

Mice lacking PLAP-1/aspurin counteracts high fat diet-induced metabolic disorder and alveolar bone loss by controlling adipose tissue expansion

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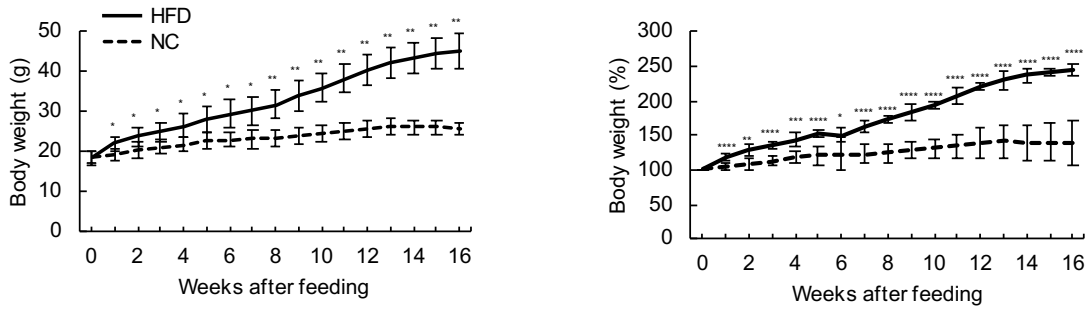
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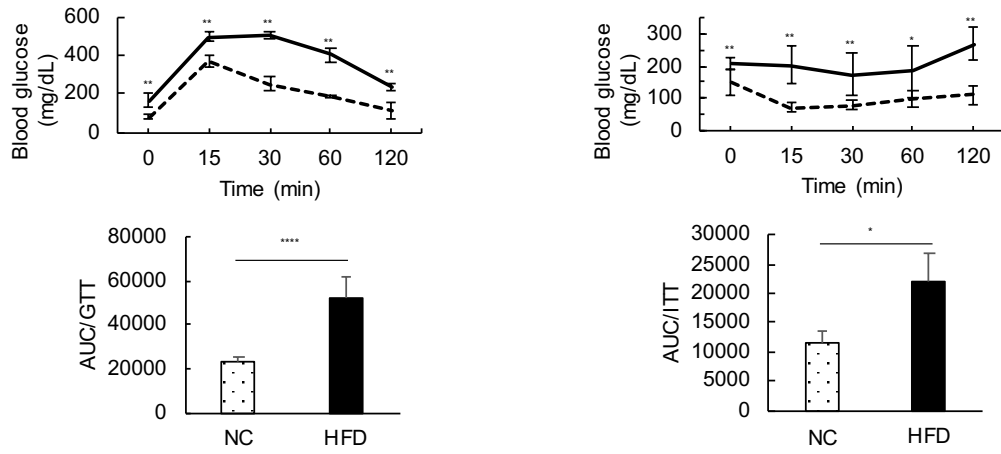
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Fig S1

A



B



C

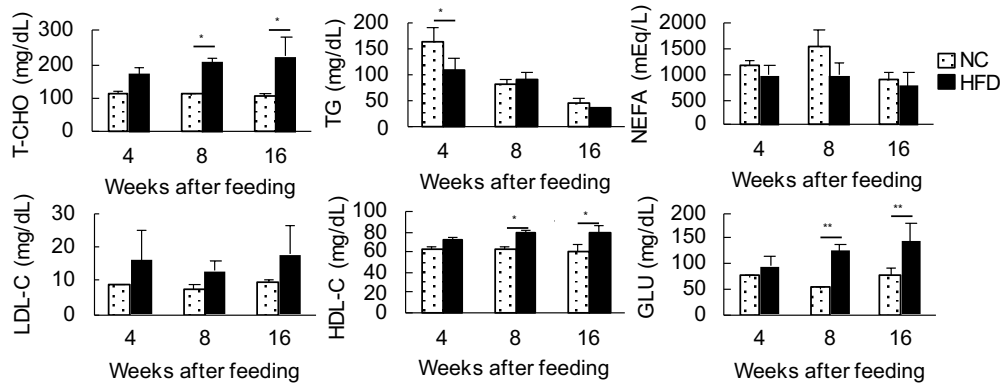


Fig S1. HFD-induced obesity and metabolic abnormalities.

