1 Supplemental Methods

Cell isolation and culture

- 3 Murine bone marrow-derived dendritic cells (BMDCs), murine T cells from the
- 4 spleen, and Lewis lung cancer (LLC), mouse melanoma B16-F10 and the
- 5 OVA-transfected clone derived from E.G7 (E.G7-OVA) cell lines were used for in
- 6 vitro studies. BMDCs and T cells were isolated and cultured in RPMI-1640 medium.
- 7 Briefly, the bone marrow cells from mice were pooled, filtered through a 70 μm nylon
- 8 mesh filter (Corning, USA), centrifuged and resuspended in ACK lysis buffer for 5
- 9 min of incubation on ice. After rinsing by medium, the bone marrow cells were
- 10 cultured in RPMI-1640 medium supplemented with 10% heat-inactivated FBS (Gibco,
- 11 USA), 100 U/mL penicillin/streptomycin (Gibco, USA) and additions of 1 mM
- sodium pyruvate, 50 μ M β -mercaptoethanol (β -ME), 20 ng/mL GM-CSF and 15
- 13 ng/mL IL-4 for BMDCs differentiation. The non-adherent and loosely adherent cells
- were collected as BMDCs. The spleen was isolated from mice, grinded and filtered
- throng a 70 μm nylon mesh filter for lymphocytes processing using a lymphocyte
- separation solution kit (Dakewe, China). The isolated lymphocytes were cultured in
- 17 RPMI-1640 medium supplemented with 10% heat-inactivated FBS, 100 U/mL
- 18 penicillin/streptomycin and additions of 1 mM pyruvate, 50 Mm β -ME and 100 U/mL
- 19 IL-2 for T cells differentiation. Murine LLC, B16-F10, and E.G7-OVA cells were
- 20 purchased from American Type Culture Collection (ATCC) and cultured in DMEM
- 21 except for E.G7-OVA cells in RPMI-1640 medium. G418 (0.4 mg/ml) was added to
- 22 the E.G7-OVA culture. All cells were maintained in a humidified atmosphere
- 23 containing 5% CO₂ at 37°C.

Tumor model

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- 25 To mimic tumor lung metastasis models, female or male Tfam-/- mice were
- intravenously injected with a total of 5×10^5 LLC cells or 2×10^5 B16-F10 cells.
- 27 Specially, for survival statistics of the lung metastasis B16-F10 model, mice were
- intravenously injected with a total of 5×10^5 B16-F10 cells. Tfam^{fl/fl} littermates and

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wild type (WT) mice were used as control. The survival rate was figured. After inoculation for 24 days (LLC models) or 14 days (B16-F10 models), the mice were 30 31 sacrificed and lung metastasis was evaluated. To investigate the immunological 32 responses, mice were subcutaneously injected with 10 µg OVA in PBS in a total volume of 100 μL (Tfam^{-/-} + OVA/Control + OVA group) or 100 μL PBS alone 33 (Tfam^{-/-} + PBS/Control + PBS group) antigen to the left flank on days of 0, 14, and 21. 34 Then subcutaneously injected with a total of 5×10^5 E.G7-OVA cells to the right flank 35 on day 28. Tumor size was monitored and measured using a vernier caliper every 36 37 third day. Mice were sacrificed between days of 16 and 19. For Survival statistics, 38 mice were humanely sacrificed when the volume of tumors reached about 2000 mm³. Further, we blocked the STING signaling in mice via C-176 (MCE, #HY-112906), 39 a strong and covalent STING inhibitor ¹. Mice were intravenously injected with a total 40 of 5×10^5 LLC cells, and then intraperitoneally administrated with 13.4 mg/kg of 41 C-176 dissolved in 10% DMSO and 90% saline containing 20% SBE-β-CD or equal 42 volume of vehicle (10% DMSO and 90% saline containing 20% SBE-β-CD) daily 43 during the course ². 44

