



Supplementary Figure 1. Analysis of the content of NIK and TRAF3 in the mock- and ECTV-infected RAW 264.7 macrophages. **(A)** Immunoblot analysis of NIK content in mock- and ECTV-infected RAW 264.7 cells that were untreated or treated with poly(I:C), Pam3CSK4, or rmIFN- γ + *Escherichia coli* LPS O111:B4 at 4 hpi. **(B)** Densitometric evaluation of NIK expression in ECTV-infected RAW 264.7 cells (4 hpi). The analysis was based on the results of three independent biological experiments. The data are presented on a histogram with a logarithmic scale. GAPDH—loading control. **(C)** Immunoblot analysis of NIK content in the mock- and ECTV-infected RAW 264.7 cells that were untreated or treated with poly(I:C), Pam3CSK4, or rmIFN- γ + *E. coli* LPS O55:B5 or O111:B4 at 18 hpi. **(D)** Densitometric evaluation of NIK expression in ECTV-infected RAW 264.7 cells (18 hpi). The analysis was based on the results of two independent biological experiments. The data are presented on a histogram with a logarithmic scale. GAPDH—loading control ($*p \leq 0.05$). **(E)** Immunoblot analysis of TRAF3 content in the mock- and ECTV-infected RAW 264.7 cells at 4 hpi. **(F)** Densitometric evaluation of TRAF3 expression in ECTV-infected RAW 264.7 cells (4 hpi). The analysis was based on the results of two independent biological experiments. The data are presented on a histogram with a logarithmic scale. GAPDH—loading control ($*p \leq 0.05$). **(G)** Immunoblot analysis of TRAF3 content in the mock- and ECTV-infected RAW 264.7 cells at 18 hpi. **(H)** Densitometric evaluation of TRAF3 expression in ECTV-infected RAW 264.7 cells (18 hpi). The analysis was based on the results of two independent biological experiments. The data are presented on a histogram with a logarithmic scale. GAPDH—loading control ($*p \leq 0.05$).

Supplementary Table 1. Expression of genes involved in NF- κ B signaling pathways in RAW 264.7 macrophages stimulated with poly(I:C) for 18 h.

Poly(I:C) vs control			
Gene	Fold change	<i>p</i> value	Log ₂ fold change
<i>Atf3</i>	0,9727	0,837472	-0,039933177
<i>Bcl2</i>	0,7186	0,448858	-0,47673916
<i>Bcl3</i>	1,5227	0,170099	0,606631732
<i>Birc2</i>	1,2397	0,39966	0,309991039
<i>Birc3</i>	2,2191	0,073059	1,149974682
<i>Ccl19</i>	0,8685	0,521313	-0,203402246
<i>Ccl25</i>	0,8766	0,963013	-0,190009416
<i>Ccr10</i>	0,8685	0,521313	-0,203402246
<i>Ccr8</i>	0,8685	0,521313	-0,203402246
<i>Cd36</i>	0,5261	0,011111	-0,926591045
<i>Cd40</i>	3,4903	0,042535	1,803351045
<i>Cd68</i>	1,0619	0,737596	0,086647913
<i>Cd86</i>	0,8685	0,521313	-0,203402246
<i>Cflar</i>	1,4845	0,282632	0,569977093
<i>Chuk</i>	1,162	0,499438	0,216610069
<i>Creb1</i>	0,9659	0,952453	-0,050054261
<i>Csf2</i>	0,8685	0,521313	-0,203402246
<i>Cxcl10</i>	2,2346	0,112797	1,160016608
<i>Cxcl5</i>	0,8685	0,521313	-0,203402246
<i>Cxcr5</i>	1,0425	0,690566	0,060047384
<i>Daxx</i>	1,4077	0,255517	0,493339909
<i>Ddx58</i>	3,7668	0,070815	1,913339435
<i>Ets2</i>	2,3511	0,033747	1,233335903
<i>Hdac9</i>	0,9772	0,980521	-0,033274231
<i>Hmgb1</i>	0,5864	0,065331	-0,770042991
<i>Icam1</i>	2,2763	0,043682	1,186690707
<i>Ido1</i>	0,8685	0,521313	-0,203402246
<i>Ifit1</i>	19,2484	0,055576	4,266666623
<i>Ifnb1</i>	0,8685	0,521313	-0,203402246
<i>Ifngr1</i>	0,9223	0,872043	-0,116691997
<i>Ikkkb</i>	0,9885	0,928764	-0,016687129
<i>Ikkke</i>	2,4453	0,065553	1,290011472
<i>Ikkkg</i>	4,0278	0,006102	2,009992048
<i>Il15</i>	2,2868	0,068131	1,193330195
<i>Il18</i>	1,0718	0,700665	0,100035721
<i>Il9</i>	0,8685	0,521313	-0,203402246
<i>Irak1</i>	0,727	0,368887	-0,459972731
<i>Irak3</i>	0,9374	0,967064	-0,0932633
<i>Irf1</i>	0,9223	0,964208	-0,116691997
<i>Irf2</i>	0,9075	0,777673	-0,140030452
<i>Irf3</i>	0,9309	0,913952	-0,103301897
<i>Irf5</i>	0,984	0,911822	-0,023269779
<i>Irf8</i>	0,6507	0,190601	-0,619935541
<i>Jak1</i>	1,162	0,591372	0,216610069
<i>Ltbr</i>	0,8586	0,674539	-0,219941922
<i>Map2k3</i>	0,8351	0,423571	-0,25997913
<i>Mapkapk2</i>	1,2864	0,324814	0,363339312
<i>Mavs</i>	1,021	0,89367	0,029982866
<i>Mmd2</i>	0,8685	0,521313	-0,203402246
<i>Mmp2</i>	0,8685	0,521313	-0,203402246

