Supplementary information

Wang et al.

Adipose group 1 innate lymphoid cells promote adipose tissue fibrosis and diabetes in obesity

were analyzed							
	Control	Obese	Obese T2D	P values			
				Overall	Control	Control	Obese
					vs.	vs.	vs.
					Obese	Obese	Obese
						T2D	T2D
Ν	24	21	20	-	-	-	-
Age (years)	42.4±6.7	41.5±13.5	38.9±11.6	0.554	-	-	-
Male, n (%)	6(25.0)	5(23.8)	6(30.0)	0.896	-	-	-
BMI (kg m ⁻²)	24.2±2.7	35.5±7.4	38.6±5.3	0.000	0.000	0.000	0.073
HbA1c (%)	5.3±0.5	5.7±0.3	8.8±1.2	0.000	0.076	0.000	0.000
Fasting glucose (mmol 1-1)	4.7±0.4	5.3±0.5	11.0±3.5	0.000	0.304	0.000	0.000
2h blood glucose (mmol l ⁻¹)	5.7±0.9ª	7.1±1.6	16.8±4.1	0.000	0.314	0.000	0.000
Fasting insulin (mIU ml ⁻¹)	5.7±7.6	21.6±13.7	28.7±14.2	0.000	0.000	0.000	0.066
HOMA-IR (units)	1.2±1.5	5.3±3.7	13.8±8.1	0.000	0.008	0.000	0.000
Triglycerides (mmol l ⁻¹)	1.2±0.5	1.8±0.8	4.3±3.6	0.000	0.394	0.000	0.000
Total cholesterol (mmol l ⁻¹)	4.4±0.7	4.7±0.8	5.1±1.1	0.038	0.390	0.012	0.095
HDL-C (mmol l ⁻¹)	1.2±0.3	1.1±0.3	1.0±0.1	0.007	0.170	0.002	0.068
LDL-C (mmol l ⁻¹)	2.6±0.6	2.9±0.6	2.6±0.6	0.208	-	-	-
Fasting FFA (mmol l ⁻¹)	0.45±0.1	0.7±0.1	0.7±0.1	0.000	0.000	0.000	0.130
Adipo-IR (mIU ml ⁻¹ ×mmol l ⁻¹)	2.4±2.9	14.7±10.6	21.0±12.6	0.000	0.000	0.000	0.039

Supplementary Tables Supplementary Table 1 Characteristics of participants whose adipose tissue samples

BMI, body mass index; HOMA-IR, homeostasis model assessment-insulin resistance; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; Adipo-IR, adipose insulin resistance index. All data are presented as mean±SD or n (%). Comparisons are by ANOVA and, when appropriate, LSD post-hoc test. ^a, 13 non-obese control subjects were missing.

	Obese				Obese-T2DM				
	pre-surgery	3 months	Р	Change	pre-surgery	3 months	Р	Change	<i>P</i> for
		post-surgery	value			post-surgery	value		change
Ν	19	-	-	-	17	-	-	-	-
Age (years)	35.2±9.9	-	-	-	36.9±10.9	-	-	-	-
Male, n (%)	6(31.6)	-	-	-	3(17.6)	-	-	-	-
BMI (kg m ⁻²)	38.4±6.9	31.8±5.2	0.002	6.6±7.3	38.7±4.9	31.8±4.0	< 0.001	6.9±1.8	0.861
HbA1c (%)	5.7±0.9	5.1±0.3	0.023	0.5±1.0	8.9±1.4	5.5±0.7	< 0.001	3.5±1.1	0.000
Fasting glucose (mmol l ⁻¹)	5.6±0.7	5.0±0.4	0.004	0.6±0.7	11.4±3.8	5.6±0.7	< 0.001	5.8±3.6	0.000
2 h post (mmol 1 ⁻¹)	7.0±1.7	NA	-	-	17.1±4.5	6.8±1.9	< 0.001	10.3±4.2	-
Fasting insulin (mIU ml ⁻¹)	26.4±11.0	11.6±3.7	< 0.001	14.7±10.2	34.7±21.4	15.0±7.2	<0.001	19.7±16.0	0.265
HOMA-IR (units)	6.8±3.3	2.6±0.9	< 0.001	4.2±3.1	16.6±9.5	3.6±1.5	<0.001	13.0±8.8	0.000
Triglycerides (mmol 1 ⁻¹)	1.8±0.7	1.1±0.3	< 0.001	0.8±0.6	4.8±3.8	1.4±0.5	0.001	3.4±3.7	0.004
Total cholesterol (mmol l^{-1})	4.7±0.8	4.1±0.7	0.000	0.6±0.6	5.1±1.2	3.9±0.7	<0.001	1.3±0.9	0.012
HDL-C (mmol	1.2±0.3	1.2±1.4	0.229	0.1±0.2	1.0±0.1	1.0±0.2	0.498	0.0±0.2	0.675
LDL-C (mmol	2.9±0.7	2.2±0.6	0.000	0.7±0.7	2.9±0.6	2.2±0.6	0.125	0.3±0.7	0.136
Fasting FFA (mmol 1 ⁻¹)	0.6±0.1	0.4±0.2	0.000	0.2±0.2	0.7±0.1	0.4±0.1	< 0.001	0.4±0.2	0.076
Adipo-IR (mIU ml ⁻¹ ×mmol l ⁻¹)	17.0±9.1	4.3±2.6	< 0.001	12.7±9.2	25.1±15.6	5.2±3.1	< 0.001	19.9±14.1	0.075

