

Supplementary Figure Legends

Suppl. Figure 1. Hepatic type I interferon signalling does not modulate susceptibility to metabolic disease induced by a high-fat diet. (A) *Ifnar1* and *Ifnar2* expression relative to β -Actin determined by qPCR of 24h-stimulated FL83B hepatocytes and RAW264.7 macrophages. (B) Expression of *Ifnar1* and *Ifnar2* in the liver, adipose tissue and intestine of indicated genotypes. Note that *Alb-Cre;Ifnar1^{fl/fl}* mice specifically delete *Ifnar1* in the liver. (C-E) Mice were exposed to a high-fat or low-fat (control) diet and were analysed after 16 weeks with (C) Body weight in percent at indicated timepoints, (D) liver-to-body-weight ratio and (E) epididymal white adipose tissue to body weight ratio. (F, G) Mice were exposed to a low-fat diet and were analysed after 16 weeks, glucose tolerance test (F), insulin tolerance test (G). (H, I) Mice were exposed to a high-fat diet and were analysed after 16 weeks. Representative images of H&E sections (H) with quantification of steatosis in (I). (J) Expression of lipogenic genes in indicated genotypes fed the MCD diet determined by qPCR and normalised to β -Actin. (K-M) Mice were exposed to a MCD diet for 24 days and body weight at indicated timepoints (K), liver-to-body-weight ratio (L) and plasma ALT (M) were monitored. Data are expressed as mean \pm SEM (A-M), n = 3 (A), n = 7-10 (B-M). One-way ANOVA with Bonferroni correction (A, M), two-way repeated measures ANOVA (C, F, G, K). Unpaired Student's *t*-test (B, D, E, I, J, L). ***P<0.001, ** P<0.01, *P<0.05.

Suppl. Figure 2. Adipose type I interferon signalling does not modulate susceptibility to hepatic disease induced by a MCD diet. (A) Adipose tissue expression of *Ifnar1* in wild-type mice after exposure to a low- or high-fat diet determined by qPCR relative to *c*- β -Actin. (B) Expression of *Ifnar1* and *Ifnar2* in adipose tissue of indicated genotypes determined by qPCR relative to *c*- β -Actin. Note that the remaining *Ifnar1* expression in adipose tissue of *Ifnar1^{Δat}* mice may be explained by non-adipose cells in adipose tissue (e.g. fibroblasts, endothelium or immune cells) that are not affected by *Ifnar1* deletion. (C) Body weight of indicated genotypes on a normal chow diet. (D, E) Non-essential fatty acids (NEFA, D) and triglyceride (E) concentrations in the serum of indicated genotypes. (F-I) Mice were exposed to a MCD diet for 24 days and body weight at indicated timepoints (F), liver-to-body-weight ratio (G) and plasma ALT (H) were monitored. (I) Quantification of steatosis based on H&E sections. Data are expressed as mean \pm SEM (A-I), n = 6-10 (A-I). Two-way repeated measures ANOVA (C, F). Unpaired Student's *t*-test (A, B, D, E, G, I). ***P<0.001, **P<0.01.

Suppl. Figure 3. Intestinal-epithelial type I interferon signalling does not modulate susceptibility to hepatic or metabolic disease on a high-fat or MCD diet. (A, B) Liver-to-body-weight ratio (A) and epididymal white adipose tissue weight (B) of indicated genotypes on a high-fat diet. (C, D) Mice were exposed to a low-fat diet and were analysed after 16 weeks, glucose tolerance test (C), insulin tolerance test (D). (E-G) Mice were exposed to a MCD diet for 24 days and body weight at indicated timepoints (E), liver-to-body-weight ratio (F) and plasma ALT (G) were monitored. Data are expressed