Poly(I:C) vs control			
Gene	Fold change	p value	Log ₂ fold change
<i>Mmp25</i>	0,8685	0,521313	-0,203402246
<i>Mmp9</i>	57,4141	0,002279	5,843333179
<i>Mtor</i>	1,0234	0,880486	0,033370138
<i>Myd88</i>	1,0281	0,828426	0,039980598
<i>Nfat5</i>	0,9482	0,923404	-0,076736702
<i>Nfix</i>	0,8685	0,521313	-0,203402246
<i>Nfkb1</i>	1,2255	0,375081	0,293370484
<i>Nfkb2</i>	2,1189	0,049813	1,083315502
<i>Nfkbia</i>	3,0738	0,042033	1,620023298
<i>Nfkbib</i>	1,8277	0,073125	0,870029285
<i>Nlrc5</i>	1,0968	0,537277	0,133300476
<i>Nlrp12</i>	0,8685	0,521313	-0,203402246
<i>Otud7b</i>	1,154	0,398019	0,206643224
<i>Rel</i>	0,7544	0,419906	-0,406598419
<i>Relb</i>	1,8361	0,088699	0,876644635
<i>Snf8</i>	0,9159	0,839811	-0,126738005
<i>Sp1</i>	0,8141	0,574751	-0,296722076
<i>Sp110</i>	2,5847	0,049534	1,36999684
<i>Stap2</i>	2,6882	0,09493	1,426640478
<i>Stat1</i>	1,398	0,27485	0,483364361
<i>Stat6</i>	1,257	0,380284	0,32998465
<i>Syk</i>	1,7371	0,123443	0,796680808
<i>Tbk1</i>	1,2658	0,309278	0,340049473
<i>Tlr4</i>	0,7937	0,445875	-0,333334289
<i>Tmem173</i>	0,5743	0,135181	-0,800123533
<i>Tnfrsf11a</i>	0,79	0,233864	-0,340075442
<i>Tnfrsf12a</i>	1,4406	0,26372	0,526669809
<i>Tnfrsf13c</i>	0,8685	0,521313	-0,203402246
<i>Traf1</i>	4,7678	0,020252	2,253323719
<i>Traf2</i>	1,1173	0,588151	0,160016608
<i>Traf3</i>	0,893	0,712742	-0,16326792
<i>Traf6</i>	1,021	0,842702	0,029982866
<i>Vegfb</i>	0,9202	0,767881	-0,119980638
<i>Xbp1</i>	0,7153	0,233946	-0,483379653
<i>Yy1</i>	0,9727	0,95349	-0,039933177
<i>Rn18s</i>	1	0	

The expression of the analyzed genes was evaluated using RT-qPCR. The cells were left untreated or treated with poly(I:C) for 18 h. Significant changes ($p \leq 0.05$, \log_2 fold change ≥ 1) in gene expression are shown in red (upregulation). *Rn18s* – reference gene. Data were obtained from three independent experiments.

Supplementary Table 2. Expression of genes involved in NF- κ B signaling pathways in RAW 264.7 macrophages stimulated with IFN+LPS for 18 h.

