Supplemental Material

Supplemental Fig I



Supplemental Figure I. Gating with wound resident macrophage marker F/4/80⁺ does not impact infiltrating CD11b⁺Ly6C^{Hi} cell percentages in wounds. Wounds were created using a 4mm punch biopsy on the back of male C57BL/6 mice. Wounds were harvested on day 5 post-injury and analyzed by flow cytometry. Gating strategy to select single, live, lin⁻,Ly6G⁻, CD11b⁺, Ly6C^{Hi} cells and stratify based on F4/80⁺ vs. F4/80⁻. (n =10 mice; tissues of 2 wounds per mouse were pooled for a single biological replicate. Data is representative of 2 independent experiments.) All data are expressed as the mean +/- the standard error of the means (SEM).

Supplemental Fig II



Supplemental Figure II. Wound CD11b⁺Ly6C^{Lo} monocyte/macrophages produce more TGF β than wound CD11b⁺Ly6C^{Hi} cells. Wounds from C57BL/6 mice were collected on post-injury day 2 for cell isolation, *ex vivo* stimulation, and intra-cellular staining for flow cytometry. Percentage of single, live, lineage⁻ [CD3, CD19, NK1.1, Ter-119]⁻, Ly6G⁻, CD11b⁺, Ly6C^{Hi} and Ly6C^{Lo} cells staining positive for TGF β . (****P* < 0.001; n =15 mice; tissues of 2 wounds per mouse were pooled for a single biological replicate. Data is representative of 3 independent experiments.) All data are expressed as mean +/- the standard error of the means (SEM).

Supplemental Fig III

2000-

1000

0



but altered proportions of Ly6C^{Hi}[Live, Ly6G⁻,CD11b⁺] and Ly6C^{Lo}[Live, Ly6G⁻,CD11b⁺] cells at late time points. Wounds were created using a 4mm punch biopsy on the back of male C57BL/6 normal diet/control and high-fat diet (HFD)/DIO mice. Wounds were harvested on days 1-7 postinjury and analyzed by flow cytometry. (A) Live, lin⁻, Ly6G, CD11b⁺ cell counts in DIO and control wounds based on Ly6C^{Hi} or Ly6C^{Lo} plotted for 7 days post-wounding. (B) Representative plot of live, lin⁻, Ly6G, CD11b⁺ cell counts in DIO and control wounds. All days post-wounding demonstrate no change in live, lin⁻, Ly6G, CD11b⁺ cell counts (data not shown). (**P* < 0.05; n =30 mice; tissues of 2 wounds per mouse were pooled for a single biological replicate. Data is representative of 3 independent experiments.) All data are expressed as the mean +/- the standard error of the means (SEM).

Supplemental Fig IV



Supplemental Figure IV. Percentage of wound resident macrophages (CD11b⁺/F/4/80⁺) are similar between diabetic and control wounds. Wounds were created using a 4mm punch biopsy on the back of male C57BL/6 normal diet/control and high-fat diet (HFD)/DIO mice. Wounds were harvested on day 3 post-injury and analyzed by flow cytometry. Gating strategy to select single, live, lin⁻,Ly6G⁻, CD11b⁺, F4/80⁺ cells. (n =10 mice; tissues of 2 wounds per mouse were pooled for a single biological replicate. Data is representative of 3 independent experiments.) All data are expressed as the mean +/- the standard error of the means (SEM).



Supplemental Figure V. Ly6C^{Lo}[Live, Ly6G⁻,CD11b⁺] cells are increased at late time points in control compared to DIO wounds. Wounds were created using a 4mm punch biopsy on the back of male C57BL/6 normal diet/control and high-fat diet (HFD)/DIO mice. Wounds were harvested on days 1-7 post-injury and analyzed by flow cytometry. The percentage of Ly6C^{Lo}[Live, Ly6G⁻,CD11b⁺] cells in DIO and control wounds were plotted over time. (*P < 0.05, ***P < 0.001; n =30 mice; tissues of 2 wounds per mouse were pooled for a single biological replicate. Data is representative of 2 independent experiments.) All data are expressed as the mean +/- the standard error of the means (SEM).

Supplemental Fig VI



Supplemental Figure VI. Peripheral blood Ly6C^{Hi}[Lin⁻ Ly6G CD11b⁺] cells are decreased in diabetic mice during the inflammatory phase of healing. C57BL/6 mice were either anesthetized alone or anesthetized and wounded and peripheral blood was collected on post-injury day 3 for flow cytometry. (A) Representative flow plots of peripheral blood Ly6C^{Hi}[Lin⁻Ly6G⁻CD11b⁺] cells in wounded and unwounded control mice on post-injury day 3 and comparison of percentage of CD11b⁺ cells. C57BL/6 mice were fed high fat chow (60%kCal) for 12-16 weeks to induce obesity and insulin resistance/glucose intolerance in the diet-induced obese model (DIO) of physiologic type 2 diabetes. Peripheral blood was collected on post-injury day 3, to correspond with the second influx of Ly6C^{Hi}[Lin-Ly6G⁻CD11b⁺] cells in DIO wounds, and analyzed by flow cytometry. (B) Representative flow plots and comparison of DIO and control Ly6C^{Hi}[Lin⁻Ly6G⁻ CD11b⁺] peripheral blood cells on day 3 (*P < 0.05; n = 10 mice. Data is representative of 2 experiments). All data are expressed as mean +/- the standard error of the mean (SEM).



