

## **SUPPORTING INFORMATION**

“The BCKDK inhibitor BT2 is a chemical uncoupler that lowers mitochondrial ROS production and *de novo* lipogenesis”

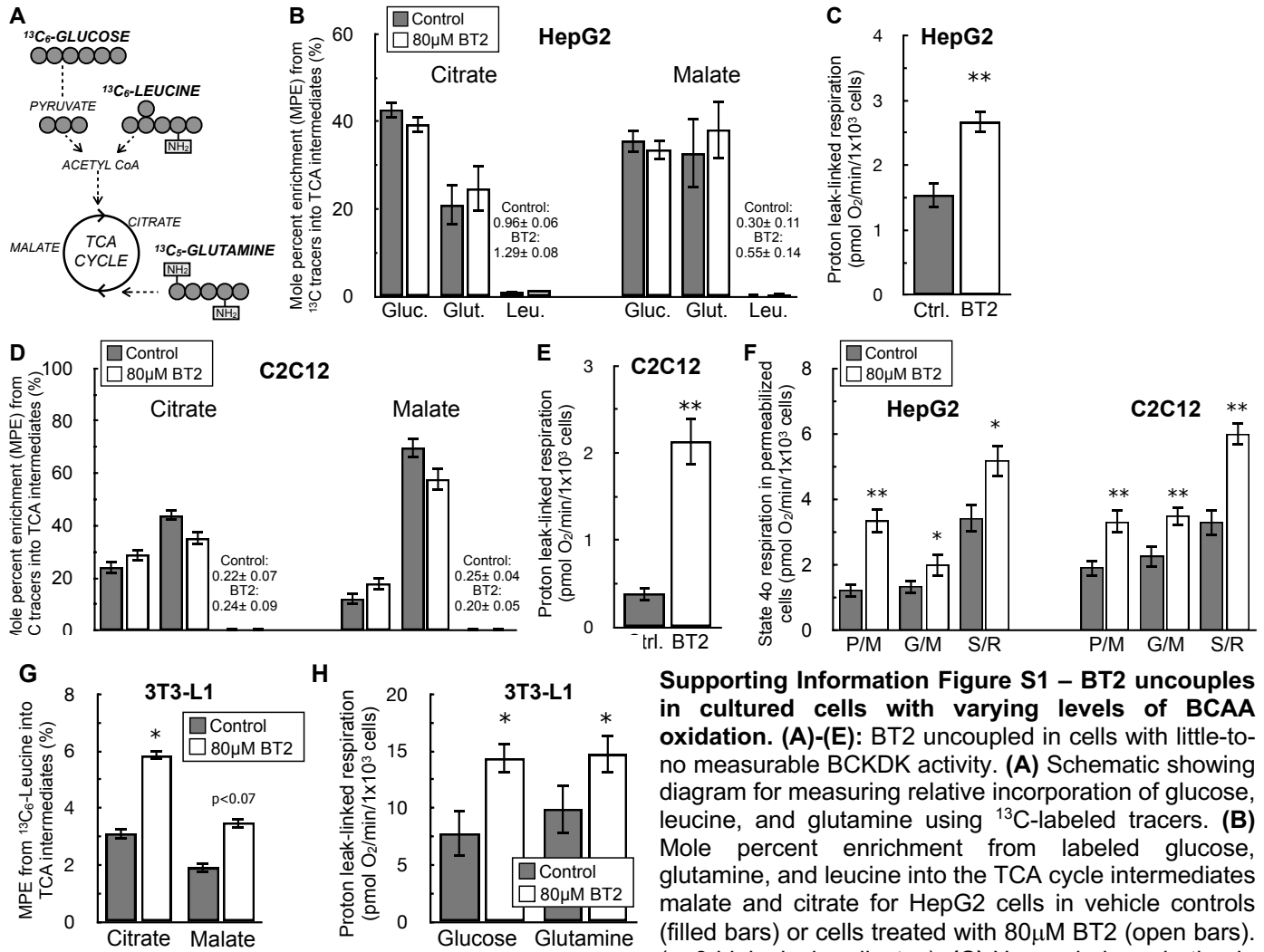
Includes:

Supporting Information Figure S1 – BT2 uncouples in cultured cells with varying levels of BCAA oxidation.

