

## SBML Model Report

### Model name: “Brannmark2010- \_InsulinSignalling\_Mifamodel”



May 6, 2016

## 1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by Ishan Ajmera<sup>1</sup> at July 14<sup>th</sup> 2011 at 2:54 p. m. and last time modified at April eighth 2016 at 5:01 p. 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 | 19       | unit definitions     | 0        |
| rules             | 25       | initial assignments  | 0        |

## Model Notes

This model is from the article:

**Mass and information feedbacks through receptor endocytosis govern insulin signaling as revealed using a parameter-free modeling framework.**

Brannmark C, Palmer R, Glad ST, Cedersund G, Stralfors P. J Biol Chem.2010 Jun 25;285(26):20171-9. [20421297](#),

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**Abstract:**

Insulin and other hormones control target cells through a network of signal-mediating molecules. Such networks are extremely complex due to multiple feedback loops in combination with redundancy, shared signal mediators, and cross-talk between signal pathways. We present a novel framework that integrates experimental work and mathematical modeling to quantitatively characterize the role and relation between co-existing submechanisms in complex signaling networks. The approach is independent of knowing or uniquely estimating model parameters because it only relies on (i) rejections and (ii) core predictions (uniquely identified properties in unidentifiable models). The power of our approach is demonstrated through numerous iterations between experiments, model-based data analyses, and theoretical predictions to characterize the relative role of co-existing feedbacks governing insulin signaling. We examined phosphorylation of the insulin receptor and insulin receptor substrate-1 and endocytosis of the receptor in response to various different experimental perturbations in primary human adipocytes. The analysis revealed that receptor endocytosis is necessary for two identified feedback mechanisms involving mass and information transfer, respectively. Experimental findings indicate that interfering with the feedback may substantially increase overall signaling strength, suggesting novel therapeutic targets for insulin resistance and type 2 diabetes. Because the central observations are present in other signaling networks, our results may indicate a general mechanism in hormonal control.

## 2 Unit Definitions

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

### 2.1 Unit `substance`

**Notes** Mole is the predefined SBML unit for substance.

**Definition** mol

### 2.2 Unit `volume`

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

**Definition** l

### 2.3 Unit `area`

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

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

## 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<br>Dimensions | Size | Unit  | Constant                            | Outside |
|---------------------|---------------|-----|-----------------------|------|-------|-------------------------------------|---------|
| compartment_0000001 | compartemnt 1 |     | 3                     | 1    | litre | <input checked="" type="checkbox"/> |         |

## 3.1 Compartment [compartment\\_0000001](#)

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

**Name** compartemnt 1

## 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 7 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                  |
|-------|-------|---------------------|----------------------------------|--------------------------|-------------------------------------|
| IR    | IR    | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| IRins | IRins | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| IRp   | IRp   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| IRip  | IRip  | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| IRi   | IRi   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| IRS   | IRS   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| IRSip | IRSip | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| X     | X     | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| Xp    | Xp    | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1a   | V1a   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1b   | V1b   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1c   | V1c   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1d   | V1d   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1e   | V1e   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1g   | V1g   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V1r   | v1r   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V2    | V2    | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| Vm2   | Vm2   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| V3    | V3    | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| Vm3   | Vm3   | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| simXP | simXp | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |

| Id             | Name           | Compartment         | Derived Unit                     | Constant                 | Boundary<br>Condi-<br>tion          |
|----------------|----------------|---------------------|----------------------------------|--------------------------|-------------------------------------|
| intamount      | intamount      | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| measIRp        | measIRp        | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input type="checkbox"/>            |
| measdoublestep | measdoublestep | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| measanna       | measanna       | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |
| measdosR       | measdosR       | compartment_0000001 | $\text{mol} \cdot \text{l}^{-1}$ | <input type="checkbox"/> | <input checked="" type="checkbox"/> |

## 5 Parameters

This model contains 19 global parameters.

