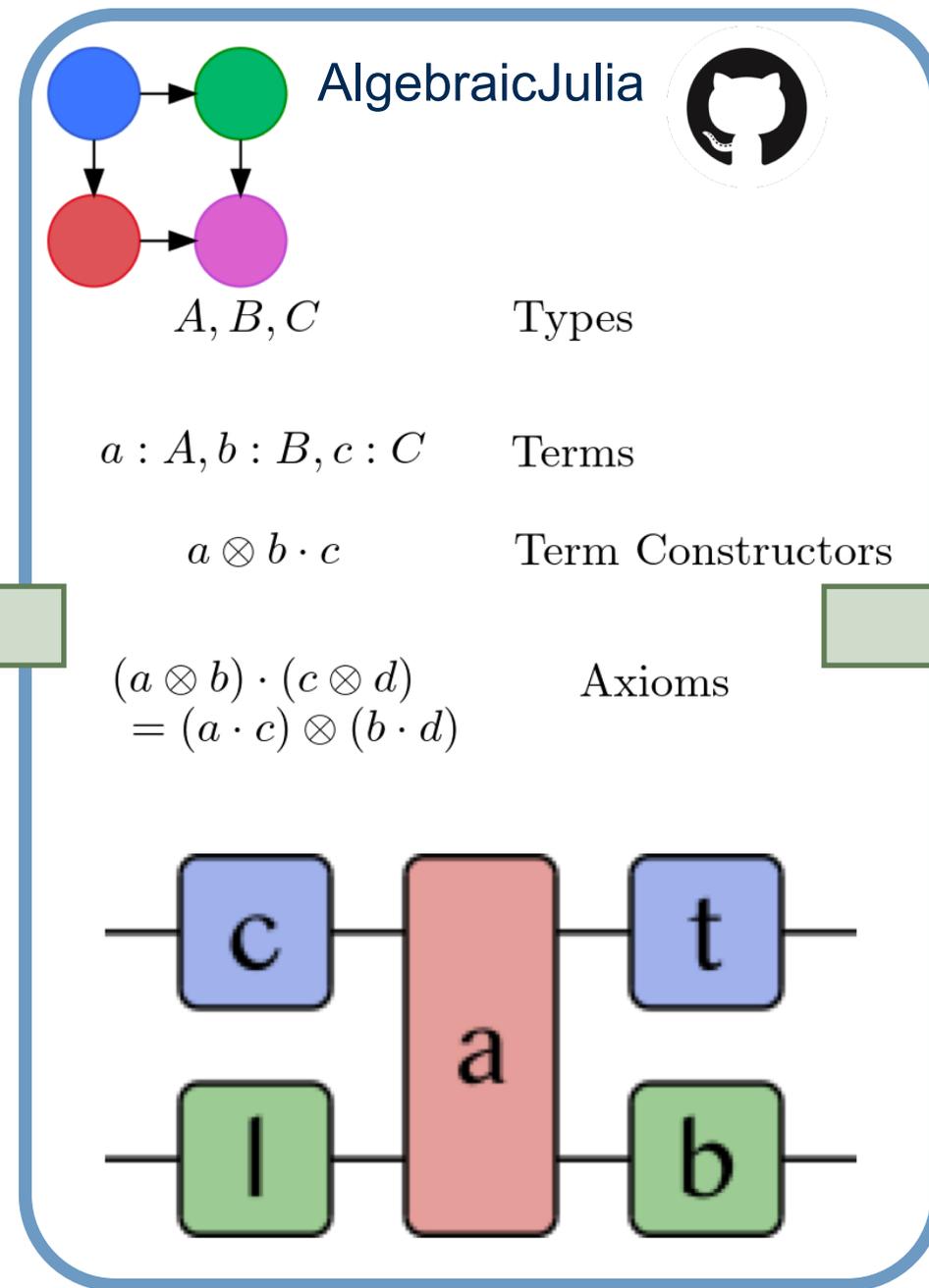
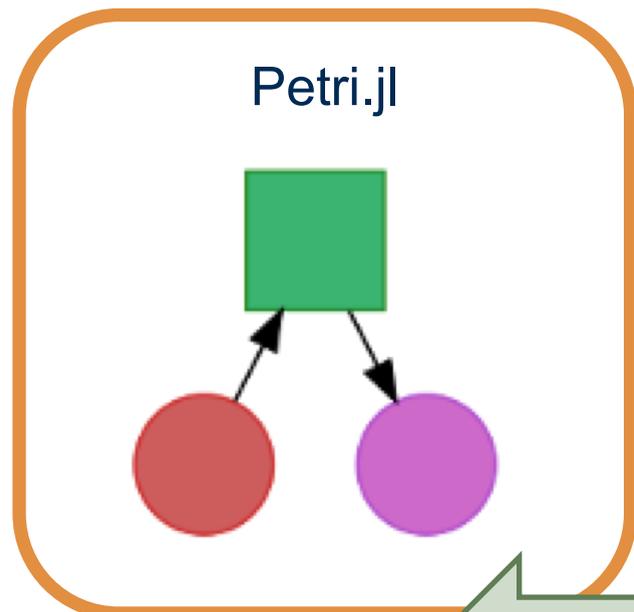
Selection



Comparison

Verification & Validation

# AlgebraicJulia Ecosystem



# Algebraic Julia Team



Evan Patterson



Micah Halter



Sophie Libkind



Andrew Baas

