

***Dcf1* deficiency attenuates the role of activated microglia during
neuroinflammation**

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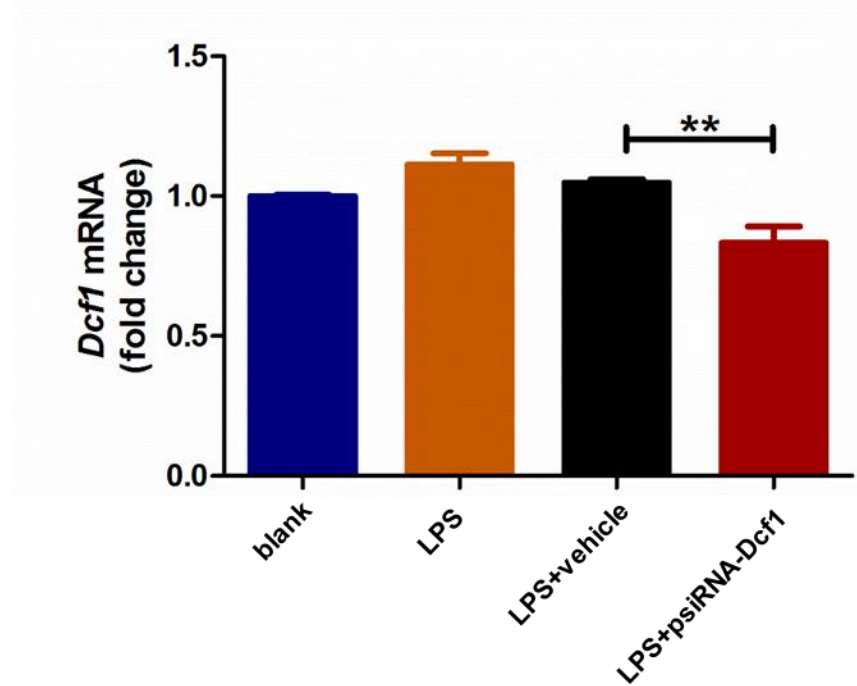
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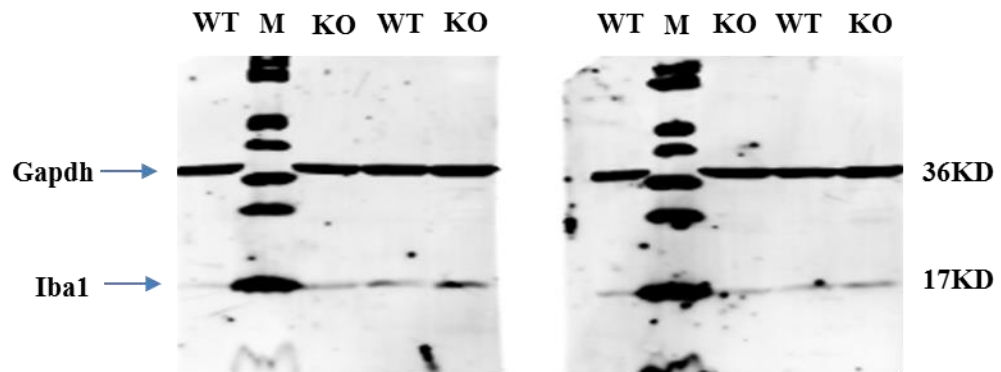
Supplementary Figures:



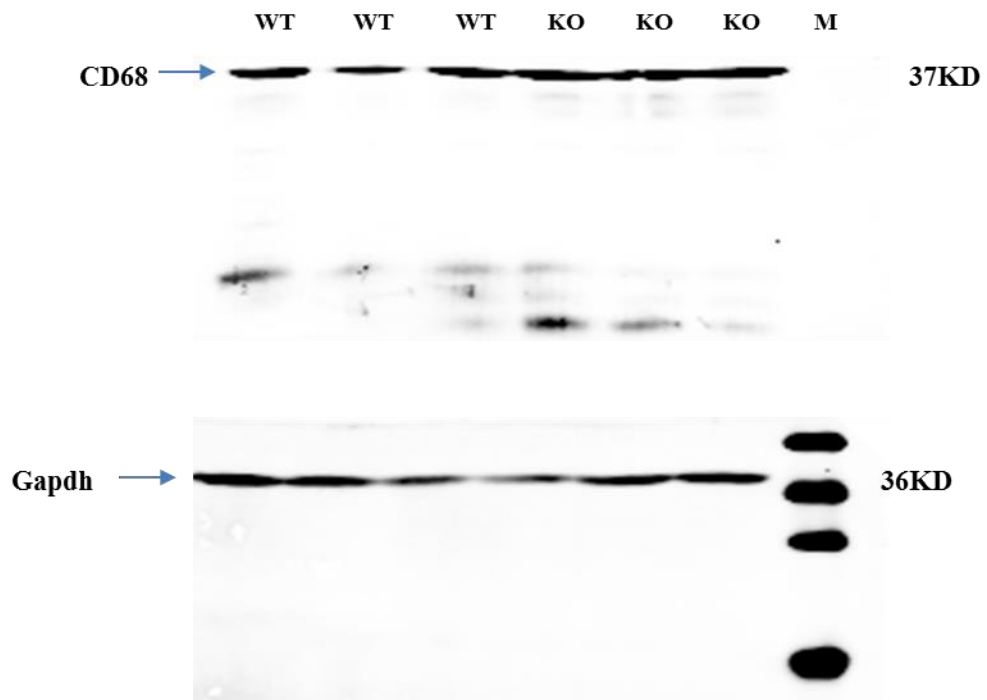
Supplementary Figure S1. *Dcf1* knockdown efficiency in BV2 cells.

Plasmids were transfected 48 h prior to treatment with LPS for 12 h. *Dcf1* knockdown identification by qPCR. Abundance of the *Dcf1* mRNA transcript is expressed relative to *Gapdh* as an internal control. Data were expressed as the mean $\pm$ SEM. n= 4. *, $p<0.05$; **, $p<0.01$; ***, $p<0.001$ (Supplementary Table 13).