IFN+LPS vs control			
Gene	Fold change	p value	Log ₂ fold change
<i>Aim2</i>	1,1755	0,376043	0,233274539
<i>Atf3</i>	8,0556	0,001672	3,009992048
<i>Bcl2</i>	1,1947	0,443994	0,25664839
<i>Bcl3</i>	4,3671	0,002542	2,126675567
<i>Birc2</i>	1,1173	0,485158	0,160016608
<i>Birc3</i>	2,0046	0,022456	1,003314389
<i>Ccl19</i>	0,79	0,277318	-0,340075442

IFN+LPS vs control			
Gene	Fold change	p value	Log ₂ fold change
<i>Ccl25</i>	0,732	0,037062	-0,450084446
<i>Ccr10</i>	0,79	0,277318	-0,340075442
<i>Ccr8</i>	0,79	0,277318	-0,340075442
<i>Cd36</i>	0,7992	0,003956	-0,323371512
<i>Cd40</i>	14,1559	0,000531	3,82333157
<i>Cd68</i>	1,0792	0,669441	0,109962253
<i>Cd86</i>	24,0284	0,004954	4,586668681
<i>Cflar</i>	3,387	0,0008	1,760007987
<i>Chuk</i>	1,0867	0,625717	0,11953718
<i>Creb1</i>	1,0693	0,712791	0,096666669
<i>Csf2</i>	29,5824	0,023173	4,886667197
<i>Cxcl10</i>	10,3627	0,099441	3,373328041
<i>Cxcl5</i>	0,79	0,277318	-0,340075442
<i>Cxcr5</i>	0,5023	0,014451	-0,99337882
<i>Daxx</i>	3,3558	0,000113	1,746656736
<i>Ddx58</i>	13,7688	0,000444	3,783330924
<i>Ets2</i>	3,793	0,001497	1,923339372
<i>Hdac9</i>	0,0976	0,000009	-3,356975042
<i>Hmgb1</i>	0,2137	0,000341	-2,226341187
<i>Icam1</i>	0,4622	0,004463	-1,113410835
<i>Ido1</i>	0,79	0,277318	-0,340075442
<i>Ifit1</i>	321,7954	0,003255	8,329999893
<i>Ifnb1</i>	0,79	0,277318	-0,340075442
<i>Ifngr1</i>	0,3669	0,019232	-1,44654119
<i>Ikkkb</i>	0,8312	0,242412	-0,266732441
<i>Ikkke</i>	2,362	0,006043	1,240008965
<i>Ikkkg</i>	3,2944	0,01148	1,720015735
<i>Il15</i>	3,5884	0,008991	1,843340717
<i>Il18</i>	3,3404	0,017744	1,74002087
<i>Il9</i>	0,79	0,277318	-0,340075442
<i>Irak1</i>	0,52	0,018042	-0,943416472
<i>Irak3</i>	8,187	0,00222	3,033334895
<i>Irf1</i>	1,2142	0,029227	0,280006078
<i>Irf2</i>	0,5535	0,035182	-0,853344778
<i>Irf3</i>	0,7614	0,108418	-0,393273525
<i>Irf5</i>	0,4137	0,00059	-1,273343137
<i>Irf8</i>	1,9908	0,004015	0,993348292
<i>Jak1</i>	2,1987	0,002813	1,13665077
<i>Ltbr</i>	1,0892	0,664786	0,123268887
<i>Map2k3</i>	0,3807	0,005289	-1,393273525
<i>Mapkapk2</i>	1,3787	0,045093	0,463308566
<i>Mavs</i>	1,2894	0,27235	0,366699889
<i>Mmd2</i>	0,79	0,277318	-0,340075442
<i>Mmp2</i>	0,79	0,277318	-0,340075442
<i>Mmp25</i>	4,2673	0,010628	2,093323539
<i>Mmp9</i>	71,3414	0,005554	6,156667622
<i>Mtor</i>	0,7055	0,147103	-0,503282012
<i>Myd88</i>	0,7087	0,10163	-0,496753046
<i>Nfat5</i>	1,3134	0,011848	0,39330636
<i>Nfix</i>	0,79	0,277318	-0,340075442
<i>Nfkb1</i>	2,0753	0,011231	1,053319904
<i>Nfkb2</i>	4,3974	0,003583	2,13665077
<i>Nfkbia</i>	9,2964	0,000836	3,216672146

