

Expanded View Figures

Figure EV1.

Figure EV1. Properties of various GM3 species as a function of clinical markers of metabolic disorders and chronic inflammation.

- A–D LCFA species (A), VLCFA species (B), unsaturated VLCFA species (C), and α -hydroxy VLCFA-GM3 (h24:0) (D). Colors indicate disease severity: light blue, no abnormal scores (n = 17); orange, early-phase obesity (n = 80); purple, severe obesity (n = 25). Data shown are mean \pm SD, analyzed by two-tailed unpaired t-test with Bonferroni's correction. *P < 0.05, **P < 0.01, and ***P < 0.001 for comparisons between indicated groups.
- E–J Spearman's correlations for GM3 h24:0 vs. BMI (E), GM3 h24:0 vs. abdominal circumference (F), total of α -hydroxy GM3 (h24:0 and h24:1) vs. HOMA-IR (G), GM3 h24:0 vs. serum CRP (H), α -hydroxylation rate (h24:0 to 24:0) vs. serum CRP (I), and α -hydroxylation rate (h24:0 and h24:1) vs. serum CRP (J).

Data information: Sample sizes: (A–G), n = 122; (H–J), n = 121.



Figure EV2.

Figure EV2. Positive and negative regulation of innate immune response by GM3 gangliosides.

- A GM3-mediated enhancement and inhibition of IL-12/23 production from LPS-stimulated monocytes (measured by ELISA).
- B Co-stimulation of monocytes by LPS plus GM3 species or complex ganglioside species. IL-6 production in culture supernatant was measured by ELISA.
- C-E Co-stimulation of monocytes by LPS plus GM3 species or precursor GSL species. The production of IL-6 (C), TNF-α (D), and IL-12/23 p40 (E) in culture supernatant was measured by ELISA.
- F, G Inhibitory effect of LCFA and unsaturated VLCFA-GM3 on VLCFA-GM3 species. The production of TNF-α (F) and IL-12/23 p40 (G) in culture supernatant was measured by ELISA.

Data information: Data shown are mean \pm SD (n = 3) analyzed by Tukey's multiple comparison test. **P < 0.01 for comparison with LPS stimulation without GM3 species (A), or co-stimulation by LPS and proinflammatory GM3 (22:0, 24:0) (F, G).



Figure EV3. VLCFA-GM3 species synergistically and selectively control human TLR4/MD-2 activation.

- A Co-stimulation of human monocytes by GM3 species plus various TLR ligands: LPS (0.13, 0.25 ng/ml), TLR4/MD-2, Pam3CSK4 (0.25, 0.5 µg/ml), TLR1/2, Flagellin (10, 50 ng/ml), TLR5, R848 (0.25, 0.5 µg/ml), TLR7/8, MALP-2 (0.5, 1.0 ng/ml), and TLR2/6. IL-6 production in culture supernatant was quantified by ELISA.
- B–D Production of IL-6 (B), TNF-α (C), and IL-12/23 p40 (D) in culture supernatant following costimulation of monocytes by GM3 species plus LA506 (synthetic TLR4 ligand) (3, 10, 30 ng/ml).
- E Relative luciferase reporter activities of NF-κB, AP-1, and ISRE in response to LPS (+, 5 ng/ml; ++, 1 μg/ml), sCD14 (1 μg/ml), GM3 22:0 (5 μM), and their combinations. Relative luciferase activity of control condition was defined as 1 for every reporter gene.

Data information: Data shown are mean \pm SD (n = 3, A–D; n = 4, E) analyzed by Tukey's multiple comparison test (A) or by two-tailed unpaired *t*-test (B–D). *P < 0.05 and **P < 0.01 for comparisons between indicated groups.



Figure EV4. GM3 ganglioside in adipose tissue showed increased abundance in early-phase obesity and short-term HFD.

- A Body weight, blood glucose, and epididymal fat pad weight of 6-week-old control C57/BL6 mice and ob/ob mice (n = 3).
- B Ganglioside species in epididymal fat pad were analyzed by TLC.
- C Body weight, weight gain, blood glucose, and epididymal fat pad weight of normal diet (ND) and high-fat diet (HFD) C57/BL6 mice (n = 4).
- D Ganglioside species in epididymal fat pad were analyzed by TLC.
- E Body weight, blood glucose, and epididymal fat pad weight of C3H/HeN mice (ND, HFD) and C3H/HeJ mice (ND, HFD) (n = 4).
- F Full-size images of Fig 7C. TLC analysis of acidic GSL fraction (equivalent to 0.1 mg protein) from epididymal fat pads of C3H/HeN and C3H/HeJ mice on ND or HFD for 8 weeks.

Data information: Data shown are mean \pm SD analyzed by two-tailed unpaired t-test (A, C) or by Tukey's multiple comparison test (E). *P < 0.05 and **P < 0.01 for comparisons between indicated groups.



Figure EV5. Binding model of VLCFA/LCFA-GM3 species on hTLR4/hMD-2 and the comparison to LPS and lipid IVa.

A-C Docking model of GM3 24:0 (A), Ra-LPS (in 3FXI) (B), and superposition of GM3 24:0 vs. core structure of Ra-LPS (lipid A) (C).

D-F Docking model of GM3 16:0 (D), lipid IVa (in complex with hMD-2 in 2E59) (E), and superposition of GM3 16:0 vs. lipid IVa (F). Basic residues of hTLR4 are colored in blue.