

ThermoMouse: an *in vivo* model to identify modulators of UCP1 expression in brown adipose tissue

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SUPPLEMENTARY INFORMATION

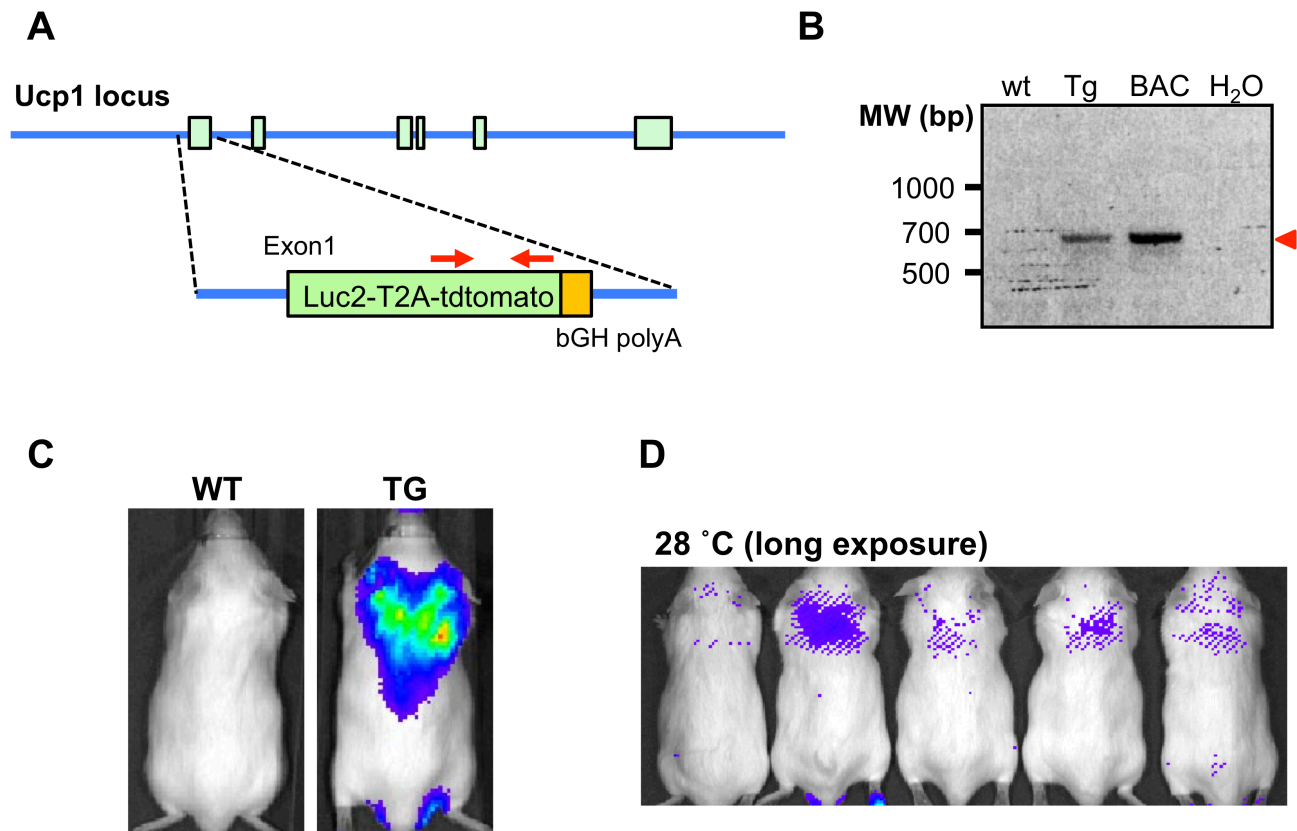


Figure S1 related to Figure 1. Generation of *Ucp1*-luciferase reporter mice

A: Genomic structure of the *Ucp1*-luciferase BAC transgene. The coding sequence of luciferase2-T2A-tdTomato followed by a bovine growth hormone (bGH)-derived polyadenylation signal (polyA) was inserted into the initiation codon of *Ucp1* located in exon 1.

