

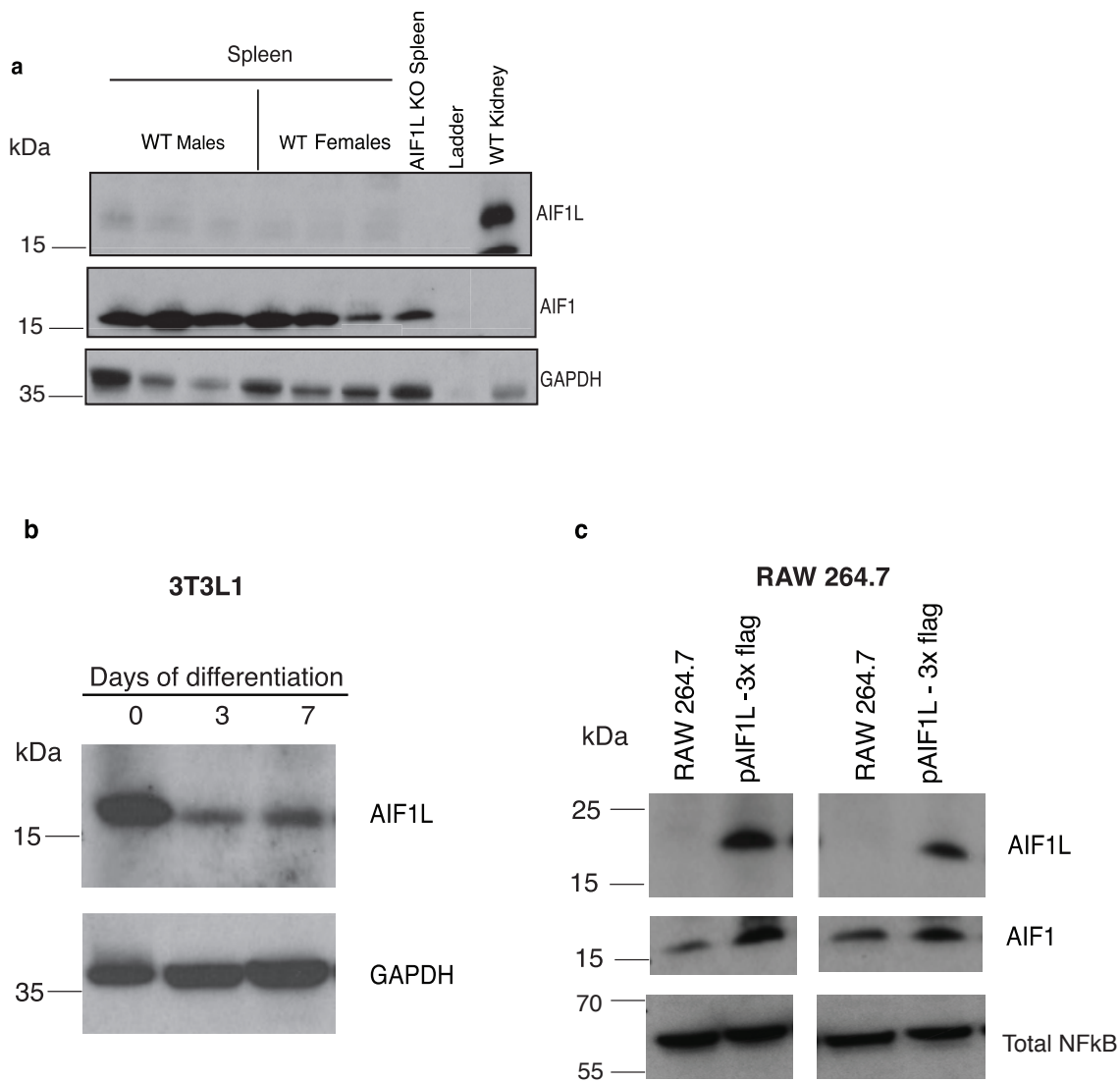
**Allograft inflammatory factor-1-like is not essential for age dependent weight gain or HFD-induced obesity and glucose insensitivity**

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<sup>1</sup>Albert Einstein College of Medicine, Wilf Family Cardiovascular Research Institute, Department of Medicine (Cardiology), and Department of Developmental and Molecular Biology. 1300 Morris Park Avenue, Bronx, New York, 10461, USA.

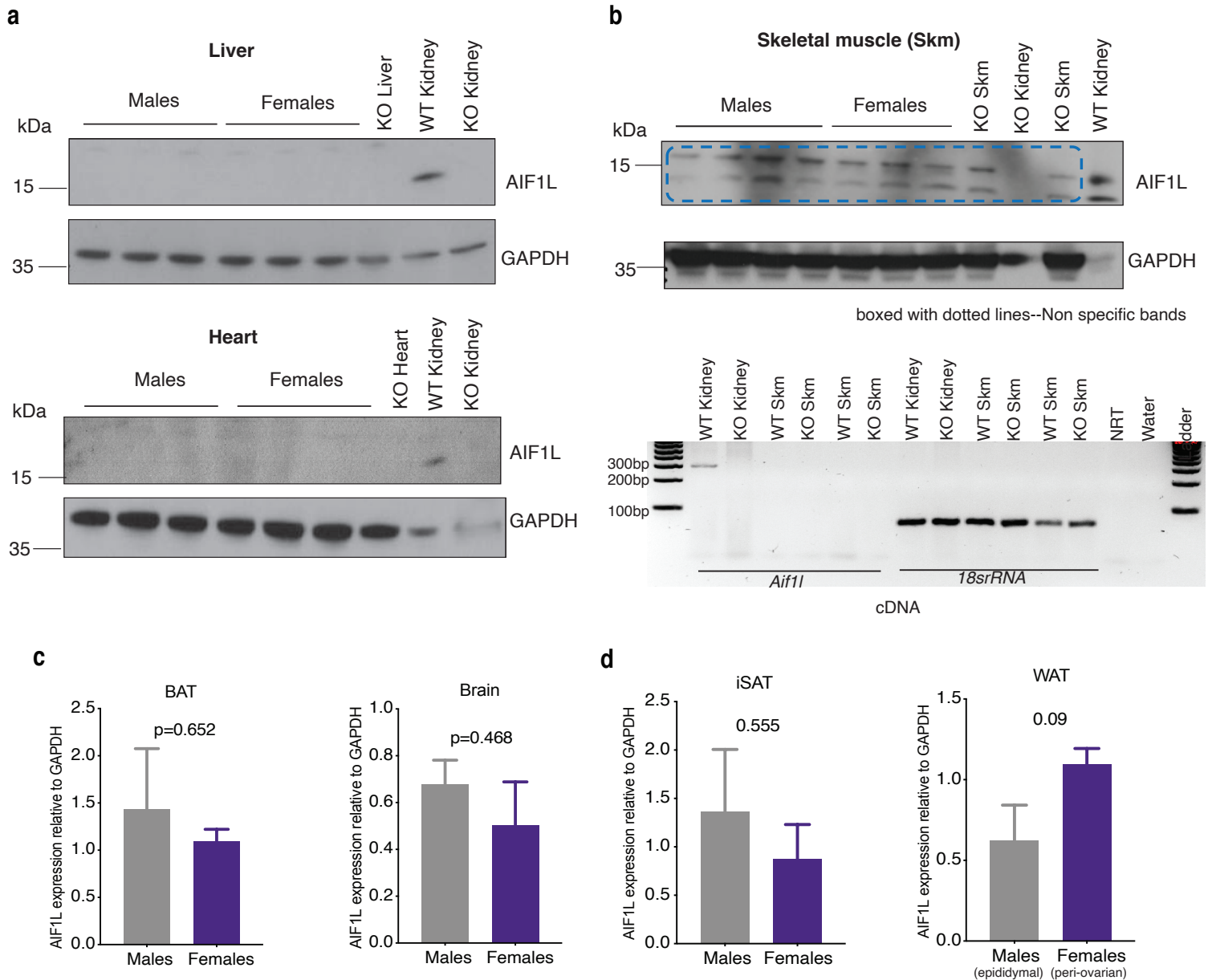
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Correspondence to Nicholas Sibinga, Albert Einstein College of Medicine, Forchheimer G46, 1300 Morris Park Avenue, Bronx, New York 10461, USA; phone 718-430-2881; fax 718-430-8989; e-mail: [nicholas.sibinga@einstein.yu.edu](mailto:nicholas.sibinga@einstein.yu.edu)



**Supplementary Figure 1. Characterization of AIF1L antibody and its expression in cellular model of adipocytes**

- Immunoblot for AIF1L and AIF1 in spleen and kidney lysates from WT adult mice with anti AIF1L antibody (40 $\mu$ g of total protein loaded)
- Immunoblot for AIF1L in 3T3L1 cells during induced adipogenic differentiation with anti AIF1L antibody (30 $\mu$ g of total protein loaded)
- Immunoblot for FLAG-tagged AIF1L and endogenous AIF1 in RAW 264.7 cells with anti AIF1L and anti AIF1 antibodies respectively



### Supplementary Figure 2. Characterization of AIF1L expression in mouse tissues

- Undetectable AIF1L expression in liver and heart
- Undetectable AIF1L protein and mRNA expression in skeletal muscle  
(immunoblotting for AIF1L using skeletal muscle lysates from both WT and AIF1L KO mice showed bands of ~14 and 17 kDa that are most likely non-specific. The global KO strategy used in this case is effective, as confirmed by the loss of signal for AIF1L in kidney, where the protein is highly expressed in WT animals — suggesting that these bands in the skeletal muscle samples are not due to AIF1L. RT-PCR analysis of skeletal muscle RNA using primers spanning exons 1 and 5 yielded the expected ~300 bp band in WT samples, but no products with KO, including potential transcripts that might arise from exon skipping events; in any case, such events in the recombined *Aif1l* allele would be expected to yield smaller products (as discussed in Methods), and would not explain the 17 kDa band.)
- Quantification of AIF1L expression in BAT and Brain
- Quantification of AIF1L expression in white adipose tissues

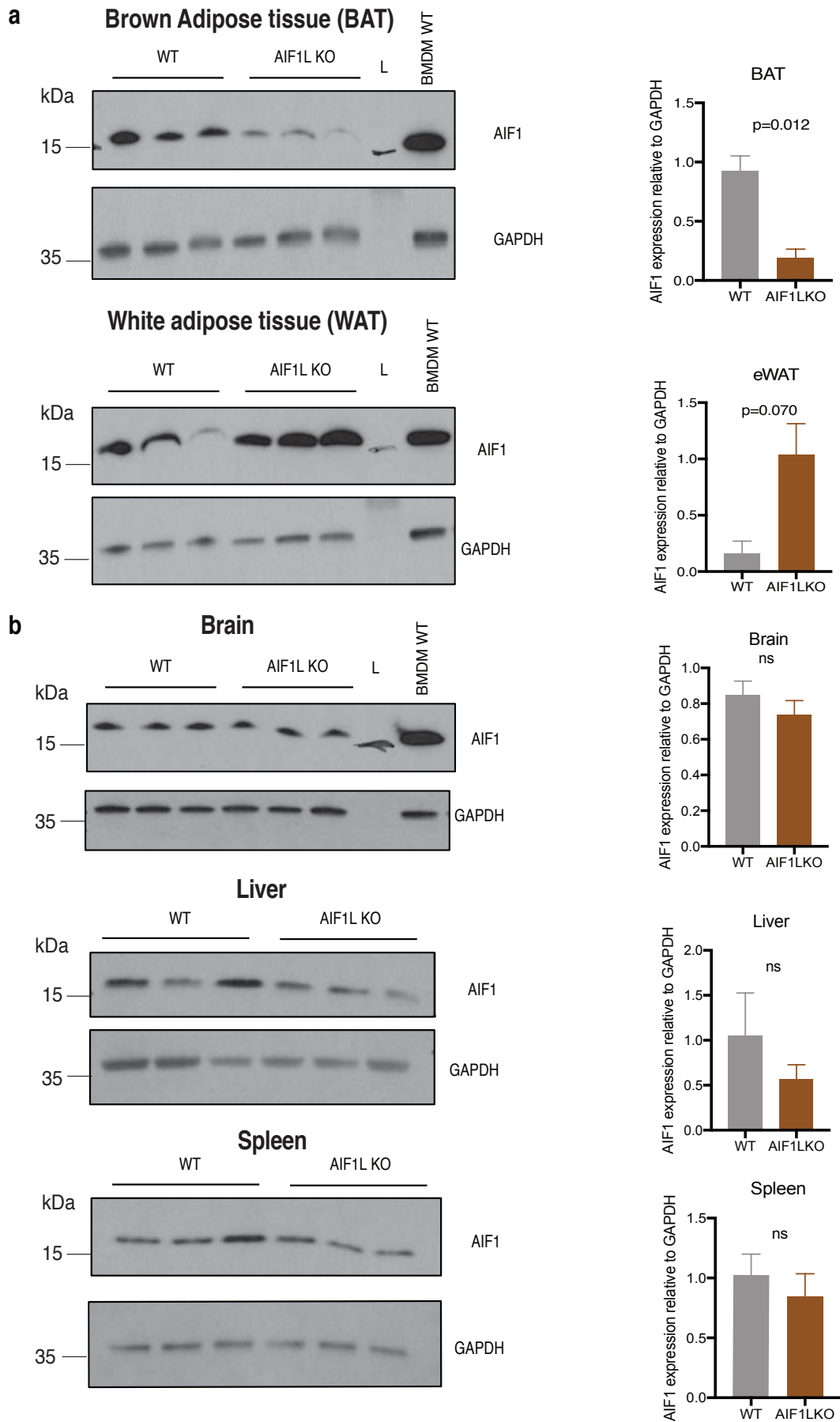
Data are represented as mean  $\pm$  SEM. ns - not significant.



### Supplementary Figure 3 Modified *Aif1* locus, tail DNA genotyping, and validation of loss of full length AIF1L

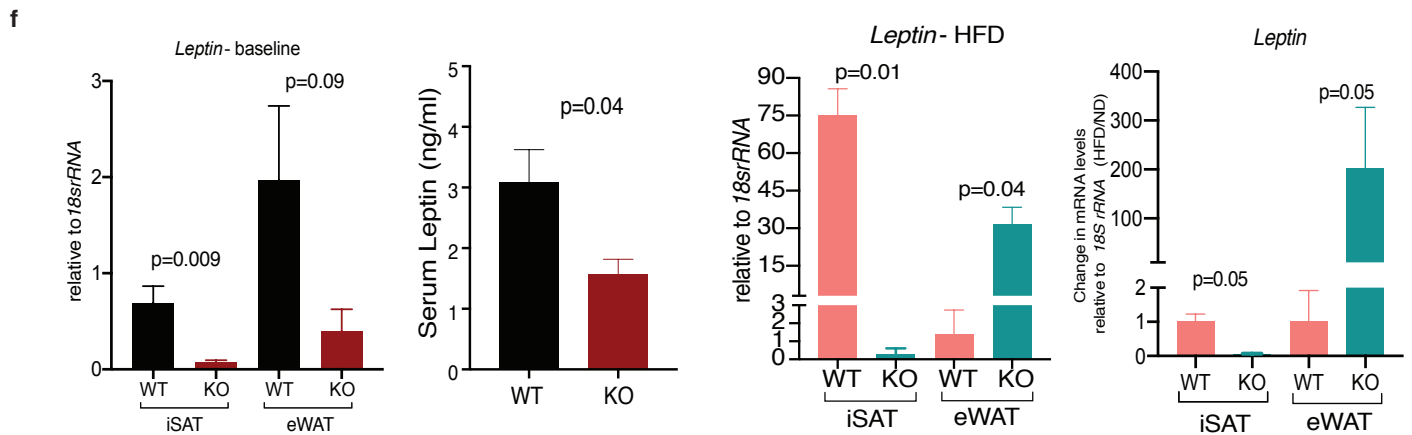
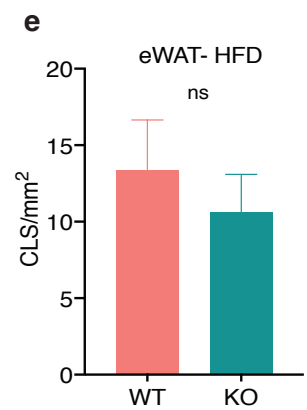
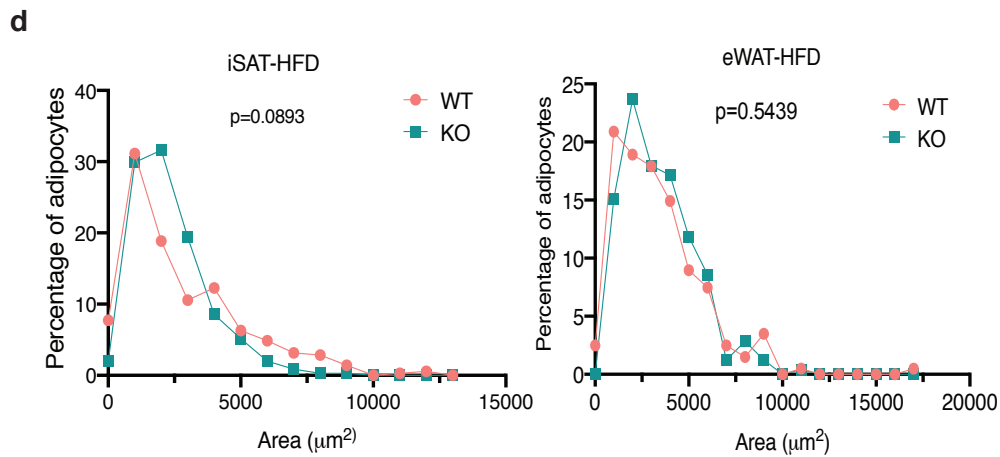
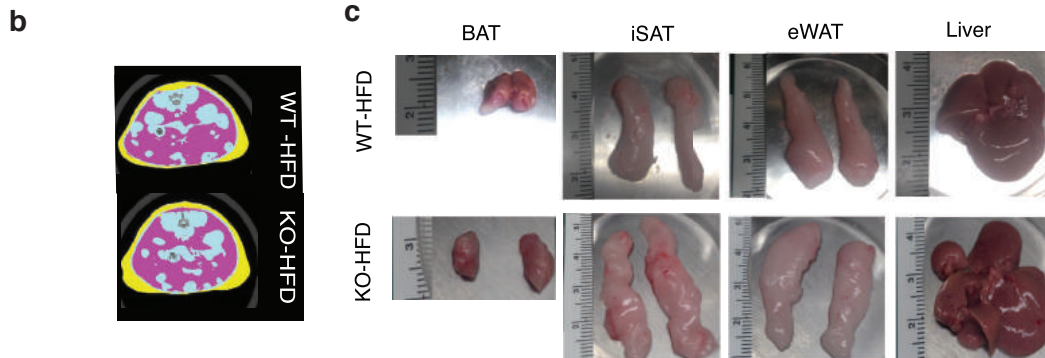
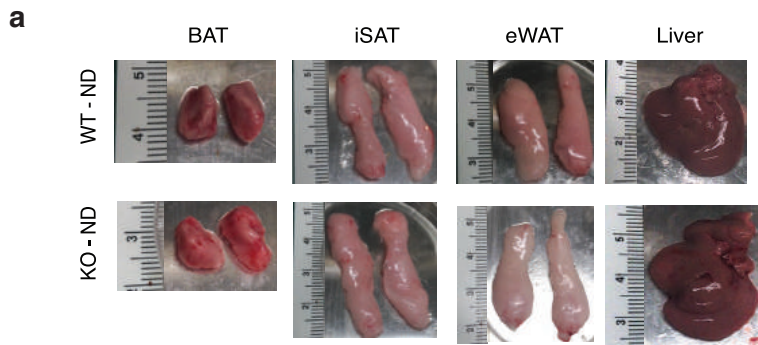
- a) As described by Wellcome Trust Sanger Institute (WTSI) -- The L1L2\_Bact\_P cassette was inserted at position 31961760 of Chromosome 2 upstream of the critical exon(s) (Build GRCm38). The cassette is composed of an FRT site followed by *lacZ* sequence and a *loxP* site. This first *loxP* site is followed by a neomycin resistance gene under the control of the human beta-actin promoter, SV40 polyadenylation cassette, a second FRT site and a second *loxP* site. A third *loxP* site is inserted downstream of the targeted exon(s) at position 31962560. The critical exon is thus flanked by *loxP* sites.
- b) Cre mediated recombination removes the neomycin cassette and exon 3, leaving one FRT site and *lacZ*, which is transcribed under the endogenous promoter of *Aif1*.
- c) Tail DNA genotyping from mice WT, heterozygous, or homozygous for the modified *Aif1* allele.
- d) End point PCR of *Aif1* cDNA from WT (n = 2) and KO kidney (n = 2) with forward and reverse primers spanning exon 1 and exon 5, respectively.
- e) End point PCR of *Aif1* cDNA from WT (n = 2) and KO kidney (n = 2) with forward and reverse primers spanning exon 1 and exon 2, respectively.
- f) Immunohistochemical (IHC) staining for AIF1L expression in kidney sections from 8-9-week-old WT and AIF1L KO male mice.

