

**Table 1: Summary of exchange rates reported in this paper.**

**1. Protein specificity of the malate/phosphate exchange**

Substrates		Exchange rate, $\mu\text{M/s}$			
		UCP3	UCP2	ANT1	None
Phosphate	Malate	$9.57 \pm 4.29$	$14.5 \pm 3.5$	$0.19 \pm 0.01$	$0.23 \pm 0.09$

**2. Substrate specificity of UCP3 and UCP2**

		Exchange rate, $\mu\text{M/s}$		Exchange rate $\text{mmol}/(\text{min} \cdot \text{g}_{\text{Protein}})$	
		UCP3	UCP2	UCP3	UCP2
<i>Homoexchange</i>					
Malate	Malate	$7.94 \pm 1.88$	$16.6 \pm 3.5$	$9.93 \pm 2.35$	$20.8 \pm 4.3$
Phosphate	Phosphate	$11.7 \pm 3.20$	$8.00 \pm 1.40$	$14.7 \pm 4.0$	$10.0 \pm 1.8$
<i>Heteroexchange</i>					
Phosphate	Asparagine	$0.40 \pm 0.04$	$0.27 \pm 0.04$	$0.50 \pm 0.05$	$0.34 \pm 0.05$
	Aspartate	$19.1 \pm 4.6$	$1.31 \pm 0.19$	$23.9 \pm 5.8$	$1.64 \pm 0.24$
	Citrate	$1.10 \pm 0.39$	$1.14 \pm 0.38$	$1.38 \pm 0.49$	$1.43 \pm 0.48$
	Glutamate	$0.29 \pm 0.07$	$0.36 \pm 0.06$	$0.36 \pm 0.08$	$0.45 \pm 0.08$
	Glutamine	$0.27 \pm 0.12$	$0.24 \pm 0.02$	$0.34 \pm 0.15$	$0.30 \pm 0.03$
	Malate	$9.57 \pm 4.39$	$14.5 \pm 3.50$	$12.0 \pm 5.5$	$18.1 \pm 4.4$
	Malonate	$1.96 \pm 0.72$	$16.5 \pm 2.0$	$2.45 \pm 0.90$	$20.6 \pm 2.5$
	None	$0.13 \pm 0.12$	$0.18 \pm 0.06$	$0.16 \pm 0.15$	$0.23 \pm 0.08$
	Oxaloacetate	$5.07 \pm 1.92$	$1.26 \pm 0.18$	$6.34 \pm 2.40$	$1.58 \pm 0.23$
	Oxoglutarate	$0.37 \pm 0.04$	$0.32 \pm 0.04$	$0.46 \pm 0.05$	$0.40 \pm 0.05$
	Succinate	$1.16 \pm 0.17$	$0.52 \pm 0.19$	$1.45 \pm 0.21$	$0.65 \pm 0.24$
	Sulfate	$14.0 \pm 4.1$	$18.4 \pm 3.7$	$17.5 \pm 5.1$	$23.0 \pm 4.7$
	Sulfite	$12.9 \pm 4.5$	$0.24 \pm 0.06$	$16.1 \pm 5.6$	$0.30 \pm 0.08$
Thiosulfate	$2.70 \pm 0.36$	$0.54 \pm 0.36$	$3.38 \pm 0.45$	$0.68 \pm 0.45$	

