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## Supplementary Materials for

### **Molecular signatures and functional analysis of beige adipocytes induced from *in vivo* intra-abdominal adipocytes**

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#### **The PDF file includes:**

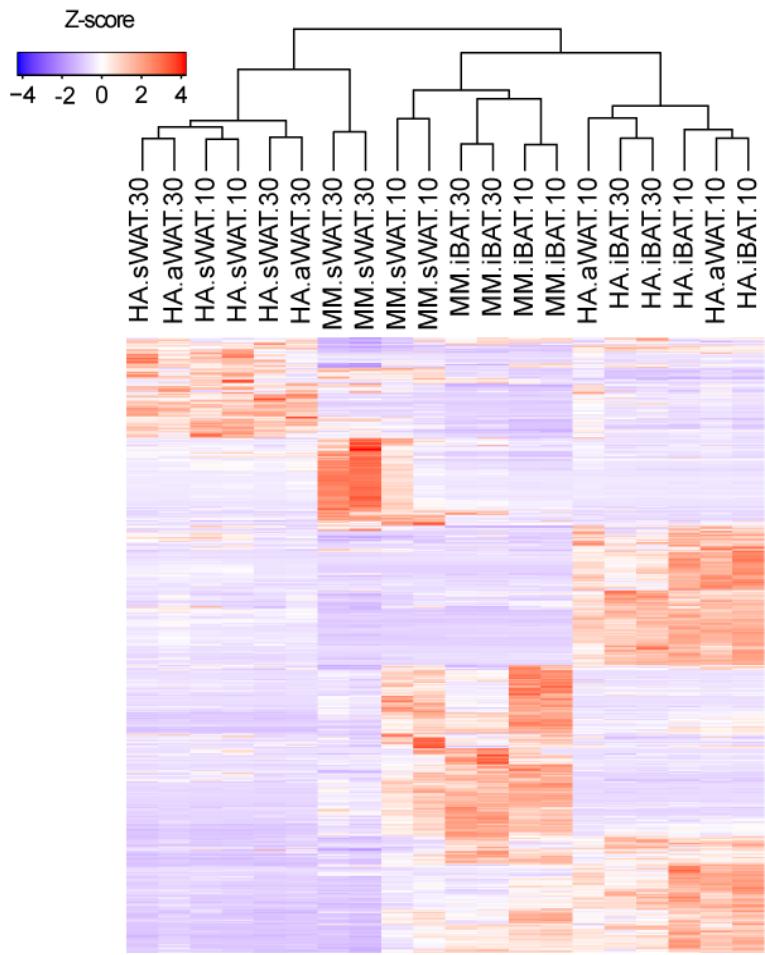
- Fig. S1. Clustered profile of all marker genes.
- Fig. S2. Expression levels of six beige marker genes assayed by RT-qPCR.
- Fig. S3. Ponceau staining as loading control in Western blots.
- Legends for Tables S1 and S2
- Table S3. Primers used for qPCR.
- References (38, 39)

#### **Other Supplementary Material for this manuscript includes the following:**

(available at [advances.sciencemag.org/cgi/content/full/4/7/eaar5319/DC1](https://advances.sciencemag.org/cgi/content/full/4/7/eaar5319/DC1))

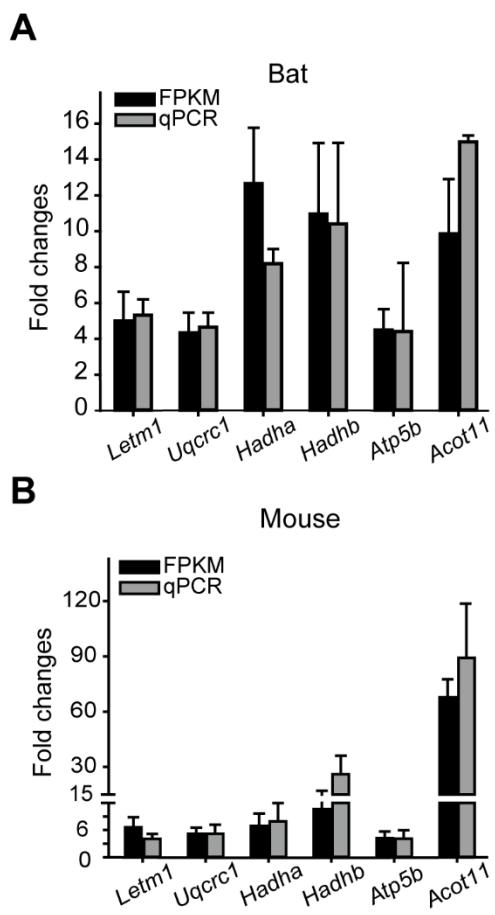
Table S1 (Microsoft Excel format). Mapped Ensembl genes and raw read counts from 20 samples.

