

**Table S1. Primers sequences for siRNA**

Gene	Sense	Antisense
Jak2	GGAACAUAUUGGGUGGAAAATT	UUUUCACCAAAUAGUUCCTT
Stat1	AGAAGGAGCUGGACAGUAATT	UUACUGUCCAGCUCCUUCUTT
Pfkp	CGUGCACUUGACAGAGAAATT	UUUCUCUGUCAAGUGCACGTT
Negative Control	UUCUCCGAACGUGUCACGUTT	ACGUGACACGUUCGGAGAATT

**Table S2. Antibodies used in Western blot (WB), IHC and IF**

Name	Vendor	Cat#	Dilution ratio
β-actin	Affinity	AF7018	1:1000 (WB)
PCK	MCE	HY-P81164	1:200 (IF)
α-SMA	Affinity	AF1032	1:200 (IF)
CD4	Immunoway	YT0762	1:200 (IF)
F4/80	Affinity	DF2789	1:200 (IF)
HMGB1	Affinity	AF7020	1:1000(WB); 1:200 (IHC, IF)
IL-18	Affinity	DF6252	1:1000 (WB)
GSDMD	Affinity	AF4012	1:1000 (WB)/1:200(IF)
Caspase1	Affinity	AF5418	1:1000 (WB)/1:200(IF)
Jak2	Immunoway	YT2426	1:1000 (WB)
p-Jak2	Immunoway	YP0155	1:1000 (WB)
Stat1	Cell Signaling Techology	9172T	1:1000 (WB)
p-Stat1	Immunoway	YP0249	1:1000 (WB)/1:200(IF)
PfkP	Proteintech	13389-1-AP	1:1000 (WB)/1:200(IF)
iNOS	Affinity	AF0199	1:1000 (WB)/1:200(IF)
Arg 1	Affinity	DF6657	1:1000 (WB)/1:200(IF)
NLRP3	Affinity	DF7438	1:1000 (WB)
ASC	Affinity	DF6304	1:1000 (WB)

**Table S3. Primers used in RT-qPCR analysis**

Gene	Forward primer sequence	Reverse primer sequence
β-actin	GGCTGTATTCCCCTCCATCG	CCAGTTGGTAACAATGCCATGT
IL-1 $\beta$	TGGGAAACAACAGTGGTCAGG	CCATCAGAGGAAGGAGGAA
IL-6	TAGTCCTCCTACCCCAATTCC	TTGGTCCTTAGCCACTCCTTC
TNF- $\alpha$	GAGTGACAAGCCTGTAGCC	CTCCTGGTATGAGATAGCAAA
iNOS	CACCAAGCTGAAC TGAGCG	CGTGGCTTGGGCTCCTC
Arg-1	CCAGAAGAATGGAAGAGTCAGTGT	GCAGATATGCAGGGAGTCACC
HMGB1	GCTGACAAGGCTCGTTATGAA	CCTTGATTTGGGCGGTA
Stat1	TCACAGTGGTTCGAGCTTCAG	CGAGACATCATAGGCAGCGTG
Pfkp	CGCCTATCCGAAGTACCTGGA	CCCCGTGTAGATTCCCATGC

**Table S4. Primers sequences for ChIP-qPCR**

Gene	Foward	Reverse
Pfkp site1	CAAAGTGAGTTCCAGGACAGC	TTGGGGGTGGCTTGTAA
Pfkp site2	AGTCTGACACCCCTCCTCC	TGATGAGCACTAAACTGGAGAA
Pfkp site3	TCAGATGTTCTCCAGTTAGTGC	TCCTGAACCTCCATGATCAAC

**Figure S1. Flow cytometry gating strategy and polarization status of M2-type macrophages.**

- (A) Flow cytometry gating strategy for M1 polarization of splenic macrophages in EAP mice;
- (B) Flow cytometry gating strategy for M1 polarization of macrophages in iBMDM;
- (C) Flow cytometry gating strategy for M2 polarization of macrophages in iBMDM;
- (D-G) The effects of dsHMGB1 and Flu on M2 polarization of iBMDM. \* $P < 0.05$ ; \*\* $P < 0.01$ ;  
\*\*\* $P < 0.001$ .

