

Supplemental Figure 1: Labeling diagrams for stable isotope tracing experiments. A Schematics of predicted labeling from one turn of the traditional TCA cycle when cells are grown in culture media either (left) with glucose uniformly labeled with <sup>13</sup>C ([<sup>13</sup>C<sub>6</sub>] glucose) and unlabeled glutamine or (right) with glutamine uniformly labeled with <sup>13</sup>C ([<sup>13</sup>C<sub>5</sub>] glutamine) and unlabeled glucose. [<sup>13</sup>C<sub>6</sub>] glucose is converted to doubly labeled citrate,  $\alpha$ -KG, succinate, and 2-HG. [<sup>13</sup>C<sub>5</sub>] glutamine is converted to fully labeled  $\alpha$ -KG, succinate, and 2-HG and to 4- or 5-carbon labeled citrate. Citrate can also be converted to acetyl-CoA and oxaloacetate by ATP citrate lyase. Dashed arrows indicate that one or more intermediate reactions have been omitted from the diagram for clarity. **B** Model to explain the observed labeling data, in which citrate generated from intramitochondrial oxaloacetate. The oxaloacetate can be re-imported into the mitochondria as malate in exchange for an  $\alpha$ -KG generated from glutaminolysis. This cytosolic  $\alpha$ -KG can be used directly by mutant IDH1 to generate 2-HG in the cytosol.