Stimulation of BMDCs and T cells

- To study the induced responses by tumor cells to BMDCs from Tfam-/- mice, the 46
- 47 tumor supernatant (TS) of LLC cells was harvested, filtrated by 0.22 µm filters, and
- 48 added to the BMDCs at a final dilution of 1:3 for 24 h of incubation. To block the
- STING signal, 2 µM H-151 (MCE, #HY-112693) was added to the diluted TS. The 49
- 50 supernatant and cells were collected for further study. To remove nucleic acid in TS,
- 51 the filtrated TS was incubated with 10 U/mL SuperNuclease (Sino Biological, China)
- for 1-2 h at 37°C. 52
- To investigate cell-mediated immune responses, T cells from spleens of Tfam^{-/-} and 53
- WT mice immunized with 10 µg OVA in PBS were obtained, and CD8⁺ T cells were 54
- sorted using a negative magnetic bead screening kit (Stemcell Technologies, #19853) 55
- following the manufacturer's instruction. Cells were cultured with OVA₂₅₇₋₂₆₄ peptides 56

- 57 $(10 \ \mu\text{g/mL})$ for 24 h or 72 h. The supernatant and cells were collected for further
- 58 study.

Cell staining for flow cytometry (FCM)

- Mice were euthanized for lungs with tumor nodules and tumor tissues processing.

 Briefly, tissues were dissected from mice, minced on ice into small pieces of less than
- 62 1 mm³ and then suspended in a 10 mL of digestion buffer consisting of 1 mg/mL
- 63 collagenase I (Gibco), 0.5 mg/mL collagenase IV (Gibco) and 40 U/mL DNase I
- 64 (Sigma) in RPMI-1640 medium (Gibco, USA). The digestion buffer was incubated
- with frequent agitation at 37°C for 1 h. Subsequently, the suspensions were passed
- through a 70 μ m nylon mesh filter and centrifuged at 400×g for 5 min. All pelleted
- 67 cells were resuspended into ice-cold PBS and counted. Besides, mice were also
- 68 euthanized for inguinal lymph nodes and spleens processing. Cells were collected
- from inguinal lymph nodes and spleens by grinding and passing through a 70 μm
- 70 nylon mesh filter. For *in vitro* studies, cultured cells were collected at the indicated
- 71 end point.
- 72 Cells were pre-treated with FcR blocking (BD Biosciences, #553142, USA) to
- 73 avoid non-specific staining. Then the cells were incubated with live/dead staining dye
- 74 (Invitrogen, #L34974) for 30 min at 4 °C. For surface staining, cells were stained with
- 75 the indicated antibodies at 4 °C. For intracellular staining, cells pre-treated with
- 76 Brefeldin A for 4-6 h were then fixed using 4% paraformaldehyde (PFA) and
- 77 permeabilized by the 1% Triton X-100, and were incubated with
- 78 fluorochrome-labeled antibodies specific for the mouse. Fluorescence minus one
- 79 (FMO) control was used for gates strategy. Samples were performed using BD
- 80 LSRFortessa, and data were analyzed by FlowJo 10.7.1 or Novoexpress. Antibodies
- 81 used for FCM analysis included BV650 anti-CD45, PerCP-Cy5.5 anti-CD11b, BV711
- anti-CD11c, PE anti-F4/80, APC anti-MHCII, BV421 anti-Ly6C, BV510 anti-Ly6G,
- 83 FITC anti-CD206, PerCP-Cy5.5 anti-CD3, BV421 anti-CD4, BV510 anti-CD8, FITC
- 84 anti-PD-1, PE anti-IFN-γ, APC anti-GzmB, PE anti-CD69, FITC anti-Foxp3, FITC

- 85 anti-CD62L, APC anti-CD44, APC anti-11c, PE anti-CD25, APC anti-CD40,
- 86 PE-anti-CD86, FITC anti-CD80, PerCP-Cy5.5-anti-CD115, and BV421-anti-CD135
- 87 (all from BD Biosciences or BioLegend, USA), and APC H-2K^b MuLV p15E
- 88 Tetramer (MBL, Japan). For TFAM staining, permeabilized cells were stained by
- rabbit anti-TFAM antibody (Abcam, UK) followed by PE anti-rabbit IgG (BioLegend,
- 90 USA). For apoptosis analysis, cells were stained by FITC Annexin V Apoptosis
- 91 Detection Kit I (BD Biosciences, USA).