**Figure S2. si-Pfkp inhibits dsHMGB1-induced M1 macrophage polarization.**

- (A-C) The expression of Pfkp in the si-NC group and the si-Pfkp group was detected by RT-qPCR and Western blot. (D) The expression of IL-1 $\beta$ , IL-6, TNF- $\alpha$ , iNOS and Arg1 at RNA level between si-NC+dsHMGB1 and si-Pfkp +dsHMGB1 groups based on macrophage; (E-F) The protein level of iNOS and Arg1 between the si-NC+dsHMGB1 and si-Pfkp +dsHMGB1 groups was detected by western blot; (G-H) Flow cytometry was used to detect the M1 polarization of macrophages between si-NC+dsHMGB1 and si-Pfkp+dsHMGB1 groups; (I) Lactate levels between the si-NC+dsHMGB1 and si-Pfkp +dsHMGB1 groups; (J) 2-NBDG was used to measure glucose uptake between the si-NC+dsHMGB1 and si-Pfkp +dsHMGB1 groups. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .

**Figure S3. si-Jak2 inhibits dsHMGB1-induced M1 macrophage polarization.**

- (A-C) The expression of Jak2 in the si-NC group and the si-Jak2 group was detected by RT-qPCR and Western blot. (D) The expression of IL-1 $\beta$ , IL-6, TNF- $\alpha$ , iNOS and Arg1 at RNA

level between si-NC+dsHMGB1 and si-Jak2+dsHMGB1 groups based on macrophage; (E-F) The protein level of p-Stat1, Stat1, Pfkp, iNOS and Arg1 between the si-NC+dsHMGB1 and si-Jak2+dsHMGB1 groups was detected by western blot; (G-H) Flow cytometry was used to detect the M1 polarization of macrophages between si-NC+dsHMGB1 and si-Jak2+dsHMGB1 groups; (I)Lactate levels between the si-NC+dsHMGB1 and si-Jak2+dsHMGB1 groups; (J) 2-NBDG was used to measure glucose uptake between the si-NC+dsHMGB1 and si-Jak2+dsHMGB1 groups.\* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .

**Figure S4. si-Stat1 inhibits dsHMGB1-induced M1 macrophage polarization.**

(A-C) The expression of Stat1 in the si-NC group and the si-Stat1 group was detected by RT-qPCR and Western blot; (D) The expression of IL-1 $\beta$ , IL-6, TNF- $\alpha$ , iNOS and Arg1 at RNA level between si-NC+dsHMGB1 and si-Stat1+dsHMGB1 groups based on macrophage; (E-F) The protein level of Pfkp, iNOS and Arg1 between the si-NC+dsHMGB1 and si-Stat1+dsHMGB1 groups was detected by western blot; (G-H) Flow cytometry was used to detect the M1 polarization of macrophages between si-NC+dsHMGB1 and si-Stat1+dsHMGB1 groups; (I)Lactate levels between the si-NC+dsHMGB1 and si-Stat1+dsHMGB1 groups; (J) 2-NBDG was used to measure glucose uptake between the si-NC+dsHMGB1 and si-Stat1+dsHMGB1 groups.\* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .

Figure S1

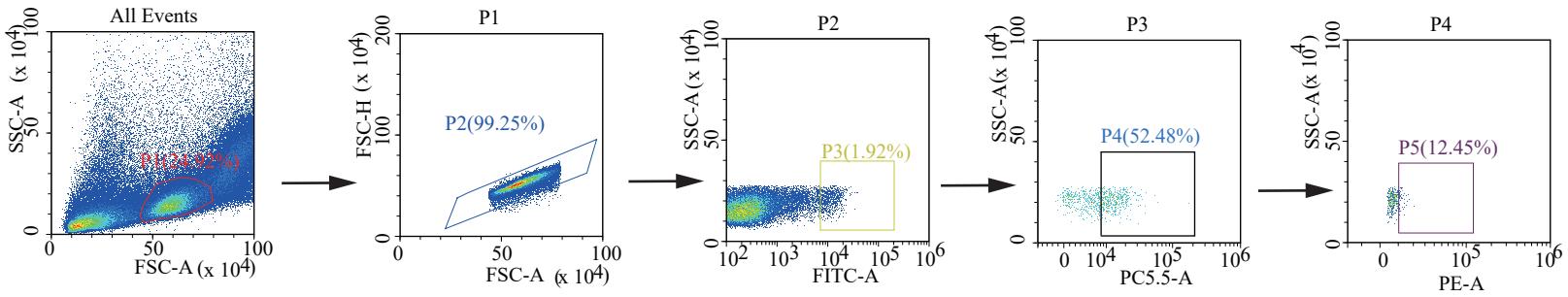
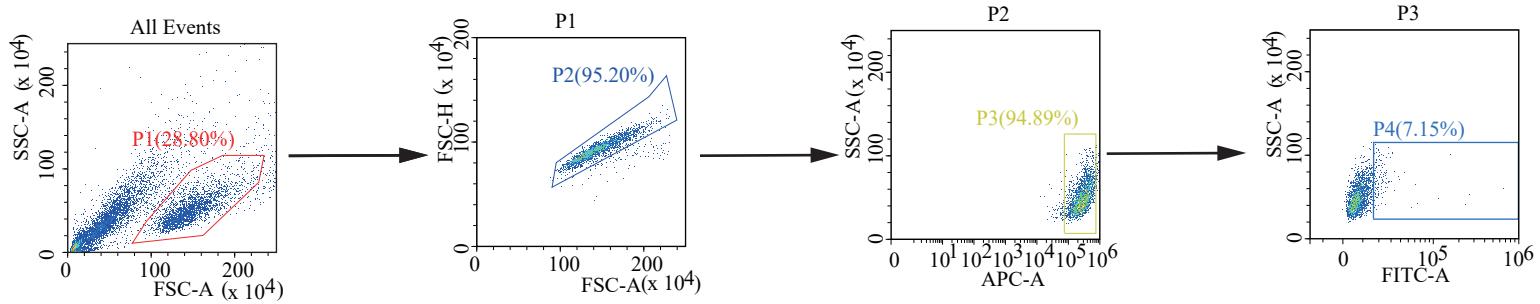
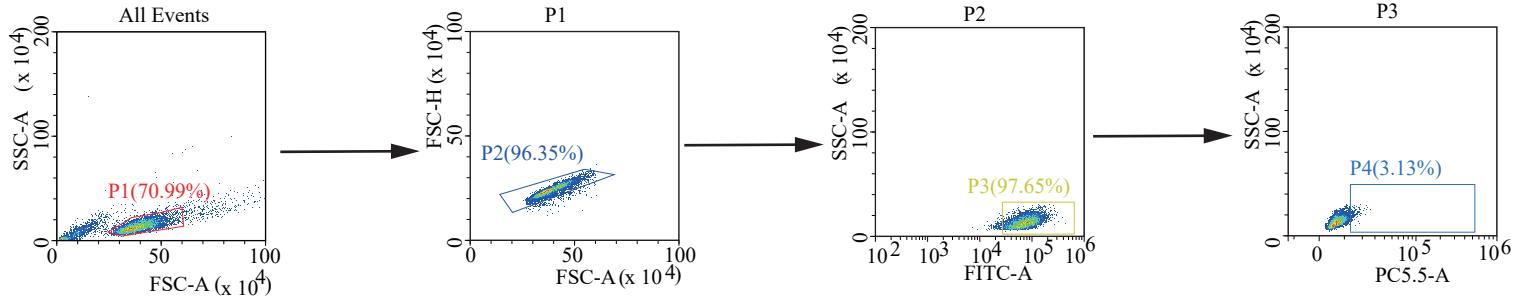
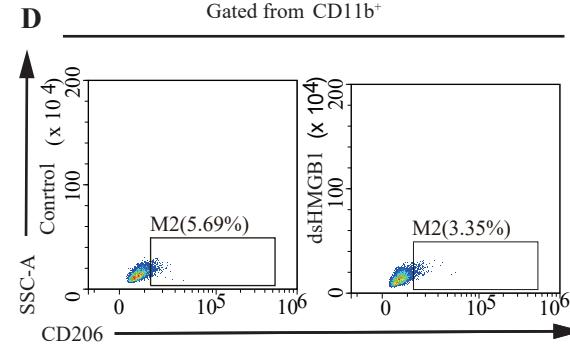
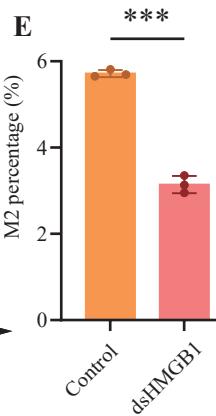
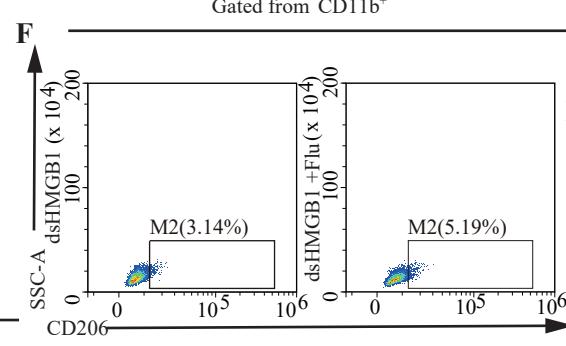
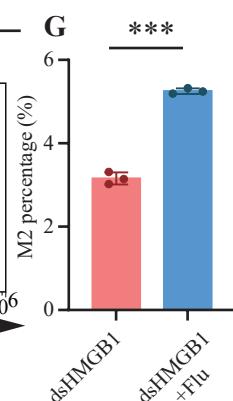
**A****B****C****D****E****F****G**