Supporting Information Figure S2 – 200 $\mu$ M BT2 results in proton conductance measurements that cannot be accurately quantified

Supporting Table 1 – ISA modeled values and 95% confidence intervals for individual technical replicates

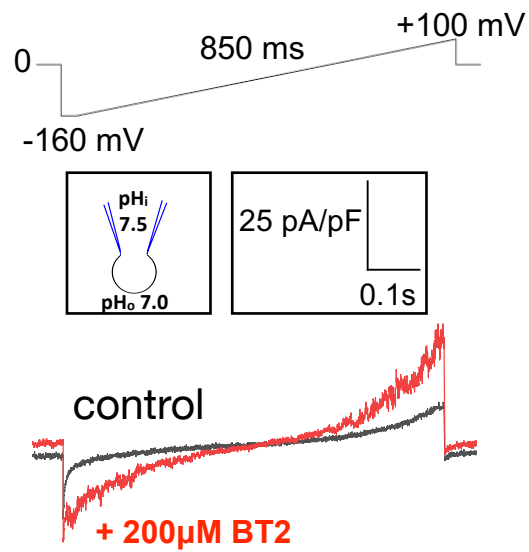
**Supporting Information Figure S1 – BT2 uncouples in cultured cells with varying levels of BCAA oxidation.**



**Supporting Information Figure S1 – BT2 uncouples in cultured cells with varying levels of BCAA oxidation. (A)-(E):** BT2 uncoupled in cells with little-to-no measurable BCKDK activity. **(A)** Schematic showing diagram for measuring relative incorporation of glucose, leucine, and glutamine using  $^{13}\text{C}$ -labeled tracers. **(B)** Mole percent enrichment from labeled glucose, glutamine, and leucine into the TCA cycle intermediates malate and citrate for HepG2 cells in vehicle controls (filled bars) or cells treated with 80  $\mu\text{M}$  BT2 (open bars). (n=3 biological replicates). **(C)** Uncoupled respiration in

HepG2 cells offered glucose, glutamine, and pyruvate in DMEM assay medium. (n=4 biological replicates). **(D)** Mole percent enrichment from labeled glucose, glutamine, and leucine into the TCA cycle intermediates malate and citrate for C2C12 myoblasts in vehicle controls (filled bars) or treated with 80  $\mu\text{M}$  BT2 (open bars). (n=4 biological replicates). **(E)** Uncoupled respiration in C2C12 cells offered glucose, glutamine, and pyruvate in DMEM assay medium. (n=3 biological replicates). **(F)** BT2 uncouples respiration in permeabilized cells with substrates that bypass BCKDK: State 4<sub>o</sub> respiration in permeabilized HepG2 (left) and C2C12 (right) cells in vehicle controls (filled bars) or cells treated with 80  $\mu\text{M}$  BT2 (open bars). Permeabilized cells were offered pyruvate/malate (P/M), glutamate/malate (G/M), or succinate/rotenone (S/R). [n=4 (C2C12) or n=5 (HepG2) biological replicates]. **(G)-(H):** As a 'positive control' for measurable leucine oxidation, 3T3-L1 adipocytes show BT2 behaves as expected, but uncouples irrespective of the respiratory substrate provided. **(G)** Mole percent enrichment from uniformly labeled  $^{13}\text{C}_6$ -leucine in differentiated 3T3-L1 adipocytes in vehicle controls (filled bars) or cells treated with 80  $\mu\text{M}$  BT2 (open bars). (n=4 biological replicates) **(H)** Uncoupled respiration in differentiated 3T3-L1 adipocytes in the presence or absence of BT2 as before. Cells were assayed in a simple-salts, Krebs-Henseleit buffer supplemented with either glucose or glutamine. (n=3 biological replicates). All data are presented as mean  $\pm$  S.E.M.