B: BAC transgenic mice (Tg) were identified by PCR analysis using the primers shown as red arrows in (A). The BAC construct (BAC) was used as a positive control. Mouse wild-type (wt) DNA and water (H₂O) were used as negative controls.

C: Luciferase signal in wild-type (wt) and *Ucp1*-luciferase transgenic mice (Tg) imaged using the IVIS Spectrum Imaging system at 9°C.

D: Long-exposure image of luciferase signal in *Ucp1*-luciferase reporter mice maintained at 28°C.

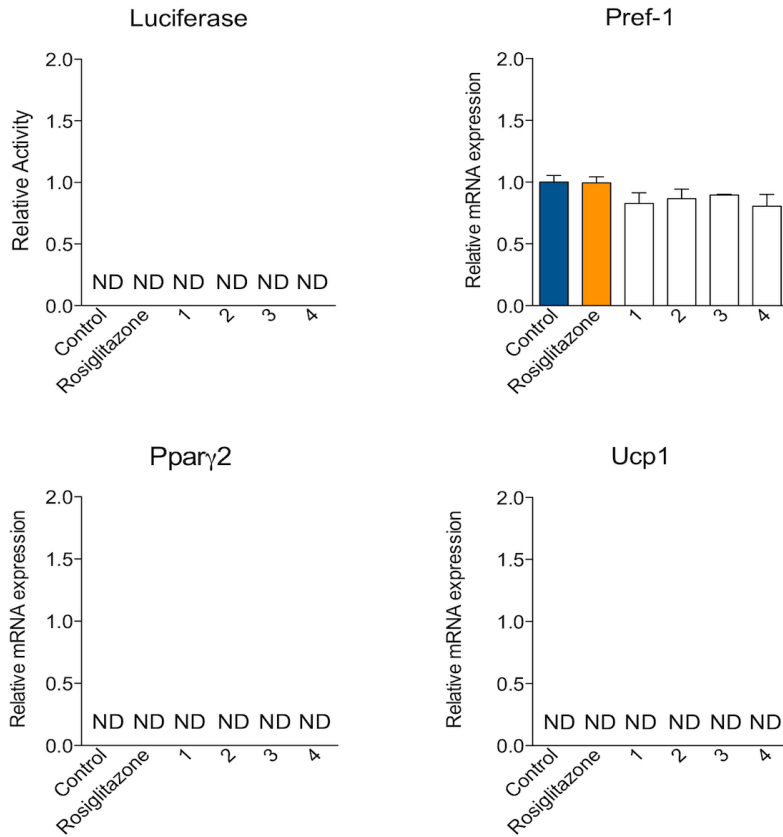


Figure S2 related to Figure 3. Effect of screening hits on *Ucp1*-luciferase preadipocytes
 Subconfluent *Ucp1*-luciferase preadipocytes were treated with selected screen hits (10 μ M) or rosiglitazone (0.5 μ M) for 24 hr prior to luciferase and gene expression analysis (n=4). ND, not detectable.