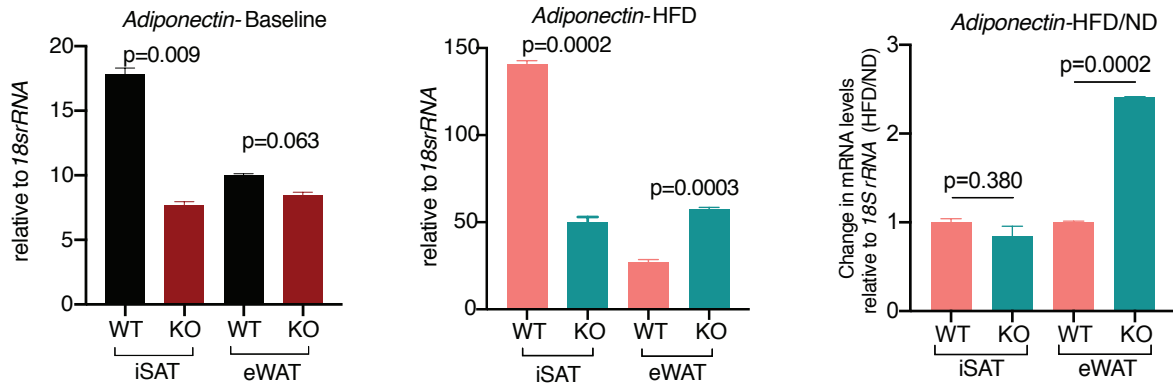
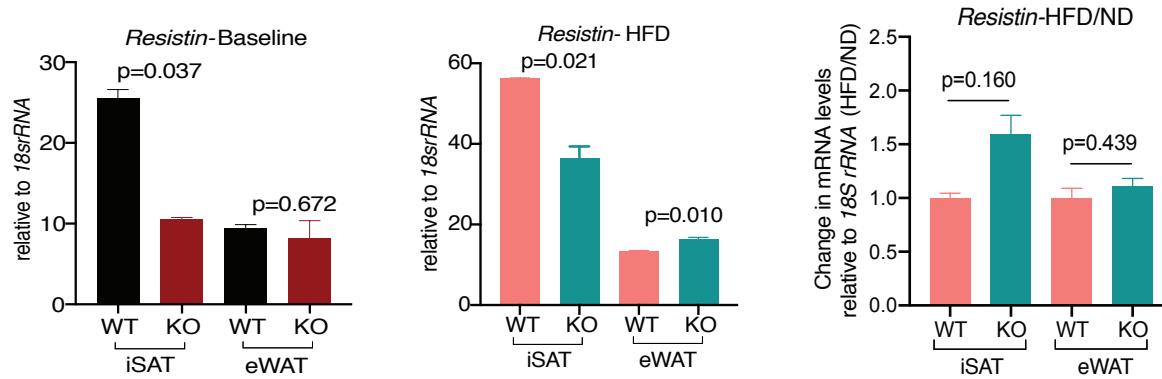
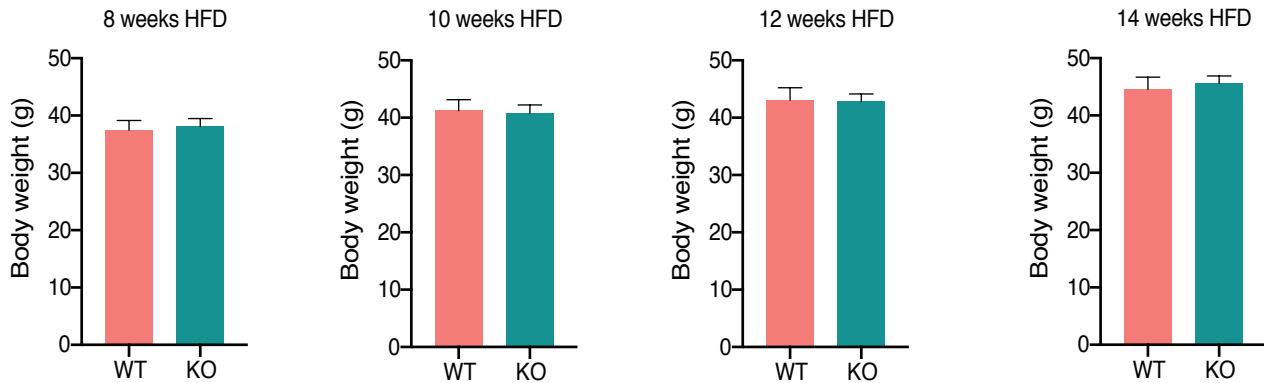
Abbreviations for genotyping PCR- Mut- presence of modified allele, *lacZ*- presence of *lacZ* cassette, Cre-presence of Cre, WT-presence of Wild-type allele, Ex3- presence of exon 3 in the modified allele.



**Supplementary Figure 4. AIF1 expression in tissues from WT and AIF1L KO 8-10 week old male mice**

- a) AIF1 expression in brown and epididymal white adipose depots (n=3)  
 b) AIF1 expression in brain, liver, and spleen (n=3)

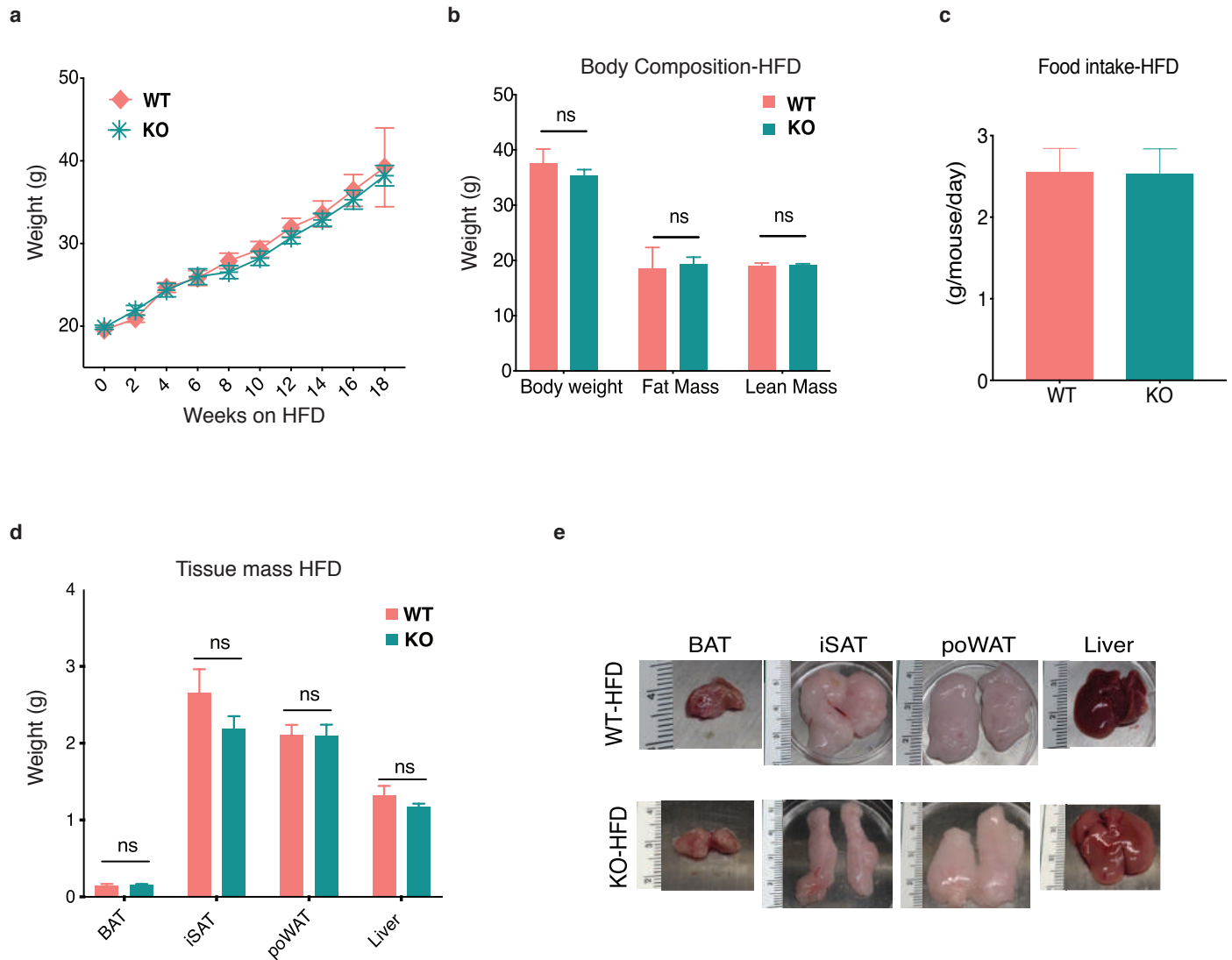


**g****h****i**

**Supplementary Figure 5 Macroscopic appearance of harvested organs from ND- and HFD-fed mice, histological analysis of eWAT sections and adipokine expression changes**

- Macroscopic appearance of tissues harvested from mice on ND for 18 weeks - representative images.
- $\mu$ CT scans (transverse plane) of WT and KO mice fed HFD for 18 weeks- representative images
- Gross appearance of tissues harvested from mice on HFD for 18 weeks - representative images.
- Frequency distribution of adipocyte size in iSAT and eWAT sections of WT and KO male mice fed HFD for 18 weeks (n = 3)
- Crown like structures quantification in eWATs from WT (n=9) and KO (n=6) mice fed HFD for 18 weeks
- Leptin* mRNA expression in adipose depots from male mice at baseline and upon long term HFD feeding and serum leptin levels at baseline (n=5 for WT and n=3 for KO)
- g,h) *Adiponectin* and *resistin* mRNA expression in adipose depots from male mice at baseline and upon long term HFD feeding (n=3)
- i) Whole body weight after 8, 10, 12, and 14 weeks of HFD feeding (n=11 for WT and n=8 for KO)



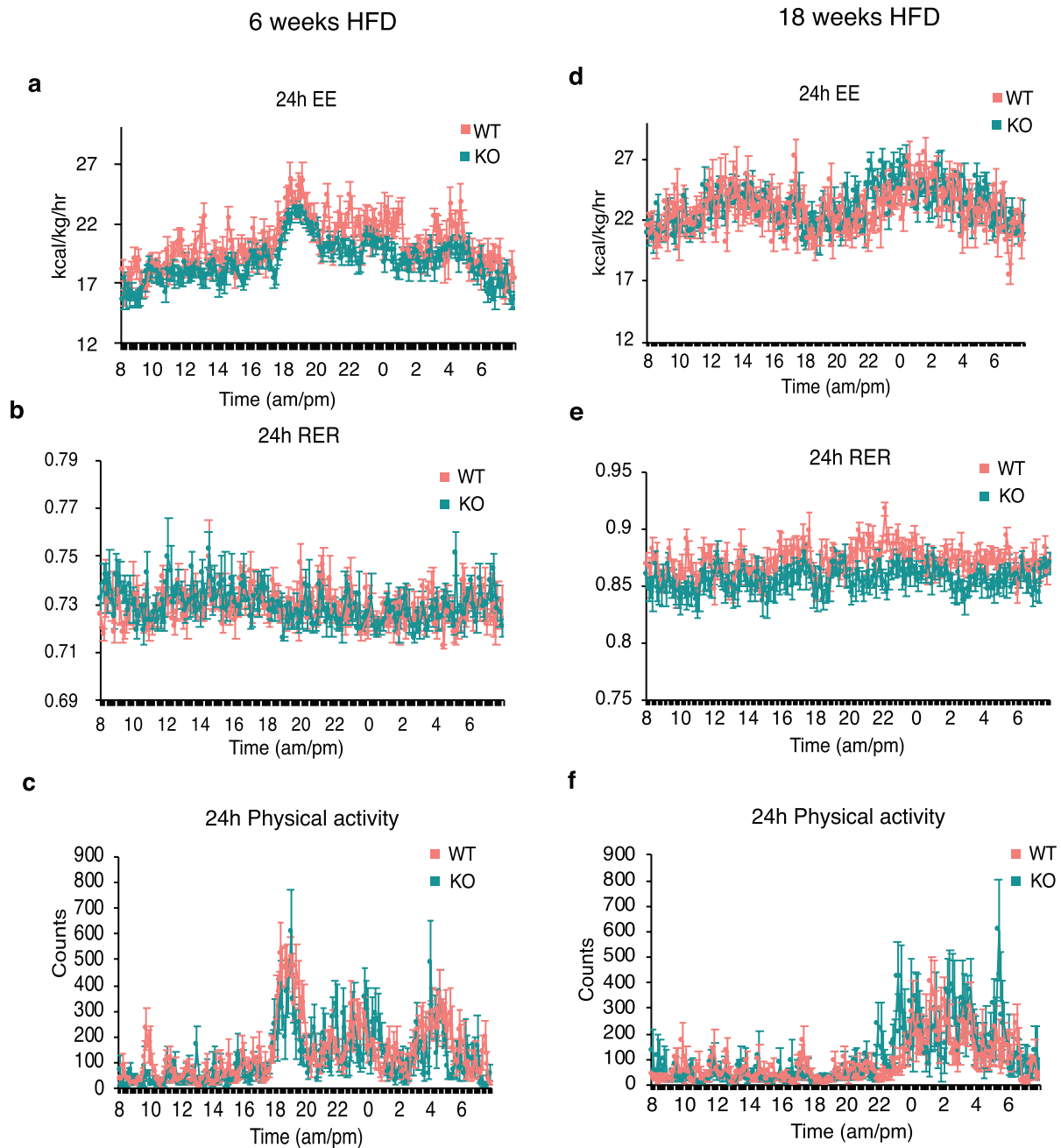


**Supplementary Figure 6 Loss of AIF1L does not affect weight gain curves or adiposity in female mice upon long-term HFD feeding.**

**Female mice on HFD for 16-18 weeks**

- Total body weight curves of WT and KO mice fed high-fat diet starting at 8 weeks of age (n = 11 for WT, n = 13 for KO)
- Body weight, fat mass and lean mass measured by MRI at the end of HFD feeding period; age - 26 weeks. (n = 7 for WT, n = 10 for KO)
- Food intake over 4 weeks (n = 5)
- Brown (BAT), inguinal subcutaneous (iSAT), peri-ovarian (poWAT) white adipose depot, and liver mass. (n = 7 for WT, n = 10 for KO)
- Gross appearance of tissues- representative images

Data are represented as mean  $\pm$  SEM. ns - not significant.

