

**Table S1** Antibodies used in Western blotting (WB), immunofluorescence, and flow cytometry

<b>Antibody</b>	<b>Origin</b>	<b>Catalog number</b>	<b>Assay</b>
CD41a monoclonal antibody (eBioMWRReg30 (MWRReg30)), APC	eBioscience	Cat# 17-0411-82	FC
CD42d monoclonal antibody (1C2), APC	eBioscience	Cat# 17-0421-82	FC
CD42d monoclonal antibody (1C2), PE	eBioscience	Cat# 12-0421-82	FC
CD41a monoclonal antibody [eBioMWRReg30 (MWRReg30)], FITC	eBioscience	Cat# 11-0411-85	FC
CD41a monoclonal antibody [eBioMWRReg30 (MWRReg30)], Biotin	eBioscience	Cat# 13-0411-82	FC
PE anti-mouse/rat CD62P (P-selectin) antibody	Biologend	Cat# 148306	FC
Integrin alpha IIb beta 3 (GPIIb/IIIa) activation, PE	Emfret Analytics	Cat# M023-2	FC
Biotin mouse hematopoietic lineage cocktail	eBioscience	Cat# 88-7774-75	FC
Brilliant Violet 510™ Streptavidin	Biologend	Cat# 405234	FC
Ly-6A/E (Sca-1) monoclonal antibody (D7), PerCP-Cyanine5.5	eBioscience	Cat# 45-5981-82	FC
CD117 (c-Kit) monoclonal antibody (2B8), APC-eFluor™ 780	eBioscience	Cat# 47-1171-82	FC
CD150 monoclonal antibody (mShad150), PE-Cyanine7	eBioscience	Cat# 25-1502-82	FC
CD45 monoclonal antibody (HI30), eFluor 450	eBioscience	Cat# 48-0459-42	FC
CD115 (c-fms) monoclonal antibody (AFS98), PE-Cyanine7	eBioscience	Cat# 25-1152-80	FC
CD11b monoclonal antibody (M1/70), APC	eBioscience	Cat# 17-0112-82	FC
Ly-6G/Ly-6C (Gr-1) monoclonal antibody (RB6-8C5), PE	eBioscience	Cat# 12-5931-82	FC
APC anti-mouse CD80	eBioscience	Cat# 17-0801-82	FC
PerCP-Cyanine5.5 anti-mouse CD11c	eBioscience	Cat# 45-0114-82	FC
Anti-human CD41 antibody, APC	Biologend	Cat# 303710	FC
Anti-human CD62P (P-Selectin) antibody, PE	Biologend	Cat# 304906	FC

<b>Antibody</b>	<b>Origin</b>	<b>Catalog number</b>	<b>Assay</b>
Annexin V- eFluor 450 apoptosis detection kit	eBioscience	Cat# 88-8006-74	FC
Bcl-xL monoclonal antibody (C.85.1)	Thermo Fisher Scientific	Cat# MA5-15142	FC
MCL1 recombinant rabbit monoclonal antibody (SI16-04)	Thermo Fisher Scientific	Cat# MA5-32060	FC
BAX monoclonal antibody (6A7)	Thermo Fisher Scientific	Cat# MA5-14003	FC
BAK1 monoclonal antibody (4C2)	Thermo Fisher Scientific	Cat# MA5-36225	FC
Cleaved caspase-3 (Asp175) (D3E9) rabbit mAb (Alexa Fluor® 647 Conjugate)	Cell Signaling Technology	Cat# 9602S	FC
Alexa Fluor® 647 anti-cytochrome c antibody	Biolegend	Cat# 612310	FC
TBK1 antibody	Abcam	Cat# ab40676	WB
p-TBK1 (Ser172) antibody	Cell Signaling Technology	Cat# 5483	WB, IF
p-IRF3 (Ser396) antibody	Cell Signaling Technology	Cat# 29047	WB, IF
IRF3 antibody	Cell Signaling Technology	Cat# 4302	WB
p-TBK1 (Ser172) antibody (Alexa Fluor 647 Conjugate)	Cell Signaling Technology	Cat# 14590	FC
p-IRF3 (Ser396) antibody (Alexa Fluor 647 Conjugate)	Cell Signaling Technology	Cat# 10327	FC
IFN- $\beta$ antibody	Abcam	Cat# ab85803	WB, IF, FC
Anti-IFN- $\beta$ (EPR22186-266)	Abcam	Cat# ab218229	WB
IFN- $\beta$ antibody (MIB-8C4.1)	Santa Cruz Biotechnology	Cat# sc-53586	FC, IF
p-STAT1 (Tyr701) antibody	Cell Signaling Technology	Cat# 9167	WB, IF
STAT1 antibody	Cell Signaling Technology	Cat# 14994	WB
p-STAT1 (Ser727) antibody (Alexa Fluor 647 Conjugate)	Biolegend	Cat# 686412	FC
Anti-GBP2 antibody	Abcam	Cat# ab203238	WB
GBP2 polyclonal antibody	Thermo Fisher Scientific	Cat# PA5-112426	FC

