

Bidirectional effect of Wnt signaling antagonist DKK1 on the modulation of anthrax toxin uptake

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Received March 4, 2014; accepted March 7, 2014; published online March 25, 2014

LRP6, a co-receptor for the morphogen Wnt, aids endocytosis of anthrax complexes. Here we report that Dickkopf1 (DKK1) protein, a secreted LRP6 ligand and antagonist, is also a modulator of anthrax toxin sensitivity. shRNA-mediated gene silencing or TALEN-mediated gene knockout of *DKK1* reduced sensitivity of cells to PA-dependent hybrid toxins. However, unlike the solely inhibitory effect on Wnt signaling, the effects of DKK1 overexpression on anthrax toxicity were bidirectional, depending on its endogenous expression and cell context. Fluorescence microscopy and biochemical analyses showed that DKK1 facilitates internalization of anthrax toxins and their receptors, an event mediated by DKK1-LRP6-Kremen2 complex. Monoclonal antibodies against DKK1 provided dose-dependent protection to macrophages from killing by anthrax lethal toxin (LT). Our discovery that DKK1 forms ternary structure with LRP6 and Kremen2 in promoting PA-mediated toxin internalization provides a paradigm for bacterial exploitation of mechanisms that host cells use to internalize signaling proteins.

DKK1, anthrax toxin, LRP6, TALENs, internalization, Kremen2, receptor, Wnt

Citation: Qian LL, Cai CZ, Yuan PF, Jeong SY, Yang XZ, DeAlmeida V, Ernst J, Costa M, Cohen SN, Wei WS. Bidirectional effect of Wnt signaling antagonist DKK1 on the modulation of anthrax toxin uptake. *Sci China Life Sci*, 2014, 57: 469–481, doi: 10.1007/s11427-014-4646-x

Anthrax is a lethal disease caused by the combined actions of three polypeptide toxins secreted by the gram-positive bacterium, *Bacillus anthracis*: protective antigen (PA, 83 kD), lethal factor (LF, 90 kD), and edema factor (EF, 89 kD) [1,2]. None of these proteins is individually toxic to the host, but bipartite complexes consisting of PA and LF or PA and EF, which form lethal toxin or edema toxin, respectively, initiate events that result in anthrax disease. LF is a zinc-dependent metalloproteinase that blocks the ERK, p38

and Jun N terminus kinase (JNK) pathways [3] by cleaving and inactivating mitogen-activated protein kinase kinases (MEKs) 1–4, 6, and 7 [4–6]. EF is calmodulin-dependent adenylate cyclase that causes rapid accumulation of cAMP, leading to multiorgan failure [1,7]. Entry of EF and LF into mammalian cells requires interaction of these toxic moieties with PA, which attaches them to receptor proteins on the surface of targeted cells: tumor endothelium marker-8 (TEM8; anthrax toxin receptor 1 (ANTXR1)), capillary morphogenesis protein-2 (CMG2; anthrax toxin receptor 2 (ANTXR2)), and integrin β 1 complexes [8–10]. CMG2 is

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the principal ligand for PA in rodents [11]. As LF lethality requires dependence of targeted cells on MAPKK [5,12], which normally is observed for macrophages but not for most other cell types, FP59, a PA-binding fusion protein containing LF amino acids 1–254 and the catalytic domain of *Pseudomonas aeruginosa* exotoxin A commonly is employed to investigate the binding and internalization actions of PA [13,14]. For similar reasons, LFnDTA (a fusion of the PA binding domain of LF to the enzymic A chain of diphtheria toxin) [15] has also been used as a surrogate for LF. At least two other host-encoded cell proteins, low density lipoprotein receptor like protein 6 (LRP6), which has been widely studied as a co-receptor for the morphogen Wnt [16–18], and ARAP3, a GTPase implicated in PI3K signaling [19], do not directly bind to PA but can promote endocytosis of PA-complexes [19–21].

DKK1 (Dickkopf-1) is a secreted, high-affinity ligand for LRP6 that antagonizes LRP6-mediated Wnt signaling by two distinct mechanisms: (i) by interacting with LRP6, DKK1 interrupts Wnt-induced binding of LRP6 to Frizzled (Fz), another LRP6 ligand implicated in Wnt signaling, interfering with signal transduction without affecting LRP6 internalization or degradation [22–24]; and (ii) DKK1 forms complexes with LRP6 and Kremen1 or Kremen2—transmembrane proteins that trigger internalization of LRP6 and reduce LRP6 availability at cell surface for Wnt signaling [25,26]. It has been proposed recently that DKK1 can also inhibit Wnt signaling by blocking Wnt binding to LRP6 [27,28]. Here we report that DKK1 is a modulator of PA-mediated anthrax toxin endocytosis, and that anthrax toxin hijacks a ternary complex consisting of DKK1, LRP6 and Kremen2 to assist with its endocytosis, further linking the developmentally-important Wnt signaling pathway to anthrax toxin internalization.

1 Materials and methods

1.1 Expression constructs

DKK1 genes were obtained through PCR-amplification from human or mouse cDNA, and cloned into pIRES-hyg2 or pEF6MycHisBsd-B-based vectors for expression.

1.2 Mammalian cell culture, transfection and lentivirus infection

The human prostate cancer cell line M2182 was maintained in RPMI 1640 medium (Invitrogen, USA) by using supplements as described in [29]. RAW264.7 mouse macrophage, HEK293T, HeLa and BHK cell lines were cultured in DMEM (Invitrogen, USA) containing 10% FBS. DNA transfections were performed with Polyethylenimine (PEI)-mediated method [30], FuGene6 (Roche, Switzerland) or Lipofectamin 2000 (Invitrogen, USA) according to the manufacturer's recommended protocols. Lentivirus were

produced by transient transfection of HEK293T cells (PEI-mediated methods) by using DNAs along with packaging and VSVG envelope constructs as described [31]. Details are the same as reported [20].

1.3 Preparation of *hDKK1*-specific shRNAs

Expression constructs carrying lentiviral shRNAs specifically targeting on human *DKK1* gene were obtained from Openbiosystems, Inc. Stable shRNA expression in the vector of pLKO.1 was achieved in M2182 cells through viral infection followed by puromycin ($1 \mu\text{g mL}^{-1}$) selection. Total of five different *DKK1*-specific shRNAs were used. Their clone IDs and targeting sequences are TRCN-0000033384 (5'-CGGGAATAAGTACCAGACCAT), TRCN-0000033385 (5'-CGGTTCTCAATTCCAACGCTA), TRCN-00000333856 (5'-CCTGTCCTGAAAGAAGGTCAA), TRCN0000033387 (5'-GCCAGTAATTCTTCTAGGCTT), and TRCN0000033388 (5'-CCAGAAGAACCACCTTGTCTT), respectively.

1.4 Antibodies

Full-length His-tagged human DKK1 protein was expressed and purified from baculovirus cell supernatants, and antibodies were raised against the protein using standard hybridoma technology. Hybridoma clone supernatants were screened by ELISA and Western analysis, and seven positive clones were identified for monoclonal selection and antibody purification. The monoclonal antibody against PA was obtained from Novus Biologicals (NB 600-407) (USA). Agarose-conjugated monoclonal antibodies of c-Myc (9E10), HA (F-7) were obtained from Santa Cruz Biotechnology (USA). Peroxidase-conjugated mouse monoclonal antibody (clone 9E10) against c-Myc (11814150001) and peroxidase-conjugated mouse monoclonal antibody (clone 12CA5) against HA (11667475001) were ordered from Roche (China) Ltd. Mouse anti-tubulin mAb and horseradish peroxidase-conjugated secondary antibodies were purchased from Santa Cruz Biotechnology (USA).

