

SUPPLEMENTARY MATERIALS

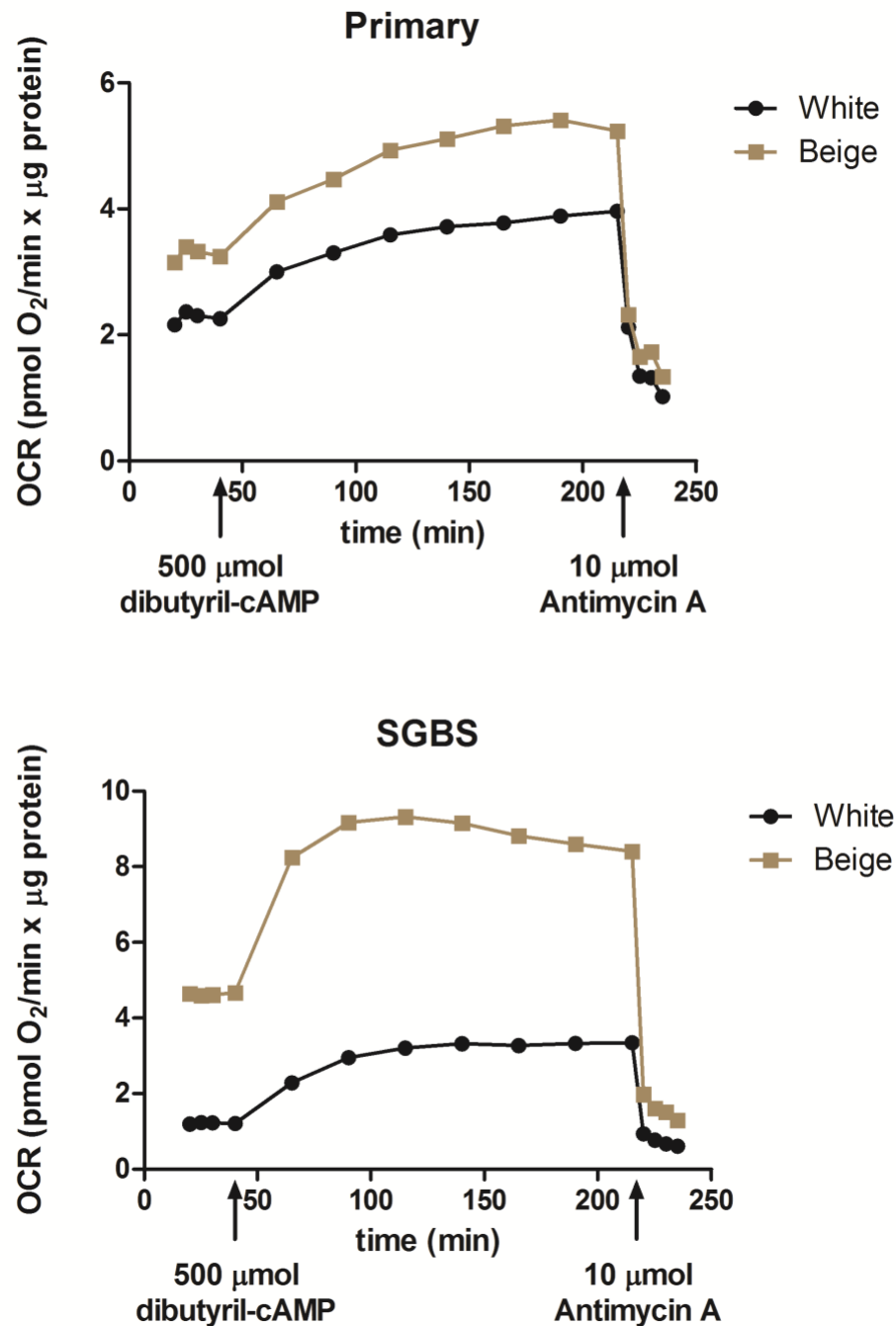
Thermogenic activation downregulates high mitophagy rate in human masked and mature beige adipocytes