<b>Antibody</b>	<b>Origin</b>	<b>Catalog number</b>	<b>Assay</b>
GAPDH (D16H11) XP <sup>®</sup> Rabbit mAb	Cell Signaling Technology	Cat# 5174S	WB
cGAS antibody	Abcam	Cat# ab224144	IF
cGAS antibody	Santa Cruz Biotechnology	Cat# sc-515777	IF
Tom20 antibody	Santa Cruz Biotechnology	Cat# sc-17764	IF
LC3B antibody	Novus Biologicals	Cat# NB100-2220	WB, IF
DNA antibody	Millipore	Cat# CBL186	IF
Monoclonal ANTI-Flag <sup>®</sup> M2 antibody	Sigma-Aldrich	Cat# F1804	ChIP
Goat anti-mouse IgG (H + L) cross-adsorbed secondary antibody, Alexa Fluor 488	Thermo Fisher Scientific	Cat# A-11001	IF
Goat anti-mouse IgG (H + L) cross-adsorbed secondary antibody, Alexa Fluor 546	Thermo Fisher Scientific	Cat# A-11003	IF
Goat anti-rabbit IgG (H + L) cross-adsorbed secondary antibody, Alexa Fluor 488	Thermo Fisher Scientific	Cat# A-11008	IF
Goat anti-rabbit IgG (H + L) cross-adsorbed secondary antibody, Alexa Fluor 546	Thermo Fisher Scientific	Cat# A-11010	IF
Goat anti-rabbit IgG (H + L) cross-adsorbed secondary antibody, Alexa Fluor 647	Thermo Fisher Scientific	Cat# A-21244	FC
Goat anti-mouse IgG (H + L) cross-adsorbed secondary Antibody, Alexa Fluor 647	Thermo Fisher Scientific	Cat# A-21235	FC
HRP labeled goat anti-rabbit IgG (H + L)	Abcam	Cat# 205718	WB
HRP labeled goat anti-Mouse IgG (H + L)	Abcam	Cat# 205719	WB

*IF* immunofluorescence, *FC* flow cytometry, *ChIP* chromatin immunoprecipitation, *LC3B* microtubule-associated protein 1 light chain 3B, *TOM20* translocase of outer mitochondrial membrane 20, *cGAS* cyclic GMP-AMP synthase, *STING* stimulator of interferon genes, *TBK1* TANK-binding kinase 1, *IRF3* interferon regulatory factor 3, *IFN-β* interferon-β, *STAT1* signal transducer and activator of transcription 1, *GBP2* guanylate-binding protein 2, *GAPDH* glyceraldehyde-3-phosphate dehydrogenase, *FITC* fluorescein isothiocyanate, *PE* phycoerythrin, *APC* allophycocyanin, *PerCP* peridinin chlorophyll protein, *IgG* Allophycocyanin, *HRP* horseradish peroxidase

**Table S2** Primer sequences for mitochondrial DNA (mtDNA) and nuclear DNA (nDNA) analysis

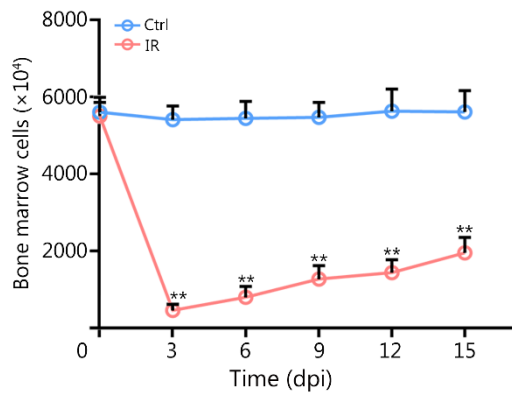
<b>Origin</b>	<b>Gene</b>	<b>Forward primer (5' - 3')</b>	<b>Reverse primer (5' - 3')</b>
mtDNA (mouse)	<i>D-loop</i>	AATCTACCATCCTCCGTGAAACC	TCAGTTTAGCTACCCCAAGTTAA
	<i>Cox1</i>	GCCCCAGATATAGCATTCCC	G TTCATCCTGTTCTGCTCC
	<i>ND4</i>	AACGGATCCACAGCCGTA	AGTCCTCGGGCCATGATT
mtDNA (human)	<i>ND1</i>	CCCTAAAACCCGCCACATCT	GAGCGATGGTGAGAGCTAAGGT
	<i>ND2</i>	ACCATCTTTGCAGGCACACT	GCTTCTGTGGAACGAGGGTT
	<i>ND4</i>	TTCCTCCGACCCCTAACAA	GATAAGTGGCGTTGGCTTGC
nDNA (human)	<i>LINE1</i>	AGAACGCCACAAAGATACTCCTCG	CTCTCTTCTGGCTTGTAGGGTTTCTG
	<i>RNA18S</i>	TGCCCTATCAACTTTCGATGGTAGTC	TTGGATGTGGTAGCCGTTTCTCA

