Supplementary material



Supplementary Figure S1: CA inhibits the browning process of human white adipocytes. White hMADS adipocytes (A, D) were converted (days 14 to 18) into brite adipocytes with 100 nM rosiglitazone (B, E) or 300 nM GW7647 (C, F) in the presence of increasing CA concentrations. mRNA levels of thermogenic (A-C) and adipogenic (D-F) markers were analyzed. Paired student, p<0.05 considered as significant: #, white vs brite adipocyte; *, untreated *vs* CA treated condition.



Supplementary Figure S2: CA inhibits the browning process of mice primary adipocytes. SVF cells derived from mice BAT were induced to differentiate into white or brown (rosiglitazone-treatment) adipocytes in the absence (-) or presence (+) of 10 μ M of CA for the last 4 days and mRNA levels of thermogenic (A) and adipogenic (B) markers were analyzed. Histograms display mean ± SEM of three independent experiments.



Supplementary Figure S3: CA inhibits thermogenic marker gene expression of human brite adipocytes. hMADS adipocytes were differentiated into white adipocytes (A, D), or brite adipocytes using rosiglitazone (B, E) or GW7647 (C, F) in the presence of increasing CA concentrations and mRNA levels of thermogenic (A-C) and adipogenic (D-F) markers were analyzed. Paired student, p<0.05 considered as significant: #, white *vs* brite adipocyte; *, untreated *vs* CA treated condition.



Supplementary Figure S4: CA antagonizes rosiglitazone-induced activation of PPAR α . Concentrationdependent PPAR α transactivation activities of GW7647 were measured in the presence of varying amounts of CA using a PPAR α -LBD-GAL4 chimera assay. Values are expressed as % of the maximal response measured with GW7647 (100 nM). Data are displayed as mean ± SEM of two to three independent experiments (three replicates for each experiments). Paired student, p<0.05 considered as significant: #, GW7647 treated *vs* untreated; * and α , CA treated *vs* untreated.

Human oligonucleotide sequences			
name	Reverse primer	Forward primer	
PPARy2	ATCAGTGAAGGAATCGCTTTCTG	CAAACCCCTATTCCATGCTGTT	
PPARa	TCCAAAACGAATCGCGTTGT	GGCGAACGATTCGACTCAAG	
PGC1a	CTGTGTCACCACCCAAATCCTTAT	TGTGTCGAGAAAAGGACCTTGA	
UCP1	CCAGGATCCAAGTCGCAAGA	GTGTGCCCAACTGTGCAATG	
FABP4	CAACGTCCCTTGGCTTATGCT	TGTGCAGAAATGGGATGGAAA	
CPT1-M	GAGCAGCACCCCAATCAC	AACTCCATAGCCATCATCTGCT	
PLN1	GATGGGAACGCTGATGCTGTT	ACCCCCTGAAAAGATTGCTT	
PLN5	CTACGAGCACTCTGTGGGGA	GGTCTATCAGCTCCAGCGTCT	
36B4	TGCATCAGTACCCCATTCTATCAT	AGGCAGATGGATCAGCCAAGA	

Supplementary Table S1. Sequence of primers used for gene expression analysis.

Mice oligonucleotide sequences			
name	Reverse primer	Forward primer	
36B4	TCC AGG CTT TGG GCA TCA	CTTTATCAGCTGCACATCACTCAGA	
Cpt1m	GGCTCCAGGGTTCAGAAAGT	TGCCTTTACATCGTCTCCAA	
Fabp4	CTTGTGGAAGTCACGCCTTT	AAGAGAAAACGAGATGGTGACAA	
Ucp1	CACCTTCCCGCTGGACACT	CCTGGCCTTCACCTTGGAT	
Pln1	AGCGTGGAGAGTAAGGATGTC	CTTCTGGAAGCACTCACAGG	