**Supporting Information Figure S2 – 200 $\mu$ M BT2 results in proton conductance measurements that cannot be accurately quantified**



**Supporting Information Figure S2:** Patch-clamp electrophysiology with cardiac mitoplasts demonstrates that 200 $\mu$ M BT2 yields a bigger current amplitude than 100 $\mu$ M, but disrupts the integrity of the inner membrane. As such, a proton conductance cannot be reliably quantified.

**Supporting Information Table S1: ISA modeled values and 95% confidence intervals for individual technical replicates.**

Sample	Measurement	Value	Lower range	Upper range	Sample	Measurement	Value	Lower range	Upper range
N1-NT1	D(M2-AcCoA)	0.309	0.264	0.353	N3-NT1	D(M2-AcCoA)	0.558	0.509	0.605
	D(M1-AcCoA)	0.027	0.006	0.052		D(M1-AcCoA)	0.034	0.010	0.066
	1-D(AcCoA)	0.665	0.621	0.708		1-D(AcCoA)	0.408	0.360	0.457
	g(t) palmitate	0.419	0.375	0.459		g(t) palmitate	0.456	0.408	0.498
	TOTAL AcCoA contribution	0.335	0.270	0.405		TOTAL AcCoA contribution	0.592	0.519	0.671
N1-NT2	D(M2-AcCoA)	0.288	0.240	0.335	N3-NT2	D(M2-AcCoA)	0.292	0.236	0.348
	D(M1-AcCoA)	0.027	0.006	0.055		D(M1-AcCoA)	0.030	0.005	0.065
	1-D(AcCoA)	0.685	0.638	0.731		1-D(AcCoA)	0.677	0.624	0.730
	g(t) palmitate	0.394	0.349	0.435		g(t) palmitate	0.354	0.307	0.396
	TOTAL AcCoA contribution	0.315	0.246	0.390		TOTAL AcCoA contribution	0.323	0.241	0.412
N1-NT3	D(M2-AcCoA)	0.244	0.194	0.291	N3-NT3	D(M2-AcCoA)	0.326	0.248	0.398
	D(M1-AcCoA)	0.032	0.009	0.064		D(M1-AcCoA)	0.030	0.000	0.078
	1-D(AcCoA)	0.724	0.678	0.769		1-D(AcCoA)	0.644	0.574	0.715
	g(t) palmitate	0.395	0.348	0.437		g(t) palmitate	0.262	0.211	0.306
	TOTAL AcCoA contribution	0.276	0.204	0.355		TOTAL AcCoA contribution	0.356	0.248	0.476
N1-BT2-1	D(M2-AcCoA)	0.322	0.255	0.385	N3-BT2-1	D(M2-AcCoA)	0.327	0.282	0.372
	D(M1-AcCoA)	0.029	0.000	0.070		D(M1-AcCoA)	0.023	0.002	0.048
	1-D(AcCoA)	0.650	0.588	0.712		1-D(AcCoA)	0.650	0.605	0.694
	g(t) palmitate	0.285	0.236	0.329		g(t) palmitate	0.366	0.325	0.403
	TOTAL AcCoA contribution	0.350	0.255	0.455		TOTAL AcCoA contribution	0.350	0.284	0.420
N1-BT2-2	D(M2-AcCoA)	0.299	0.221	0.370	N3-BT2-2	D(M2-AcCoA)	0.295	0.223	0.362
	D(M1-AcCoA)	0.030	0.000	0.079		D(M1-AcCoA)	0.030	0.000	0.075