**Table S3** siRNA sequences used for human RNA interference

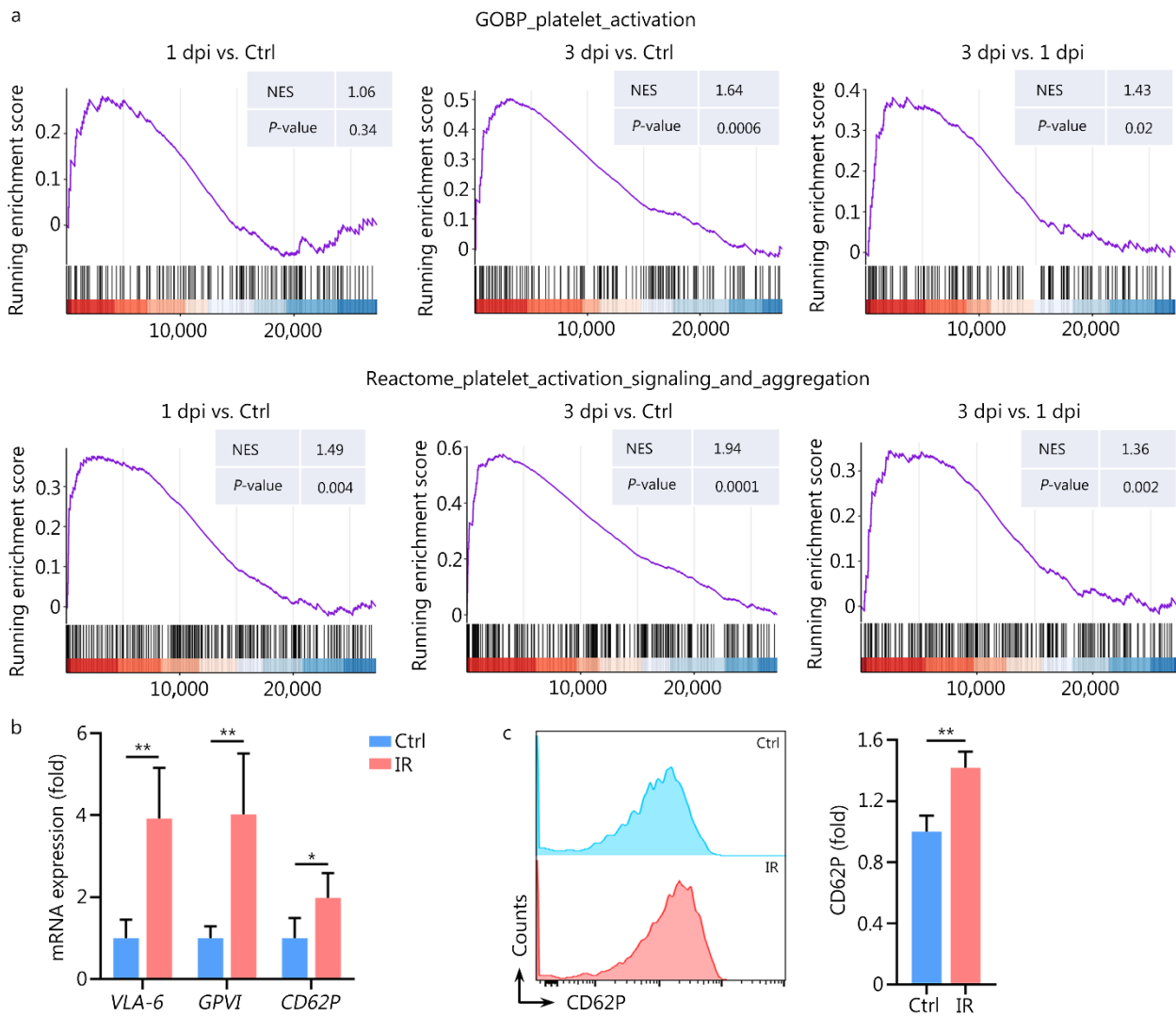
<b>Gene</b>	<b>Sense</b>	<b>Antisense</b>
<i>cGAS</i>	CCAACACUCGUGCAUAAUUATT	UAAUAUGCACGAGUGUUGGTT
<i>STING</i>	GCAUCAAGGAUCGGGUUUATT	UAAACCCGAUCCUUGAUGCTT
<i>IFNAR1</i>	CCUACUUCCUCCAGUCUUUTT	AAAGACUGGAGGAAGUAGGTT
<i>GBP2</i>	GCCCUUUAGAAGAAGAUGUTT	ACAUCUUCUUCUAAAGGGCTT
Negative control	UCCUCCGAACGUGUCACGUTT	ACGUGACACGUUCGGAGAATT

**Table S4** Primer sequences for mRNA expression analysis

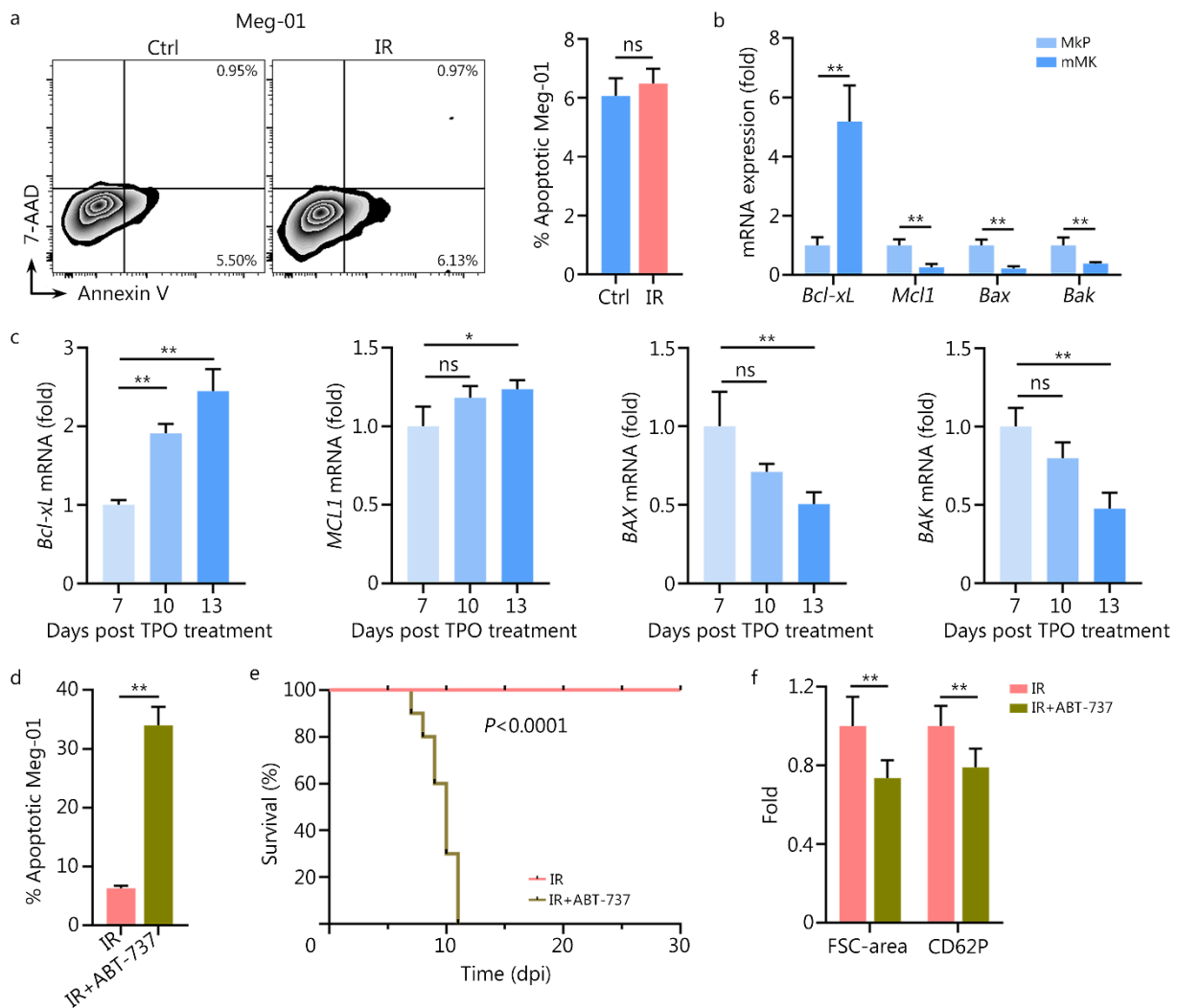
Source	Gene	Forward primer (5' - 3')	Reverse primer (5' - 3')
Human	<i>Bcl-xL</i>	AAAAGATCTTCCGGGGGCTG	TCCACAAAAGTATCCTGTTCAAAGC
	<i>MCL</i>	CGACTTTTGGCCACCGGC	TAGCCAGTCCCGTTTTGTCC
	<i>BAX</i>	TCATGGGCTGGACATTGGAC	GCGTCCCAAAGTAGGAGAGG
	<i>BAK</i>	GGACCCATGCTGGAGTAAGAATAA	TTCTGCTGATGGCGGTAAA
	<i>VLA-6</i>	GGCGGTGTTATGTCCTGAGTC	AATCGCCCATCACAAAAGCTC
	<i>GPVI</i>	TCCCGGCCATGAAGAGAAGT	TTACGTCCCCTCCTGACGAC
	<i>CD62P</i>	ACTGCCAGAATCGCTACACAG	CACCCATGTCCATGTCTTATTGT
	<i>cGAS</i>	ACATGGCGGCTATCCTTCTCT	GGTTCTGGGTACATACGTGAAA
	<i>STING</i>	CCAGAGCACACTCTCCGGTA	CGCATTGGGGAGGGAGTAGTA
	<i>IFNAR1</i>	AACAGGAGCGATGAGTCTGTC	TGCGAAATGGTGTAATGAGTCA
	<i>GBP2</i>	ATTGTGGGCCTCTATCGCAC	CCAGGTGAGGAGTTTGCCTT
	<i>GAPDH</i>	AGCCTCAAGATCATCAGCAA	GTCATGAGTCCTTCCACGATAC
	Mouse	<i>IFN-β</i>	TGGGTGGAATGAGACTATTGTTG
<i>Bcl-xL</i>		GACAAGGAGATGCAGGTATTGG	TCCCGTAGAGATCCACAAAAGT
<i>MCL</i>		AAAGGCGGCTGCATAAGTC	TGGCGGTATAGGTCGTCCTC
<i>BAX</i>		CCGGCGAATTGGAGATGAACT	CCAGCCCATGATGGTTCTGAT
<i>BAK</i>		GTGACCTGCTTTTTGGCTGAT	GGTCTCTACGCAAATTCAGGG
<i>GAPDH</i>		AGGTCGGTGTGAACGGATTTG	TGTAGACCATGTAGTTGAGGTCA