Table S2 (Microsoft Excel format). All beige adipocyte, BAT, and WAT marker genes.

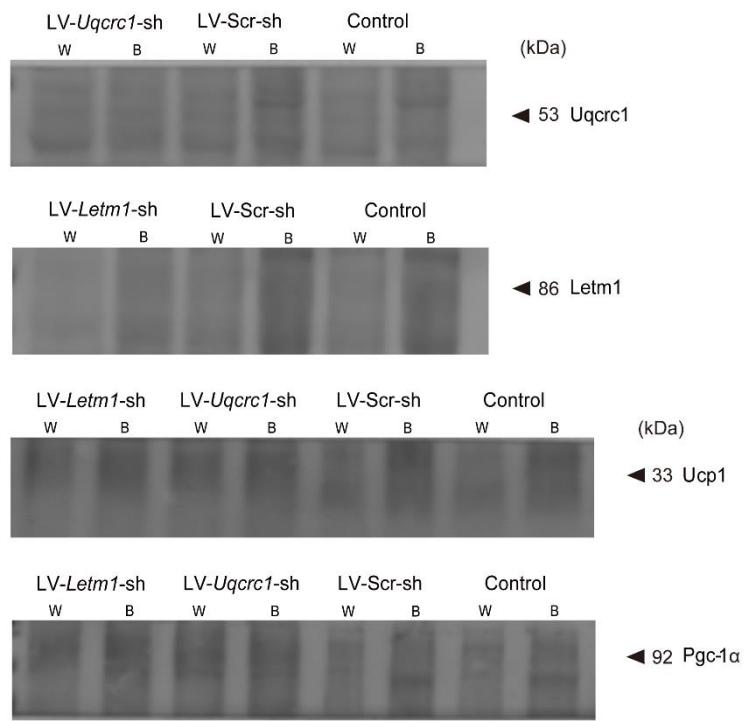


**Fig. S1. Clustered profile of all marker genes.**

HA: *Hipposideros armiger*; MM: *Mus musculus*.



**Fig. S2. Expression levels of six beige marker genes assayed by RT-qPCR. (A) Bat markers. (B) Mouse markers. n = 3 for all groups.**



**Fig. S3. Ponceau staining as loading control in Western blots.** W and B represents white and beige adipocytes respectively. n = 3 for all groups.

**Table S1. Mapped Ensembl genes and raw read counts from 20 samples.** Please see the Excel file.

**Table S2. All beige adipocyte, BAT, and WAT marker genes.** Please see the Excel file.

**Table S3. Primers used for qPCR.**

Gene	Forward primer (5' → 3')	Reverse primer (5' → 3')
<i>Letm1</i>	GTGAAGGACATCCAGCCAGAAG	CCTTCCAGTACAGGTGCCGTAT
<i>Uqerc1</i>	CAAGGGAACAAAGAACATCGGC	ACAGTGCTTGATGAGGTAAGC
<i>Pparγ</i> *	GTGCCAGTTTCGATCCGTAGA	GGCCAGCATCGTGTAGATGA
<i>Ucp1</i> *	CTGCCAGGACAGTACCCAAG	TCAGCTGTTCAAAGCACACA
<i>Pgc-1α</i> *	CCCTGCCATTGTTAACGACC	TGCTGCTGTTCCCTGTTTC
<i>Tbp</i> *	GCTGTAAACTTGACCTAAAGACCAT	AACGCAGTTGTCCGTGGCTCT
<i>Fabp4</i> *	AAGGTGAAGAGCATCATAACCCCT	TCACGCCTTCATAACACATTCC
<i>Cytb</i>	CGCAGTCATAGCCA CAGCATT	CTTGACCCGATTCTCGCTT
<i>Cebp/β</i> *	GCAAGAGCCGCGACAAG	GGCTCGGGCAGCTGCTT
<i>Tbx1</i> *	GGCAGGCAGACGAATGTT	TTGTCATCTACGGGCACAAAG
<i>Prdm16</i> *	ACCTGCCACAGCAAAGAA	CCATCCAAGCAGAGAAGTAGAC
<i>Cd137</i> *	GAGGTAGAAGAGAAAGGGTTG	GTAGAGGACCCAGGTTGATT
<i>Cited1</i> *	AACCTGGAGTGAAGGATCGC	GTAGGAGAGCCTATTGGAGATGT

\*Primers for mice were from previous publications (2, 10, 38, 39).