Supplemental Figure VII. MCP-1 expression is similar in diabetic and control wound lymphocytes, neutrophils, T cells, B cells and structural cells. C57BL/6 mice were fed high fat chow (60%kCal) for 12-16 weeks to induce obesity and insulin resistance/glucose intolerance in the diet-induced obese model (DIO) of physiologic type 2 diabetes. Wounds were harvested on day 5 and neutrophils (Ly6G+), lymphocytes (CD3+), T cells (CD4+), B cells (CD19+) and structural cells CD11b-CD3-CD19-Ly6G- were isolated using magnetic-activated cell sorting (MACs). MCP-1 gene expression was compared between DIO and control wound cells (n = 5, repeated one time). All data are expressed as mean +/- the standard error of the means (SEM).



Supplemental Figure VIII. Peripheral blood Ly6C^{Hi}[Live, Ly6G⁻,CD11b⁺] cells are decreased and Ly6C^{Lo}[Live, Ly6G⁻,CD11b⁺] cells increased following anti-MCP1 injection. C57BL/6 mice were fed high fat chow (60%kCal) for 12-16 weeks to induce obesity and insulin resistance/glucose intolerance in the diet-induced obese model (DIO) of physiologic type 2 diabetes. Peripheral blood was collected on post-injury day 5, to correspond with the second influx of Ly6C^{Hi}[Live, Ly6G⁻,CD11b⁺] cells in DIO wounds, and analyzed by flow cytometry. Ly6C^{Hi}[Live, Ly6G⁻,CD11b⁺] and Ly6C^{Lo}[Live, Ly6G⁻,CD11b⁺] peripheral blood cells were plotted in DIO and control mice on day 5 post-wounding. (*P <0.05; n = 10 mice; tissues of 2 wounds per mouse were pooled for a single biological replicate. Data is representative of 2 independent experiments.) All data are expressed as mean +/- the standard error of the means (SEM).

Supplemental Table I

Major Resources Tables

Animals

Species/Strain	Vendor or Source	Background Strain	Sex	
Wild-type	Jackson Laboratory,	C57BL/6	Male	
	Bar Harbor, ME			
mT/mG mice	Jackson Laboratory,	C57BL/6	Male	
(Gt(ROSA)26Sor ^{tm4(ACTB-}	Bar Harbor, ME			
tdTomato,-EGFP)Luo/J)				

Antibodies

Target antigen	Vendor or Source	Catalog #	Working concentration
Anti-CD16/32	BioXCell	CUS-HB-197	1:200 dilution
Anti-CD3	Biolegend	100304	1:400 dilution
Anti-CD19	Biolegend	115504	1:400 dilution
Anti-Ter-119	Biolegend	116204	1:400 dilution
Anti-NK1.1	Biolegend	108704	1:400 dilution
Anti-Ly6G	Biolegend	127604	1:400 dilution
Anti-CD11b	Biolegend	101230	1:400 dilution
Anti-Ly6C	Biolegend	128035	1:400 dilution
Anti-F4/80	Biolegend	123121	1:400 dilution
Streptavidin	Biolegend	405208	1:1,000 dilution
Anti-IL1β	eBioscience	25-7114-82	1:200 dilution
Anti-TNFα	Biolegend	506308	1:200 dilution

Supplemental Table I.

List of major experimental resources and associated vendors.

