

Table 1. siRNA Sequences

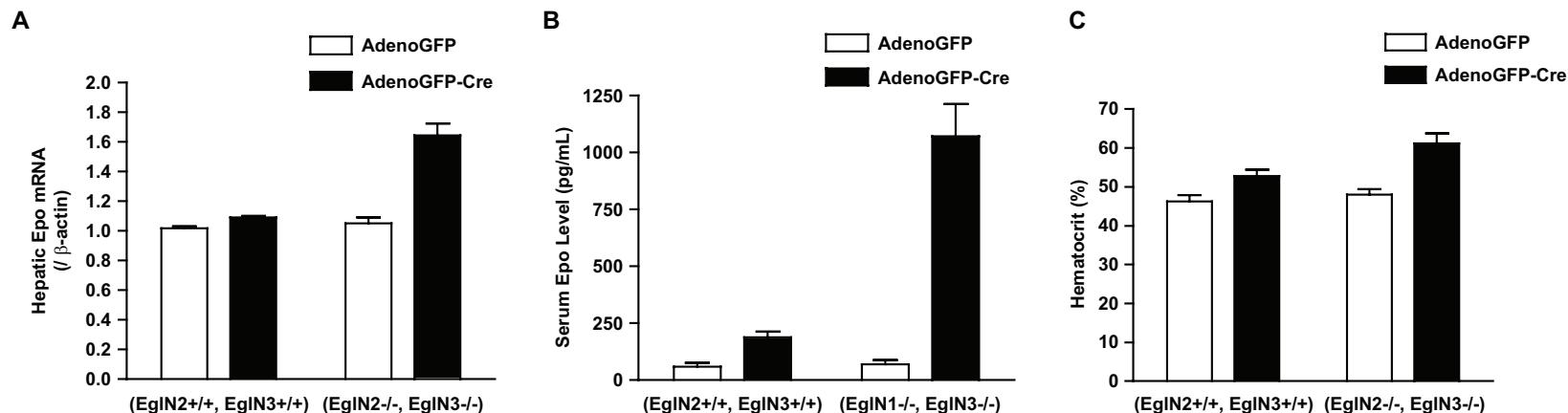
	Sequence 5' to 3'
LUC-S	cuuAcGcuGAGuA <u>c</u> uuucGAdTsdT
LUC-AS	UCGAAGuACUcAGCGuAA <u>G</u> dTsdT
mEGLN1-S	cAAGGuAcG <u>c</u> AAuAA <u>c</u> uGudTsdT
mEGLN1-AS	AcAGUuAUUGCGuACCUUGdTsdT
m/rEGLN2-S	AGAcAA <u>G</u> uA <u>c</u> AG <u>c</u> uAG <u>c</u> AdTsdT
m/rEGLN2-AS	UGCuAGCUGAuACUUGUC <u>d</u> TsdT
m/rEGLN3-S	GuAGuAA <u>c</u> AG <u>c</u> u <u>c</u> GGAA <u>d</u> TsdT
m/rEGLN3-AS	UUUCCGGAACUGUuACuACd <u>T</u> sdT
rEGLN1-S	cG <u>c</u> AcAAGGuAcG <u>c</u> AAuAdTsdT
rEGLN1-AS	uAUUGCGuACCUUGUGGCGdTsdT
GFP-S	AcAuGAAG <u>c</u> AG <u>c</u> ACGACuU <u>d</u> TsdT
GFP-AS	AAGUCGUGCUGCUUCAUGU <u>d</u> TsdT

Lowercase base=2'OMe, s=phosphorothioate, m=mouse, r=rat

Table 2. 5'RACE**primers**

	Sequence 5' to 3'
Adaptor oligo	CGACTGGAGCACGAGGACACTGACATGG
Nested adaptor oligo	GGACACTGACATGGACTGAAGGAGTAG
EGLN1 GSP	AGAGATGAAATGAACTCAGTTAGGTGACAGGTCTG
EGLN1 PCR Round 1	TTGTTTGTGTCCAGATGGAAAAGCTACTCTCCTC
EGLN1 PCR Round 2	GGCTTGAGTTCAACCCTCACACCTTCTCACCTG
EGLN2 GSP	TATTTCAGGCTGGCAGAACCTCCATAC
EGLN2 PCR Round 1	CAGACAGTGGCAGCCCAGTCCATACACTG
EGLN2 PCR Round 2	CAGCAGAGGTCTCTCCTTGTGCTCCTCAGTG
EGLN3 GSP	GATGTGGAAGAACTCCAATAGCTCTGAGGTC
EGLN3 PCR Round 1	CAGTGCTGAATTACCAGGAAGCTTCTATCCTCTG
EGLN3 PCR Round 2	GCAAGAAAACATGAAGTACCACAAACAAG

