

## Supplementary Materials for

### ***Drosophila* Dicer-2 has an RNA interference–independent function that modulates Toll immune signaling**

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Fig. S1. Overexpression of Dcr-2 enhances Toll signaling activation in *Drosophila* S2 cells.

Fig. S2. *Toll* 3'UTR is important for Toll signaling.

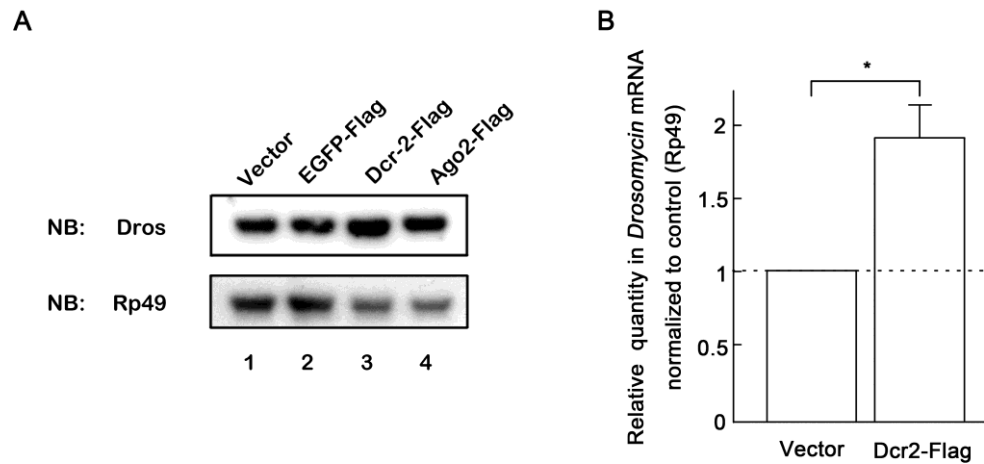
Fig. S3. The PAZ domain of Dcr-2 has no interaction with the 3'UTR of DIF or Dorsal in vitro.

Fig. S4. Toll signaling can be induced by FHV or VSV.

Fig. S5. The prediction of the secondary structure of *Toll* 3'UTR.

Table S1. Primers used in this work.

Figure S1

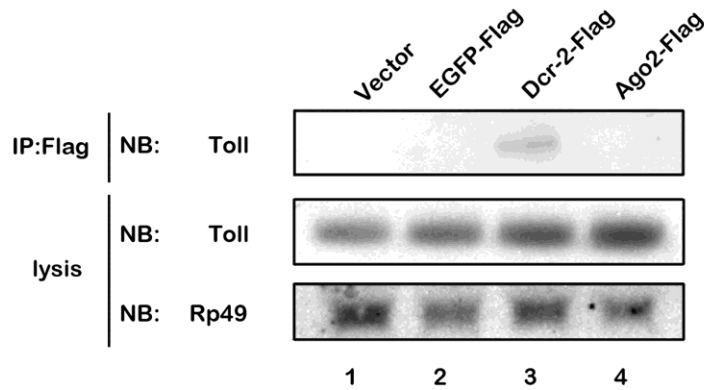


**Fig. S1. Overexpression of Dcr-2 enhanced Toll signaling activation in *Drosophila* S2 cells.**

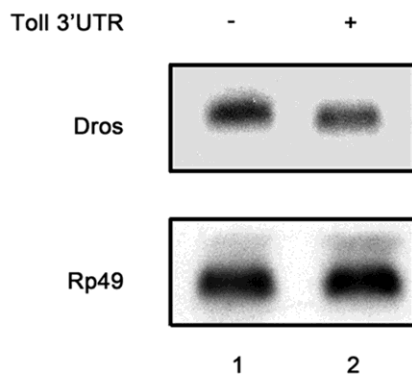
(A) Cultured S2 cells were transfected with empty vector (lane 1) or the plasmid expressing EGFP-Flag (lane 2), Dcr-2-Flag (lane 3), or AGO2-Flag (lane 4) as indicated. Total RNA extracts were prepared, followed by Northern blots using the indicated probes. (B) Cultured S2 cells were transfected with empty vector or the plasmid expressing Dcr-2-Flag as indicated. After that, total RNA extracts were prepared for qRT-PCR assay of *Dros* mRNA (normalized to Rp49;  $n=3$ ; \*,  $P < 0.05$  by two-tailed Student's t test; error bars, s.d.).

