

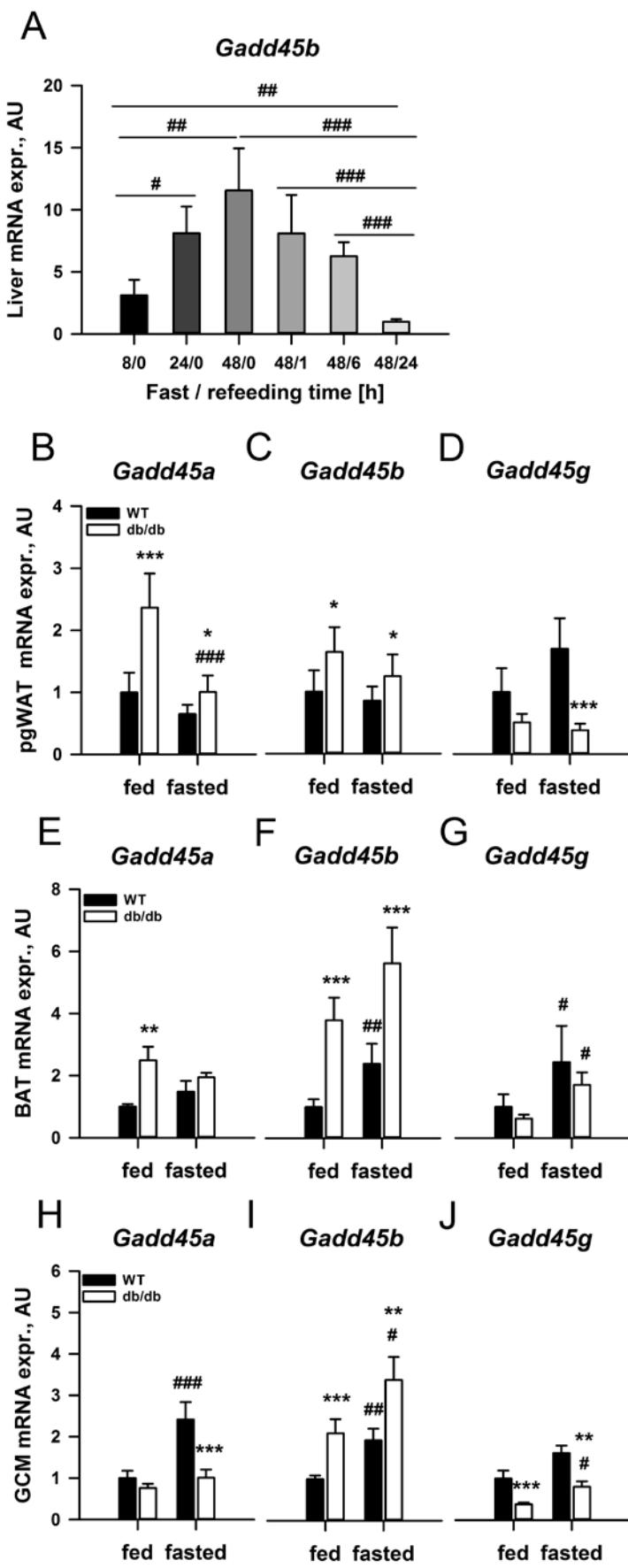
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