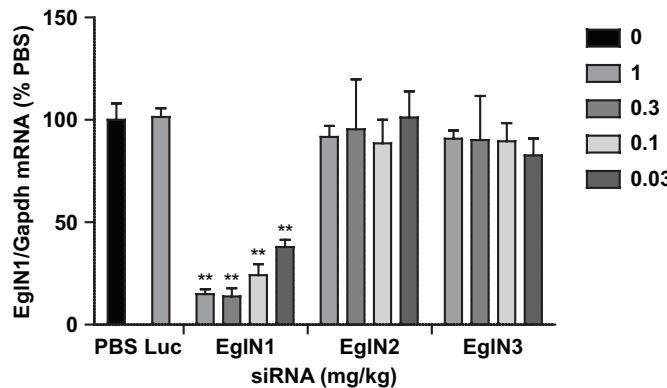
Supplemental Figure 1



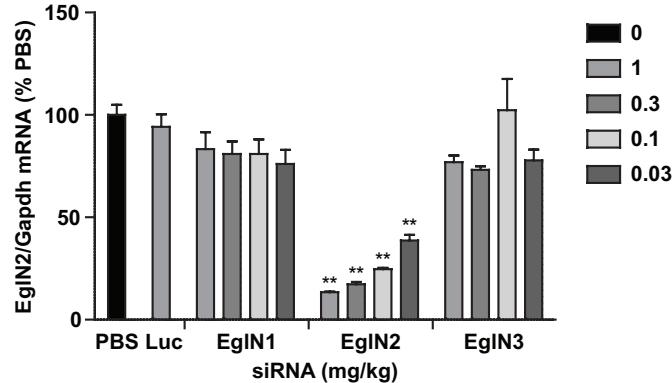
Supplemental Figure 1. Control of hepatic EPO production by EgIN1 and its paralogs. Hepatic mRNA, serum EPO, and Hematocrit values in *EgIN1* f/f mice with the indicated EgIN2 and EgIN3 genotypes two weeks after receiving adenoviruses encoding a GFP-Cre fusion protein or GFP alone by tail vein injection.