Table 4: Properties of each parameter.

| Id       | Name     | SBO     | Value                   | Unit | Constant |
|----------|----------|---------|-------------------------|------|----------|
| k1a      | k1a      | 0000009 | 0.389                   |      | ✓        |
| k1abasic | k1abasic | 0000009 | 0.012                   |      | ✓        |
| k1b      | k1b      | 0000009 | 0.020                   |      | ✓        |
| k1e      | k1e      | 0000009 | $4.38334 \cdot 10^{-5}$ |      | ✓        |
| k1f      | k1f      | 0000009 | 20.073                  |      | ✓        |
| k1g      | k1g      | 0000009 | 286.699                 |      | ✓        |
| k1r      | k1r      | 0000009 | 3.633                   |      | ✓        |
| k21      | k21      | 0000009 | 1.672                   |      | ✓        |
| k22      | k22      | 0000009 | 0.036                   |      | ✓        |
| km2      | km2      | 0000009 | 32.594                  |      | ✓        |
| k3       | k3       | 0000009 | 1.629                   |      | ✓        |
| km3      | km3      | 0000009 | 0.113                   |      | ✓        |
| ky1      | ky1      | 0000009 | 152.963                 |      | ✓        |
| ky2      | ky2      | 0000009 | 8936.220                |      | ✓        |
| kyanna   | kyanna   | 0000009 | 16760.120               |      | ✓        |
| kyDosR   | kyDosR   | 0000009 | 13740.432               |      | ✓        |
| ins      | ins      | 0000196 | 100.000                 |      | ✓        |
| k1c      | k1c      | 0000009 | 0.364                   |      | ✓        |
| k1d      | k1d      | 0000009 | 1580.678                |      | ✓        |

## 6 Rules

This is an overview of 25 rules.

### 6.1 Rule measanna

Rule measanna is an assignment rule for species measanna:

$$\text{measanna} = \text{kyanna} \cdot [\text{IRSip}] \quad (1)$$

### 6.2 Rule measdosR

Rule measdosR is an assignment rule for species measdosR:

$$\text{measdosR} = \text{kyDosR} \cdot [\text{IRSip}] \quad (2)$$

### 6.3 Rule `measdoubstep`

Rule `measdoubstep` is an assignment rule for species `measdoubstep`:

$$\text{measdoubstep} = k_{y2} \cdot [\text{IRSip}] \quad (3)$$

### 6.4 Rule `V1a`

Rule `V1a` is an assignment rule for species `V1a`:

$$\text{V1a} = k_{1a} \cdot \text{ins} \cdot [\text{IR}] + k_{1a\text{basic}} \cdot [\text{IR}] \quad (4)$$

### 6.5 Rule `V1b`

Rule `V1b` is an assignment rule for species `V1b`:

$$\text{V1b} = k_{1b} \cdot [\text{IRins}] \quad (5)$$

### 6.6 Rule `V1c`

Rule `V1c` is an assignment rule for species `V1c`:

$$\text{V1c} = k_{1c} \cdot [\text{IRins}] \quad (6)$$

### 6.7 Rule `V1d`

Rule `V1d` is an assignment rule for species `V1d`:

$$\text{V1d} = k_{1d} \cdot [\text{IRp}] \quad (7)$$

### 6.8 Rule `V1e`

Rule `V1e` is an assignment rule for species `V1e`:

$$\text{V1e} = [\text{IRip}] \cdot \left( k_{1e} + \frac{k_{1f} \cdot [\text{Xp}]}{1 + [\text{Xp}]} \right) \quad (8)$$