(A) 5-week-old male C57BL/6J mice were fed with high fat diet (HFD) or normal chow diet (NC) and weighted weekly (n = 7 in each group). (B) Glucose tolerance test (GTT) and insulin tolerance test (ITT) were performed in mice after NC or HFD feeding (n = 4 in each group). (C) Serum levels of total cholesterol (T-CHO), triglyceride (TG), LDL-cholesterol (LDL-C), HDL-cholesterol (HDL-C), non-esterified fatty acid (NEFA), glucose (GLU) were measured during NC or HFD feeding (n = 3 in each group). *: $p < 0.05$, **: $p < 0.01$, ****: $p < 0.0001$.

Fig S2

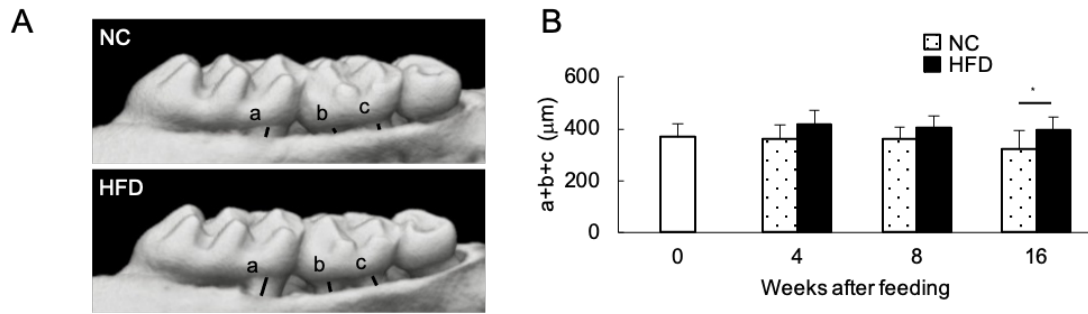


Fig S2. HFD-induced alveolar bone resorption.

(A) Alveolar bone resorption in mice fed with NC or HFD was evaluated by μ CT. (B) Distance between alveolar bone crest and cement-enamel junction was measured at distal root of first molar (a), mesial (b) and distal (c) root of second molar (n = 14 in each group). Results show the mean \pm SD. *: $p < 0.05$