Supplemental Figure 2

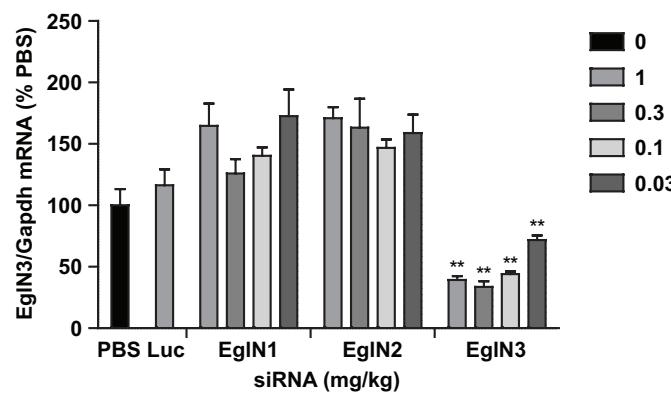
A



B

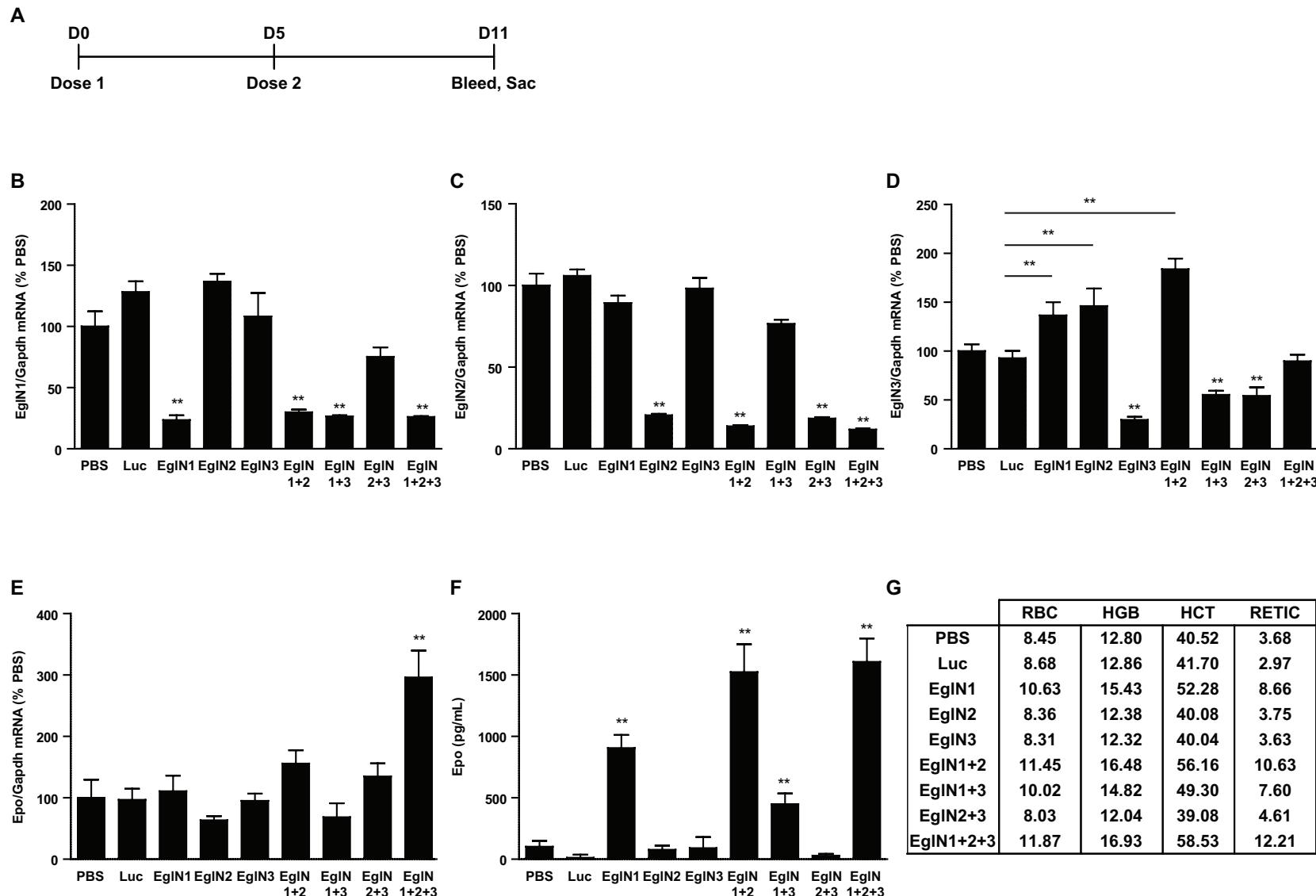


C



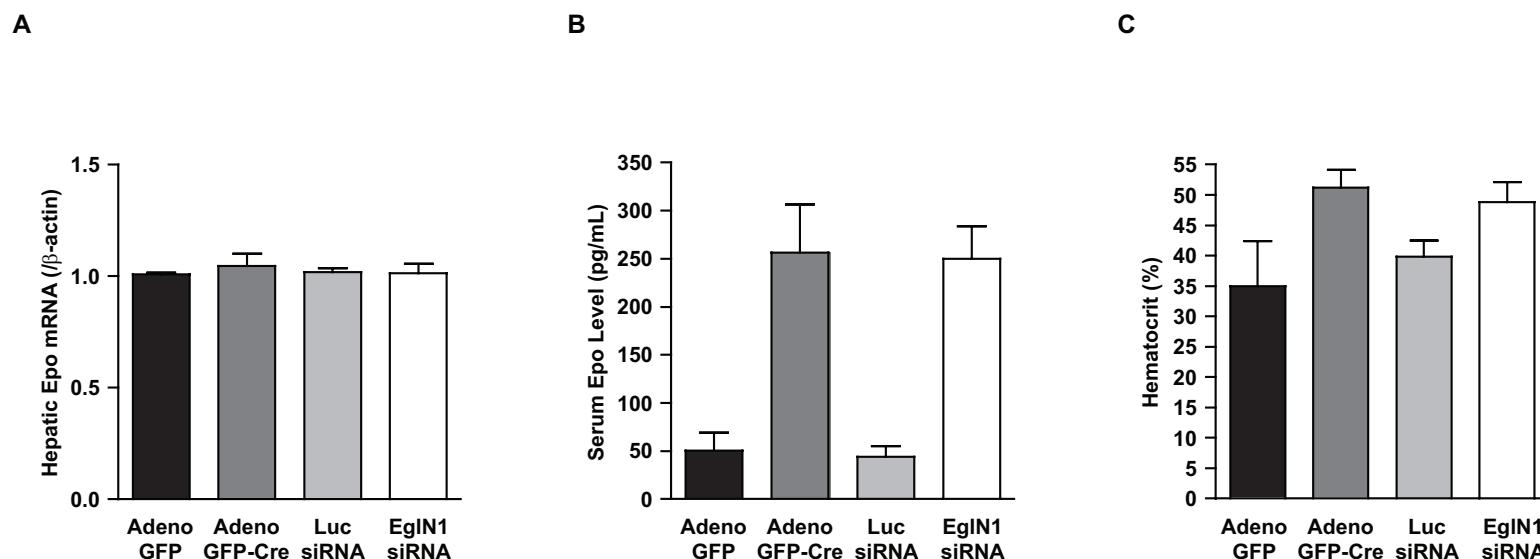
Supplemental Figure 2. Validation of EgIN siRNAs. *EgIN1*, *EgIN2*, and *EgIN3* hepatic mRNA levels in mice 72 hours after receiving increasing amounts of LNPs containing the indicated siRNAs by tail vein injection (0.03, 0.1, 0.3, or 1 mg/kg). LUC = firefly luciferase. mRNA levels were normalized to GAPDH and then to the corresponding value for PBS-treated mice. n=3. Error Bars represent 1 std. dev. **P<0.01.

