## **Supplementary Material**

Immunomodulatory glycan LNFPIII alleviates hepatosteatosis and insulin resistance through direct and indirect control of metabolic pathways

Prerna Bhargava<sup>1</sup>, Changlin Li<sup>2</sup>, Kristopher J. Stanya<sup>1</sup>, David Jacobi<sup>1,3</sup>, Lingling Dai<sup>1,4</sup>, Sihao Liu<sup>1</sup>, Matthew R. Gangl<sup>1</sup>, Donald A. Harn<sup>2</sup>, Chih-Hao Lee<sup>1</sup>.

<sup>1</sup>Department of Genetics and Complex Diseases, Division of Biological Sciences, Harvard School of Public Health, 665 Huntington Ave, Boston, MA 02115, USA <sup>2</sup>Department of Infectious Diseases, College of Veterinary Medicine, University of Georgia, Athens, GA 30602, USA.

<sup>3</sup>CHRU de Tours, Service de Médecine Interne-Nutrition, INSERM U 1069, Université François Rabelais, Tours, France

<sup>4</sup>Pharmacogenetics Research Institute, Institute of Clinical Pharmacology, Central South University, Changsha, Hunan, People's Republic of China

Correspondence should be addressed:

Chih-Hao Lee, PhD. e-mail: clee@hsph.harvard.edu

Donald A. Harn, PhD. e-mail: dharn@uga.edu



**Supplementary Figure 1** Induction of II-10 by LNFPIII is independent of Th2 cytokines. (a) Real-time q-PCR analyses examining the expression of *II-10* and *Arg1* in bone marrow derived macrophages treated with 10, 20 or 40 ng mL<sup>-1</sup> of II-4 or 10, 20 or 40 µg mL<sup>-1</sup> of LNFPIII. (b) The expression (left panel) and production (right panel) of II-10 by LNFPIII (20 µg mL<sup>-1</sup>) in dendritic cells. (c) *II-10* expression determined by real-time q-PCR in wild-type (WT), *Ppar-* $\delta^{-/-}$  and *Stat6*<sup>-/-</sup> macrophages given 20 µg mL<sup>-1</sup> LNFPIII overnight. (d) Serum II-10 and II-4 concentrations in vehicle or SEA treated mice (*n* = 4–6 per treatment). (e) Glucose tolerance test and (f) insulin tolerance test in high fat fed mice (*n* = 4–6) treated with vehicle or SEA (25 µg, twice a week). Inset: area under the curve (AUC). (g) Fasting insulin concentrations and HOMA-IR to assess insulin sensitivity in vehicle and SEA treated mice. Values are expressed as means ± SEM. For *in vitro* assays, the mean and SEM were determined from 3–4 biological replicates for a representative experiment. Experiments were repeated three times. *In vivo* studies were reproduced in three mouse cohorts (*n* = 4–6 per treatment). \**P* < 0.05 (SEA versus vehicle control).



Supplementary Figure 2 SEA reduces inflammation and improves metabolic homeostasis in WAT. (a) WAT histology showing crown-like structures (indicated with arrows). Scale bar = 200  $\mu$ m. (b) Real-time q-PCR analyses examining the expression of M1 and M2 genes in WAT from vehicle or SEA treated mice (n = 5 per treatment). (c) WAT metabolic gene expression. Values are expressed as means ± SEM. Metabolic studies were reproduced in three mouse cohorts (n = 4-6 per treatment). Histology and expression analyses were examined in one and two of the three cohorts, respectively. \*P < 0.05 (SEA versus vehicle control).



**Supplementary Figure 3** The insulin sensitizing activity of LNFPIII and SEA is mediated by II-10. (**a** and **b**) The expression of inflammatory and metabolic genes in WAT of  $II-10^{-/-}$  mice treated with vehicle or LNFPIII (n = 6 per treatment). (**c**) Insulin tolerance test in wild-type (WT, left panel) and  $II-10^{-/-}$  mice (middle panel) treated with vehicle or SEA (n = 6 per treatment). Right panel: area under the curve of ITT. (**d**) Glucose tolerance and (**e**) insulin tolerance test in high fat fed male C57BL/6J mice given PBS (vehicle) or rII-10 (1 µg, every other day, 3 doses, n = 6 per treatment). (**f**) Serum II-10 and II-4 concentrations in vehicle or rII-10 treated mice. (**g** and **h**) The expression of inflammatory and metabolic genes in WAT of mice treated with vehicle or rII-10 (n = 6 per treatment). Values are expressed as means ± SEM. Studies using  $II-10^{-/-}$  and control mice (n = 6) and in rII-10 treatment (n = 6) were conducted in 1 cohort. \*P < 0.05 (treatment versus vehicle control); #P < 0.05 ( $II-10^{-/-}$  versus wt).