### 6.9 Rule `V1g`

Rule `V1g` is an assignment rule for species `V1g`:

$$\text{V1g} = k_{1g} \cdot [\text{IRp}] \quad (9)$$

### 6.10 Rule `V1r`

Rule `V1r` is an assignment rule for species `V1r`:

$$\text{V1r} = k_{1r} \cdot [\text{IRi}] \quad (10)$$

### 6.11 Rule V2

Rule V2 is an assignment rule for species V2:

$$V2 = k21 \cdot ([IRp] + k22 \cdot [IRip]) \cdot [IRS] \quad (11)$$

### 6.12 Rule Vm2

Rule Vm2 is an assignment rule for species Vm2:

$$Vm2 = km2 \cdot [IRSip] \quad (12)$$

### 6.13 Rule V3

Rule V3 is an assignment rule for species V3:

$$V3 = k3 \cdot [X] \cdot [IRSip] \quad (13)$$

### 6.14 Rule Vm3

Rule Vm3 is an assignment rule for species Vm3:

$$Vm3 = km3 \cdot [Xp] \quad (14)$$

### 6.15 Rule simXP

Rule simXP is an assignment rule for species simXP:

$$\text{simXP} = [Xp] \quad (15)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

### 6.16 Rule intamount

Rule intamount is an assignment rule for species intamount:

$$\text{intamount} = \frac{[IRi] + [IRip]}{10} \quad (16)$$

### 6.17 Rule IR

Rule IR is a rate rule for species IR:

$$\frac{d}{dt}IR = [V1a] + [V1b] + [V1r] + [V1g] \quad (17)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$



### 6.18 Rule IRins

Rule IRins is a rate rule for species IRins:

$$\frac{d}{dt}\text{IRins} = [\text{V1a}] - [\text{V1b}] - [\text{V1c}] \quad (18)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

### 6.19 Rule IRp

Rule IRp is a rate rule for species IRp:

$$\frac{d}{dt}\text{IRp} = [\text{V1c}] - [\text{V1d}] - [\text{V1g}] \quad (19)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

### 6.20 Rule IRip

Rule IRip is a rate rule for species IRip:

$$\frac{d}{dt}\text{IRip} = [\text{V1d}] - [\text{V1e}] \quad (20)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

### 6.21 Rule IRi

Rule IRi is a rate rule for species IRi:

$$\frac{d}{dt}\text{IRi} = [\text{V1e}] - [\text{V1r}] \quad (21)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

### 6.22 Rule IRS

Rule IRS is a rate rule for species IRS:

$$\frac{d}{dt}\text{IRS} = [\text{V2}] + [\text{Vm2}] \quad (22)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

### 6.23 Rule IRSip

Rule IRSip is a rate rule for species IRSip:

$$\frac{d}{dt}\text{IRSip} = [\text{V2}] - [\text{Vm2}] \quad (23)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

## 6.24 Rule X

Rule X is a rate rule for species X:

$$\frac{d}{dt}X = [V3] + [Vm3] \quad (24)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

## 6.25 Rule Xp

Rule Xp is a rate rule for species Xp:

$$\frac{d}{dt}Xp = [V3] - [Vm3] \quad (25)$$

**Derived unit**  $\text{mol} \cdot \text{l}^{-1}$

# 7 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.

## 7.1 Species IR

**Name** IR

**SBO:0000244** receptor

**Initial amount** 10 mol

**Involved in rule** IR

One rule determines the species' quantity.

## 7.2 Species IRins

**Name** IRins

**SBO:0000240** material entity

**Initial amount** 0 mol

**Involved in rule** IRins

One rule determines the species' quantity.

### 7.3 Species IRp

**Name** IRp

**SBO:0000240** material entity

**Initial amount** 0 mol

**Involved in rule** IRp

One rule determines the species' quantity.

### 7.4 Species IRip

**Name** IRip

**SBO:0000240** material entity

**Initial amount** 0 mol

**Involved in rule** IRip

One rule determines the species' quantity.

### 7.5 Species IRi

**Name** IRi

**SBO:0000240** material entity

**Initial amount** 0 mol

**Involved in rule** IRi

One rule determines the species' quantity.

