

Supplemental Figures

Figure S1

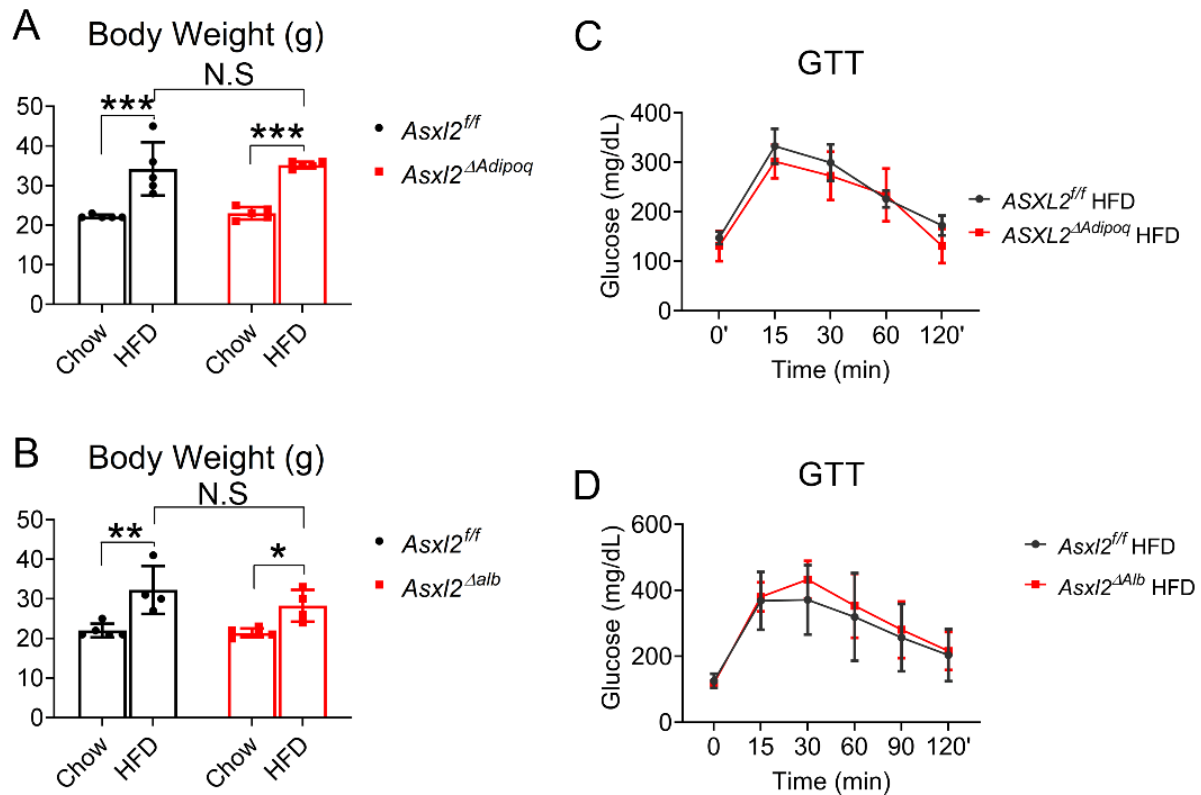


Fig S1: *Asx12* deletion in adipose tissue or liver does not impact HFD-induced weight gain and glucose tolerance. A, B) Body weights of mice with selective deletion of *Asx12* in A) adipose tissue or B) liver, fed chow diet or HFD for 8 weeks. C, D) Glucose tolerance test of mice with selective deletion of *Asx12* in C) adipose tissue or D) liver, fed HFD for 8 weeks. Data are presented as mean ± SD. **p < 0.01; *** p < 0.001 as determined by 2 way ANOVA with Holm-Sidak's post hoc analysis for multiple comparisons test.

