

## SBML Model Report

# Model name: “Vizan2013 - TGF pathway long term signaling”



February 24, 2014

## 1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by the following two authors: Pedro Vizan<sup>1</sup> and Nick Juty<sup>2</sup> at December 16<sup>th</sup> 2013 at 11:13 a. m. and last time modified at February 24<sup>th</sup> 2014 at 9:47 a. m. Table 1 provides an overview of the quantities of all components of this model.

Table 1: Number of components in this model, which are described in the following sections.

Element	Quantity	Element	Quantity
compartment types	0	compartments	1
species types	0	species	26
events	0	constraints	0
reactions	0	function definitions	0
global parameters	29	unit definitions	3
rules	30	initial assignments	6

## 2 Unit Definitions

This is an overview of five unit definitions of which two are predefined by SBML and not mentioned in the model.

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## 2.1 Unit volume

**Name** volume

**Definition** dimensionless

## 2.2 Unit time

**Name** time

**Definition** 3600 s

## 2.3 Unit substance

**Name** substance

**Definition** dimensionless

## 2.4 Unit area

**Notes** Square metre is the predefined SBML unit for area since SBML Level 2 Version 1.

**Definition** m<sup>2</sup>

## 2.5 Unit length

**Notes** Metre is the predefined SBML unit for length since SBML Level 2 Version 1.

**Definition** m

# 3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
compartment_1	Cell		3	1	dimensionless	<input checked="" type="checkbox"/>	

## 3.1 Compartment compartment\_1

This is a three dimensional compartment with a constant size of one dimensionless.

**Name** Cell

## 4 Species

This model contains 26 species. The boundary condition of 25 of these species is set to `true` so that these species' amount cannot be changed by any reaction. Section 8 provides further details and the derived rates of change of each species.

Table 3: Properties of each species.

Id	Name	Compartment	Derived Unit	Constant	Boundary Condition
species_1	S22	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_2	S24	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_3	pS2tot	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_4	TGF	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_5	R	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_6	S2c	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_7	Rcom	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_8	pS2c	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_9	Rcom_S	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓
species_10	S2n	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	✓

Id	Name	Compartment	Derived Unit	Constant	Boundary Condition
species_11	S22n	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_12	S4n	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_13	S22c	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_14	pS2n	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_15	pS2fn	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_16	S24n	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_17	S24c	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_18	S4fc	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_19	S4c	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_20	pS2fc	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_21	S4fn	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_22	SBI	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input type="checkbox"/>
species_23	Rtot	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>

Id	Name	Compartment	Derived Unit	Constant	Boundary Condition
species_24	RT	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_25	Rcom_I	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>
species_26	Ract	compartment_1	dimensionless dimensionless <sup>-1</sup>	· <input type="checkbox"/>	<input checked="" type="checkbox"/>

## 5 Parameters

This model contains 29 global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
parameter_1	kd	0000009	0.320		<input checked="" type="checkbox"/>
parameter_2	kex	0000009	20.000		<input checked="" type="checkbox"/>
parameter_3	kin	0000009	9.360		<input checked="" type="checkbox"/>
parameter_4	alpha	0000009	0.080		<input checked="" type="checkbox"/>
parameter_5	CHX	0000390	0.000		<input checked="" type="checkbox"/>
parameter_6	kp	0000009	21.372		<input checked="" type="checkbox"/>
parameter_7	kdp	0000009	24.000		<input checked="" type="checkbox"/>
parameter_8	koff	0000282	60.000		<input checked="" type="checkbox"/>
parameter_9	kon	0000337	350.877		<input type="checkbox"/>
parameter_10	KDiss	0000282	0.171		<input type="checkbox"/>
parameter_11	CIF	0000380	5.700		<input checked="" type="checkbox"/>
parameter_12	D	0000360	4.000		<input checked="" type="checkbox"/>
parameter_13	a	0000360	2.270		<input checked="" type="checkbox"/>
parameter_14	S2tot	0000361	1.000		<input checked="" type="checkbox"/>
parameter_15	S4tot	0000361	1.000		<input checked="" type="checkbox"/>
parameter_16	rc0	0000540	0.050		<input checked="" type="checkbox"/>
parameter_17	KSBI	0000282	0.197		<input checked="" type="checkbox"/>
parameter_18	k'T	0000009	100.000		<input checked="" type="checkbox"/>
parameter_19	kex4	0000009	9.360		<input type="checkbox"/>
parameter_20	Total Nuc S2 for fit	0000360	1.000		<input type="checkbox"/>
parameter_21	Ktr	0000281	0.711		<input type="checkbox"/>
parameter_22	k'act	0000009	24.538		<input checked="" type="checkbox"/>
parameter_23	Tmax in ng/ml	0000470	2.000		<input checked="" type="checkbox"/>
parameter_24	TSca	0000470	2.000		<input checked="" type="checkbox"/>
parameter_25	k'cc	0000009	0.350		<input checked="" type="checkbox"/>
parameter_26	k'synT	0000009	0.000		<input checked="" type="checkbox"/>
parameter_27	k'synTbas	0000009	0.000		<input checked="" type="checkbox"/>
parameter_28	MG132	0000009	0.000		<input checked="" type="checkbox"/>
Metabolite_9	Initial for S2n	0000360	0.559		<input checked="" type="checkbox"/>

