

Models and molecular mechanisms for trade-offs in the context of metabolism

Seirana Hashemi¹, Roosa Laitinen², and Zoran Nikoloski³

¹Universität Potsdam

²University of Helsinki

³Max-Planck-Institut für molekulare Pflanzenphysiologie

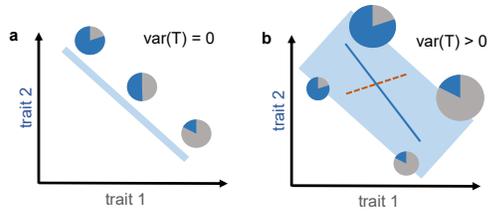
February 22, 2024

Abstract

Accumulating evidence for trade-offs involving metabolic traits has demonstrated their importance in evolution of organisms. Metabolic models with different level of complexity have already been considered when investigating mechanisms that explain various metabolic trade-offs. Here we provide a systematic review of modelling approaches that have been used to study and explain trade-offs between: (i) kinetic properties of individual enzymes, (ii) rates of metabolic reactions, (iii) rate and yield of metabolic pathways and networks, (iv) different metabolic objectives in single organisms and in metabolic communities, and (v) metabolic concentrations. In providing insights into mechanisms underlying these five types of metabolic trade-offs obtained from constraint-based metabolic modelling, we emphasize the relation of metabolic trade-offs to the classical black box Y-model that provides conceptual explanation for resource acquisition-allocation trade-offs. In addition, we identify several pressing concerns and offer a perspective for future research in the identification and manipulation of metabolic trade-offs by relying on the toolbox provided by constraint-based metabolic modelling for single organisms and microbial communities.

Hosted file

trade_offs_models_final.docx available at <https://authorea.com/users/500570/articles/581317-models-and-molecular-mechanisms-for-trade-offs-in-the-context-of-metabolism>



c

a

$\max v_{bio}$
s.t.
 $\mathbf{Nv} = \mathbf{0}$ (steady state)
 $\mathbf{v}_{\min} \leq \mathbf{v} \leq \mathbf{v}_{\max}$ (flux capacity)
(other constraints)

$v_i \leq k_{cat}^{ij} E_j$ (enzyme constraints)
 $\sum_j MW_j E_j \leq E_{tot}$

translation constraints
considering ribosomes
transcription constraints
(growth constraints, $v_{bio} = \mu$)