**Supplementary Figure 4** SEA protects against hepatic steatosis and suppresses lipogenic gene expression. (a) Liver histology and triglyceride content analyses to determine hepatic fat accumulation in vehicle and SEA treated mice. Scale bar = 100  $\mu$ m. (b) Circulating AST and ALT concentrations to assess liver function. (c and d) Expression analyses of metabolic genes and nuclear receptor signaling pathways known to regulate lipogenesis in livers from vehicle or SEA treated mice (n = 5) by real-time q-PCR. (e) Gene expression in muscle from vehicle or SEA treated mice determined by real-time q-PCR. Values are expressed as means ± SEM. Metabolic studies were reproduced in 3 mouse cohorts (n = 4-6 per treatment). Histology was examined in one and lipid and expression analyses were examined in two of the three cohorts. \*P < 0.05 (SEA versus vehicle control).



**Supplementary Figure 5** LNFPIII, but not II-10, suppresses de novo lipogenesis in the liver. (a) *Ex vivo* lipogenesis assays in primary hepatocytes treated with vehicle, LNFPIII ( $20 \ \mu g \ ml^{-1}$ ) or rII-10 ( $10 \ ng \ ml^{-1}$ ) for 24 hr. (b and c) Triglyceride content and lipogenic gene expression in the liver from vehicle or rII-10 treated mice (n = 6). (d and e) Triglyceride content and lipogenic gene expression in the liver of *Il-10<sup>-/-</sup>* mice treated with vehicle or LNFPIII (n = 6) (f) Gene expression analyses in livers of *Fxr-\alpha^{-/-}* mice treated with vehicle or LNFPIII (n = 6) treated with vehicle, SEA or LNFPIII. (i) *Ex vivo* insulin stimulated glucose uptake in adipose tissue slices. Values are expressed as means ± SEM. For *in vitro* assays, the mean and SEM were determined from 3–4 biological replicates for a representative experiment. Experiments were repeated three times. Studies using rII-10 (n = 6), *Il-10<sup>-/-</sup>* (n = 6), *Fxr-\alpha^{-/-}* (n = 6) and corresponding controls were performed in 1 mouse cohort. \**P* < 0.05 (treatment versus control).



Supplementary Figure 6 Fxr- $\alpha$  is a molecular target of LNFPIII and SEA. (a) Sequence comparison of the 5' proximal regulatory region of human, rat and mouse  $FXR-\alpha$ downstream promoter (promoter 2). The transcriptional start site is indicated with an arrow. Putative binding sites for C/EBP (site 1) and AP1 (site 2 and site 3) are highlighted. Sequences mutated for the reporter assays are indicated above the highlighted binding sites. (b) Reporter assays showing that the induction of human FXR- $\alpha$  promoter 2 is mediated by AP1 binding sequences in site 3. Similar results were obtained with LNFPIII treatment (data not shown). (c) Lipogenic and fatty acid oxidation assays conducted in hepatocytes isolated from wild type (WT) or  $Fxr-\alpha^{-/-}$  mice treated with vehicle, LNFPIII (20  $\mu$ g ml<sup>-1</sup>) or SEA (2  $\mu$ g ml<sup>-1</sup>). (d) Left panel: Western blot analyses showing Erk phosphorylation in hepatocytes ± SEA ± PD98059. p-Erk: phospho-Erk; t-Erk; total Erk. Actin was included as a loading control. Right panel: de novo lipogenesis assays in hepatocytes. Values are expressed as means ± SEM. Mean and SEM were determined from 3-4 biological replicates for a representative experiment. Experiments were repeated three times. \*P < 0.05 (SEA or LNFPIII versus vehicle control).

