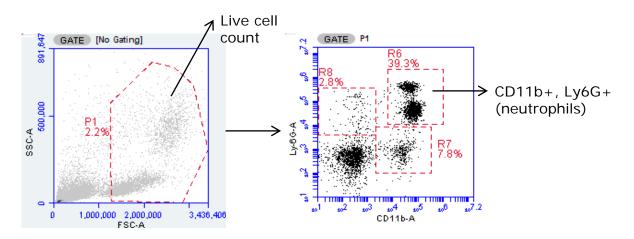
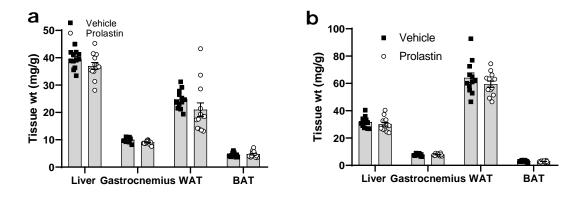
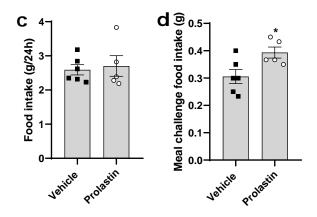
SUPPLEMENTAL DATA

Supplementary figure 1

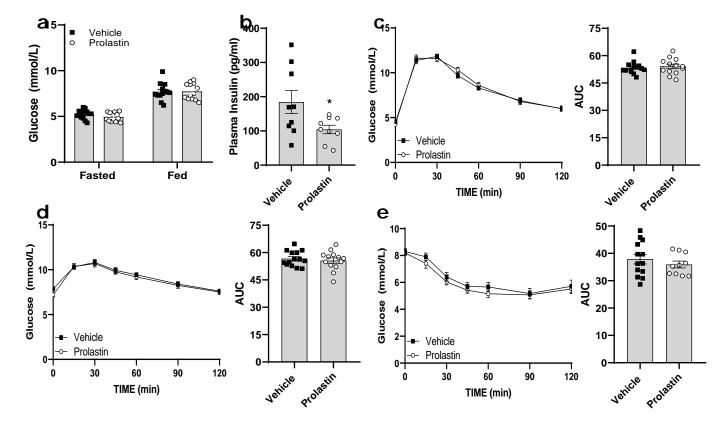


Supplementary Figure 1. FACS gating strategy for identification of neutrophil content in blood cells. Gate P1 contains all cells that were deemed acceptable immune cell events as per forward and side scatter criteria. Within P1, the R7 gate contains all cells that were Cd11b+, Ly6G-whilst the R8 gate reflects all events that were Cd11b-, Ly6G+. The R6 gate contained all cell events that were both Cd11b+, Ly6G+ which were identified as neutrophils for the present analyses.

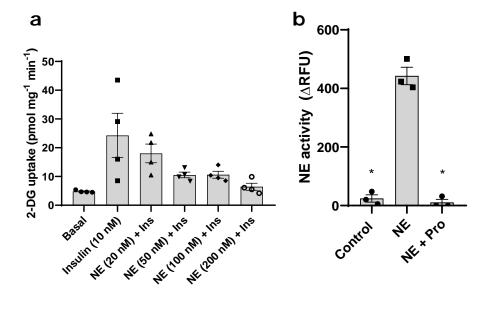




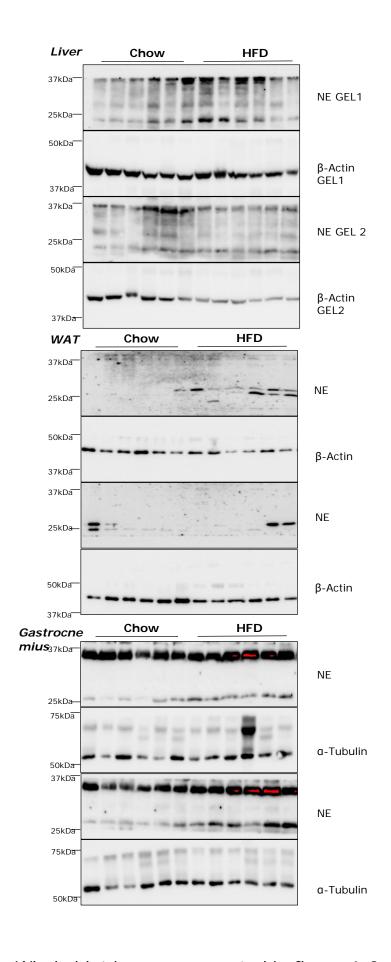
Supplementary Figure 2: Effect of Prolastin on body composition, diet consumption responses to a single meal. Body composition for chow (a) and high fat diet fed mice respectively (b). (c) High fat diet fed mice feed consumption per day, and in an acute 3 hour feed (d) following an overnight fast. Data shown as means \pm SE. (a-d) n=12 per group except HFD + Prolastin group n=13. Significance for (a-d) was determined using two tailed student's t-test. *p < 0.05.