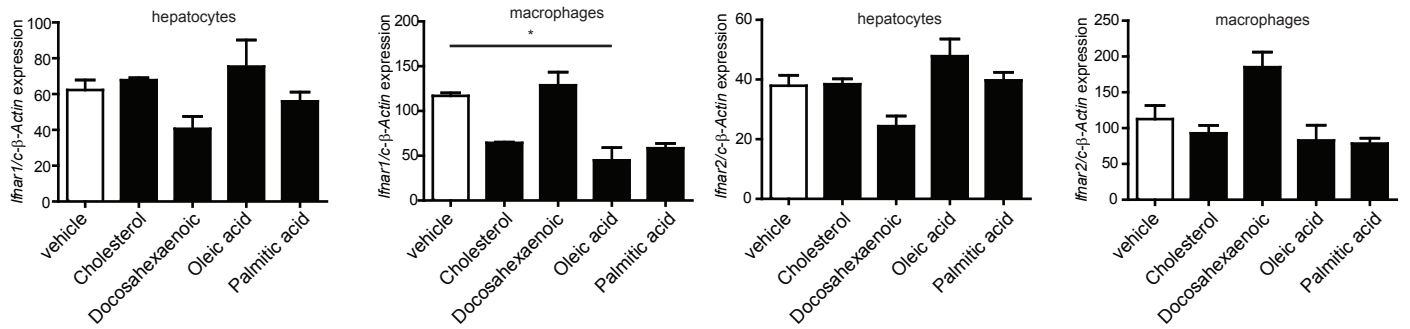
as mean \pm SEM (A-G), n = 9-10 (A-G). Two-way repeated measures ANOVA (C-E). Unpaired Student's *t*-test (A, B, F, G). ***P<0.001.

Suppl. Figure 4. Myelocyte type I interferon signalling does not modulate susceptibility to hepatic or metabolic disease on a high-fat or MCD diet. (A, B) Liver-to-body-weight ratio (A) and epididymal white adipose tissue weight (B) of indicated genotypes on a high-fat diet. (C, D) Mice were exposed to a low-fat diet and were analysed after 16 weeks, glucose tolerance test (C), insulin tolerance test (D). (E-G) Mice were exposed to a MCD diet for 24 days and body weight at indicated timepoints (E), liver-to-body-weight ratio (F) and plasma ALT (G) were monitored. Data are expressed as mean \pm SEM (A-G), n = 9-10 (A-G). Two-way repeated measures ANOVA (C-E). Unpaired Student's *t*-test (A, B, F, G). ***P<0.001.

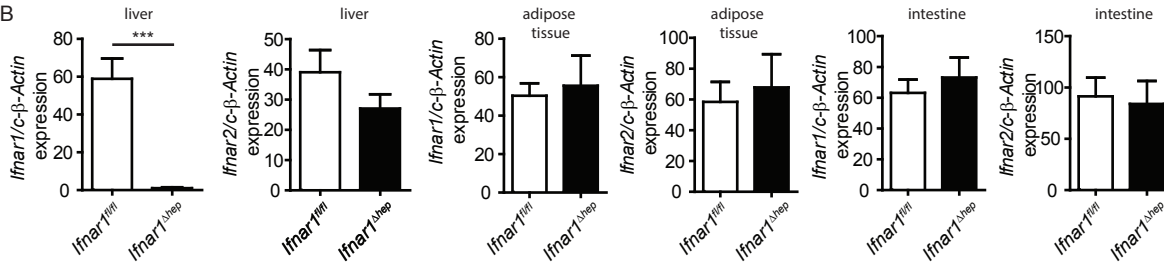
Suppl. Figure 5. Type I interferon regulated gene expression correlates with TNF and IL-6 expression in the liver. (A, B) Expression of type I interferon regulated genes (*OAS1*, *MX1*, *GIP2*, *IFIT1*) were correlated with *IL-6* (A) and *TNF α* (B) expression, respectively, in liver biopsies from obese patients before and 6 months after LAGB. Pearson correlation was performed with indicated r_s coefficient and p-values in the figure. Note that *TNF* expression positively correlates with type I interferon responses (B).