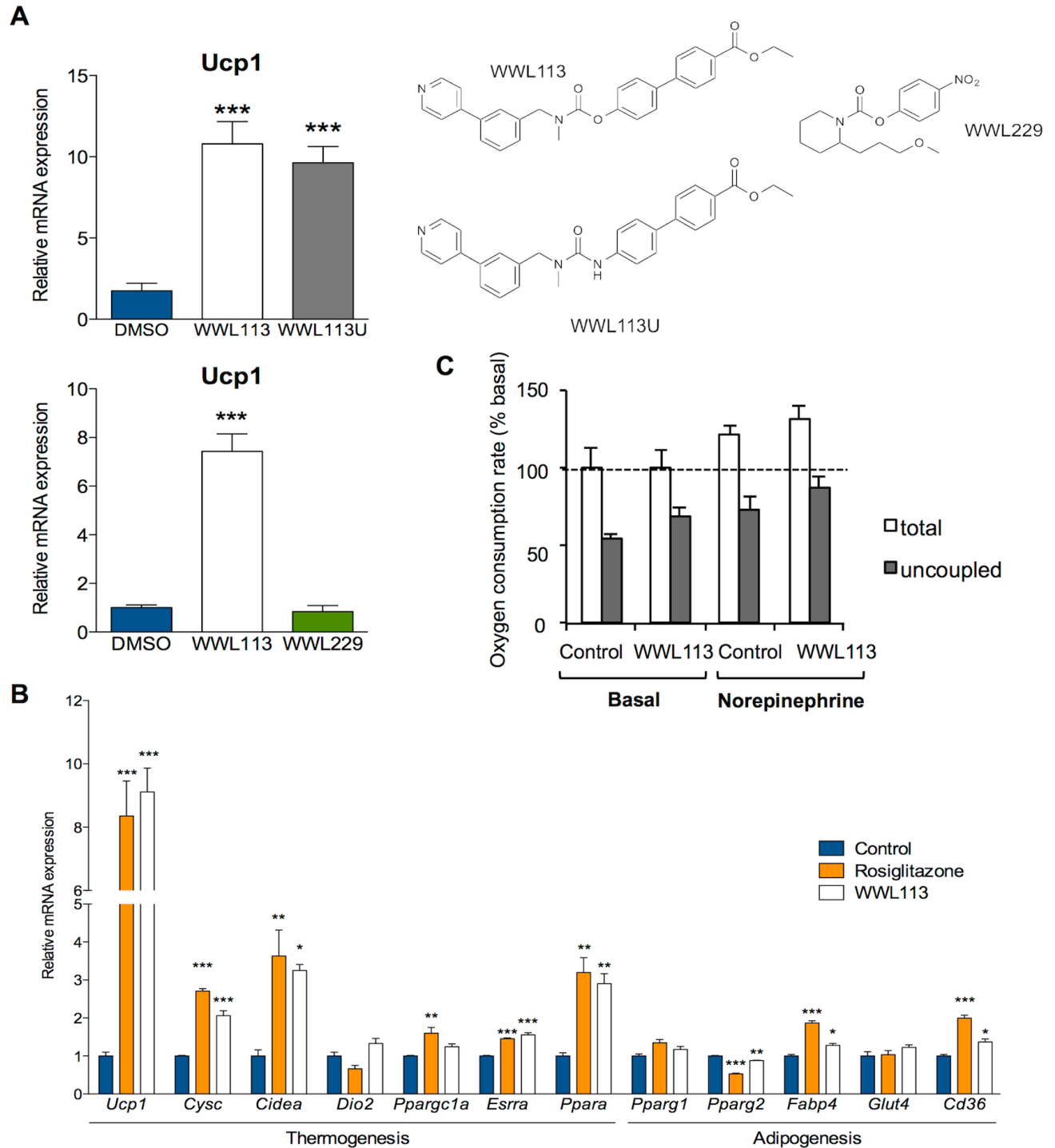


Figure S3 related to Figure 3. Induction of *Ucp1* expression and cellular respiration by WWL113 is not mediated by *Ces3* inhibition

A. *Ucp1* mRNA in differentiated primary brown adipocytes treated with WWL113, WWL113U or WWL229 (all at 10 μ M) for 24 hr. WWL113 and WWL229 inhibit *Ces3*, while WWL113U does not (n=4). Structures are shown on the right. *** $P < 0.001$ vs. control.

B: Gene expression in differentiated primary brown adipocytes treated with vehicle, WWL113 (10 μ M) or rosiglitazone (0.5 μ M) for 48 hr (n=3). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ vs. control.