## 6 Initialassignments

This is an overview of six initialassignments.

### 6.1 Initialassignment species\_4

**Derived unit** contains undeclared units

**Math**  $\text{parameter\_23} \cdot \text{parameter\_24}$

### 6.2 Initialassignment species\_5

**Derived unit** contains undeclared units

**Math**  $\frac{1 - \text{parameter\_16}}{\text{parameter\_4} + 1}$

### 6.3 Initialassignment species\_6

**Derived unit** contains undeclared units

**Math**  $\frac{\text{parameter\_14} \cdot \text{parameter\_2} \cdot (1 + \text{parameter\_13})}{\text{parameter\_3} + \text{parameter\_13} \cdot \text{parameter\_2}}$

### 6.4 Initialassignment species\_7

**Derived unit** contains undeclared units

**Math**  $\frac{\text{parameter\_16} + \text{parameter\_4}}{1 + \text{parameter\_4}}$

### 6.5 Initialassignment species\_19

**Derived unit** contains undeclared units

**Math**  $\text{parameter\_15}$

### 6.6 Initialassignment Metabolite\_9

**Derived unit**  $\text{dimensionless}^{-1}$

**Math**  $[\text{species\_10}]$

## 7 Rules

This is an overview of 30 rules.

### 7.1 Rule species\_16

Rule species\_16 is an assignment rule for species species\_16:

$$\text{species\_16} = (\text{parameter\_13} + 1) \cdot [\text{species\_2}] - \text{parameter\_13} \cdot [\text{species\_17}] \quad (1)$$

## 7.2 Rule species\_18

Rule species\_18 is an assignment rule for species species\_18:

$$\text{species\_18} = [\text{species\_19}] - [\text{species\_17}] \quad (2)$$

**Derived unit** dimensionless<sup>-1</sup>

## 7.3 Rule parameter\_10

Rule parameter\_10 is an assignment rule for parameter parameter\_10:

$$\text{parameter\_10} = 0.171 \quad (3)$$

## 7.4 Rule parameter\_9

Rule parameter\_9 is an assignment rule for parameter parameter\_9:

$$\text{parameter\_9} = \frac{\text{parameter\_8}}{\text{parameter\_10}} \quad (4)$$

## 7.5 Rule parameter\_19

Rule parameter\_19 is an assignment rule for parameter parameter\_19:

$$\text{parameter\_19} = \text{parameter\_3} \quad (5)$$

## 7.6 Rule parameter\_21

Rule parameter\_21 is an assignment rule for parameter parameter\_21:

$$\text{parameter\_21} = \frac{\text{parameter\_16}}{1 - \text{parameter\_16}} \cdot \frac{\text{parameter\_4} + 1}{\text{parameter\_4}} \quad (6)$$

## 7.7 Rule species\_20

Rule species\_20 is an assignment rule for species species\_20:

$$\text{species\_20} = [\text{species\_8}] - 2 \cdot [\text{species\_13}] - [\text{species\_17}] \quad (7)$$

## 7.8 Rule species\_23

Rule species\_23 is an assignment rule for species species\_23:

$$\text{species\_23} = [\text{species\_5}] + [\text{species\_7}] + [\text{species\_24}] + [\text{species\_26}] \quad (8)$$