Figure S2

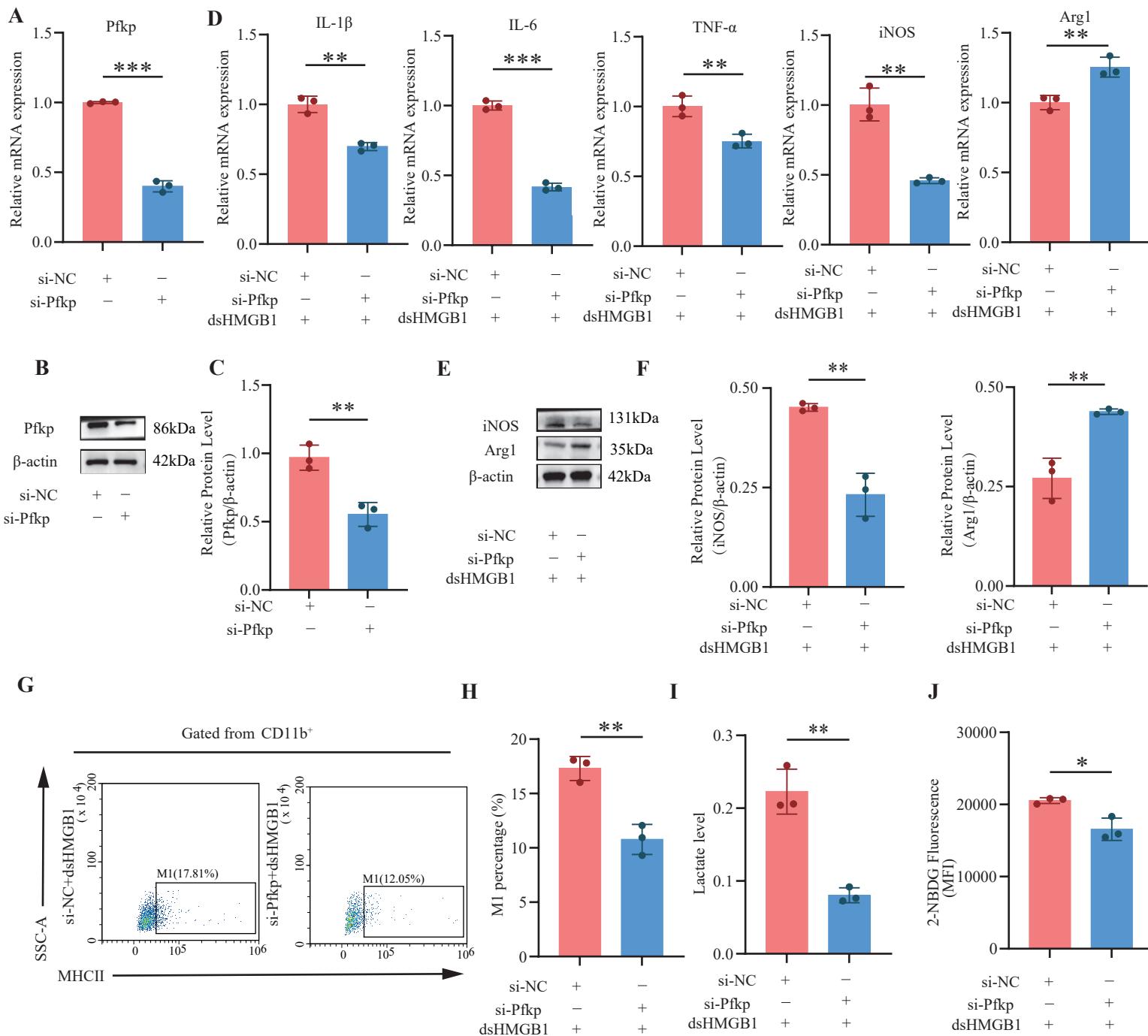