## SUPPORTING INFORMATION

**Uncoupling protein 3 catalyzes the exchange of C4 metabolites similar to UCP2.**

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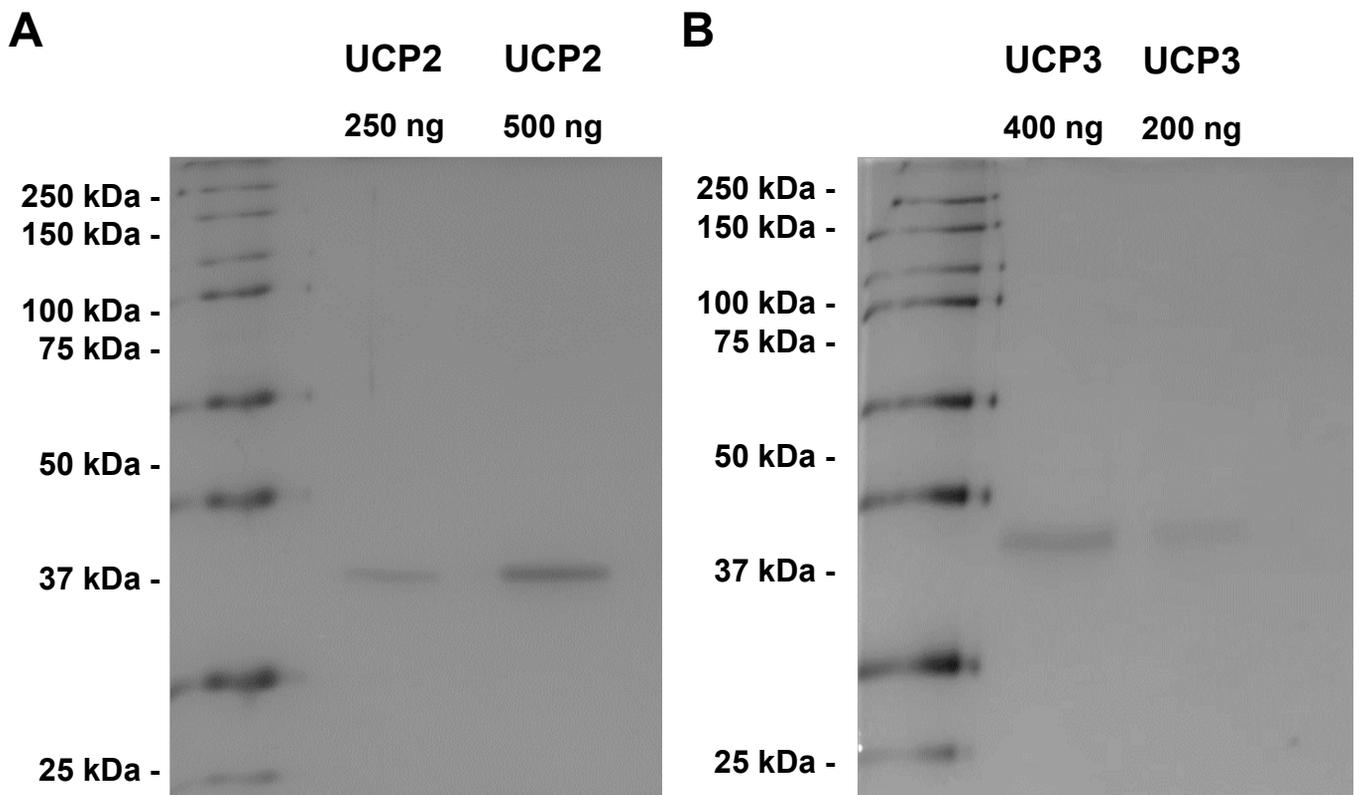
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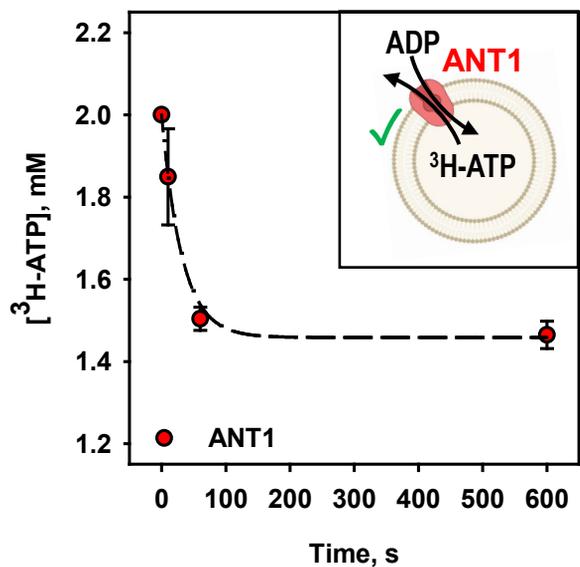
<i>mUCP3</i>	MVGLQPSEVPPTTVVKFLGAGTAACFADLLTF <b>PLDTAK</b> VRLQIQGENPG----AQSVQYR	56
<i>mUCP2</i>	MVGFKATDVPPTATVKFLGAGTAACIADLITF <b>PLDTAK</b> VRLQIQGESQGLVRTAASAQYR	60
<i>mANT1</i>	-----MGDQALSFLKDFLAGGIAAAV <b>SKTAVAP</b> <b>PIERVK</b> LLLQVQHASKQ---ISA EKQYK	52
<i>mUCP3</i>	GVLGTILTMVRTEGPRSPYSGLVAGL <b>HR</b> QMSFASIRIGLYDSVKQFYTPKGADHSSV---	113
<i>mUCP2</i>	GVLGTILTMVRTEGPRSLYNGLVAGL <b>QR</b> QMSFASVIRIGLYDSVKQFYT-KGSEHAGI---	116
<i>mANT1</i>	GIIDCVVRI <b>PKE</b> QGFLSFWRGNLANV <b>IR</b> YFPTQALNFAFKDKYKQIFLGGVDRHKQFWRY	112
<i>mUCP3</i>	-AIRILAGCTTGAMAVTCAQ <b>PTDVVK</b> VRFQAMIRLGTGGERKYRGTMDAYRTIAREEGVR	172
