

CD36 regulates lipopolysaccharide-induced signaling pathways and mediates the internalization of *Escherichia coli* in cooperation with TLR4 in goat mammary gland epithelial cells

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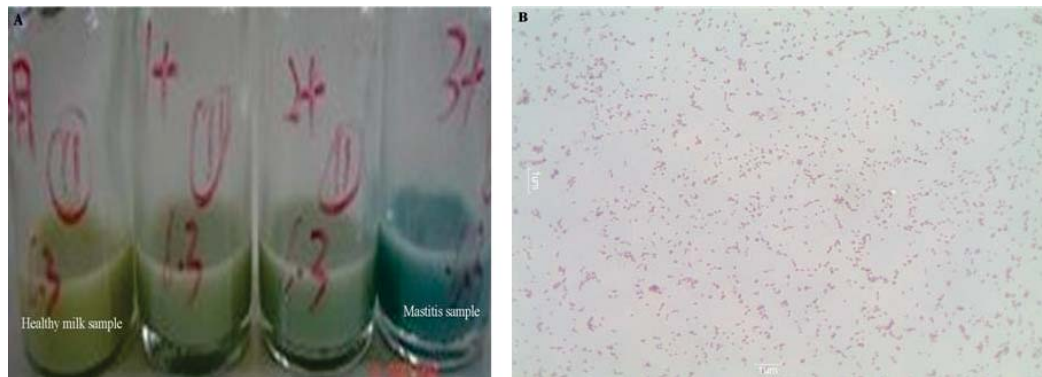
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Supplementary Figure S1 (A) Milk samples were collected and tested for mastitis using LMT (LanZhou Mastitis Test, China). (B) Gram stain for identification of *Escherichia coli* from clinical mastitis.

