

## SBML Model Report

# Model name: “Phillips2003 - The Mechanism of Ras GTPase Activation by Neurofibromin”



May 17, 2018

## 1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by Matthew Grant Roberts<sup>1</sup> at March 19<sup>th</sup> 2018 at 3:37 p. m. and last time modified at March 19<sup>th</sup> 2018 at 4:08 p. m. Table 1 gives 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	8
events	0	constraints	0
reactions	5	function definitions	5
global parameters	2	unit definitions	1
rules	2	initial assignments	0

## Model Notes

Phillips2003 - The Mechanism of Ras GTPaseActivation by NeurofibrominA mathematical model for Ras-GTPactivation by neurofibromin and the kinetic rates of the relevantreactions.

This model is described in the article:[The mechanism of Ras GTPase activation by neurofibromin](#).Phillips RA, Hunter JL, Eccleston JF, Webb MR.Biochemistry 2003 Apr; 42(13): 3956-3965

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#### Abstract:

Individual rate constants have been determined for each step of the Ras.GTP hydrolysis mechanism, activated by neurofibromin. Fluorescence intensity and anisotropy stopped-flow measurements used the fluorescent GTP analogue, mantGTP (2'(3')-O-(N-methylanthraniloyl)GTP), to determine rate constants for binding and release of neurofibromin. Quenched flow measurements provided the kinetics of the hydrolytic cleavage step. The fluorescent phosphate sensor, MDCC-PBP was used to measure phosphate release kinetics. Phosphate-water oxygen exchange, using (18)O-substituted GTP and inorganic phosphate (P(i)), was used to determine the extent of reversal of the hydrolysis step and of P(i) binding. The data show that neurofibromin and P(i) dissociate from the NF1.Ras.GDP.P(i) complex with identical kinetics, which are 3-fold slower than the preceding cleavage step. A model is presented in which the P(i) release is associated with the change of Ras from „GTP,, to „GDP,, conformation. In this model, the conformation change on P(i) release causes the large change in affinity of neurofibromin, which then dissociates rapidly.

This model is hosted on [BioModels Database](#) and identified by: [BIOMD0000000692](#).

To cite BioModels Database, please use: [Chelliah V et al. BioModels: ten-year anniversary. Nucl. Acids Res. 2015, 43\(Database issue\):D542-8.](#)

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## 2 Unit Definitions

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

### 2.1 Unit `substance`

**Name** substance

**Definition**  $\mu\text{mol}$

### 2.2 Unit `volume`

**Notes** Litre is the predefined SBML unit for `volume`.

**Definition** 1

### 2.3 Unit `area`

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

**Definition**  $\text{m}^2$

## 2.4 Unit length

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

**Definition** m

## 2.5 Unit time

**Notes** Second is the predefined SBML unit for time.

**Definition** s

# 3 Compartment

This model contains one compartment.

Table 2: Properties of all compartments.

Id	Name	SBO	Spatial Dimensions	Size	Unit	Constant	Outside
geometry	E. coli		3	$5.236 \cdot 10^{-13}$	l	<input checked="" type="checkbox"/>	

## 3.1 Compartment geometry

This is a three dimensional compartment with a constant size of  $5.236 \cdot 10^{-13}$  litre.

**Name** E. coli

## 4 Species

This model contains eight species. Section 9 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
Pi	Pi	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGTP	RasGTP	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGTP_minus_NF1	RasGTP_minus_NF1	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGTP_minus_NF1-_star_	RasGTP_minus_NF1_star_	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGDP_minus_NF1-_Pi	RasGDP_minus_NF1_Pi	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGDP_NF1	RasGDP_NF1	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
RasGDP	RasGDP	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>
NF1	NF1	geometry	$\mu\text{mol} \cdot \text{l}^{-1}$	<input type="checkbox"/>	<input type="checkbox"/>

## 5 Parameters

This model contains two global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
Pi_curve	Pi_curve		0.0		<input type="checkbox"/>
hplc_curve	hplc_curve		100.0		<input type="checkbox"/>

## 6 Function definitions

This is an overview of five function definitions.