Fig S3

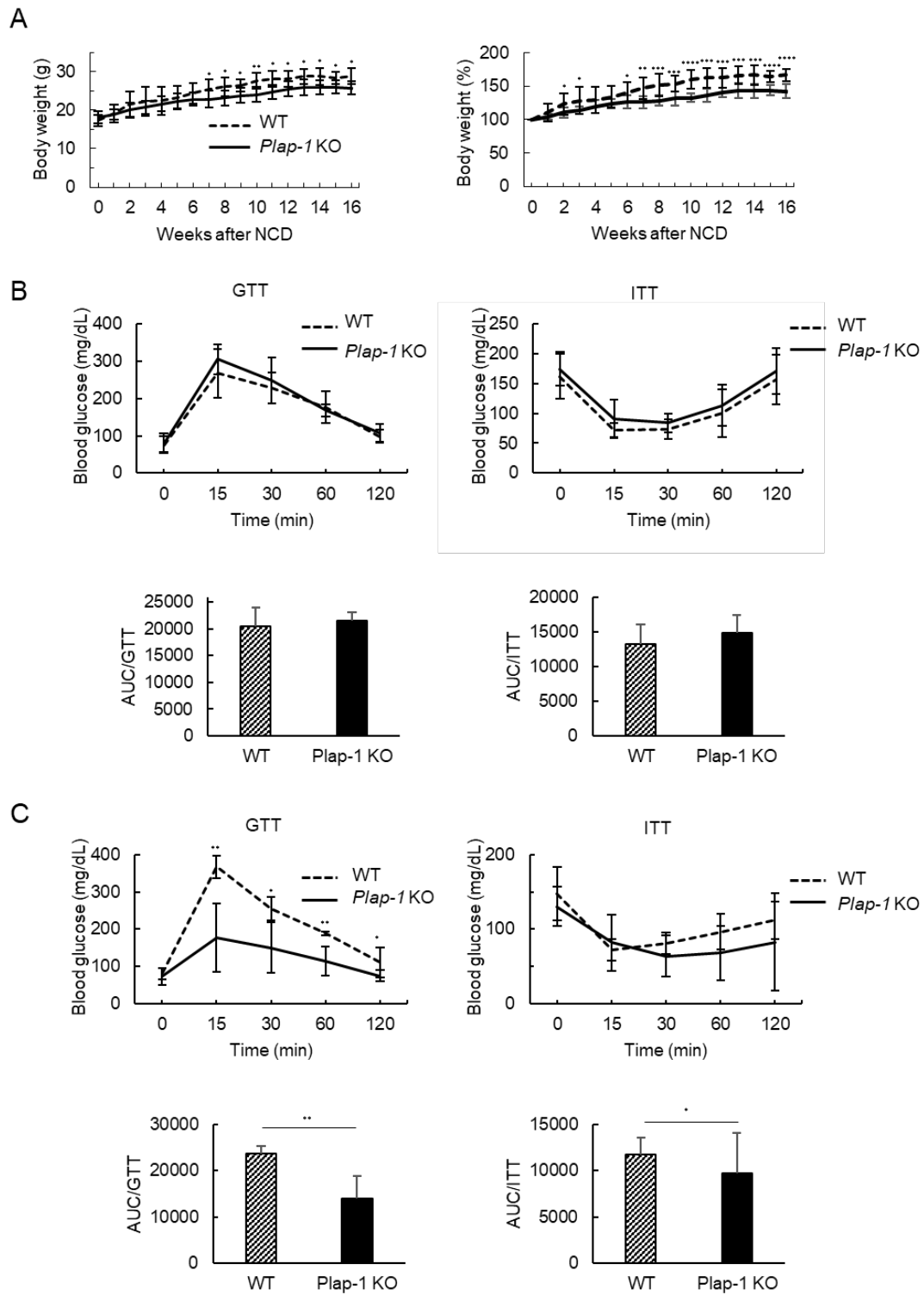


Fig S3. Body weight change and glucose homeostasis in *Plap-1* KO mice with NC feeding.

(A) Body weight changes in WT and *Plap-1* KO mice during NC feeding. 5-week-old male WT and *Plap-1* KO mice were fed with NC and weighted weekly. WT ($n = 11$), *Plap-1* KO ($n = 8$). Results show the mean \pm SD. (B) GTT and ITT were performed in 5-week-old male WT and *Plap-1* KO mice ($n = 9$ in each group). (C) GTT and ITT were performed in WT and *Plap-1* KO mice after 16 weeks NC feeding. For GTT, WT ($n = 8$), *Plap-1* KO ($n = 7$) and for ITT, WT ($n = 8$), *Plap-1* KO ($n = 8$). Results show the mean \pm SD. *: $p < 0.05$, **: $p < 0.01$, ***: $p < 0.001$, ****: $p < 0.0001$.