### 7.6 Species IRS

**Name** IRS

**SBO:0000240** material entity

**Initial concentration** 10 mol · l<sup>-1</sup>

**Involved in rule** IRS

One rule determines the species' quantity.

### 7.7 Species [IRSiP](#)

**Name** [IRSiP](#)

**SBO:0000240** material entity

**Initial amount** 0 mol

**Involved in rule** [IRSiP](#)

One rule determines the species' quantity.

### 7.8 Species [X](#)

**Name** [X](#)

**SBO:0000240** material entity

**Initial amount** 10 mol

**Involved in rule** [X](#)

One rule determines the species' quantity.

### 7.9 Species [Xp](#)

**Name** [Xp](#)

**SBO:0000240** material entity

**Initial amount** 0 mol

**Involved in rule** [Xp](#)

One rule determines the species' quantity.

### 7.10 Species [V1a](#)

**Name** [V1a](#)

**Initial concentration** 389.41271264 mol · l<sup>-1</sup>

**Involved in rule** [V1a](#)

One rule determines the species' quantity.

### 7.11 Species V1b

**Name** V1b

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

**Involved in rule** V1b

One rule determines the species' quantity.

### 7.12 Species V1c

**Name** V1c

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

**Involved in rule** V1c

One rule determines the species' quantity.

### 7.13 Species V1d

**Name** V1d

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

**Involved in rule** V1d

One rule determines the species' quantity.

### 7.14 Species V1e

**Name** V1e

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

**Involved in rule** V1e

One rule determines the species' quantity.

### 7.15 Species V1g

**Name** V1g

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

**Involved in rule** V1g

One rule determines the species' quantity.

### 7.16 Species $V_{1r}$

**Name**  $v_{1r}$

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

**Involved in rule**  $V_{1r}$

One rule determines the species' quantity.

### 7.17 Species $V_2$

**Name**  $V_2$

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

**Involved in rule**  $V_2$

One rule determines the species' quantity.

### 7.18 Species $V_{m2}$

**Name**  $V_{m2}$

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

**Involved in rule**  $V_{m2}$

One rule determines the species' quantity.

### 7.19 Species $V_3$

**Name**  $V_3$

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

**Involved in rule**  $V_3$

One rule determines the species' quantity.

### 7.20 Species $V_{m3}$

**Name**  $V_{m3}$

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

**Involved in rule**  $V_{m3}$

One rule determines the species' quantity.

### 7.21 Species `simXP`

**Name** `simXp`

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

**Involved in rule** `simXP`

One rule determines the species' quantity.

### 7.22 Species `intamount`

**Name** `intamount`

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

**Involved in rule** `intamount`

One rule determines the species' quantity.

### 7.23 Species `measIRp`

**Name** `measIRp`

**SBO:0000196** concentration of an entity pool

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

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

$$\frac{d}{dt} \text{measIRp} = 0 \quad (26)$$

### 7.24 Species `measdoubstep`

**Name** `measdoubstep`

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

**Involved in rule** `measdoubstep`

One rule determines the species' quantity.

### 7.25 Species `measanna`

**Name** `measanna`

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

**Involved in rule** `measanna`

One rule determines the species' quantity.

## 7.26 Species `measdosR`

**Name** `measdosR`

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

**Involved in rule** `measdosR`

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:0000196 concentration of an entity pool:** The amount of an entity per unit of volume.

**SBO:0000240 material entity:** A real thing that is defined by its physico-chemical structure.

**SBO:0000244 receptor:** Participating entity that binds to a specific physical entity and initiates the response to that physical entity. The original concept of the receptor was introduced independently at the end of the 19th century by John Newport Langley (1852-1925) and Paul Ehrlich (1854-1915). Langley JN. On the reaction of cells and of nerve-endings to certain poisons, chiefly as regards the reaction of striated muscle to nicotine and to curari. J Physiol. 1905 Dec 30;33(4-5):374-413

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