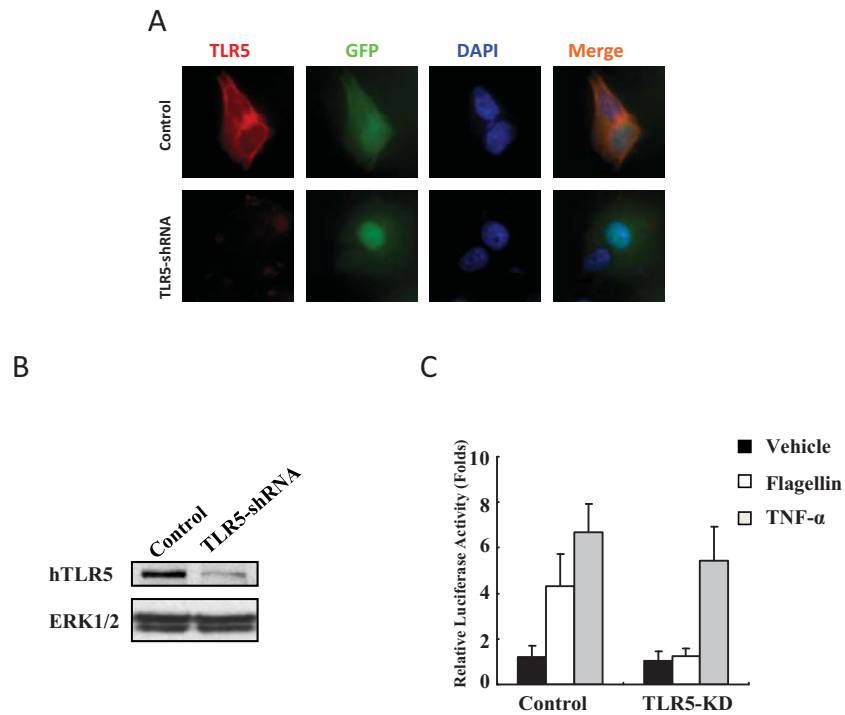
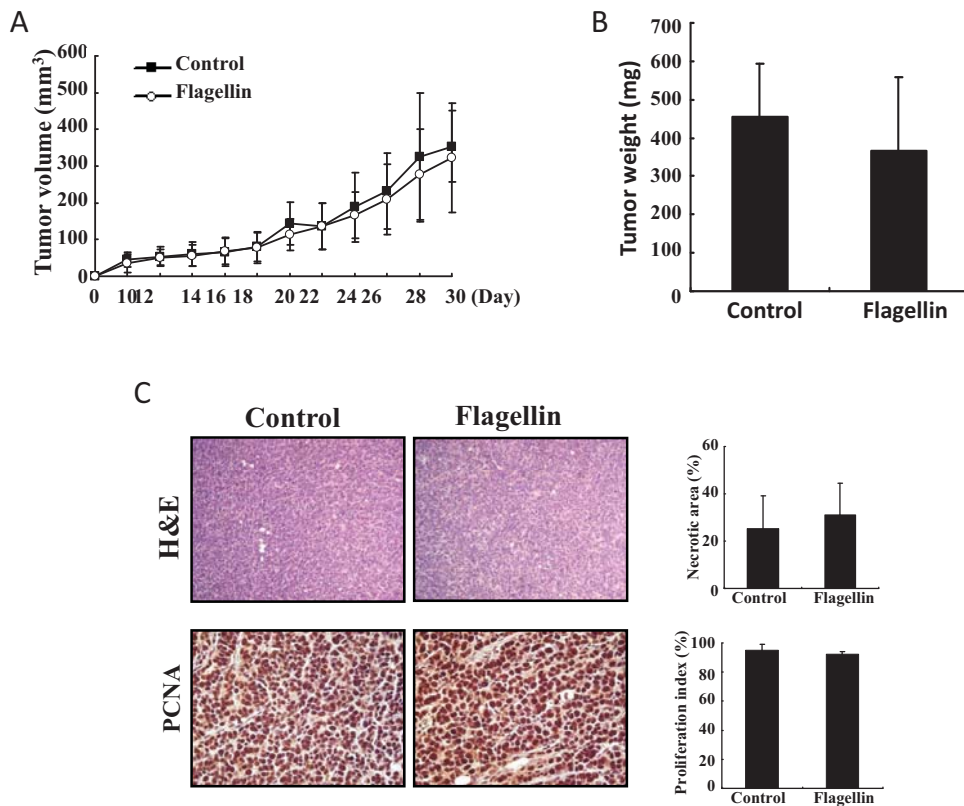


