

Supplemental Figure Legends

Supplemental Figure 1: Characterization of Tadipo mice on chow diet

a-b) qPCR analysis of TLR4 and IL1 β mRNA expression in various tissue normalized to TLR4 $^{fl/fl}$. **c)** 10 week body weight follow up (n=10~15). **d)** oral glucose tolerance test (2.5 mg/g BW). **e)** insulin tolerance test (0.75 mU/g BW). **f)** triglyceride clearance test (15ml/kg BW of 20% intralipid) (n=4). **g)** glucose infusion rate during hyperinsulinemic euglycemic clamp (n=4~5). **h)** glucose infusion rate after acute lard infusion during hyperinsulinemic euglycemic clamp (n=4~5). *p<0.05 compared to TLR4 $^{fl/fl}$. Data are presented as means \pm SEM

Supplemental Figure 2: β 3 adrenergic receptor stimulation and average adipocyte size

- a)** Circulating NEFA (non-esterified fatty acid) (left) and triglyceride concentrations (right) in male TLR4 ff and Tadipo mice during a β -3 adrenergic agonist sensitivity test (1 mg per kg body weight) CL316,243 intraperitoneally, n = 5 TLR4 ff , n=10 Tadipo) on chow diet.
- b)** Circulating NEFA (left) and triglyceride concentrations (right) in male TLR4 ff and Tadipo mice during a β -3 adrenergic agonist sensitivity test (n = 6 TLR4 ff , n=7 Tadipo) after chronic HFD.
- c)** Average adipocytes sized (n \geq 105).

Supplemental Figure 3: Insulin levels during hyperinsulinemic euglycemic clamps

- a-c)** plasma insulin measured at basal and steady state during hyperinsulinemic euglycemic clamp after 3 week HFD, chronic HFD, and lard infusion. **d-f)** hepatic glucose production during hyperinsulinemic euglycemic clamp after 3 week HFD, chronic HFD, and lard infusion. *p<0.05 compared to TLR4 $^{fl/fl}$. Data are presented as means \pm SEM

Supplemental Figure 4: Plasma and tissue sphingolipid analysis with and without HFD challenge

- a-b)** plasma. **c-d)** liver. **e-f)** sWAT ceramide level after 8 weeks Dox chow or HFD induction.