1.5 Immunofluorescence microscopy

Chimeric LF-EGFP protein purification was the same as reported [32]. To monitor cell surface binding, cells were first treated with PA83 (600 nmol L^{-1}) for 12 min at 37°C , washed, incubated with LF-EGFP (200 nmol L^{-1}) for 2 min at 37°C and immediately washed and fixed before examined by confocal microscopy. To monitor PA/LF-EGFP internalization, cells were incubated with PA83 (600 nmol L^{-1}) and LF-EGFP (200 nmol L^{-1}) for 15 min at 37°C , followed by confocal microscopy. Immunostaining was performed according to a standard protocol [33]. The cover slips were mounted onto slides with a mounting medium containing DAPI (Vector Laboratories, USA), and the cells were

examined by using an LSM 710&NLO confocal microscope (Zeiss, Germany). The detailed procedure was the same as reported [32].

1.6 Cytotoxicity assay

PA and LF were purchased from List Biological Laboratories, Inc., and FP59 was produced as described previously [34]. LFnDTA, a surrogate of LF consisting of the N-terminal domain of LF fused to the catalytic subunit of diphtheria toxin, was produced by using plasmid pET-15b-LFNDA (Addgene plasmid 11075) [35]. Cytotoxicity assays for M2182, HeLa, HEK293T, and BHK were performed by seeding the cells in 96-well plates (100 μL /well) at a concentration of 5×10^4 cells mL^{-1} one day prior to the toxin treatment; the cell concentration used was 1×10^5 cells mL^{-1} for RAW264.7 cell line. Various concentrations of PA combined with a fixed concentration of FP59 (50 ng mL^{-1}) or LF (500 ng mL^{-1} for RAW264.7 line), or LFnDTA (50 ng mL^{-1}) was added to wells, and cells were incubated at 37°C for 48 or 18 h (for RAW264.7 line). Cell viability was measured by MTT assay as described [20].

1.7 Western blotting and co-immunoprecipitation

Western blotting was performed essentially as described by Harlow and Lane [33] using 10% or 3%–8% gels (NuPAGE Novex Tris-Acetate Gels, EA0378BOX, Invitrogen, USA) for electrophoresis and standard conditions. The soluble cell proteins were quantitated using BCA protein assay kit (Pierce Biotechnology, USA). For co-immunoprecipitation assays, HEK293T cells, 48 h after transfection, were lysed in NP40 buffer containing a protease inhibitor cocktail (Complete Mini, Roche China, Ltd.). The lysates were centrifuged at $16000 \times g$, and the soluble fraction was immunoprecipitated with the desired antibody-conjugated agarose beads (Santa Cruz Biotechnology, USA).

1.8 Biochemical assay of PA binding and internalization

Methods were as described [20]. The concentration and its incubation time for binding assay were 1 $\mu\text{g mL}^{-1}$ and 1 h, respectively.

1.9 Construction of stable knockout cell lines using TALEN technique

The design and assembly of the two pairs of TALENs constructs used for *DKK1* gene-knockout were based on our own ULtiMATE protocol [36]. More specifically, the two targeting sequences for *DKK1* loci are 5'-TCCAACGCT-ATCAAG-3' for TALEN^L and 5'-GCCCCGCAGCGCCGC-3' for TALEN^R, with a spacer sequence (5'-AACCTGCC-

CCACCGCTGG-3'). The identification and verification of gene knockout events were based on both sequencing analysis of genome PCR fragments of targeting loci and functional assay of Wnt signaling using a beta-catenin-regulated promoter on the Super8XTOPflash reporter plasmid [37].

2 Results

2.1 Effect of *DKK1* knockdown on PA-mediated toxicity

Earlier work has shown that reduced activity of LRP6 gene mediated by either genetic or immunological interference leads to a decrease in the cell killing by anthrax toxin and also by a surrogate toxin that is dependent on PA for endocytosis [20,21]. LRP6 was found to facilitate internalization of PA and toxin moieties bound to it in the human prostate cancer cell line M2182 and in mouse macrophage RAW264.7 cells [20]. As interference with LRP6 function decreases toxin lethality, we wished to learn whether inhibition of LRP6 function by its biological antagonist *DKK1* has the converse effect. To elucidate the role of *DKK1* in PA-mediated toxicity, five TRC lentiviral borne shRNAs specifically targeting the human *DKK1* gene (Open Biosystems, USA) were introduced individually into M2182 cells by viral infection. The effects of one such shRNA (TRCN0000033384) that corresponds to an mRNA sequence (5'-CGGGAAUAAGUACCAGACCAU) of the human *DKK1* gene are shown in Figure 1 for two randomly selected clones, shRNA384-2 and -6. Stable expression of the shRNA under control of the hU6 promoter resulted in both a decrease in abundance of *DKK1* mRNA (Figure 1A) in M2182 cells and a decrease in *DKK1* protein in the culture media (Figure 1B) to 30% and 50% of normal for the M2182/shRNA384-6 and M2182/shRNA384-2 clones, respectively. Associated with the observed decreased abundance of *DKK1* mRNA and protein, was sharply increased expression of a TCF/beta-catenin-mediated reporter gene as assayed using a beta-catenin-regulated promoter on the Super8XTOPflash reporter plasmid [37] (Figure 1C), consistent with the known inhibitory role of *DKK1* in Wnt signaling [38]. shRNA (TRCN0000033387) that corresponds to an mRNA sequence (5'-GCCAGUAAUUCUUCUAGGCUU) of the human *DKK1* gene had little effect on the abundance of *DKK1* mRNA (Figure 1A) or *DKK1* protein (Figure 1B), and consequently had no effect on the Wnt signaling (Figure 1C). Data for one such clone (M2182/shRNA387-5) are included in this and other figures as a negative control.

Lethality mediated by PA+FP59 in M2182 clones shNRA384-2 and -6 was assessed by MTT assay [20] using a series of PA concentrations and a fixed concentration of FP59 (50 ng mL^{-1}). Paradoxically, given the observed