IFN+LPS vs control			
Gene	Fold change	p value	Log ₂ fold change
<i>Nfkbib</i>	4,7459	0,000054	2,246681702
<i>Nlrc5</i>	3,6217	0,000655	1,856667047
<i>Nlrp12</i>	0,79	0,277318	-0,340075442
<i>Otud7b</i>	1,617	0,016423	0,693319679
<i>Rel</i>	2,0326	0,015182	1,023326332
<i>Relb</i>	1,4641	0,04962	0,550014095
<i>Snf8</i>	0,9033	0,197377	-0,146722886
<i>Sp1</i>	0,7509	0,151336	-0,413307303
<i>Sp110</i>	5,3517	0,000013	2,419997245
<i>Stap2</i>	2,8481	0,004973	1,509999802
<i>Stat1</i>	4,5842	0,001479	2,196669988
<i>Stat6</i>	1,0473	0,726421	0,066674763
<i>Syk</i>	8,0556	0,000924	3,009992048
<i>Tbk1</i>	2,114	0,002412	1,079975377
<i>Tlr4</i>	1,4175	0,051654	0,503348735
<i>Tmem173</i>	0,8179	0,402	-0,290003631
<i>Tnfrsf11a</i>	0,3703	0,000024	-1,433233545
<i>Tnfrsf12a</i>	0,8586	0,212509	-0,219941922
<i>Tnfrsf13c</i>	0,79	0,277318	-0,340075442
<i>Traf1</i>	9,736	0,000331	3,283329168
<i>Traf2</i>	1,2834	0,043059	0,359970888
<i>Traf3</i>	0,6537	0,051457	-0,613299398
<i>Traf6</i>	1,8025	0,001163	0,849999259
<i>Vegfb</i>	0,7614	0,21571	-0,393273525
<i>Xbp1</i>	0,6043	0,068131	-0,726663153
<i>Yy1</i>	0,7614	0,089189	-0,393273525
<i>Rn18s</i>	1	0	

The expression of the analyzed genes was evaluated using RT-qPCR. The cells were left untreated or treated with rmIFN- γ + *Escherichia coli* LPS O111:B4 for 18 h. Significant changes ($p \leq 0.05$, log₂ fold change ≤ -1 or ≥ 1) in gene expression are shown in colors (blue-downregulation, red-upregulation). *Rn18s* – reference gene. Data were obtained from three independent experiments.

Supplementary Table 3. Expression of genes involved in NF- κ B signaling pathways in RAW 264.7 macrophages infected with ECTV (18 hpi).

ECTV vs control			
Gene	Fold change	p value	Log ₂ fold change
<i>Aim2</i>	1,1173	0,602205	0,160016608
<i>Atf3</i>	1,3755	0,29558	0,45995614
<i>Bcl2</i>	0,9482	0,770468	-0,076736702
<i>Bcl3</i>	0,8526	0,326947	-0,23005904
<i>Birc2</i>	0,8706	0,339986	-0,199918075
<i>Birc3</i>	0,4147	0,002874	-1,269860048
<i>Ccl19</i>	0,8217	0,336514	-0,283316328
<i>Ccl25</i>	1,2454	0,204151	0,316609184
<i>Ccr10</i>	0,8217	0,336514	-0,283316328
<i>Ccr8</i>	0,8217	0,336514	-0,283316328
<i>Cd36</i>	0,8274	0,534445	-0,273343137
<i>Cd40</i>	0,9615	0,786884	-0,056641237
<i>Cd68</i>	0,9772	0,975382	-0,033274231
<i>Cd86</i>	0,9461	0,788723	-0,079935415
<i>Cflar</i>	1,2114	0,482786	0,276675317
<i>Chuk</i>	0,8766	0,541925	-0,190009416