**Fig. S1** Number of bone marrow cells in mice at indicated dpi.  $n = 6$ . Data represent mean  $\pm$  standard deviation. \*\* $P < 0.01$ , compared to Ctrl. Two-tailed unpaired student's  $t$ -test. Ctrl control, IR ionizing radiation, dpi day post IR

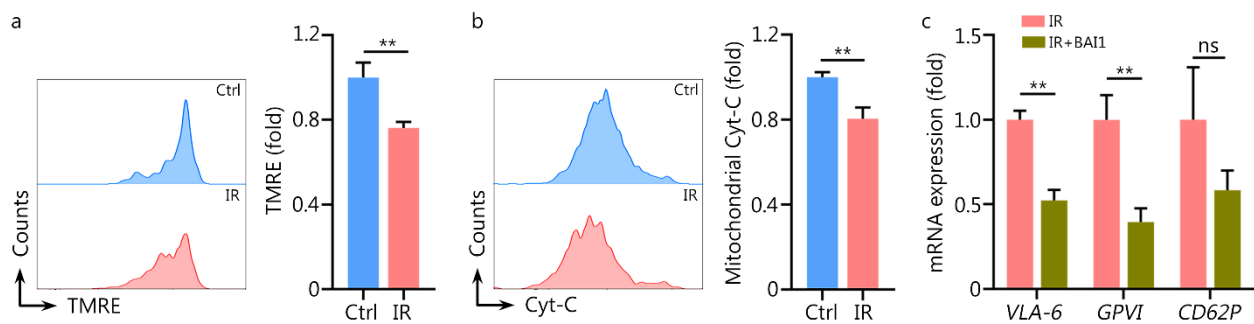


**Fig. S2** Platelets are molecularly and functionally hyperreactive post IR. **a** Gene set enrichment analysis of indicated gene sets in BM mMKs of mice at 1 and 3 dpi. **b** Relative mRNA expression of *VLA-6*, *GPVI*, and *CD62P* in Meg-01 cells at 3 dpi ( $n = 4$ ). **c** Flow cytometric analysis and quantification of CD62P expression in response to thrombin on PLPs from Meg-01 cultures at 3 dpi ( $n = 5$ ). Data represent mean  $\pm$  standard deviation. \* $P < 0.05$ , \*\* $P < 0.01$ , compared to Ctrl. Two-tailed unpaired student's  $t$ -test. mMK mature megakaryocyte, BM bone marrow, Ctrl control, IR ionizing radiation, dpi day post IR, VLA-6 integrin  $\alpha 6\beta 1$ , GPVI glycoprotein VI, PLP platelet-like particle, NES normalized enrichment score, GOBP gene ontology-biological process