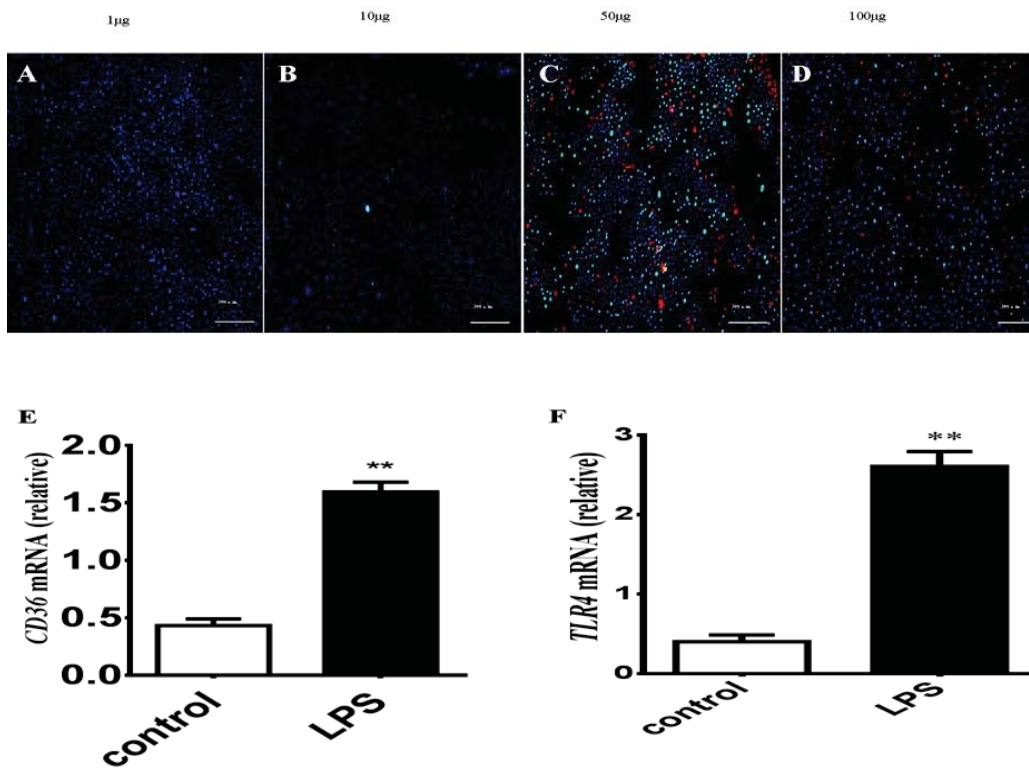


Supplementary Table S1 Biochemical identification of *Escherichia coli*

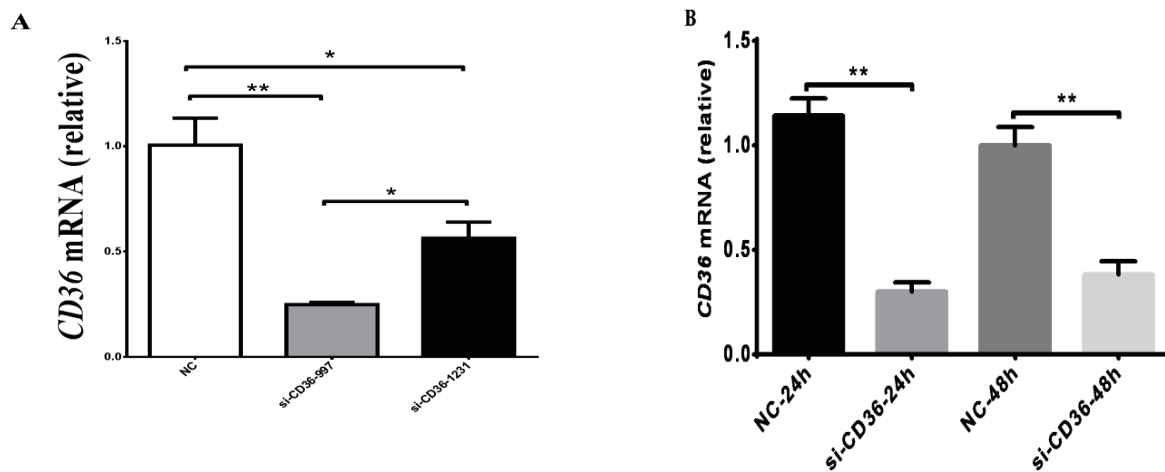
	MacConkey medium	Eosin and methylene blue medium	Triple sugar iron medium		Glucose		Lactose		Maltose		Sucrose	
			acid production	aerogenesis	acid production	aerogenesis	acid production	aerogenesis	acid production	aerogenesis	acid production	aerogenesis
1	+	+	+	+	+	+	+	+	+	+	-	-
2	+	+	+	+	+	+	+	+	+	+	-	-
3	+	+	+	+	+	+	+	+	+	+	-	-

Supplementary Figure S2 (A–D) Incubation of cells in different concentrations of LPS (1, 10, 50, and 100 µg/ml) for 12 hours. Cells were stained with Hoechst33342 and PI to determine whether they induced apoptosis or necrosis. (E, F) Cells were incubated with LPS for 12

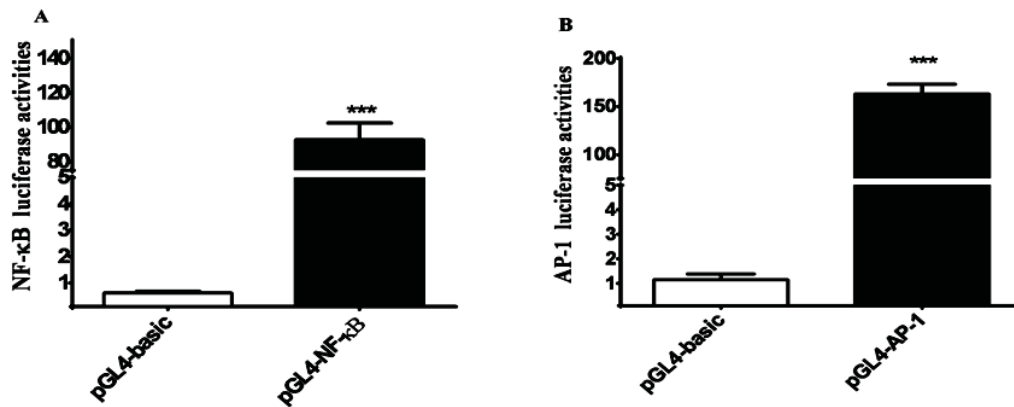
hours, and the relative mRNA levels of CD36 and TLR4 increased after stimulation. Quantitative PCR data were normalized to GAPDH, UXT, and MRPL39. Data are means \pm SEM from three experiments. $**P < 0.01$, and not significant (NS).