Supplemental Figure 3



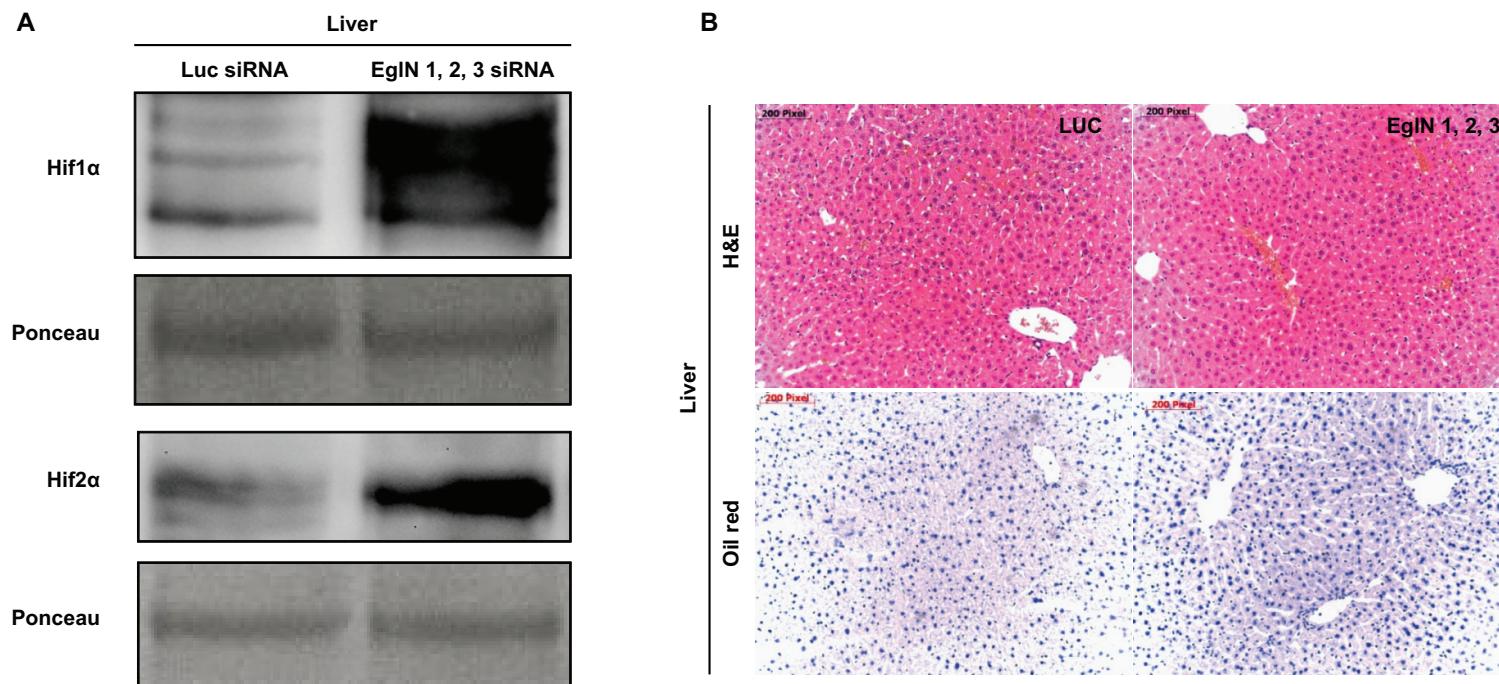
Supplemental Figure 3. Combinatorial effects of EgIN siRNAs on hepatic EPO production. (A) Dosing Schema. (B-E). Hepatic EgIN (B-D) and EPO (E) mRNA levels for mice treated with combinations of siRNA nanoparticles targeting the indicated EgIN family members (0.5 mg/kg per EgIN family member). mRNA levels were normalized to GAPDH and then to the corresponding value for PBS-treated mice. (F and G). Serum EPO (F) and hematologic (G) values for mice treated as above. n=3. Error Bars represent 1 std. dev. **P<0.01

