

SUPPLEMENTAL MATERIAL

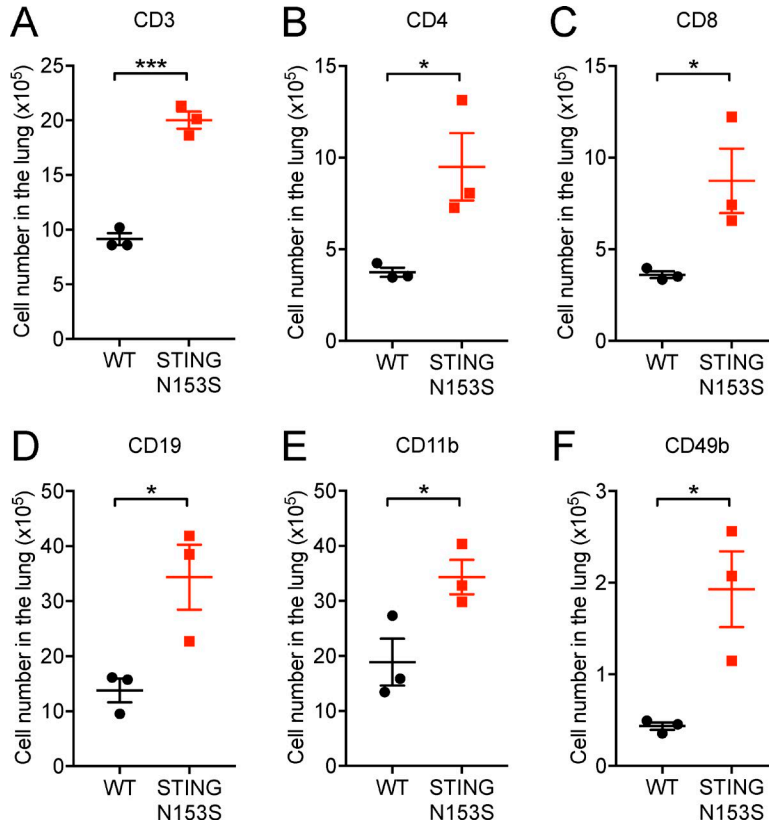
Warner et al., <https://doi.org/10.1084/jem.20171351>

Figure S1. **Pulmonary immune cell analysis of WT and STING N153S mice.** (A–F) Flow cytometry analysis of immune cells isolated from the lungs of WT and STING N153S mice. Total numbers were determined for the following immune cell subsets: CD3⁺ (A), CD4⁺ (B), CD8⁺ (C), CD19⁺ (D), CD11b⁺ (E), and CD49b⁺ populations (F). Data represent the mean \pm SEM of $n = 3$ mice per group. *, $P < 0.05$, ***, $P < 0.0005$ by unpaired t test.