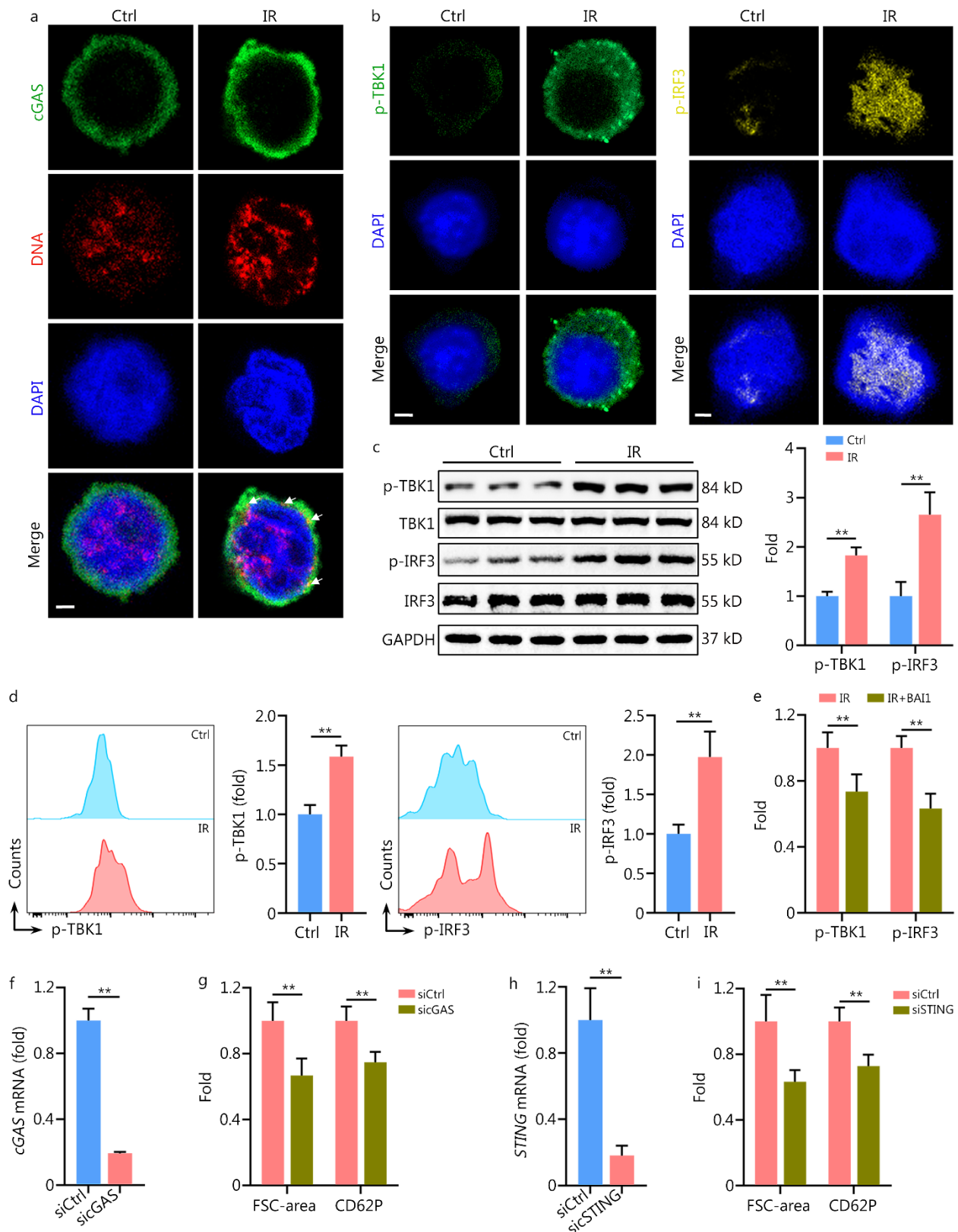


**Fig. S3** The inherently high pro-survival threshold confers radioresistance onto mMKs. **a** Flow cytometric analysis and quantification of Meg-01 apoptosis at 1 dpi ( $n = 5$ ). **b** Relative mRNA expression of pro-survival (*Bcl-xL* and *Mcl1*) and pro-apoptotic (*Bax* and *Bak*) Bcl-2 family in BM MkpPs and mMkPs of mice ( $n = 4$ ). **c** Relative mRNA expression of pro-survival (*Bcl-xL* and *MCL1*) and pro-apoptotic (*BAX* and *BAK*) Bcl-2 family in human primary MKs during TPO-induced maturation ( $n = 3$ ). **d** Flow cytometric quantification of Meg-01 apoptosis with or without ABT-737 treatment in vitro at 1 dpi ( $n = 5$ ). **e** Survival analysis of mice with or without ABT-737 treatment post IR ( $n = 10$ ). **f** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures at 3 dpi ( $n = 5$ ). Data represent mean  $\pm$  standard deviation. ns, non-significance. \* $P < 0.05$ , \*\* $P < 0.01$ , compared to Mkp, Day 7, or IR. Two-tailed unpaired student's  $t$ -test unless stated otherwise. One-way analysis of variance (**b**). Log-rank test (**e**). MK megakaryocyte, mMk mature MK, Mkp MK progenitor, TPO thrombopoietin, Ctrl control, IR ionizing radiation, dpi day post IR, 7-AAD 7-amino-actinomycin D, PLP platelet-like particle, FSC forward scatter