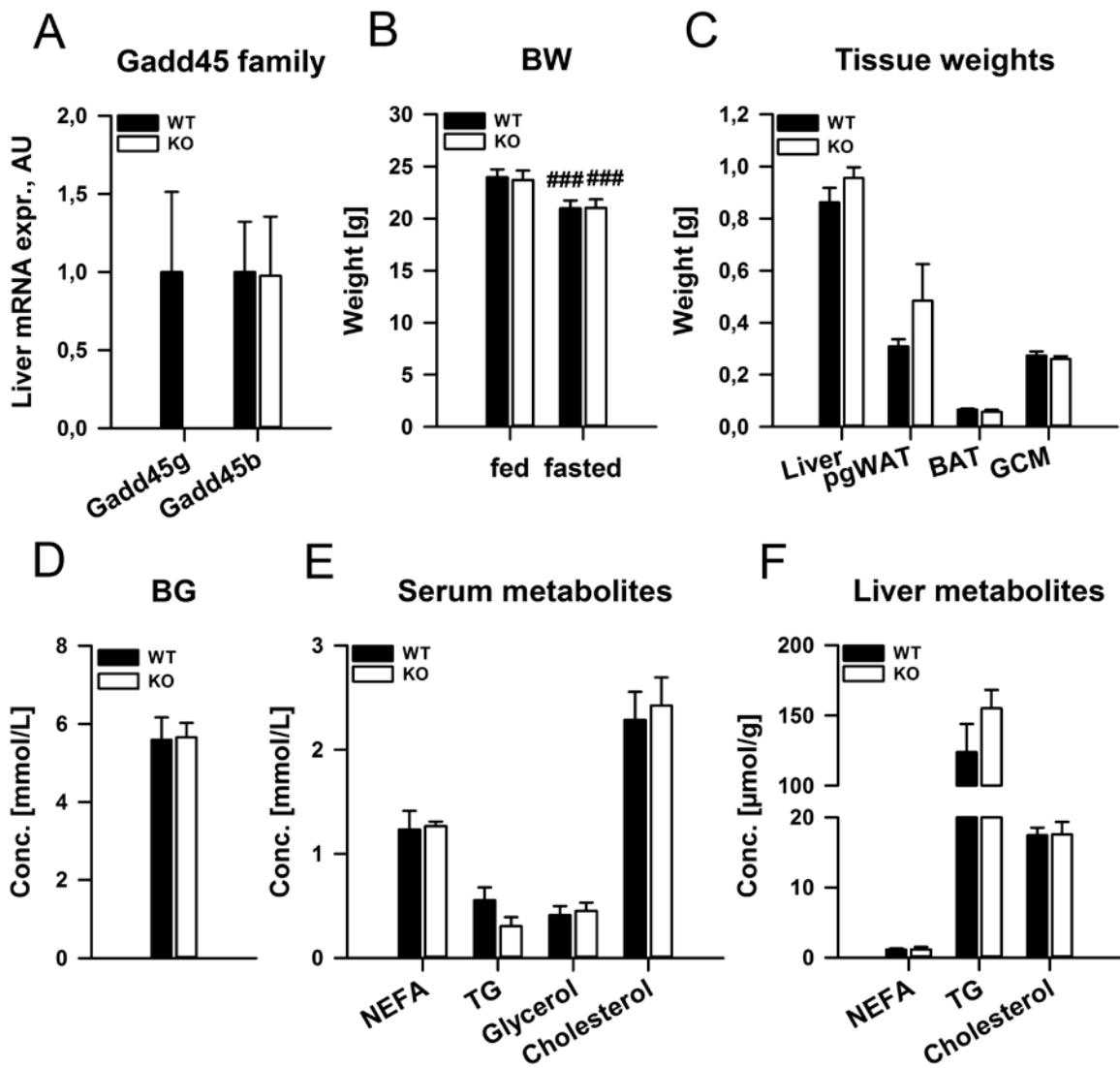
Appendix Figure S2