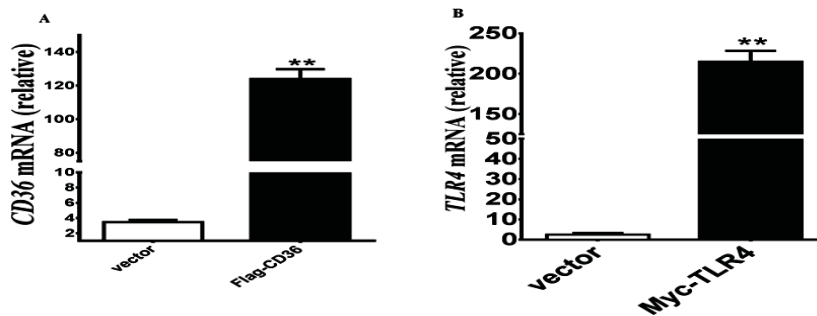
Supplementary Figure S3. Small interference RNA of CD36 in pGMECs efficiency test (A) The si-CD36-977 and si-CD36-1231 interference efficiency in pGMECs for 24 hours. (B) Changes of CD36 mRNA expression were influenced by knockdown CD36 expression in pGMECs for 24 and 48 hours. Quantitative PCR data were normalized to GAPDH, UXT, and MRPL39. Data are means \pm SEM from three experiments. $**P < 0.01$, and not significant (NS).

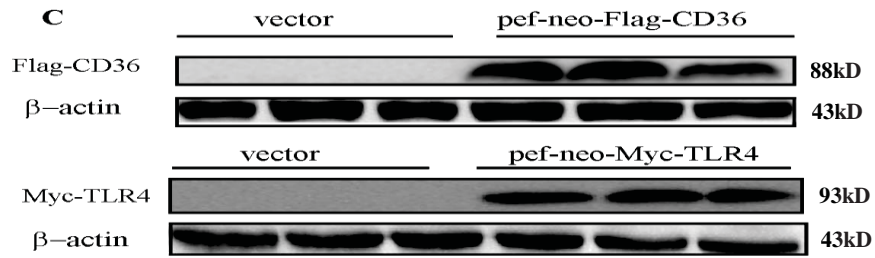


Supplementary Figure S4 Identification of the luciferase activity of pGL4-NF- κ B-RE (A) and pGL4-AP-1-RE (B) vector constructs in pGMECs. Luciferase activity is compared with pGL4 luciferase vectors. Data are means \pm SEM from three experiments. *** $P < 0.001$.



Supplementary Figure S5 The efficiency of pef-NEO-FLAG-CD36 and pef-NEO-FLAG-CD36 were cotransfected in pGMECs. CD36 (A) and TLR4 (B) mRNA levels increased significantly compared with vector groups. (C) The anti-Myc and anti-Flag were used for detection Myc-TLR4 and Flag-CD36 protein levels in pGMECs. Quantitative PCR data were normalized to GAPDH, UXT, and MRPL39. Data are means \pm SEM from three experiments (** $P < 0.01$). Immunoblotting was used to test the fusion protein (Flag-CD36 and Myc-TLR4) expressions.





Supplement Table S2. Quantitative PCR Primers

q-PCR Primer	Sequence
GAPDH-F	GCAAGTCCACGGCACAG
GAPDH-R	GGTTCACGCCATCACA
MRPL39-F	AGGTTCTCTTTGTGGCATCC
MRPL39-R	TTGGTCAGAGCCCCAGAAGT
UXT-F	TGTGGCCCTGGATATGGTT
UXT-R	GGTTGTCGCTGAGCTCTGTG
IL6-F	AGATATACCTGGACTTCCT
IL6-R	TGTTCTGATACTGCTCTG
TLR4-F	AGATGGCAACACTAGAA
TLR4-R	ATACTGAAGGCTGGTAG
TNfa-F	TGGTTCAGACACTCAGGT
TNfa-R	CGCTGATGTTGGCTACAA
MYD88-F	CGGATGGTGGTGTGTCT
MYD88-R	GGAACCTCTTCTTCATTGGCTTGT
IL-1 β -F	GCGACTCACTTTCACCTTTTCACTT
IL-1 β -R	TGTGTGACCCCATAGACGGTAG
IL-8-F	AAGCTGGCTGTTGCTCTCTTG
IL-8-R	GGGTGGAAAGGTGTGGAATG
NF-Kb-F	TGGCAGCTCTTCTCAAAGCA
NF-Kb-R	GACCCCTTCATCTCTCCATC
TGF- β -F	TTCCGTGGGATACCGAGA
TGF- β -R	CTGTTGCGGGAGAGTTG
CD36-F	CGCCATAATTGACACATACA
CD36-R	CTCCGAACACAGCATAGA