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Supplemental Information

Consumption of NADPH for 2-HG Synthesis

Increases Pentose Phosphate Pathway Flux

and Sensitizes Cells to Oxidative Stress

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Figure S1. *IDH1* **mutants have increased PPP flux, Related to Figure 3**. (A-C) *IDH1* mutants have increased PPP flux when cultured in DMEM with 10% FBS and 5 mM glucose. (A) Isotopologue distribution of lactate in wild-type cells and HCT116 R132H/+ mutants. The M+1 isotopologue is a result of glucose carbon that passed through the PPP. The M+2 isotopologue corresponds to glucose carbon that was metabolized to lactate through glycolysis directly. (B) Relative PPP flux, as determined by lactate labeling from $1,2-{}^{13}C_2$ glucose and glucose uptake. (C) Uptake of glucose by wildtype HCT116 cells and *IDH1* mutants. (D-E) The flux of the oxidative phase of the PPP is increased in HCT116 *IDH1* mutants relative to wild-type cells. (D) Decay of the unlabeled fraction of glucose 6-phosphate (G6P) and the unlabeled fraction of 6-phosphogluconate (6PG) in wild-type and R132H/+ cells. (E) Oxidative PPP flux was determined by fitting G6P and 6PG data with Newtonian type minimization, using the measured 6PG concentrations of 2.9 pmol per μ L of cells and 3.5 pmol per μ L of cells for wild-type and R132H/+ cells, respectively. Data shown are mean values \pm s.d. (*n*=3). ** indicates a *p*-value < 0.01, and *** indicates a *p*-value < 0.001.



Figure S2. Evaluating the activity of malic enzyme (ME) with U- 13 C glutamine, Related to Experimental Procedures. (A) Schematic to show labeling when U- 13 C glutamine is transformed to lactate through ME. The ratios of labeled malate and lactate were analyzed. (B) Labeling of malate and lactate from U- 13 C glutamine. The data do not support increased ME activity in *IDH1* mutants relative to wild-type cells. It is important to note that this method does not differentiate between ME subtypes or compartmentalization of the NADPH produced. Data shown are mean values \pm s.d. (n=3).



Figure S3. Exposing cells to 2-HG does not increase PPP flux, Related to Figure 3. (A) LC/MS measurements show that the intracellular concentration of 2-HG is comparable between HCT116 R132H/+ cells and wild-type cells treated with 0.1 mM octyl 2-HG. Wild-type cells alone display very low levels of 2-HG. (B) PPP flux of HCT116 wild-type cells exposed to 0.1 mM octyl 2-HG for 72 hrs. Compared to control wild-type cells, no significant change in PPP flux is detected upon treatment with octyl 2-HG. R132H/+ cells, however, have increased PPP flux relative to both wild-type and wild-type + octyl 2-HG. Data shown are mean values \pm s.d. (*n*=3). *** indicates a *p*-value < 0.001.



Figure S4. Effects of 6-aminonicotinamide, an inhibitor of 6-phosphogluconate dehydrogenase, Related to Figure 3. (A) Intracellular levels of 2-HG decrease after treating HCT116 R132H/+ cells with 6-aminonicotinamide for 24 hr. (B) The intracellular levels of other central carbon metabolites, such as lactate shown here, also decrease upon treatment with 6-aminonicotinamide for 24 hr. Data shown are mean values \pm s.d. (*n*=3). *** indicates a *p*-value < 0.001.



Figure S5. Assessing the effects of increased NADPH consumption, Related to Experimental Procedures. (A) Expression of G6PD is not statistically different between wildtype cells and *IDH1* mutants. $\Delta\Delta$ CT values were determined by qPCR using beta actin as a housekeeping gene. (B) AGI-5198 protects *IDH1* mutants during H₂O₂ exposure. Cells were exposed to 1 mM H₂O₂ and cell viability was measured with a trypan blue exclusion assay. Data shown are mean values \pm s.d. (*n*=3). * indicates a *p*-value < 0.05.

Supplemental Tables

Supplemental Table S1. Fluxes of PPP and 2-HG production, Related to Figure 3.

	WT	R132H/+
PPP flux (fmol per glucose per cell per hour)	4.85 ± 0.17	6.92 ± 0.23
2-HG production flux (fmol/cell/hr)	-	6.50 ± 0.43

Supplemental Table S2. ISA values from cells labeled with U-¹³C glucose in the presence of exogenous acetate show that R132H/+ mutants use more acetate for palmitate synthesis compared to wildtype cells, Related to Figure 5.

	WT	R132H/+
	acetate	acetate
Dglucose	0.62 ± 0.02	0.57 ± 0.02
g (24 hr)	0.68 ± 0.03	0.61 ± 0.04

Supplemental Experimental Procedures

Kinetic Flux Profiling

Kinetic flux profiling of the oxidative phase of the PPP was performed with U-¹³C glucose. Both wildtype and R132H cells were exposed to labeled glucose for 0.5, 1, 3, and 5 min. Media was then aspirated and cells were immediately quenched with 80:20 methanol:water at 4 °C. Cells were extracted as previously described (Yuan et al., 2008). The decay function of the unlabeled glycolytic and pentose phosphate intermediates were fitted with Newtonian minimizations and 95% confidence intervals were determined from t-values (OriginLab).

Quantitative RT-PCR

Total RNA was isolated with TRIzol Reagent (Invitrogen) and single-strand cDNA was synthesized with a First Strand Synthesis Kit (Origene). PCR was performed with PowerUp SYBR Green Master Mix (Thermo Fisher) and StepOnePlus PCR instrument at 60 °C.

Malic enzyme measurement

Cells were given U-¹³C glutamine for 24 hr to assess malic enzyme activity. The M+3 isotopologue of lactate and the M+4 isotopologue of malate were used to evaluate malic enzyme activity (Fan et al., 2014).