Supplementary Table 2 Clinical parameters of subjects pre-surgery and 3 months post-surgery

BMI, body mass index; HOMA-IR, homeostasis model assessment-insulin resistance; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; Adipo-IR, adipose insulin resistance index. All data are presented as mean±SD or n (%). Comparisons are by ANOVA and, when appropriate, LSD post-hoc test. NA, not applicable.

	Trichrome positive area (% of WAT area)		
	r	Р	
BMI (kg m ⁻²)	0.785	0.000	
HOMA-IR (units)	0.714	0.000	
Adipo-IR (mIU ml ⁻¹ ×mmol l ⁻¹)	0.658	0.000	

Supplementary Table 3 Correlations between the percentage of positively stained area indicated for adipose tissue fibrosis and BMI, HOMA-IR, and Adipo-IR

Supplementary Table 4 Primers used in this study

For human	studies:				
Number	Gene	Forward (5'-3')	Reverse (5'-3')		
1	COL1A1	GAGGGCCAAGACGAAGACATC	CAGATCACGTCATCGCACAAC		
2	COL3A1	GGAGCTGGCTACTTCTCGC	GGGAACATCCTCCTTCAACAG		
3	MMP-2	TACAGGATCATTGGCTACACACC	GGTCACATCGCTCCAGACT		
4	MMP-9	TGTACCGCTATGGTTACACTCG	GGCAGGGACAGTTGCTTCT		
5	TIMP-1	CTTCTGCAATTCCGACCTCGT	ACGCTGGTATAAGGTGGTCTG		
6	LOX	CGGCGGAGGAAAACTGTCT	TCGGCTGGGTAAGAAATCTGA		
7	MINCLE	AAGAACTGCTCAGCCATGGG	CCTGCTCCTCCTGTGAGTTGA		
8	iNOS	TTCAGTATCACAACCTCAGCAAG	TGGACCTGCAAGTTAAAATCCC		
9	TGFB1	GGCCAGATCCTGTCCAAGC	GTGGGTTTCCACCATTAGCAC		
10	β -ACTIN	GTTGCTATCCAGGCTGTG	TGATCTTGATCTTCATTGTG		

For animal studies:

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Number	Gene	Forward (5'-3')	Reverse (5'-3')
1	Collal	TGCTGGTCCCAAAGGTTC	CAGGGCGACCATCTTGAC
2	Col3a1	ATGCCCACAGCCTTCTACAC	ACCAGTTGGACATGATTCACAG
3	Mincle	ACCAAATCGCCTGCATCC	CACTTGGGAGTTTTTGAAGCATC
4	iNOS	CCAAGCCCTCACCTACTTCC	CTCTGAGGGCTGACACAAGG
5	Tgfb1	CCTGACTGGCTGTCTTTTGACG	AGTGAGCGCTGAATCGAAAGC
6	Pdgfb	CCATCCGCTCCTTTGATGAT	TCAGCCCCATCTTCATCTACG
7	Acta2	GTTCAGTGGTGCCTCTGTCA	ACTGGGACGACATGGAAAAG
8	Il-6	CCTTCTTGGGACTGATGCTGGTG	AGGTCTGTTGGGAGTGGTATCCTC
10	TNF - α	AGCCGATGGGTTGTACCTTG	GTGGGTGAGGAGCACGTAGTC
11	β -actin	CCACAGCTGAGAGGGAAATC	AAGGAAGGCTGGAAAAGAGC

Supplementary Figures



Supplementary Figure 1. a Flow cytometry analysis of cells stained with isotype controls for CD45, CD127, CD117, CRTH2, and NKP44. **b** Representative Masson's trichrome C and Sirius red staining in omental adipose tissue samples from control subjects and obese subjects (scale bar: $100\mu m$). Quantitation of Masson's trichrome C and Sirius red staining by Image J software from NIH was performed (n=24 and 41,

respectively). ** P < 0.01 vs. control group (unpaired two-way Student's *t* test). **c** Expression levels of fibrosis related genes in control subjects and obese subjects (n=24 per group). ** P < 0.01 vs. control group (unpaired two-way Student's *t* test). **d** Sorting strategy by flow cytometry. **e** Analysis of macrophages in human adipose tissue by flow cytometry. **f** Representative flow cytometric analysis for adipose ILC1s in SVFs. **g,h** Hepatic TG levels and serum FFA levels. n.s., no significance (two-way ANOVA tests followed by Bonferroni post hoc test). Data are representative of three independent experiments, with n = 4-6 mice per group. Error bars indicate s.d..