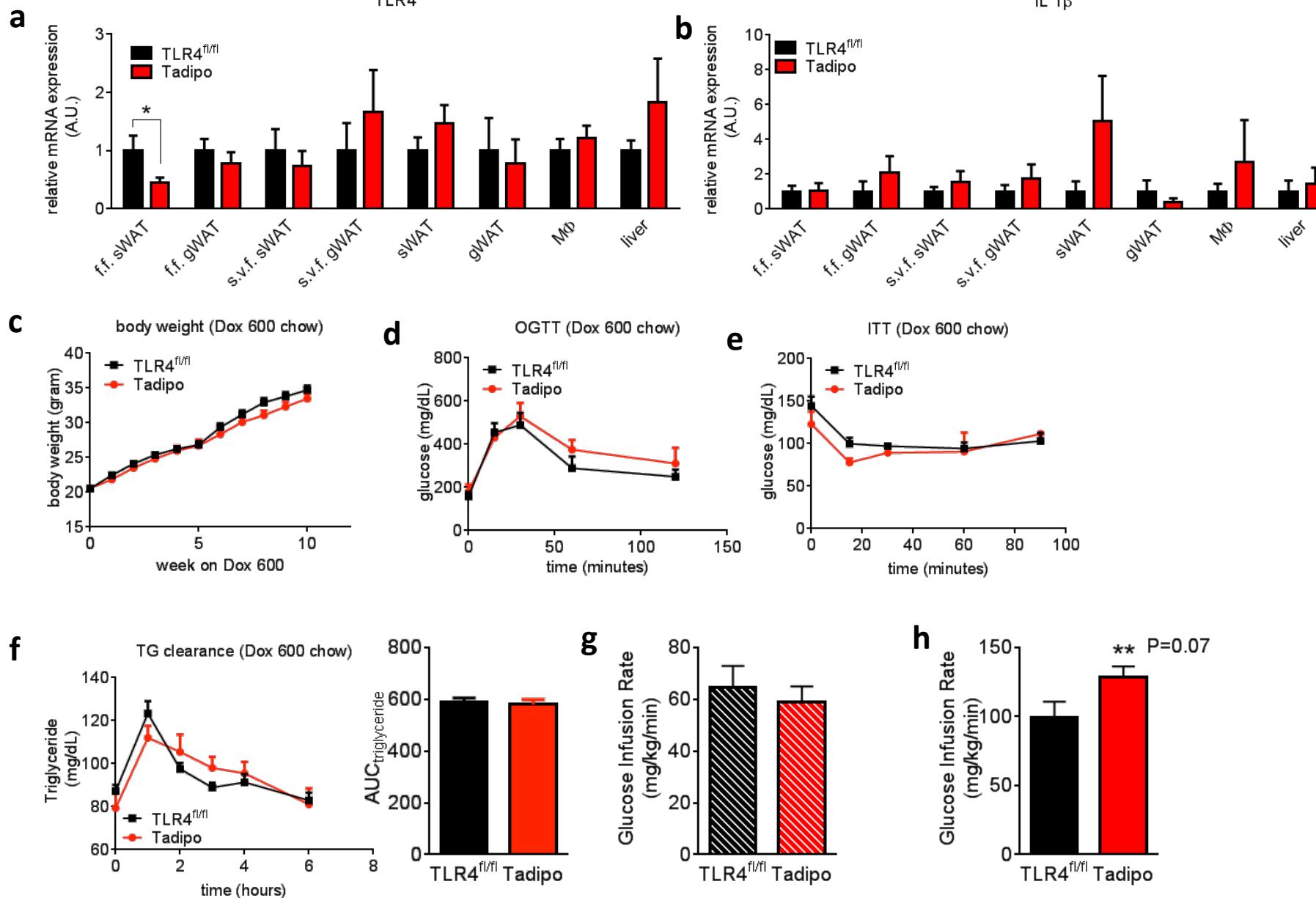
Supplemental Table 1: Primer sets for qPCR analysis

| Gene | Forward primer | Reverse primer |
|--------------|---------------------------|---------------------------|
| Acc1 | TGGACAGACTGATCGCAGAGAAAG | TGGAGAGCCCCACACACA |
| Acox 1 | TCCCGATCTGCGCAAGGAGC | CTGGTGAAGCAAGGTGGCA |
| β-actin | TACCACAGGCATTGTGATGG | TTTGATGTCACGCACGATT |
| Fas | GCTGCGGAAACTTCAGGAAAT | AGAGACGTGTCACTCCTGGACTT |
| IL 1 β | CAACCAACAAGTGATATTCTCCATG | GATCCACACTCTCCAGCTGCA |
| Scd1 | CCGGAGACCCCTAGATCGA | TAGCCTGTAAAAGATTCTGCAAACC |
| SREBP1-c | GGAGCCATGGATTGCACATT | CAGGAAGGCTTCCAGAGAGG |
| TLR4 | GACACCAGGAAGCTTGAATCC | TGATCCATGCATTGGTAGGTAA |