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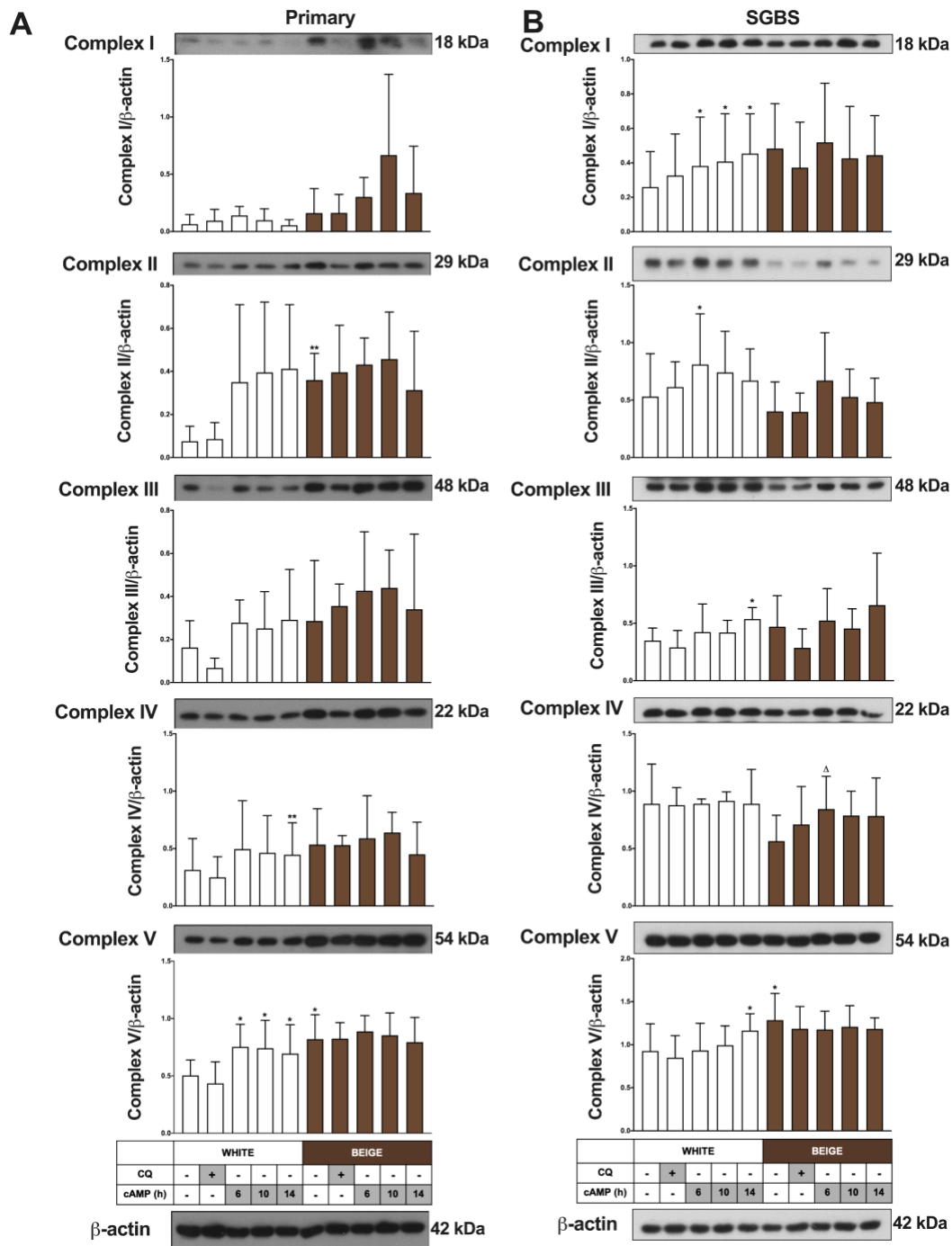
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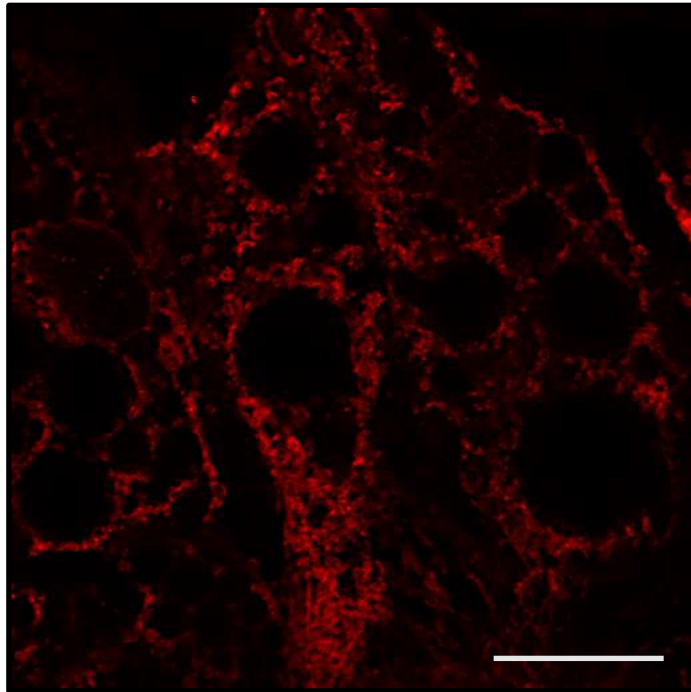