### 6.1 Function definition [Function\\_for\\_GTP\\_hydrolysis\\_sbo\\_2\\_sbc](#)

**Name** Function for GTP\_hydrolysis\_sbo\_2\_sbc\_

**Arguments** [Pi], [RasGDP\_NF1], [RasGDP\_minus\_NF1\_Pi], vol(geometry), kb, kf

**Mathematical Expression**

$$\frac{kf \cdot [\text{RasGDP\_minus\_NF1\_Pi}] - kb \cdot [\text{Pi}] \cdot [\text{RasGDP\_NF1}]}{\text{vol}(\text{geometry})} \quad (1)$$

### 6.2 Function definition [Function\\_for\\_NF1\\_binding](#)

**Name** Function for NF1\_binding

**Arguments** [NF1], [RasGTP], [RasGTP\_minus\_NF1], vol(geometry), kb, kf

**Mathematical Expression**

$$\frac{kf \cdot [\text{RasGTP}] \cdot [\text{NF1}] - kb \cdot [\text{RasGTP\_minus\_NF1}]}{\text{vol}(\text{geometry})} \quad (2)$$

### 6.3 Function definition [Function\\_for\\_Ras\\_activation](#)

**Name** Function for Ras\_activation

**Arguments** [RasGTP\_minus\_NF1], [RasGTP\_minus\_NF1\_star\_], vol(geometry), kb, kf

**Mathematical Expression**

$$\frac{kf \cdot [\text{RasGTP\_minus\_NF1}] - kb \cdot [\text{RasGTP\_minus\_NF1\_star\_}]}{\text{vol}(\text{geometry})} \quad (3)$$

#### 6.4 Function definition `Function_for_GAP_dissociation`

**Name** Function for GAP\_dissociation

**Arguments** `[NF1]`, `[RasGDP]`, `[RasGDP_NF1]`, `vol(geometry)`, `kb`, `kf`

**Mathematical Expression**

$$\frac{kf \cdot [RasGDP\_NF1] - kb \cdot [RasGDP] \cdot [NF1]}{vol(geometry)} \quad (4)$$

#### 6.5 Function definition `Function_for_GTP_hydrolysis_sbo_1_sbc`

**Name** Function for GTP\_hydrolysis\_sbo\_1\_sbc

**Arguments** `[RasGDP_minus_NF1_Pi]`, `[RasGTP_minus_NF1_star_]`, `vol(geometry)`, `kb`, `kf`

**Mathematical Expression**

$$\frac{kf \cdot [RasGTP\_minus\_NF1\_star\_] - kb \cdot [RasGDP\_minus\_NF1\_Pi]}{vol(geometry)} \quad (5)$$

### 7 Rules

This is an overview of two rules.

#### 7.1 Rule `Pi_curve`

Rule `Pi_curve` is an assignment rule for parameter `Pi_curve`:

$$Pi\_curve = \frac{100 \cdot ([Pi] - 1)}{1} \quad (6)$$

#### 7.2 Rule `hplc_curve`

Rule `hplc_curve` is an assignment rule for parameter `hplc_curve`:

$$hplc\_curve = \frac{[RasGTP] \cdot 100}{1} \quad (7)$$

## 8 Reactions

This model contains five reactions. All reactions are listed in the following table and are subsequently described in detail. If a reaction is affected by a modifier, the identifier of this species is written above the reaction arrow.