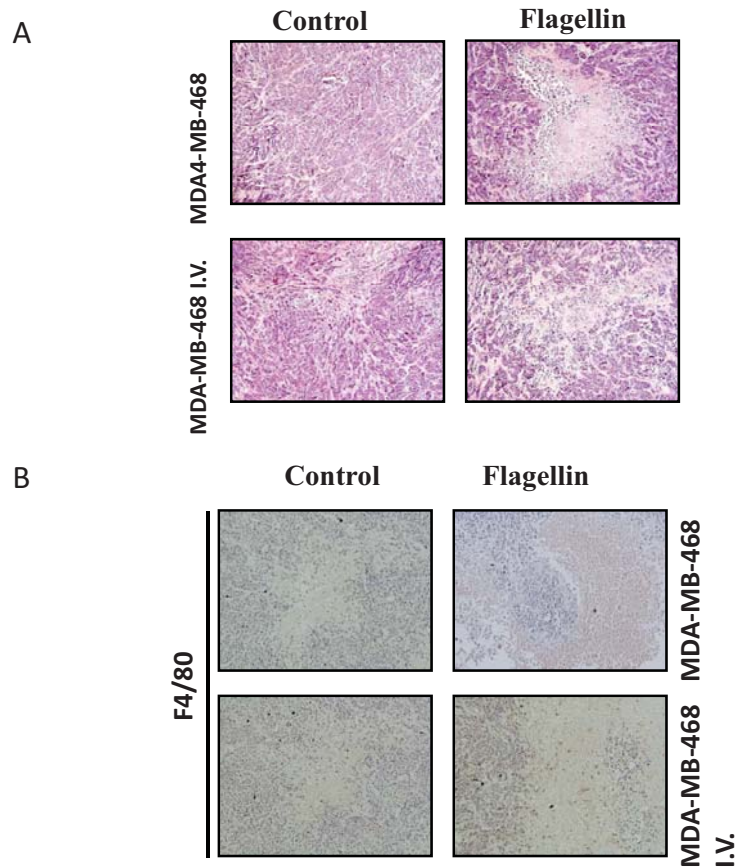
**Supplementary Figure 1.** Expression of TLR5 and MyD88 in breast cancer cells. With western blotting analysis, the expression of TLR5 and its adaptor MyD88 was detected in MCF-7, MDA-MB-468, SKBR3, T47D, MDA-MB-231 and MDA-MB-435 cells.



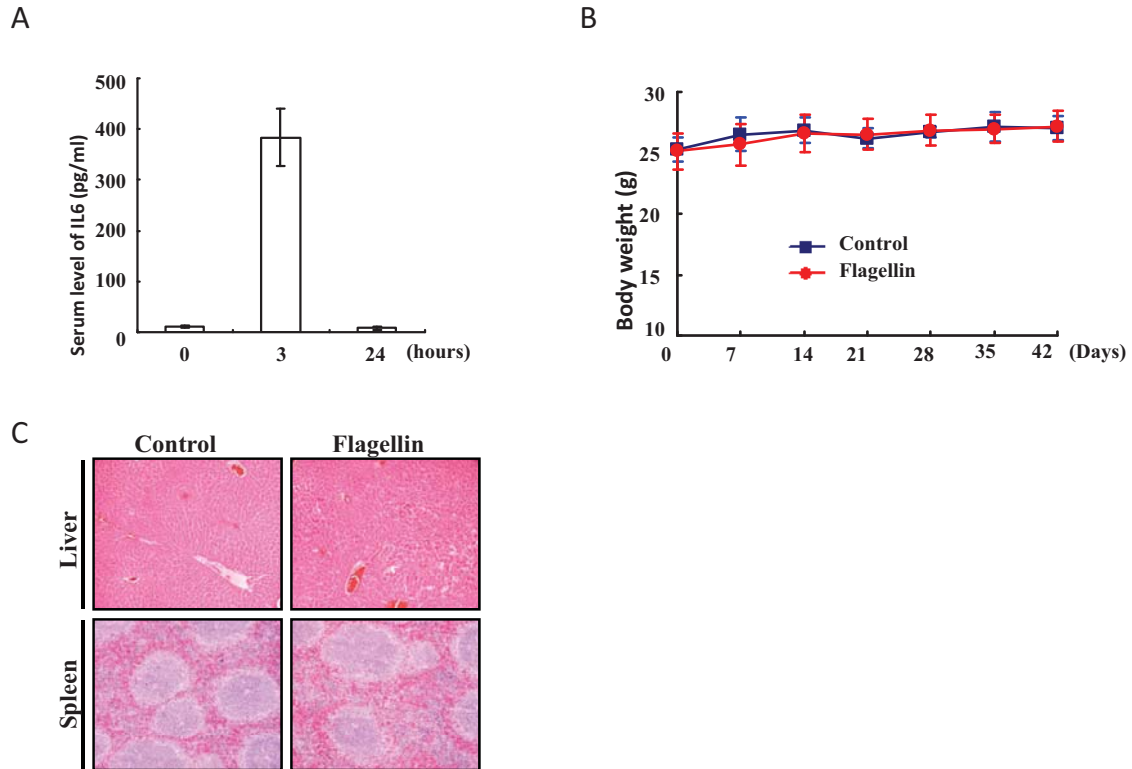
**Supplementary Figure 2.** Generating TLR5 shRNA knocked down cells. A, Detection of TLR5 expression in MCF-7 control and knocked down cells by immunofluorescence with the same exposure time. Because these vectors encode a GFP protein, stably transfected cells were identified by GFP expression. B, TLR5 protein expression was examined by Western blot in MCF-7 control and knocked down cells. And total ERK1/2 level as a loading control. C, MCF-7 knocked down cells and scramble control were transfected with NF $\kappa$ B reporter plasmid and analyzed for luciferase activity induced by flagellin (0.1  $\mu$ g/ml) and TNF- $\alpha$  (100U/ml).