Supplemental Table II

LV6CL0 down	n in HEI	,		Ly6C ^{Hi}	down in HE	D.	
Lyoc down	loaFC	p value	FDR	Lyoc	logFC	p value	FDR
7fn963	33	0.003256	0 9991 33332	Cd163	25	0.033831588	1
Hist1h2bb	3.6	0.020722	1	Cd5l	2.5	0.204814934	1
Col14a1	3.7	0.059306	1	Hbb-bs	2.8	0.031606452	1
8030453O22Rik	3.8	0.006858	1	Prq4	3.0	0.000583966	0.560786763
Gm42879	3.8	0.003071	0.999133332	Cd300ld2	3.2	0.004771237	1
XIr	4.0	0.004173	1	Lvve1	3.6	0.061650162	1
Gm10643	4.1	0.000839	0.498757072	Sparc	4.4	0.015034378	1
Kif20b	5.2	0.000005	0.029665105	Col3a1	5.5	6.82E-06	0.028364945
				Col1a1	5.6	1.37E-05	0.042774789
				Gm1821	5.8	0.000777029	0.692888023
				Hi			
суос ир п	JogEC	n value	FDR	Ly6C'''	up in HFD		
Cm1921	6 1	0.000406972	0.492046771		- logFC	p value	FDR
Gm1821 Cm12229	-0.1	0.000496872	0.483046771	Ltf	-5.6	3.10E-06	0.025536425
GIII15250 Kr+7	-4.5	0.000002217	0.493004309	Krt7	-5.4	0.001084455	0.846146287
Eafr	-4.5	0.000977739	0.000122222	Aox3	-5.4	0.031953751	1
Light1a2	-3.0	0.005261557	1	Gm37949	-5.3	0.00026188	0.412374168
Ogt122	-5.4	0.005566676	0.095256072	Gm17233	-5.0	0.012716922	1
GIII11950 Cm26415	-5.4	0.00262436	0.965550972	Gm43787	-4.5	0.006127952	1
GI120415	-5.2	0.002590899	0.960145091	Gbp4	-4.2	0.000270884	0.412374168
LOX	-3.2	0.008640524	1	Cecr6	-4.0	0.010596233	1
Aum Dufu4	-5.2	0.0391392	1	Ngp	-4.0	4.14E-05	0.103386371
Kuly4 Tup22a	-5.0	0.019000325	1	Grin3a	-3.9	0.017373863	1
Sko2	-3.0	0.01230331	1	Boc	-3.9	0.003593002	1
Gm37/100	-2.9	0.007397023	1	Gm10612	-3.7	0.013121808	1
Gm/52//	-2.9	0.010037073	1	Gm14212	-3.7	0.00017858	0.371565154
Svcn2	-2.0	0.031300032	1	9130230N0	9Rik -3.7	0.001776599	1
Col3a1	-2.0	0.020037332	1	Zfp229	-3.6	0.000992756	0.826237643
Gm25613	-2.8	0.002781079	0 985356972	Ms4a4b	-3.6	0.004683427	1
Gm43924	-2.8	0.00223108	0.928426824	Tktl1	-3.6	0.006588717	1
Gm5483	-2.8	0.018366815	1	Zfp760	-3.4	0.002508534	1
4933433G15Rik	-2.7	0.001209824	0.580901837	Gm37745	-3.3	0.021465555	1
Chil1	-2.7	0.019059383	1	Gm38020	-3.3	0.017924297	1
Gm22478	-2.7	0.056445751	1	Gm37655	-3.3	0.039229686	1
A730017L22Rik	-2.6	0.003542641	1	Sgo1	-3.2	0.001601303	1
II23a	-2.5	2.92E-06	0.029665105	Gm13/85	-3.2	0.013531473	1
				Gm43961	-3.2	0.00886///3	1
				Angpti3	-3.2	0.004600298	1
				Ugt1a6a	-3.1	0.006962539	
				Gm234/9	-3.1	0.014062972	1
				Capiti	-3.1	0.030575049	1
				GIII37700	-3.1	0.004005987	1
				AW140134	-3.0	0.004871870	0 412274169
				A620072L1	-3.0 OBil: 2.0	0.00029729	1
				Gm44227	2 O	0.034333499	1
				Gm44237	-3.0	0.013400721	1
				Gm42701	-3.0	0.010217434	1
ntral ways	4			Drr33	-2.9	0.020005277	1
muloi wouli	u			Gm13146	-2.9	0.013396669	1
16G- CD11h	+1 cells	Different	ial	Gm26530	-2.9	0.052921919	1
			Hnad	-2.8	0.019591682	1	
using the common dispersion			Gm14984	-2.7	0.006645571	1	
			Gen1	-2.7	0.004783536	1	
itially expressed genes between			Ptas2os	-2.7	0.026641216	1	
highor and n	متنادين	of <0.05 -	aro	II12a	-2.6	0.033812844	1
inginer ann p	value	01 \0.03 a		Cep72	-2.6	0.016313818	1
				Ppl	-2.6	0.009891927	1
				Mast4	-2.6	0.014963024	1
				Gm6542	-2.6	0.011599255	1
				Mest	-2.5	0.044752346	1
				BC035044	-2.5	0.09639372	1

Supplemental Table II.

Comparison was made between DIO and control wound

Ly6C^{Hi}[Live,Ly6G,CD11b⁺] and Ly6C^{Lo}[Live, Ly6G⁻,CD11b⁺] cells. Differential gene expression was determined in edgeR using the common dispersion coefficient. Due to a low number of differentially expressed genes between groups, genes with a fold change of 2.5 or higher and p value of <0.05 are shown.