Table 5: Overview of all reactions

Nº	Id	Name	Reaction Equation	SBO
1	GTP_hydrolysis- _sbo_1_sbc_	GTP_hydrolysis_sbo_1_sbc_	$\text{RasGTP\_minus\_NF1\_star\_} \rightleftharpoons \text{RasGDP\_minus\_NF1\_Pi}$	
2	GTP_hydrolysis- _sbo_2_sbc_	GTP_hydrolysis_sbo_2_sbc_	$\text{RasGDP\_minus\_NF1\_Pi} \rightleftharpoons \text{Pi} + \text{RasGDP\_NF1}$	
3	Ras_activation	Ras_activation	$\text{RasGTP\_minus\_NF1} \rightleftharpoons \text{RasGTP\_minus\_NF1\_star\_}$	
4	GAP- _dissociation	GAP_dissociation	$\text{RasGDP\_NF1} \rightleftharpoons \text{RasGDP} + \text{NF1}$	
5	NF1_binding	NF1_binding	$\text{RasGTP} + \text{NF1} \rightleftharpoons \text{RasGTP\_minus\_NF1}$	

## 8.1 Reaction GTP\_hydrolysis\_sbo\_1\_sbc\_

This is a reversible reaction of one reactant forming one product.

**Name** GTP\_hydrolysis\_sbo\_1\_sbc\_

**Notes** First step in hydrolysis of GTP bound to Ras complexed with NF1 - a mammalian GAP  
Kf = 19.5 /sec Kb = 0.22 /sec Table 3, Phillips RA et al 2003 Biochemistry 3956-3965

### Reaction equation



### Reactant

Table 6: Properties of each reactant.

Id	Name	SBO
RasGTP_minus_NF1_star_	RasGTP_minus_NF1_star_	

### Product

Table 7: Properties of each product.

Id	Name	SBO
RasGDP_minus_NF1_Pi	RasGDP_minus_NF1_Pi	

### Kinetic Law

**Derived unit** contains undeclared units

$$v_1 = \text{vol}(\text{geometry}) \cdot \text{Function\_for\_GTP\_hydrolysis\_sbo\_1\_sbc}([\text{RasGDP\_minus\_NF1\_Pi}], [\text{RasGTP\_minus\_NF1\_star\_}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) \quad (9)$$

$$\begin{aligned} &\text{Function\_for\_GTP\_hydrolysis\_sbo\_1\_sbc}([\text{RasGDP\_minus\_NF1\_Pi}], \\ &[\text{RasGTP\_minus\_NF1\_star\_}], \text{vol}(\text{geometry}), \text{kb}, \\ &\text{kf}) = \frac{\text{kf} \cdot [\text{RasGTP\_minus\_NF1\_star\_}] - \text{kb} \cdot [\text{RasGDP\_minus\_NF1\_Pi}]}{\text{vol}(\text{geometry})} \end{aligned} \quad (10)$$



$$\begin{aligned} &\text{Function\_for\_GTP\_hydrolysis\_sbo\_1\_sbc} ([\text{RasGDP\_minus\_NF1\_Pi}], \\ &[\text{RasGTP\_minus\_NF1\_star\_}], \text{vol}(\text{geometry}), \text{kb}, \\ &\text{kf}) = \frac{\text{kf} \cdot [\text{RasGTP\_minus\_NF1\_star\_}] - \text{kb} \cdot [\text{RasGDP\_minus\_NF1\_Pi}]}{\text{vol}(\text{geometry})} \end{aligned} \quad (11)$$

Table 8: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kb	kb		$1.15192 \cdot 10^{-13}$		<input checked="" type="checkbox"/>
kf	kf		$1.02102 \cdot 10^{-11}$		<input checked="" type="checkbox"/>

## 8.2 Reaction `GTP_hydrolysis_sbo_2_sbc_`

This is a reversible reaction of one reactant forming two products.

**Name** `GTP_hydrolysis_sbo_2_sbc_`

**Notes** Second step in hydrolysis of GTP bound to Ras is complexed with NF1 - a mammalian GAP  
 $K_f = 40 \text{ /sec}$   $K_b = 108 \text{ /M/sec} = 1.08\text{e-}04 \text{ /uM/sec}$  Phillips RA et al 2003 Biochemistry 3956-3965

### Reaction equation



### Reactant

Table 9: Properties of each reactant.