A

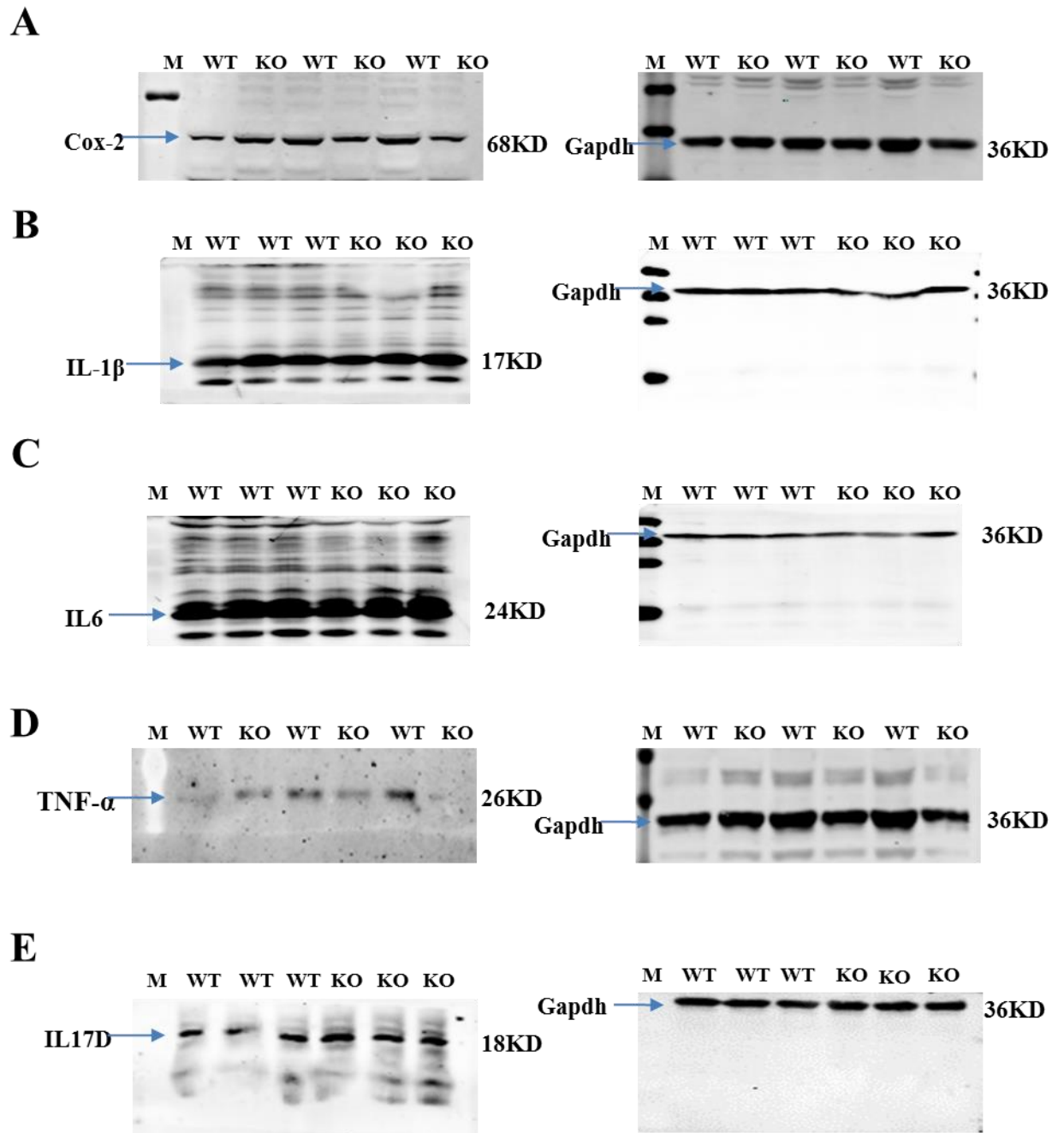


B



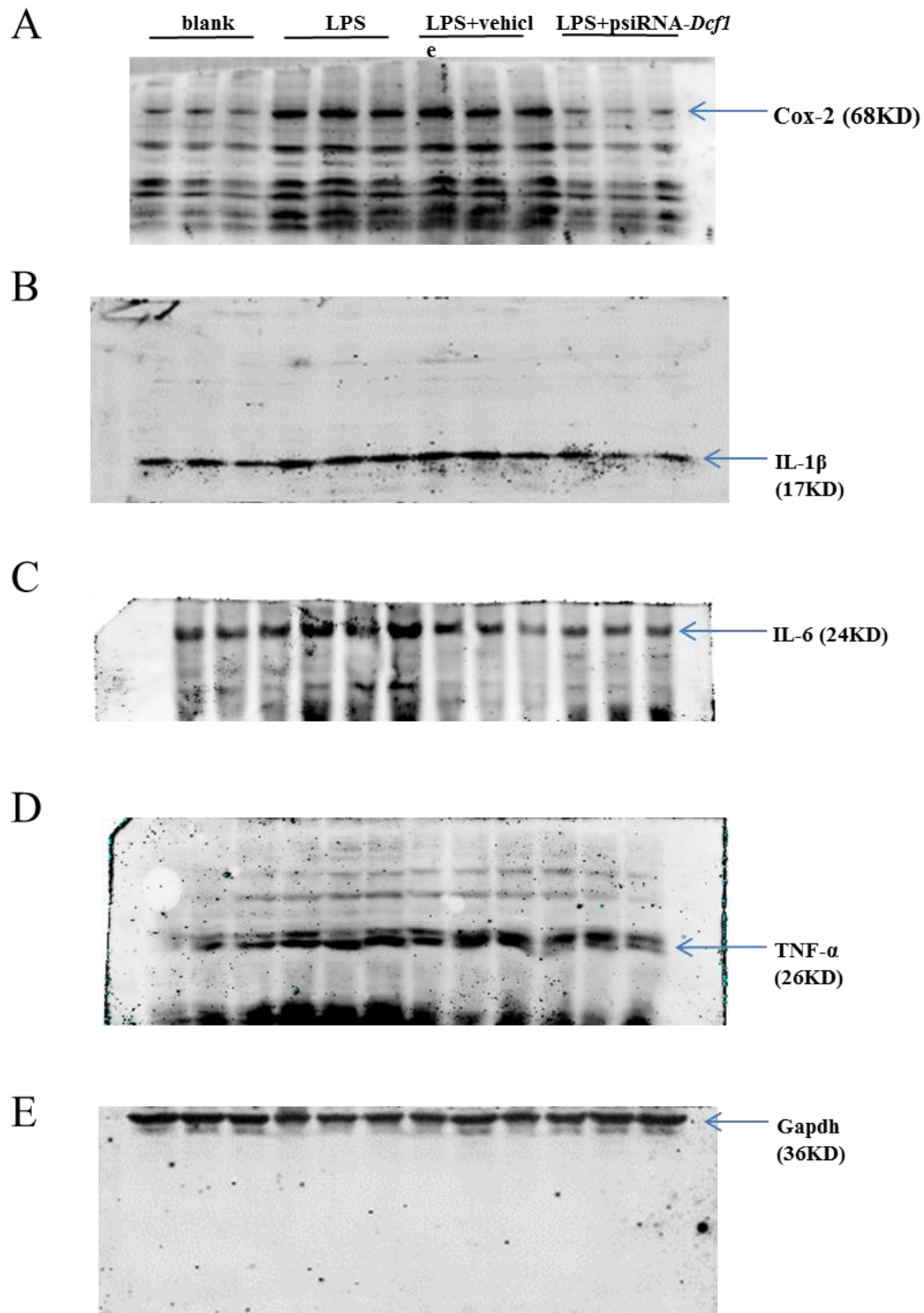
Supplementary Figure S2. *Dcf1* deletion induces upregulation of microglial activation markers *in vivo*