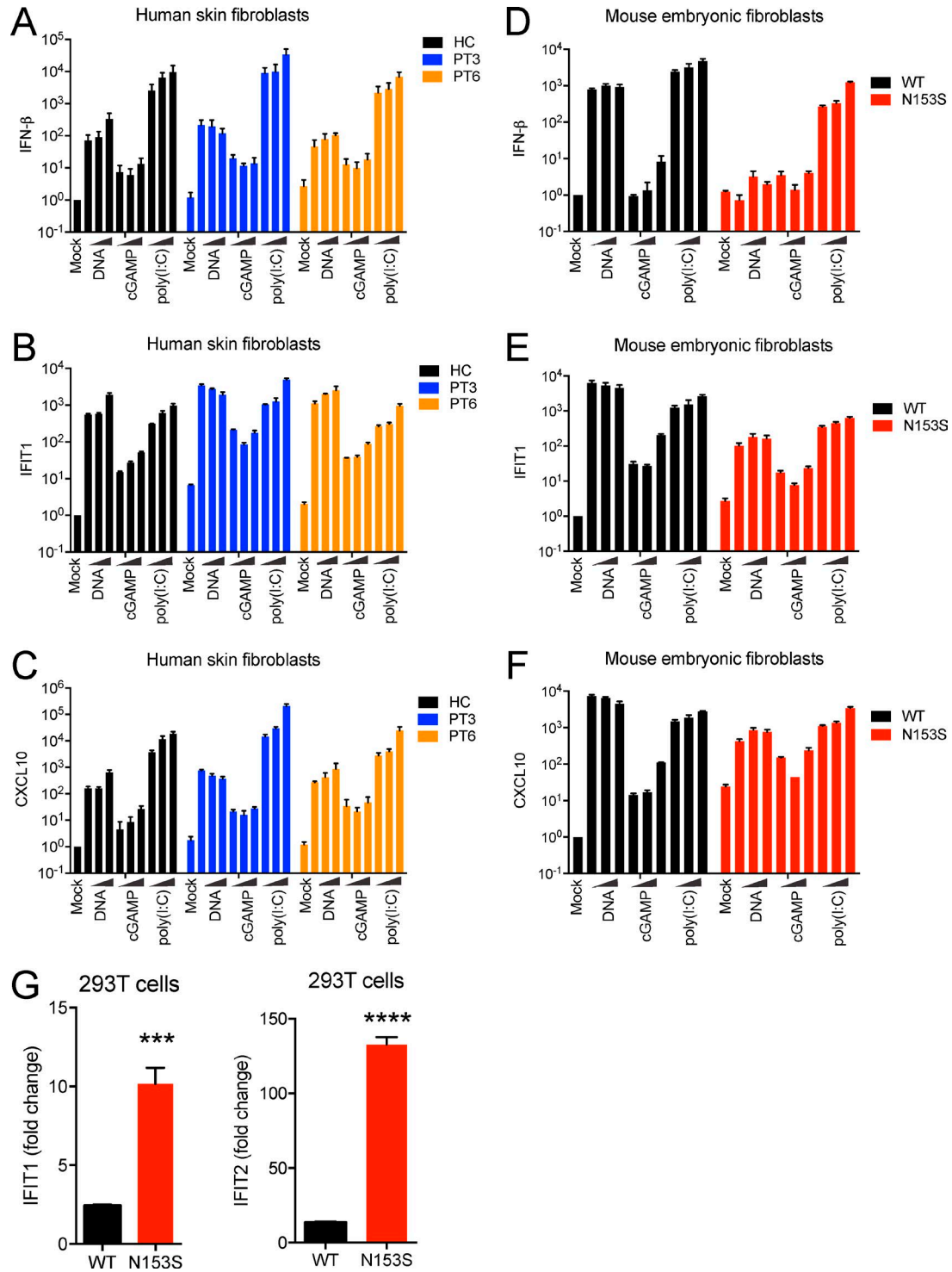


Figure S2. **IFN- β and ISG expression in primary STING N154S patient skin fibroblasts, STING N153S MEFs, and transfected 293T cells.** (A–F) Cells were stimulated with increasing amounts of 2'3'-cGAMP (2, 4, and 8 μ g), herring testes DNA (htDNA; 0.1, 0.5, and 1 μ g) and polyinosinic-polycytidylic acid (poly I:C; 0.1, 0.5, and 1 μ g) for 6 h. There was no significant difference cGAMP-transfected WT and STING N153S MEFs or between healthy control and STING N154S skin fibroblasts. Data were analyzed by ANOVA. (G) ISG expression in 293T cells 24 h after overexpression of WT mouse or mutant STING N153S as previously described (Dobbs et al., 2015). ***, $P < 0.0005$; ****, $P < 0.0001$ by t test. Stimulations were performed in biological triplicates, and qRT-PCR was performed in technical triplicates. Data represent the mean \pm SEM.