ECTV vs control

Gene	Fold change	p value	Log ₂ fold change
<i>Creb1</i>	0,8467	0,357689	-0,240077206
<i>Csf2</i>	0,8217	0,336514	-0,283316328
<i>Cxcl10</i>	0,4147	0,010591	-1,269860048
<i>Cxcl5</i>	0,8217	0,336514	-0,283316328
<i>Cxcr5</i>	0,8123	0,149095	-0,299915451
<i>Daxx</i>	0,9054	0,650766	-0,143372788
<i>Ddx58</i>	1,0644	0,958966	0,090040415
<i>Ets2</i>	0,8546	0,209773	-0,226678778
<i>Hdac9</i>	0,7974	0,34211	-0,32662449
<i>Hmgb1</i>	1,0305	0,789881	0,043344505
<i>Icam1</i>	0,5824	0,008824	-0,779917739
<i>Ido1</i>	0,8217	0,336514	-0,283316328
<i>Ifit1</i>	1,1755	0,531604	0,233274539
<i>Ifnb1</i>	0,8217	0,336514	-0,283316328
<i>Ifngr1</i>	0,9862	0,901123	-0,020047842
<i>Ikbkb</i>	0,8971	0,566751	-0,156659283
<i>Ikbke</i>	0,4527	0,010334	-1,143372788
<i>Ikbkg</i>	0,4965	0,008475	-1,010134377
<i>Il15</i>	1,0093	0,953586	0,013355059
<i>Il18</i>	0,8625	0,647426	-0,213403638
<i>Il9</i>	0,8217	0,336514	-0,283316328
<i>Irak1</i>	0,9593	0,88127	-0,059946038
<i>Irak3</i>	0,839	0,206032	-0,253257284
<i>Irf1</i>	0,8606	0,193443	-0,216585255
<i>Irf2</i>	0,8766	0,396661	-0,190009416
<i>Irf3</i>	0,9181	0,617204	-0,123276793
<i>Irf5</i>	0,7614	0,178038	-0,393273525
<i>Irf8</i>	1,2716	0,328404	0,346644922
<i>Jak1</i>	1,0994	0,646682	0,136716384
<i>Ltbr</i>	1,1277	0,608658	0,173383321
<i>Map2k3</i>	0,9593	0,770533	-0,059946038
<i>Mapkapk2</i>	1,0163	0,920971	0,023326332
<i>Mavs</i>	1,0521	0,89971	0,073271836
<i>Mmd2</i>	0,8217	0,336514	-0,283316328
<i>Mmp2</i>	0,8217	0,336514	-0,283316328
<i>Mmp25</i>	0,8217	0,336514	-0,283316328
<i>Mmp9</i>	0,8217	0,336514	-0,283316328
<i>Mtor</i>	1,0449	0,923134	0,063364879
<i>Myd88</i>	0,9395	0,768999	-0,090034933
<i>Nfat5</i>	0,9374	0,532508	-0,0932633
<i>Nfix</i>	0,8217	0,336514	-0,283316328
<i>Nfkb1</i>	0,8507	0,597675	-0,233277641
<i>Nfkb2</i>	0,395	0,01653	-1,340075442
<i>Nfkbia</i>	0,4434	0,005844	-1,173319325
<i>Nfkbib</i>	0,6814	0,026913	-0,553426147
<i>Nlrc5</i>	0,9548	0,921309	-0,066729528
<i>Nlrp12</i>	0,8217	0,336514	-0,283316328
<i>Otud7b</i>	0,9461	0,743349	-0,079935415
<i>Rel</i>	1,0281	0,895858	0,039980598
<i>Relb</i>	0,5129	0,012834	-0,963250524
<i>Snf8</i>	0,9309	0,930681	-0,103301897
<i>Sp1</i>	1,0305	0,858416	0,043344505
<i>Sp110</i>	0,8448	0,437815	-0,24331826

ECTV vs control			
Gene	Fold change	p value	Log ₂ fold change
<i>Stap2</i>	0,7774	0,373436	-0,363270987
<i>Stat1</i>	0,9117	0,541575	-0,133368919
<i>Stat6</i>	0,9615	0,858482	-0,056641237
<i>Syk</i>	1,3787	0,209777	0,463308566
<i>Tbk1</i>	0,9931	0,975308	-0,009989098
<i>Tlr4</i>	0,8428	0,304096	-0,246737781
<i>Tmem173</i>	1,1783	0,51165	0,236706902
<i>Tnfrsf11a</i>	0,5	0,008603	-1
<i>Tnfrsf12a</i>	0,5743	0,048308	-0,800123533
<i>Tnfrsf13c</i>	0,8217	0,336514	-0,283316328
<i>Traf1</i>	0,2031	0,001663	-2,299737855
<i>Traf2</i>	0,7337	0,262713	-0,44673781
<i>Traf3</i>	0,9013	0,484798	-0,149920704
<i>Traf6</i>	1,1728	0,275573	0,229957009
<i>Vegfb</i>	1,2628	0,382737	0,336626166
<i>Xbp1</i>	0,8274	0,366386	-0,273343137
<i>Yy1</i>	0,9885	0,990923	-0,016687129
<i>Rn18s</i>	1	0	

The expression of the analyzed genes was evaluated using RT-qPCR. The cells were uninfected or infected with ECTV for 18 h. Significant changes ($p \leq 0.05$, log₂ fold change ≤ -1) in gene expression are shown in colors (blue-downregulation). *Rn18s* – reference gene. Data were obtained from three independent experiments.

Supplementary Table 4. Expression of genes involved in NF- κ B signaling pathways in RAW 264.7 macrophages infected with ECTV and stimulated with poly(I:C) (18 hpi).