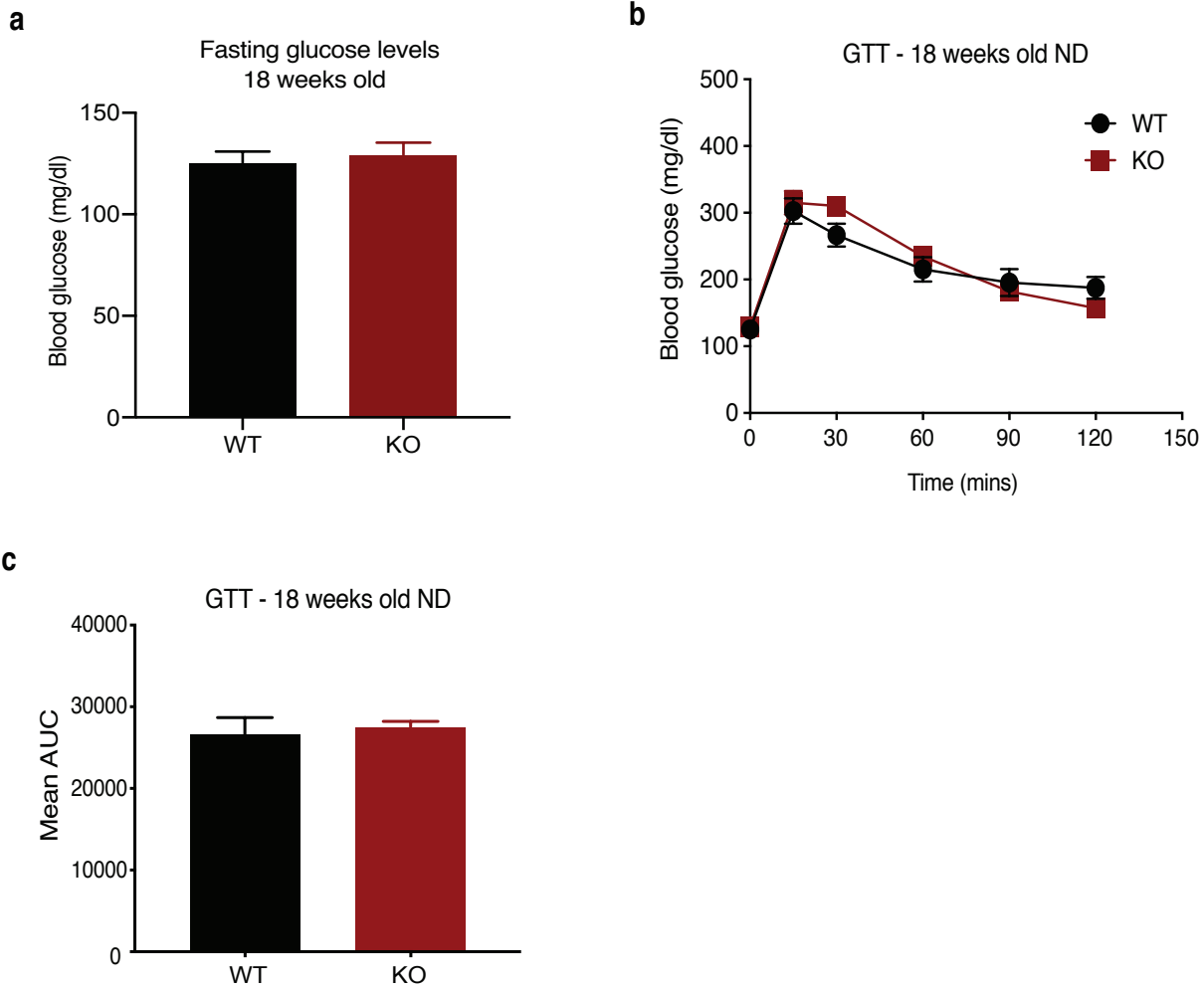


**Supplementary Figure 7 Metabolic profile of WT and AIF1L deficient mice over a 24h period, upon short- and long-term HFD feeding**

a-c) Energy expenditure, RER, and Physical activity measurements over a 24 h period in metabolic cages for WT and KO mice fed HFD for 6 weeks (n = 3 for each genotype)

d-f) Energy expenditure, RER, and Physical activity measurements over a 24 h period in metabolic cages for WT and KO mice fed HFD for 18 weeks (n = 3 for each genotype)

Data are represented as mean  $\pm$  SEM. ns- not significant.



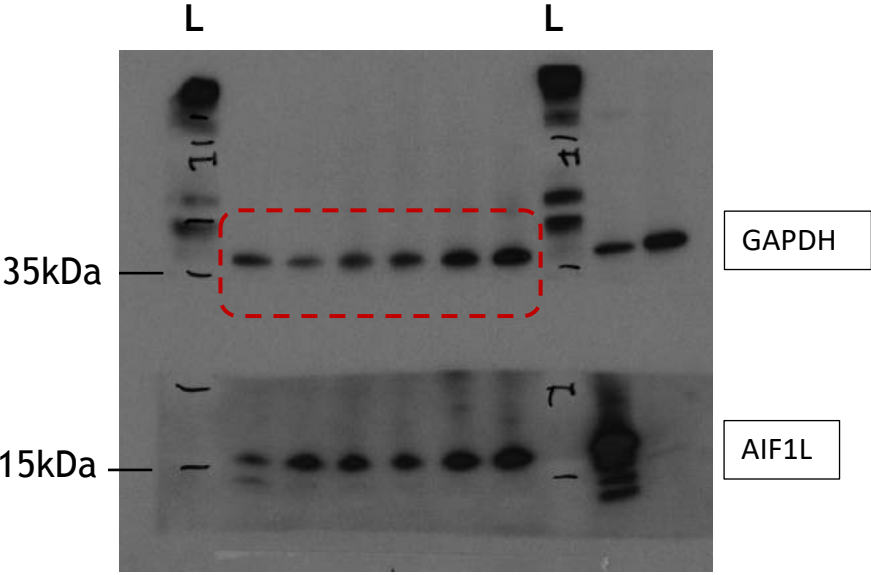
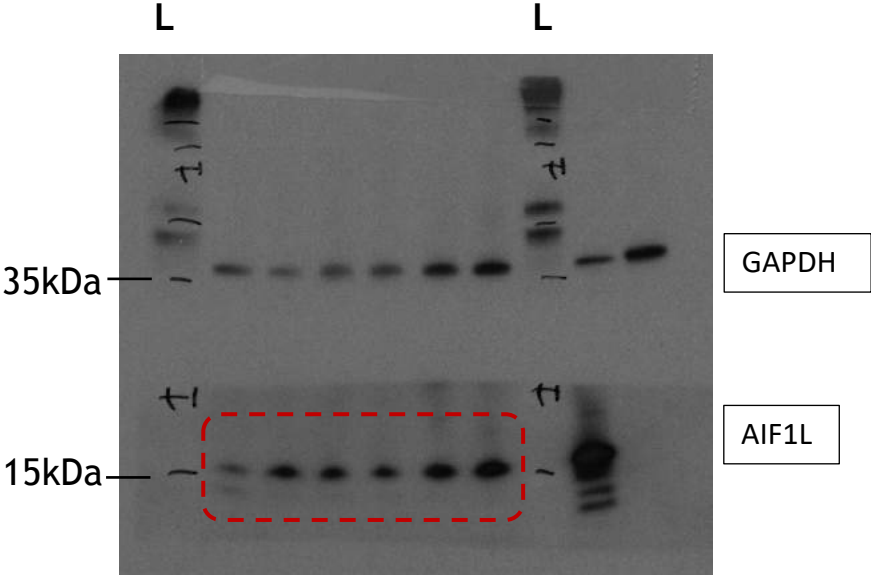
**Supplementary Figure 8. 18 weeks old WT and AIF1L deficient mice have similar response to glucose challenge**

**(a–c) 18 weeks old male mice on ND**

- a) Blood glucose levels in WT (n = 3) and KO (n = 4) mice after overnight fasting.
- b) IPGTT (intra peritoneal glucose tolerance test) with 1g/kg (of body weight) of glucose.
- c) Quantification of GTT as mean area under curve (AUC).

Data are represented as mean  $\pm$  SEM. ns - not significant.

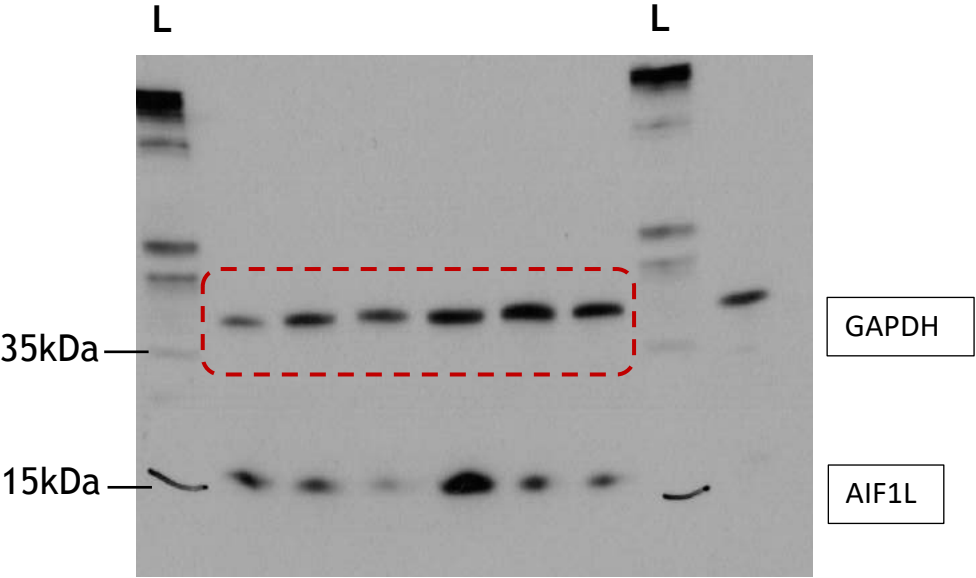
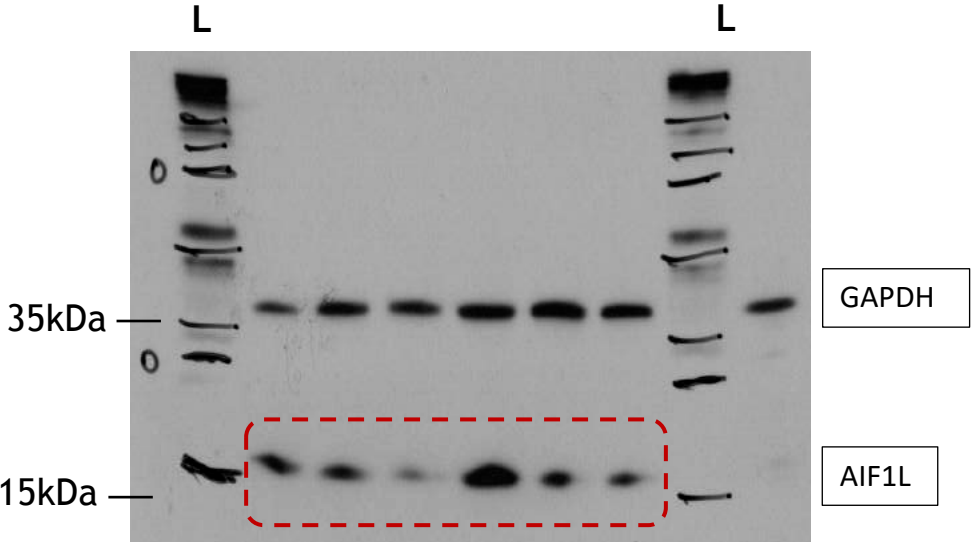
Uncropped blots for Figure 1a- BAT




Cropped along the dotted lines

L Ladder

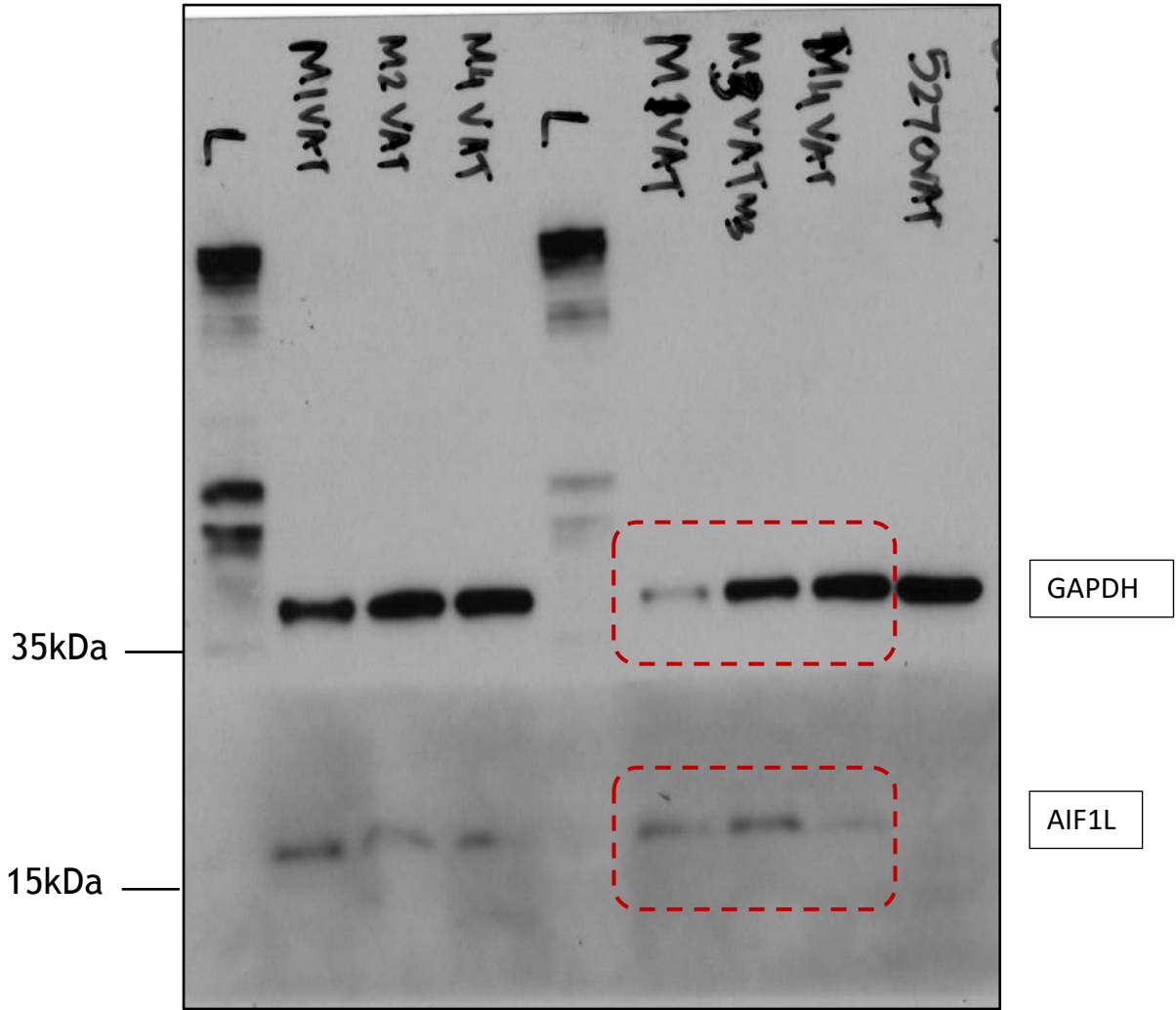
Uncropped blots for Figure 1a- SAT




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**L** Ladder

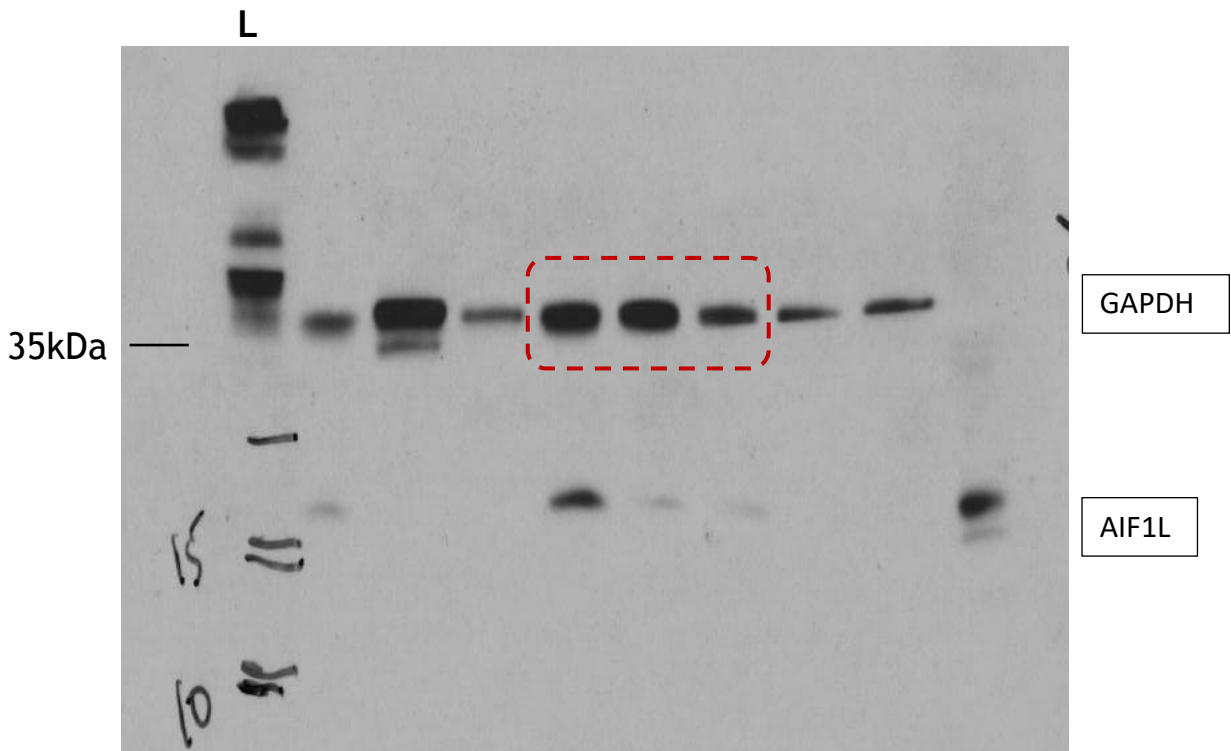
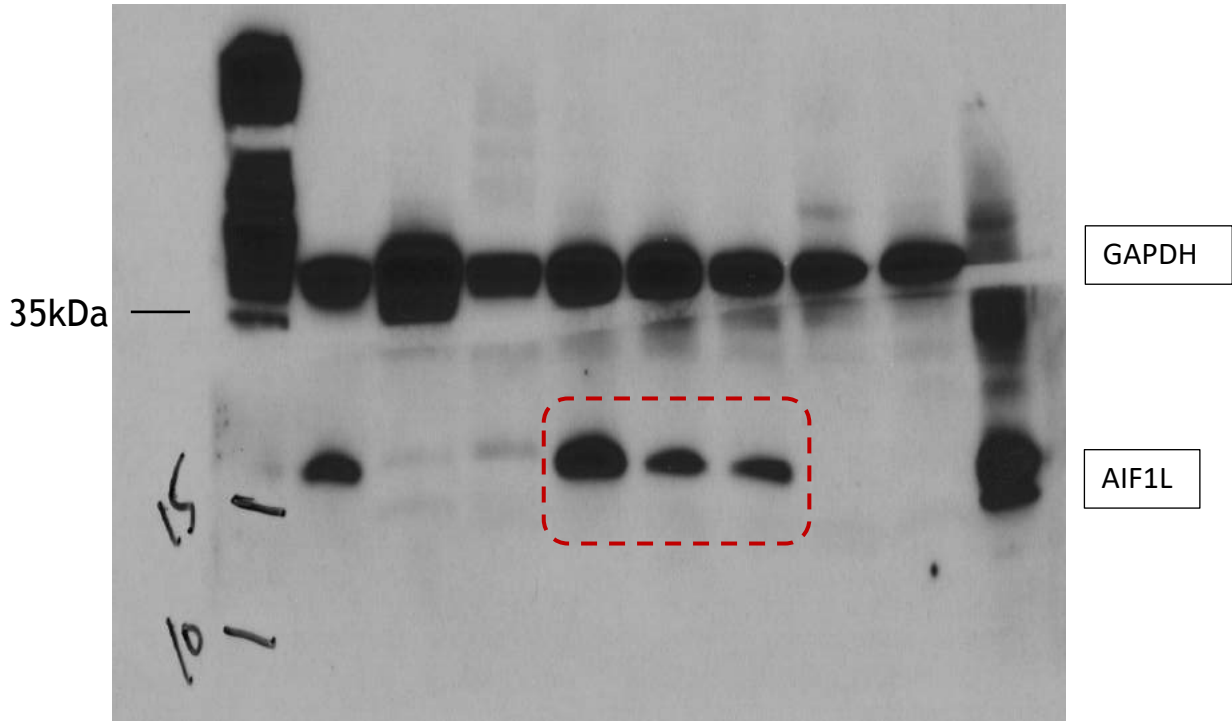
Uncropped blots for Figure 1a- eWAT Males