	1-D(AcCoA)	0.671	0.603	0.739		1-D(AcCoA)	0.675	0.611	0.739
	g(t) palmitate	0.258	0.209	0.303		g(t) palmitate	0.235	0.191	0.274
	TOTAL AcCoA contribution	0.329	0.221	0.449		TOTAL AcCoA contribution	0.325	0.223	0.437
N1-BT2-3	D(M2-AcCoA)	0.231	0.076	0.320	N3-BT2-3	D(M2-AcCoA)	0.316	0.245	0.383
	D(M1-AcCoA)	0.046	0.004	0.169		D(M1-AcCoA)	0.028	0.000	0.072
	1-D(AcCoA)	0.723	0.646	0.826		1-D(AcCoA)	0.655	0.591	0.720
	g(t) palmitate	0.275	0.218	0.325		g(t) palmitate	0.248	0.202	0.289
	TOTAL AcCoA contribution	0.277	0.079	0.489		TOTAL AcCoA contribution	0.345	0.245	0.455
N1-DNP-1	D(M2-AcCoA)	0.355	0.236	0.455	N3-DNP-1	D(M2-AcCoA)	0.551	0.483	0.615
	D(M1-AcCoA)	0.024	0.000	0.095		D(M1-AcCoA)	0.023	0.000	0.062
	1-D(AcCoA)	0.621	0.524	0.721		1-D(AcCoA)	0.426	0.360	0.495
	g(t) palmitate	0.197	0.141	0.247		g(t) palmitate	0.111	0.088	0.134
	TOTAL AcCoA contribution	0.379	0.236	0.550		TOTAL AcCoA contribution	0.574	0.483	0.677
N1-DNP-2	D(M2-AcCoA)	0.387	0.243	0.505	N3-DNP-2	D(M2-AcCoA)	0.338	0.000	0.483
	D(M1-AcCoA)	0.022	0.000	0.105		D(M1-AcCoA)	0.031	0.000	0.287
	1-D(AcCoA)	0.591	0.476	0.713		1-D(AcCoA)	0.631	0.492	0.993
	g(t) palmitate	0.171	0.114	0.221		g(t) palmitate	0.175	0.110	0.231
	TOTAL AcCoA contribution	0.409	0.243	0.610		TOTAL AcCoA contribution	0.369	0.000	0.770
N1-DNP-3	D(M2-AcCoA)	0.346	0.201	0.456	N3-DNP-3	D(M2-AcCoA)	0.295	0.178	0.386
	D(M1-AcCoA)	0.025	0.000	0.109		D(M1-AcCoA)	0.029	0.000	0.100
	1-D(AcCoA)	0.629	0.524	0.743		1-D(AcCoA)	0.677	0.592	0.766
	g(t) palmitate	0.178	0.123	0.228		g(t) palmitate	0.172	0.127	0.212
	TOTAL AcCoA contribution	0.371	0.201	0.565		TOTAL AcCoA contribution	0.323	0.178	0.486
N2-NT1	D(M2-AcCoA)	0.224	0.187	0.260	N4-NT1	D(M2-AcCoA)	0.346	0.306	0.386
	D(M1-AcCoA)	0.031	0.012	0.054		D(M1-AcCoA)	0.023	0.004	0.045
	1-D(AcCoA)	0.746	0.711	0.780		1-D(AcCoA)	0.631	0.591	0.672
	g(t) palmitate	0.395	0.358	0.430		g(t) palmitate	0.378	0.340	0.412
	TOTAL AcCoA contribution	0.254	0.198	0.314		TOTAL AcCoA contribution	0.369	0.309	0.431
N2-NT2	D(M2-AcCoA)	0.387	0.337	0.436	N4-NT2	D(M2-AcCoA)	0.523	0.475	0.571
	D(M1-AcCoA)	0.024	0.002	0.052		D(M1-AcCoA)	0.029	0.007	0.059
