Transcription 2nd day

Overview

→ Defined i/o for ¾ modules in new format

→ Defined parameters that are exchanged

→ Improved our modules ...

Transcription module

created python script to build up loci of each polymerase

 to improve performance we will work on a 2nd version (more abstract)

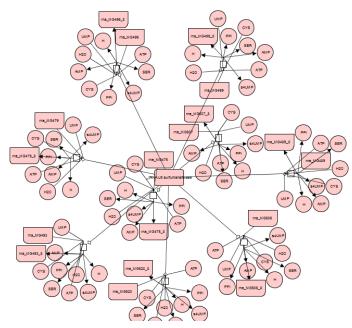
Transcriptional regulation

- SBML model with all reactions

- Problem: reactions happen with poisson distr. (original is multinomial)

RNA modification

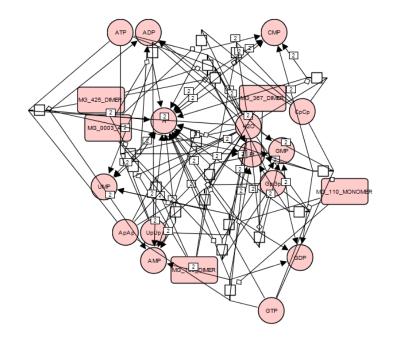
- built SBML/SBGN



RNA processing

- started to encode the Model

- current status,
 - details have to be added



RNA decay

mark down the RNA decay reactions to its products (nucleotides)

coding in SBML will be finished tomorrow

Thats it =)