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**L** Ladder

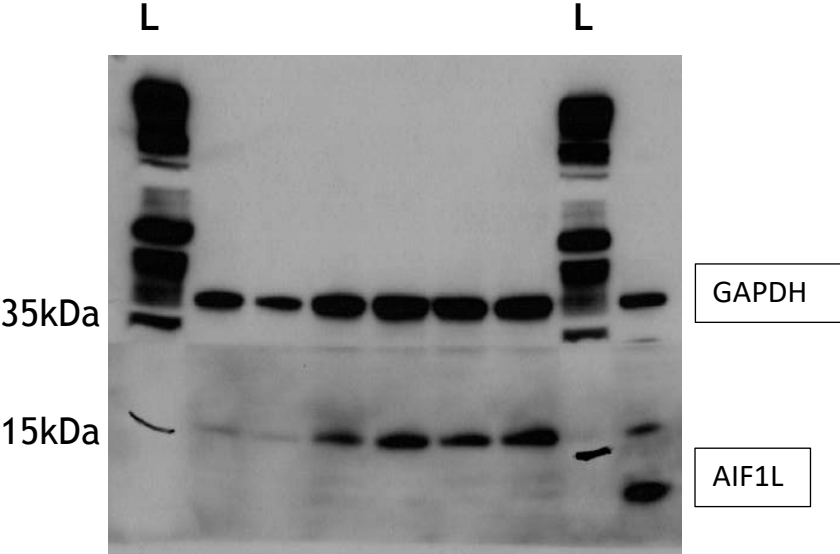
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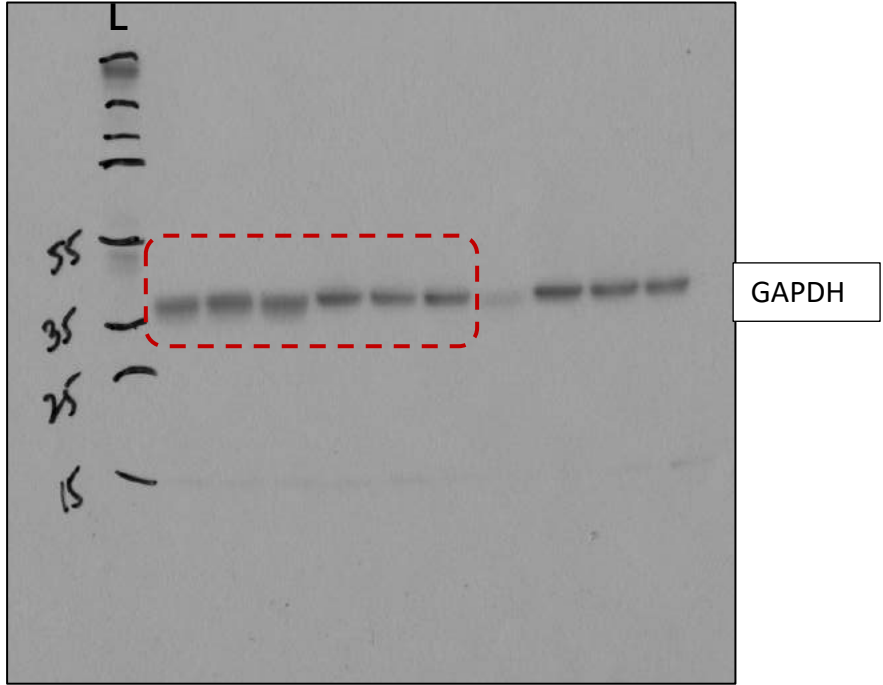
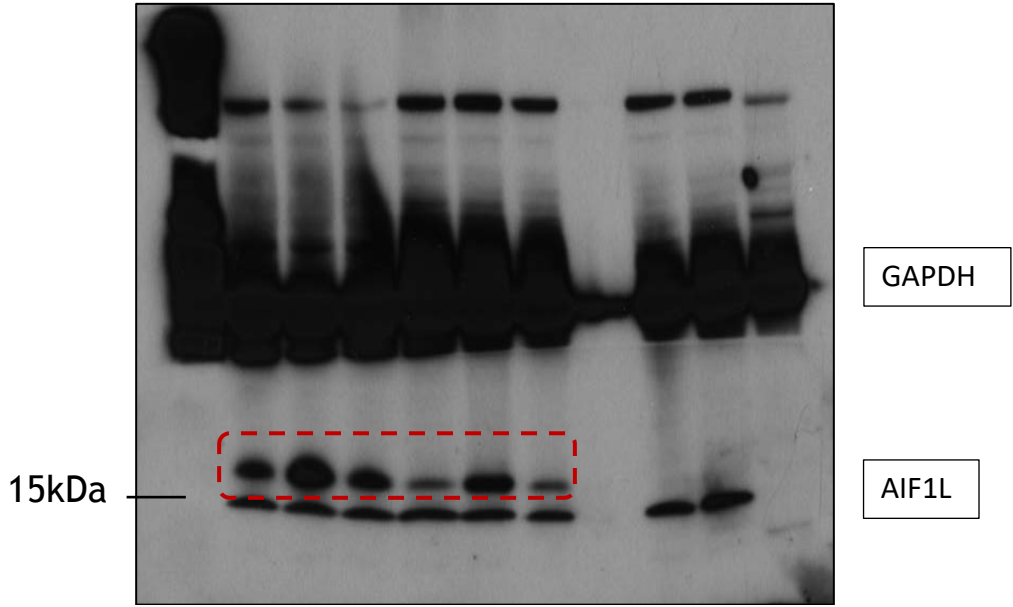
L Ladder


**Uncropped blot #2 used for VAT quantification and comparison between males and females -**





Uncropped blots for Figure 1b- Kidney

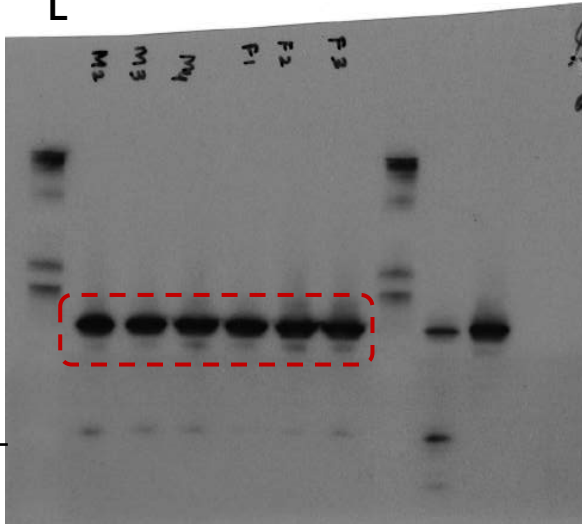


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**L** Ladder

Uncropped blots for Figure 1b- Brain

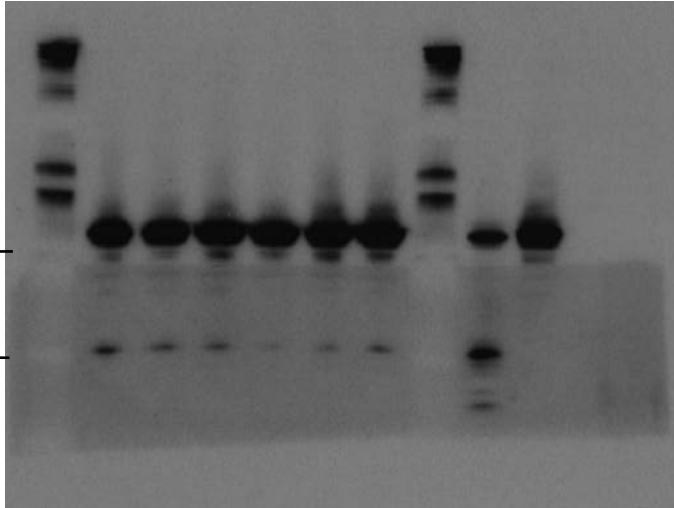
L



GAPDH

AIF1L

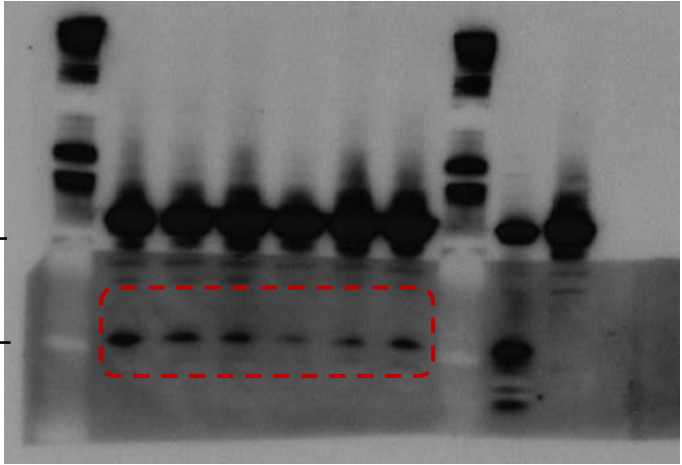
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GAPDH

AIF1L

L



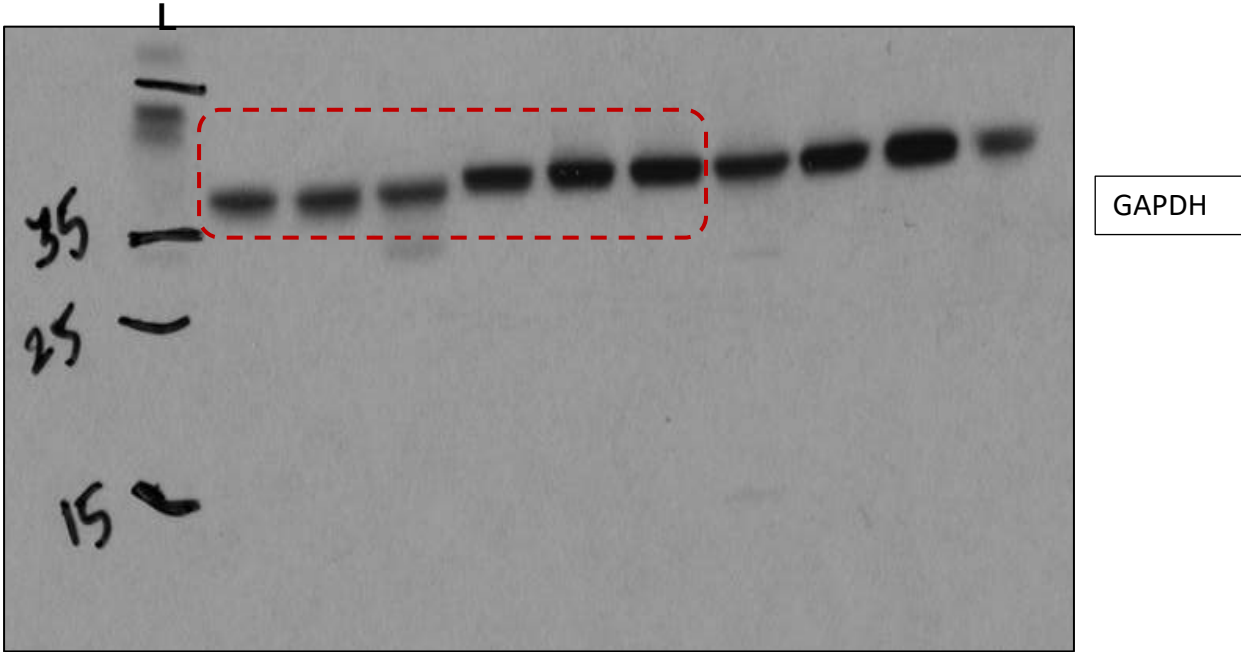
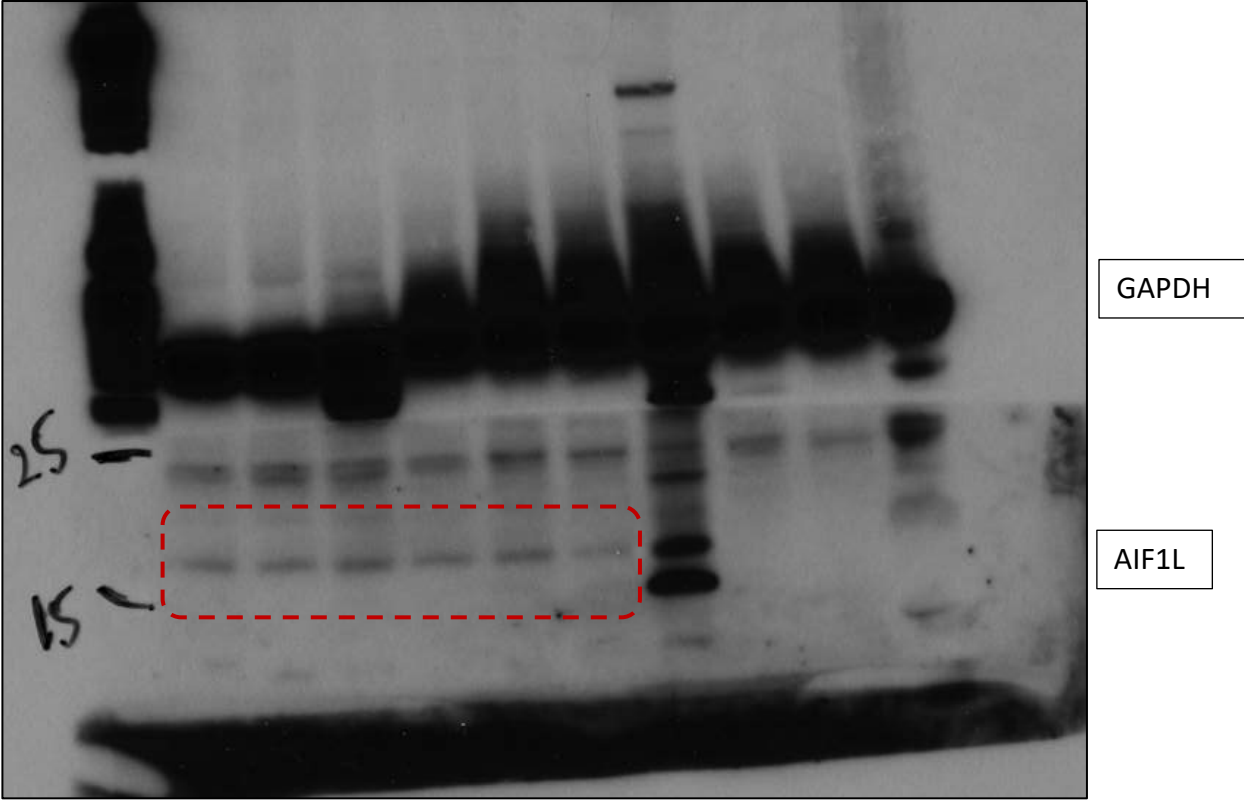
GAPDH