<i>mUCP2</i>	-GSRLLAGSTTGALAVAVAQ <b>PTDVVK</b> VRFQAQARA--GGRRYQSTVEAYKTIAREEGIR	173
<i>mANT1</i>	FAGNLASGGAAGATSLCFVY <b>PLDFAR</b> TRLAADVKGK-SSQREFNGLGDCLTKIFKSDGLK	171
<i>mUCP3</i>	GLWKGTWPNIT <b>R</b> NAIVNCAEMVTYDIIKEKLLSHLFTDNFPCHFVSAFGAGFCATVVAS	232
<i>mUCP2</i>	GLWKGTSPNVA <b>R</b> NAIVNCAELVTYDLIKDTLLKANLMTDDLPCHFVSAFGAGFCTTVIAS	233
<i>mANT1</i>	GLYQGFSSVSVQGI <b>I</b> IYRAAYFGVYDTAKGMLPDPKNVHIIV-SWMIAQSVTAVAGLVSY-	229
<i>mUCP3</i>	<b>PVDVVK</b> TRYMNAP-----LGRYRSPLHCMLKMVAQEGPTAFYKGFVPSFL <b>R</b> LGAWNVMMF	287
<i>mUCP2</i>	<b>PVDVVK</b> TRYMNSA-----LGQYHSAGHCALTMLRKEGPRAFYKGFMPFSL <b>R</b> LGSWNVVMF	288
<i>mANT1</i>	<b>PFDTVR</b> RRMMM <b>Q</b> SGRKGADIMYTGTLDCWRKIAKDEGANAF <b>F</b> KGAWSNV <b>L</b> <b>R</b> G <b>M</b> -GGAFVL	288
<i>mUCP3</i>	VTYEQLKRALMKVQVLRES <b>PF</b>	308
<i>mUCP2</i>	VTYEQLKRALMAAYQSRE <b>APF</b>	309
<i>mANT1</i>	VLYDEIKKYV-----	298