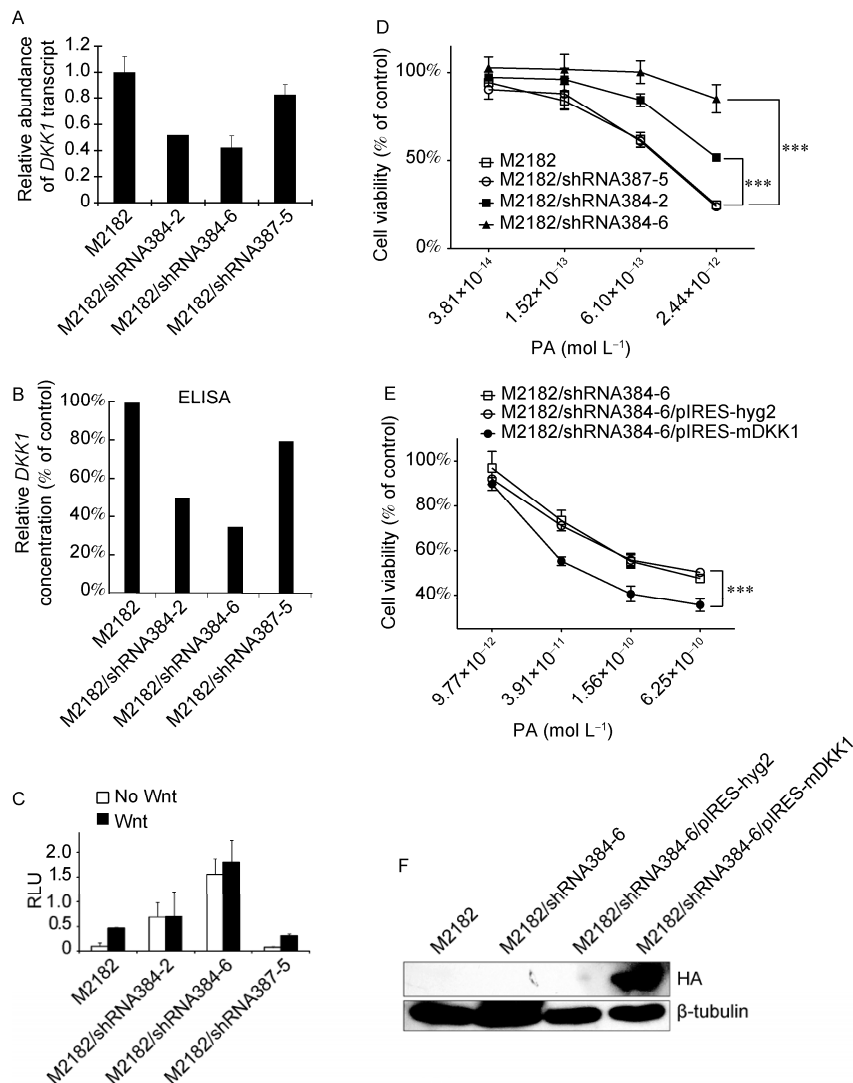


Figure 1 Effect of human *DKK1* down-regulation on M2182 to PA/FP59. TRC lentiviral shRNA (TRCN0000033384) specifically targeting on human *DKK1* gene was introduced into cells through viral infection. Two such clones (shRNA384-2 and -6) from M2182 were assayed for their cytotoxicity and associated gene expression level. Lentiviral shRNA (TRCN0000033387) was used as a control. A, Effect of *DKK1*-specific shRNA on transcript abundance. *DKK1* transcripts were measured by quantitative RT-PCR and normalized to 18S rRNA. The primers used to amplify a region of *hDKK1* mRNA and 18S rRNA were (5'-CCGAGGAGAAATTGAGGAAA and 5'-CCGAGAGCAAACAGAACCTT) and (5'-TTTGACTCAACACGGGAAAC and 5'-ATCGCTCCACCAACTAAGAAC), respectively. Data are mean±SD, $n=3$. B, Effect of *DKK1*-specific shRNA on secreted *DKK1* protein abundance. *DKK1* protein concentration was measured by Assay Designs' human *DKK1* TiterZyme Immunometric Assay (ELA) kit (Assay Design, USA). C, Effect of *DKK1* deficiency on cellular response to Wnt activation in M2182/shRNA384-2 and -6. Wnt3a-stimulated Wnt signaling was assayed in cells carrying Super8XTOPflash reporter plasmid and pcDNA3.1-lacZ plasmid for normalization through transient transfection [20]. Data are mean±SD, $n=3$. D, Cytotoxicity assay of M2182/shRNA384-2 and -6 to PA/FP59. Data are mean±SD, $n=6$; ***, $P<0.005$. E, Cytotoxicity assay for the effect of exogenous expression of mDkk1 in M2182/shRNA384-6 to PA/FP59; mean±SD, $n=6$; ***, $P<0.005$. F, Western blot analysis of exogenous expressed mouse *DKK1* in M2182/shRNA384-6 was conducted as described [20]. The cells carrying corresponding cloning vectors were used as control. β -tubulin abundance was measured as an internal control

effects of interference with LRP6 activity, we observed that an shRNA-induced decrease in *DKK1* expression was associated with increased cell survival in the presence of toxin (Figure 1D). Reversal of increased cell survival by shRNA directed against human *DKK1* in M2182 cells was achieved by adventitious expression of mouse *DKK1* in these cells (Figure 1E and F) (clone shRNA384-6), confirming that the effect is specifically a consequence of *DKK1* deficiency.

2.2 Effect of *DKK1* knockout on PA-mediated toxicity

To further determine the role of *DKK1* in PA-mediated toxicity, *DKK1* gene knockout clones were obtained in both HEK293T and M2182 cells using the TALEN technique [36,39], which enables the introduction of double strand DNA breaks at specific chromosomal sites. Analysis of the sequence of the TALEN-targeted genome region confirmed that the selected HEK293T/*DKK1*^{-/-} clone contained at

least three types of indels (insertions or deletions) that caused a frame shift in the coding sequence of *DKK1* (Figure 2A and B). Similarly, the M2182/*DKK1*^{-/-} clone was confirmed to have identical 17-nt deletion in *DKK1* alleles that completely disrupted gene expression (Figure 2C). The loss of *DKK1* gene expression in both HEK293T and M2182 cells was functionally confirmed by the elevated basal or induced level of Wnt signaling in these clones (Figure 2D and F). Importantly, the absence of *DKK1* expression in both lines was associated with less sensitivity to PA/LFnDTA (Figure 2E and G), and exogenous expression of *mDKK1*, but not of the mock control (cells transfected with only the expression vector), restored normal toxin sensitivity to the HEK293T/*DKK1*^{-/-} clone (Figure 2D–E). Our gene knockout data confirm that *DKK1* deficiency in HEK293T and M2182 cells results in decreased PA-mediated toxin lethality.

2.3 Bidirectional effect of *DKK1* expression on cell susceptibility to PA/FP59 or PA/LFnDTA

Collectively, the above results and those reported previously [20,21] indicate that the actions of both LRP6 and its antagonist *DKK1* are necessary for normal PA-mediated toxicity in HEK293T and M2182 cells. To better understand the mechanism(s) underlying such conjoint regulation by Wnt signaling proteins, we examined the effect of adventitious expression of *DKK1* on naïve cells on PA-mediated toxicity. By introducing a pIRES-hyg2 based construct expressing mouse *DKK1* into the M2182 cell line and selecting a stably transfected pooled population using an appropriate antibiotic marker (500 µg mL⁻¹ of hygromycin), we tested the effects of such overexpression on both Wnt signaling and PA-mediated toxicity. As seen in Figure 3A, we found by measuring beta-catenin activity that the Wnt signaling pathway is constitutively active in M2182 cells, consistent with the published effects of *DKK1* on Wnt signaling mediated through its known antagonism of LRP6 [25]. Adventitious *DKK1* expression in M2182 cells also reduced sensitivity to PA+FP59, enabling M2182 cells to survive normally lethal toxin concentrations (Figure 3A, right), consistent with earlier evidence that down-regulation of LRP6 in M2182 cells decreases cytotoxicity of the PA+FP59 complex [20,21].