Suppl. Figure 1

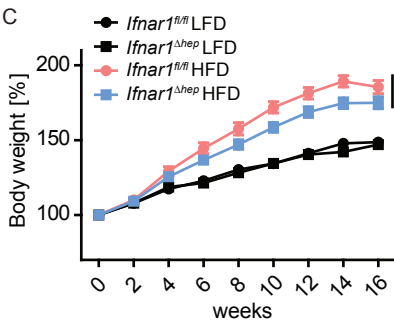
A



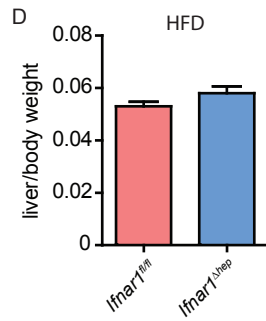
B



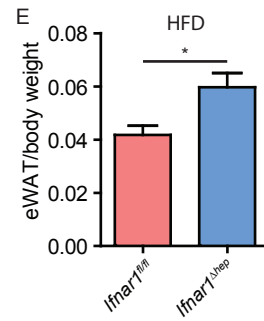
C



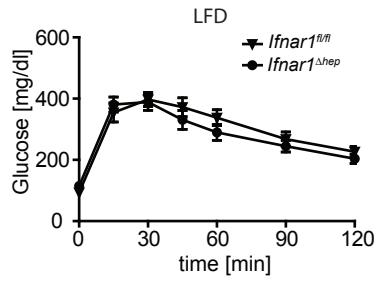
D