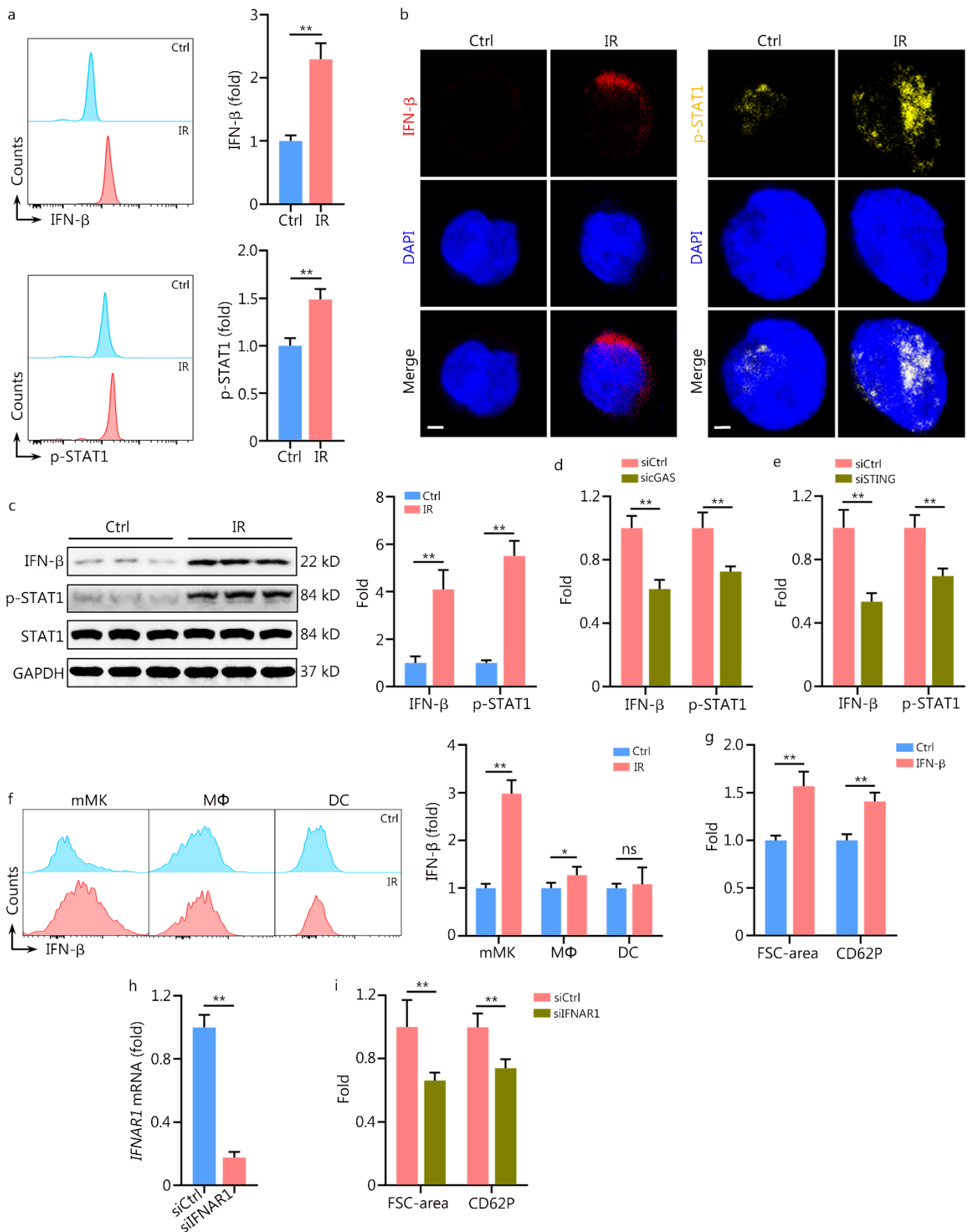


**Fig. S4** Minority MOMP is triggered in Meg-01 post IR. **a** Flow cytometric analysis and quantification of MMP in Meg-01 cells at 1 dpi ( $n = 5$ ). **b** Flow cytometric analysis and quantification of mitochondrial Cyt-C release in Meg-01 cells at 1 dpi. **c** Relative mRNA expression of *VLA-6*, *GPVI*, and *CD62P* in Meg-01 cells with or without BAI1 treatment at 3 dpi ( $n = 3$ ). Data represent mean  $\pm$  standard deviation. \*\* $P < 0.01$ , compared to Ctrl or IR. Two-tailed unpaired student's *t*-test. ns non-significance, Ctrl control, IR ionizing radiation, dpi day post IR, MOMP mitochondrial outer membrane permeabilization, MMP mitochondrial membrane potential, Cyt-C cytochrome C, TMRE tetramethylrhodamine ethyl ester, VLA-6 integrin  $\alpha\beta 1$ , GPVI glycoprotein VI



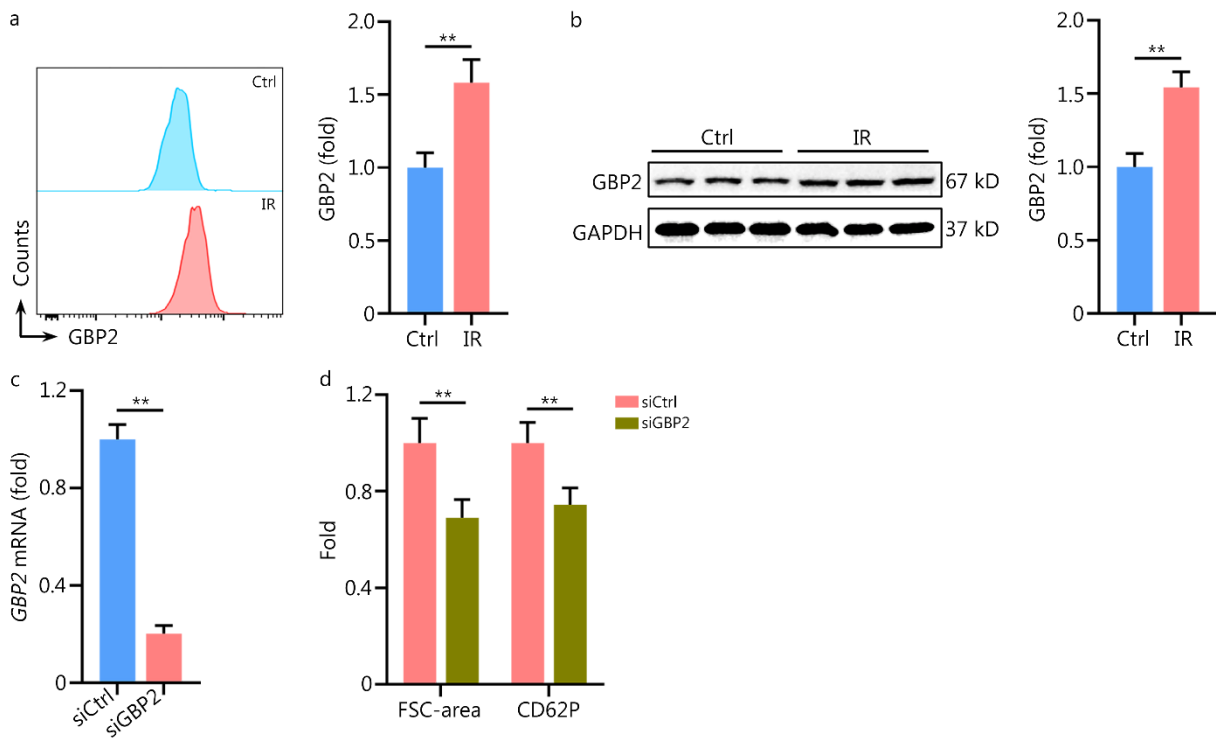
**Fig. S5** Minority MOMP stimulates cGAS/STING in Meg-01 post IR. **a** Colocalization of cytosolic DNA with cGAS in Meg-01 cells at 1 dpi was assessed by immunofluorescence staining with antibodies specific for DNA and cGAS. Scale bar = 20  $\mu$ m. The arrow indicates cGAS and DNA colocalization. **b** p-TBK1 and p-IRF3 expression in Meg-01 cells at 1 dpi was assessed by immunofluorescence staining. Scale bar = 20  $\mu$ m. **c** Western blotting analysis and quantification of p-TBK1 and p-IRF3 expression in Meg-01 cells at 1 dpi ( $n = 3$ ). **d** Flow cytometric analysis and