AIF1L

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L Ladder

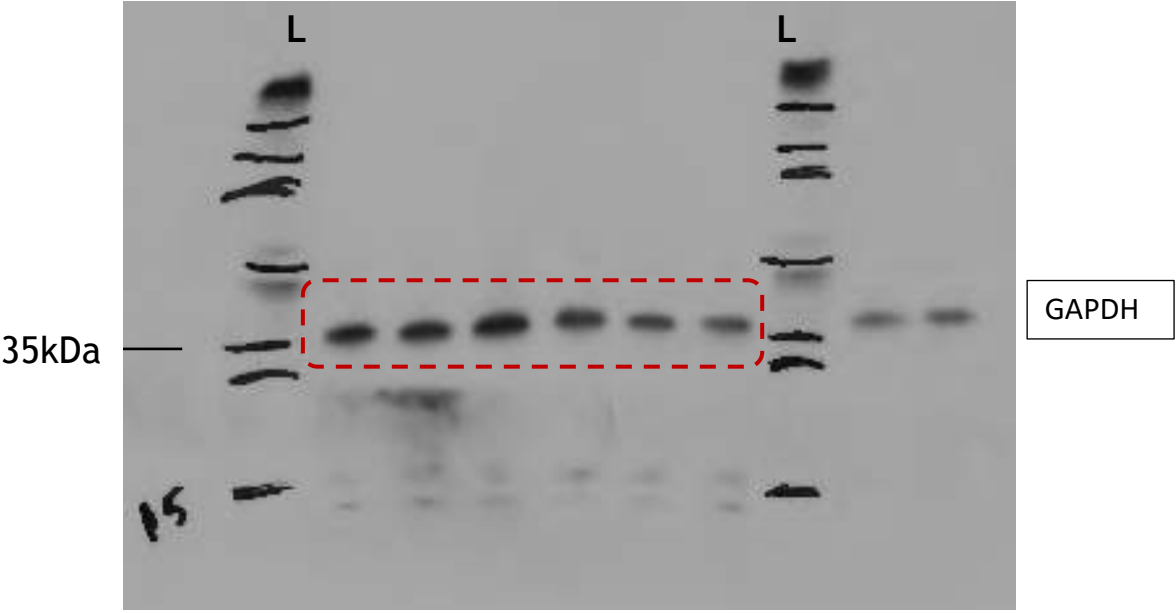
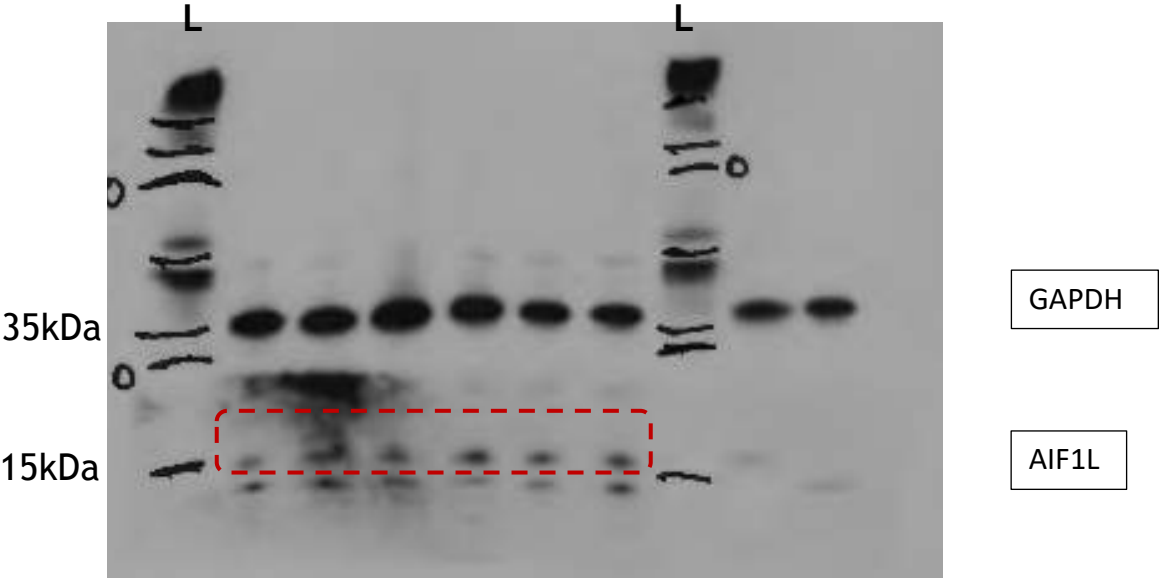
Uncropped blots for Figure 1b- Lung




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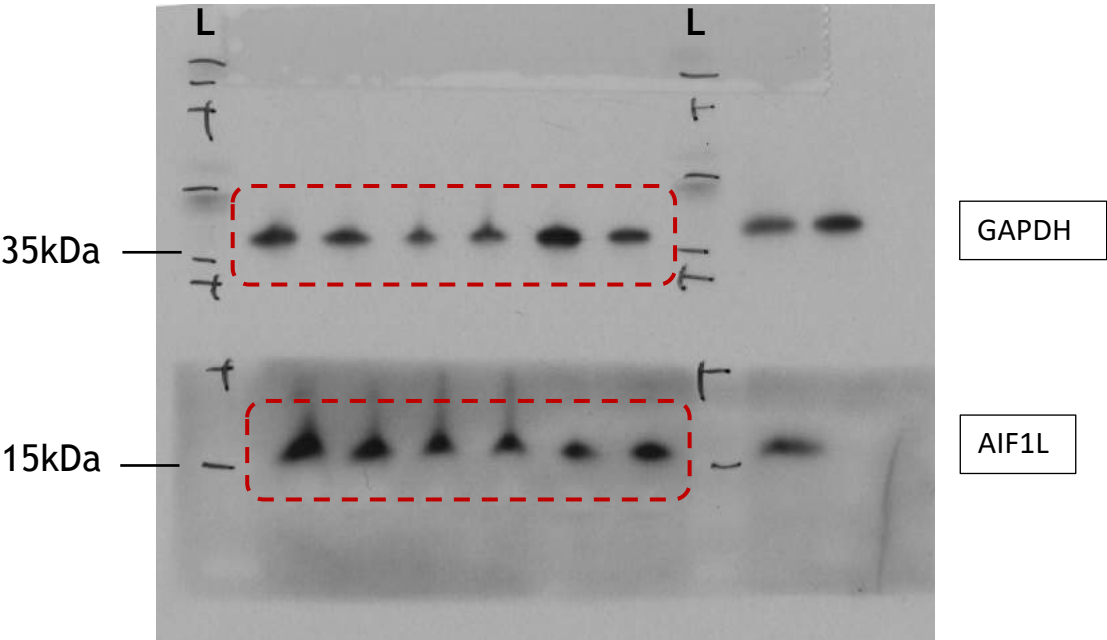
L Ladder

Uncropped representative blots used in Figure 1c- BAT

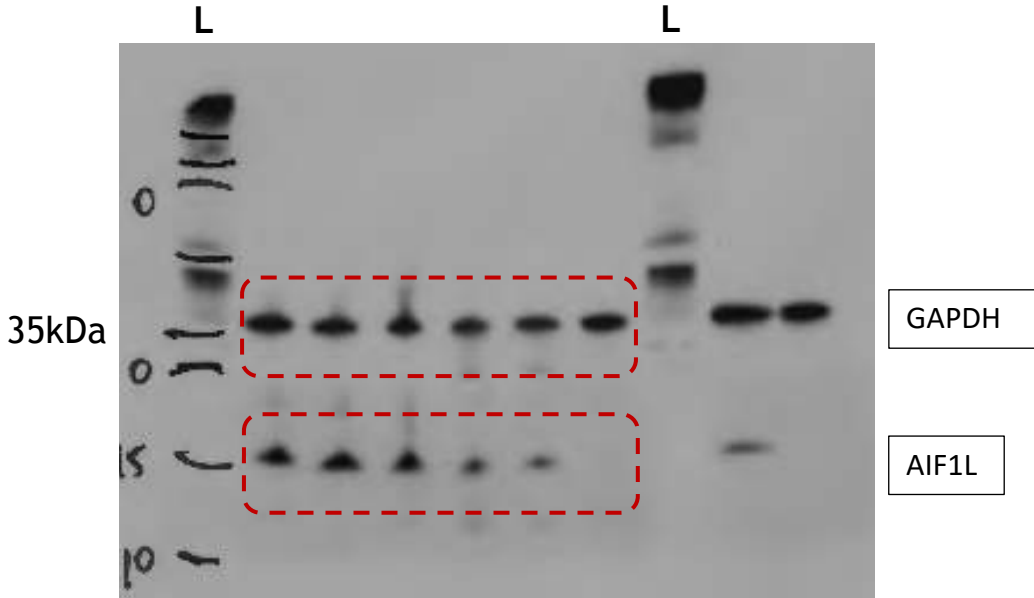



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**Uncropped blots for Figure 1c- iSAT**

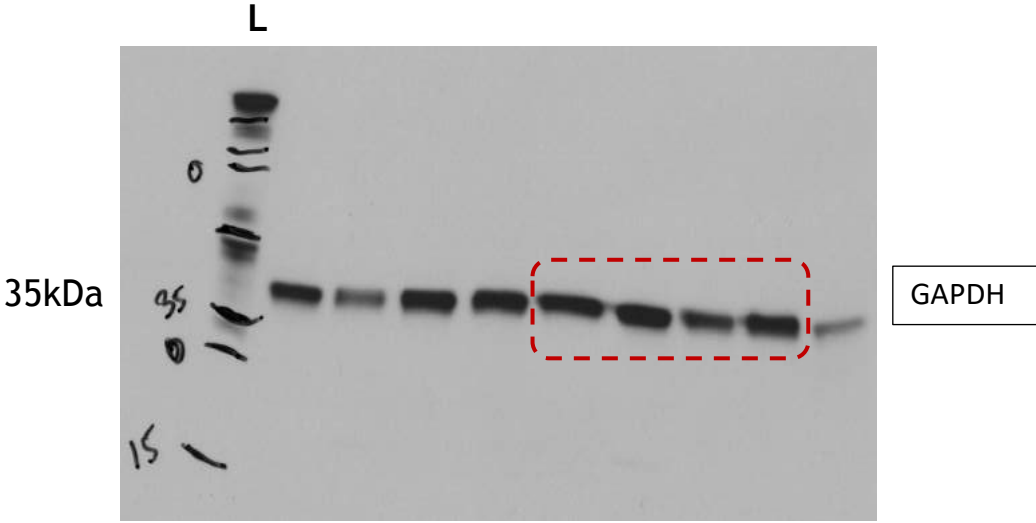
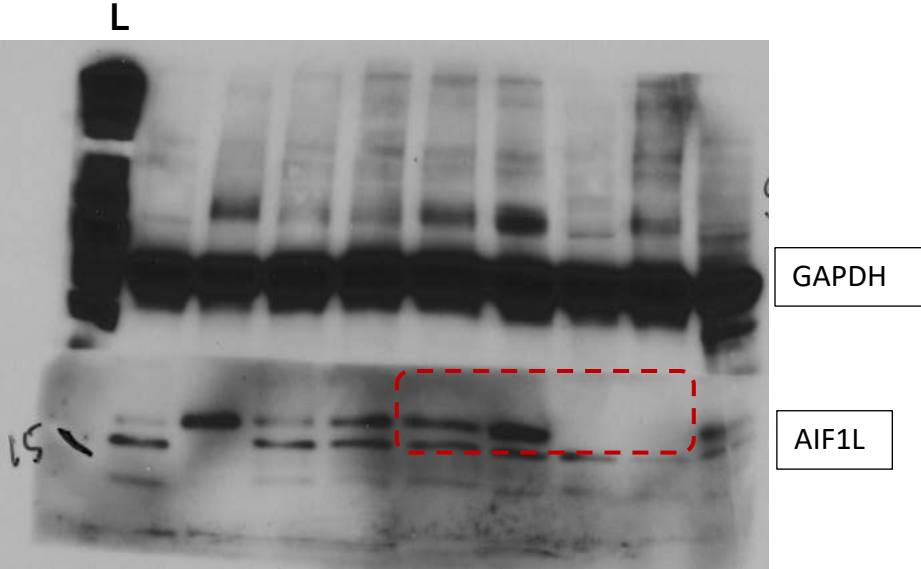


**Uncropped blots for Figure 1c- eWAT**



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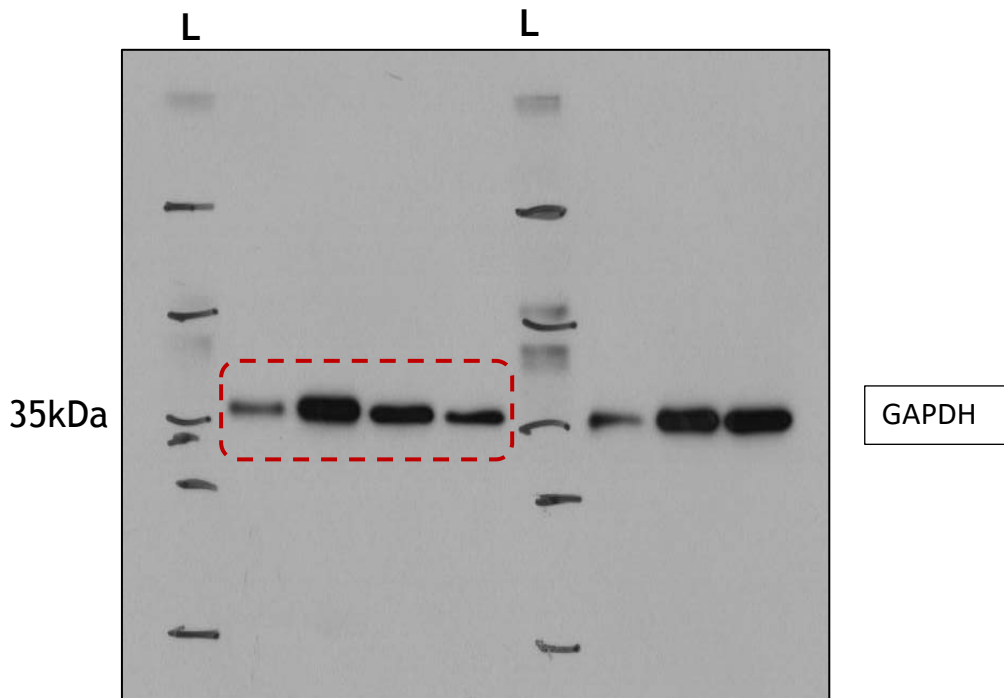
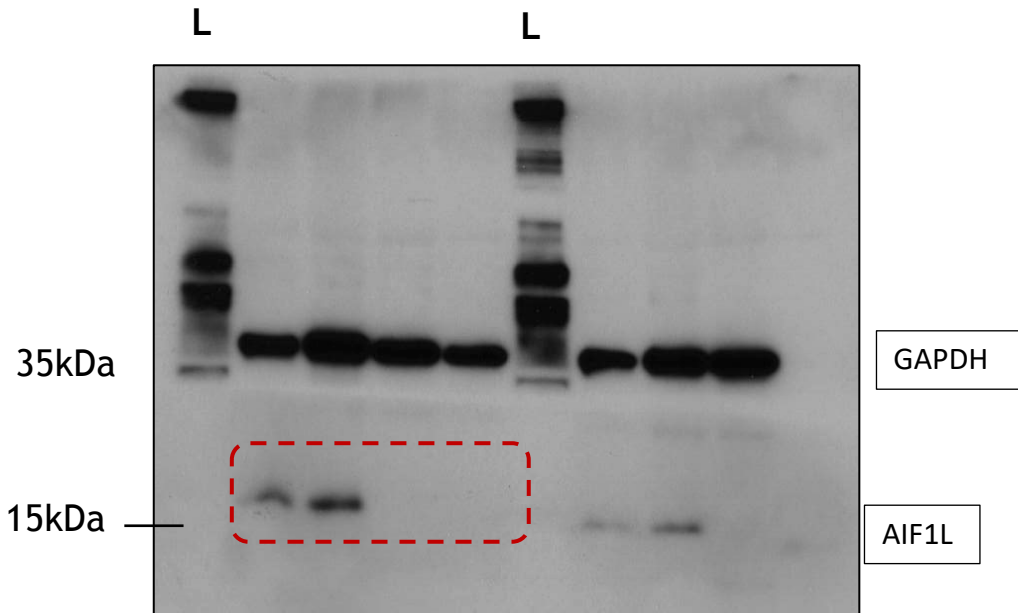
Uncropped blots for Figure 2a- BAT



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L Ladder

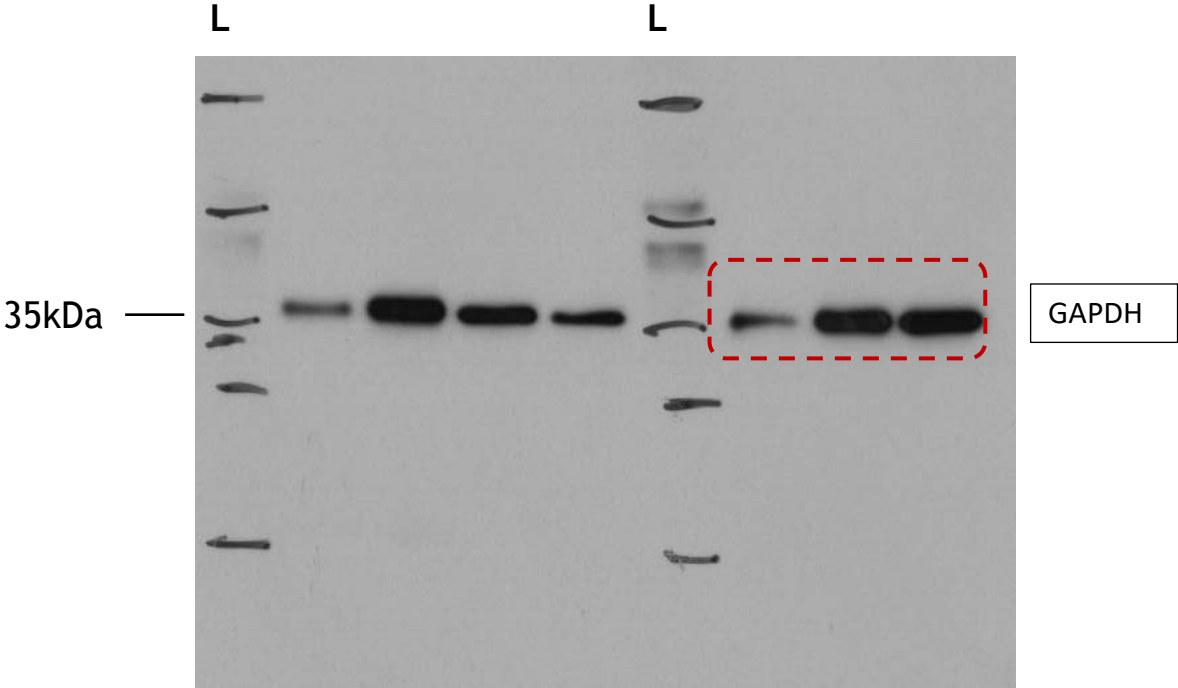
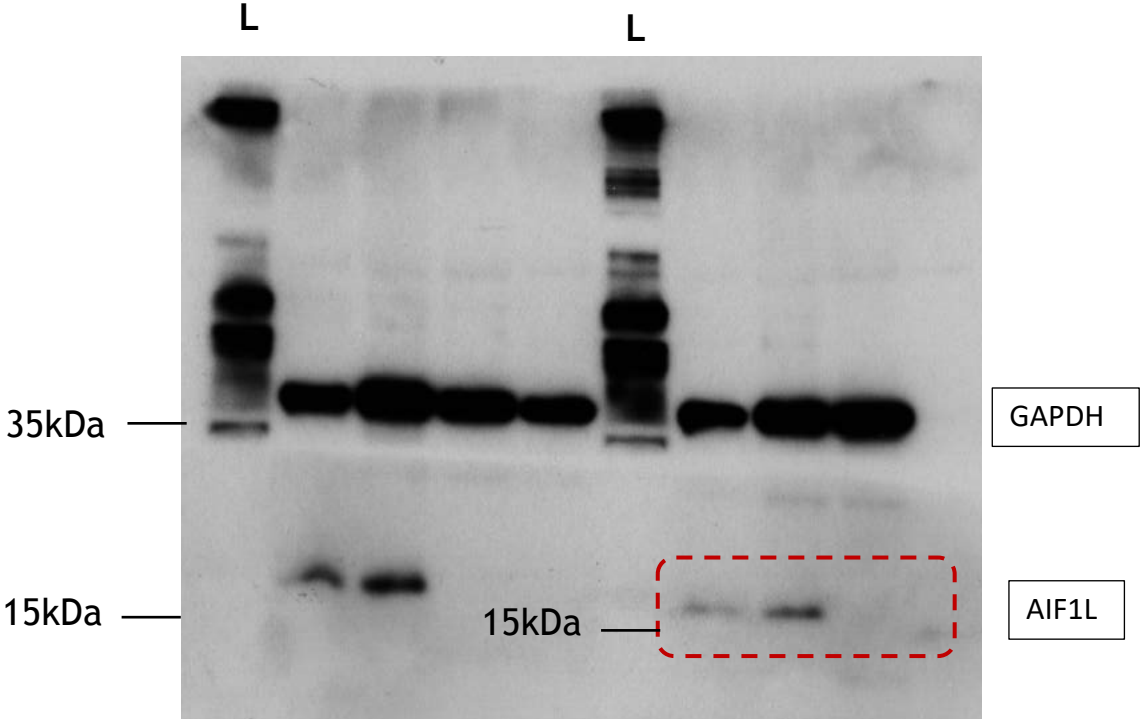
Uncropped blots for Figure 2a- iSAT