Supplemental Figure 4



Supplemental Figure 4. Acute inactivation of EgIN1 with either Cre Recombinase or siRNA induces erythropoiesis. Hepatic EPO mRNA, serum EPO, and Hematocrit values in *EgIN1* f/f; *EgIN2* +/++; *EgIN3*+/+ mice two weeks after receiving either an adenovirus encoding a GFP-Cre fusion protein (or GFP alone) or one dose of EgIN1 siRNA LNP (or luciferase siRNA LNP) (1 mg/kg) by tail vein injection. Error bars represent 1 std. dev.

Supplemental Figure 5

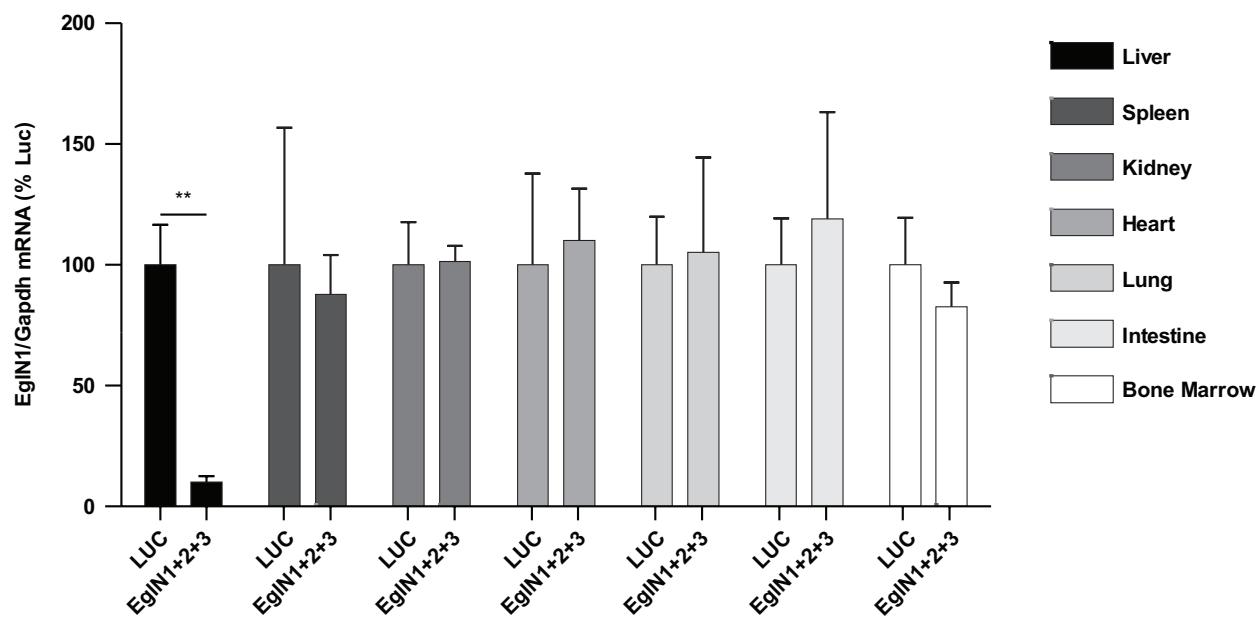


Supplemental Figure 5. Four weeks of EgIN mRNA suppression with LNPs does not result in fatty liver or histological abnormalities. (A).