	1-D(AcCoA)	0.589	0.540	0.638		1-D(AcCoA)	0.447	0.399	0.496
	g(t) palmitate	0.420	0.373	0.462		g(t) palmitate	0.429	0.383	0.470
	TOTAL AcCoA contribution	0.411	0.339	0.488		TOTAL AcCoA contribution	0.553	0.481	0.630
N2-NT3	D(M2-AcCoA)	0.249	0.203	0.293	N4-NT3	D(M2-AcCoA)	0.435	0.381	0.489
	D(M1-AcCoA)	0.030	0.008	0.059		D(M1-AcCoA)	0.023	0.000	0.052
	1-D(AcCoA)	0.721	0.679	0.763		1-D(AcCoA)	0.541	0.487	0.595
	g(t) palmitate	0.393	0.350	0.432		g(t) palmitate	0.412	0.363	0.455
	TOTAL AcCoA contribution	0.279	0.212	0.352		TOTAL AcCoA contribution	0.459	0.381	0.541
N2-BT2-1	D(M2-AcCoA)	0.398	0.295	0.492	N4-BT2-1	D(M2-AcCoA)	0.300	0.246	0.352
	D(M1-AcCoA)	0.029	0.000	0.094		D(M1-AcCoA)	0.026	0.001	0.058
	1-D(AcCoA)	0.573	0.480	0.669		1-D(AcCoA)	0.674	0.624	0.724
	g(t) palmitate	0.231	0.172	0.281		g(t) palmitate	0.266	0.226	0.302
	TOTAL AcCoA contribution	0.427	0.295	0.586		TOTAL AcCoA contribution	0.326	0.247	0.411
N2-BT2-2	D(M2-AcCoA)	0.282	0.203	0.354	N4-BT2-2	D(M2-AcCoA)	0.317	0.266	0.367
	D(M1-AcCoA)	0.032	0.000	0.084		D(M1-AcCoA)	0.023	0.000	0.053
	1-D(AcCoA)	0.685	0.619	0.753		1-D(AcCoA)	0.660	0.611	0.709
	g(t) palmitate	0.279	0.227	0.325		g(t) palmitate	0.232	0.198	0.263
	TOTAL AcCoA contribution	0.315	0.203	0.437		TOTAL AcCoA contribution	0.340	0.266	0.420
N2-BT2-3	D(M2-AcCoA)	0.369	0.271	0.459	N4-DNP-1	D(M2-AcCoA)	0.365	0.247	0.468
	D(M1-AcCoA)	0.030	0.000	0.091		D(M1-AcCoA)	0.026	0.000	0.093
	1-D(AcCoA)	0.601	0.514	0.690		1-D(AcCoA)	0.609	0.508	0.714
	g(t) palmitate	0.252	0.193	0.303		g(t) palmitate	0.183	0.133	0.229
	TOTAL AcCoA contribution	0.399	0.271	0.549		TOTAL AcCoA contribution	0.391	0.247	0.561
N2-DNP-1	D(M2-AcCoA)	0.464	0.319	0.590	N4-DNP-2	D(M2-AcCoA)	0.340	0.271	0.408
	D(M1-AcCoA)	0.021	0.000	0.117		D(M1-AcCoA)	0.019	0.000	0.056
	1-D(AcCoA)	0.515	0.390	0.650		1-D(AcCoA)	0.641	0.576	0.706
	g(t) palmitate	0.150	0.094	0.201		g(t) palmitate	0.163	0.129	0.195
	TOTAL AcCoA contribution	0.485	0.319	0.707		TOTAL AcCoA contribution	0.359	0.271	0.464
					N4-DNP-3	D(M2-AcCoA)	0.316	0.239	0.388
						D(M1-AcCoA)	0.021	0.000	0.064
						1-D(AcCoA)	0.663	0.594	0.732
						g(t) palmitate	0.162	0.126	0.194
						TOTAL AcCoA contribution	0.337	0.239	0.452