Supplementary figure 1. Oxygen consumption of one representative hASC-derived adipocyte donor and SGBS adipocyte sample measured by XF96 oximeter as described earlier [29,30]. After recording the baseline oxygen consumption, cells received a single bolus dose of dibutyryl-cAMP (500 μM final concentration) modelling adrenergic stimulation. Then, stimulated oxygen consumption was recorded every 30 min. The oxygen consumption rate was normalized to protein content and normalized readings were displayed.

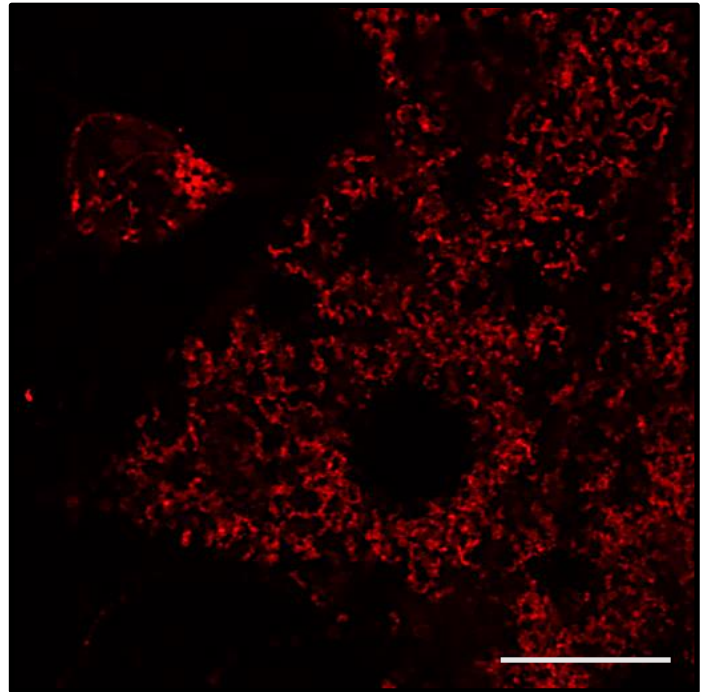


Supplementary figure 2. Expression of mitochondrial complex subunits in adipocytes upon cAMP treatment. Adipocytes were differentiated and treated as in Figure 1. Representative immunoblots and densitometry analysis of mitochondrial complex subunits protein expression (A) in hASC-derived (n=4) and (B) in SGBS cells (n=5) normalized to β -actin. Results are expressed as mean \pm SD. Statistics: two-tailed paired student t-test, *p<0.05, **p<0.01, *: significant as compared to white untreated sample, Δ p<0.05, $\Delta\Delta$ p<0.01, Δ : significant as compared to beige untreated sample.