Supplementary Figure 2. a Representative flow cytometric analysis for ATMs in mice. **b** Absolute density of CD11c⁺CD206⁺ and CD206⁺ macrophages in VAT of mice with different treatment. Data are representative of three independent experiments, with n = 4-6 mice per group. ***P*<0.01 *vs*. CHOW group; ##*P*<0.01 *vs*. HFD+PBS group (two-way ANOVA tests followed by Bonferroni post hoc test). **c** Adipose ILC1s (2×10⁴ cells well⁻¹) sorted from VAT of lean and obese mice were co-cultured with BMDMs (2×10⁵ cells well⁻¹) in 24-well plates using a transwell system. Recombinant mouse IL-12 (20 ng ml⁻¹) and IL-18 (20 ng ml⁻¹) were added in the upper chamber, with neutralizing IFN- γ antibody (20 ng ml⁻¹) or isotype control IgG (20 ng ml⁻¹) supplemented in separate group. ** *P*<0.01 *vs*. control group; ^{§§}

P<0.01 vs. Lean ILC1s+Control IgG group; ## P<0.01 vs. Lean ILC1s+Control IgG group; ^{††} P<0.01 vs. Obese ILC1s+Control IgG group (two-way ANOVA tests followed by Bonferroni post hoc test). Data are representative of three independent experiments. **d** Absolute density of CD11c⁺CD206⁺ and CD206⁺ macrophages in VAT of mice with different treatment. **P < 0.01 vs. CHOW group; $^{\#\#}P < 0.01$ vs. HFD+Control IgG group (two-way ANOVA tests followed by Bonferroni post hoc test). Data are representative of two independent experiments, with n = 5 mice per group. e-g C57BL/6 mice were fed with normal diet or HFD, while mice in the HFD group were randomized into two groups of HFD anti-IL-12 mAb group and HFD control IgG group and continued with HFD feeding for another 4 weeks. HFD anti-IL-12 mAb group received rat anti-murine anti-IL-12p35 mAb administration via intraperitoneal injection at 250 µg per mouse every 3 days for 4 weeks; HFD control IgG group received the same dose of control IgG. (e) Glucose tolerance test. (f) Expression levels of fibrosis related genes, assessed by q-PCR. (g) Representative Masson's trichrome C and Sirius red staining in VAT of mice with different treatment. ** P<0.01 vs. respective CHOW group; # P<0.05 vs. HFD+control IgG group; ## P<0.01 vs. HFD+control IgG group (two-way ANOVA tests followed by Bonferroni post hoc test). Data are representative of two independent experiments, with n = 5-6mice per group. Error bars indicate s.d..



Supplementary Figure 3. Gating strategies used for cell identification and sorting. a Gating strategy to identify human adipose and circulating ILC1 (Lin⁻CD45⁺CD127⁺CD117⁻CRTH2⁻NKP44⁻) cells for analyses presented on Fig. 1b-k,2a. **b** Gating strategy to sort adipose CD45⁺ cells or ILCs depleted CD45⁺ cells from the SVFs of obese T2D patients for in-vitro co-culture assay presented on Fig. 2d,e. c Gating strategy to identify human adipose macrophages (CD45⁺CD14⁺CD11c⁺) for analyses presented on Fig. 2d. d Gating strategy to identify adipose ILC1s (Lin-NK1.1+NKP46+T-bet+Eomes-DX5-) from Prkdc-/-IL2rg-/- and Rag1-/- mice for analyses presented on Fig. 3b,6a. e Gating strategy to identify adipose ATMs

(CD45⁺CD11b⁺F4/80⁺) from $Prkdc^{-/-}IL2rg^{-/-}$ and $Rag1^{-/-}$ mice for analyses presented on Fig. 4b,6c. **f** Gating strategy to sort adipose ILC1s from C57BL/6 and *Ifng*^{-/-} mice for adoptive transfer assay presented on Fig. 3a,5a.

Supplementary Methods

Bone Marrow Derived Macrophages Isolation

BMDMs were isolated from C57BL/6 mice fed a normal diet as previously described¹. In brief, bone marrow cells were flushed out from femur, resuspended and grown in DMEM/F12-10 medium (10% FBS and 1% penicillin and streptomycin) containing 10 ng ml⁻¹ M-CSF (PeproTech). Fresh growth medium was added on day 3. Macrophages were harvested on day 7 and replated for further experiment.

Co-culture experiments

As for the co-culture experiment, a transwell system (BD Biosciences) was used. Adipose ILC1s were sorted from C57BL/6 mice fed a normal diet or HFD diet for 4 weeks by flow cytometry. About 2×10^5 BMDMs were seeded in 24-well plate and co-cultured with adipose ILC1s (2×10^4 cells well⁻¹) in the presence of neutralizing IFN- γ antibody (BioXCell, #BP0055) or IgG isotype control antibody (BioXCell, #BP0290). Meanwhile, recombinant mouse IL-12 (R&D Systems, #419-ML) and IL-18 (R&D Systems, #9139-IL) were added in the upper chamber. After co-culture for 24 hours, BMDMs were collected for further detection.

Supplementary References

1. Zhang X, Goncalves R, Mosser DM. The isolation and characterization of murine macrophages. *Current protocols in immunology* Chapter 14, Unit 14 11 (2008).