**Supplementary Table 1**. Metabolic parameters of mouse cohorts used in the study. Serum samples were collected after 6 h fasting. \*P < 0.05.

Treatment	Vehicle	SEA	Vehicle	LNFPIII
Body weight (g)	43.63 ± 1.74	41.62 ± 1.36	45.26 ± 0.94	46.86 ± 1.09
WAT/body weight	0.044 ± 0.0044	0.068 ± 0.002*	0.031 ± 0.015	0.039 ± 0.001
Liver/body weight	0.042 ± 0.004	0.041 ± 0.003	0.038 ± 0.002	0.043 ± 0.002
Triglyceride (mg dL <sup>-1</sup> )	102.38 ± 3.89	130.78 ± 7.52	67.17 ± 2.01	69.79 ± 1.18
Free fatty acid (mmol $L^{-1}$ )	1.06 ± 0.05	1.14 ± 0.04	1.08 ± 0.08	1.02 ± 0.03
Cholesterol (mg dL <sup>-1</sup> )	165.90 ± 8.59	166.78 ± 12.35	154.72 ± 14.06	144.24 ± 15.99
Glucose (mg dL <sup>-1</sup> )	221.67 ± 15.17	221.17 ± 8.53	236.2 ± 18.45	217 ± 12.01
Insulin (ng mL <sup>-1</sup> )	1.002 ± 0.156	0.615 ± 0.081*	1.545 ± 0.065	1.288 ± 0.059*
Adiponectin ( $\mu g m L^{-1}$ )	5.92 ± 0.16	6.35 ± 0.13	5.59 ± 0.32	5.84 ± 0.10

a. WT mice treated with SEA and LNFPIII

b. *Il-10<sup>-/-</sup>* mice treated with LNFPIII

Treatment	Vehicle	LNFPIII
Body weight (g)	33.67 ± 3.84	37.03 ± 4.14
WAT/body weight	0.019 ± 0.004	0.028 ± 0.005
Liver/body weight	0.038 ± 0.002	0.032 ± 0.003
Triglyceride (mg dL <sup>-1</sup> )	29.11 ± 1.11	36.14 ± 3.61
Free fatty acid (mmol $L^{-1}$ )	0.49 ± 0.05	0.54 ± 0.08
Cholesterol (mg dL <sup>-1</sup> )	136.36 ± 10.87	153.89 ± 19.95
Glucose (mg dL <sup>-1</sup> )	132.17 ± 17.36	142.5 ± 16.42
Insulin (ng mL <sup>-1</sup> )	0.260 ± 0.033	0.259 ± 0.048

c. WT mice treated with rIl-10

Treatment	Vehicle	rll-10
Body weight (g)	31.93 ± 1.75	32.67 ± 1.03
WAT/body weight	0.025 ± 0.001	0.032 ± 0.001*
Liver/body weight	0.036 ± 0.001	0.035 ± 0.001
Triglyceride (mg dL <sup>-1</sup> )	76.69 ± 5.04	84.83 ± 4.2
Free fatty acid (mmol $L^{-1}$ )	0.57 ± 0.04	0.61 ± 0.06
Cholesterol (mg dL <sup>-1</sup> )	106 ± 3.80	104.21 ± 4.31
Glucose (mg dL <sup>-1</sup> )	226.5 ± 11.93	228 ± 8.47

## d. *Fxr-* $\alpha^{-/-}$ mice treated with LNFPIII

Treatment	Vehicle	LNFPIII
Body weight (g)	40.73 ± 3.23	36.33 ± 3.54
WAT/body weight	0.029 ± 0.003	0.041 ± 0.003
Liver/body weight	0.046 ± 0.004	0.053 ± 0.011
Triglyceride (mg dL <sup>-1</sup> )	46.12 ± 5.22	44.11 ± 4.90
Free fatty acid (mmol $L^{-1}$ )	0.40 ± 0.02	0.49 ± 0.03
Cholesterol (mg dL <sup>-1</sup> )	303.18 ± 27.10	315.47 ± 8.16
Glucose (mg dL <sup>-1</sup> )	175 ± 4.38	150 ± 18.78
Insulin (ng mL <sup>-1</sup> )	0.450 ± 0.025	0.198 ± 0.026*

\_\_\_\_