Appendix

Infiltrative and drug-resistant slow-cycling cells support metabolic heterogeneity in glioblastoma

LB Hoang-Minh, FA Siebzehnrubl, C Yang, S Suzuki-Hatano, K Dajac, T Loche, N Andrews, M Schmoll, J Patel, K Amin, A Vuong, A Jimenez-Pascual, P Kubilis, TT Garrett, C Moneypenny, CA Pacak, J Huang, EJ Sayour, DA Mitchell, MR Sarkisian, BA Reynolds, LP Deleyrolle

Table of contents

Appendix Figure S1	p. 2
Appendix Figure S2	p. 3
Appendix Figure S3	p. 4

Appendix Figure Legends

Appendix Figure S1: (**A**) SCCs and FCCs were cultured in high glucose (HG, > 500 mg/dL) or physiological glucose (PG, 90-110 mg/dL) conditions for 24 hours. Cell death was quantified by flow cytometry through propidium iodide (PI) incorporation (L1, n=2; L2, n=4) (**A**) and cleaved caspase 3 expression (L1, n=2; L2, n=3) (**B**) SCCs and FCCs were cultured in 0, 5, or 20 mM 2DG for 24 hours and cell death quantified by flow cytometry through propidium iodide (PI) incorporation (L1, n=2; L2, n=4) and cleaved caspase 3 expression (L1, n=2; L2, n=3). (**C**) Using RT-qPCR, expression levels of the isoforms A, B, and C of lactate dehydrogenase were compared between SCCs and FCCs (n=3 technical replicates for each line) of the three hGBM cell lines. Changes in transcript levels are given as the mean percent change relative to FCCs. * p < 0.05, *** p < 0.001, one sample t-test. The combinatorial effect of administering glucose restriction and mitochondrial targeting with rotenone (n=16, **D**) or metformin (n=16, **E**) was measured using the CyQUANT assay after 24 hours of treatment for L0 and L2 hGBM cell lines (* p < 0.05, *** p < 0.001, one-way ANOVA with Tukey post-test, ^{##} p < 0.01, t-test).

Appendix Figure S2: List of all the pathways up-regulated in the SCCs, based on metabolites that are over-represented in SCCs compared to FCCs (fold change \geq 2).

Appendix Figure S3: (A) Raw values of mean fluorescence intensity (MFI) in SCCs and FCCs following LipidTox staining. GSEA of SCC and FCC RNA-seq data sets for enrichment of the autophagosome-lysosome signature, as defined by Perera *et. al.* (Perera et al., 2015) (B) and Jegga *et. al.* (Jegga et al., 2011) (C). FDR, false discovery rate; NES, normalized enrichment score; Nom., nominal. Raw values of mean fluorescence intensity (MFI) in SCCs and FCCs following immunostaining for LC3B (D) and LAMP2 (E).

Appendix Figure S1



Appendix Figure S2







Appendix Figure S3



4