Unlike the effects of over-expression of *DKK1* in M2182 cells, such overexpression in HEK293T, HeLa and BHK revealed an increase, rather than decrease in toxin sensitivity (Figures 3B, 3C and 4, right), while still showing the expected ability of *DKK1* to down-regulate either constitutive or induced Wnt signaling (Figures 3B, 3C and 4, left). Quantitative PCR analysis revealed that basal expression of *DKK1* transcripts was significantly different in the above cell lines (Figure 3D, left). Additionally, this analysis showed that the abundance of *DKK1* in M2182 cells was 30× and 2× greater than that in HEK293T and

HeLa cells, respectively (Figure 3D, right).

The above findings raised the possibility that differences in the level of endogenous expression of *DKK1* may determine the consequences of adventitious overexpression of the gene and that overall *DKK1* concentration above or below the optimal range diminishes toxicity. To test this notion, we adventitiously expressed increasing amounts of *DKK1* in HEK293T, which natively expresses a low level of *DKK1* (Figure 3D), by transfecting cells with increasing amounts of a *DKK1* expressing plasmid (pEF-h*DKK1*). Whereas increasing amounts of adventitious expression of *DKK1* resulted in incremental inhibition of Wnt signaling (Figure 3E), the effects of incremental *DKK1* expression on PA-dependent toxicity were bidirectional (Figure 3F). Similar effects were observed in *DKK1* deficient M2182/*DKK1*^{-/-} cells (Figure 3G–J). Control experiments showed no effect of *DKK1* on cell viability in the absence of toxin (Figure 3H). Collectively, our results suggest that the effect of *DKK1* on cellular susceptibility to PA-mediated toxicity is both bidirectional and dependent on cell context.

Although we showed that *DKK1* promoted PA-mediated toxicity of the hybrid toxin FP59 or LFnDTA, we wanted to confirm this finding for native anthrax toxin. Accordingly, we introduced pIRES-hyg2 and pEF6MycHisBsd-B-based constructs expressing human or mouse *DKK1* into RAW264.7 cells, which are killed by PA/LF [40], and selected the stably transfected pooled population using appropriate antibiotic markers (500 µg mL⁻¹ of hygromycin for pIRES-Hyg2 clones, and 2 µg mL⁻¹ of Blasticidin for pEF6MycHisBsd-B clones). We observed that *DKK1* expression from the introduced *DKK1*-overexpression constructs reduced basal Wnt signaling (Figure 5A), as expected, and that adventitious *DKK1* expression in RAW264.7 cells reduced sensitivity to PA/LF, leading to cell survival at normally lethal toxin concentrations (Figure 5B). This result parallels to the observations we had made for M2182 cells exposed to PA+FP59. Moreover, exposure of the RAW264.7 macrophage cell line to three of seven monoclonal antibodies directed against the full-length human *DKK1* decreased PA/LF toxicity in a dose-dependent manner (Figure 5C). Control experiments showed that none of the seven antibodies tested affected cell viability per se, even at the highest antibody dosage (20 µg mL⁻¹) tested.

2.4 *DKK1* deficiency reduces PA internalization in M2182 cells

The known ability of *DKK1*, which is a secreted protein [41] to modulate LRP6 function, and the previously shown ability of LRP6 deficiency to reduce toxin internalization in M2182 cells [20] suggested that perturbation of the cellular *DKK1* level may alter internalization of toxin. As shown in Figure 6, Western blot analysis of the lysates from M2182 cells made resistant to toxin by shRNA384-6, but not by an anti-*DKK1* shRNA (i.e., shRNA387-5) that was found to