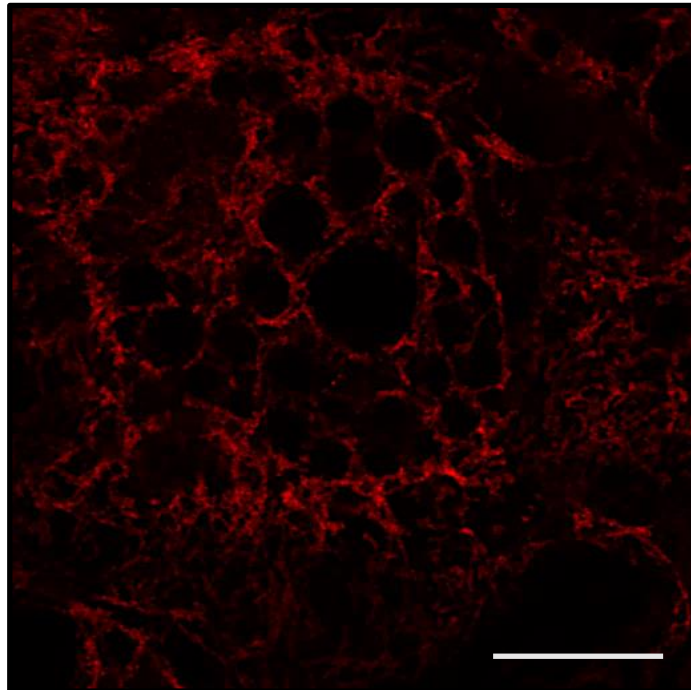
Primary White Untreated



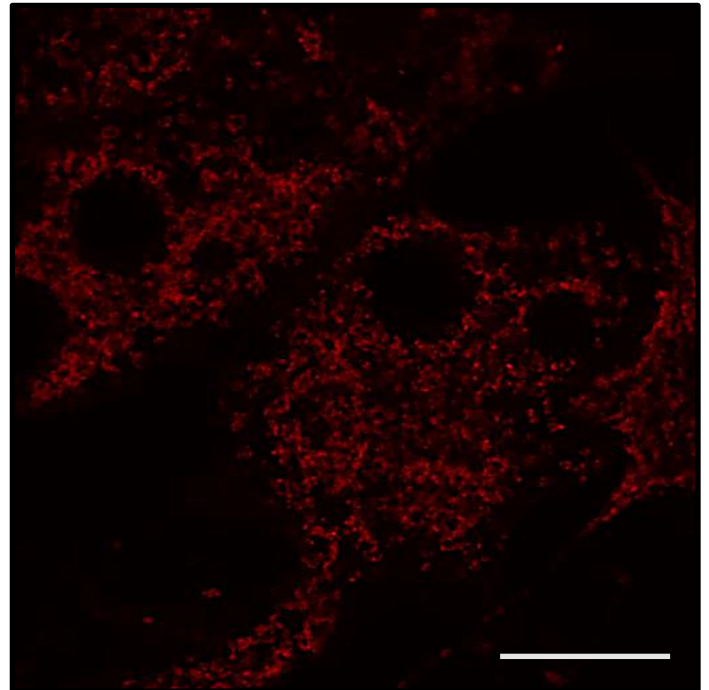
Primary White cAMP treated



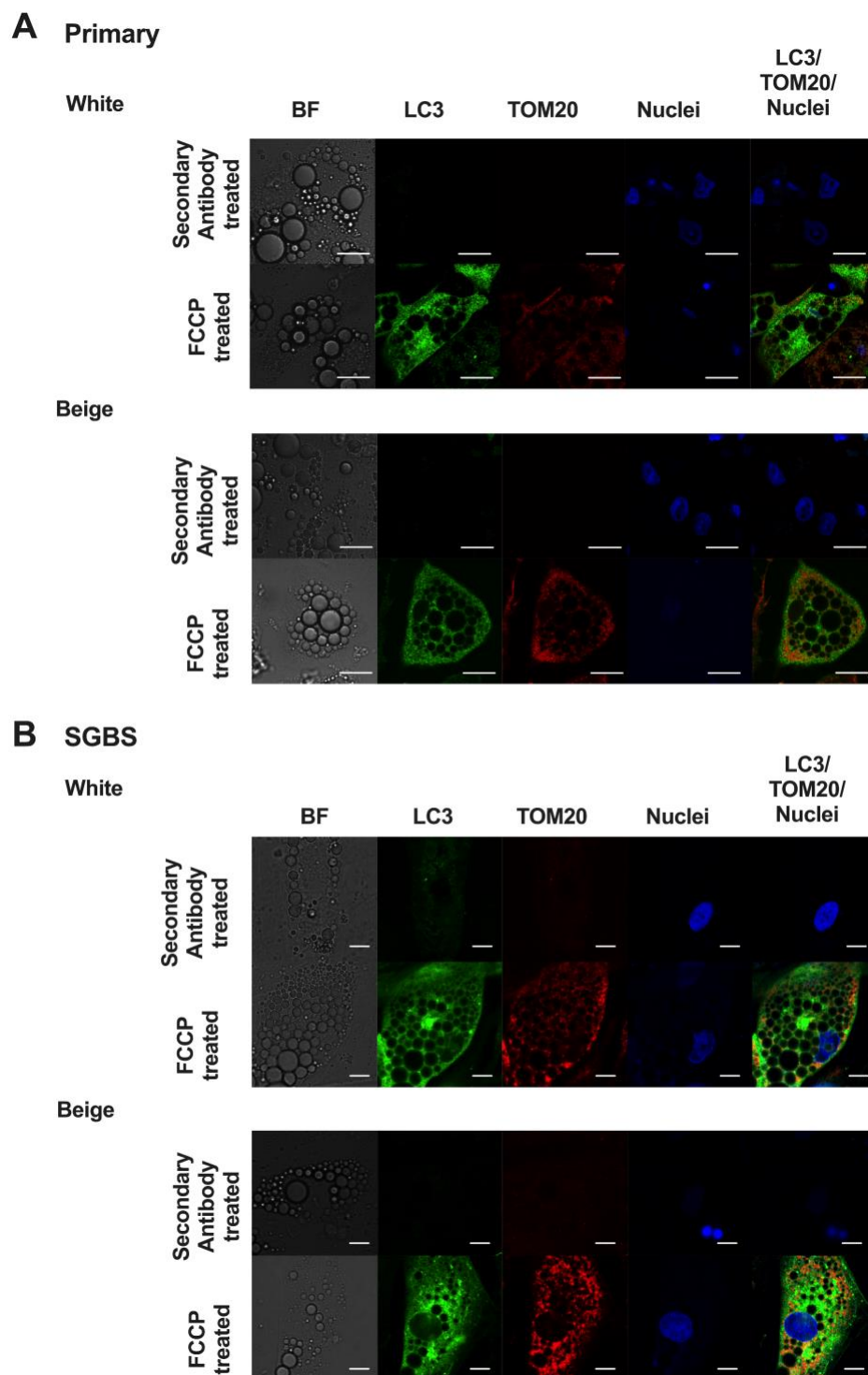
Primary Beige Untreated



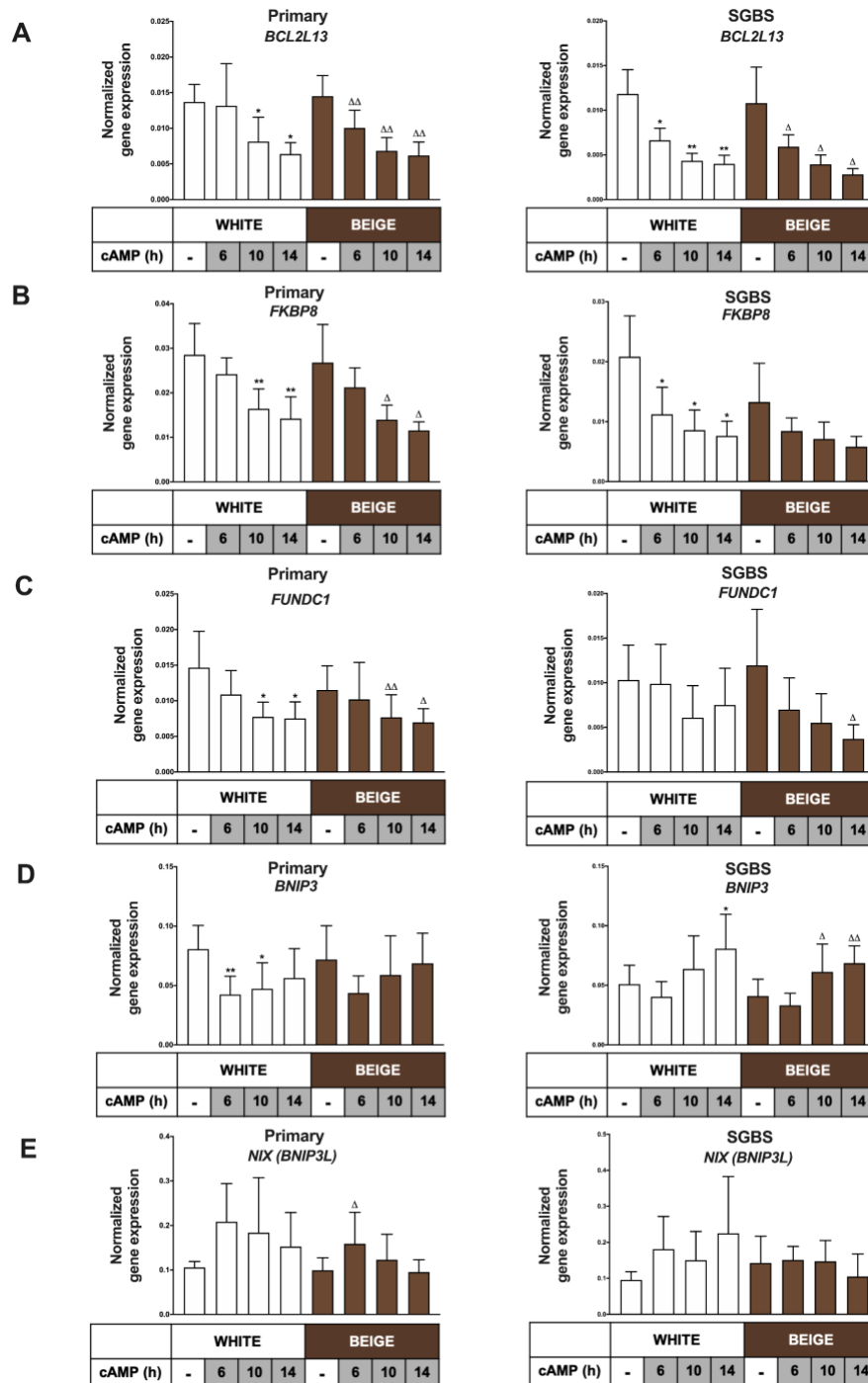
Primary Beige cAMP treated



Supplementary figure 3. High-resolution confocal microscopy images of TOM20 immunostaining. Scale bars represent 5 μ m.



Supplementary figure 4. Confocal images of LC3 and TOM20 immunostaining in control and FCCP treated samples. Representative confocal microscopy images of secondary antibody controls for LC3 and TOM20 immunostaining in white (up) and beige (down) adipocytes and upon FCCP treatment (10 μ M, 6 h) (A) in primary and (B) in SGBS cells.



Supplementary figure 5. Expression of genes involved in Parkin-independent mitophagy is decreased as a result of cAMP stimulus. Adipocytes were differentiated and treated as in Figures 1 and 2. Gene expression of (A) *BCL2L13*, (B) *FKBP8*, (C) *FUNDC1*, (D) *BNIP3*, (E) *NIX* (*BNIP3L*) in hASC-derived (left) and SGBS cells (right) (n=5). Gene expression was determined by RT-qPCR and target genes were normalized to GAPDH. Results are expressed as mean±SD. Statistics: two-tailed paired student t-test, *p<0.05, **p<0.01, *: significant as compared to white untreated sample, Δp<0.05, ΔΔp<0.01, Δ: significant as compared to beige untreated sample.