**Figure S1. Alignment of the amino acid sequences of murine UCP3, UCP2 and ANT1.** The highly conserved PX[DE]XX[KR] signature motif of mitochondrial carriers is marked in blue. The three amino acids involved in the putative substrate binding site in the central cavity of *mUCP3*, *mUCP2* and *mANT1* are colored in red.



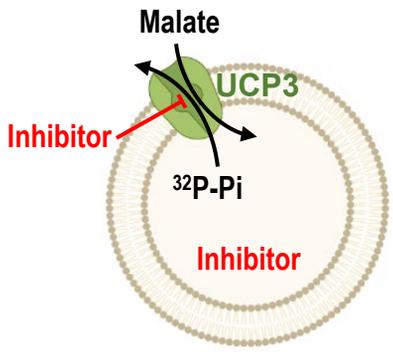
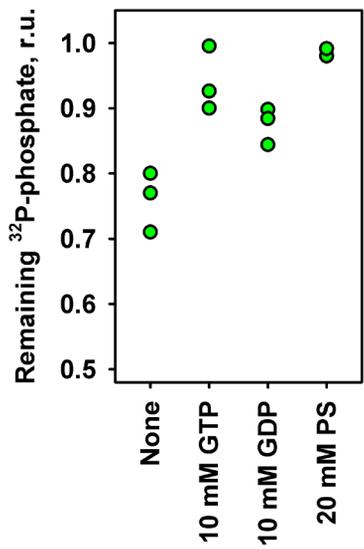
**Figure S2. Representative silver staining of reconstituted UCP2 and UCP3.**

For quality control, 250 ng or 500 ng of UCP2 (A) and 400 ng and 200 ng of UCP3 (B) containing proteoliposomes were loaded onto a 15% acrylamide gel and SDS-PAGE was conducted. Subsequently, proteins were visualized by silver staining.



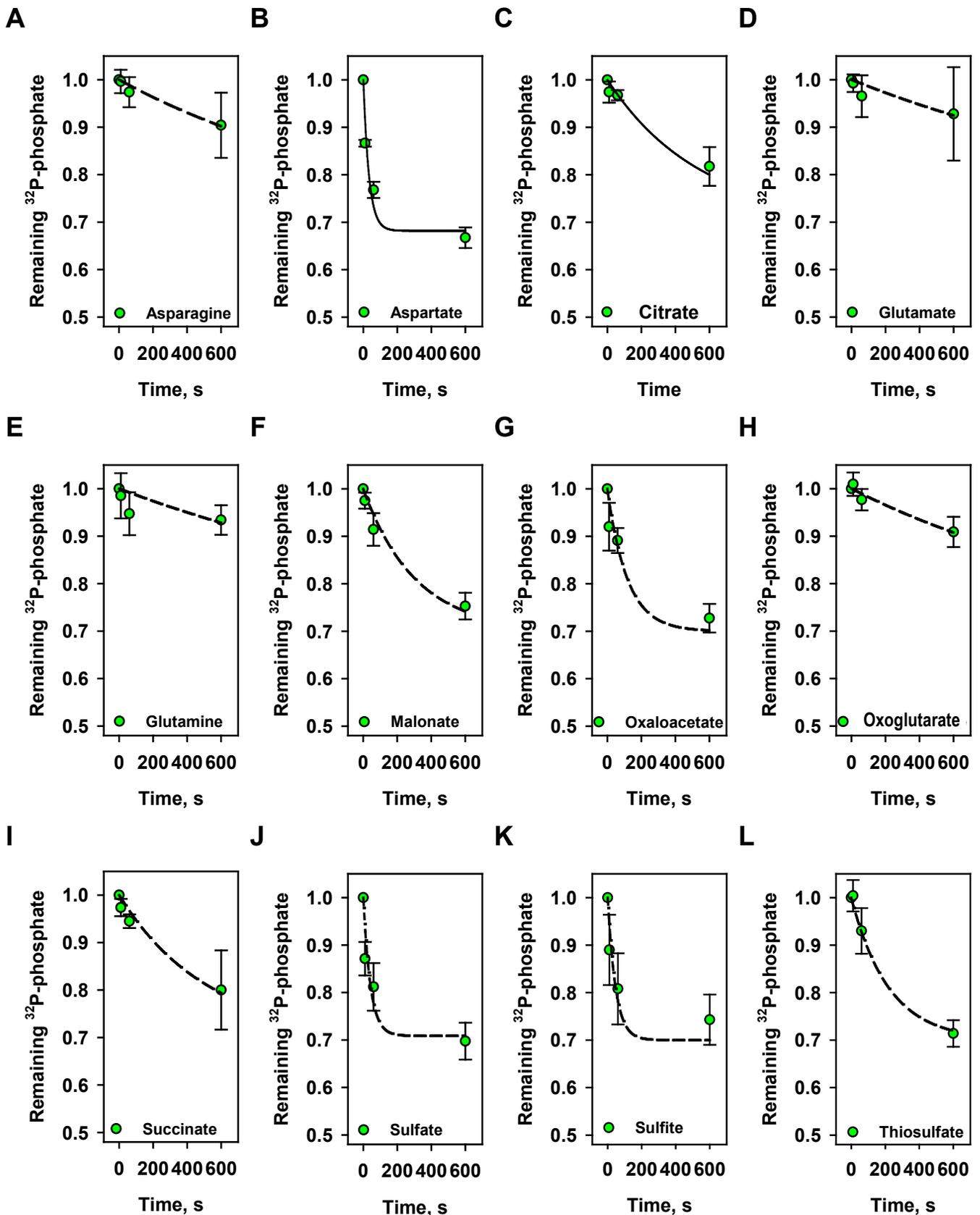
**Figure S3. ANT1-mediated ADP/ATP exchange as a control for ANT1 activity.**