**Supplementary Figure 3.** Flagellin treatment did not alter the growth of breast cancer xenograft mouse model generated from MCF-7- TLR5-shRNA cells. A, tumor growth curves of control. B, tumor weight of control or flagellin-treated groups were measured. C H&E staining of tumor sections (*Top panel*). The percentage necrosis area was calculated from five sections per tumor. Mean  $\pm$  s.d. values of data from five tumors are shown (*Top right*). Cell proliferation marker PCNA staining of tumor sections (*Bottom panel*). The ratio of PCNA-positive to total tumor cells was calculated from five sections per tumor. Mean  $\pm$  s.d. values of data from five tumors are shown (*Bottom right*). Magnification: H&E, 10X; PCNA, 40X.



**Supplementary Figure 4.** Histology study of tumor sections from MDA-MB-468 xenografts. A, H&E staining of MDA-MB-468 xenograft tumor sections showed more prevalent necrosis and leucocytes infiltration in tumors either by peritumoral and i.v. flagellin treatment. Magnification: 10X. B, Tumor sections from MDA-MB-468 xenografts were stained by neutrophil-specific marker Gr-1. Representative images from tumor sections with flagellin peritumoral or i.v. treatment are shown. Red indicates positive staining. Magnification: 10X.



**Supplementary Figure 5.** Toxicity assessment following intravenous delivery of flagellin in nude mice. A, Serum level of IL-6 in mice (n=6) following flagellin (2  $\mu$ g per mouse) administration at baseline (0 hour), 3 and 24 hours. B, Body weight curves of mice in control and flagellin-treated groups during the period of i.v. injection. C, H&E staining of liver and spleen sections in control and flagellin-treated groups. Magnification: 10X.

**Supplementary Table 1. Clinicopathologic features of TLR5 in patients with breast cancer \***

	TLR5	
	Positive	Negative
	N=60 (80% )	N=15 (20% )
<b>Age</b>		
< 50	32 (42.7%)	11(14.7%)
≥ 50	28 (37.3%)	4 (5.3%)
<b>Tumor type</b>		
Invasive ductal carcinoma	53(70.7%)	8(10.7%)
Ductal carcinoma <i>in situ</i>	1 (1.3%)	1(1.3%)
Medullary carcinoma	2 (2.7%)	0
Mucinous carcinoma	0	3(4%)
Invasive lobular carcinoma	1(1.3%)	2(2.7%)
Mixed lobular and ductal carcinoma	1(1.3%)	0
Ductal papillary carcinoma	1(1.3%)	0
Invasive papillary carcinoma	1(1.3%)	0
Neuroendocrine carcinoma	0	1(1.3%)
<b>Tumor grade</b>		
I	2(2.7%)	0
I- II	2(2.7%)	3(4%)
II	12(16%)	5(6.7%)
II- III	27(36%)	6(8%)
III	17(22.7%)	1(1.3%)
<b>Tumor size</b>		
T1	8(10.7%)	0
T2	36(48%)	15(20%)
T3	12(16%)	0
T4	3(4%)	0
Tis (Carcinoma <i>in situ</i> )	1(1.3%)	0
<b>Lymph node metastasis</b>		
N0	42(56%)	14(18.7%)
N1	16(21.3%)	1(1.3%)
N2	3(4%)	0
<b>Distant metastasis</b>		
M0	58(77.3%)	15(20%)
M1	2(2.7%)	0

\* One adenosquamous carcinoma sample and one squamous cell carcinoma sample were not counted in this study.

**Supplemental Table 2 Primers used in this study**

<b>Oligonucleotide</b>	<b>Sequence (5'→3')</b>	<b>Purpose</b>
Human IL-1 $\beta$ -F	GACACATGGGATAACGAGGC	RT-PCR
Human IL-1 $\beta$ -R	ACGCAGGACAGGTACAGATT	RT-PCR
Human IL-6-F	GGAGACTTGCCTGGTGAA	RT-PCR
Human IL-6-R	GCATTTGTGGTTGGGTCA	RT-PCR
Human IL-8-F	TTGGCAGCCTTCCTGATTTC	RT-PCR
Human IL-8-R	AACTTCTCCACAACCCTCTG	RT-PCR
Human IL-12-F	TGGAGTGCCAGGAGGACAGT	RT-PCR
Human IL-12-R	GATGATGTCCCTGATGAAGAAGC	RT-PCR
Human TNF- $\alpha$ -F	CGAGTGACAAGCCTGTAGC	RT-PCR
Human TNF- $\alpha$ -R	GGTGTGGGTGAGGAGCACAT	RT-PCR
Human $\beta$ -actin F	TCCCTGGAGAAGAGCTACG	RT-PCR
Human $\beta$ -actin R	GTAGTTTCGTGGATGCCACA	RT-PCR