Protein expression of microglial activation markers Iba1 (F) and CD68 (G) in WT and *Dcf1*-KO mouse brain tissue. Quantification of protein expression levels normalized to Gapdh.



Supplementary Figure S3. Expression of proinflammatory cytokines in WT and *Dcf1*-KO mice

Protein expression of Cox-2(A), IL-1 β (B), IL-6(C), TNF- α (D) and IL17D(E) were assessed by Western blotting in WT and KO mice.



Supplementary Figure S4. Effects of *Dcf1* downregulation on proinflammatory cytokines expression in LPS-stimulated BV2 microglia

BV₂ microglial cells were transfected with the psiRNA-hH1neo plasmid or the psiRNA-*Dcf1* plasmid. 24 h post-transfection, BV2 microglia were stimulated with LPS (1000 ng/ml) and incubated for 12 h. Protein expression of COX-2(A), IL-1 β (B), TNF- α (C) and IL-6(D) were assessed by Western blotting.

Table 1: t-test analysis of Figure. 2C

Unpaired t-test	p value	Significant? p<0.05	Summary
WT vs KO	0.0084	Yes	**

Data are expressed as the mean ± SEM. n = 3.

*, $p<0.05$; **, $p<0.01$; ***, $p<0.001$ vs. WT.

Table 2: t-test analysis of Figure. 2E

Unpaired t-test	p value	Significant? p<0.05	Summary
WT vs KO	0.0151	Yes	*

Data are expressed as the mean ± SEM. n = 4.

*, $p<0.05$; **, $p<0.01$; ***, $p<0.001$ vs. WT.

Table 3: t-test analysis of Figure. 2F

Unpaired t-test	p value	Significant? p<0.05	Summary
WT vs KO	0.0353	Yes	*

Data are expressed as the mean ± SEM. n = 3.

*, $p<0.05$; **, $p<0.01$; ***, $p<0.001$ vs. WT.

Table 4: t-test analysis of Figure. 2G

Unpaired t-test	p value	Significant? p<0.05	Summary
WT vs KO	0.0264	Yes	*

Data are expressed as the mean $\pm$ SEM. n = 3.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. WT.

Table 5: t-test analysis of Figure. 3A

Unpaired t-test	p value	Significant? $p < 0.05$	Summary
● <i>Cox2</i> :			
WT vs KO	0.0433	Yes	*
● <i>IL-1beta</i> :			
WT vs KO	0.0441	Yes	*
● <i>Tnfsf111</i> :			
WT vs KO	0.7862	no	ns
● <i>Cxcl1</i> :			
WT vs KO	0.2350	no	ns
● <i>Ccl7</i> :			
WT vs KO	0.0333	Yes	*
● <i>IL-6</i> :			
WT vs KO	0.0132	Yes	*
● <i>IL17D</i> :			
WT vs KO	0.0238	Yes	*
● <i>TNF-alpha</i> :			
WT vs KO	0.0364	Yes	*
● <i>Csf1</i> :			
WT vs KO	0.0200	Yes	*

Data are expressed as the mean $\pm$ SEM. n = 3.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. WT.

Table 6: t-test analysis of Figure. 3B

Unpaired t-test	p value	Significant? $p < 0.05$	Summary
● <i>Cox2</i> :			
WT vs KO	0.047751	Yes	*
● <i>IL-1beta</i> :			
WT vs KO	0.048958	Yes	*
● <i>IL-6</i> :			

WT vs KO	0.047189	Yes	*
● IL17D:			
WT vs KO	0.035709	Yes	*
● TNF-alpha:			
WT vs KO	0.012969	Yes	*

Data are expressed as the mean $\pm$ SEM. n = 3.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. WT.

