

## SBML Model Report

# Model name: “Baker2013 - Cytokine Mediated Inflammation in Rheumatoid Arthritis”



May 6, 2016

## 1 General Overview

This is a document in SBML Level 2 Version 4 format. This model was created by Vincent Knight-Schrijver<sup>1</sup> at September 25<sup>th</sup> 2014 at 1:48 p.m. and last time modified at December twelveth 2014 at 3:01 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	2
events	0	constraints	0
reactions	0	function definitions	0
global parameters	5	unit definitions	3
rules	2	initial assignments	0

## Model Notes

Baker2013 - Cytokine Mediated Inflammation in Rheumatoid Arthritis This model by Baker M. 2013, describes the interaction between pro and anti-inflammatory cytokine signalling in rheumatoid arthritis.

---

<sup>1</sup>EMBL-EBI, [vknight@ebi.ac.uk](mailto:vknight@ebi.ac.uk)

Using two ordinary differential equations, the first model [\[BIOMD0000000550\]](#) analyses bifurcation and describes different pathological states by altering inflammatory regulation parameters. The second model [\[BIOMD0000000549\]](#) includes the effect that ageing has on pro-inflammatory signalling, allowing for time-dependant properties and disease progression to be observed. The author also describes potential dosing for reversal of the disease state.

This model is described in the article: [Mathematical modelling of cytokine-mediated inflammation in rheumatoid arthritis](#). Baker M, Denman-Johnson S, Brook BS, Gaywood I, Owen MR. Math Med Biol 2013 Dec; 30(4): 311-337

Abstract:

Rheumatoid arthritis (RA) is a chronic inflammatory disease preferentially affecting the joints and leading, if untreated, to progressive joint damage and disability. Cytokines, a group of small inducible proteins, which act as intercellular messengers, are key regulators of the inflammation that characterizes RA. They can be classified into pro-inflammatory and anti-inflammatory groups. Numerous cytokines have been implicated in the regulation of RA with complex up and down regulatory interactions. This paper considers a two-variable model for the interactions between pro-inflammatory and anti-inflammatory cytokines, and demonstrates that mathematical modelling may be used to investigate the involvement of cytokines in the disease process. The model displays a range of possible behaviours, such as bistability and oscillations, which are strongly reminiscent of the behaviour of RA e.g. genetic susceptibility and remitting-relapsing disease. We also show that the dose regimen as well as the dose level are important factors in RA treatments.

This model is hosted on [BioModels Database](#) and identified by: [BIOMD0000000550](#).

To cite BioModels Database, please use: [BioModels Database: An enhanced, curated and annotated resource for published quantitative kinetic models](#).

To the extent possible under law, all copyright and related or neighbouring rights to this encoded model have been dedicated to the public domain worldwide. Please refer to [CC0 Public Domain Dedication](#) for more information.

## 2 Unit Definitions

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

### 2.1 Unit volume

**Name** volume

**Definition** dimensionless

### 2.2 Unit time

**Name** time

**Definition** dimensionless

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

### 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	Synovium		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** Synovium

**Notes** Compartment notes: `{\textquotestraightdblbase}`The synovium is modelled as a spatial

## 4 Species

This model contains two species. The boundary condition of two 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
species_1	a	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	<input checked="" type="checkbox"/>
species_2	p	compartment_1	dimensionless dimensionless <sup>-1</sup>	· ⊞	<input checked="" type="checkbox"/>

## 5 Parameters

This model contains five global parameters.

Table 4: Properties of each parameter.

Id	Name	SBO	Value	Unit	Constant
parameter_1	alpha1		0.025		✓
parameter_2	alpha2		1.000		✓
parameter_3	alpha3		0.500		✓
parameter_4	alpha4		3.500		✓
parameter_5	gamma		1.250		✓

## 6 Rules

This is an overview of two rules.

### 6.1 Rule `species_1`

Rule `species_1` is a rate rule for species `species_1`:

$$\frac{d}{dt}\text{species}_1 = [\text{species}_1] + \text{parameter}_4 \cdot \frac{[\text{species}_2]^2}{\text{parameter}_3^2 + [\text{species}_2]^2} \quad (1)$$

### 6.2 Rule `species_2`

Rule `species_2` is a rate rule for species `species_2`:

$$\begin{aligned} \frac{d}{dt}\text{species}_2 = & \text{parameter}_5 \cdot [\text{species}_2] + \frac{1}{1 + [\text{species}_1]^2} \\ & \cdot \left( \text{parameter}_1 + \text{parameter}_2 \cdot \frac{[\text{species}_2]^2}{1 + [\text{species}_2]^2} \right) \end{aligned} \quad (2)$$

## 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 `species_1`

**Name** a

**Notes** Anti-inflammatory cytokine concentration (dimensionless)

**Initial concentration** 0.00577667577789773 dimensionless · dimensionless<sup>-1</sup>

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

One rule determines the species' quantity.

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

**Name** p

**Notes** Pro-inflammatory cytokine concentration (dimensionless)

**Initial concentration** 0.0203298264712407 dimensionless · dimensionless<sup>-1</sup>

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

One rule determines the species' quantity.

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.

<sup>a</sup>Center for Bioinformatics Tübingen (ZBIT), Germany

<sup>b</sup>California Institute of Technology, Beckman Institute BNMC, Pasadena, United States

<sup>c</sup>European Bioinformatics Institute, Wellcome Trust Genome Campus, Hinxton, United Kingdom

<sup>d</sup>EML Research gGmbH, Heidelberg, Germany