C: Total and uncoupled (oligomycin-insensitive) respiration measured in differentiated brown adipocytes (5×10^5 cells/sample) treated with WWL113 (10 μ M) in the presence or absence of norepinephrine (0.1 μ M) (n=3-4). Data is expressed as relative to basal for each condition. * $P < 0.05$ vs. control.

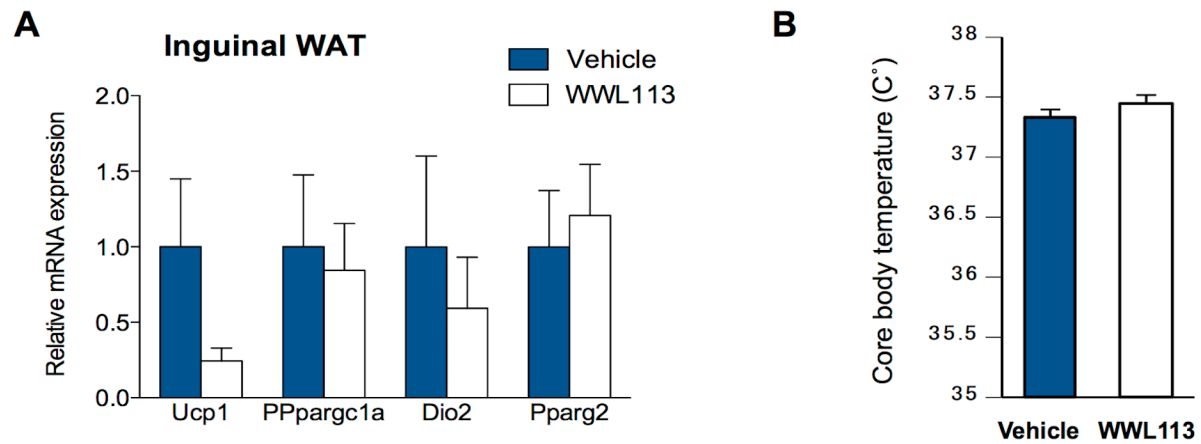


Figure S4 related to Figure 5. WWL113 does not enhance *Ucp1* expression in inguinal WAT

A: Expression of thermogenic genes in inguinal WAT of C57BL/6 mice treated daily with WWL113 (50 mg/kg) or vehicle for 5 days (n=5).

B: Core body temperature in wild-type mice treated with vehicle or WWL113 (50 mg/kg) for 7 days (n=6).

Supplementary Table related to Experimental Procedures: Primer sequences

Gene Id	Forward primer	Reverse Primer
<i>Adipoq</i>	GCACTGGCAAGTTCTACTGCAA	GTAGGTGAAGAGAACGGCCTTGT
<i>Cidea</i>	ATCACAACTGGCCTGGTTACG	TACTACCCGGTGTCCATTTCT
<i>Cox7a1</i>	CAGCGTCATGGTCAGTCTGT	AGAAAACCGTGTGGCAGAGA
<i>Dio2</i>	CAGTGTGGTGCACGTCTCCAATC	TGAACCAAAGTTGACCACCAG
<i>Fabp4</i>	ACACCGAGATTTTCCTTCAAACCTG	CCATCTAGGGTTATGATGCTCTTCA
<i>Pparg</i>	GTGCCAGTTTCGATCCGTAGA	GGCCAGCATCGTGTAGATGA
<i>Ppargc1a</i>	AGCCGTGACCACTGACAACGAG	GCTGCATGGTTCTGAGTGCTAAG
<i>Prdm16</i>	GGCGAGGAAGCTAGCCAAA	GGTCTCCTCCTCGGCACTCT
<i>Tbp</i>	ACCCTTCACCAATGACTCCTATG	TGACTGCAGCAAATCGCTTGG
<i>Ucp1</i>	CACCTTCCCGCTGGACACT	CCCTAGGACACCTTTATACCTAATGG
<i>Ucp1-luciferase genotyping</i>	CATGAACGGCCACGAGTT	TGACGGCCATGTTGTTGT