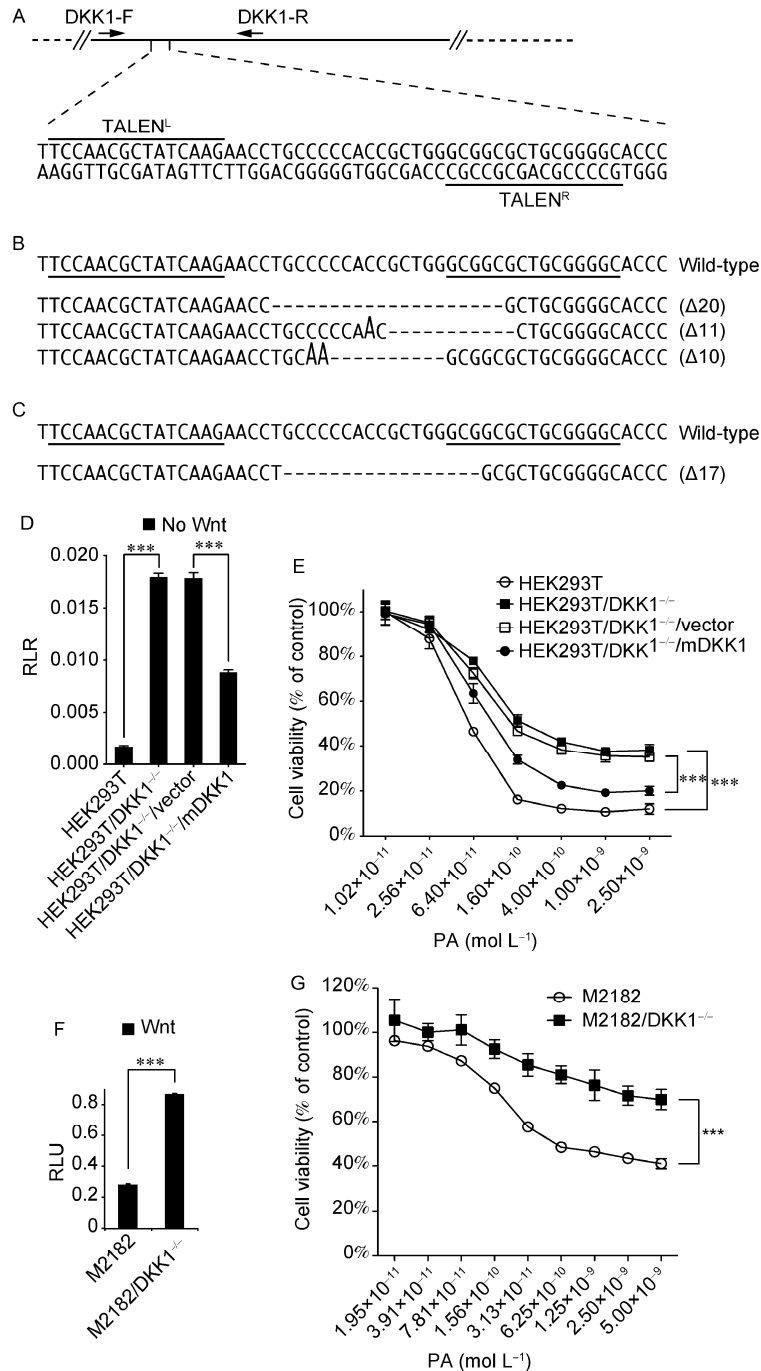


Figure 2 Effect of TALEN-mediated *DKK1* knockout on cell sensitivity of HEK293T and M2182 to PA/LFnDTA. **A**, Partial sequence of wild-type *hDKK1* gene in genome containing TALENs binding regions (overlined for TALEN^L and underlined for TALEN^R). DKK1-F and DKK1-R are primers used to amplify the TALEN-targeting region of *hDKK1* loci. **B**, Sequencing analysis of mutated alleles from eight randomly selected genome PCR clones (in pMD19-T vector) isolated from a single *DKK1*-targeting-TALEN clone (in HEK293T cells). The primers used to amplify the TALENs targeting region for sequencing analysis were 5'-CCTTCTGAGATGATGGCTCTGGGCG-3' (DKK1-F) and 5'-CCAGACGTCCAGGTACCCGCC-3' (DKK1-R). Compared with the wild-type, four clones had 20-nt deletion (Δ20), two clones had 11-nt deletion (Δ11), and two clones had 10-nt deletion (Δ10). The TALENs binding sites (underlined) are highlighted. Dashes and tall letters indicate deletions and insertions, respectively. **C**, Sequencing analysis of mutated alleles from 10 randomly selected genome PCR clones (in pMD19-T vector) isolated from a single *DKK1*-targeting-TALEN clone (in M2182 cells). All 10 clones had 17-nt deletion in the TALEN-targeting region as indicated by dashes. **D**, Effect of DKK1 knockout and exogenous expressed mouse DKK1 on cellular response to native Wnt signaling in HEK293T/DKK1^{-/-} and HEK293T/DKK1^{-/-}/mDKK1 cells. Wnt signaling assay was conducted as described as in Figure 1. Data are mean±SD, *n*=3; ***, *P*<0.005. **E**, Cytotoxicity assay of HEK293T/DKK1^{-/-} and HEK293T/DKK1^{-/-}/mDKK1 to PA/LFnDTA. The cells carrying corresponding cloning vectors were used as controls. Data are mean±SD, *n*=6; ***, *P*<0.005 for HEK293T/DKK1^{-/-} (vs. HEK293T), and ***, *P*<0.005 for HEK293T/DKK1^{-/-}/mDKK1 (vs. HEK293T/DKK1^{-/-}/vector). **F**, Effect of DKK1 knockout on cellular response to native Wnt signaling in M2182 cells. Data are mean±SD, *n*=3; ***, *P*<0.005. **G**, Cytotoxicity assay of M2182/DKK1^{-/-} to PA/LFnDTA; mean±SD, *n*=6; ***, *P*<0.005.