Appendix Figure S3

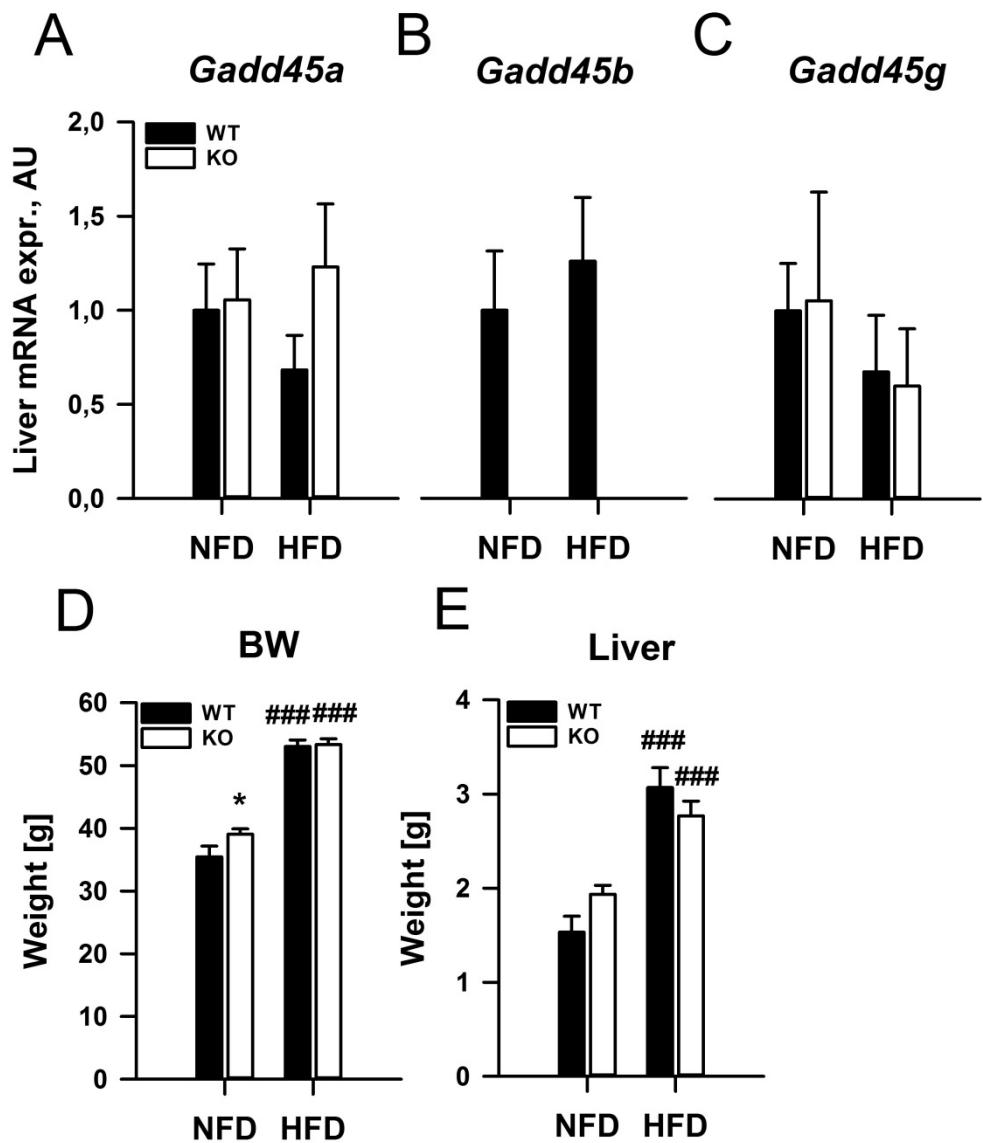
Appendix Table 1



**Figure S1. Liver hepatocyte GADD45b expression is consistently dysregulated upon starvation in several mouse models of metabolic dysfunction.** Male, 12wk old C57Bl/6J mice were fasted for 8, 24, or 48h or fasted for 48h and subsequently refed for 1, 6, or 24h. Liver *Gadd45b* mRNA expression was measured (A) (n = 4/group). Male, 12wk wildtype (WT; C57bl/6J) or obese/diabetic (*db/db*; BKS.Cg-m<sup>+/+</sup> Lepr DB/J) were fed *ad libitum* (fed) or fasted for 24h (fasted) (n=4/group). mRNA expression of Growth Arrest and DNA Damage inducible 45 alpha (*Gadd45a*; B, E, H), beta (*Gadd45b*; C, F I) and gamma (*Gadd45g*; D, G, J) was measured from perigonadal white adipose (B-D), intrascapular brown adipose (E-G) and gastrocnemius complex muscle (H-J). Data are mean ± SEM. Effect of genotype/age/fraction, \* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001. Effect of nutritional state: # p < 0.05, ## p < 0.01, ### p < 0.001. The statistical test used and respective p-value outputs can be found in Appendix Table S1.