Supplemental Table 2: Primer sets to screen TLR4 deletion

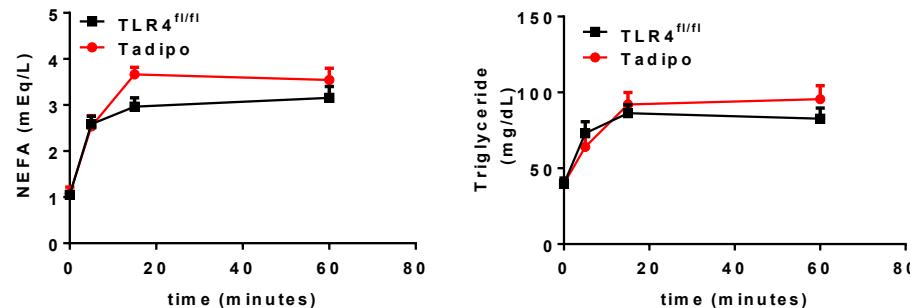
| Gene | Forward primer | Reverse primer |
|--------|--------------------------|--------------------------|
| fl/fl | GACTTGAGAGTTATGTAACACCTG | TCCTATAGACCAGTCTGGCCTTAA |
| Tadipo | GACTTGAGAGTTATGTAACACCTG | GTGGCTATGTTCCAGTTGAATG |

Supplemental Figure 1

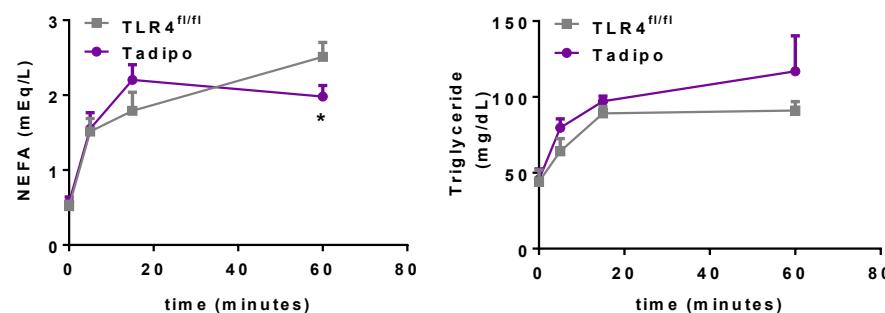


Supplemental Figure 2

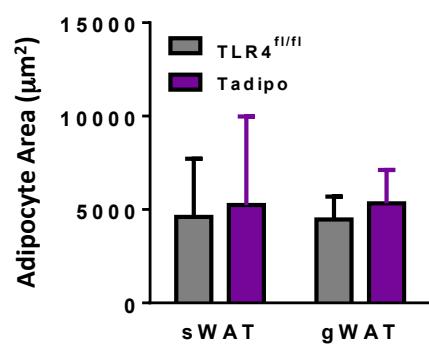
a β 3 adrenergic receptor agonist-induced lipolysis
(Dox 600 Chow)