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**L** Ladder

Uncropped blots for Figure 2a- eWAT

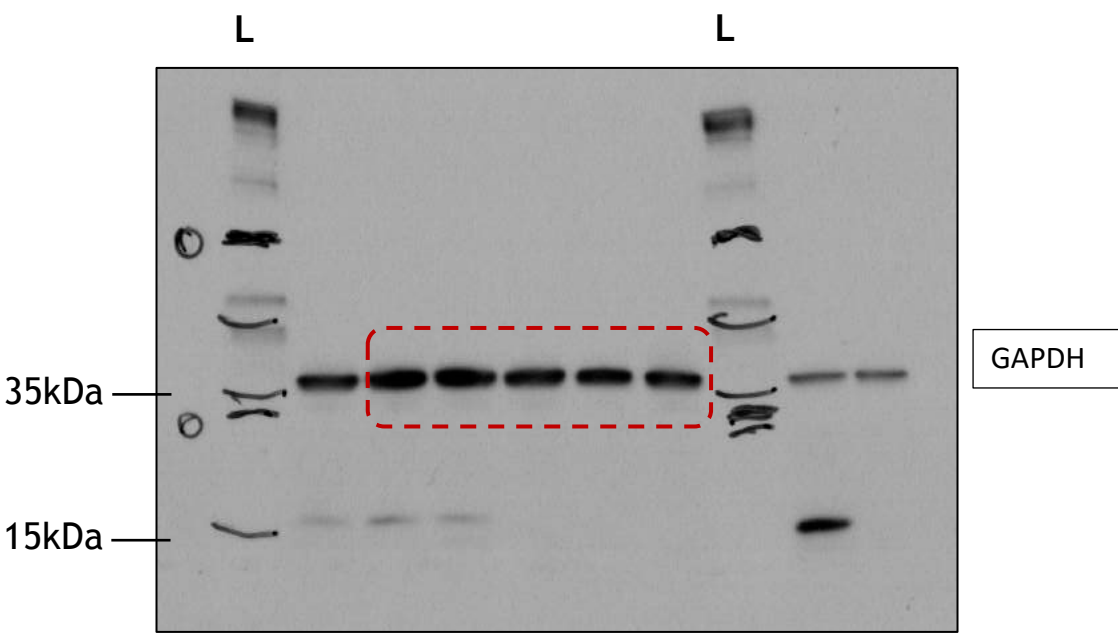
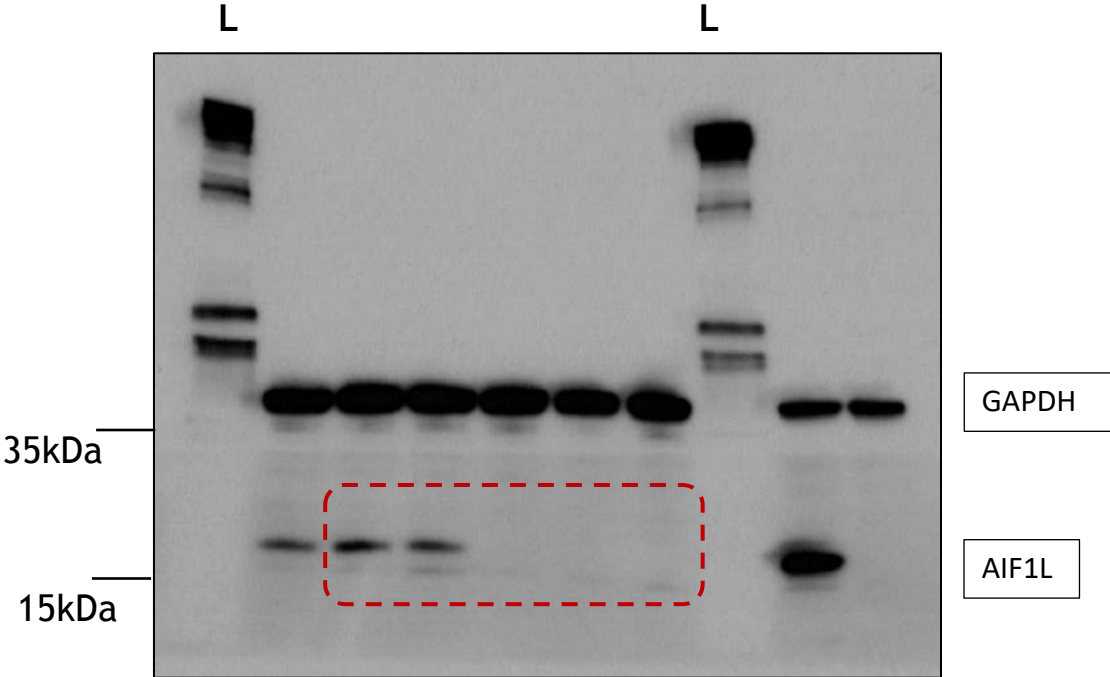



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**L** Ladder

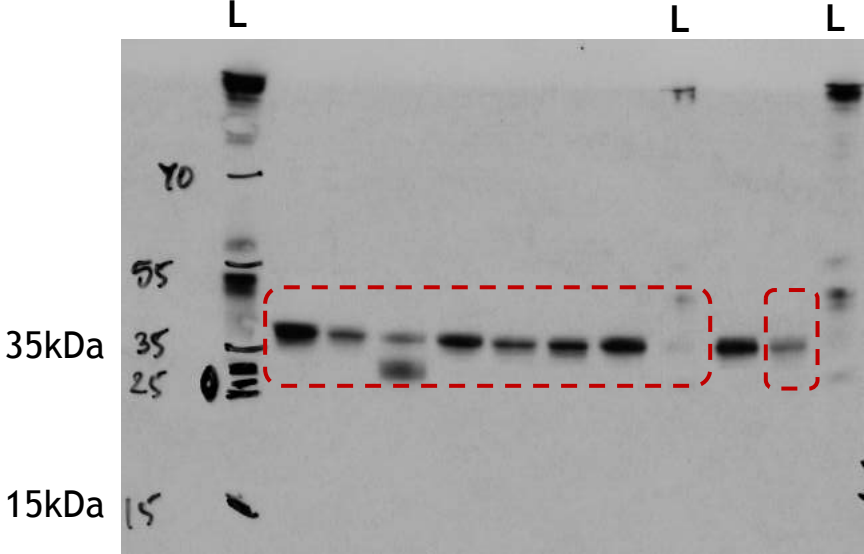
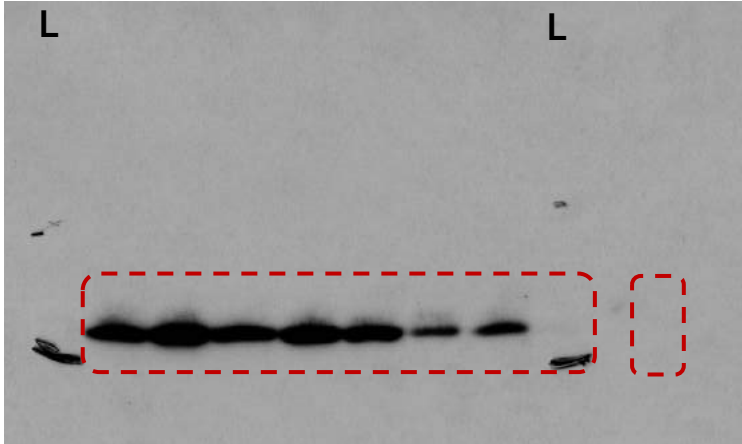
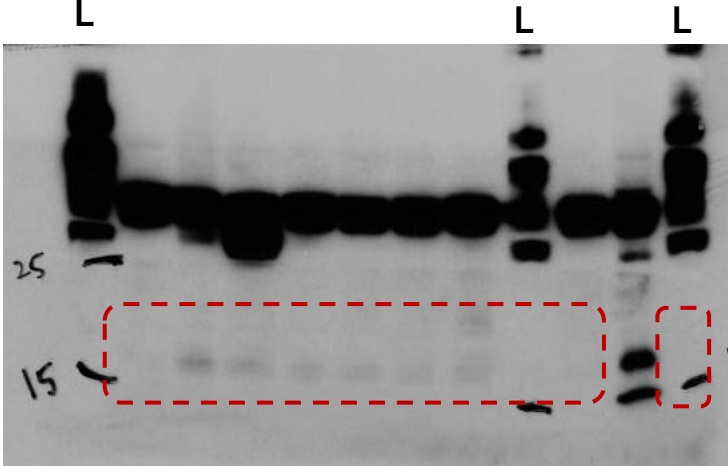



Uncropped blots for Figure 2a- Brain



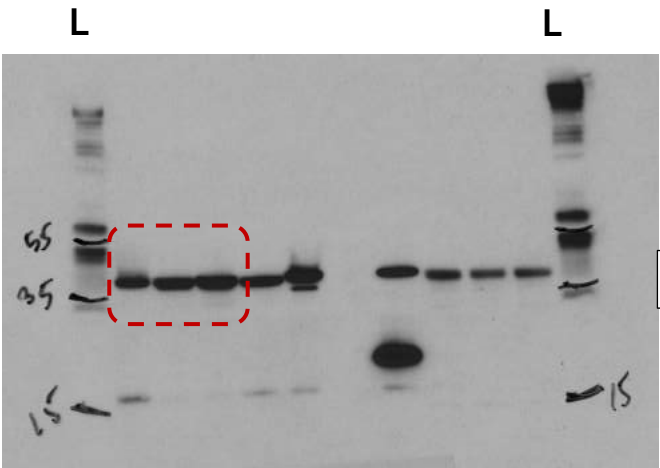
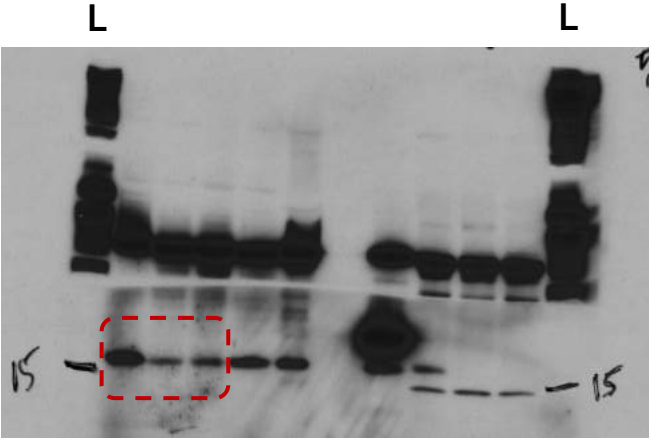
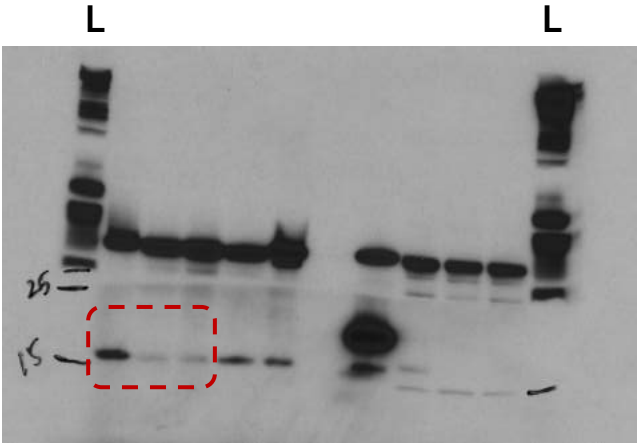
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Uncropped blots for Supplementary Figure 1a



 Cropped along the dotted lines      L Ladder

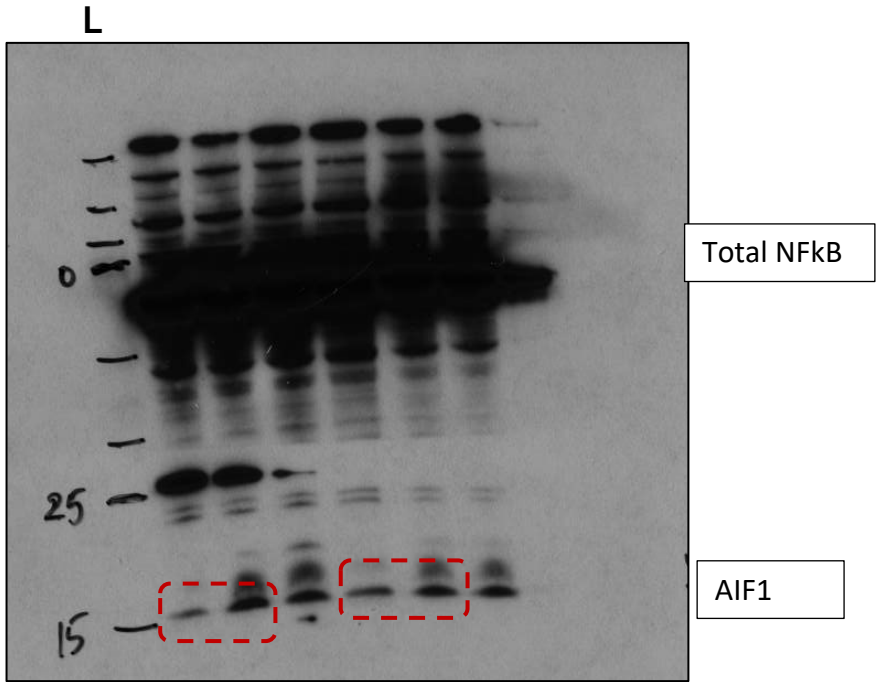
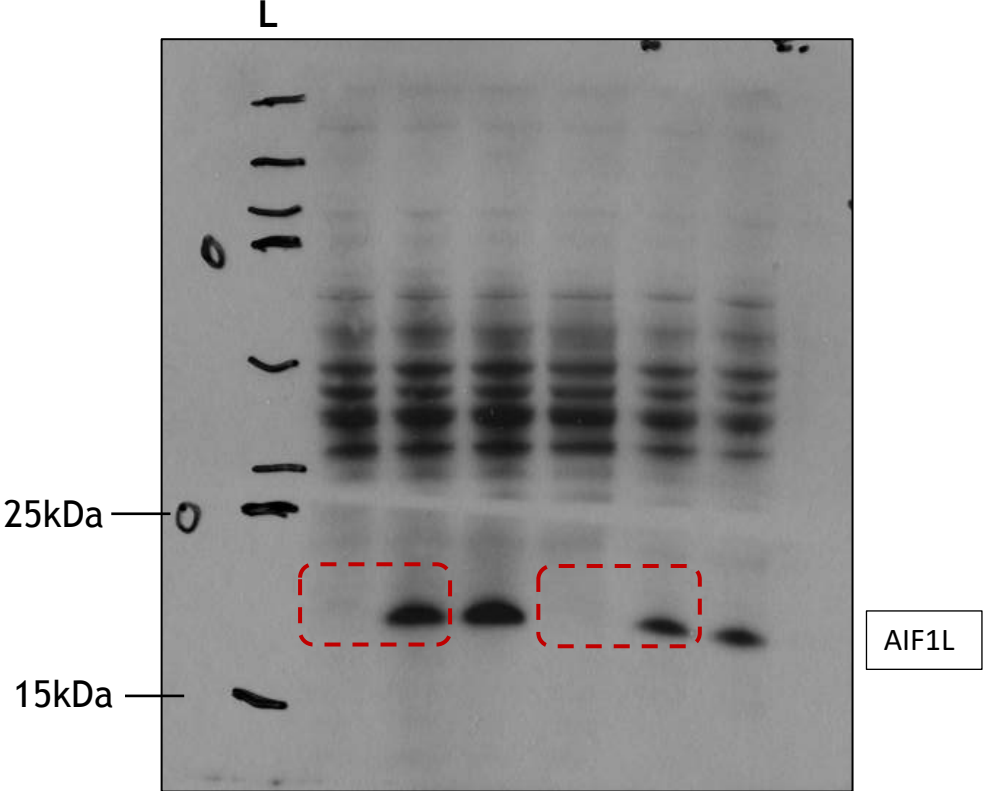
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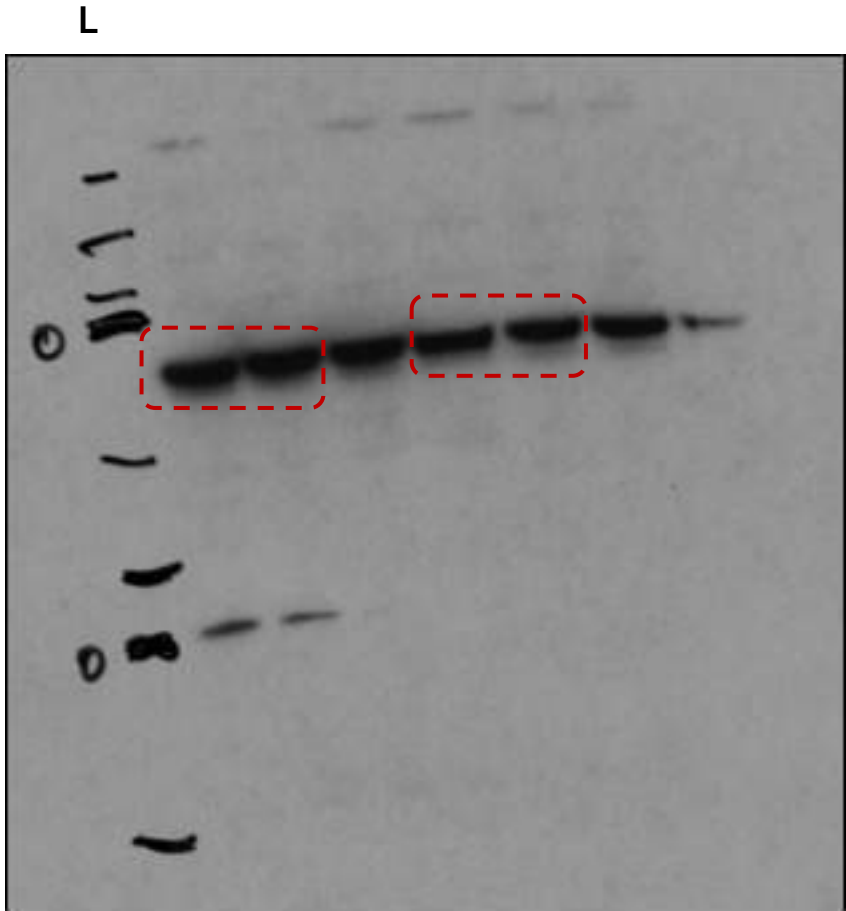
**L** Ladder

