SUPPLEMENTARY DATA

Supplementary Table 1. Quantitative PCR Primer sequences:

Acc	TAA CTC				
Acox	GTC GGT				
Arg-1	ATG GCT				
36b4	AGA GTT				
Cd11c	ACA GCC				
Chga	AGG CAG				
Cpt1	CAG GAG				;
Gapdh	TAT GTC				
<i>G6Pase</i>	GTT CGA				
IL-1b	AAA CTT				
IL-6	CCA ACT				
IL-10	TGA TCA				
IL-12p40	CCA AGA				
iNos	GAG TCC				
Mcp-1	AGG GCT				

SUPPLEMENTARY DATA

Mg1-1	F:	ATG	ATG	TCT	GCC	AGA	GAA	CC		
	R:	ATC	ACA	GAT	TTC	AGC	AAC	CTT	A	
Mg1-2	F:	CAG	AAC	TTG	GAG	CGG	GAA	GAG		
	R:	TTC	TTG	TCA	CCA	TTT	CTC	ATC	TCC	Т
Pepck	F:	CTG	CAT	AAC	GGT	CTG	GAC	TTC		
	R:	CAG	CAA	CTG	CCC	GTA	CTC	С		
Srebp-1c	F:	GGA	GCC	ATG	GAT	TGC	ACA	TT		
	R:	CTT	CCA	GAG	AGG	AGG	CCA	G		
${f Tnf}lpha$	F:	CCA	GAC	CCT	CAC	ACT	CAG	ATC		
	R:	CAC	TTG	GTG	GTT	TGC	TAC	GAC		
Ym1	F:	GGG	CAT	ACC	TTT	ATC	CTG	AG		
	R:	CCA	CTG	AAG	TCA	TCC	ATG	TC		

Supplementary Figure 1. GTT and ITT in mice with C57BL/6 genetic background. Three month old WT and $\it Chga$ -KO mice were fed HFD for 12 weeks. Body weight-matched mice were fasted for 12 hrs and subjected to IP-GTT (A) and ITT (C) and the corresponding AUCs for glucose excursions were determined (B&D). (E) Four month old $\it Chga$ -KO mice were fed HFD for 12 months, fasted for 12 hrs, injected with PST (5 μ g/g BW, IP) at -30 min and subjected to GTT after injecting glucose at 0 min. *: P<0.05, **: p<0.01, ***: p<0.001.

SUPPLEMENTARY DATA

