

# Integrative Modeling of Mitochondrial Energy Metabolism

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## 1 Introduction

We are in the process of constructing a kinetic model of mitochondrial energy metabolism including the electron transport (respiratory chain), the TCA cycle, the fatty acid metabolism ( $\beta$  oxidation), and the inner-membrane metabolite carriers. In this work, we report a model of these metabolic pathways that has been recently completed using the E-CELL system [2], a generic simulation environment we have developed for cellular simulation.

## 2 Material and Method

### 2.1 Model

The model consists of the following enzymatic reactions: four (4) for the respiratory chain, twelve (12) along with some ancillary reactions for the TCA cycle, seven (7) for the  $\beta$  oxidation, and eight (8) for the metabolite carrier system. All of the enzymatic reactions are modeled based on rate equations and kinetic parameters found in the literature or estimated by genetic algorithm (GA) and various parameter estimating methods [1].

Table 1: The reactions in the model

Group	Enzyme	Group	Enzyme
respiratory chain	NADH dehydrogenase	$\beta$ oxidation	Acyl-CoA dehydrogenase
	Ubiquinol:Cytochrome <i>c</i> oxidoreductase		Enoyl-CoA hydratase
	Cytochrome <i>c</i> oxidase		3-hydroxyacyl-CoA dehydrogenase
	ATP synthase		3-ketoacyl-CoA thiolase
TCA cycle	Pyruvate dehydrogenase complex		ETF:Q oxidoreductase
	Malic enzyme		CPT I
	Pyruvate carboxylase		CPT II
	Citrate synthase	Metabolite carriers	Pyruvate carrier
	Aconitase		Dicarboxylate carrier
	Isocitrate dehydrogenase ( $\times 2$ )		Tricarboxylate carrier
	2-oxoglutarate dehydrogenase complex		Oxoglutarate carrier
	Succinyl-CoA synthetase		Adenine nucleotide carrier
	Succinate dehydrogenase		Phosphate carrier
	Fumarase		Carnitine carrier
Malate dehydrogenase	Asp/Glu carrier		

## 2.2 Parameter Estimation

Here, we show 2 examples of kinetic parameter estimation. Each graph shows fitness of experimental values in the literature and computer-simulated values.

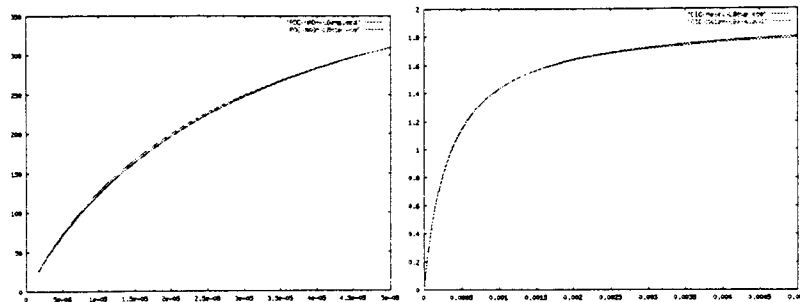


Figure 1. Pyruvate Dehydrogenase Complex (left), Tricarboxylate Carrier (right)  
vertical axis: turnover (1/sec), horizontal axis: metabolite concentration (M)

## 3 Future Works

We will continue to work on this mitochondrial model, with the eventual goal of applying it to pathological analyses of mitochondrial diseases such as Leigh syndrome and mitochondrial myopathy, encephalopathy lactic acidosis and stroke-like episodes (MELAS).

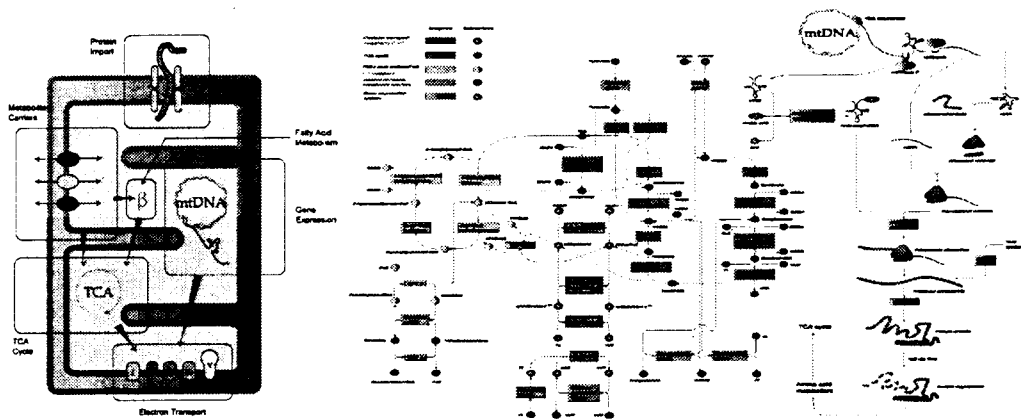


Figure 2: a) An overview of the model and b) the modeled reactions

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## References

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- [2] Tomita, M., Hashimoto, K., Takahashi, K., Shimizu, T., Matsuzaki, Y., Miyoshi, F., Saito, K., Tanida, S., Yugi, K., Venter, J.C. and Hutchison, C., E-CELL : Software environment for whole cell simulation, *Bioinformatics*, 15(1):72-84, 1999.