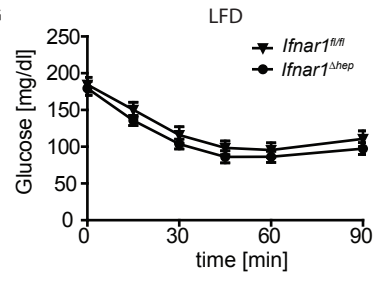
E



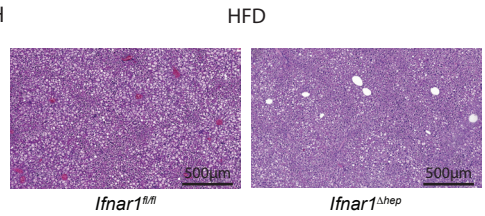
F



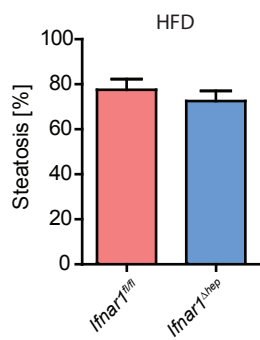
G



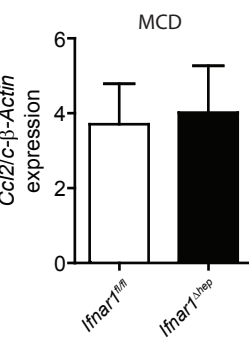
H



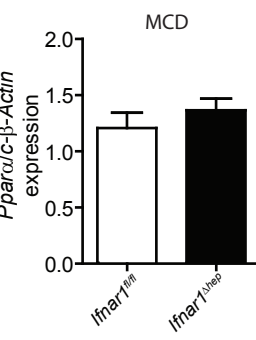
I