Figure S3

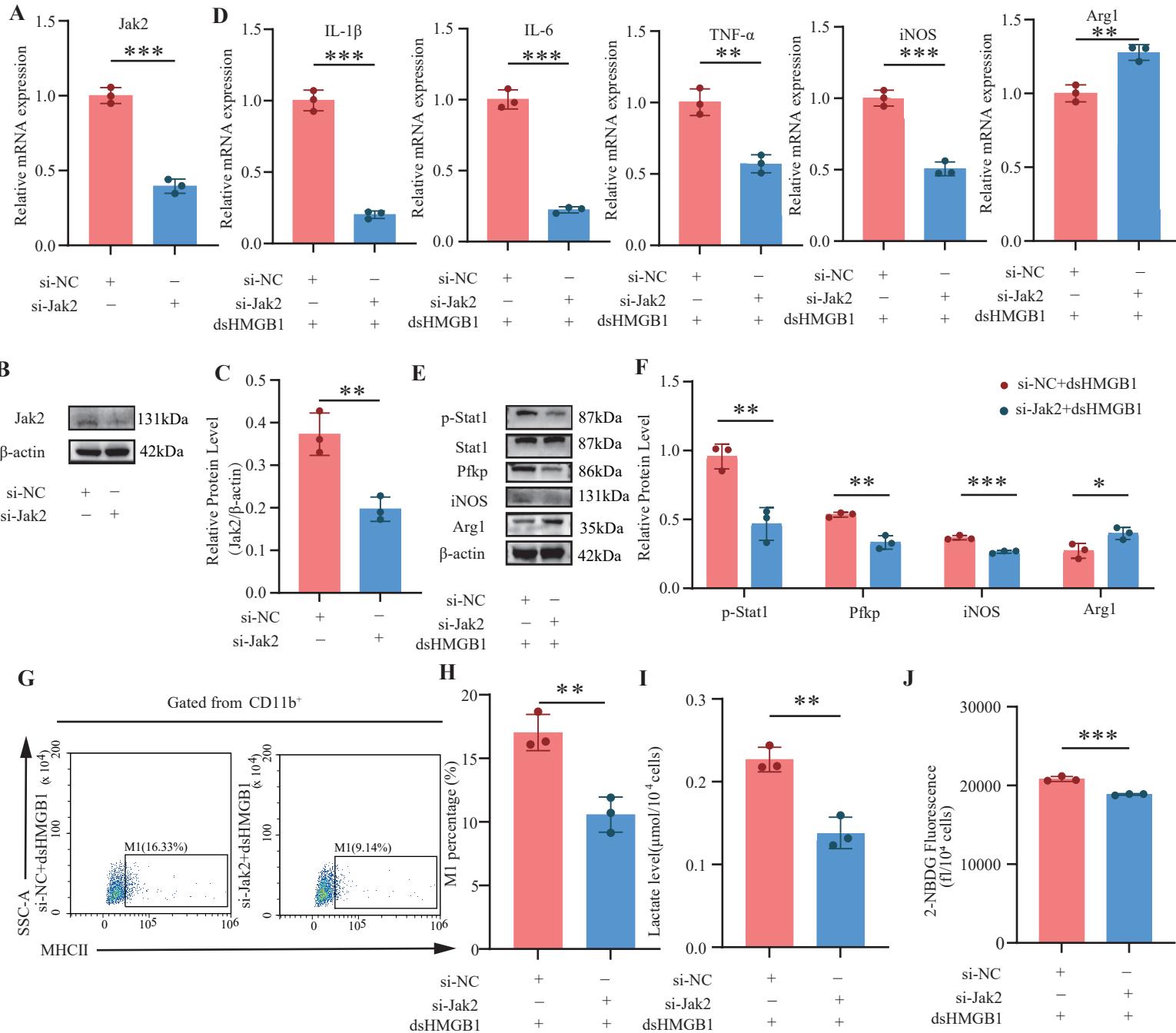


Figure S4