**Figure S2. Systemic GADD45 $\gamma$  deletion does not affect metabolic homeostasis during fasting.** Female, GADD45 $\gamma$ +/+ (WT) or GADD45 $\gamma$ -/- (KO) were fasted for 24h (fasted) ( $n = 5$ /group) with blood samples taken before and during fasting. Liver mRNA expression of *Gadd45* family members (A). Body weight (B) before and after fasting and fasted tissue weights (C) including perigonadal white adipose, brown adipose and gastrocnemius complex muscle tissues. Blood glucose (D) as well as serum (E) and liver (F) metabolite concentrations, including non-esterified fatty acids, triglycerides, glycerol and cholesterol. Data are mean  $\pm$  SEM. Effect of genotype, \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$ . Effect of nutritional state: #  $p < 0.05$ , ##  $p < 0.01$ , ###  $p < 0.001$ . The statistical test used and respective p-value outputs can be found in Appendix Table S1.



**Figure S3. Effects of chronic dietary treatment on metabolic homeostasis in mice with systemic GADD45 $\beta$  deletion.** GADD45 $\beta$   $+/+$  (WT) or GADD45 $\beta$   $-/-$  (KO) were chronically fed a normal- (NFD) or high (HFD) fat diet (n= 6-9/group) and sacrificed in the *ad libitum* fed state. Liver mRNA expression of *Gadd45* family members (A-C). Body weight (D), and liver mass (F). Data are mean  $\pm$  SEM. Effect of genotype, \* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001. Effect of nutritional state: # p < 0.05, ## p < 0.01, ### p < 0.001. The statistical test used and respective p-value outputs can be found in Appendix Table S1.

Appendix Table S1. P values from statistical tests. NP: not performed.

Figure 1	nutrition	Genotype/strain/age	Post-hoc testing				test performed	
			fed		fasted			
			wt/NZB/ young vs db/NZO/aged	wt/NZB/ young vs db/NZO/aged	wt/NZB/young	db/NZO/aged		
A	0.004	0.043	0.033	0.233	0.033	0.014	0.031	
B	0.158	0.243	0.012	0.042	0.016	0.038	0.025	
C	>0.001	0.078	0.046	0.672	0.043	>0.001	>0.001	
D	0.077	0.011	0.003	0.088	>0.001	>0.001	0.342	
E	0.006	0.044	>0.001	0.279	>0.001	>0.001	0.149	
F	0.047	0.027	0.044	0.389	0.033	0.023	0.231	
G	0.090	0.113	0.009	0.044	0.028	0.031	0.428	
H	>0.001	0.222	0.031	0.385	0.024	>0.001	0.008	
I	0.088	0.197	0.027	0.179	0.008	0.009	0.527	
J	0.012	0.333	0.021	0.428	>0.001	>0.001	0.200	
K	0.009	0.198	0.012	0.371	>0.001	>0.001	>0.001	
L	0.021	0.033	0.028	>0.001	>0.001	0.008	0.006	
M	>0.001	0.081	0.009	0.287	>0.001	>0.001	>0.001	
N	0.031	0.019	0.003	0.471	>0.001	0.006	0.197	
O	0.002	0.208	0.019	0.309	>0.001	>0.001	0.06	
P	0.005	0.081	0.044	0.618	0.007	0.021	0.009	
Q	0.002	0.194	0.049	0.777	0.418	0.033	0.028	
R	0.018	0.090	0.037	0.447	>0.001	0.088	0.006	
NEFA	Genotype		BSA	BSA-NEFA	wt	db/db		
			wt vs db/db	wt vs db/db	BSA vs NEFA	BSA vs NEFA		
S	0.033	0.011	0.032	0.009	>0.001	0.038	0.008	