Figure S2

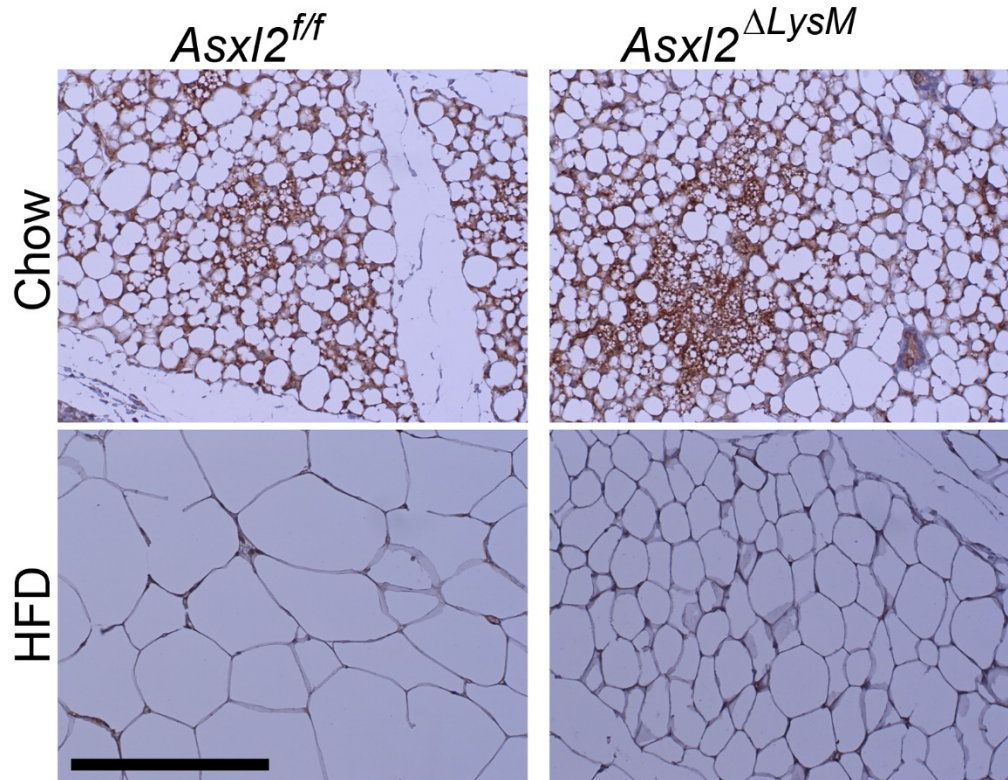


Fig S2: Beiging is not evident in WAT of *Asx12^{ΔLysM}* mice fed HFD. Histological sections of inguinal WAT of *Asx12^{f/f}* and *Asx12^{ΔLysM}* mice fed chow diet or HFD immunostained for UCP1; Scale bar: 200μm.