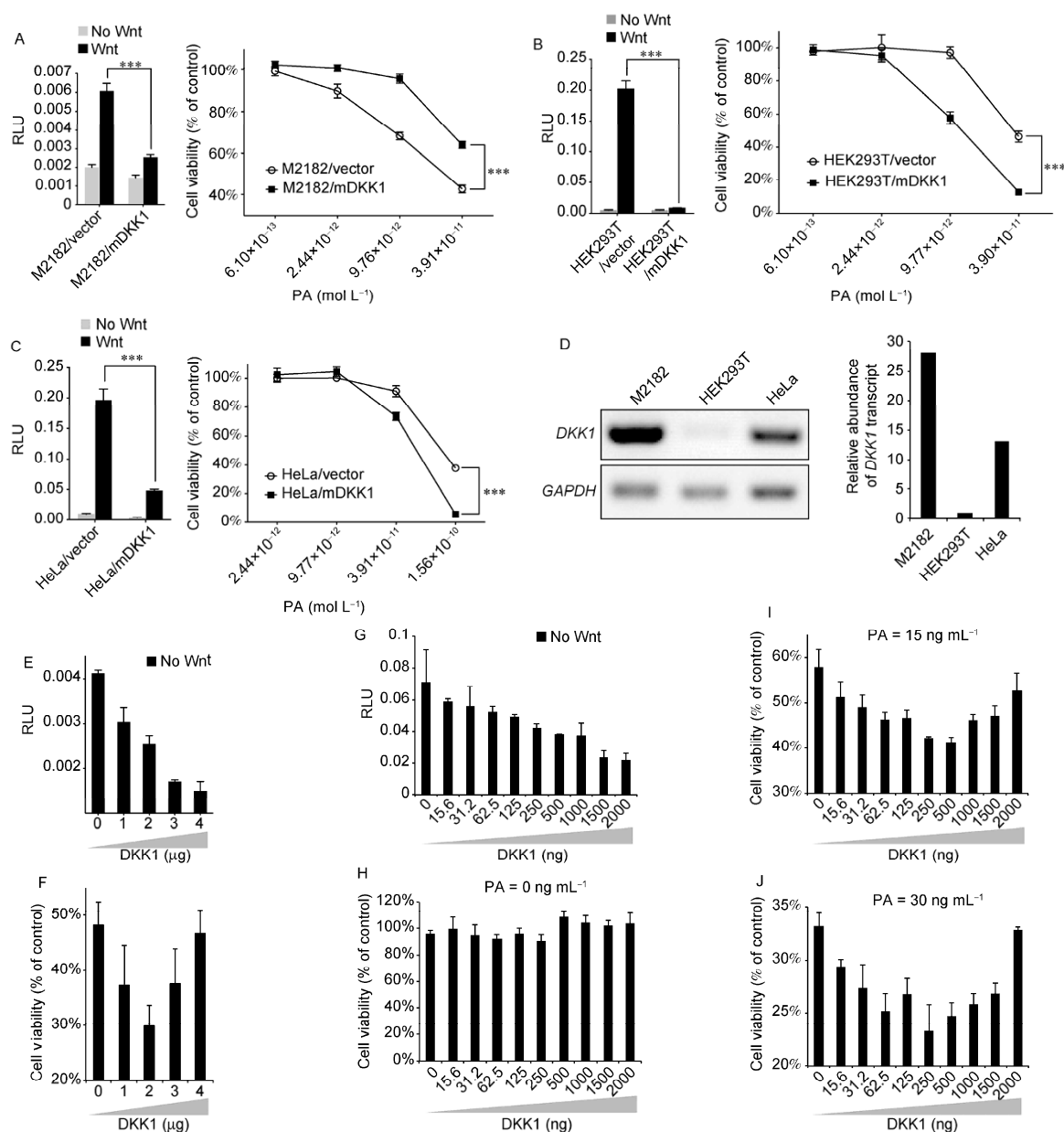


Figure 3 Bidirectional effect of adventitious expression of DKK1 on cell sensitivity to PA/LFnDTA. *DKK1* genes were cloned into two kinds of expression vectors, pIRES-hyg2 and pEF6MycHisBsd-B. A–C (left), Effect of DKK1 over-expression on cellular response to Wnt activation in M2182 (A), HEK293T (B), and HeLa (C) cells. Wnt1-stimulated Wnt signaling was assayed in cells carrying Super8XTOPflash reporter plasmid and plasmid carrying SV40-Renilla luciferase for normalization through transient transfection. Data are mean±SD, $n=3$; ***, $P<0.005$. A–C (right), Cytotoxicity of PA/FP59 in DKK1-overexpressed M2182 (A), HEK293T (B) and HeLa (C) cells. Cells carrying corresponding cloning vectors were used as controls. Cells were seeded at a concentration of 5×10^3 cells/well (96-well plate), and treated with serially diluted PA and fixed concentration of FP59 (50 ng mL^{-1}) for two days before being assayed by MTT. Data are mean±SD, $n=6$; ***, $P<0.005$. D, Level of *DKK1* transcript in M2182, HEK293T and HeLa cells was measured by RT-PCR and normalized to *GAPDH* (D, left). The primers used in PCR reactions were (5'-GACCATTGACAACACTACCAGCCGTAC-3' and 5'-CCTGAGGCACAGTCTGTATGACC-3') for *DKK1*, and (5'-ACGGATTTGGTCGTATTGGG-3' and 5'-CGCTCCTGGAAGATGGTGAT-3') for *GAPDH*. The relative abundance of *DKK1* transcript in these cells was quantified using ImageJ (<http://rsbweb.nih.gov/ij/>) (D, right). E and F, Effect of increasing amount of DKK1 on cellular response to native Wnt signaling (E) and cell survivability to PA/LFnDTA (F) in HEK293T cells. HEK293T cells were transfected with serially increased amount of pEF-hDKK1 plasmid DNA (0, 1, 2, 3, 4 μg) using Lipofectamine 2000. The corresponding cloning vector pEF was used to ensure that the total amount of DNA used in the transfection was equal (4 μg in total). Forty-eight hours post transfection, cells were assessed for both the basal level of Wnt signaling (E) and the cell survivability to PA (6.48 ng mL^{-1})/LFnDTA (50 ng mL^{-1}) (F). Data are mean±SD, $n=3$ (E) and 6 (F). G–J, Effect of increasing amount of DKK1 on cellular response to native Wnt signaling (G) and cell survivability to PA/LFnDTA in M2182/DKK1^{-/-} cells (H–J). M2182/DKK1^{-/-} cells were transfected with serially increased amount of pEF-mDKK1 plasmid DNA (0, 15.6, 31.2, 62.5, 125, 250, 500, 1000, 1500, and 2000 ng) using X-tremeGENE HP transfection reagent. The corresponding cloning vector pEF was used to ensure that the total amount of DNA used in the transfection was equal (2000 ng in total). Forty-eight hours post transfection, cells were split into four portions. One was used to test the basal level of Wnt signaling (G), and the other three were used to test the cell survivability to LFnDTA (50 ng mL^{-1}) plus variable concentration of PA (0, 15 and 30 ng mL⁻¹) (H–J). Data are mean±SD, $n=3$ (G) and 6 (H–J).

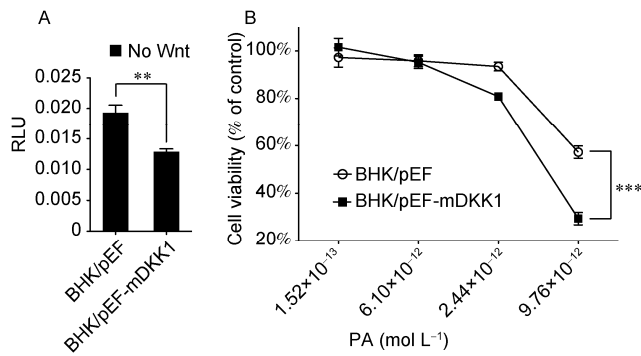


Figure 4 Effect of adventitious expression of DKK1 on cell sensitivity of BHK to PA/FP59. A, Effect of DKK1 over-expression on cellular response to native Wnt signaling in BHK cells. Data are mean±SD, $n=3$; **, $P<0.01$. B, Cytotoxicity of PA/FP59 in DKK1-overexpressed BHK cells. Data are mean±SD, $n=6$; ***, $P<0.005$.

not significantly reduce DKK1 expression (Figure 1), showed no detectable difference on 83 kD PA monomer bound to the cell surface (Figure 6A). However, treatment of cells with the *DKK1*-inhibiting shRNA was associated with reduction of an SDS-resistant oligomeric form of PA that characteristically survives proteolytic treatment when cell cultures exposed to toxin complexes at 4°C are shifted to 37°C and toxin complexes are internalized (Figure 6B). Confocal fluorescence microscopy using chimeric LF-EGFP [32] and PA indicated that the LF detected in the cytoplasm, but not at the cell surface, was reduced in the DKK1-knockdown clone M2182/shRNA384-6 (Figure 6C), consistent with our biochemical evidence of decreased PA internalization. We conclude that DKK1 affects anthrax toxicity by modulating internalization of the cell surface complex containing LRP6, PA, and PA receptors rather than by altering the binding of the PA to its receptors.

2.5 DKK1 forms a ternary complex with LRP6 and Kremen2 to internalize PA-mediated toxins

We have previously shown that LRP6 interacts with PA receptors (ANTXR), ANTXR1/TEM8 and ANTXR2/CMG2 [20]. Co-immunoprecipitation (co-IP) experiments showed that DKK1 is also a member of the complexes that PA forms with both of these ATRs (Figure 7A and B). In these experiments, the C-terminal Myc-tagged hDKK1 or HA-tagged mDKK1 was harvested from HEK293T cells incubated with lysates containing HA-tagged ANTXR1/TEM8 (ANTXR1-HA) or Myc-tagged ANTXR2ΔC/CMG2ΔC (ANTXR2ΔC-Myc, containing only the extracellular and transmembrane region of ANTXR2); DKK1 was detected in precipitates of ANTXR1-HA (Figure 7A, left), and ANTXR1 was observed in precipitates of DKK1-Myc (Figure 7A, right). Similarly, we observed ANTXR2ΔC in precipitates containing DKK1-HA (Figure 7B, left) and found DKK1 in precipitates containing ANTXR2ΔC-Myc (Figure 7B, right). Although we observed no interaction between DKK1 and PA, hDKK1-Myc was able to precipitate PA through ANTXR1 (Figure 7C). In addition, the presence of PA enhanced the interaction between DKK1 and ANTXR1 (Figure 7C). This finding plus our observation in confocal microscopy that red fluorescence-labelled DKK1-Myc co-localized with internalized ANTXR2-EGFP triggered by PA (1 μg mL⁻¹) (Figure 7D), suggested that DKK1 is a component of PA-ANTXR1/2-LRP6 complex.

The above result, together with our data showing that DKK1 and LRP6 interact functionally to facilitate the endocytosis of anthrax toxin, led us to hypothesize that the previously described ternary complex consisting of DKK1, LRP6 and Kremen2 [26] may facilitate endocytosis of the toxin—together with its receptor [20,21,42]. To test this