Immunoblot analysis of liver extracts prepared from mice (day 9) that received two doses (day 0 and day 4) of LNPs targeting EgIN1+EgIN2+EgIN3 or luciferase (1 mg/kg siRNA/dose). (B) Hematoxylin and Eosin staining (top) and oil red staining (bottom) of liver tissue sections from mice treated with 4 weekly doses (1.5 mg/kg siRNA) of LNPs targeting EgIN1+EgIN2+EgIN3(0.5 mg/kg each)(right) or luciferase (left).

Supplemental Figure 6

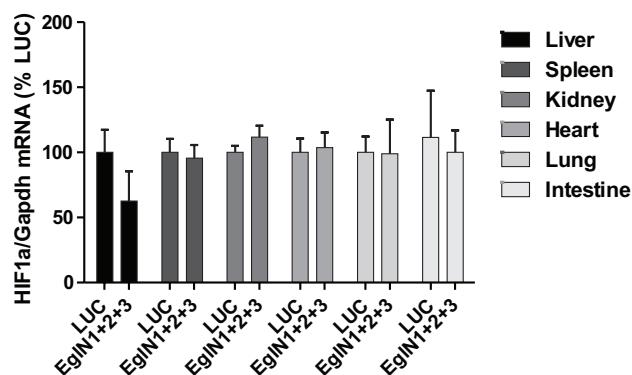
A



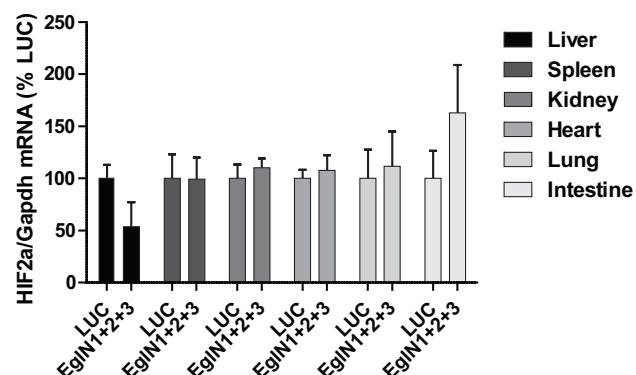
Supplemental Figure 6. EgIN mRNA silencing in different tissues. *EgIN1* mRNA levels in various mouse tissues 48 hours after receiving LNPs containing EgIN or LUC control siRNAs by tail vein injection at 1 mg/kg dose. LUC = firefly luciferase. mRNA levels were normalized to GAPDH and then to the corresponding value for LUC-treated mice. n=3. Error Bars represent 1 std. dev. **P<0.01.