Figure S3

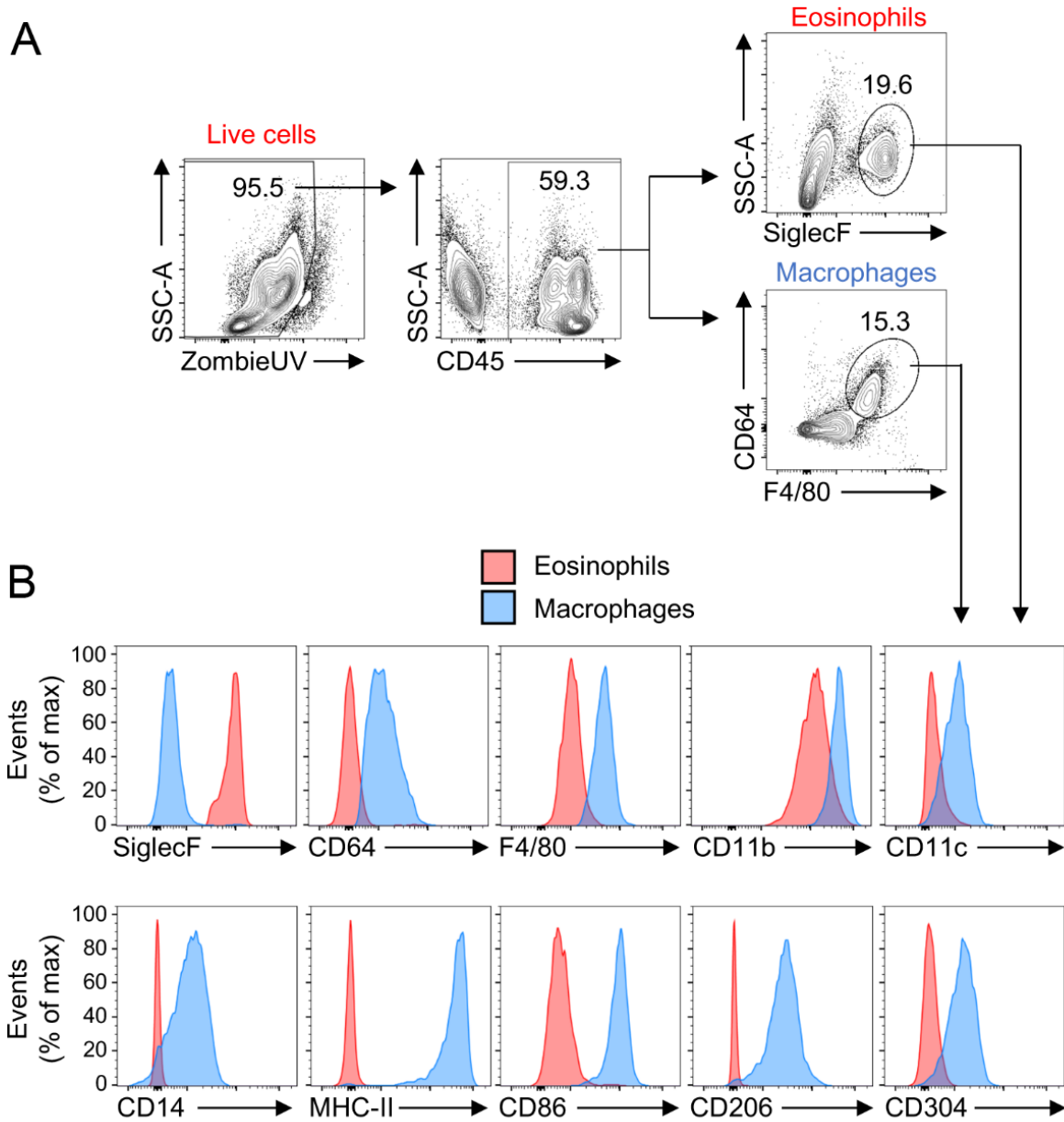


Fig S3: Identification of macrophages and eosinophils in adipose tissue. The stromal vascular fraction of gonadal adipose tissue of HFD- fed *Asx12^{ff}* and *Asx12^{ALysM}* mice was isolated and stained for flow cytometric analysis. A) Gating strategy to define SSC^{hi} SiglecF⁺ eosinophils (red) and F4/80⁺ CD64⁺ macrophages (blue). B) These eosinophil and macrophage populations were gated and overlaid to examine expression of the indicated surface markers.