Supplementary Figure 3. The effect of Prolastin on blood glucose homeostasis, insulin sensitivity, and liver function in standard chow fed mice. (a) Overnight fasting and fed blood glucose, (b) 3h fasted plasma insulin, (c-e) temporal glucose in response to GTT (overnight fast followed by 1mg/g oral gavage), PTT (4h fast followed by 1mg/g sodium pyruvate ip) and ITT (4h fast followed by 0.5mU/g insulin ip) respectively. Data shown as means \pm SE, n=12 per group. Significance for (a, b) was determined using two tailed student's t-test. *p < 0.05. Significance for (c-e) was determined using two-way repeated measures ANOVA with Prolastin as a between-group factor and time as repeated factor. *p <0.05 between groups at respective timepoints.

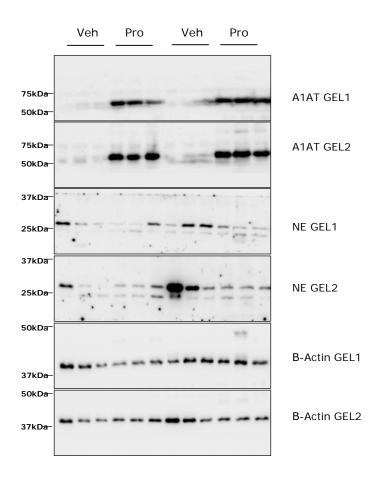


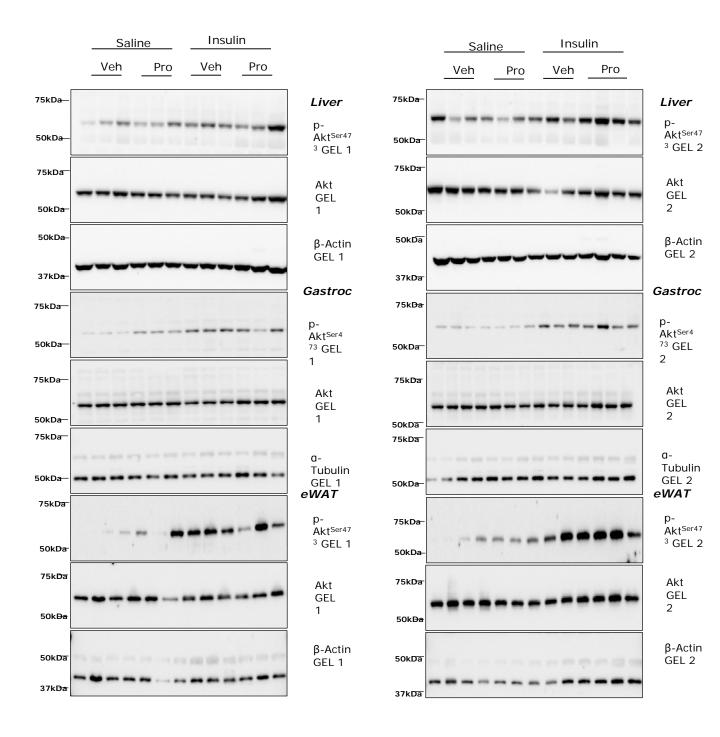
Supplementary Figure 4: NE impairs insulin stimulated glucose uptake in 3T3L1 adipocytes. (a) Dose-response of NE impairing glucose uptake and (b) NE maintains activity in cell media. Data shown as means \pm SE. (b) Significance was determined via one way ANOVA. *p<0.05 between NE and other groups.

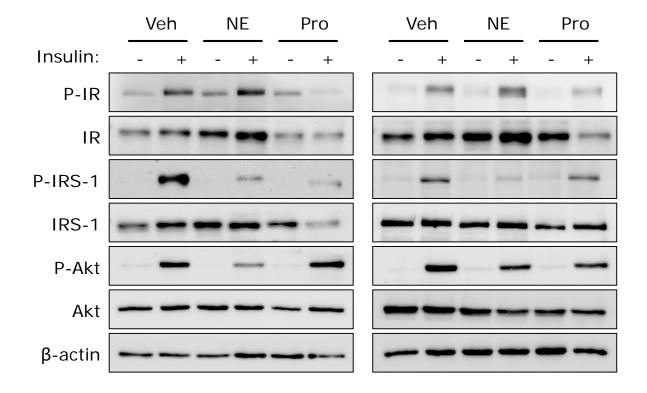


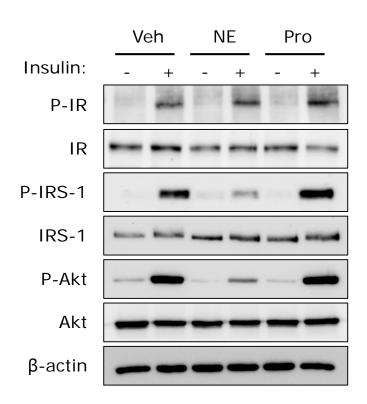
Supplementary Figure 5-9: Whole blot images presented in figure 1-2 and 4-6 respectively