Efflux of ATP from liposomes over time measured by remaining intraliposomal <sup>3</sup>H-ATP radioactivity of ANT1-containing liposomes. Concentration of intraliposomal <sup>3</sup>H-ATP and extraliposomal ADP was 2 mM. **Inset:** Experimental setup to test ADP/ATP exchange of ANT1.

**A****B**

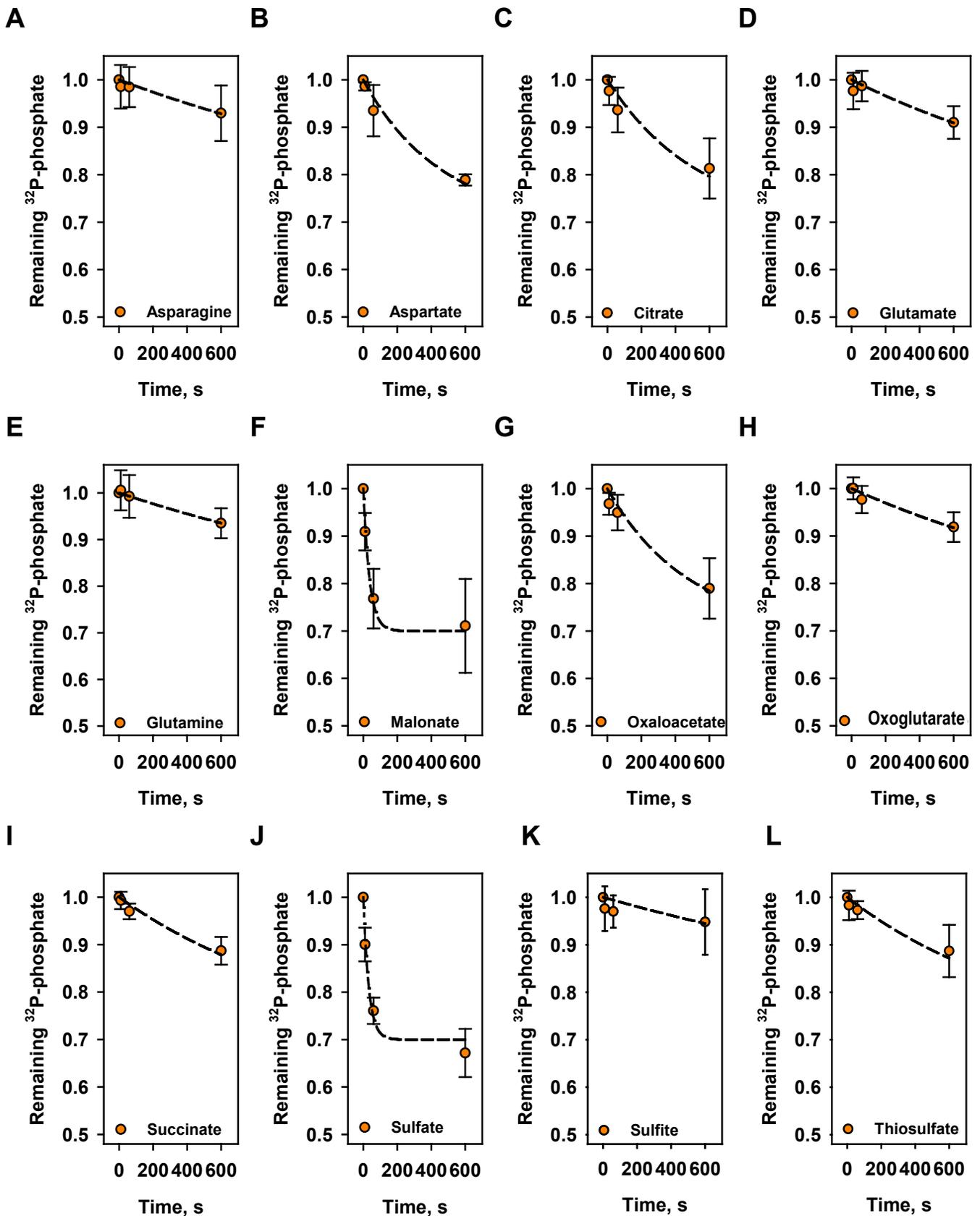
**Figure S4. Inhibition of UCP3-mediated phosphate/malate exchange.**

