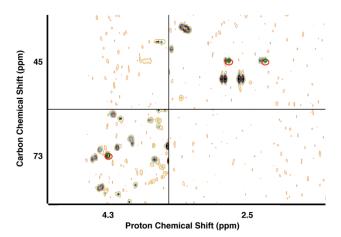
Supplementary Material

(A) Malate Annotated in Endpoint Extract by COLMARm

Peak	database 1H	database 13C	exp 1H	exp 13C
0	2.36	45.32	2.35	45.15
1	2.66	45.32	2.66	45.16
2	4.29	73.06	4.29	72.97

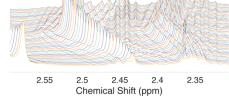


(B) Putative Glutamine Annotation in *in-vivo* data

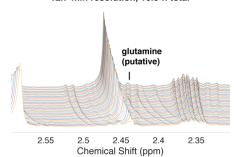
Sample 4 (aerobic)

12.7-min resolution, 10.8-h total

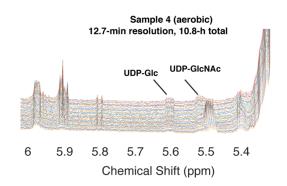
glutamine
(putative)

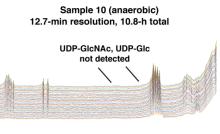


Sample 10 (anaerobic) 12.7-min resolution, 10.8-h total



(C) UDP-GlcNAc and UDP-Glc qualitative changes in *in-vivo* data





6 5.9 5.8 5.7 5.6 5.5 5.4 Chemical Shift (ppm) **Supplementary Figure 6.** Qualitative assessment for unquantifiable metabolites in this study. **(A)** Malate was annotated in the 2D data for endpoint extracts by peak matching to databases using COLMARm¹. Positions of matched peaks are indicated with a red ellipse and in the peak table. **(B)** Peaks consistent with glutamine increased in the aerobic and may be present in the anaerobic condition. **(C)** UDP-GlcNAc (UDP-N-Acetyl Glucosamine; UDP-NAG) and UDP-Glc (UDP-Glucose) were annotated in the aerobic condition but could not be traced reliably.