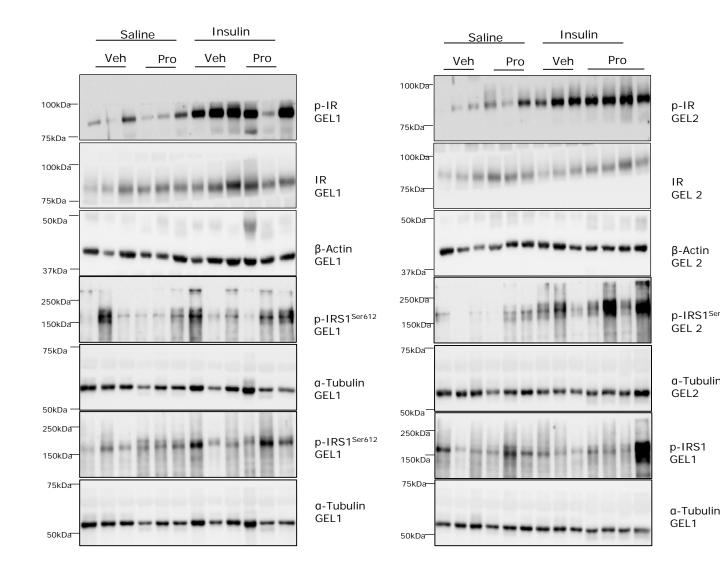
WAT

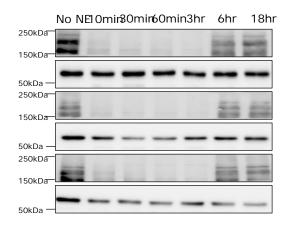












Supplementary table 1: Antibody list

Antibody	Company	Catalog	Dilution
CD11b	BD Biosciences	564985	1:100
LY6G	Biolegend	127605	1:50
Neutrophil Elastase	Abcam	ab68672	1:1000
A1AT	ThermoFisher Scientific	PA5-16661	1:2500
Akt ^{Ser473}	Cell Signalling	CST9271	1:1000
Akt	Cell Signalling	CST9279	1:1000
p-ERK1/2	Cell Signalling	CST9101	1:2000
ERK1/2	Cell Signalling	CST9107	1:2000
p-P38	Cell Signalling	CST4511	1:2000
P38	Cell Signalling	CST9212	1:2000
p-IR1β	Cell Signalling	CST3024	1:1000
IR	Cell Signalling	CST3020	1:1000
p-IRS-1	Cell Signalling	CST3203	1:1000
IRS-1	Cell Signalling	CST2382	1:1000
α-Tubulin	Sigma-Aldrich	T9026	1:5000
β-Actin	Sigma-Aldrich	A2228	1:5000
Secondary Mouse	ThermoFisher Scientific	G21240	1:10000
Seocondary Rabbit	ThermoFisher Scientific	G21234	1:10000
Secondary Rat	ThermoFisher Scientific	A10549	1:5000

Supplementary table 2: qPCR primer sequences

Gene name	Forward primer	Reverse primer
36B4	GGCCCTGCACTCTCGCTTTC	TGCCAGGACGCGCTTGT
B2m	TGACCGGCCTGTATGCTATC	GGCGGGTGGAACTGTGTTAC
Gapdh	CTTTGGCATTGTGGAAGGGC	CAGGGATGATGTTCTGGGCA
Itgam (Cd11b)	TGGCCTATACAAGCTTGGCTTT	AAAGGCCGTTACTGAGGTGG
Ly6G	GGAGGGCTGAGAGAAAGTAAA	GCTGCACAGATAAAACTTCCTC
Elane	AGGCGTGGAGGTCATTTCTG	TGACCGGAAATTTAGGCCGT
Cd68	ACTGGTGTAGCCTAGCTGGT	CCTTGGGCTATAAGCGGTCC
F4/80	CTTTGGCTATGGGCTTCCAGTC	GCAAGGAGGACAGAGTTTATCGTG
ChREBP	CTGGGGACCTAAACAGGAGC	GAAGCCACCCTATAGCTCCC
G6P	TGCAAGGGAGAACTCAGCAA	GGACCAAGGAAGCCACATG
GK	CCCTGAGTGGCTTACAGTTC	ACGGATGTGAGTGTTGAAGC
PEPCK	GTGTTTGTAGGAGCAGCCATGAGA	GCCAGTGGGCCAGGTATTTG
Tnf-α	CCCACACCGTCAGCCGATTT	GTCTAAGTACTTGGGCAGATTGACC
Il-6	TCCTCTCTGCAAGAGACTTCC	TTGTGAAGTAGGGAAGGCCG
Mcp-1	AGGTCCCTGTCATGCTTCTGG	AGGAGCTGTCATTAGGGACATC
IL-1β	GCCACCTTTTGACAGTGATGAG	GACAGCCCAGGTCAAAGGTT
Cd206	GGCTGATTACGAGCAGTGGA	CATCACTCCAGGTGAACCCC
Cd163	TCTCCTGGTTGTAAAAGGTTTGT	CAGTTGTTTTCACCACCCGC