Figure EV1	nutrition	Genotype/strain/age	Post-hoc testing				test performed	
			fed		fasted			
			wt/NZB/ young vs db/NZO/aged	wt/NZB/ young vs db/NZO/aged	wt/NZB/young	db/NZO/aged		
A	0.001	0.005	0.008	0.011	0.023	0.021	0.018	
B	0.009	0.222	0.014	0.398	0.121	>0.001	>0.001	
C	0.030	0.281	0.039	0.444	0.319	0.007	0.018	
D	0.039	0.315	0.049	0.199	0.551	0.051	0.009	
E	0.009	0.008	0.007	0.040	0.044	0.033	0.028	
F	0.033	0.071	0.088	NP	NP	NP	NP	
G	0.055	0.071	0.033	0.388	0.021	0.081	0.050	
H	0.221	0.038	0.111	NP	NP	NP	NP	
I	0.002	0.091	0.021	0.424	>0.001	>0.001	0.033	
J	0.379	0.559	0.394	NP	NP	NP	NP	
K	0.021	0.189	0.172	NP	NP	NP	NP	
L	0.465	0.444	0.451	NP	NP	NP	NP	

Figure 2	nutrition	Genotype/strain/age	Post-hoc testing				test performed	
			fed		fasted			
			wt/NZB/ young vs db/NZO/aged	wt/NZB/ young vs db/NZO/aged	wt/NZB/young	db/NZO/aged		
A	0.033	0.066	0.031	0.111	>0.001	0.008	0.189	
B	0.003	0.004	0.008	0.089	>0.001	>0.001	0.009	
C	0.069	0.077	0.002	0.491	>0.001	>0.001	0.321	
D	0.444	0.111	0.044	0.099	0.048	0.122	0.366	
E	0.033	0.008	0.021	0.088	0.033	0.007	0.086	
F	0.005	0.199	0.033	0.077	0.336	0.021	0.009	
G	>0.001	0.111	0.028	0.217	>0.001	>0.001	>0.001	
H	0.033	0.041	0.015	0.039	>0.001	>0.001	0.199	
I	0.044	0.056	0.027	0.035	0.009	>0.001	0.289	
J	0.098	0.181	>0.001	0.389	>0.001	>0.001	0.225	
K	0.153	0.210	>0.001	0.255	0.008	0.005	0.351	
L	0.073	0.222	>0.001	0.400	0.007	0.006	0.492	

Figure S1	nutrition	Genotype/strain/age	Post-hoc testing				test performed	
			fed		fasted			
			wt/NZB/ young vs db/NZO/aged	wt/NZB/ young vs db/NZO/aged	wt/NZB/young	db/NZO/aged		
8/0 vs 24/0	8/0 vs 48/0	8/0 vs 48/24	48/24 vs 48/0	48/24 vs 48/1	48/24 vs 48/6			
A	0.039	0.007	0.002	>0.001	>0.001	>0.001	One-way ANOVA, Holm-Sidak post-hoc test	
B	0.033	0.008	0.021	>0.001	0.038	0.222	>0.001	
C	0.111	0.012	0.049	0.038	0.044	0.219	0.200	
D	0.288	0.231	0.046	0.051	>0.001	0.191	0.099	
E	0.121	0.037	0.039	0.009	0.318	0.298	0.194	
F	0.030	>0.001	0.022	>0.001	>0.001	0.008	0.099	
G	0.007	0.055	0.044	0.111	0.192	0.035	0.032	
H	0.288	0.391	0.014	0.399	>0.001	>0.001	0.404	
I	0.010	0.009	0.009	>0.001	0.008	0.007	0.020	
J	0.066	0.004	0.017	>0.001	0.007	0.097	0.046	