**Derived unit** dimensionless<sup>-1</sup>



### 7.9 Rule species\_25

Rule species\_25 is an assignment rule for species species\_25:

$$\text{species\_25} = [\text{species\_7}] \cdot \frac{1}{1 + \text{parameter\_21}} \quad (9)$$

### 7.10 Rule species\_9

Rule species\_9 is an assignment rule for species species\_9:

$$\text{species\_9} = [\text{species\_7}] \cdot \frac{\text{parameter\_21}}{1 + \text{parameter\_21}} \quad (10)$$

### 7.11 Rule species\_10

Rule species\_10 is an assignment rule for species species\_10:

$$\text{species\_10} = (\text{parameter\_13} + 1) \cdot (\text{parameter\_14} - [\text{species\_3}]) - \text{parameter\_13} \cdot [\text{species\_6}] \quad (11)$$

### 7.12 Rule species\_11

Rule species\_11 is an assignment rule for species species\_11:

$$\text{species\_11} = (\text{parameter\_13} + 1) \cdot [\text{species\_1}] - \text{parameter\_13} \cdot [\text{species\_13}] \quad (12)$$

### 7.13 Rule species\_12

Rule species\_12 is an assignment rule for species species\_12:

$$\text{species\_12} = (\text{parameter\_13} + 1) \cdot \text{parameter\_15} - \text{parameter\_13} \cdot [\text{species\_19}] \quad (13)$$

### 7.14 Rule species\_21

Rule species\_21 is an assignment rule for species species\_21:

$$\text{species\_21} = [\text{species\_12}] - [\text{species\_16}] \quad (14)$$

**Derived unit** dimensionless<sup>-1</sup>

### 7.15 Rule species\_14

Rule species\_14 is an assignment rule for species species\_14:

$$\text{species\_14} = (\text{parameter\_13} + 1) \cdot [\text{species\_3}] - \text{parameter\_13} \cdot [\text{species\_8}] \quad (15)$$

### 7.16 Rule species\_15

Rule species\_15 is an assignment rule for species species\_15:

$$\text{species\_15} = [\text{species\_14}] - 2 \cdot [\text{species\_11}] - [\text{species\_16}] \quad (16)$$

### 7.17 Rule parameter\_20

Rule parameter\_20 is an assignment rule for parameter parameter\_20:

$$\text{parameter\_20} = \frac{[\text{species\_10}] + [\text{species\_14}]}{\text{Metabolite\_9}} \quad (17)$$

### 7.18 Rule species\_1

Rule species\_1 is a rate rule for species species\_1:

$$\begin{aligned} \frac{d}{dt}\text{species\_1} = & \frac{1}{1 + \text{parameter\_13}} \cdot (\text{parameter\_9} \cdot (\text{parameter\_13} \cdot [\text{species\_20}]^2 + [\text{species\_15}]^2) \\ & - \text{parameter\_8} \cdot (\text{parameter\_13} \cdot [\text{species\_13}] + [\text{species\_11}])) \end{aligned} \quad (18)$$

### 7.19 Rule species\_2

Rule species\_2 is a rate rule for species species\_2:

$$\begin{aligned} \frac{d}{dt}\text{species\_2} = & \frac{1}{\text{parameter\_13} + 1} \cdot (\text{parameter\_9} \\ & \cdot (\text{parameter\_13} \cdot [\text{species\_18}] \cdot [\text{species\_20}] + [\text{species\_15}] \cdot [\text{species\_21}]) \\ & - \text{parameter\_8} \cdot (\text{parameter\_13} \cdot [\text{species\_17}] + [\text{species\_16}])) \end{aligned} \quad (19)$$

### 7.20 Rule species\_3

Rule species\_3 is a rate rule for species species\_3:

$$\begin{aligned} \frac{d}{dt}\text{species\_3} = & \frac{1}{1 + \text{parameter\_13}} \cdot \left( \text{parameter\_13} \cdot \text{parameter\_6} \cdot [\text{species\_26}] \right. \\ & \left. \cdot \frac{\text{parameter\_17}}{\text{parameter\_17} + [\text{species\_22}]} \cdot [\text{species\_6}] - \text{parameter\_7} \cdot [\text{species\_15}] \right) \end{aligned} \quad (20)$$

