

# Dynamic Flux Balance Analysis (DFBA) with SBML (core, comp, fbc)

Matthias König, Leandro Watanabe & Chris Meyers

LiSyM Systems Medicine of the Liver Humboldt University Berlin, Institute for Theoretical Biology livermetabolism.com



# Dynamic FBA (DFBA)

# Coupling of dynamical model parts to steady state networks (FBA)

- Subset of general problem of coupling models with different simulation frameworks
  - Logical/boolean models
  - Stochastic simulations

### **Approaches**

- Resource allocation (creating large optimization problem based)
  - Very large optimization problems
  - Dynamic achieved via optimization boundary conditions like biomass(24h) = k \*biomass(0h) in combination with resource variables
  - Coupling to arbitrary ODEs not possible

### Coupling of ODE to FBA model

- **Direct embedding** of LP-Solver in ODE solver
  - execution in every time step
  - complex implementation (high level expertise on ODE solver design)
  - Stiffness of system at borders of flux-cone
- Stationary Optimization Approach (SOA)
- Dynamic Optimization Approach (DOA)



## Examples

### Whole-cell model (Karr et al)

**Table I.** Overview of existing DFBA simulation studies (sorted by date).

Refs.	FBA model	Met.	Fluxes	Method/solver
Varma and Palsson (1994)	Based on Majewski and Domach (1990)	24	34	SOA/—
Mahadevan et al. (2002)	Based on Schilling et al. (2000)	3	4	SOA/CPLEX
				DOA/fmincon
Sainz et al. (2003)	_	43	38	SOA/—
Luo et al. (2006) (MDFBA)	_	7	8	DOA/fmincon
Hjersted and Henson (2006, 2009)	iGH99	98	82	DA/CONOPT
Pizarro et al. (2007)	Based on Sainz et al. (2003)	38	39	SOA/—
Hjersted et al. (2007)	iND750	1,059	1,265	DA/MOSEK
Anesiadis et al. (2008)	iJR904	625	931	SOA/CPLEX
Lee et al. (2008) (idFBA)	_	_	13	SOA/—
Luo et al. (2009) (MDFBA)	_	8	5	DOA/fmincon
Oddone et al. (2009)	IL1403	422	621	SOA/Mathematica
Lequeux et al. (2010) (MDFA)	<del>_</del>	24	34	polynomial fitting
Salimi et al. (2010)	iFS2007	679	712	DA/—
	iFS431	603	621	
Zhuang et al. (2011)	G. sulfurreducens	541	522	DA/LINDO
	R. ferrireducens	790	762	
Meadows et al. (2010)	Based on Varma and Palsson (1994)	30	123	ODE15S/linprog
Vargas et al. (2011)	idFV715(iFF708)	590	1,181	SOA/LINDO
Nolan and Lee (2011) (MDFA)	_	150	136	SOA/—
Hanly and Henson (2011)	iRJ904	625	931	DA/MOSEK
Hanly et al. (2012)	iND750	1,059	1,265	



## Applications & Use cases

#### Circadian Liver metabolism

 Coupling circadian blood metabolite & gene/protein expression patterns to FBA model of liver metabolism (HepatoNet1)

#### Whole-body PKPD

- Coupling of tissue specific FBA models to whole body PKPD
- Glucose regulation
- (liver, muscle, fat, pancreas, stomach)
- Cori cycle & other multi-tissue physiological cycles
  - Glucose alanine/pyruvate shuttle

### Model embedding

 Coupling ODE pathway models to genome-scale metabolic models (liver)



### **SBML**

# De facto standard for Systems Biology Models (ODE, FBA, mixed compartments)

- Good description for kinetic and FBA models, but no implementation of DFBA in standard
- Core language (core) & extension packages
  - core

#### **Kinetic models**

- Compartments, Parameters, Species, Reactions, RateRules, AssignmentRules, Events, FunctionDefinitions)
- fbc

#### **FBA** encoding

- objective functions, upper & lower bounds, GPR encoding
- · comp

#### **Coupling of models**

 ExternalModelDefinitions, ModelDefinitions replacements, replacedBy, deletions, ports, submodels,



# Encoding

https://github.com/matthiaskoenig/dfba https://github.com/matthiaskoenig/dfba/blob/ master/DFBA%20models%20in%20SBML.md

#### **TOP**

- kinetic part & coupling of submodelsBOUNDS
- kinetic bounds calculation

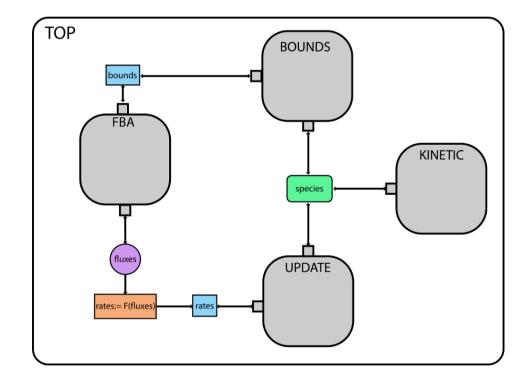
#### **UPDATE**

kinetic update of species from FBA
 FBA

FBA problem (interface via exchange reactions)

#### **Interface**

- exchange reactions (boundaryCondition=True species)
  - What is taken up, what is imported, how unbalanced are metabolites?
- update kinetics
  - How does flux effect metabolites (scaling by biomass, ...)
- bound kinetics
  - How do kinetic players effect bounds
    & resource limitation of FBA





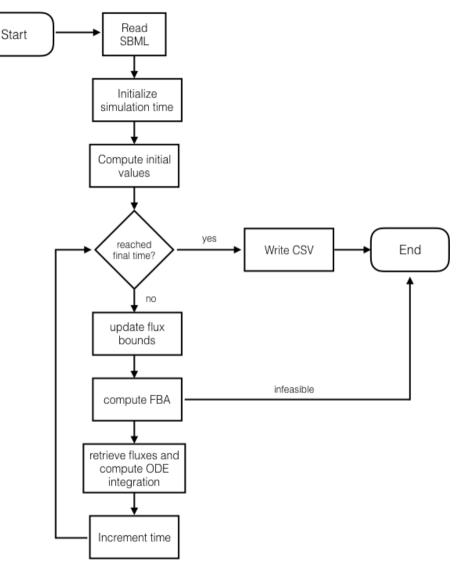
# Simulation Algorithm

#### Implementations

- sbmlutils
  - https://github.com/matthiaskoenig/sbmliils
  - iBioSim
  - http://www.async.ece.utah.edu/ibiosim

#### Challenges

- standardized encoding of DFBA models
- uniqueness of FBA solutions
- bounds limitations based on species amounts/concentrations
- the (in)famous hack
- if (c<0): c=0





## Example Models

### Toy model

3 reactions, linear chain, 2 exchange reactions

### **Diauxic growth**

4 effective reactions, 4 exchange reactions

#### E.coli core

- 95 reactions, 72 metabolites, 137 genes, 4 exchange reactions)
  - http://bigg.ucsd.edu/models/e\_coli\_core
  - https://escher.github.io/builder/index.h tml?enable\_editing=true&map\_name=e\_coli\_ core.Core%20metabolism