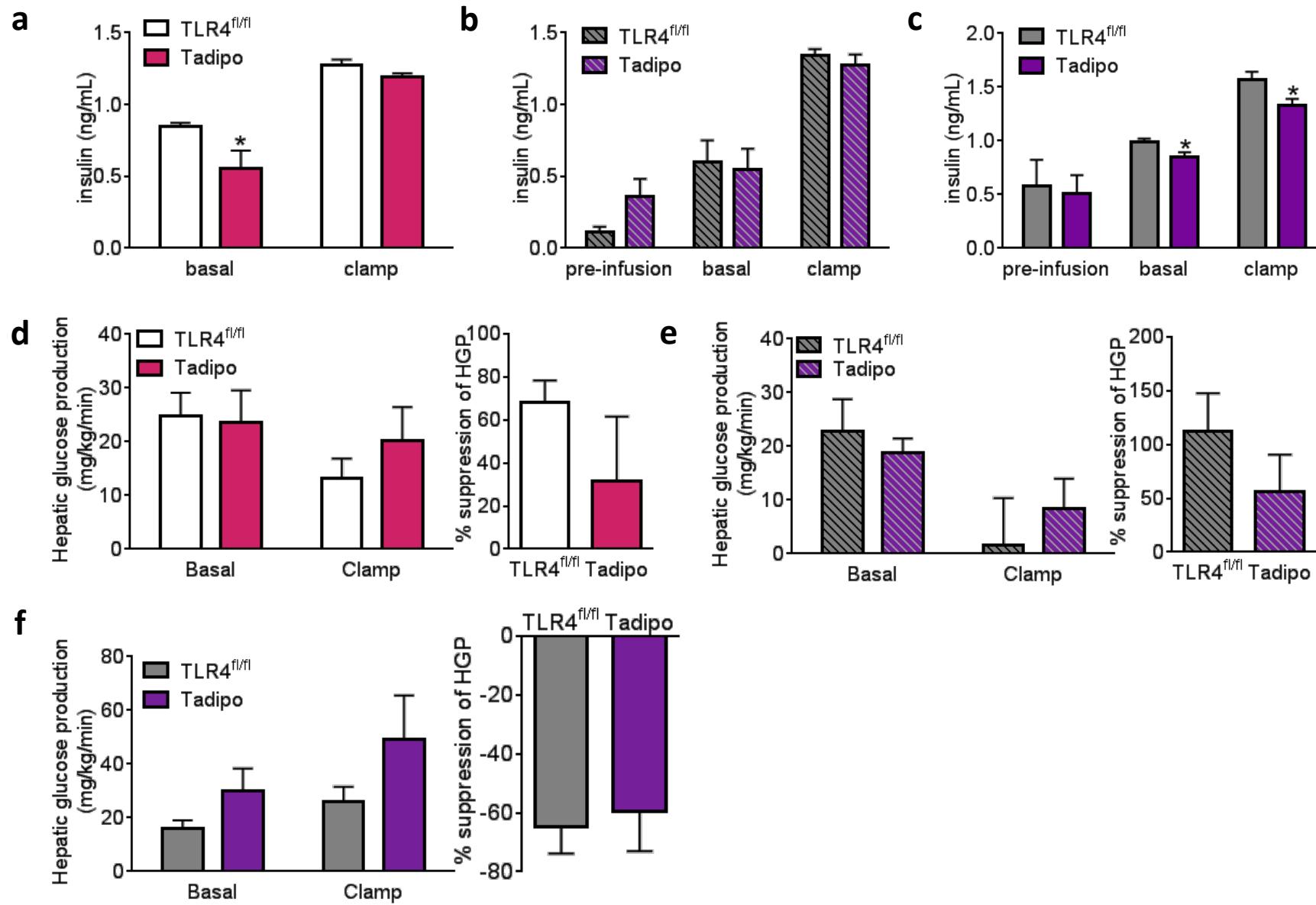
b β 3 adrenergic receptor agonist-induced lipolysis
(Dox 600 HFD)

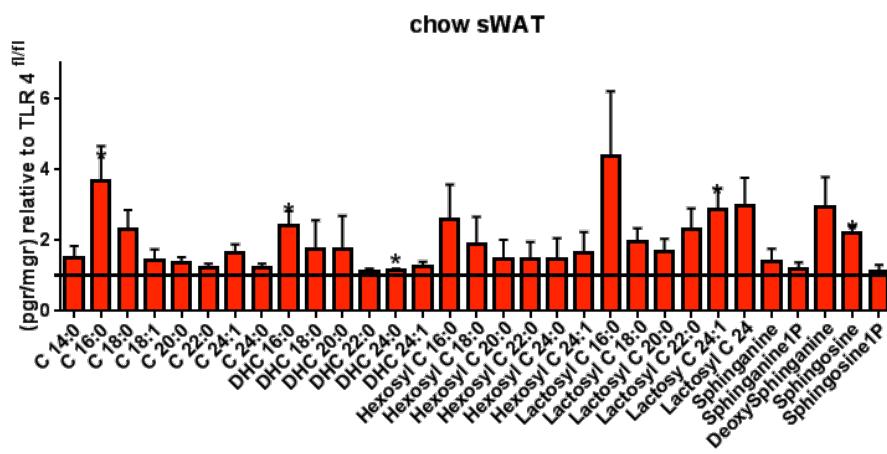


c Adipocyte Area (μm^2)



Supplemental Figure 3



e**f**