Id	Name	SBO
<code>RasGDP_minus_NF1_Pi</code>	<code>RasGDP_minus_NF1_Pi</code>	

### Products

Table 10: Properties of each product.

Id	Name	SBO
<code>Pi</code>	<code>Pi</code>	
<code>RasGDP_NF1</code>	<code>RasGDP_NF1</code>	

Id	Name	SBO
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### Kinetic Law

**Derived unit** contains undeclared units

$$v_2 = \text{vol}(\text{geometry}) \cdot \text{Function\_for\_GTP\_hydrolysis\_sbo\_2\_sbc}([\text{Pi}], [\text{RasGDP\_NF1}], [\text{RasGDP\_minus\_NF1\_Pi}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) \quad (13)$$

$$\text{Function\_for\_GTP\_hydrolysis\_sbo\_2\_sbc}([\text{Pi}], [\text{RasGDP\_NF1}], [\text{RasGDP\_minus\_NF1\_Pi}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) = \frac{\text{kf} \cdot [\text{RasGDP\_minus\_NF1\_Pi}] - \text{kb} \cdot [\text{Pi}] \cdot [\text{RasGDP\_NF1}]}{\text{vol}(\text{geometry})} \quad (14)$$

$$\text{Function\_for\_GTP\_hydrolysis\_sbo\_2\_sbc}([\text{Pi}], [\text{RasGDP\_NF1}], [\text{RasGDP\_minus\_NF1\_Pi}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) = \frac{\text{kf} \cdot [\text{RasGDP\_minus\_NF1\_Pi}] - \text{kb} \cdot [\text{Pi}] \cdot [\text{RasGDP\_NF1}]}{\text{vol}(\text{geometry})} \quad (15)$$

Table 11: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kb	kb		$5.65482 \cdot 10^{-17}$		<input checked="" type="checkbox"/>
kf	kf		$2.0944 \cdot 10^{-11}$		<input checked="" type="checkbox"/>

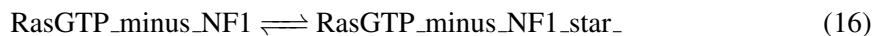
### 8.3 Reaction Ras\_activation

This is a reversible reaction of one reactant forming one product.

**Name** Ras\_activation

**Notes** Activation of Ras by GAP (i.e NF1) Kf = 418 /sec Kb = 5.5 /sec Table 3, Phillips RA et al 2003 Biochemistry 3956-3965

#### Reaction equation



#### Reactant

Table 12: Properties of each reactant.

Id	Name	SBO
RasGTP_minus_NF1	RasGTP_minus_NF1	

## Product

Table 13: Properties of each product.

Id	Name	SBO
RasGTP_minus_NF1_star_	RasGTP_minus_NF1_star_	

## Kinetic Law

**Derived unit** contains undeclared units

$$v_3 = \text{vol}(\text{geometry}) \cdot \text{Function\_for\_Ras\_activation}([\text{RasGTP\_minus\_NF1}], [\text{RasGTP\_minus\_NF1\_star\_}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) \quad (17)$$

$$\text{Function\_for\_Ras\_activation}([\text{RasGTP\_minus\_NF1}], [\text{RasGTP\_minus\_NF1\_star\_}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) = \frac{\text{kf} \cdot [\text{RasGTP\_minus\_NF1}] - \text{kb} \cdot [\text{RasGTP\_minus\_NF1\_star\_}]}{\text{vol}(\text{geometry})} \quad (18)$$

$$\text{Function\_for\_Ras\_activation}([\text{RasGTP\_minus\_NF1}], [\text{RasGTP\_minus\_NF1\_star\_}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) = \frac{\text{kf} \cdot [\text{RasGTP\_minus\_NF1}] - \text{kb} \cdot [\text{RasGTP\_minus\_NF1\_star\_}]}{\text{vol}(\text{geometry})} \quad (19)$$

Table 14: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kb	kb		$2.8798 \cdot 10^{-12}$		<input checked="" type="checkbox"/>
kf	kf		$2.18865 \cdot 10^{-10}$		<input checked="" type="checkbox"/>

## 8.4 Reaction GAP\_dissociation

This is a reversible reaction of one reactant forming two products.