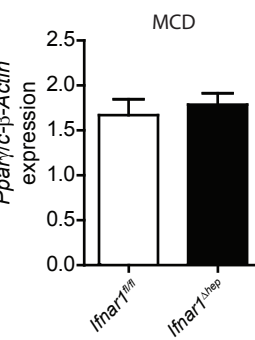
J



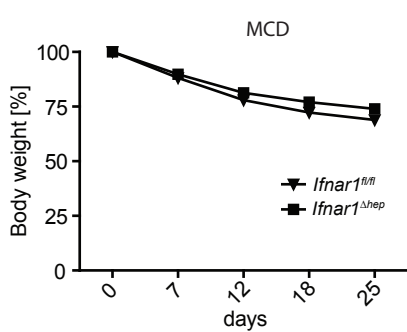
K



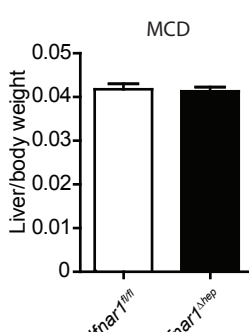
L



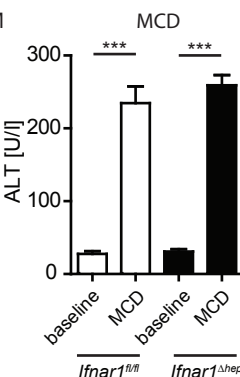
K



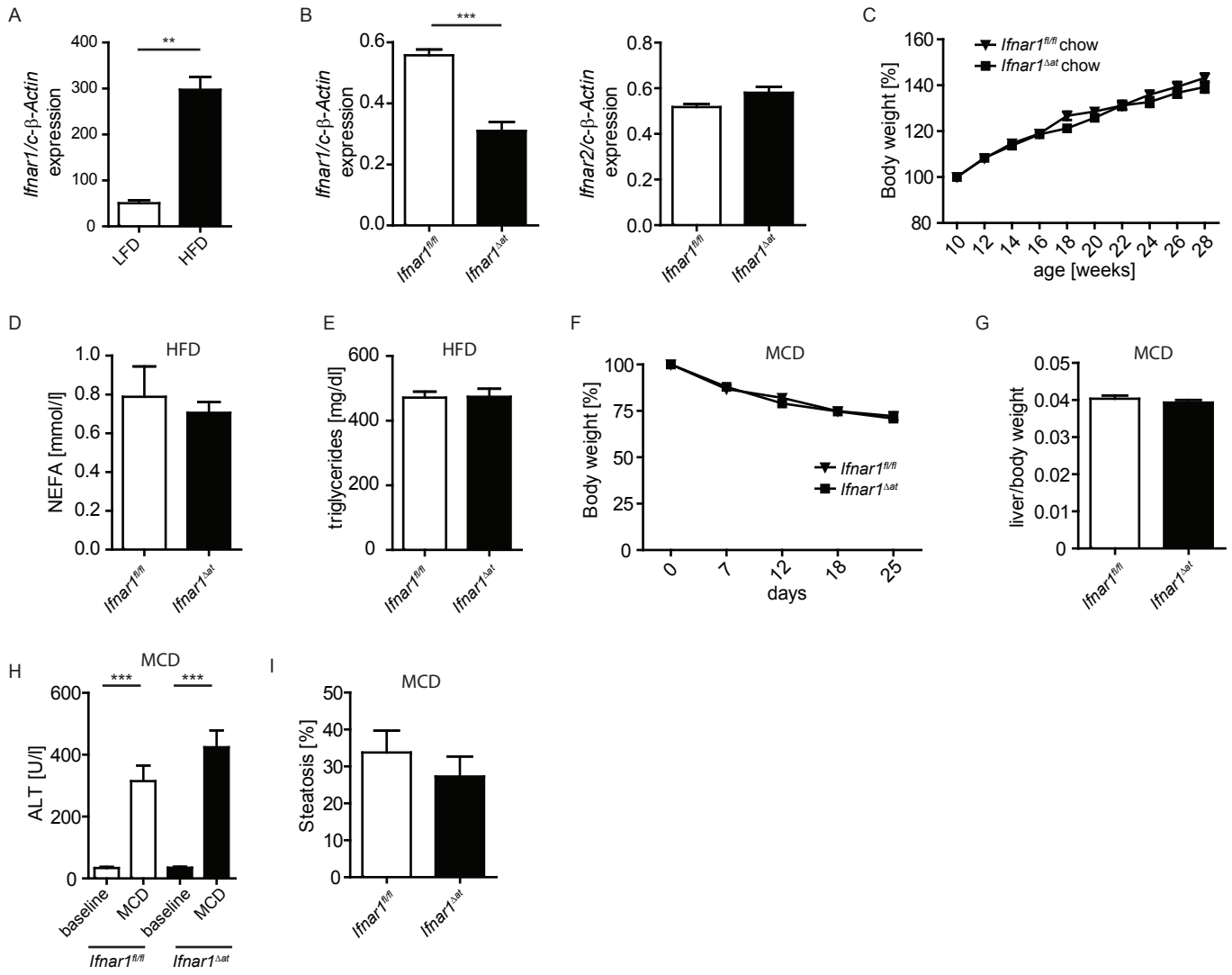