Figure S2

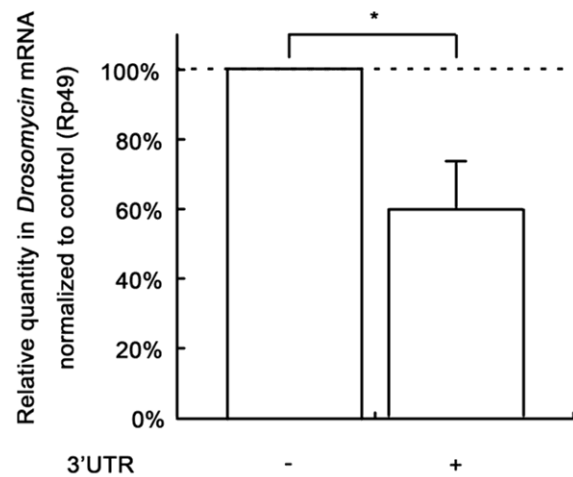
A



B



C



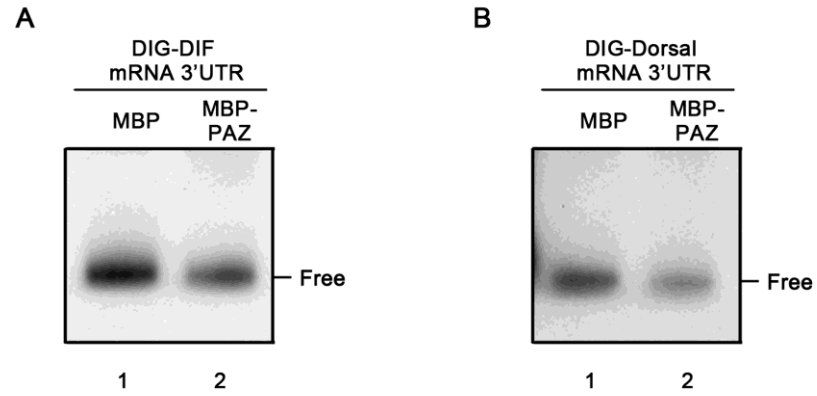
D

		Toll	dMyD88	DIF	Dorsal	PGRP-LC	IMD	Relish
Dcr-2-Flag	input	++	++	++	+++	++	+++	++++
	IP:Flag	+	-	-	-	-	-	-
EGFP-Flag	input	++	++	++	+++	++	+++	++++
	IP:Flag	-	-	-	-	-	-	-

**Fig. S2. *Toll* 3'UTR is important for Toll signaling.** (A) Cultured S2 cells were transfected with empty vector (lane 1), or the plasmid expressing EGFP-Flag (lane 2), Dcr-2-Flag (lane 3), or AGO2-Flag (lane 4) as indicated. The crude cell lysates were prepared and subjected to RNA-IP using anti-Flag antibody. The RNA extracts prepared from the precipitates and total cell

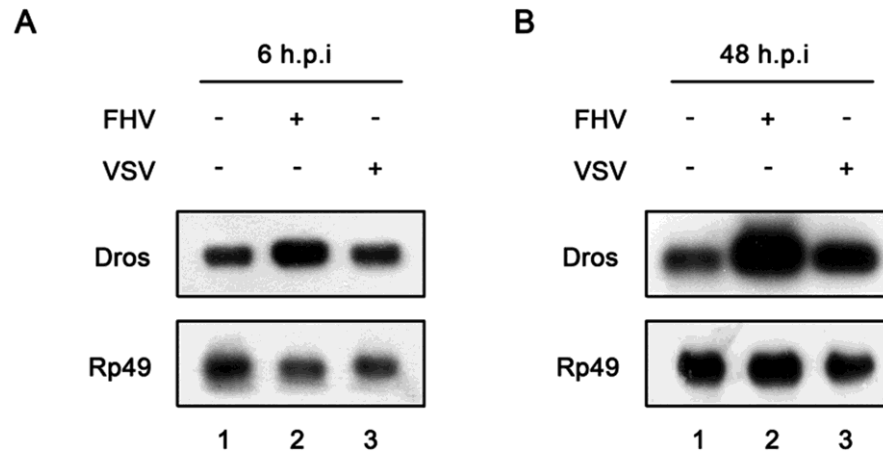
lysates were subjected to Northern blots as indicated. (B-C) Cultured S2 cells were transfected with *in vitro* transcribed *Toll* 3'-UTR as indicated. Total RNA extracts were prepared for Northern blots using the indicated probes (B) or for qRT-PCR assay of *Dros* mRNA (normalized to Rp49;  $n=3$ ; \*,  $P < 0.05$  by two-tailed Student's t test; error bars, s.d.) (C). (D) Cultured S2 cells were transfected with the plasmid expressing EGFP-Flag or Dcr-2-Flag. The crude cell lysates were prepared and subjected to RNA-IP using anti-Flag antibody. The RNA extracts prepared from the precipitates were subjected to qRT-PCR using the indicated pairs of primers. -, undetectable or Cq (quantification cycle) value  $\geq 37$ ; +,  $37 > \text{Cq value} \geq 30$ ; ++,  $30 > \text{Cq value} \geq 25$ ; +++,  $25 > \text{Cq value} \geq 20$ ; +++++, Cq value  $< 20$ .

