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Supplementary Materials for

Viral cGAMP nuclease reveals the essential role of DNA sensing in protection against acute lethal virus infection

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Figs. S1 to S6 Tables S1 and S2 **Supplementary Materials**

Activation of DNA sensing is essential for protection against acute lethal virus infection.



Fig. S1. vSlfn does not inhibit RNA virus-induced IFN β or NF- κ B reporter activation. (A-C) HEK293T cells expressing firefly luciferase under the control of the IFNβ (A) or NF-κB (B-C) promoters were transfected with vectors encoding vSlfn or VACV virus proteins C6 and A49, or the corresponding empty vector as controls. C6 and A49 are known inhibitors of IRF3 and NF-kB signaling included here as controls. (A) 24 h post-transfection, cells were infected or not with Sendai virus 1 and luciferase activity measured at 24 hpi. (B) Similarly, cells were stimulated or not with interleukin (IL)-1 β for 8 h before luciferase activity determination. (C) Cells were transfected as above and simultaneously transfected with TNF receptor associated factor 6 (TRAF6). After 16 h, luciferase activity was determined. In all cases, luciferase activity was measured, normalized and presented as a fold increase over unstimulated EV-transfected cells. Data are presented as mean \pm SD and represent one of at least three experiments performed in triplicate. ***, p < 0.001 (unpaired Student's *t* test comparing with empty vector). (D) Expression of FLAG-tagged versions of vSlfn. HEK293T cells were independently transfected with expression vectors encoding full-length vSlfn, vSlfn lacking p26 domain (vSlfn∆p26) or p26 domain, fused with a C-terminal 3xFLAG epitope. Whole cell lysates were analyzed 24 h after transfection by SDS-PAGE and immunoblotting using an antibody against FLAG.



Fig. S2. Absence of vSlfn expression in ECTV Δ vSlfn infected cells. (A) Analysis of vSlfn expression by western blot in BSC-1 cells after infection with 5 pfu/cell of ECTV-WT and ECTV Δ vSlfn at the indicated times post-infection. (B) Analysis of vSlfn and vSlfn Δ p26 expression from recombinant ECTVs expressing C-terminally Strep-Tag tagged versions of vSlfn Δ p26 and full length vSlfn (ECTV-vSlfn Δ p26StrepTag and ECTV-vSlfnStrepTag, respectively). BSC-1 cells were infected as above and lysates analyzed by immunoblotting using an anti-StrepTag antibody at 24 hpi, confirming expression of vSlfn Δ p26. ECTV protein CrmD was used as control of infection and β -actin as protein loading control. One representative experiment from two performed is shown. Representation of the genomic organization of vSlfn locus and surrounding loci in the recombinant viruses used is shown.



Fig. S3. Absence of vSlfn in ECTV Δ vSlfn does not affect viral replication. A) To analyze ECTV replication diverse types of cells were infected with the indicated viruses. Results showed that all recombinant viruses had similar kinetics of replication *in vitro* to ECTV-WT. B) Similarly, ECTV replication in the presence or absence of vSlfn was analyzed in THP-1 cells and THP-1 cells lacking STING or cGAS expression. The multiplicity of infection of the viral inoculum is indicated in parenthesis. BMDM, Bone marrow derived macrophages.



Fig. S4. vSlfn prevents STING activation in response to cGAMP during ECTV infection. PMA-differentiated THP-1 cells were infected with 2 pfu/cell of the indicated viruses for 6 h before stimulation with cGAMP. Cell lysates were harvested at indicated times after cGAMP addition and analyzed by SDS-PAGE and immunoblotting with antibodies against the indicated proteins. The levels of cGAMP-induced STING dimerization as well as TBK1 and IRF3 phosphorylation were reduced during ECTV-WT infection, but remained similar to those in mock-infected cells during ECTVΔvSlfn and ECTV-vSlfnΔp26 infection. D8 is an unrelated ECTV protein used as infection marker.



Fig. S5. Pathway enrichment analysis from DPLNs at 5 dpi. Similar to Fig. 4., DPLNs from ECTV-WT and ECTV Δv Slfn infected animals were collected at 5 dpi for RNA-sequencing and further differential expression analysis to compare wild type with ECTV Δv Slfn infection. Analysis of the corresponding differentially expressed genes revealed the enrichment of diverse pathways related to innate immunity (highlighted). Z-score values for activated (orange) or repressed (blue) predicted pathways are indicated.



Fig. S6. Gating strategy for data in Fig. 6 and a representative example of each experimental condition are shown. The numbers of events positively stained in corresponding fluorescence minus one control samples were 0.069 % and 0.001 % positives in the case of CD49b and granzyme b, respectively.