Supplemental Figure 7

A

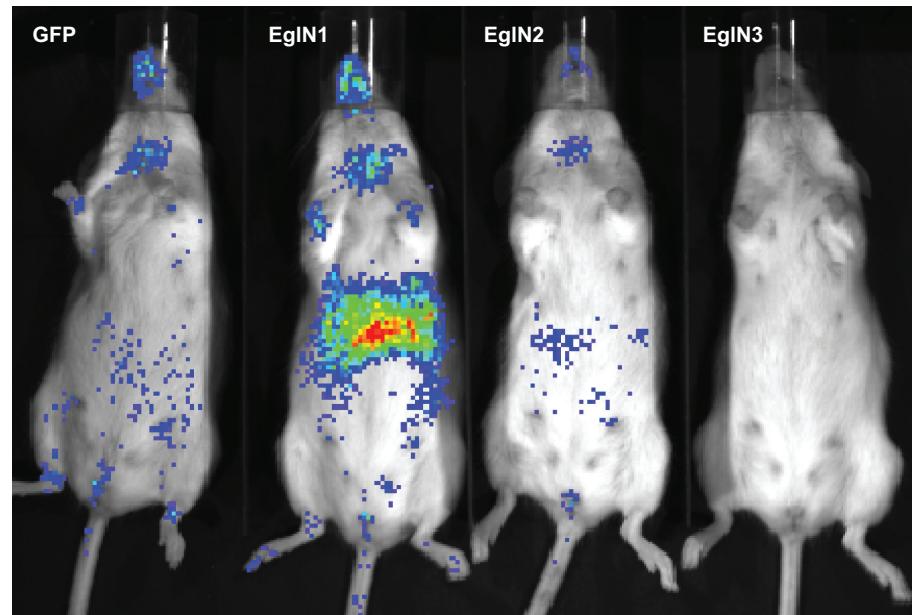


B



Supplemental Figure 7. HIF mRNA changes in different tissues after EgIN mRNA knockdown. HIF1 α (A) and HIF2 α (B) mRNA in the indicated tissues of mice treated 48 hours earlier with LNPs containing EgIN or LUC control siRNAs (1mg/kg) by tail vein injection. LUC= firefly luciferase. mRNA levels were normalized to GAPDH and then to the corresponding value for LUC-treated tissue. n=3. Error Bars represent 1 std. dev.

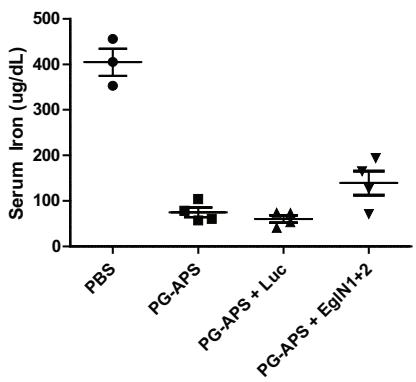
Supplemental Figure 8



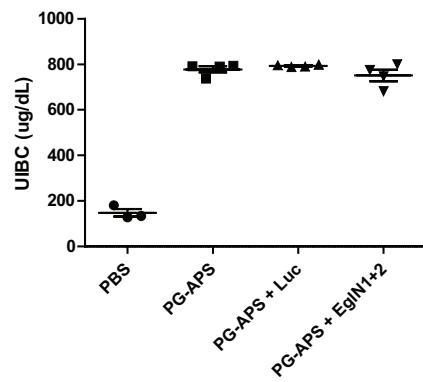
Supplemental Figure 8. Non-invasive imaging of HIF stabilization after EgIN1 inactivation. Bioluminescent images of HIF1 α -Luc mice 72 hours after a single intravenous dose of LNPs with siRNA targeting the indicated EgIN family members or, as a negative control, green fluorescent protein (GFP) (1 mg/kg).

Supplemental Figure 9

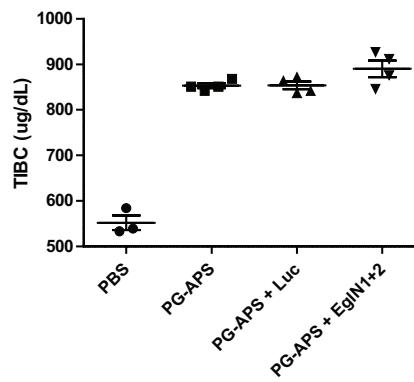
A



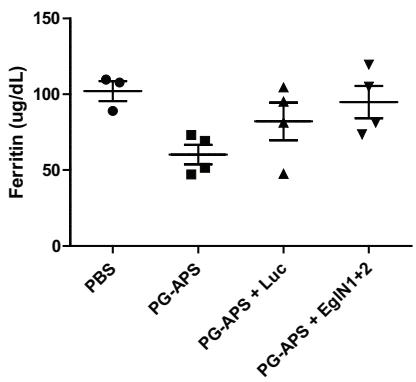
B



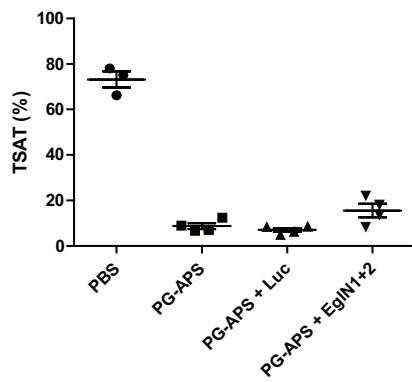
C



D



E



Supplemental Figure 9. Iron parameters in rat anemia of inflammation model. Serum iron (A), UIBC (B), TIBC (C), Ferritin (D) and Transferrin saturation (TSAT) (E) at day 37 after 3 weekly 1 mg/kg doses of LNPs targeting EgIN1/2 or Luciferase. N=3 (PBS), N=4 (other groups). Error Bars represent 1 std. dev.