Figure S3



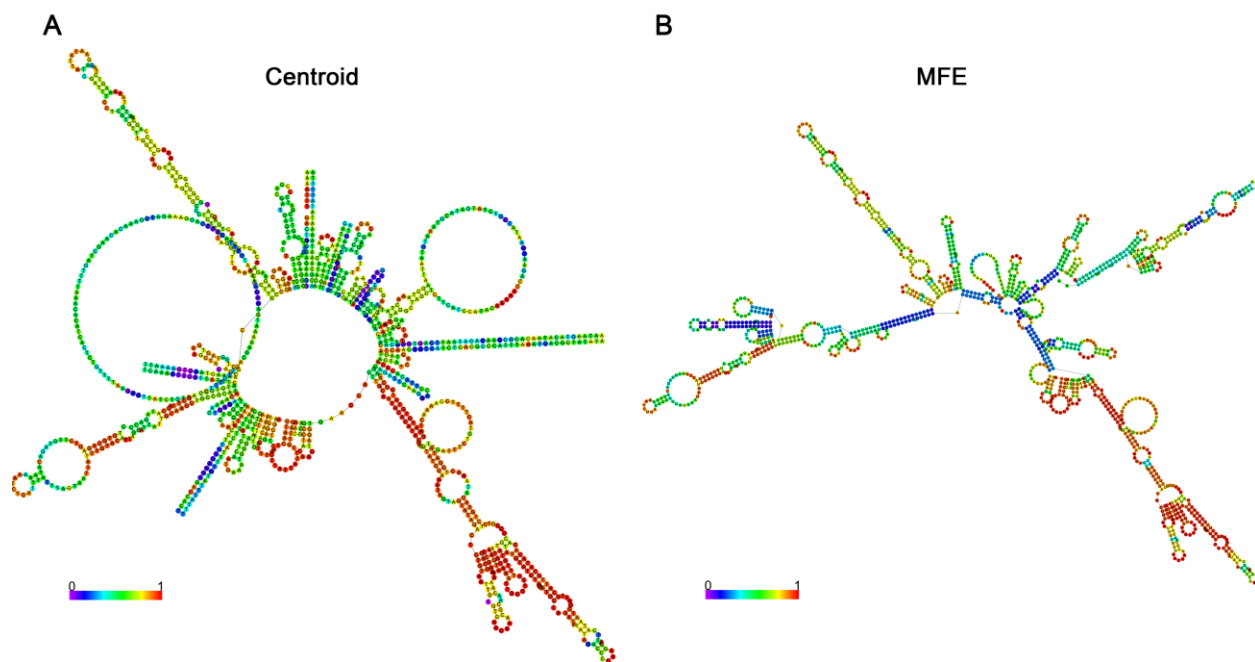
**Fig. S3. The PAZ domain of Dcr-2 has no interaction with the 3'UTR of DIF or Dorsal in vitro.** (A-B) Gel shift assays were performed to evaluate the capacity of MBP-PAZ to bind in vitro transcribed *dif* (A) or *dorsal* (B) 3'UTR as indicated.

