

## Independent Section

Contains tests that are independent of the class of modeled organism, a model's

### Consistency

Stoichiometric Consistency

Mass Balance

Charge Balance

Metabolite Connectivity

Unbounded Flux In Default Medium

Sub Total

### Annotation - Metabolites

Presence of Metabolite Annotation

Metabolite Annotations Per Database

pubchem.compound

kegg.compound

seed.compound

inchikey

inchi

chebi

hmdb

reactome

metanetx.chemical

bigg.metabolite

biocyc

Metabolite Annotation Conformity Per Database

pubchem.compound

kegg.compound

seed.compound

inchikey

inchi

chebi

hmdb

## Specific Section









Covers general statistics and specific aspects of a metabolic network that are not

### SBML





SBML Level and Version **Errored** 

FBC enabled **Errored** 






### Basic Information

Model Identifier	ChemoModelGrowth	
Total Metabolites	1,114	
Total Reactions	1,149	
Total Genes	535	
Total Compartments	3	
Metabolic Coverage	2.15	
Uncoserved Metabolites	0	
Minimal Inconsistent Net Stoichiometries	Skipped	

### Metabolite Information

Unique Metabolites	818	
Duplicate Metabolites in Identical Compartments	0	
Metabolites without Charge	0	
Metabolites without Formula	0	
Medium Components	22	

### Reaction Information

Purely Metabolic Reactions	760	
Purely Metabolic Reactions with Constraints	2	
Transport Reactions	275	
Transport Reactions with Constraints	0	
Reactions With Partially		

bigg.metabolite

biocyc

Uniform Metabolite Identifier Namespace

Sub Total

## Annotation - Reactions

Presence of Reaction Annotation

Reaction Annotations Per Database

rhea

kegg.reaction

seed.reaction

metanetx.reaction

bigg.reaction

reactome

ec-code

brenda

biocyc

Reaction Annotation Conformity Per Databases

rhea

kegg.reaction

seed.reaction

metanetx.reaction

bigg.reaction

reactome

ec-code

brenda

biocyc

Uniform Reaction Identifier Namespace

Sub Total

## Annotation - Genes

Presence of Gene Annotation

Gene Annotations Per Database

rhea

Reactions with Identical  
Genes

0.59



## Gene-Protein-Reaction (GPR) Associations

Reactions without GPR

69

Fraction of Transport  
Reactions without GPR

0.15



Enzyme Complexes

131



## Biomass

Biomass Reactions  
Identified

1



Biomass Consistency

0.95

Biomass Production In  
Default Medium

0.09

Unrealistic Growth Rate  
In Default Medium

false

Biomass Production In  
Complete Medium

57.25

Blocked Biomass  
Precursors In Default  
Medium

0

Blocked Biomass  
Precursors In Complete  
Medium

0

Ratio of Direct  
Metabolites in Biomass  
Reaction

0.11

Number of Missing  
Essential Biomass  
Precursors

1



## Energy Metabolism

Non-Growth Associated  
Maintenance Reaction

1

Growth-associated  
Maintenance in Biomass  
Reaction

true

Number of Reversible  
Oxygen-Containing  
Reactions

3

Erroneous Energy-  
generating Cycles

Info



MNXM3

Skipped



MNXM63

Skipped



ecogene

MNXM423

Skipped



kegg.genes

MNXM6

Skipped



ncbigi

MNXM10

Skipped



ncbigene

MNXM38

Skipped



ncbiprotein

MNXM208

Skipped



ccds

MNXM191

Skipped



hprd

MNXM223

Skipped



asap

MNXM7517

Skipped

**Gene Annotation Conformity Per Database**

MNXM12233

Skipped



refseq

MNXM558

Skipped



uniprot

MNXM21

Skipped



ecogene

MNXM89557

Skipped



kegg.genes

**Network Topology**

ncbigi

Universally Blocked  
Reactions

481



ncbigene

Orphan Metabolites

168



ncbiprotein

Dead-end Metabolites

157



ccds

Stoichiometrically  
Balanced Cycles

41



hprd

Metabolite Production In  
Complete Medium

502



asap

Metabolite Consumption  
In Complete Medium

518



Sub Total

**Annotation - SBO Terms**

Metabolite General SBO Presence

Metabolite SBO:0000247 Presence

Reaction General SBO Presence

Metabolic Reaction SBO:0000176 Presence

Transport Reaction SBO:0000185 Presence

Exchange Reaction SBO:0000627 Presence

Demand Reaction SBO:0000628 Presence

Sink Reactions SBO:0000632 Presence

Gene General SBO Presence

Gene SBO:0000243 Presence

Biomass Reactions SBO:0000629 Presence

**Matrix Conditioning**Ratio Min/Max Non-Zero  
Coefficients

0.00

Independent  
Conservation Relations

128



Rank

986



Degrees Of Freedom

163

**Experimental Data  
Comparison**

Growth Prediction

Skipped

Gene Essentiality  
Prediction

Skipped



Sub Total

# 51%

Score per Category



## Environment

Python Version	3.8.12
Platform	Linux
Memote Version	0.13.0