Figure S4

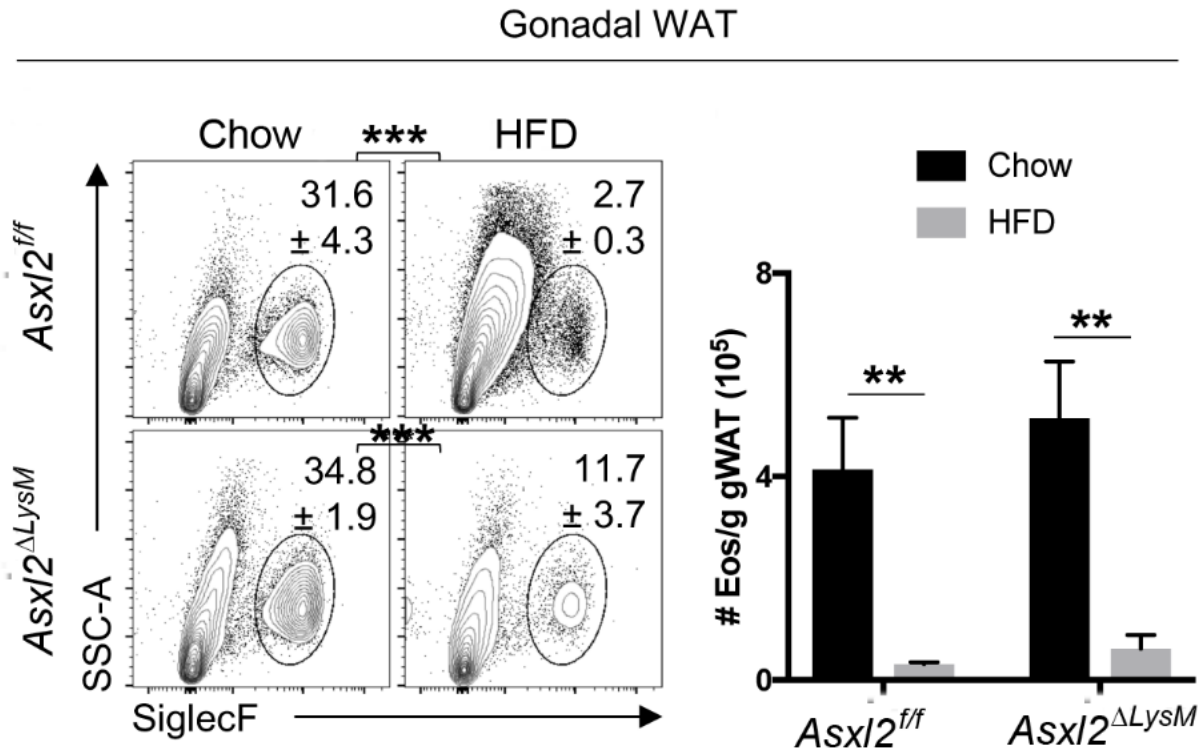


Fig S4: Eosinophils are decreased in obese gonadal WAT in an ASXL2-independent manner. Female *Asxl2^{ff}* and *Asxl2^{ΔLysM}* mice were fed either a chow diet or HFD for 8 weeks. A) Frequencies and B) numbers of SSC^{hi} SiglecF⁺ eosinophils in gonadal white adipose tissue (WAT). Pre-gated on singlet, live, CD45⁺ cells. Two-way analysis of variance with Tukey post-hoc test.