**(A)** Experimental setup to measure the exchange of <sup>32</sup>P-phosphate against malate in the presence of various inhibitors. **(B)** Remaining <sup>32</sup>P-phosphate in liposomes containing UCP3 after T = 600 s as the ratio to the initial <sup>32</sup>P-phosphate amount at T = 0 s. Concentration of <sup>32</sup>P-phosphate and malate was 2 mM, respectively. In the experiments, the lipid concentration was 4 mg/ml and the protein concentration was 4 μg/mg of lipid. Membranes were made of 45:45:10 mol% PC:PE:CL. Buffer solution contained 50 mM Na<sub>2</sub>SO<sub>4</sub>, 10 mM Tris, 10 mM MES and 0.6 mM EGTA at pH = 7.34 and T = 296 K.



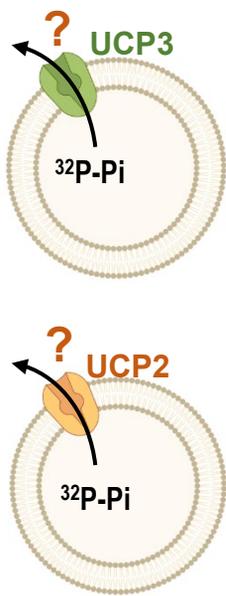
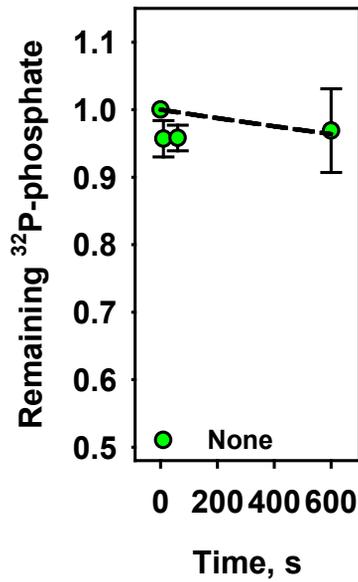
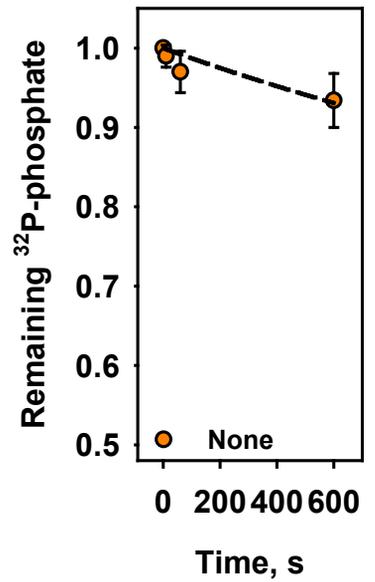
**Figure S5. Evaluation of different substrates as possible metabolites transported by UCP3**

Time course of intraliposomal phosphate concentration in the presence of UCP3 and extraliposomal asparagine (A), aspartate (B), citrate (C), glutamate (D), glutamine (E), malonate (F), oxaloacetate (G), oxoglutarate (H), succinate (I), sulfate (J), sulfite (K) or thiosulfate (L). In the experiments, the lipid concentration was 4 mg/ml and the protein concentration was 4  $\mu\text{g}/\text{mg}$  of lipid. Membranes were made of 45:45:10 mol% PC:PE:CL. Buffer solution contained 50 mM  $\text{Na}_2\text{SO}_4$ , 10 mM Tris, 10 mM MES and 0.6 mM EGTA at pH = 7.34 and T = 296 K. Data are the mean  $\pm$  SD of at least three independent experiments.