ECTV/Poly(I:C) vs Poly(I:C)			
Gene	Fold change	p value	Log ₂ fold change
<i>Aim2</i>	0,6285	0,212908	-0,67001535
<i>Atf3</i>	0,4774	0,022207	-1,066729528
<i>Bcl2</i>	0,5023	0,172089	-0,99337882
<i>Bcl3</i>	0,7405	0,301668	-0,433428359
<i>Birc2</i>	0,4434	0,074796	-1,173319325
<i>Birc3</i>	0,2343	0,030834	-2,093571141
<i>Ccl19</i>	0,9977	0,912319	-0,00332202
<i>Ccl25</i>	0,8332	0,585547	-0,263265256
<i>Ccr10</i>	0,9977	0,912319	-0,00332202
<i>Ccr8</i>	0,9977	0,912319	-0,00332202
<i>Cd36</i>	0,8236	0,381239	-0,279984265
<i>Cd40</i>	0,2387	0,035577	-2,066729528
<i>Cd68</i>	0,4393	0,084845	-1,186721595
<i>Cd86</i>	0,9977	0,912319	-0,00332202
<i>Cflar</i>	0,5012	0,091022	-0,99654168
<i>Chuk</i>	0,4665	0,048302	-1,100051014
<i>Creb1</i>	0,5212	0,069633	-0,940091011
<i>Csf2</i>	0,9977	0,912319	-0,00332202
<i>Cxcl10</i>	0,2535	0,062131	-1,979942348
<i>Cxcl5</i>	0,9977	0,912319	-0,00332202
<i>Cxcr5</i>	0,6926	0,549072	-0,529905707
<i>Daxx</i>	0,4147	0,061215	-1,269860048
<i>Ddx58</i>	0,1032	0,038097	-3,276485124
<i>Ets2</i>	0,483	0,047249	-1,049904906
<i>Hdac9</i>	0,5421	0,056488	-0,883369088

ECTV/Poly(I:C) vs Poly(I:C)			
Gene	Fold change	p value	Log ₂ fold change
<i>Hmgb1</i>	0,8971	0,572062	-0,156659283
<i>Icam1</i>	0,3914	0,034135	-1,353284339
<i>Ido1</i>	0,9977	0,912319	-0,00332202
<i>Ifit1</i>	0,08	0,060512	-3,64385619
<i>Ifnb1</i>	0,9977	0,912319	-0,00332202
<i>Ifngr1</i>	0,4931	0,104808	-1,020047842
<i>Ikbkb</i>	0,4137	0,05596	-1,273343137
<i>Ikbke</i>	0,2553	0,034131	-1,969734557
<i>Ikbkg</i>	0,3157	0,008588	-1,663373834
<i>Il15</i>	0,1715	0,018517	-2,543719518
<i>Il18</i>	0,2606	0,165576	-1,940091011
<i>Il9</i>	0,9977	0,912319	-0,00332202
<i>Irak1</i>	0,531	0,138692	-0,913216234
<i>Irak3</i>	0,5664	0,156333	-0,820106829
<i>Irf1</i>	0,52	0,146167	-0,943416472
<i>Irf2</i>	0,5421	0,120486	-0,883369088
<i>Irf3</i>	0,5285	0,122142	-0,920024623
<i>Irf5</i>	0,5212	0,092894	-0,940091011
<i>Irf8</i>	0,5797	0,142175	-0,786621609
<i>Jak1</i>	0,4988	0,23652	-1,00346663
<i>Ltbr</i>	0,6492	0,190703	-0,623265095
<i>Map2k3</i>	0,5371	0,080589	-0,896737373
<i>Mapkapk2</i>	0,4313	0,038549	-1,213236379
<i>Mavs</i>	0,5535	0,138558	-0,853344778
<i>Mmd2</i>	0,9977	0,912319	-0,00332202
<i>Mmp2</i>	0,9977	0,912319	-0,00332202
<i>Mmp25</i>	0,9977	0,912319	-0,00332202
<i>Mmp9</i>	0,0387	0,002466	-4,691522623
<i>Mtor</i>	0,5285	0,14745	-0,920024623
<i>Myd88</i>	0,5082	0,075377	-0,976531719
<i>Nfat5</i>	0,5757	0,186766	-0,796610883
<i>Nfix</i>	0,9977	0,912319	-0,00332202
<i>Nfkb1</i>	0,4475	0,058497	-1,160040413
<i>Nfkb2</i>	0,2558	0,014396	-1,966911831
<i>Nfkbia</i>	0,2398	0,030497	-2,060096436
<i>Nfkbib</i>	0,3415	0,027322	-1,550042516
<i>Nlrc5</i>	0,5837	0,055837	-0,776701027
<i>Nlrp12</i>	0,9977	0,912319	-0,00332202
<i>Otud7b</i>	0,6142	0,061027	-0,703219583
<i>Rel</i>	0,8179	0,45206	-0,290003631
<i>Relb</i>	0,4774	0,065263	-1,066729528
<i>Snf8</i>	0,577	0,112781	-0,793356776
<i>Sp1</i>	0,6085	0,174744	-0,716670832
<i>Sp110</i>	0,2248	0,022922	-2,153286059
<i>Stap2</i>	0,1466	0,033616	-2,770042991
<i>Stat1</i>	0,2845	0,038976	-1,813499442
<i>Stat6</i>	0,473	0,075463	-1,080087911
<i>Syk</i>	0,3439	0,026521	-1,539938979
<i>Tbk1</i>	0,4852	0,048226	-1,043348544
<i>Tlr4</i>	0,5561	0,111615	-0,846583758
<i>Tmem173</i>	0,6358	0,247556	-0,653355078
<i>Tnfrsf11a</i>	0,5023	0,038571	-0,99337882
<i>Tnfrsf12a</i>	0,4323	0,090839	-1,209895259