Figure S5

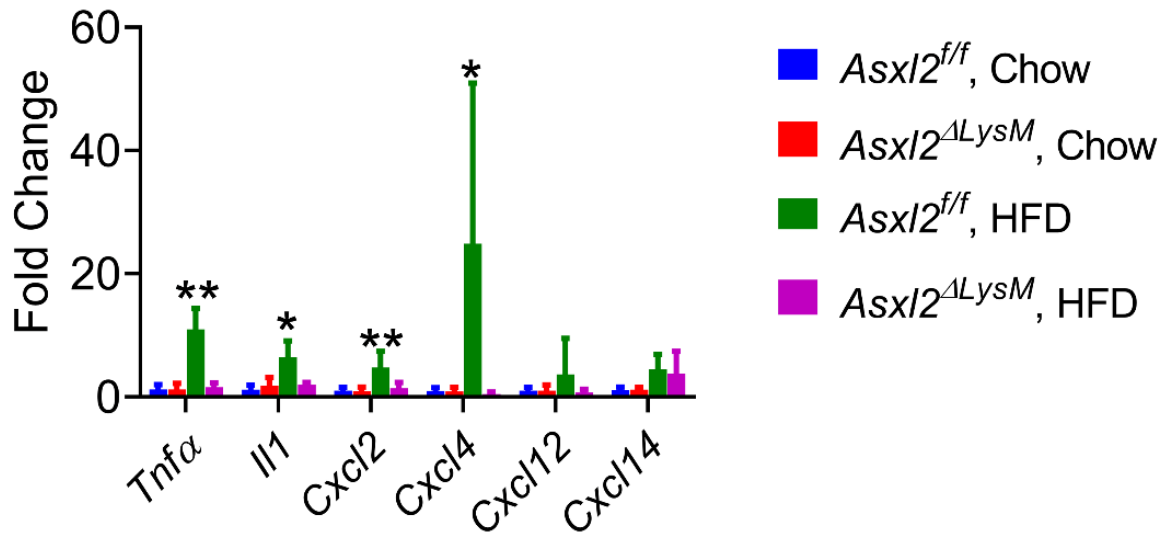


Fig S5: *Asxl2* expression in myeloid cells is required for macrophage accumulation in WAT and BAT in obesity. Inflammatory cytokine and chemokine mRNA expression in stromal vascular fraction of gonadal WAT of *Asxl2^{f/f}* or *Asxl2^{ΔLysM}* mice after 8 weeks fed with Chow diet or HFD. Data are presented as mean \pm SD. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$ as determined 2 way ANOVA with Holm-Sidak's post hoc analysis for multiple comparisons test.