Uncropped blots for Supplementary Figure 1c- RAW 264.7 cells



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**L** Ladder



Total NFkB

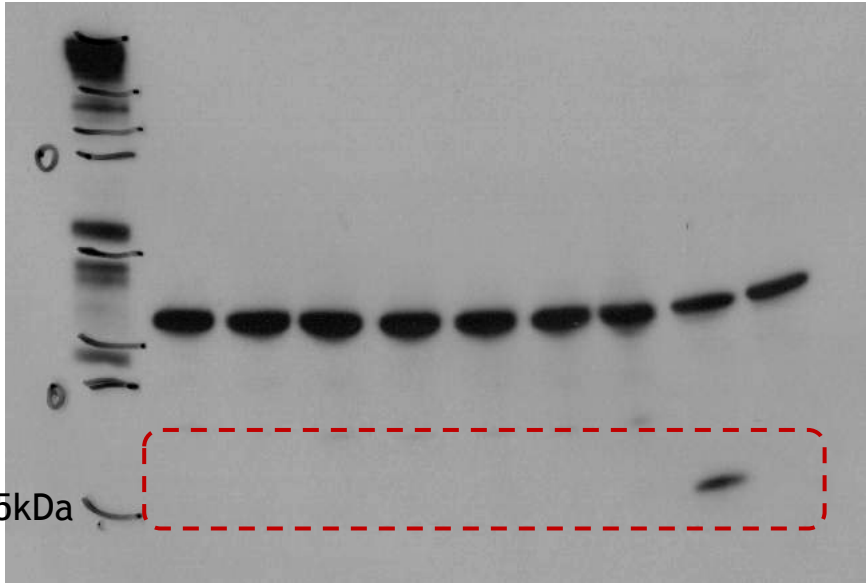


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L Ladder

Uncropped blots for Supplementary Figure 2a- Liver

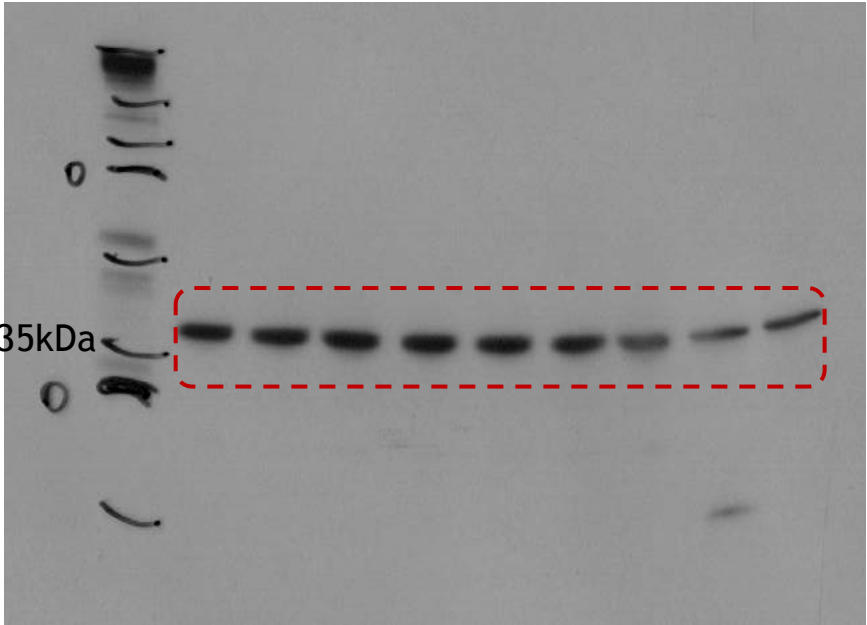
L



15kDa

AIF1L

L



35kDa

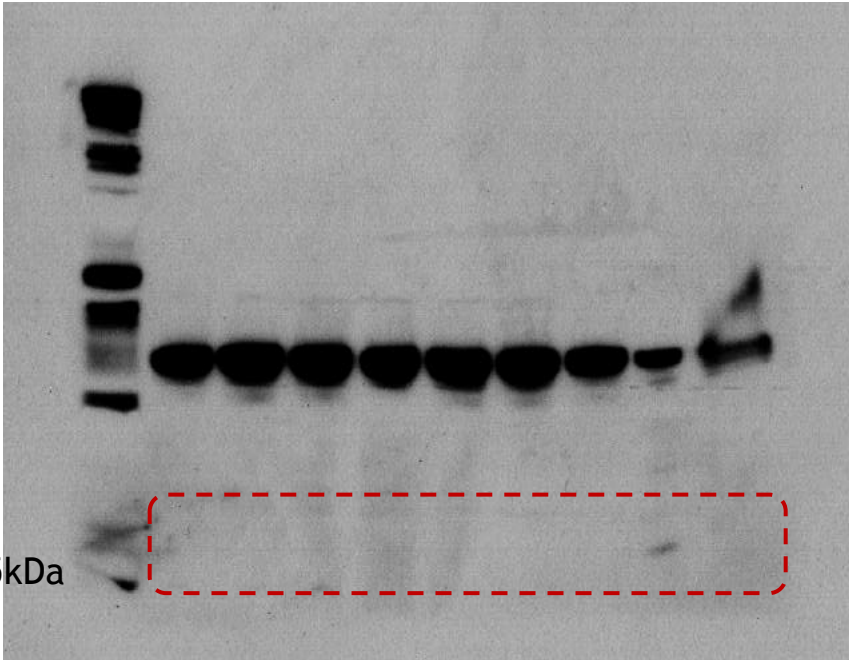
GAPDH

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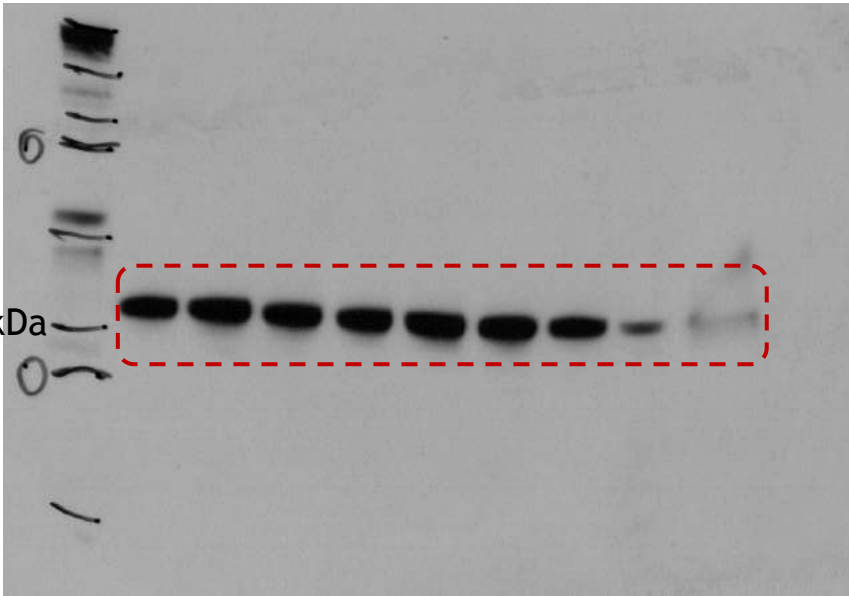
L Ladder

Uncropped blots for Supplementary Figure 2a- Heart

L



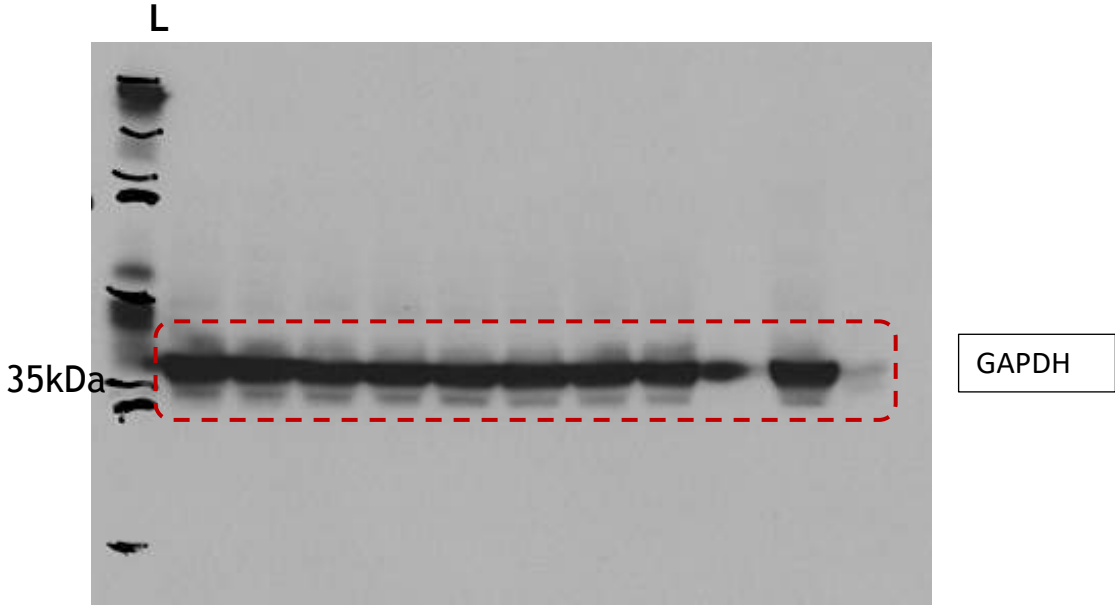
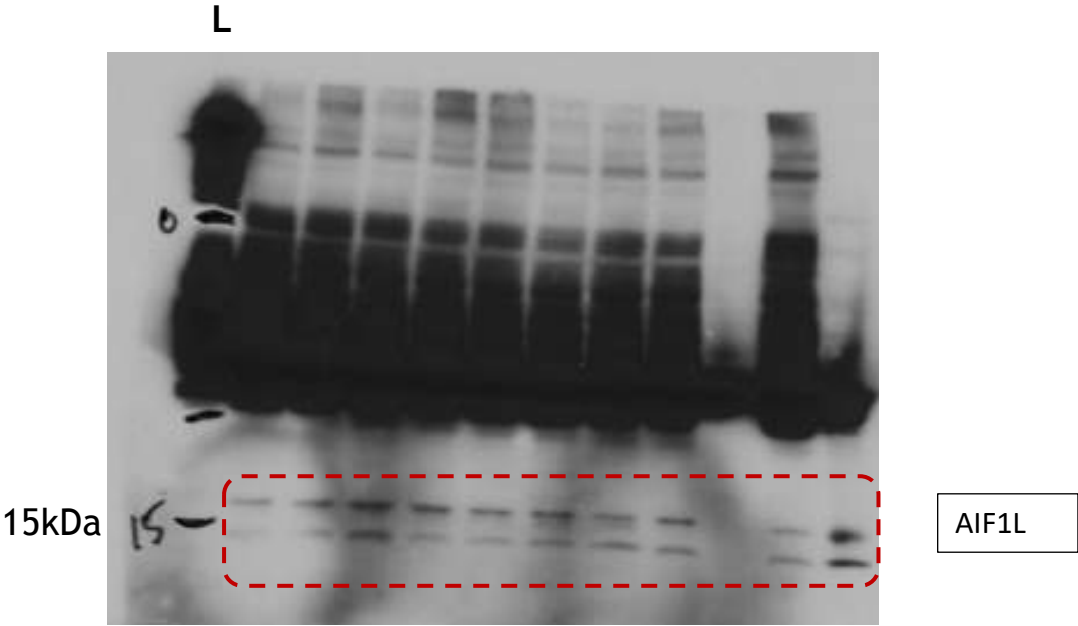
L



Cropped along the dotted lines

L Ladder

Uncropped blots for Supplementary Figure 2a- Skeletal muscle

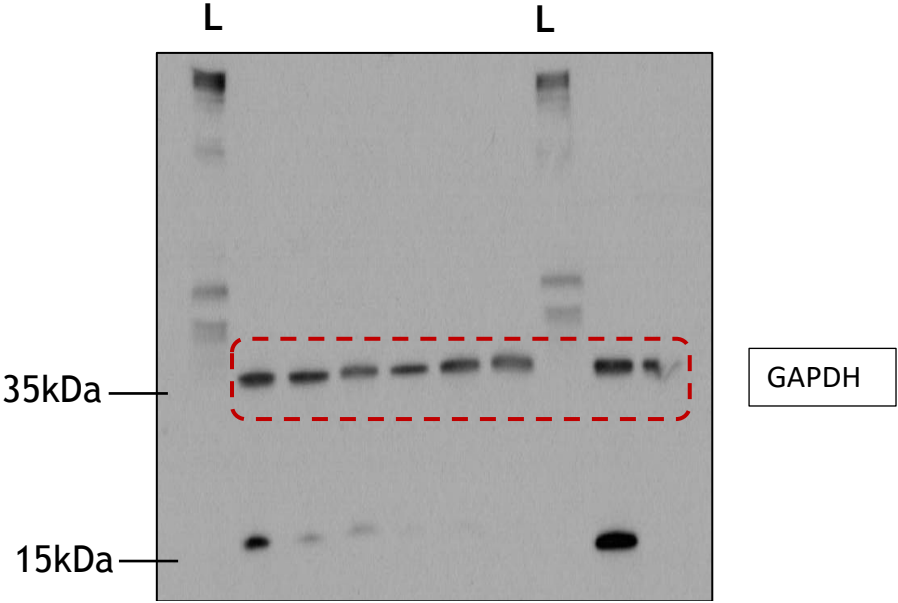
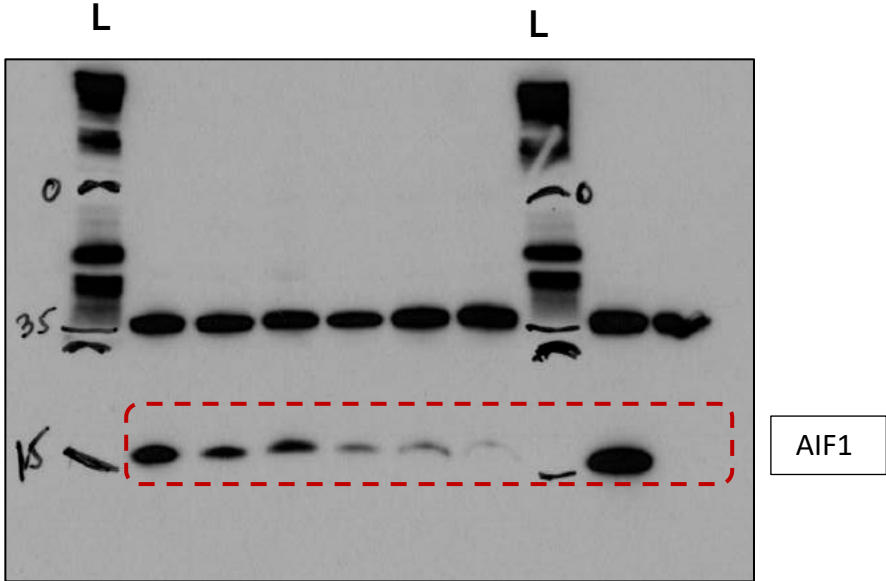



