

Bidirectionality and compartmentation of metabolic fluxes are revealed in the dynamics of isotopomer networks
Schryer, David; Peterson, Pearu; Paalme, Toomas; Vendelin, Marko International journal of molecular sciences 2009 / 4, p. 1697-1718 <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2680642/>

Isotopomeric ¹³C labeling of amino acids reveal compartmentation in *Saccharomyces uvarum*
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Metabolic flux analysis of compartmentalized systems using dynamic isotopologue modeling = Isotopoloogilise modelleerimise rakendamine heterogeensete bioloogiliste süsteemide ainevahetusvoo analüüsis
Schryer, David 2012 https://www.eester.ee/record=b2776763*est

Sensitivity analysis of flux determination in heart by H(2) (18)O-provided labeling using a dynamic isotopologue model of energy transfer pathways
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