Figure S6

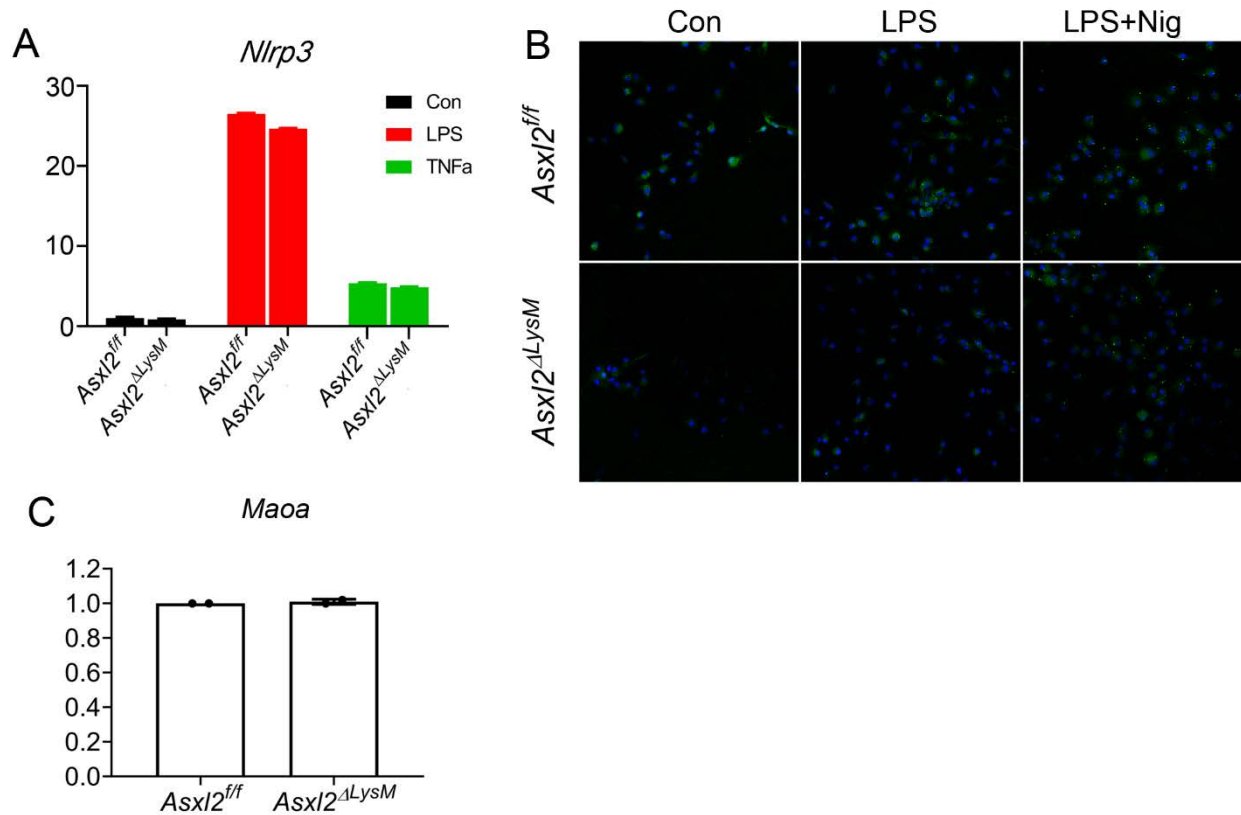


Fig S6: ASXL2 does not affect NLRP3 activation in macrophages. A) BMMs of *Asxl2^{ff}* and *Asxl2 Δ LysM* mice were stimulated with LPS (100ng/ml) or TNF α (10 ng/ml) for 3 hours, *Nlrp3* mRNA expression were measured by qPCR; B) *Asxl2^{ff}* or *Asxl2 Δ LysM* BMMc were incubated for 3 hours with 100 ng/ml LPS and stimulated with 15 μ M nigericin for 30 minutes. Cells were then incubated with FLICATM FAM-YVAD-FMK probe and analyzed by fluorescence microscopy. C) *Maa* mRNA abundance in BMMs of *Asxl2^{ff}* and *Asxl2 Δ LysM* mice were analyzed by qPCR. (n= 2 independent experiments).

Figure S7

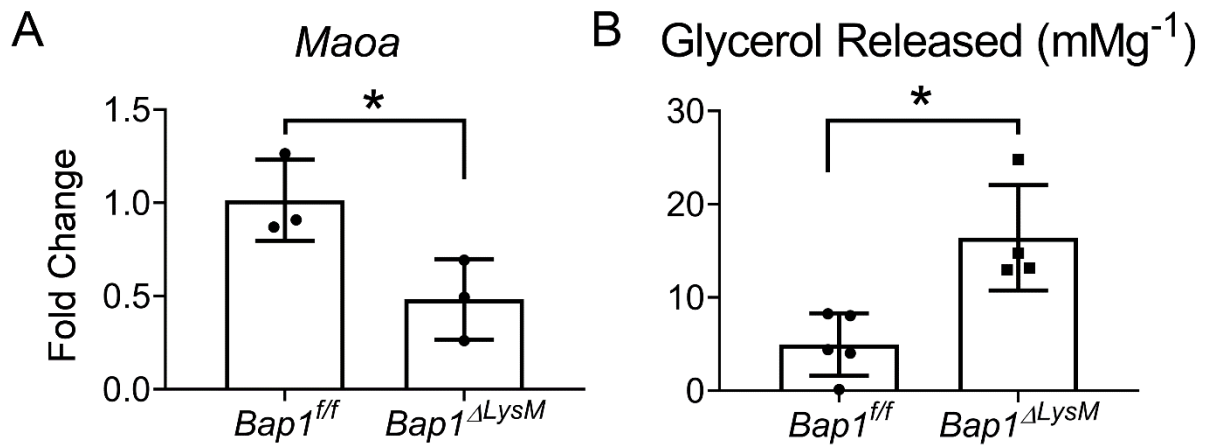


Fig S7: A) *Maoa* mRNA abundance in BAT stromal vascular fraction of HFD-fed control and *Bap1^{ΔLysM}* mice; B) Lipolysis analysis in BAT explants derived from HFD-fed control and *Bap1^{ΔLysM}* mice. Data are presented as mean \pm SD. * $p < 0.05$ as determined by unpaired t test.