**Name** GAP\_dissociation

**Notes** Dissociation of NF1 from Ras.GDP NF1 is a mammalian GAP Kf = 46.5 /sec Kb = 1.2 /sec/uM Table 3, Phillips RA et al 2003 Biochemistry 3956-3965

### Reaction equation



### Reactant

Table 15: Properties of each reactant.

Id	Name	SBO
RasGDP_NF1	RasGDP_NF1	

### Products

Table 16: Properties of each product.

Id	Name	SBO
RasGDP NF1	RasGDP NF1	

### Kinetic Law

**Derived unit** contains undeclared units

$$v_4 = \text{vol}(\text{geometry}) \cdot \text{Function\_for\_GAP\_dissociation}([NF1], [\text{RasGDP}], [\text{RasGDP\_NF1}], \text{vol}(\text{geometry}), kb, kf) \quad (21)$$

$$\text{Function\_for\_GAP\_dissociation}([NF1], [\text{RasGDP}], [\text{RasGDP\_NF1}], \text{vol}(\text{geometry}), kb, kf) = \frac{kf \cdot [\text{RasGDP\_NF1}] - kb \cdot [\text{RasGDP}] \cdot [NF1]}{\text{vol}(\text{geometry})} \quad (22)$$

$$\text{Function\_for\_GAP\_dissociation}([NF1], [\text{RasGDP}], [\text{RasGDP\_NF1}], \text{vol}(\text{geometry}), kb, kf) = \frac{kf \cdot [\text{RasGDP\_NF1}] - kb \cdot [\text{RasGDP}] \cdot [NF1]}{\text{vol}(\text{geometry})} \quad (23)$$

Table 17: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kb	kb		$6.28318 \cdot 10^{-13}$		<input checked="" type="checkbox"/>
kf	kf		$2.43474 \cdot 10^{-11}$		<input checked="" type="checkbox"/>

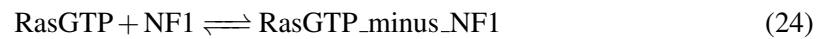
## 8.5 Reaction NF1\_binding

This is a reversible reaction of two reactants forming one product.

**Name** NF1\_binding

**Notes** Binding of NF1 to Ras.GTP NF1 is a mammalian GAP Kd = 5.3 uM Table 3, Phillips RA et al 2003 Biochemistry 3956-3965

### Reaction equation



### Reactants

Table 18: Properties of each reactant.

Id	Name	SBO
RasGTP	RasGTP	
NF1	NF1	

### Product

Table 19: Properties of each product.

Id	Name	SBO
RasGTP_minus_NF1	RasGTP_minus_NF1	

### Kinetic Law

**Derived unit** contains undeclared units

$$v_5 = \text{vol}(\text{geometry}) \cdot \text{Function\_for\_NF1\_binding}([\text{NF1}], [\text{RasGTP}], [\text{RasGTP\_minus\_NF1}], \text{vol}(\text{geometry}), \text{kb}, \text{kf}) \quad (25)$$

$$\text{Function\_for\_NF1\_binding}([NF1], [RasGTP], [RasGTP\_minus\_NF1], \text{vol}(\text{geometry}), kb, kf) = \frac{kf \cdot [RasGTP] \cdot [NF1] - kb \cdot [RasGTP\_minus\_NF1]}{\text{vol}(\text{geometry})} \quad (26)$$

$$\text{Function\_for\_NF1\_binding}([NF1], [RasGTP], [RasGTP\_minus\_NF1], \text{vol}(\text{geometry}), kb, kf) = \frac{kf \cdot [RasGTP] \cdot [NF1] - kb \cdot [RasGTP\_minus\_NF1]}{\text{vol}(\text{geometry})} \quad (27)$$