Dose (pfu/animal)								
	WT	∆vSlfn		vSlfn∆p26	mSlfn1		mSlfn2	
1	2/5	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
10	0/5	5/5	n.d.	n.d.	n.d.	n.d.	n.d.	n.d.
102	n.d.	3/5	n.d.	5/5	n.d.	n.d.	n.d.	n.d.
103	n.d.	5/5	5/5	5/5	5/5	4/5	5/5	5/5
104	n.d.	n.d.	5/5	5/5	5/5	n.d.	5/5	n.d.
105	n.d.	n.d.	5/5	5/5	5/5	n.d.	5/5	n.d.
106	n.d.	n.d.	4/5	4/5	5/5	n.d.	4/5	n.d.

Table S1. Mortality rates of vSlfn recombinant ECTVs. Groups of 5 BALB/c mice were s.c. infected in the footpad with increasing doses of the indicated viruses and the number of survivors at 21 dpi determined for each condition are shown. Each column corresponds to an independent experiment. n.d., not determined.

Recombinant virus	Parental virus	Vector	5´flanking region oligos (5´-3´)	3´flanking region oligos (5´-3´)	Insert between Flanking Regions-		
ECTV∆vSlfn	ECTV p33 Naval strain		GGAATTCATCAGATG GATAAGAATA GGGATCCGTTTTCGG TTCCACTCCC	GGGATCCGTTGTGCTGTG TTTGCAG GGCTGCAGCTAGATAAAA			
ECTV- vSlfn∆p26	ECTV Naval strain	pBH4		GCGCGGATCCATGAAAGG TGCGGTGCTACA GGCTGCAGCTAGATAAAA AATCAACT			
ECTV-mSlfn1	ECTV Naval strain	p35	GGAATTCATCAGATG GATAAGAATA GGGATCCGTTTTCGG TTGCACTCGC	GGGATCCGTTGTGCTGTG TTTGCAG GGCTGCAGCTAGATAAAA AATCAACT	CGAGCTCATAATGAAC ATCACCGAT CGAGCTCCTAAGACAT GAGGAGCTT		
ECTV-mSlfn2	ECTV Naval strain	pBH2	GGAATTCATCAGATG GATAAGAATA GGGATCCGTTTTCGG TTGCACTCGC	GGGATCCGTTGTGCTGTG TTTGCAG GGCTGCAGCTAGATAAAA AATCAACT	CGGATCCATGGGTACT AGACTTGAGGC CGGATCCTCAACCTGA TGGGGCATTCATC		
ECTV- vSlfn∆p26Stre pTag	ECTV- vSlfn∆p26	pBH5	GCGGAATTCCGGCGG TGGCACAATCGAGG GCGGGATCCTCACTT CTCGAATTGAGGGTG	GCGGGATCCTTAATGTAA CTATAGAG GCGCTGCAGTGGATGATA ACCCGAGATTA			
ECTV- vSlfnStrepTag	ECTV Naval strain	pBH5	GCGGAATTCCGGCGG TGGCACAATCGAGG GCGGGATCCTCACTT CTCGAATTGAGGGTG	GCGGGATCCTTAATGTAA CTATAGAG GCGCTGCAGTGGATGATA ACCCGAGATTA	CGGATCCATGGGTACT AGACTTGAGGC CGGATCCTCAACCTGA TGGGGCATTCATC		
Expression Vector	Description			Oligos for InFusion cloning			
pvSlfn-FLAG (pBH21)	Expresses F expression i	LAG-taggeon n mammalia	d vSlfn optimized for in cells	n/a			
pvSlfn∆p26- FLAG (pBH32)	Expresses F optimized fo	LAG-tagged	d vSlfn lacking p26 n in mammalian cells	GCGGCCGCCACCATGAAAGGCGCCGTGCTGCAGA TCCCCAA CTGCAGCACGGCGCCTTTCATGGTGGCGCGGCGC			
pp26-FLAG (pBH33)	Expresses F optimized fo	LAG-tagged r expression	d vSlfn p26 domain n in mammalian cells	CTGCAGATCCCCAACGTGTCCTACATCAAAGTGATCG ATGAC TTTGATGTAGGACACGTTGGGGGATCTGCAGCACGGCG CCTTT			

Table S2. Vectors generated in the present work. Oligonucleotides used to generate therecombination vector together with the parental virus used in the construction of eachrecombinant mutant virus are detailed. Every recombination vector is based on previous pMS30.Below, mammalian expression vectors generated in this study and used in the reporter geneassays are shown.