### 7.21 Rule species\_4

Rule species\_4 is a rate rule for species species\_4:

$$\begin{aligned} \frac{d}{dt}\text{species\_4} = & \text{parameter\_1} \cdot (\text{parameter\_27} + \text{parameter\_26} \cdot [\text{species\_16}] \\ & - (\text{parameter\_18} \cdot [\text{species\_9}] + \text{parameter\_25}) \cdot [\text{species\_4}]) \end{aligned} \quad (21)$$

### 7.22 Rule `species_5`

Rule `species_5` is a rate rule for species `species_5`:

$$\begin{aligned} \frac{d}{dt} \text{species\_5} = & \text{parameter\_1} \cdot ((1 - \text{parameter\_5}) \cdot (1 - \text{parameter\_16}) \\ & - (\text{parameter\_4} + (1 - \text{parameter\_28})) \cdot [\text{species\_5}]) \end{aligned} \quad (22)$$

### 7.23 Rule `species_6`

Rule `species_6` is a rate rule for species `species_6`:

$$\begin{aligned} \frac{d}{dt} \text{species\_6} = & \text{parameter\_2} \cdot [\text{species\_10}] - \left( \text{parameter\_3} + \text{parameter\_6} \cdot [\text{species\_26}] \right. \\ & \left. \cdot \frac{\text{parameter\_17}}{\text{parameter\_17} + [\text{species\_22}]} \right) \cdot [\text{species\_6}] \end{aligned} \quad (23)$$

### 7.24 Rule `species_7`

Rule `species_7` is a rate rule for species `species_7`:

$$\begin{aligned} \frac{d}{dt} \text{species\_7} = & \text{parameter\_1} \cdot (\text{parameter\_4} \cdot [\text{species\_5}] - (1 - \text{parameter\_28}) \\ & \cdot [\text{species\_25}] - \text{parameter\_18} \cdot [\text{species\_4}] \cdot [\text{species\_9}]) \end{aligned} \quad (24)$$

### 7.25 Rule `species_8`

Rule `species_8` is a rate rule for species `species_8`:

$$\begin{aligned} \frac{d}{dt} \text{species\_8} = & \text{parameter\_6} \cdot [\text{species\_26}] \cdot \frac{\text{parameter\_17}}{\text{parameter\_17} + [\text{species\_22}]} \\ & \cdot [\text{species\_6}] + \text{parameter\_2} \cdot [\text{species\_15}] - \text{parameter\_3} \\ & \cdot ([\text{species\_20}] + \text{parameter\_11} \cdot ([\text{species\_17}] + 2 \cdot [\text{species\_13}])) \end{aligned} \quad (25)$$

### 7.26 Rule `species_13`

Rule `species_13` is a rate rule for species `species_13`:

$$\begin{aligned} \frac{d}{dt} \text{species\_13} = & \text{parameter\_9} \cdot [\text{species\_20}]^2 \\ & - (\text{parameter\_8} + \text{parameter\_3} \cdot \text{parameter\_11}) \cdot [\text{species\_13}] \end{aligned} \quad (26)$$

### 7.27 Rule species\_17

Rule species\_17 is a rate rule for species species\_17:

$$\begin{aligned} \frac{d}{dt} \text{species\_17} = & \text{parameter\_9} \cdot [\text{species\_18}] \cdot [\text{species\_20}] \\ & - (\text{parameter\_8} + \text{parameter\_3} \cdot \text{parameter\_11}) \cdot [\text{species\_17}] \end{aligned} \quad (27)$$

### 7.28 Rule species\_19

Rule species\_19 is a rate rule for species species\_19:

$$\begin{aligned} \frac{d}{dt} \text{species\_19} = & \text{parameter\_19} \cdot [\text{species\_21}] - \text{parameter\_3} \\ & \cdot ([\text{species\_18}] + \text{parameter\_11} \cdot [\text{species\_17}]) \end{aligned} \quad (28)$$

### 7.29 Rule species\_24

Rule species\_24 is a rate rule for species species\_24:

$$\begin{aligned} \frac{d}{dt} \text{species\_24} = & \text{parameter\_1} \cdot (\text{parameter\_18} \cdot [\text{species\_4}] \cdot [\text{species\_9}] \\ & - (\text{parameter\_22} + \text{parameter\_12} \cdot (1 - \text{parameter\_28})) \cdot [\text{species\_24}]) \end{aligned} \quad (29)$$

### 7.30 Rule species\_26

Rule species\_26 is a rate rule for species species\_26:

$$\begin{aligned} \frac{d}{dt} \text{species\_26} = & \text{parameter\_1} \cdot (\text{parameter\_22} \cdot [\text{species\_24}] - \text{parameter\_12} \\ & \cdot (1 - \text{parameter\_28}) \cdot [\text{species\_26}]) \end{aligned} \quad (30)$$

## 8 Derived Rate Equations

When interpreted as an ordinary differential equation framework, this model implies the following set of equations for the rates of change of each species.