ECTV/Poly(I:C) vs Poly(I:C)			
Gene	Fold change	<i>p</i> value	Log ₂ fold change
<i>Tnfrsf13c</i>	0,9977	0,912319	-0,00332202
<i>Traf1</i>	0,2181	0,022437	-2,196938325
<i>Traf2</i>	0,4414	0,066839	-1,179841465
<i>Traf3</i>	0,7423	0,252861	-0,429925726
<i>Traf6</i>	0,7137	0,206748	-0,486610322
<i>Vegfb</i>	0,5046	0,064417	-0,986787889
<i>Xbp1</i>	0,4676	0,048256	-1,096653165
<i>Yy1</i>	0,6752	0,206195	-0,566613191
<i>Rn18s</i>	1	0	

The expression of the analyzed genes was evaluated using RT-qPCR. The cells were uninfected and or infected with ECTV and were left untreated or treated with poly(I:C) for 18 h. Significant changes ($p \leq 0.05$, log₂ fold change ≤ -1) in gene expression are shown in blue (downregulation). *Rn18s* – reference gene. Data were obtained from three independent experiments.

Supplementary Table 5. Expression of genes involved in NF- κ B signaling pathways in RAW 264.7 macrophages infected with ECTV and stimulated with IFN+LPS (18 hpi).

ECTV/IFN+LPS vs IFN+LPS			
Gene	Fold change	<i>p</i> value	Log ₂ fold change
<i>Aim2</i>	0,5409	0,017001	-0,886566197
<i>Atf3</i>	0,4741	0,010668	-1,076736702
<i>Bcl2</i>	0,2365	0,001898	-2,080087911
<i>Bcl3</i>	0,6227	0,037635	-0,683390816
<i>Birc2</i>	0,3618	0,006656	-1,466735687
<i>Birc3</i>	0,1839	0,004017	-2,443006615
<i>Ccl19</i>	1,0234	0,81205	0,033370138
<i>Ccl25</i>	0,6057	0,001724	-0,723324683
<i>Ccr10</i>	1,0234	0,81205	0,033370138
<i>Ccr8</i>	1,0234	0,81205	0,033370138
<i>Cd36</i>	0,1856	0,000045	-2,429731384
<i>Cd40</i>	0,1975	0,000941	-2,340075442
<i>Cd68</i>	0,1684	0,001443	-2,570035956
<i>Cd86</i>	0,1672	0,008808	-2,580353247
<i>Cflar</i>	0,4024	0,000294	-1,31329779
<i>Chuk</i>	0,3172	0,004748	-1,656535324
<i>Creb1</i>	0,2117	0,004114	-2,239906826
<i>Csf2</i>	0,0504	0,024532	-4,310432456
<i>Cxcl10</i>	0,0167	0,083843	-5,904008087
<i>Cxcl5</i>	1,0234	0,81205	0,033370138
<i>Cxcr5</i>	1,4473	0,191448	0,533363998
<i>Daxx</i>	0,1715	0,000065	-2,543719518
<i>Ddx58</i>	0,0357	0,000326	-4,807932116
<i>Ets2</i>	0,483	0,010502	-1,049904906
<i>Hdac9</i>	0,722	0,067664	-0,469929258
<i>Hmgb1</i>	0,4538	0,005938	-1,139871486
<i>Icam1</i>	1,4044	0,174043	0,489953901
<i>Ido1</i>	1,0234	0,81205	0,033370138
<i>Slfit1</i>	0,0164	0,003422	-5,930160375
<i>Ifnb1</i>	1,0234	0,81205	0,033370138
<i>Ifngr1</i>	0,3959	0,000768	-1,336792028
<i>Ikkb</i>	0,386	0,002503	-1,373327247
<i>Ikbke</i>	0,1966	0,001655	-2,346664773
<i>Ikkkg</i>	0,1627	0,006376	-2,619713844