Figure S8

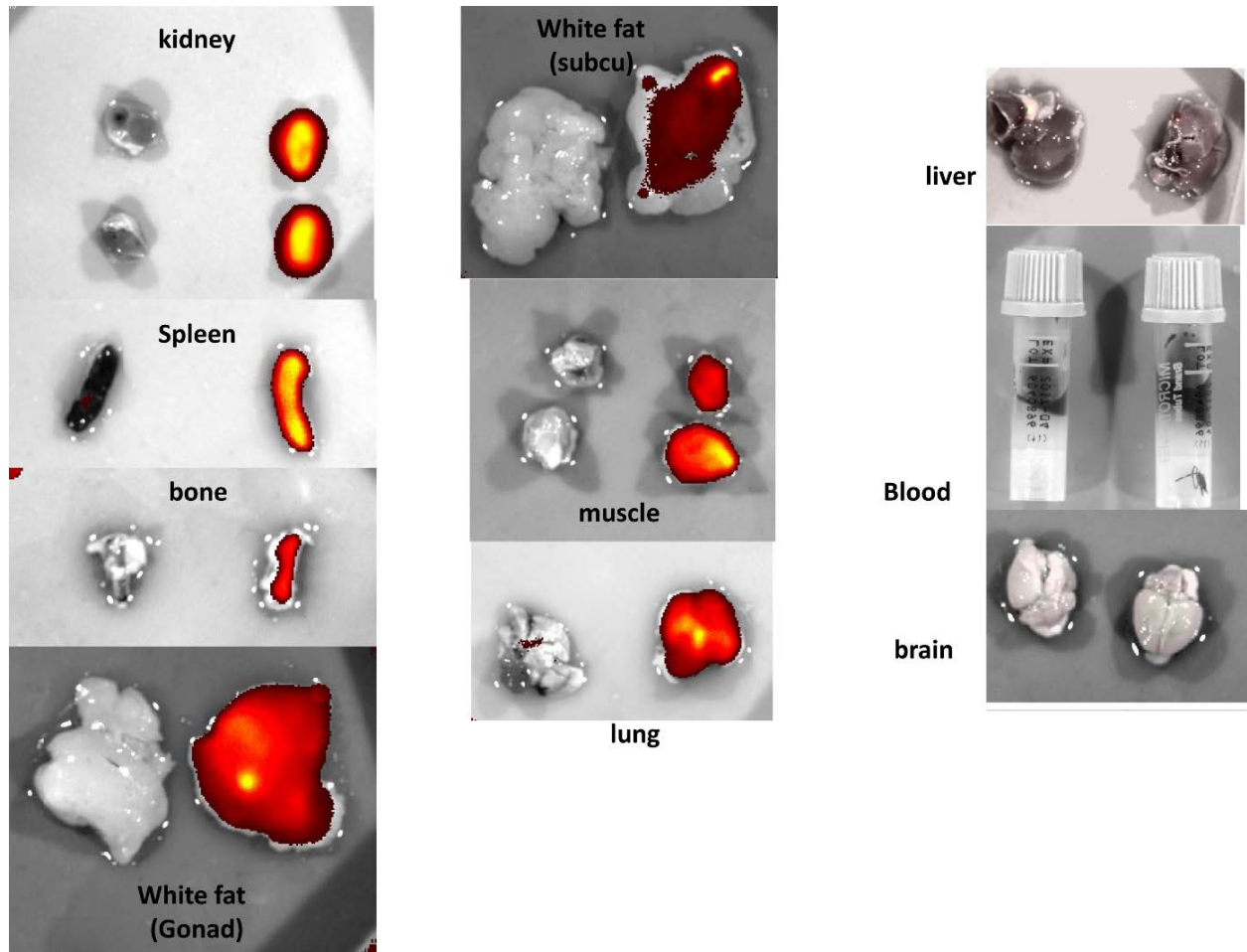


Fig S8: Nanoparticle-associated *Asx12*-siRNA targets macrophages in numerous tissues. Fluorescent scan of organs of WT HFD-fed WT mice administered Cy5.5-labeled nanoparticle free (left) or nanoparticle-associated (right) *Asx12*-siRNA