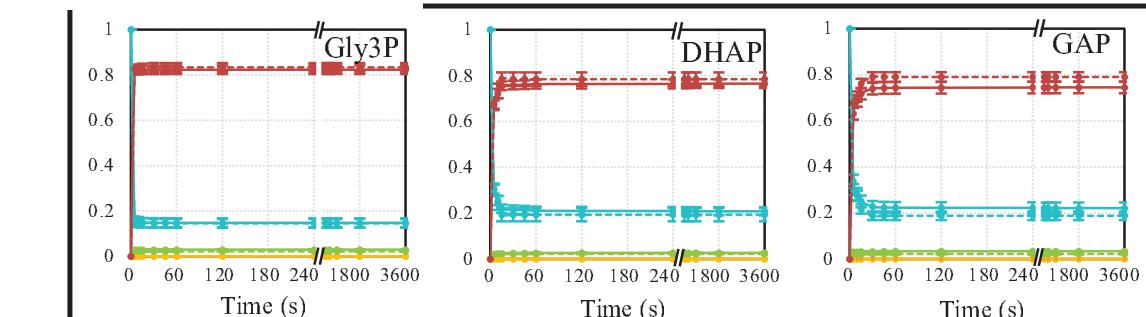
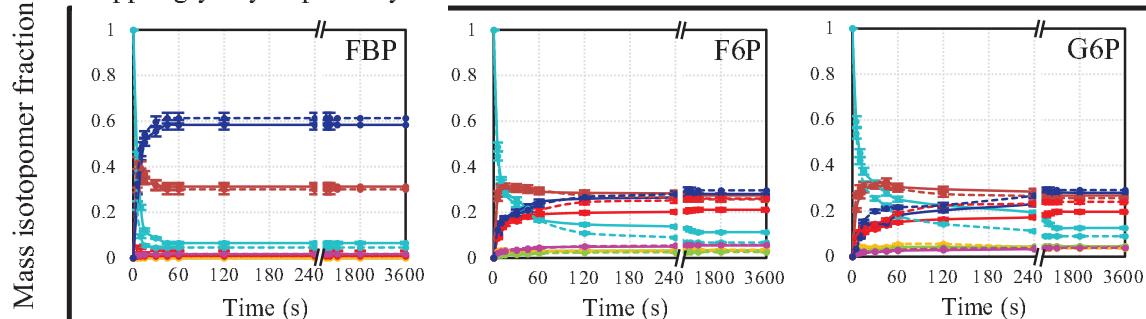


**Additional file 2.**  $^{13}\text{C}$ -labeling trajectories of selected intracellular metabolites after the introduction of [U- $^{13}\text{C}$ ]glycerol at the exponential phase. **a** Mass isotopomer abundances after correction for natural isotopic abundances. **b** Average  $^{13}\text{C}$ -enrichments calculated using the formula  $(\frac{1}{N}) \sum_{i=1}^N M_i \times i$ , where  $N$  is the number of carbon atoms in the metabolite and  $M_i$  is the fractional abundance of the  $i$ th mass isotopomer. The solid lines (HT66) and dashed lines (HT66LSP) illustrate the measured labeling trends.

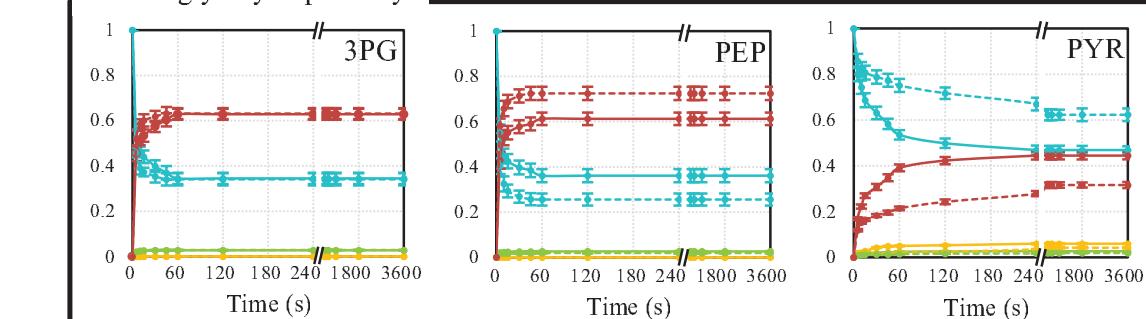
**a Glycerol uptake pathway**



**Upper glycolytic pathway**



**Lower glycolytic pathway**



● M+0	○ M+1	● M+2	● M+3	● M+4	● M+5	● M+6	● M+7	(HT66)
● M+0	○ M+1	● M+2	● M+3	● M+4	● M+5	● M+6	● M+7	(HT66LSP)