ECTV/IFN+LPS vs IFN+LPS			
Gene	Fold change	p value	Log ₂ fold change
<i>Il15</i>	0,0994	0,003178	-3,330610338
<i>Il18</i>	0,1101	0,006546	-3,183113626
<i>Il9</i>	1,0234	0,81205	0,033370138
<i>Irak1</i>	0,3471	0,000436	-1,52657673
<i>Irak3</i>	0,1822	0,002975	-2,456405136
<i>Irf1</i>	0,2987	0,000473	-1,743230857
<i>Irf2</i>	0,2264	0,002904	-2,143054137
<i>Irf3</i>	0,2269	0,000139	-2,139871486
<i>Irf5</i>	0,3172	0,001794	-1,656535324
<i>Irf8</i>	0,1634	0,000034	-2,613520111
<i>Jak1</i>	0,2859	0,000175	-1,806417475
<i>Ltbr</i>	0,408	0,010363	-1,293358943
<i>Map2k3</i>	0,5522	0,161258	-0,856737207
<i>Mapkapk2</i>	0,2636	0,000016	-1,923577725
<i>Mavs</i>	0,25	0,000927	-2
<i>Mmd2</i>	1,0234	0,81205	0,033370138
<i>Mmp2</i>	1,0234	0,81205	0,033370138
<i>Mmp25</i>	0,2012	0,008294	-2,31329779
<i>Mmp9</i>	0,3353	0,024132	-1,576475612
<i>Mtor</i>	0,5129	0,003578	-0,963250524
<i>Myd88</i>	0,4434	0,009862	-1,173319325
<i>Nfat5</i>	0,2655	0,000185	-1,913216234
<i>Nfix</i>	1,0234	0,81205	0,033370138
<i>Nfkb1</i>	0,3754	0,006904	-1,413499445
<i>Nfkb2</i>	0,1525	0,002281	-2,713118852
<i>Nfkbia</i>	0,0838	0,000774	-3,576905946
<i>Nfkbib</i>	0,1917	0,000055	-2,383077758
<i>Nlrc5</i>	0,15	0,000383	-2,736965594
<i>Nlrp12</i>	1,0234	0,81205	0,033370138
<i>Otud7b</i>	0,7153	0,15712	-0,483379653
<i>Rel</i>	0,3643	0,008452	-1,4568011
<i>Relb</i>	0,6522	0,129553	-0,616613654
<i>Snf8</i>	0,3345	0,019971	-1,579921884
<i>Sp1</i>	0,3987	0,001931	-1,32662449
<i>Sp110</i>	0,1568	0,000013	-2,673002535
<i>Stap2</i>	0,1116	0,000085	-3,163591068
<i>Stat1</i>	0,0273	0,000541	-5,194955239
<i>Stat6</i>	0,3322	0,000669	-1,589876022
<i>Syk</i>	0,198	0,001391	-2,336427665
<i>Tbk1</i>	0,3106	0,000991	-1,686870265
<i>Tlr4</i>	0,2349	0,00018	-2,089881382
<i>Tmem173</i>	0,4414	0,024423	-1,179841465
<i>Tnfrsf11a</i>	0,1805	0,00039	-2,469929258
<i>Tnfrsf12a</i>	0,6001	0,040125	-0,736725165
<i>Tnfrsf13c</i>	1,0234	0,81205	0,033370138
<i>Traf1</i>	0,0921	0,000318	-3,440655033
<i>Traf2</i>	0,2932	0,000177	-1,770042991
<i>Traf3</i>	0,5141	0,006526	-0,959879083
<i>Traf6</i>	0,5	0,001708	-1
<i>Vegfb</i>	0,3577	0,008124	-1,483177977
<i>Xbp1</i>	0,1642	0,001041	-2,606473968
<i>Yy1</i>	0,6814	0,087199	-0,553426147
<i>Rn18s</i>	1	0	

The expression of the analyzed genes was evaluated using RT-qPCR. The cells were uninfected or infected with ECTV and were left untreated or treated with rmIFN- γ + *Escherichia coli* LPS O111:B4 for 18 h. Significant changes ($p \leq 0.05$, \log_2 fold change ≤ -1) in gene expression are shown in blue (downregulation). *Rn18s* – reference gene. Data were obtained from three independent experiments.