Histology, immunohistochemistry (IHC) and immunofluorescence

- 93 Lungs were fixed in 4% PFA at RT, embedded in paraffin and sectioned at 3 μm, or
- 94 frozen in OCT compound (Sakura Finetek, Japan) and sectioned at 10 µm.
- 95 Haematoxylin and eosin (H&E) staining was used to assess pathology and lung tumor
- 96 metastasis. The lung metastatic area was calculated using ImageJ software.
- Paraffin-embedded sections were incubated with 3% H₂O₂ to block endogenous
- 98 peroxidases, and then subjected to an EDTA buffer for antigen retrieval.
- 99 OCT-embedded sections were post-fixed with ice-cold acetone and incubated with 3%
- H₂O₂. Cell slides were fixed with 4% PFA. The samples were then incubated with 5%
- 101 goat serum for blocking. Primary antibodies used for immunohistochemistry or
- immunofluorescence analysis included rabbit anti-Hsp60 (Abcam, #ab137706),
- hamster anti-CD11c (BioLegend, #117301), rabbit anti-CD8α (CST, #98941), rabbit
- anti-CD45 (Servicebio, #GB11066), rabbit anti-CD3 (Servicebio, #GB13014), rabbit
- anti-CD31 (Abcam, #ab28364) and rabbit anti-cleaved caspase-3 (CST, #9664). The
- 106 immunofluorescent images were captured using a Zeiss LSM880 laser confocal
- 107 microscope, and immunohistochemistry and H&E images were captured using the
- 108 Pannoramic MIDI scanner (3DHISTECH, Hungary).

109 ELISA assay

- 110 To investigate the serum antibodies against the OVA, mice immunized with 10 μg
- 111 OVA on days of 0,14 and 21 and sera were collected by retro-orbital puncture on day
- 112 28. OVA was performed to coat flat-bottom 96-well plates (NUNC-MaxiSorp,

113 Thermo Fisher Scientific) at a final concentration of 10 µg/mL in 50 mM carbonate coating buffer (pH 9.6) at 4 °C overnight. Then blocking solution containing 5% BSA 114 115 in PBST was added for 1 h of incubation at RT. Diluted sera were added and incubated at 37 °C for 1 h. Antibodies, including anti-mouse IgG, IgG1, IgG2b, IgG2c 116 117 and IgG3 horseradish peroxidase (HRP)-conjugated antibody, were diluted 1:5,000 in blocking solution and added to wells (100 µl per well) for 1 h of incubation at RT. 118 119 After development using 3,3',5,5'-tetramethylbiphenyldiamine (TMB) and subsequent 120 stop of reactions by 50 µL/well of 1 M H₂SO₄ solution, the absorbance was measured 121 at 450 nm using a microplate reader (Biotek, USA). 122 Cytokines in vitro were measured following the manufacturer's instructions, 123 including IL-6 Mouse Uncoated ELISA Kit (Thermo Fisher Scientific, #88-7064-88), 124 TNF alpha Mouse Uncoated ELISA Kit (Thermo Fisher Scientific, #88-7324-88), 125 IL-1 beta Mouse Uncoated ELISA Kit (Thermo Fisher Scientific, #88-7013-88), IFN gamma 'Femto-HS' High Sensitivity Mouse Uncoated ELISA Kit (Thermo Fisher 126 127 Scientific, #88-8314-88), and Mouse IL-12 p40 ELISA Kit (Abcam, #ab236717). 128 Antigen uptake function detection of BMDCs 129 BMDCs were treated with 1mg/mL FITC-Dextran (Sigma, #60842-46-8) at 37°C for 130 1 h, rinsed with ice-cold PBS and then fixed with 4% PFA. The fluorescent images 131 were captured using a fluorescence microscopy. For FCM analysis, BMDCs treated 132 with FITC-Dextran at 4°C were performed as control of the spontaneous 133 FITC-dextran phagocytic. 134 T cell proliferation and cross-presentation assay To investigate the antigen cross-presentation, the CD8⁺ T cell proliferation assay was 135 performed ³. Briefly, BMDCs were isolated from *Tfam*^{-/-} and control mice on day 28 136 137 pre-immunized with OVA (10 µg) on days of 0, 14 and 21. After incubation for 7 days 138 for BMDCs maturation, 10 µg/mL of OVA₂₅₇₋₂₆₄ peptide was added to cells following by 24 h of incubation at 37 °C, and then BMDCs were collected and counted. 139