Figure 3									
		nutrition	genotype		pre-fast vs fast	pre-fast vs. refed	fast vs refed		
A	>0.001	0.398	0.021		>0.001	0.044	>0.001		Two-way RM ANOVA, Holm-Sidak post-hoc test
B	>0.001	0.289	0.012		>0.001	0.181	>0.001		Two-way RM ANOVA, Holm-Sidak post-hoc test
C	>0.001	0.455	0.007		>0.001	0.233	>0.001		Two-way RM ANOVA, Holm-Sidak post-hoc test
		nutrition	genotype		fed	fasted	genotype	genotype	
D	>0.001	0.033	0.044		0.321	0.008	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
E	>0.001	0.358	0.007		0.243	0.198	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
F	0.055	0.256	0.169		NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test
G			0.009						
		time	genotype		0	90	180	270	
H	>0.001	0.007	>0.001		0.288	>0.001	0.098	0.213	Two-way RM ANOVA, Holm-Sidak post-hoc test
I	0.009	0.032	>0.001		0.555	0.008	>0.001	0.121	Two-way RM ANOVA, Holm-Sidak post-hoc test
J	>0.001	0.013	>0.001		0.423	>0.001	0.009	0.333	Two-way RM ANOVA, Holm-Sidak post-hoc test
		time	genotype-diet		0 (WT vs KO HFD)	20 (WT vs KO HFD)	40 (WT vs KO HFD)	60 (WT vs KO HFD)	
K	>0.001	0.041	0.023		0.099	0.008	0.021	0.056	Two-way RM ANOVA, Holm-Sidak post-hoc test
		nutrition	genotype		NFD	HFD	genotype	genotype	
L	>0.001	0.121	0.040		0.222	0.087	0.021	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
M	>0.001	0.044	0.012		0.121	0.032	0.008	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
N	>0.001	0.031	0.009		0.257	0.006	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test

Figure EV2									
		nutrition	genotype		pre-fast vs fast	pre-fast vs. refed	fast vs refed		
A	>0.001	0.350	0.011		>0.001	0.033	0.222		Two-way RM ANOVA, Holm-Sidak post-hoc test
B	>0.001	0.189	0.008		>0.001	0.027	0.005		Two-way RM ANOVA, Holm-Sidak post-hoc test
C	>0.001	0.675	0.023		0.007	0.009	0.229		Two-way RM ANOVA, Holm-Sidak post-hoc test
D				Gadd45a	Gadd45b	Gadd45g			t-test
				0.448	>0.001	0.664			
		nutrition	genotype		fed	fasted	WT	KO	
E	>0.001	0.385	0.012		0.444	0.398	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
H	>0.001	0.251	0.009		0.210	0.206	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
I	0.077	0.222	0.438		NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test
F				Liver	Fat	GCM			t-test
				0.329	0.287	0.401			
G				NEFA	Cholesterol	Glycogen			t-test
				0.201	0.363	0.191			
J				C0	C2	SCAC	MCAC	LCAC	t-test
				0.298	0.329	0.189	0.210	0.427	

Figure S2										
				Gadd45g	Gadd45b					
A				>0.001	0.456					t-test
		nutrition	genotype		fed	fasted	genotype	genotype		
B	>0.001	0.284	0.007		0.429	0.321	fed vs fasted	fed vs fasted		Two-way ANOVA, Holm-Sidak post-hoc test
				Liver	pgWAT	BAT	GCM			
C				0.222	0.198	0.267	0.335			t-test
D				BG						t-test
				0.333						
E				NEFA	TG	Glycerol	Cholesterol			t-test
				0.222	0.397	0.442	0.597			
F				NEFA	TG	Cholesterol				t-test
				0.189	0.398	0.523				

Figure S3									
		nutrition	genotype		NFD	HFD	genotype	genotype	
A	0.345	0.231	0.667		NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test
B	0.287	>0.001	>0.001		>0.001	>0.001	0.324	0.920	Two-way ANOVA, Holm-Sidak post-hoc test
C	0.138	0.409	0.567		NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test
D	>0.001	0.125	>0.001		0.033	0.230	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
E	>0.001	0.238	>0.001		0.099	0.239	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test