L



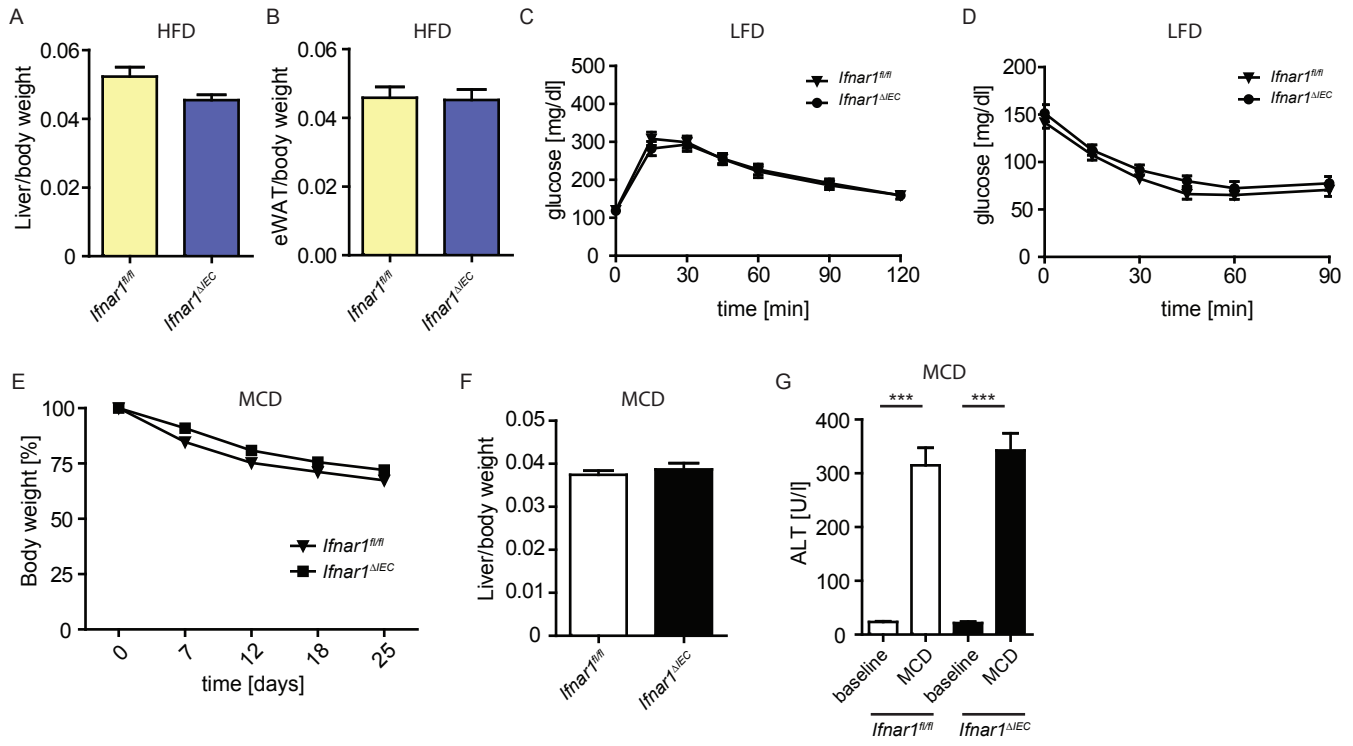
M



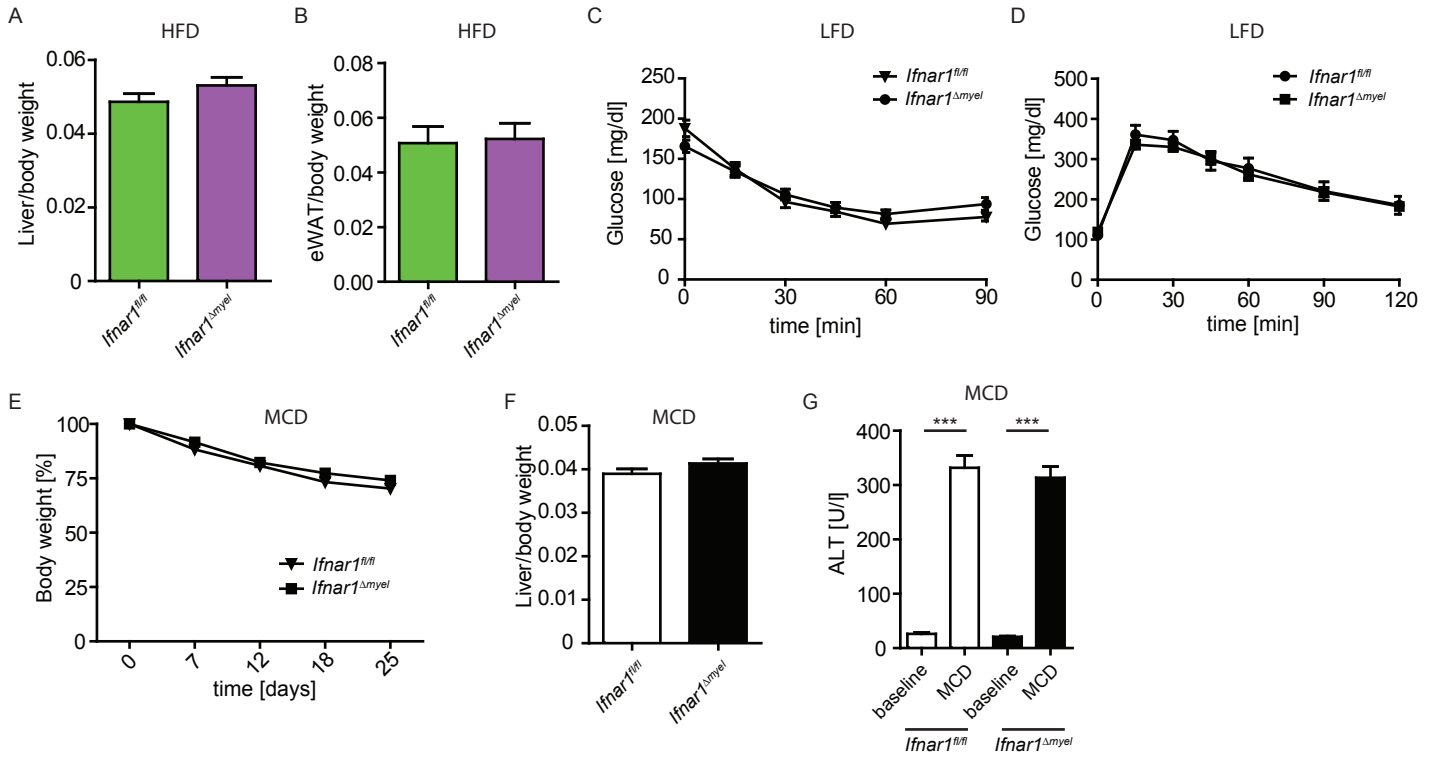
Suppl. Figure 2



Suppl. Figure 3

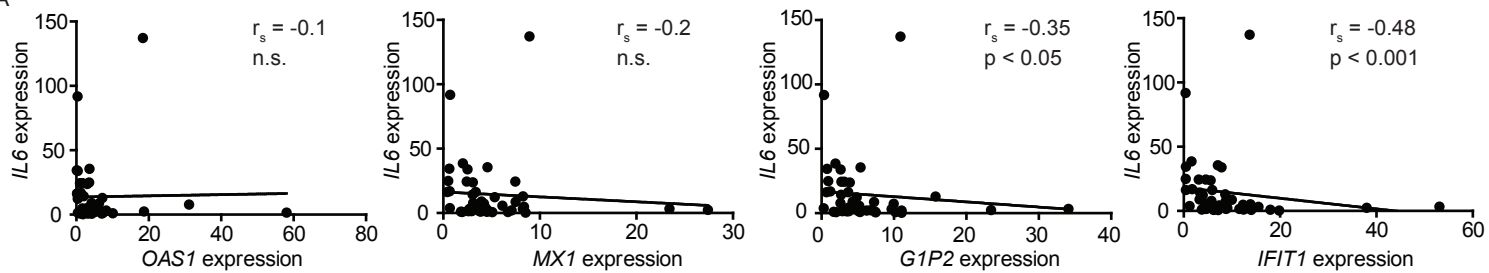


Suppl. Figure 4



Suppl. Figure 5

A



B