Figure S4



**Fig. S4. Toll signaling can be induced by FHV or VSV.** (A-B) Cultured S2 cells were infected with FHV or VSV for 6 hrs (A) or 48 hrs (B). After that, total RNA extracts were prepared, followed by Northern blots using the indicated probes.

Figure S5



**Fig. S5. The prediction of the secondary structure of *Toll* 3'UTR.** (A-B) The secondary structure of *Toll* 3' UTR was predicted using two different algorithms, MFE (minimum free energy) (A) and Centroid (B) by RNAfold software.

primers	Sequence (5' to 3') <sup>a,b</sup>	Purpose
Dicer2 dsRNA #1 For	TAATACGACTCACTATAGGATGGAAGATGTGGAAATCAAGCCTC	Construction of <i>in vitro</i> -transcription templates for preparing dsRNAs for RNAi
Dicer2 dsRNA #1 Rev	TAATACGACTCACTATAGGACAATCGCATGAACTCACGAAATGG	
Dicer2 dsRNA #2 For	TAATACGACTCACTATAGGGAAACATCATTACCGTGTCGACACC	
Dicer2 dsRNA #2 Rev	TAATACGACTCACTATAGGAGGCGTAGACTGGATGTAGTTGAGCAG	
AGO2 dsRNA #1 For	TAATACGACTCACTATAGGATGGGAAAAAAGATAAGAACAAGA	
AGO2 dsRNA #1 Rev	TAATACGACTCACTATAGGCACCTTGTTGACCTTGTTTAGTCCAC	
AGO2 dsRNA #2 For	TAATACGACTCACTATAGGACGATCCCTACGGAGCCAGTTATAAC	Construction of <i>in vitro</i> -transcription templates for preparing probes for Northern blots analysis
AGO2 dsRNA #2 Rev	TAATACGACTCACTATAGGGATTCAACCCATAAGATGAGC	
Toll dsRNA For	TAATACGACTCACTATAGGATGAGTCGACTAAAGGCCGCTCCGAGC	
Toll dsRNA Rev	TAATACGACTCACTATAGGCGGCAACTGTTTCGATGCCATTATCGTG	
Dicer2 probe For	ACAAACGTGGATGTGCCAAAGGCGTTG	
Dicer2 probe Rev	TAATACGACTCACTATAGGTTAGGCGTCGCATTTGCTTAGCTGC	
AGO2 probe For	GGGCAAGAAGGTGGCTACCAACAGCG	Construction of <i>in vitro</i> -transcription templates for preparing Toll 3'UTR RNA
AGO2 probe Rev	TAATACGACTCACTATAGGCGCTGTTGGTAGCCACCTTCTTGCCC	
Drosomycin probe For	ATAATTCAAACAGAAATCATTTACCAAGCTCCGTGAGAAC	
Drosomycin probe Rev	TAATACGACTCACTATAGGAATGTACATTAGTTTTGTTTATTAG	
Toll ORF probe For	TGCGTGAACGCCGAGATGCCAACACG	
Toll ORF probe Rev	TAATACGACTCACTATAGGGAGCTGGAACCTCTGAGGACCGTGCTC	
Toll 3'UTR probe For	TATGTGTATCGCACAGCAGATTCAAGGACCTTACAC	Primers used for qRT-PCR
Toll 3'UTR probe Rev	TAATACGACTCACTATAGGAAATTTTAAATATATTTTATTTGCGTTGTATG	
Toll 3'UTR RNA For	TAATACGACTCACTATAGGGACTGGGAGAAGGCGGAGCTGTTTCAG	
Toll 3'UTR RNA Rev	AAATTTTAAATATATTTTATTTGCGTTGTATG	
Rp49 qRT For	AAGAAGCGCACCAAGCACTTCATC	
Rp49 qRT Rev	TCTGTTGTCGATACCCTTGGGCTT	
Toll qRT For	TGCGTGCAGTGAGATGAGCATAGA	
Toll qRT Rev	TGATCTGCACGTAGTCTTTGGGCT	
Drosomycin qRT For	CGTGAGAACCTTTTCCAATATGATG	
Drosomycin qRT Rev	TCCCAGGACCACCAGCAT	