quantification of p-TBK1 and p-IRF3 expression in Meg-01 cells at 1 dpi ( $n = 5$ ). **e** Flow cytometric quantification of p-TBK1 and p-IRF3 expression in Meg-01 cells with or without BAI1 treatment at 1 dpi ( $n = 5$ ). **f** Relative *cGAS* mRNA expression in Meg-01 cells with or without sicGAS treatment ( $n = 3$ ). **g** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures with or without sicGAS treatment at 3 dpi ( $n = 5$ ). **h** Relative *STING* mRNA expression in Meg-01 cells with or without siSTING treatment ( $n = 3$ ). **i** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures with or without siSTING treatment at 3 dpi ( $n = 5$ ). Data represent mean  $\pm$  standard deviation.  $**P < 0.01$ , compared to Ctrl, IR or siCtrl. Two-tailed unpaired student's *t*-test. MOMP mitochondrial outer membrane permeabilization, cGAS cyclic GMP-AMP synthase, STING stimulator of interferon genes, Ctrl control, IR ionizing radiation, dpi day post IR, TBK1 TANK-binding kinase 1, IRF3 interferon regulatory factor 3, PLP platelet-like particle, FSC forward scatter, DAPI 4',6-diamidino-2-phenylindole, GAPDH glyceraldehyde-3-phosphate dehydrogenase, sicGAS siRNA of cGAS, siSTING siRNA of STING