Figure 4								
	nutrition	genotype		fed/BSA	fasted/BSA-NEFA	WT	KO	
				WT vs KO	WT vs KO	fed vs fasted/BSA vs NEFA	fed vs fasted/BSA vs NEFA	
A	>0.001	0.087	>0.001	0.234	>0.001	0.022	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
B	>0.001	0.543	>0.001	0.309	0.290	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
C	0.043	0.285	>0.001	0.444	>0.001	0.127	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
D	>0.001	0.105	>0.001	0.287	>0.001	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
	nutrition	fraction		Hepas	NP	fed	fed	
				fasted vs fed	fasted vs fed	Hepas vs NP	Hepas vs NP	
E	>0.001	0.106	0.025	0.030	0.179	0.088	0.197	Two-way ANOVA, Holm-Sidak post-hoc test
	nutrition	miR		fed	fasted	NC miR	G45b miR	
				NC miR vs G45b miR	NC miR vs G45b miR	fed vs fasted	fed vs fasted	
F	>0.001	0.174	>0.001	0.418	0.026	>0.001	0.005	Two-way ANOVA, Holm-Sidak post-hoc test
G	>0.001	0.201	>0.001	0.317	0.011	0.007	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
	AD	genotype		NC	G45b OE	WT	KO	
				WT vs KO	WT vs KO	NC vs G45b	NC vs G45b	
H	0.031	0.055	0.012	0.068	0.266	0.087	0.008	Two-way ANOVA, Holm-Sidak post-hoc test
	nutrition	miR		NFD	HFD	NC miR	G45b miR	
				NC miR vs G45b miR	NC miR vs G45b miR	NFD vs HFD	NFD vs HFD	
I	>0.001	0.197	0.023	0.314	0.451	0.005	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
J	>0.001	0.182	>0.001	0.202	0.028	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
K	>0.001	0.213	>0.001	0.438	0.024	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test

Figure EV3									
	nutrition	fraction	Hepas		NP		fed	fed	
			fasted vs fed		fasted vs fed	Hepas vs NP			
A	0.444	0.014	0.105	NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test	
B	0.600	>0.001	0.104	NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test	
C	0.537	>0.001	0.283	NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test	
D	0.329	>0.001	0.406	NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test	
	nutrition	miR	fed		fasted		NC miR	G45b miR	
			NC miR vs G45b miR		NC miR vs G45b miR				
E	>0.001	>0.001	>0.001	0.003	>0.001	>0.001	>0.001	0.037	Two-way ANOVA, Holm-Sidak post-hoc test
	AD	genotype	NC		G45b OE		WT	KO	
			WT vs KO		WT vs KO				
F	>0.001	0.105	>0.001	>0.001	0.211	>0.001	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
G	0.229	0.381	0.449	NP	NP	NP	NP	NP	Two-way ANOVA, Holm-Sidak post-hoc test
H	0.294	0.087	0.009	0.223	>0.001	0.276	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
	nutrition	miR	NFD		HFD		NC miR	G45b miR	
			NC miR vs G45b miR		NC miR vs G45b miR				
I	>0.001	0.041	>0.001	0.319	0.033	>0.001	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
J	>0.001	0.143	>0.001	0.252	0.128	>0.001	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test
K	>0.001	0.261	>0.001	0.278	0.124	>0.001	>0.001	>0.001	Two-way ANOVA, Holm-Sidak post-hoc test

Figure 5							
	AD	genotype	NC	G45b OE	WT	db/db	
			WT vs db/db	WT vs db/db	NC vs G45b	NC vs G45b	
A	0.030	>0.001	>0.001	>0.001	0.038	0.081	>0.001
B	0.201	>0.001	>0.001	>0.001	0.061	0.286	0.028
C	0.044	>0.001	>0.001	>0.001	0.092	0.099	>0.001
<b>NGT vs T2D</b>							
D		>0.001					Mann-Whitney Rank Sum Test

Figure EV4							
	AD	genotype	NC	G45b OE	WT	db/db	
			WT vs db/db	WT vs db/db	NC vs G45b	NC vs G45b	
A	0.005	0.150	0.065	NP	NP	NP	NP
B	0.122	>0.001	>0.001	>0.001	>0.001	0.124	0.048
C	0.228	0.356	0.437	NP	NP	NP	NP
D	0.264	>0.001	0.008	>0.001	>0.001	0.187	0.235

Figure EV5							
		AD					
A	0.524						t-test
B	0.412						t-test
D genotype							
LC3B	0.242						t-test
p-S6K1	0.358						t-test
p-ERK	0.153						t-test
p-p38	0.204						t-test
p-PERK	0.123						t-test
p-eIF2a	0.423						t-test
FATP2	0.334						t-test
CD36	0.267						t-test
FABP1	0.346						t-test