Diptericin qRT For	ACCGCAGTACCCACTCAATC	
Diptericin qRT Rev	CCCAAGTGCTGTCCATATCC	
dMyd88 qRT For	ATCTGGAACACTTCCTGGGC	
dMyd88 qRT Rev	CCACGAGAGCAGTCTGTCC	
Cactus qRT For	CTCACTAGCCACTAGCGGTAA	
Cactus qRT Rev	CCCGAATCACTGGTTTCGTTT	
Dorsal qRT For	ATCCGTGTGGATCCGTTTAA	
Dorsal qRT Rev	AATCGCACCGAATTCAGATC	
PGRP-LC qRT For	ACCCACCAATTTGTCCTTTT	
PGRP-LC qRT Rev	CAGTACGATACCCAGCAGGAA	
DIF qRT For	GGAGCCGACAAGCAATATAATCC	
DIF qRT Rev	GTAGTTGCACACTTCGATGGT	
Relish qRT For	TGGATACCATCAAAAATGGCCTG	
Relish qRT Rev	CTTGTACCGAAAGCGGAACCT	
Imd qRT For	AGGGACGCCTGGAAAAGGA	
Imd qRT Rev	GGATTCGGTCAGATCCGAGGA	
pMT Toll For	<u>GGGGTACCGCC</u> ACCATGGATTACAAGGATGACGACGATAAGATGAGTCGACTAAAGGC	Construction of plasmids for expressing proteins in S2 cells
	CGC ( <i>Kpn I</i> )	
pMT Toll Rev	<u>GCTCTAGATTATACGTCGCTCTGTTGGCATT</u> C ( <i>Xba I</i> )	
pMT Dicer2 For	ATAAGAAT <u>GCGGCCGCGGCATGGAAGATGTGGAAATCAAGCCTC</u> ( <i>Not I</i> )	
pMT Dicer2 Rev	<u>CGGGGTACCTT</u> ACTTATCGTCGTCATCCTTGTAATCGCCGGCGTCGCATTTGCTTAGCTG CTGA ( <i>Kpn I</i> )	
pAc PAZ For	<u>CGGAATTCGCC</u> ACCATGGAGTACACGGAACACATGTATTTAAATC ( <i>EcoR I</i> )	
pAc PAZ Rev	<u>GCTCTAGATT</u> ACTTATCGTCGTCATCCTTGTAATCGTTGAACTTCCACGATTATGAACG ( <i>Xba I</i> )	
pAc dsRBD For	<u>CGGAATTCGCC</u> ACCATGGTGCCCATCAATCATATACGACAGC ( <i>EcoR I</i> )	
pAc dsRBD For	<u>GCTCTAGATT</u> ACTTATCGTCGTCATCCTTGTAATCGGCGTCGCATTTGCTTAGCTGCTGA AGG ( <i>Xba I</i> )	
pGEX DExD/H For	<u>GGAATTCAGCAATGGCATTGTCTACCTGCCAC</u> ( <i>EcoR I</i> )	Construction of plasmids for expressing proteins in <i>E. coli</i>
pGEX DExD/H Rev	<u>CGCTCGAGTTAAGCCCGACCCTTTGACTGCACGTAC</u> ( <i>Xho I</i> )	
pMAL PAZ For	<u>CGGAATTCGAGTACACGGAACACATGTATTTAAATC</u> ( <i>EcoR I</i> )	
pMAL PAZ Rev	<u>GCGTCGACTTAGTTGAACTTCCACGATTATGAACG</u> ( <i>Sal I</i> )	
pMAL RNase III For	<u>CCGGAATTCGATATTTGTTGCAGGCCCTCACCCATCCCTC</u> ( <i>EcoR I</i> )	
pMAL RNase III Rev	<u>CCCAAGCTTTTAGAGCTTTGCGTTCTCCGCT</u> ( <i>Hind III</i> )	
pMAL dsRBD For	<u>CGGAATTCGTCGCCATCAATCATATACGACAGC</u> ( <i>EcoR I</i> )	
pMAL dsRBD Rev	<u>GCGTCGACTTAGGCGTCGCATTTGCTTAGCTGCTGAAGG</u> ( <i>Sal I</i> )	

<sup>a</sup>Underlined letters indicate restriction endonuclease sites, and the types of the restriction endonucleases are shown in parentheses. The T7 polymerase promoter sequence is shown in italics.

<sup>b</sup>Sequence-specific primers are designed according to the sequences with GenBank accession numbers NM079054 (Dicer2), NM140518 (AGO2), NM079794 (Toll) and Y13939 (Rp49).