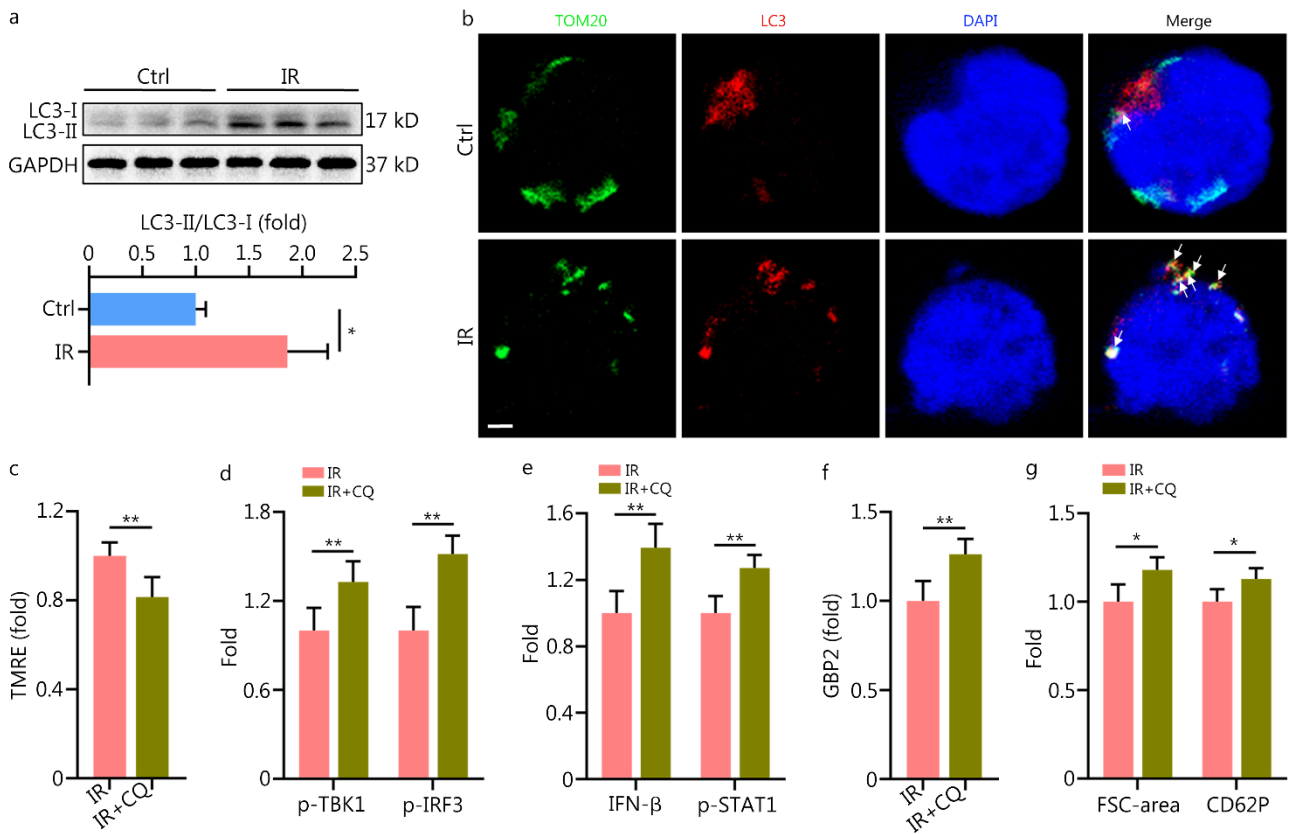


**Fig. S6** IFN- $\beta$  response is triggered in mMKs post IR. **a** Flow cytometric analysis and quantification of IFN- $\beta$  and p-STAT1 expression in Meg-01 cells at 1 dpi ( $n = 5$ ). **b** IFN- $\beta$  and p-STAT1 expression in Meg-01 cells at 1 dpi was assessed by immunofluorescence staining. Scale bar = 20  $\mu$ m. **c** Western blotting analysis and quantification of IFN- $\beta$  and p-STAT1 expression in Meg-01 cells at 1 dpi ( $n = 3$ ). **d** Flow cytometric quantification of IFN- $\beta$  and p-STAT1

expression in Meg-01 cells with or without sicGAS treatment at 1 dpi ( $n = 5$ ). **e** Flow cytometric quantification of IFN- $\beta$  and p-STAT1 expression in Meg-01 cells with or without siSTING treatment at 1 dpi ( $n = 5$ ). **f** Flow cytometric analysis and quantification of IFN- $\beta$  expression in mMK, M $\Phi$  and dendritic cell DC at 1 dpi ( $n = 5$ ). **g** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures with or without IFN- $\beta$  treatment at 3 dpi ( $n = 5$ ). **h** Relative *IFNAR1* mRNA expression in Meg-01 cells with or without siIFNAR1 treatment ( $n = 3$ ). **i** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures with or without siIFNAR1 treatment at 3 dpi ( $n = 5$ ). Data represent mean  $\pm$  standard deviation. \* $P < 0.05$ , \*\* $P < 0.01$ , compared to Ctrl or siCtrl. Two-tailed unpaired student's *t*-test. ns non-significance, IFN- $\beta$  interferon- $\beta$ , Ctrl control, IR ionizing radiation, dpi day post IR, STAT1 signal transducer and activator of transcription 1, cGAS cyclic GMP-AMP synthase, STING stimulator of interferon genes, sicGAS siRNA of cGAS, siSTING siRNA of STING, IFNAR1 IFN- $\alpha/\beta$  receptor 1, siIFNAR1 siRNA of IFNAR1, FSC forward scatter, PLP platelet-like particle, DAPI 4',6-diamidino-2-phenylindole, GAPDH glyceraldehyde-3-phosphate dehydrogenase, M $\Phi$  macrophage, DC dendritic cell



**Fig. S7** IFN-inducible gene GBP2 mediates production of large and hyperreactive platelets post IR. **a** Flow cytometric analysis and quantification of GBP2 expression in Meg-01 cells at 3 dpi ( $n = 5$ ). **b** Western blotting analysis and quantification of GBP2 expression in Meg-01 cells at 3 dpi ( $n = 3$ ). **c** Relative *GBP2* mRNA expression in Meg-01 cells with or without siGBP2 treatment ( $n = 3$ ). **d** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures with or without siGBP2 treatment in the presence of IFN- $\beta$  ( $n = 5$ ). Data represent mean  $\pm$  standard deviation.  $*P < 0.05$ ,  $**P < 0.01$ , compared to Ctrl or siCtrl. Two-tailed unpaired student's *t*-test. IFN interferon, GBP2 guanylate-binding protein 2, Ctrl control, IR ionizing radiation, dpi day post IR, FSC forward scatter, PLP platelet-like particle, siGBP2 siRNA of GBP2, GAPDH glyceraldehyde-3-phosphate dehydrogenase



**Fig. S8** Autophagy restrains minority MOMP in Meg-01 post IR. **a** Western blotting analysis and quantification of LC3 lipidation in Meg-01 cells at 1 dpi ( $n = 3$ ). **b** Autophagic clearance of mitochondria in Meg-01 cells at 1 dpi was assessed by immunofluorescence staining with antibodies specific for LC3 and TOM20. Scale bar = 20  $\mu$ m. **c** Flow cytometric quantification of MMP in Meg-01 cells with or without CQ treatment at 1 dpi ( $n = 5$ ). **d** Flow cytometric quantification of p-TBK1 and p-IRF3 expression in Meg-01 cells with or without CQ treatment at 1 dpi ( $n = 5$ ). **e** Flow cytometric quantification of IFN- $\beta$  and p-STAT1 expression in Meg-01 cells with or without CQ treatment at 1 dpi ( $n = 5$ ). **f** Flow cytometric quantification of GBP2 expression in Meg-01 cells with or without CQ treatment at 3 dpi ( $n = 5$ ). **g** Flow cytometric quantification of FSC-area and CD62P expression in response to thrombin of PLPs from Meg-01 cultures with or without CQ treatment at 3 dpi ( $n = 5$ ). Data represent mean  $\pm$  standard deviation. \* $P < 0.05$ , \*\* $P < 0.01$ , compared to Ctrl or IR. Two-tailed unpaired student's  $t$ -test. MOMP mitochondrial outer membrane permeabilization, Ctrl control, IR ionizing radiation, dpi day post IR, LC3 microtubule-associated protein 1 light chain 3, LC3-I non-lipidized LC3, LC3-II lipidized LC3, TOM20 translocase of outer mitochondrial membrane 20, MMP mitochondrial membrane potential, CQ chloroquine, TBK1 TANK-binding kinase 1, IRF3 interferon regulatory factor 3, IFN- $\beta$  interferon- $\beta$ , STAT1 signal transducer and activator of transcription 1, GBP2 guanylate-binding protein 2, FSC forward scatter, PLP platelet-like particle, TMRE tetramethylrhodamine ethyl ester, DAPI 4',6-diamidino-2-phenylindole, GAPDH glyceraldehyde-3-phosphate dehydrogenase