November 27, 2014

‘iMK1208.xml’:

* Original model causes an error, and FBA cannot be performed.

‘iMK1208Edited’:

* First, cobra\_model was generated from iMK1208.xml using “create\_cobra\_model\_from\_sbml\_file”.
* Second, cobra\_model was written into SBML using “write\_cobra\_model\_to\_sbml\_file”.
* Finally, rewritten SBML was loaded and worked fine.

May 10, 2015

‘iMK1208Edited2’:

* Initial version had an error for Heme A. No metabolite ID was given to it. Hence, its ID was: “M\_\_C”
* This metabolite appears in: HEMEAS, Biomass\_SCO
* This error was corrected: “hemeA” given for “heme A”

July 7, 2015

“iMK1208Edited3”:

* This is the version without secondary metabolite biosynthetic reactions.
* This version was prepared by HKS.

October 25, 2015

“iMK1208Edited4”:

* Pathway names for “Subsystem” were modified to be consistent.
* This version was prepared by HKS.
* Also see:

D:\Research\Progress\antiSMASH\_development\module\_modeling\Source\HKS\151022\_template\_model\_pathway\_name\_updated