Table 7: t-test analysis of Figure. 4C

Unpaired t-test	p value	Significant? $p < 0.05$	Summary
blank vs LPS	3.45×10^{-5}	Yes	***

Data are expressed as the mean $\pm$ SEM. n = 6.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. blank.

Table 8: One-way ANOVA analysis of Figure. 4D

Bonferroni's Multiple Comparison Test	Significant? $p < 0.05$	Summary
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	*
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	*

Data are expressed as the mean $\pm$ SEM. n = 6.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$

Table 9: One-way ANOVA analysis of Figure. 5A

Bonferroni's Multiple Comparison Test	Significant? $p < 0.05$	Summary
● <i>Cox2</i> :		

blank vs LPS	Yes	**
blank vs LPS+vehicle	Yes	***
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	*
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	##
● <i>IL-1β</i> :		
blank vs LPS	Yes	*
blank vs LPS+vehicle	Yes	*
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#
● <i>Tnfsf111</i> :		
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	No	ns
● <i>Cxcl1</i> :		
blank vs LPS	Yes	*
blank vs LPS+vehicle	Yes	*
blank vs LPS+psiRNA- <i>Dcf1</i>	Yes	***
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	***
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	###
● <i>Ccl7</i> :		
blank vs LPS	Yes	***
blank vs LPS+vehicle	Yes	**
blank vs LPS+psiRNA- <i>Dcf1</i>	Yes	***
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#
● <i>IL-6</i> :		
blank vs LPS	Yes	*
blank vs LPS+vehicle	Yes	*
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns

LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#
● <i>IL17D</i> :		
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	No	ns
● <i>TNF-α</i> :		
blank vs LPS	Yes	*
blank vs LPS+vehicle	Yes	**
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#
● <i>Csf1</i> :		
blank vs LPS	Yes	**
blank vs LPS+vehicle	Yes	**
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	**
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	##

Data are expressed as the mean $\pm$ SEM. n = 4.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. blank.

#, $p < 0.05$; ##, $p < 0.01$; ###, $p < 0.001$ vs. LPS+vehicle.

Table 10: One-way ANOVA analysis of Figure. 5B

Bonferroni's Multiple Comparison Test	Significant? $p < 0.05$	Summary
● <i>Cox2</i> :		
blank vs LPS	Yes	***
blank vs LPS+vehicle	Yes	**
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	**
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	##

● IL-1 β :		
blank vs LPS	Yes	**
blank vs LPS+vehicle	Yes	*
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	*
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#
● IL6:		
blank vs LPS	Yes	**
blank vs LPS+vehicle	Yes	*
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	**
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#
● TNF- α :		
blank vs LPS	Yes	**
blank vs LPS+vehicle	Yes	**
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	*
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	#

Data are expressed as the mean $\pm$ SEM. n = 3.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. blank.

#, $p < 0.05$; ##, $p < 0.01$; ###, $p < 0.001$ vs. LPS+vehicle.

Table 11: One-way ANOVA analysis of Figure. 6B

Bonferroni's Multiple Comparison Test	Significant? $p < 0.05$	Summary
● 12h:		
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	**
● 24h:		
blank vs LPS	No	ns

blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	No	ns
● 36h:		
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	No	ns
● 48h:		
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	Yes	***
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	***
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	###

Data are expressed as the mean $\pm$ SEM. n = 8.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. blank.

#, $p < 0.05$; ##, $p < 0.01$; ###, $p < 0.001$ vs. LPS+vehicle.

Table 12: One-way ANOVA analysis of Figure. 7B

Bonferroni's Multiple Comparison Test	Significant? $p < 0.05$	Summary
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	No	ns
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	*

Data are expressed as the mean $\pm$ SEM. n = 4.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. blank.

#, $p < 0.05$; ##, $p < 0.01$; ###, $p < 0.001$ vs. LPS+vehicle.

Table 13: One-way ANOVA analysis of Supplement Figure. 1

Bonferroni's Multiple Comparison Test	Significant? $p < 0.05$	Summary
blank vs LPS	No	ns
blank vs LPS+vehicle	No	ns
blank vs LPS+psiRNA- <i>Dcf1</i>	Yes	*
LPS vs LPS+vehicle	No	ns
LPS vs LPS+psiRNA- <i>Dcf1</i>	Yes	***
LPS+vehicle vs LPS+psiRNA- <i>Dcf1</i>	Yes	**

Data are expressed as the mean $\pm$ SEM. n = 4.

*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ vs. LPS+vehicle.