862	Supplementary Data
863	Self assembling short immunostimulatory duplex RNAs with broad spectrum antiviral
864	activity
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891 Figure S1. Profiling the effects of RNA-2 by RNA-seq and TMT mass spectrometry. A549

cells were transfected with RNA-2 or scrambled RNA control, cell lysates were collected at 48 h,

and analyzed by RNA-seq (left) or TMT Mass Spec (right). Differentially expressed genes

(DEGs) or proteins are shown in volcano plots (top) and GO Enrichment analysis was

performed for the DEGs (bottom) (N = 3). Plot (top) and GO Enrichment analysis was performed

for the differentially expressed proteins (bottom) (N = 3).



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899 Figure S2. Heat maps showing the effects of immunostimulatory RNAs on IFN pathway-

900 relevant gene levels. DEGs from RNA-seq (A) and differentially expressed proteins from TMT

Mass Spec analyses (B) shown in Fig. 1B and fig. S1 are presented here as heat maps (gene

902 levels of the scrambled RNA control were set as 1; N = 3).





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Figure S3. RNA-induced gene expression associated with type I interferon pathway. (A) Venn diagram showing differentially expressed ISGs from TMT Mass Spec by RNA-1 belong to type I or type II interferon stimulated genes. (B) Heat map of qPCR results showing RNA-I preferentially activates type I interferon pathway. A549 cells were transfected with RNA-1 or scrambled dsRNA control, collected at 48 hr and analyzed by qPCR (expression levels were normalized to GAPDH; gene levels induced by the RNA control were set as 1; N = 3).



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- 914 **Figure S4. The levels of IFN-β protein induced by RNA-1.** A549 cells were transfected with
- 915 RNA-1 (34 nM) for 48 h, and then supernatants were collected for detection of IFN-β using
- 916 ELISA. Scrambled RNA control NC-1 is used as negative (N = 3).

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Figure S5. IRF3 knockout abolished the ability of immunostimulatory RNAs to induce
IFN-I pathway associated genes. Wild-type (WT) HAP1 cells, IRF3 knockout HAP1 cells, or
IRF7 knockout HAP1 cells were transfected with RNA-1 or a scrambled RNA control and
STAT1, IL4L1, TRAIL, and IFI6 mRNA levels were quantified by qPCR at 48 h post transfection.
Data are presented as fold change relative to RNA control (N = 3).



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926 Figure S6. RIG-I knockout abolished the induction effects of the immunostimulatory

927 **RNAs on IFN-β.** Wild-type (WT) A549-Dual cells, RIG-I knockout A549-Dual cells, MDA5

928 knockout A549-Dual cells, or TLR3 knockout A549 cells were transfected with RNA-1, RNA-2,

929 or a scramble RNA control and IFN-β mRNA levels were detected by Quanti-Luc assay in WT,

- 930 RIG-I KO, and MDA5 KO A549-Dual cells or qPCR in TLR3 KO A549 cells at 48 h post
- 931 transfection. Data are shown as fold change relative to the scrambled RNA control (N = 6).

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935 Figure S7. TLR7/8 knockout or overexpression did not have effect on the

immunostimulatory activity of RNA-1. (A) Graph showing that the overexpression of TLR7 in
HEK cells had no effect on production of IFN-β induced by RNA-1. (B) Graph showing that the
knockout of TLR8 in THP1 cells had no effect on IFN production induced by RNA1. These cell
lines are commercial and could be purchased from InvivoGen.





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943 Figure S8. 'CCC' motif is widely distributed in human genome. (A) Graph showing the

distribution of the number of CCC sequences in human mRNAs (retrieved from UCSC hg38

refGene with prefix NM). (B) Graph showing the distribution of the number of CCC sequences in

human lncRNAs (retrieved from Incipedia). (C) Table showing the percentage of human mRNAs

and IncRNAs containing the CCC motif and their average density.





A549 cells. IFN- β and ISG15 levels were detected in cells transfected with RNA-1, RNA-2, or

951 scramble dsRNA control by qPCR at 48 h post-transfection. The IFN-β or ISG15 level induced

- by the scramble dsRNA control was set as 1. Data are shown as fold change relative to the
- 953 control (N = 3).
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960 Table S1. Summary of characteristics of reported immunostimulatory RNAs.

Characteristic	Signaling pathway	Cytokines
5'-UGUGU-3' motif	Toll-like receptor (TLR)8	IFN-alfa
5'-GUCCUUCAA-3' motif	TLR7/8	IFN-alfa
GU or AU rich	TLR7/8	IFN-alfa, TNF-alfa
Uracil repeats	TLR7	IFN-alfa, IL-6, TNF-alfa
Blunt ended dsRNA	RIG-I	Type I IFN, p56
5'-triphosphate; 5'-diphosphate	RIG-I	IFN-alfa, IFN-beta
MicroRNA-like siRNA	TLR7/8	IFN-alfa, TNF-alfa
Long dsRNA	MDA5	Type I IFN
Long dsRNA	TLR3	Type I IFN
Single stranded (ss)RNA	TLR7	Type I IFN

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