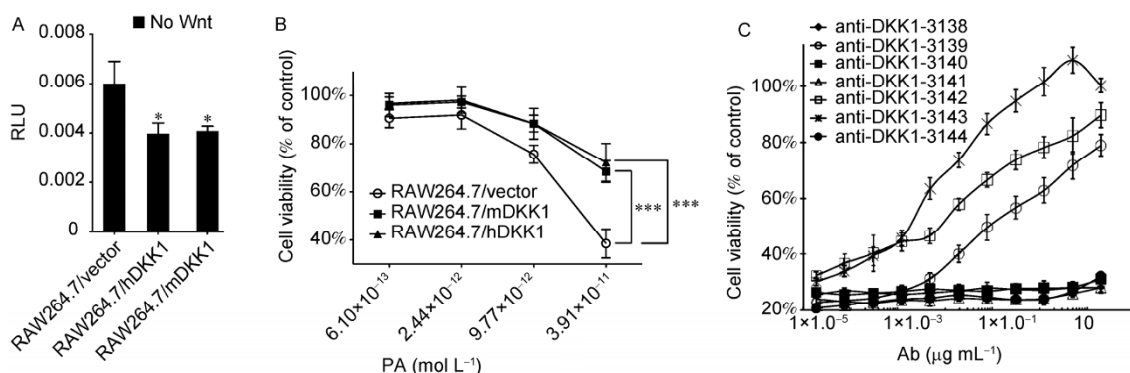


Figure 5 Effects of exogenous expression of DKK1 and anti-hDKK1 monoclonal antibodies on killing of RAW264.7 cells by PA/LF. A, Effect of DKK1 over-expression on cellular response to native Wnt signaling in RAW264.7 cells. Data are mean±SD, $n=3$; *, $P<0.05$. B, Cytotoxicity of PA/LF in DKK1-overexpressed RAW264.7 cells. Cells carrying corresponding cloning vectors were used as controls. Data are mean±SD, $n=6$; ***, $P<0.005$. C, Effect of monoclonal antibodies against hDKK1 on the sensitivity of RAW264.7 cells to PA/LF. 5×10^4 mL⁻¹ cells of RAW264.7 were first seeded in 96-well plates (100 μL/well) the day prior to the assay. The cells were pre-incubated with serially diluted antibodies (anti-DKK1-3138–3144) for 1 h, followed by the addition of DMEM growth medium containing or lacking a fixed concentration of toxins (PA and LF, each 50 ng mL⁻¹). The plates were incubated at 37°C and with 10% CO₂ for 24 h prior to the cell viability assay as described [20]. Data are mean±SD, $n=6$.

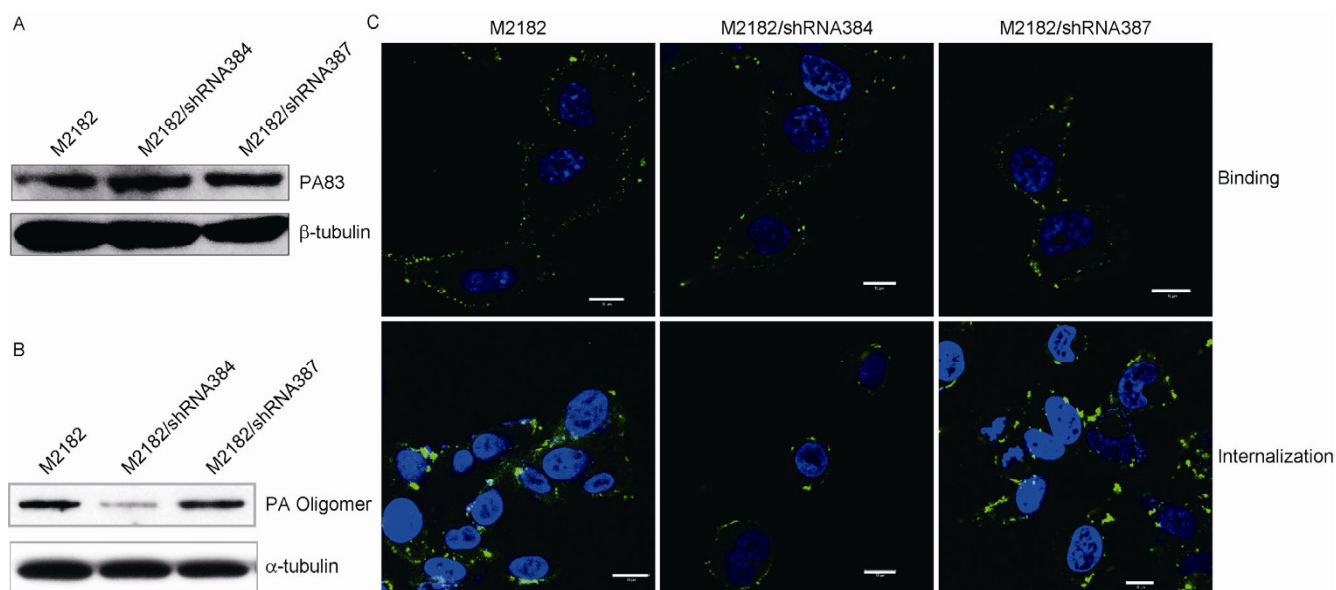


Figure 6 Effect of DKK1 deficiency in M2182 cells on PA binding and internalization. A and B, Western blot analysis of bound (A) and internalized (B) PA in different cells was conducted following the protocol as described [20]. PA83 is the full-length form of PA (83 kD), and the oligomer represents the heptamer of PA63. α -tubulin or β -tubulin abundance was measured as an internal control. C, Immunofluorescence microscopy of PA/EGFP binding and internalization in different cells. To monitor cell surface binding (top), cells were first treated with PA83 (600 nmol L^{-1}) for 12 min at 37°C , washed, incubated with LF-EGFP (200 nmol L^{-1}) for 2 min at 37°C and immediately washed and fixed before examined by confocal microscopy. To monitor PA/EGFP internalization (bottom), cells were incubated with PA83 (600 nmol L^{-1}) and LF-EGFP (200 nmol L^{-1}) for 15 min at 37°C and followed by confocal microscopy [32]. Scale bar, $10 \mu\text{m}$.