### 8.1 Species species\_1

**Name** S22

**SBO:0000297** protein complex

**Notes** total cellular homomeric S22 complexes

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_1](#)

One rule determines the species' quantity.

## 8.2 Species [species\\_2](#)

**Name** S24

**SBO:0000297** protein complex

**Notes** total cellular heteromeric S24 complexes

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_2](#)

One rule determines the species' quantity.

## 8.3 Species [species\\_3](#)

**Name** pS2tot

**Notes** total cellular pS2

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_3](#)

One rule determines the species' quantity.

## 8.4 Species [species\\_4](#)

**Name** TGF

**Notes** TGFbeta

**Initial concentration** 4 dimensionless · dimensionless<sup>-1</sup>

**Initial assignment** [species\\_4](#)

**Involved in rule** [species\\_4](#)

One rule determines the species' quantity.

### 8.5 Species `species_5`

**Name** R

**Notes** nascent receptors

**Initial concentration** 0.87962962962963 dimensionless · dimensionless<sup>-1</sup>

**Initial assignment** `species_5`

**Involved in rule** `species_5`

One rule determines the species' quantity.

### 8.6 Species `species_6`

**Name** S2c

**Notes** cytoplasmic, unphosphorylated Smad2

**Initial concentration** 1.19430241051863 dimensionless · dimensionless<sup>-1</sup>

**Initial assignment** `species_6`

**Involved in rule** `species_6`

One rule determines the species' quantity.

### 8.7 Species `species_7`

**Name** Rcom

**Notes** TGFb bound receptors

**Initial concentration** 0.12037037037037 dimensionless · dimensionless<sup>-1</sup>

**Initial assignment** `species_7`

**Involved in rule** `species_7`

One rule determines the species' quantity.

### 8.8 Species `species_8`

**Name** pS2c

**Notes** Total cytoplasmic pS2

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** `species_8`

One rule determines the species' quantity.

### 8.9 Species [species\\_9](#)

**Name** Rcom\_S

**Notes** mature, competent receptors

**Initial concentration** 0.05 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_9](#)

One rule determines the species' quantity.

### 8.10 Species [species\\_10](#)

**Name** S2n

**Notes** nuclear unphosphorylated Smad2

**Initial concentration** 0.558933528122717 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_10](#)

One rule determines the species' quantity.

### 8.11 Species [species\\_11](#)

**Name** S22n

**SBO:0000297** protein complex

**Notes** nuclear homomeric S22 complexes

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_11](#)

One rule determines the species' quantity.

### 8.12 Species [species\\_12](#)

**Name** S4n

**Notes** total nuclear Smad4

**Initial concentration** 1 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_12](#)

One rule determines the species' quantity.

### 8.13 Species [species\\_13](#)

**Name** S22c

**SBO:0000297** protein complex

**Notes** cytoplasmic homomeric S22 complexes

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_13](#)

One rule determines the species' quantity.

### 8.14 Species [species\\_14](#)

**Name** pS2n

**Notes** total nuclear pS2

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_14](#)

One rule determines the species' quantity.

### 8.15 Species [species\\_15](#)

**Name** pS2fn

**Notes** monomeric nuclear pS2

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_15](#)

One rule determines the species' quantity.

### 8.16 Species [species\\_16](#)

**Name** S24n

**SBO:0000297** protein complex

**Notes** nuclear heteromeric S24 complexes

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_16](#)

One rule determines the species' quantity.



### 8.17 Species [species\\_17](#)

**Name** S24c

**SBO:0000297** protein complex

**Notes** cytoplasmic heteromeric S24 complexes

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_17](#)

One rule determines the species' quantity.

### 8.18 Species [species\\_18](#)

**Name** S4fc

**Notes** monomeric cytoplasmic Smad4

**Initial concentration** 1 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_18](#)

One rule determines the species' quantity.