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L Ladder

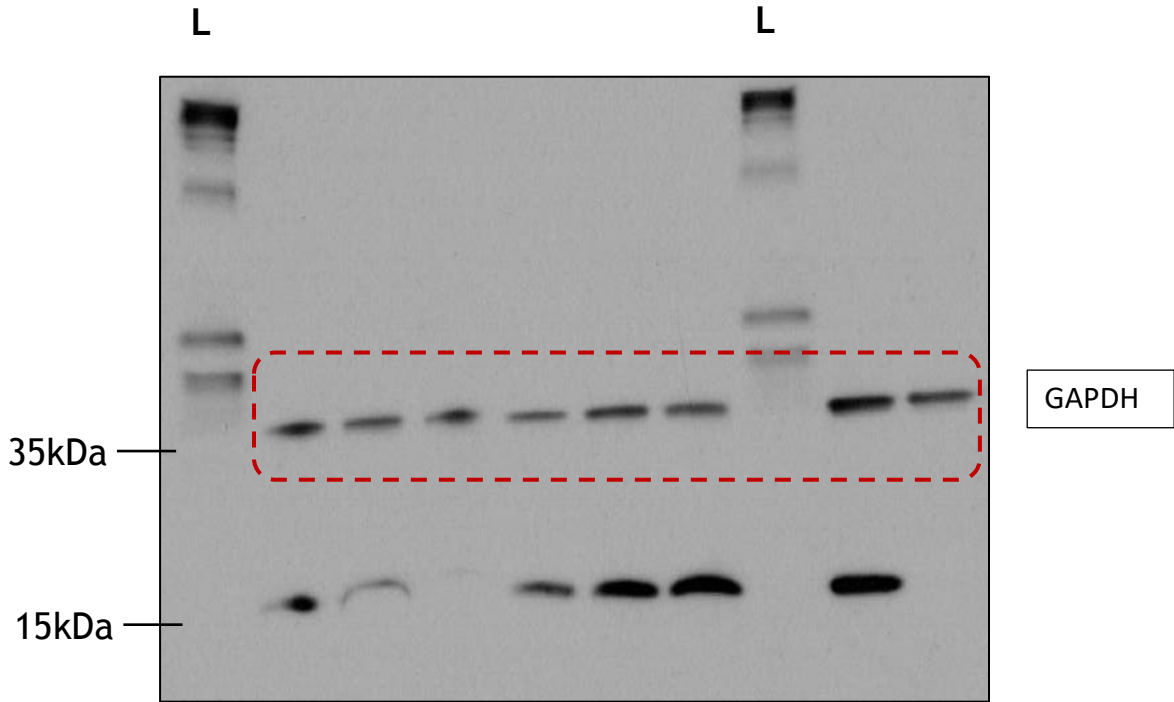
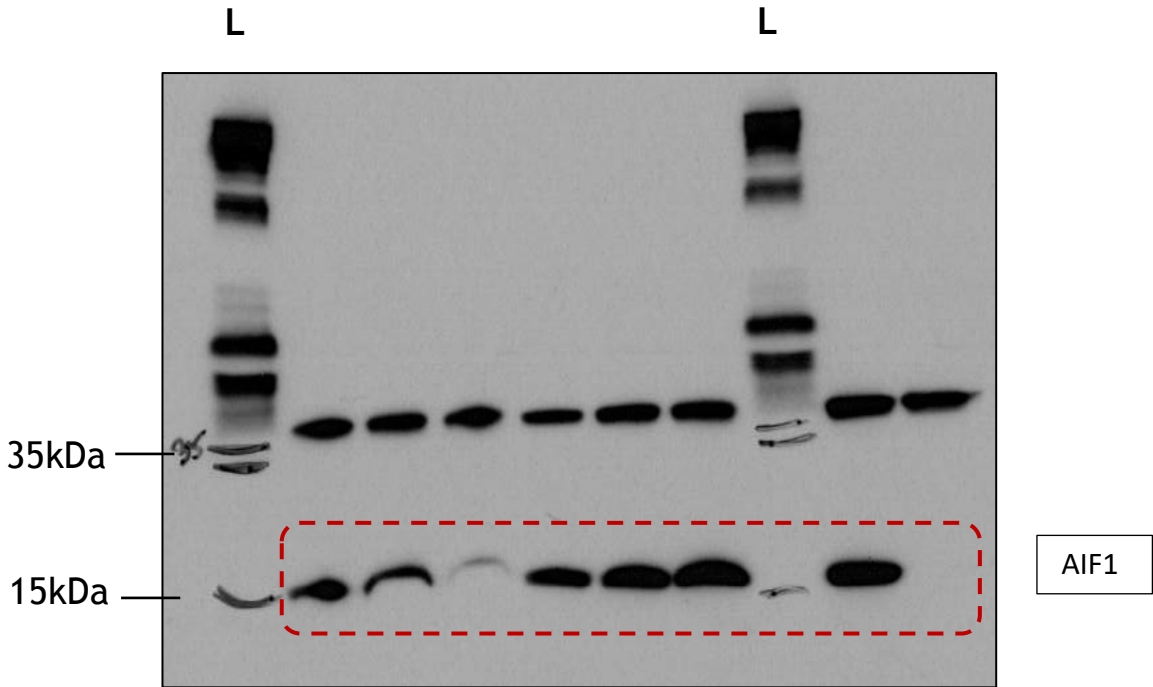



Uncropped blots for Supplementary Figure 4a- BAT



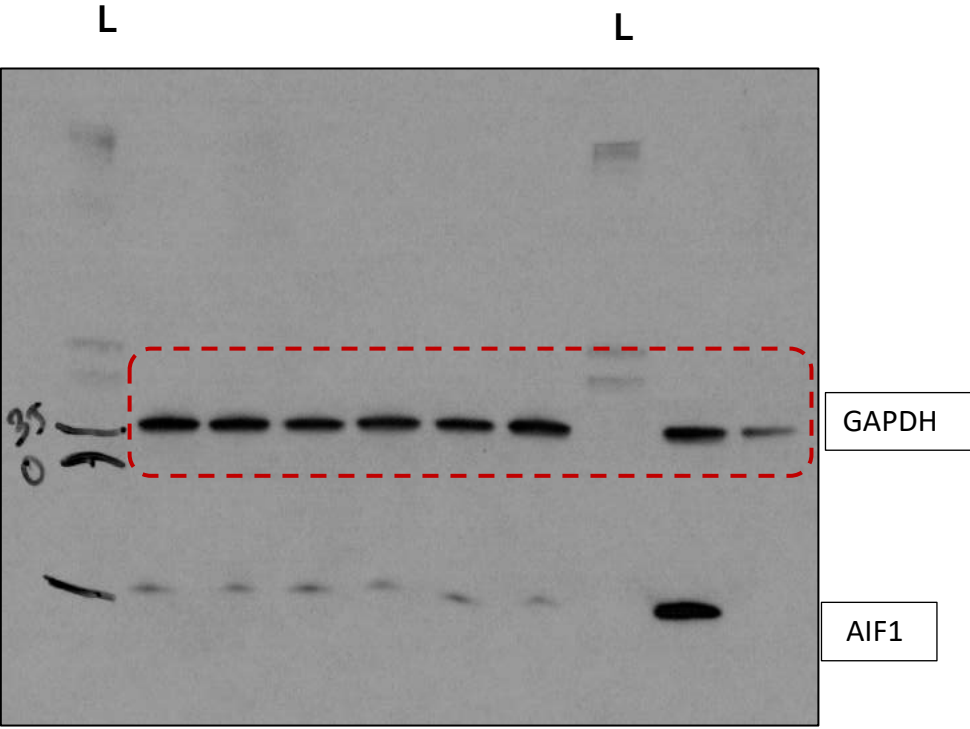
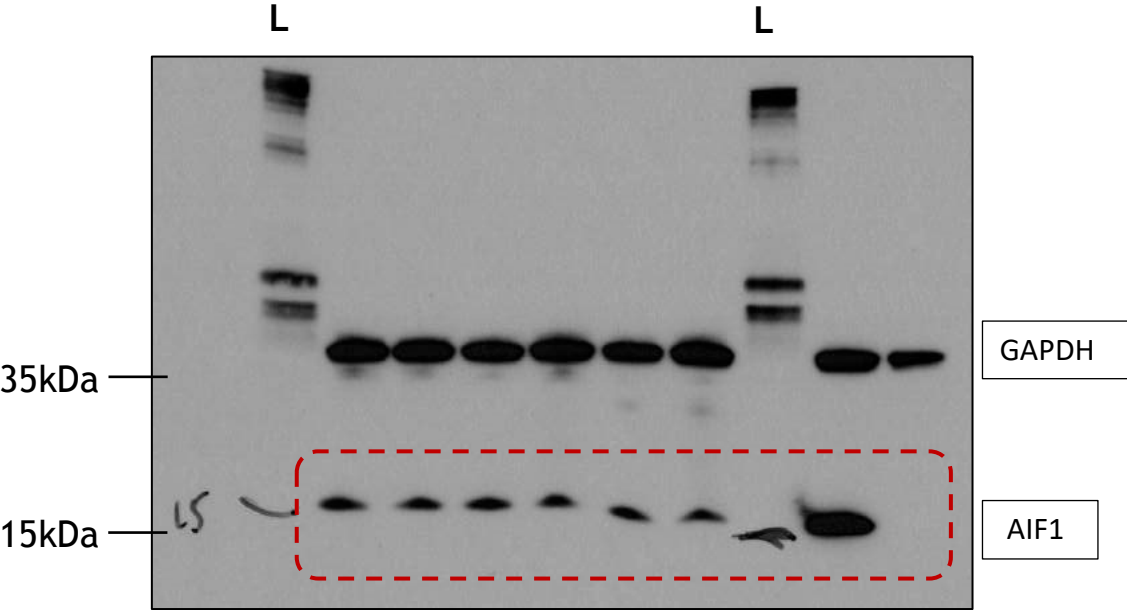
 Cropped along the dotted lines      **L** Ladder

Uncropped blots for Supplementary Figure 4a- VAT



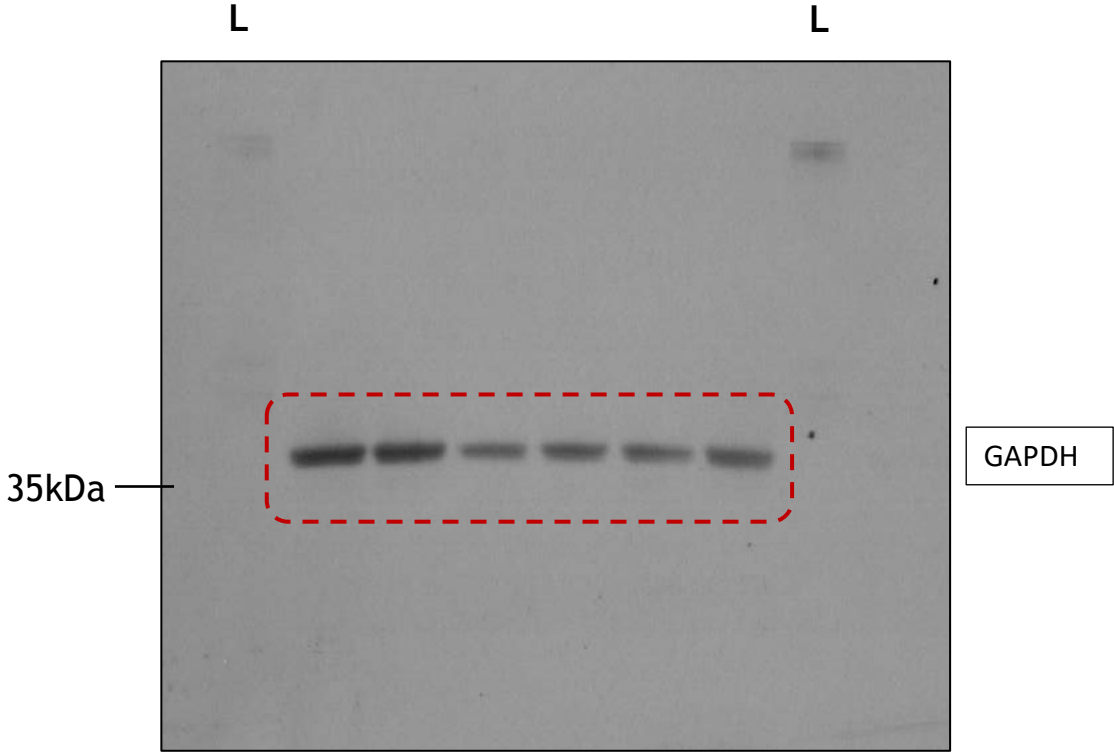
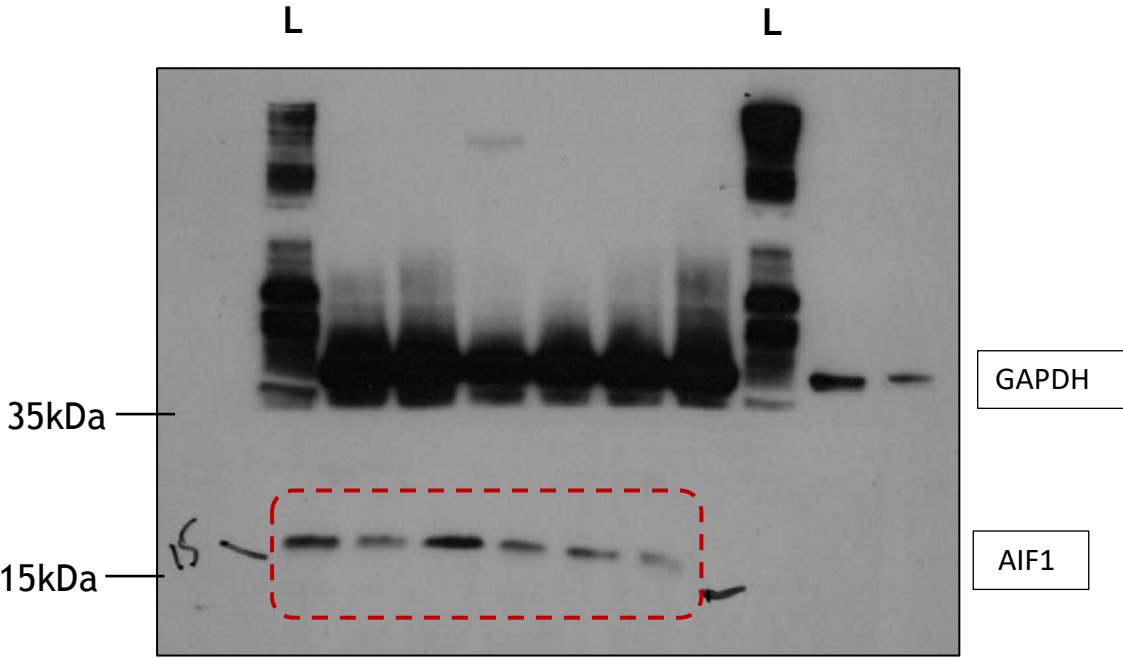
 Cropped along the dotted lines      L Ladder

Uncropped blots for Supplementary Figure 4a- Brain



Cropped along the dotted lines      L Ladder

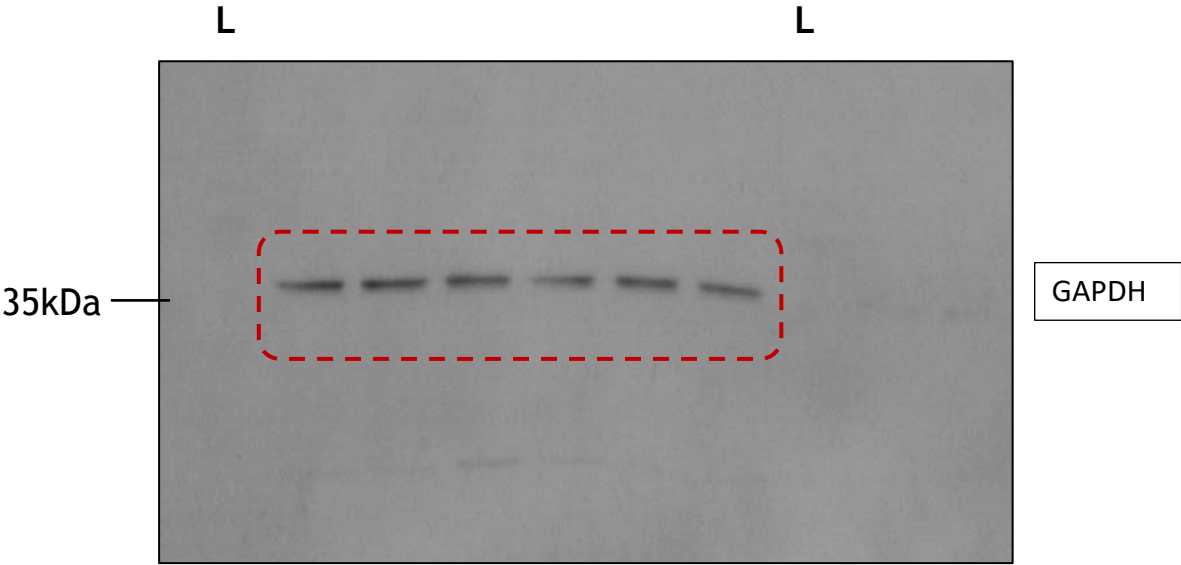
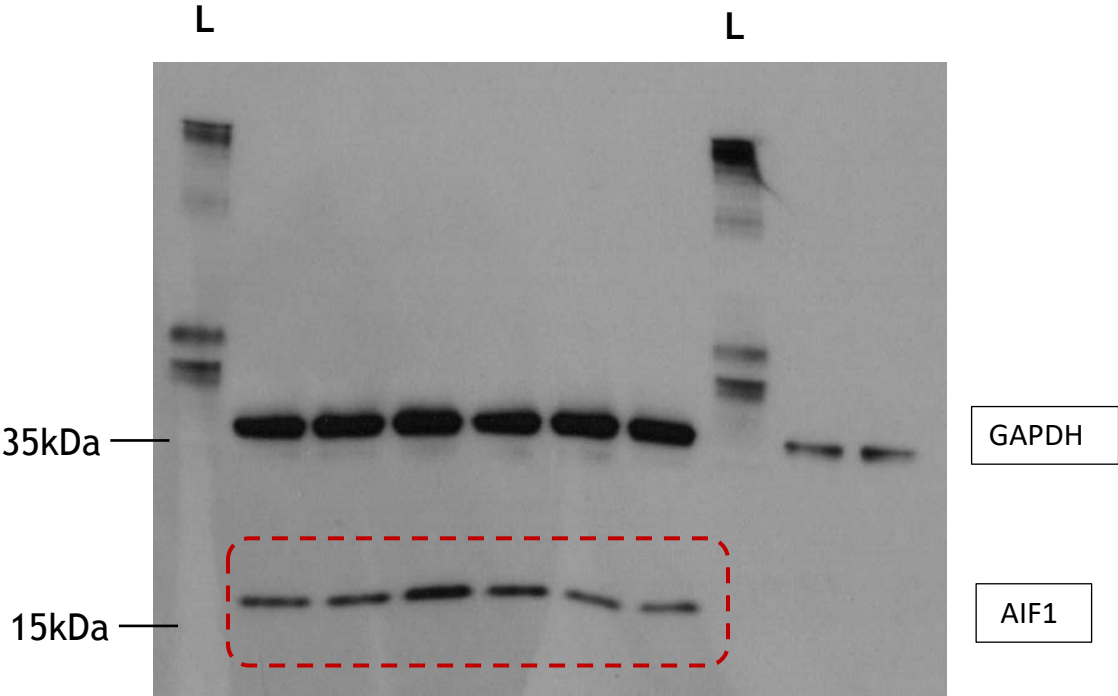
Uncropped blots for Supplementary Figure 4a- Liver




Cropped along the dotted lines

L Ladder

Uncropped blots for Supplementary Figure 4a- Spleen



 Cropped along the dotted lines

**L** Ladder