**Figure S6. Evaluation of different substrates as possible metabolites transported by UCP2**

Time course of intraliposomal phosphate concentration in the presence of UCP2 and extraliposomal asparagine (A), aspartate (B), citrate (C), glutamate (D), glutamine (E), malonate (F), oxaloacetate (G), oxoglutarate (H), succinate (I), sulfate (J), sulfite (K) or thiosulfate (L). In the experiments, the lipid concentration was 4 mg/ml and the protein concentration was 4  $\mu\text{g}/\text{mg}$  of lipid. Membranes were made of 45:45:10 mol% PC:PE:CL. Buffer solution contained 50 mM  $\text{Na}_2\text{SO}_4$ , 10 mM Tris, 10 mM MES and 0.6 mM EGTA at pH = 7.34 and T = 296 K. Data are the mean  $\pm$  SD of at least three independent experiments.

**A****B****C**

### Figure S7. Uniport transport of phosphate of UCP3 and UCP2

**(A)** Release of intraliposomal  $^{32}\text{P}$ -phosphate in the absence of an external substrate. **(B, C)** Time course of intraliposomal  $^{32}\text{P}$ -phosphate efflux concentration in the presence of liposomes containing UCP3 **(B)** or UCP2 **(C)**. Concentration of  $^{32}\text{P}$ -phosphate was 2 mM. In the experiments, the lipid concentration was 4 mg/ml and the protein concentration was 4  $\mu\text{g}/\text{mg}$  of lipid. Membranes were made of 45:45:10 mol% PC:PE:CL. Buffer solution contained 50 mM  $\text{Na}_2\text{SO}_4$ , 10 mM Tris, 10 mM MES and 0.6 mM EGTA at pH = 7.34 and T = 296 K.

**Table S1: Overview of transport rates of various mitochondrial carriers reconstituted into liposomes.** Abbreviations are: cit – citrate, mal – malate, 2-oxog – oxoglutarate, Pi – phosphate, asp – aspartate, glu – glutamate and cys - cysteinsulfinate

Protein	Species	Transport rate, mmol/min/g protein	Substrates	Ref.
<b>Citrate carrier (CiC – SLC25A1)</b>	Rat	2.05 ± 0.09	cit/cit	[1]
		1.87 ± 0.11	mal/cit	
		2.06 ± 0.13		
		1.98 ± 0.16	mal/mal	
<b>Phosphate carrier (PiC – SLC25A3)</b>	Bovine	70 - 90	Pi/Pi	[2]
		30	Pi/OH <sup>-</sup>	
<b>Dicarboxylate carrier (DiC – SLC25A10)</b>	Rat	6.4 – 6.7	mal/Pi	[3]
		7.3	mal/mal	
<b>Oxoglutarate carrier (OGC – SLC25A11)</b>	Bovine	2.96 - 22	oxog/oxog	[4] – [6]
		10.34 ± 0.37		[7]
<b>Aspartate/glutamate carrier 1 (AGC1 – SLC25A12)</b>	Human	25.5 – 40.5	asp/asp	[8]
		16.1 – 23.7	asp/glu	
		21.8	glu/glu	
		17.7 – 27.6	asp/cys	
		14.2	glu/cys	
<b>Aspartate/glutamate carrier 2 (AGC2 – SLC25A13)</b>	Human	92.8 – 178.4	asp/asp	[8]
		55.3 – 118	asp/glu	
		90.3	glu/glu	
		81.2 – 133.8	asp/cys	
		65.9	glu/cys	

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