### 8.19 Species [species\\_19](#)

**Name** S4c

**Notes** total cytoplasmic Smad4

**Initial concentration** 1 dimensionless · dimensionless<sup>-1</sup>

**Initial assignment** [species\\_19](#)

**Involved in rule** [species\\_19](#)

One rule determines the species' quantity.

### 8.20 Species [species\\_20](#)

**Name** pS2fc

**Notes** monomeric cytoplasmic pS2

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_20](#)

One rule determines the species' quantity.

### 8.21 Species [species\\_21](#)

**Name** S4fn

**Notes** monomeric nuclear Smad4

**Initial concentration** 1 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_21](#)

One rule determines the species' quantity.

### 8.22 Species [species\\_22](#)

**Name** SBI

**SBO:0000390** boolean switch

**Notes** Receptor inhibitor, either present or absent (1, 0)

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

This species does not take part in any reactions. Its quantity does hence not change over time:

$$\frac{d}{dt}\text{species\_22} = 0 \quad (31)$$

### 8.23 Species [species\\_23](#)

**Name** Rtot

**Notes** total receptors

**Initial concentration** 1 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_23](#)

One rule determines the species' quantity.

### 8.24 Species [species\\_24](#)

**Name** RT

**Notes** active receptors

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** [species\\_24](#)

One rule determines the species' quantity.

## 8.25 Species `species_25`

**Name** Rcom\_I

**Initial concentration** 0.0703703703703704 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** `species_25`

One rule determines the species' quantity.

## 8.26 Species `species_26`

**Name** Ract

**Initial concentration** 0 dimensionless · dimensionless<sup>-1</sup>

**Involved in rule** `species_26`

One rule determines the species' quantity.

# A Glossary of Systems Biology Ontology Terms

**SBO:0000009 kinetic constant:** Numerical parameter that quantifies the velocity of a chemical reaction

**SBO:0000281 equilibrium constant:** Quantity characterizing a chemical equilibrium in a chemical reaction, which is a useful tool to determine the concentration of various reactants or products in a system where chemical equilibrium occurs

**SBO:0000282 dissociation constant:** Equilibrium constant that measures the propensity of a larger object to separate (dissociate) reversibly into smaller components, as when a complex falls apart into its component molecules, or when a salt splits up into its component ions. The dissociation constant is usually denoted  $K_d$  and is the inverse of the affinity constant.

**SBO:0000297 protein complex:** Macromolecular complex containing one or more polypeptide chains possibly associated with simple chemicals. CHEBI:3608

**SBO:0000337 association constant:** Equilibrium constant that measures the propensity of two objects to assemble (associate) reversibly into a larger component. The association constant is usually denoted  $K_a$  and is the inverse of the dissociation constant.

**SBO:0000360 quantity of an entity pool:** The enumeration of co-localised, identical biochemical entities of a specific state, which constitute a pool. The form of enumeration may be purely numerical, or may be given in relation to another dimension such as length or volume

**SBO:0000361 amount of an entity pool:** A numerical measure of the quantity, or of some property, of the entities that constitute the entity pool.

**SBO:0000380 biochemical coefficient:** number used as a multiplicative or exponential factor for quantities, expressions or function

**SBO:0000390 boolean switch:** A parameter that has precisely two discrete values which may be switched between. Usually for the boolean parameter these are indicated as '0 or 1' or 'True or False'

**SBO:0000470 mass fraction:** For a given substance, A, its mass fraction ( $x_A$ ) is defined as the ratio of its mass ( $m_A$ ) to the total mass ( $m_{\text{total}}$ ) in which it is present, where the sum of all mass fractions is equal to 1. This provides a means to express concentration in a dimensionless size.

**SBO:0000540 fraction of an entity pool:** A ratio that represents the quantity of a defined constituent entity over the total number of all constituent entities present.

SBML<sup>2</sup>TeX was developed by Andreas Dräger<sup>a</sup>, Hannes Planatscher<sup>a</sup>, Dieudonné M Wouamba<sup>a</sup>, Adrian Schröder<sup>a</sup>, Michael Hucka<sup>b</sup>, Lukas Endler<sup>c</sup>, Martin Golebiewski<sup>d</sup> and Andreas Zell<sup>a</sup>. Please see <http://www.ra.cs.uni-tuebingen.de/software/SBML2LaTeX> for more information.

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