140 Simultaneously, CD8⁺ T cells were isolated from spleens of OT-I mice, sorted using a 141 negative magnetic bead screening kit (Stemcell Technologies, #19853), and then 142 labelled with CFSE (Invitrogen, #C34554) by incubation with CFSE (2.5 µM, 143 prepared with 0.1% FBS/PBS solution) for 15 min in the dark at 37°C. For co-culture system preparation, 2×10⁵ BMDCs and 1×10⁶ CD8⁺ T cells were mixed in 1 mL of 144 medium and seeded into a 24-well plate following by 72 h of incubation at 37 °C. 145 146 Terminally, the cells were collected for flow cytometric analysis of CFSE 147 fluorescence value of T cells. 148 T cell activation assay To detect the activation of OVA-specific cytotoxic T cells (CTLs), splenic 149 150 lymphocytes were isolated from mice on day 28 pre-immunized with OVA (10 µg) on 151 days of 0, 14 and 21. Subsequently, 10 µg/mL of OVA₂₅₇₋₂₆₄ peptide was added to 152 activate cells following by 72 h of incubation at 37 °C. Terminally, the cells were 153 collected and labeled with T-select MHC Tetramer /H-2Kb-OVA (SIINFEKL) (MBL, 154 #TS-5001-1C) and anti-CD8 antibody for flow cytometric analysis. 155 Adoptive immunity assay To investigate the adoptive immunity, control and *Tfam*^{-/-} mice were immunized with 156 10 μg of OVA on days of 0, 14 and 21, and then CD8⁺ T cells were isolated from the 157 immunized spleen on days of 28, 30 and 32 and sorted using a negative magnetic bead 158 screening kit (Stemcell Technologies, #19853). A total of 5×10^5 E.G7-OVA cells was 159 subcutaneously injected to the right flank of WT mice. Meanwhile, a total of 5×10^6 160 CD8⁺ T cells from control or *Tfam*^{-/-} mice was intravenously injected to the mice on 161 162 1-day ahead-inoculation of tumor cells, and on 1-day and 3-days post-inoculation of 163 tumor cells, respectively. All the tumour grew in the WT. 164 For adoption of DCs and macrophages, BMDCs and macrophages were differentiated from bone marrow cells with 100 ng/ml FLT-3L (R&D Systems, USA) or 20 ng/ml 165 M-CSF (novoprotein, China) respectively. Totally 1×10^6 Tfam^{-/-}/control BMDCs or 166

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macrophages were intravenously injected into WT LLC lung metastasis models on day 0, 1, 4, 7, and 14. Mice were sacrificed on day 24 to collect the lungs. The gross appearance of lungs were photographed, and the lung metastatic nodules were counted. Three representative lungs were selected from each group for formaldehyde fixation, paraffin embedding, and HE staining. The lung metastatic area was quantified by ImageJ software.