Table S1. Antibodies used in CyTOF

Tag	Target	Clone	Manufacturer	Catalog #
089Y	CD45	30-F11	Fluidigm	3089005B
141Pr	pSHP2	D66F10	Fluidigm	3141002A
142Nd	Caspase 3 (Cleaved)	D3E9	Fluidigm	3142004A
143Nd	TCRβ	H57-597	Fluidigm	3143010B
145Nd	CD4	RM4-5	Fluidigm	3145002B
146Nd	CD8α	53-6.7	Fluidigm	3146003B
147Sm	pSTAT5	47	Fluidigm	3147012A
148Nd	CD11B (Mac-1)	M1/70	Fluidigm	3148003B
149Sm	p4E-BP1	236B4	Fluidigm	3149005A
150Nd	CD25	3C7	Fluidigm	3150002B
151Eu	Ly-6G	1A8	Fluidigm	3151010B
153Eu	pSTAT1	58D6	Fluidigm	3153003A
154Sm	TER119	TER-119	Fluidigm	3154005B
155Gd	pTBK1/NAK	D52C2	CST	5483BF
158Gd	pSTAT3	4/P-Stat3	Fluidigm	3158005A
159Tb	TCRγδ	GL3	Fluidigm	3159012B
160Gd	CD45R (B220)	RA3-6B2	Fluidigm	3160012B
161Dy	pBAD	40A9	Fluidigm	3161006A
164Dy	IκBα	L35A5	Fluidigm	3164004A
165Ho	CD3e	145-2C11	Fluidigm	3165020B
166Er	CD19	6D5	Fluidigm	3166015B
168Er	Ki67	B56	Fluidigm	3168007B
170Er	NK1.1	PK136	Fluidigm	3170002B
171Yb	CD44	IM7	Fluidigm	3171003B
172Yb	pS6	N7-548	Fluidigm	3172008A
173Yb	CD117 (c-kit)	2B8	Fluidigm	3173004B
174Yb	MHCII	M5/114.15.2	Fluidigm	3174003B
176Yb	pCREB	87G3	Fluidigm	3176005A

Table S2. Definition of populations used in CyTOF analysis

Cell populations	Marker expression
Fig. 5	
CD4 (CD4 T cells)	CD45 ⁺ TER119 ⁻ CD3 ⁺ CD4 ⁺ TCRβ ⁺
CD8 (CD8 T cells)	CD45 ⁺ TER119 ⁻ CD3 ⁺ CD8 ⁺ TCRβ ⁺
NK (NK cells)	CD45 ⁺ TER119 ⁻ NK1.1 ⁺ CD3 ⁻
CD19 (B cells)	CD45 ⁺ TER119 ⁻ CD19 ⁺ B220 ⁺ MHCII ^{int/lo}
Ly6G ⁺ (CD11b ^{hi} Ly6G ^{hi})	CD45 ^{+/lo} TER119 ⁻ CD11b ^{hi} Ly6G ^{hi}
Monocytes	CD45 ⁺ TER119 ⁻ CD11b ⁺ Ly6G ⁻ MHCII ⁻
Immature myeloid	CD45 ^{lo} TER119 ⁻ CD117 ^{+/lo} CD11b ⁺ Ly6G ^{int/lo}
CD3 double-negative (DN)	CD45 ⁺ TER119 ⁻ CD3 ⁺ CD4 ⁻ CD8 ⁻
Fig. 6	
CD4 ⁺ CD8 ⁻	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁺ CD8 ⁻
CD8 ⁺ CD4 ⁻	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD8 ⁺ CD4 ⁻
CD4 ⁺ CD8 ⁺	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁺ CD8 ⁺
CD4 ⁻ CD8 ⁻	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁻ CD8 ⁻
CD19 ⁺ B220 ⁺	CD45 ⁺ TER119 ⁻ B220 ⁺ CD19 ⁺
NK1.1 ⁺	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁺
DN1	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁺ CD8 ⁺ CD44 ⁺ CD25 ⁻
DN2	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁺ CD8 ⁺ CD44 ⁺ CD25 ⁺
DN3	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁺ CD8 ⁺ CD25 ⁺ CD44 ^{lo/-}
DN4	CD45 ⁺ TER119 ⁻ B220 ⁻ CD19 ⁻ NK1.1 ⁻ CD4 ⁺ CD8 ⁺ CD44 ^{lo/-} CD25 ⁻
Fig. 7	
B cells	CD45 ⁺ TER119 ⁻ CD3 ⁻ CD19 ⁺
T cells	CD45 ⁺ TER119 ⁻ CD19 ⁻ CD3 ⁺
Monocytes	CD45 ⁺ TER119 ⁻ CD3 ⁻ CD19 ⁻ CD11b ⁺ Ly6G ⁻ MHCII ⁻
NK cells	CD45 ⁺ TER119 ⁻ CD19 ⁻ CD3 ⁻ NK1.1 ⁺
Immature myeloid	CD45 ⁺ TER119 ⁻ CD3 ⁻ CD19 ⁻ CD11b ⁺ MHCII ⁻

REFERENCE

Dobbs, N., N. Burnaevskiy, D. Chen, V.K. Gonugunta, N.M. Alto, and N. Yan. 2015. STING activation by translocation from the ER is associated with infection and autoinflammatory disease. *Cell Host Microbe*. 18:157–168. <http://dx.doi.org/10.1016/j.chom.2015.07.001>