Table 20: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
kb	kb		$3.3301 \cdot 10^{-12}$		<input checked="" type="checkbox"/>
kf	kf		$6.28318 \cdot 10^{-13}$		<input checked="" type="checkbox"/>

## 9 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.

Identifiers for kinetic laws highlighted in gray cannot be verified to evaluate to units of SBML substance per time. As a result, some SBML interpreters may not be able to verify the consistency of the units on quantities in the model. Please check if

- parameters without an unit definition are involved or
- volume correction is necessary because the `hasOnlySubstanceUnits` flag may be set to `false` and `spacialDimensions` > 0 for certain species.

### 9.1 Species Pi

**Name** Pi

**Initial concentration**  $1 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in one reaction (as a product in [GTP\\_hydrolysis\\_sbo\\_2\\_sbc\\_](#)).

$$\frac{d}{dt} \text{Pi} = v_2 \quad (28)$$

### 9.2 Species RasGTP

**Name** RasGTP

**Initial concentration**  $1 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in one reaction (as a reactant in [NF1\\_binding](#)).

$$\frac{d}{dt}\text{RasGTP} = -v_5 \quad (29)$$

### 9.3 Species [RasGTP\\_minus\\_NF1](#)

**Name** [RasGTP\\_minus\\_NF1](#)

**Initial concentration**  $0 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in [Ras\\_activation](#) and as a product in [NF1\\_binding](#)).

$$\frac{d}{dt}\text{RasGTP\_minus\_NF1} = v_5 - v_3 \quad (30)$$

### 9.4 Species [RasGTP\\_minus\\_NF1\\_star\\_](#)

**Name** [RasGTP\\_minus\\_NF1\\_star\\_](#)

**Initial concentration**  $0 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in [GTP\\_hydrolysis\\_sbo\\_1\\_sbc\\_](#) and as a product in [Ras\\_activation](#)).

$$\frac{d}{dt}\text{RasGTP\_minus\_NF1\_star\_} = v_3 - v_1 \quad (31)$$

### 9.5 Species [RasGDP\\_minus\\_NF1\\_Pi](#)

**Name** [RasGDP\\_minus\\_NF1\\_Pi](#)

**Initial concentration**  $0 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in [GTP\\_hydrolysis\\_sbo\\_2\\_sbc\\_](#) and as a product in [GTP\\_hydrolysis\\_sbo\\_1\\_sbc\\_](#)).

$$\frac{d}{dt}\text{RasGDP\_minus\_NF1\_Pi} = v_1 - v_2 \quad (32)$$

### 9.6 Species [RasGDP\\_NF1](#)

**Name** [RasGDP\\_NF1](#)

**Initial concentration**  $0 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in [GAP\\_dissociation](#) and as a product in [GTP\\_hydrolysis\\_sbo\\_2\\_sbc\\_](#)).

$$\frac{d}{dt}\text{RasGDP\_NF1} = v_2 - v_4 \quad (33)$$

## 9.7 Species RasGDP

**Name** RasGDP

**Initial concentration**  $0 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in one reaction (as a product in [GAP\\_dissociation](#)).

$$\frac{d}{dt}\text{RasGDP} = v_4 \quad (34)$$

## 9.8 Species NF1

**Name** NF1

**Initial concentration**  $9.99936338171632 \mu\text{mol} \cdot \text{l}^{-1}$

This species takes part in two reactions (as a reactant in [NF1\\_binding](#) and as a product in [GAP\\_dissociation](#)).

$$\frac{d}{dt}\text{NF1} = v_4 - v_5 \quad (35)$$

SBML2<sup>LaTeX</sup> 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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