Quantitative PCR (qPCR)

- 174 Total RNA was extracted using the RNA Extraction Kit (TianGen, #DP419), and
- 175 first-strand cDNA was synthesized using a PrimeSriptTM RT reagent kit (Takara,
- 4RR036) following manufacturer's protocols. qPCR was performed in triplicate at a
- 177 20 μL total volume using SsoFast EvaGreen (Bio-Rad, #1725202) on a Bio-Rad
- 178 iCycler RT-PCR detection system. Mitochondrial DNA copy number was measured
- using primers specific to nuclear Tert and the D-loop region of mtDNA. The
- 180 expression levels of target genes were normalized to β -actin using $2^{-\Delta\Delta CT}$ method. All
- primers used in this study are listed in Supplementary Table 2.

Measurement of cytosolic mitochondrial DNA (mtDNA)

- 183 Cytosolic mtDNA was extracted from BMDCs using the mitochondrial DNA isolation
- 184 kit (Abcam, #ab65321), purified using the DNeasy Blood & Tissue Kit (Qiagen,
- #USA), and quantified by PCR using a TaqMan probe as previously described ⁴.

ROS measurement

- 187 The ROS level was detected using a DCF-DA (Sigma, #35845) probe. Briefly, cells
- seeded in 96-well plates were incubated with or without the tumor supernatant (TS)
- 189 for 24 h, and then treated with 10 μM DCF-DA (prepared with phenol red-free
- 190 medium) for 30 min. The fluorescence intensity was determined by a microplate
- reader (Biotek, USA) at the excitation/emission wavelengths of 485/530 nm. All
- 192 procedures were carried out in the dark.

Extracellular oxygen consumption measurement

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- 194 The mitochondrial metabolism was measured using the Cell Metabolism kit (Abcam),
- 195 and the extracellular oxygen consumption rates (OCR) were measured using the
- 196 Extracellular Oxygen Consumption Assay kit (Abcam). All procedures were
- 197 performed according to the manufacturer's protocols, and the fluorescence intensity
- was determined by a microplate reader (Biotek, USA).

Characterization of mitochondrial structure

- 200 Transmission electron microscopy was used to characterize the mitochondrial
- 201 structure. Briefly, murine BMDCs were fixed with 2.5% glutaraldehyde, dehydrated
- and then embedded in resin. The ultrathin section (~50 nm) was obtained and
- 203 performed on a copper grid by staining with uranyl acetate and lead citrate. The
- images were captured using a JEM-2100PLUS electron microscope (JEOL, Japan).

Western blot

- 206 Total protein was extracted using the ice-cold enhanced RIPA lysis buffer
- 207 supplemented with a phosphatase inhibitor cocktail (Millipore, #524629) and
- 208 measured using the Bradford dye (BIO-RAD, #5000205) by Eppendorf
- 209 Bio-photometer Plus (Eppendorf). Western blotting was performed using standard
- 210 protocols according to the manufacturer's recommendations. Primary antibodies used
- 211 for western blot analysis included rabbit, rabbit anti-Stat1 (CST, #9172), rabbit
- 212 anti-STING (CST, #13647), rabbit anti- Phospho-TBK1 (CST, #5483), rabbit
- anti-TBK1 (CST, #3504), rabbit anti-IRF-3 (CST, #4302), rabbit anti-Phospho-IRF-3
- 214 (CST, #79945) and mouse anti- β -actin (Santa Cruz, #sc-47778). The β -actin protein
- 215 was used as the loading control.

RNA sequencing

- 217 Total RNA was isolated from BMDCs (treated with or without TS) of Tfam^{-/-} and
- 218 control mice using the TRIzol reagent kit. RNA integrity was assessed using the RNA
- Nano 6000 Assay Kit by the Bioanalyzer 2100 system (Agilent Technologies, CA,
- USA), and samples with RIN scores above 6 and a minimum total RNA of 100 ng

- 221 were used for further library construction. PCR products were purified using the
- 222 AMPure XP system, and library quality was assessed using the Agilent Bioanalyzer
- 223 2100 system. Libraries were sequenced on Illumina Novaseq6000 (Illumina)
- 224 following PE150 sequencing strategy. Differential expression analysis was performed
- using the DESeq2 R package (1.20.0), and mRNAs with a FDR <0.001 and
- 226 fold-change >2 were considered as significant differential expression between two
- groups. GSEA (gene set enrichment analysis) were used for functional annotation of
- the differentially expressed genes. The experiment was conducted in triplicate.