Fig S4

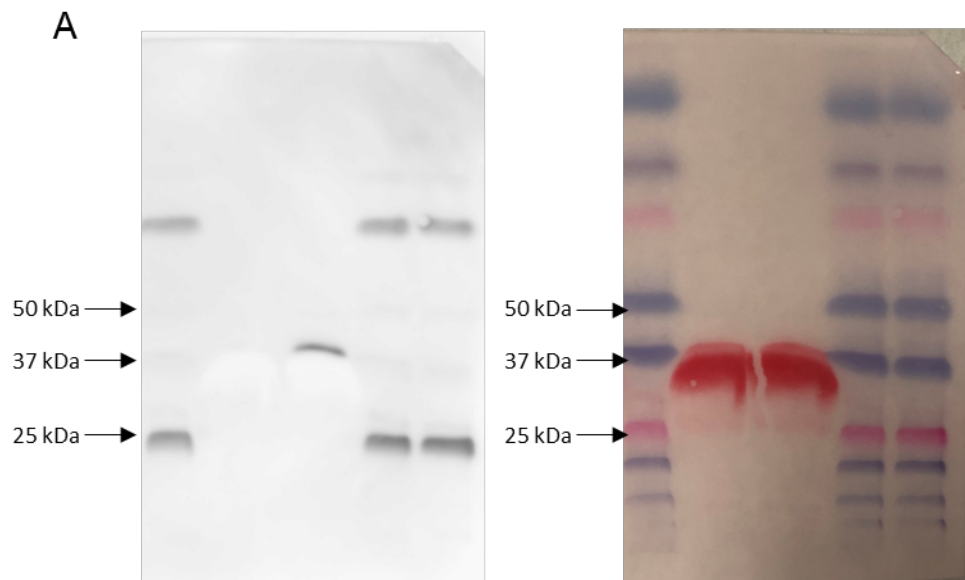


Fig S3. Immunoblots performed in this study

(A) anti-FLAG blot (left) and Ponceau S staining (right).

Table S1

Gene of interest		Forward primers sequences	Reverse primers sequences
<i>Aspn</i>	NM_025711	atgatgacgataacgatgatgacga	tgttgtttggaaccgatgacaga
<i>Adipoq</i>	NM_009605	ttctgtctgtacgattgtcagtggga	ggcatgactgggcaggatta
<i>Lep</i>	NM_008493	gaacctgtctactcatgccagcac	ctggctctgcagcctgtttg
<i>Adgre1</i>	NM_010130	gctatgggcttccagtctggtggt	gaaggcagcaacctcgtgccttg
<i>Tnf</i>	NM_013693	aagcctgtagccacgtcgta	ggcaccactagttggttgcctttg
<i>Il6</i>	NM_031168	ccacttcacaagtcggaggctta	gcaagtgcacatcatcgttgcatac
<i>Ccl3</i>	NM_011337	catgacactctgcaaccaagtcttc	gagcaaaggctgttggttca
<i>Arg1</i>	NM_007482	agctctgggaatctgcatgg	atgtacacgatgtctttggcagata
<i>Chil3</i>	NM_009892	gatggcctcaacctggactg	cgtaaatgattcctgctcctg
<i>Clec7a</i>	NM_001309637	ggtaccctgactcaaattcacaac	caccacagtaagtaggctgattctc
<i>Fabp4</i>	NM_024406	tgggaacctggaagcttgtctc	gaattccacgcccagtttga
<i>Pparg</i>	NM_011146	ggagcctaagttgagtttgcctgtg	tgcagcaggttgccttggatg
<i>Cebpa</i>	NM_007678	ttgaagcacaatcgatccatcc	gcacactgccattgcacaag
<i>Coll1a1</i>	NM_007742	cagggtattgctggacaacgtg	ggacctgtttgccaggttca
<i>Col3a1</i>	NM_009930	caggccagtggcaatgtaaaga	ctcattgccttgcgtgtttgata-
<i>Col6a1</i>	NM_009933	gagcttctcgtatgacggcttcc	gagctgtctagcaggatggtgatg
<i>Dcn</i>	NM_007833	ctgggctggcacagcataagta	cggacaggggtgccgtaaag
<i>Bgn</i>	NM_007542	gatgattgagaatgggagcctga	tccgaagcccataggacagaag
<i>Hprt</i>	NM_013556	ttgttgggatatgcccttgacta	aggcagatggccacaggacta