Integrated 13C metabolic flux analysis of 14 parallel labeling experiments in Escherichia coli

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**Table S1.** Metabolic network model of *E. coli* used for 13C metabolic flux analysis.

**Table S2.** Biomass and glucose measurements for parallel labeling experiments with *E. coli*.

**Table S3.** Mass isotopomer distributions of biomass amino acids from parallel labeling experiments with *E. coli*.

**Table S4.** Results of 13C-metabolic flux analysis for *E. coli* grown in parallel batch cultures with different 13C-labeled glucose tracers.