230 References

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- 231 1. Haag SM, Gulen MF, Reymond L, et al. Targeting STING with covalent
- 232 small-molecule inhibitors. Nature. 2018;559(7713):269-273. doi:
- 233 10.1038/s41586-018-0287-8.
- 234 2. Wei B, Xu L, Guo W, et al. SHP2-Mediated Inhibition of DNA Repair Contributes
- 235 to cGAS-STING Activation and Chemotherapeutic Sensitivity in Colon Cancer.
- 236 Cancer Res. 2021;81(12):3215-3228. doi: 10.1158/0008-5472.CAN-20-3738.
- 237 3. Moroishi T, Hayashi T, Pan WW, et al. The Hippo Pathway Kinases LATS1/2
- 238 Suppress Cancer Immunity. Cell. 2016;167(6):1525-1539 e17. doi:
- 239 10.1016/j.cell.2016.11.005.
- 240 4. Wei X, Shao B, He Z, et al. Cationic nanocarriers induce cell necrosis through
- impairment of Na(+)/K(+)-ATPase and cause subsequent inflammatory response. Cell
- 242 Res. 2015;25(2):237-53. doi: 10.1038/cr.2015.9.

244 .Table.1 Primer used in PCR for genotyping

Gene name	Forward sequences	Reaction
	Reverse sequences	
m <i>Tfam</i>	CTT GTA GGT CCT CCC CAC TG	A
	ACA GCA CCA ACC CAA AGT GT	A
mLyz2-cre	CCC AGA AAT GCC AGA TTA CG	Mutant (A)
	CTT GGG CTG CCA GAA TTT CTC	Common (A, B)
	TTA CAG TCG GCC AGG CTG AC	Wild type (B)

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246 Table.2 Primer used in qPCR

	Forward sequences	
Gene name	Reverse sequences	
	CTGAGAGGGAAATCGTGCGT	
β-actin	CCACAGGATTCCATACCCAAGA	
and Tife and	AAGGATGATTCGGCTCAGG	
m <i>Tfam</i>	GGCTTTGAGACCTAACTGG	
	CTTTCCTCATGATCCTGGTAATGAT	
mIfna4	AATCCAAAATCCTTCCTGTCCTTC	
ma I.C. l. 1	CCCTATGGAGATGACGGAGA	
mIfnb1	CCCAGTGCTGGAGAAATTGT	
L.67	CAATTCAGGGGATCCAGTTG	
m <i>Irf</i> 7	AGCATTGCTGAGGCTCACTT	
mCxcl10	CCAAGTGCTGCCGTCATTTTC	
mcxci10	GGCTCGCAGGGATGATTTCAA	
ma Luf 2	CGGAAAGAAGTGTTGCGGTTAGC	
mIrf3	CAGGCTGCTTTTGCCATTGGTG	
mDdx58	GAGTACCACTTAAAGCCAGAG	
IIIDaxo	AATCCATTTCTTCAGAGCATCC	

mTnfa	CATCTTCTCAAAATTCGAGTGACAA	
mTnfa	CCAGCTGCTCCTCCACTTG	
m////b	TGGACCTTCCAGGATGAGGACA	
	GTTCATCTCGGAGCCTGTAGTG	
m///2	CCATTGAACTGGCGTTGGAAG	
1111112	ACTTGAGGGAGAAGTAGGAATGG	
m mtDNA Dlaam 2	TCCTCCGTGAAACCAACAA	
m.mtDNA Dloop 3	AGCGAGAAGAGGGGCATT	
m.nucDNA Tert	CTAGCTCATGTGTCAAGACCCTCTT	
III.IIUCDNA Tert	GCCAGCACGTTTCTCTCGTT	