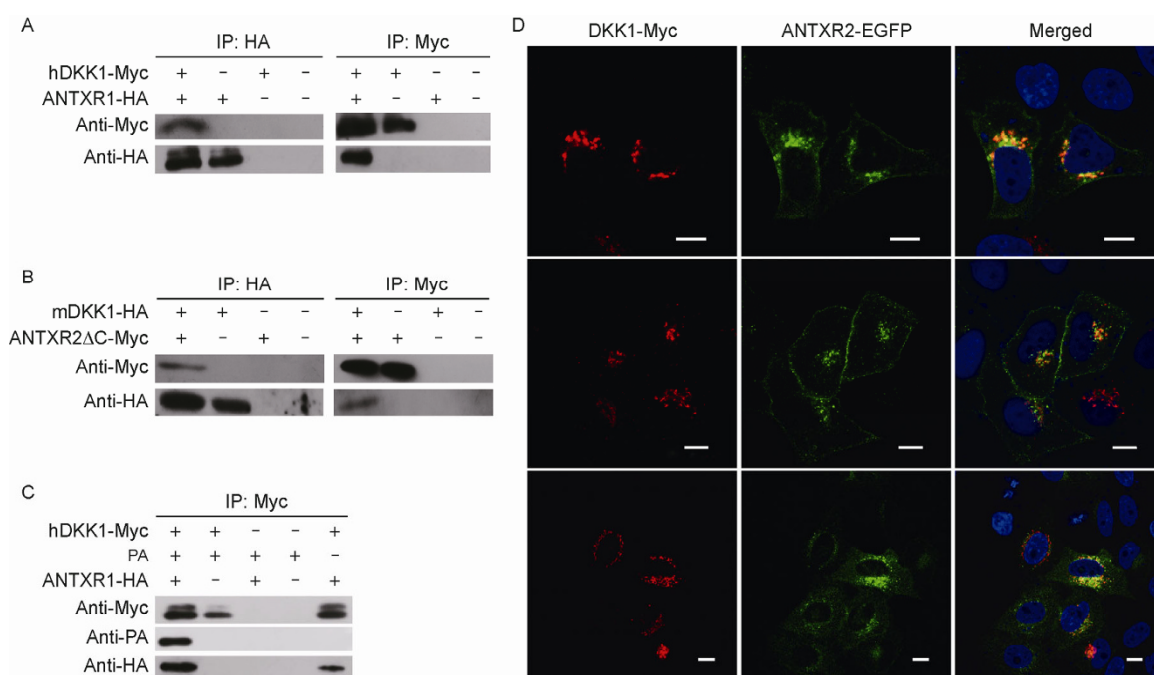


Figure 7 Binding of DKK1 to ANTXR1, ANTXR2 and PA. A, HEK293T cells were transfected with hDKK1-Myc, ANTXR1-HA, or both. Cell lysates were treated by addition of anti-HA-conjugated agarose beads (left) or anti-c-Myc-conjugated agarose beads (right). The precipitated proteins were then subjected to immunoblotting analysis probed with anti-c-Myc antibody conjugated with horseradish peroxidase (HRP) or anti-HA-HRP. B, HEK293T cells were transfected with mDKK1-HA, ANTXR2 Δ C-Myc, or both. Cell lysates were treated by addition of anti-HA-conjugated agarose beads (left) or anti-c-Myc-conjugated agarose beads (right). The precipitated proteins were then subjected to immunoblot analysis probed with anti-c-Myc-HRP or anti-HA-HRP. C, HEK293T cells were transfected with hDKK1-Myc, ANTXR1-HA, or both. Cell lysates were then incubated with or without PA83 on ice for 2 h before being mixed with anti-c-Myc-conjugated agarose beads. The precipitated proteins were then subjected to immunoblotting analysis probed with anti-c-Myc-HRP, anti-HA-HRP or anti-PA antibodies. D, Confocal microscopy images showing co-localization of DKK1-Myc (red) and ANTXR2-EGFP (green) in the presence of PA ($1 \mu\text{g mol L}^{-1}$). Three randomly selected images were used. Scale bar, $10 \mu\text{m}$.

hypothesis, we over-expressed LRP6, Kremen2, or LRP6 plus Kremen2 in M2182 cells under two extreme situations: one in which over-expression of DKK1 was constitutive, M2182/DKK1 (Figure 8A), and the other one in which DKK1 was absent, M2182/DKK1^{-/-} (Figure 8B). In the DKK1-high-expression background, adventitious expression of LRP6 significantly decreased cell sensitivity to PA/LFnDTA, and the addition of Kremen2 had little effect. However, adventitious expression of LRP6 plus Kremen2 in the presence of abundant DKK1 made M2182 much more sensitive to the toxin (Figure 8A). In contrast, in the absence of DKK1, the adventitious expression of LRP6, Kremen2 or the combination of these two showed no effect on cell sensitivity to PA/LFnDTA (Figure 8B).

3 Discussion

To exert its toxic effects, anthrax toxins must first enter the host cell cytoplasm. Three cell surface proteins can serve as receptors for PA [8–10]. Although the binding of PA to only one of the receptors is sufficient for toxin entry [43–45], loss or reduction in any of the receptors decreases toxicity [8–10], suggesting that these cell surface proteins can interact functionally or physically. LRP6, initially discovered as a co-receptor for Wnt [16–18], is a compo-

nent of complexes containing ANTXR1/TEM8 and/or ANTXR2/CMG2 and can promote internalization of anthrax toxin complex [20,21]. Our discovery that DKK1, a secreted LRP6 ligand and regulator [22,25], also affects anthrax toxin internalization provides additional evidence of the role of Wnt signaling proteins on anthrax toxicity.

Although all three ATRs contain VWA/I domains (von Willebrand factor type A domains, also called integrin-like domains), their sequences share only 40% overall amino acid identity and their primary biological roles in mammalian cells are highly disparate [9]. By generating ANTXR1/TEM8-, ANTXR2/CMG2-null and TEM8/CMG2 double null mice, Leppla and colleagues recently have shown that ANTXR1/TEM8 plays a relatively minor role in mediating anthrax toxicity in rodents. ANTXR1/TEM8-null mice were as susceptible as wild type mice to lethal toxin, edema toxin, or anthrax spore challenge [11].

DKK1 is a secreted protein that binds to LRP6, as well as to Kremen (Kremen1 or Kremen2), another cell surface receptor protein, and the ternary structure formed by the interaction of these three proteins triggers rapid internalization of the complex [26]—inhibiting Wnt signaling. The data presented here and previously [20] suggest that the anthrax toxin complex hijacks this ternary structure to accomplish toxin endocytosis. Increased DKK1 production in our experiments resulted in increased

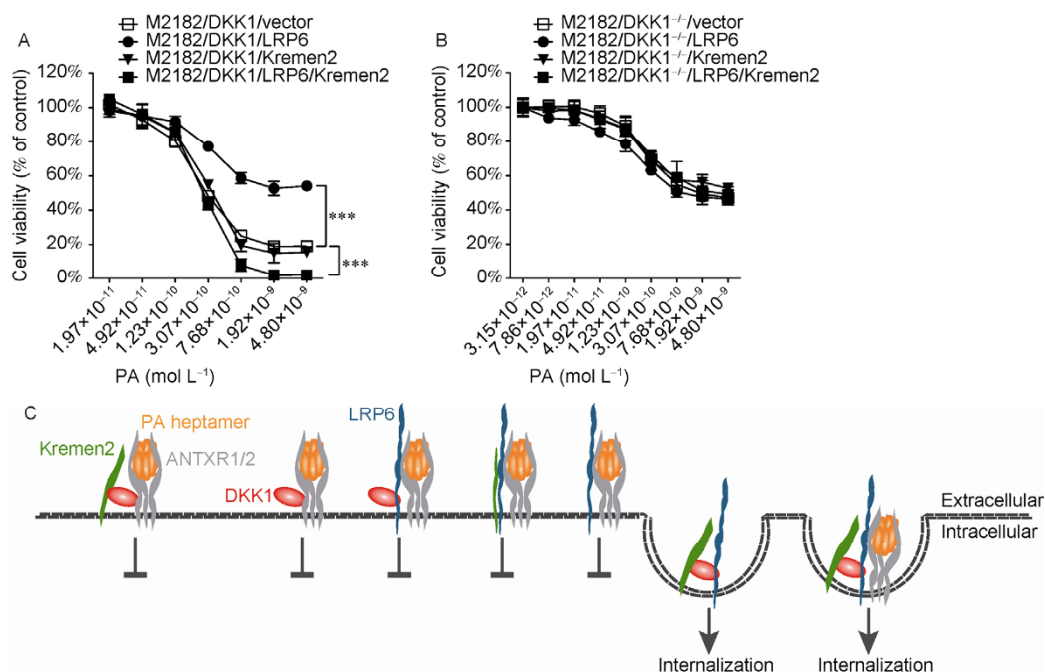


Figure 8 Effect of the ternary structure of LRP6/DKK1/Kremen2 on cell sensitivity to anthrax toxin. A and B, Effect of exogenous expression of LRP6, Kremen2 or LRP6+Kremen2 in DKK1-overexpressed M2182 (A) or DKK1-deficient M2182/DKK1^{-/-} (B) on the cytotoxicity of PA/LFnDTA. Cells carrying corresponding cloning vectors were used as controls. Data are mean±SD, *n*=6; ***, *P*<0.005. C, Model of ternary complex of LRP6/DKK1/Kremen2 in the endocytosis of anthrax toxin. Individual or any two-protein combination of LRP6, DKK1 and Kremen2 could potentially bind to the anthrax toxin complex, but none of them is capable of mediating the endocytosis process. Therefore, overexpression of DKK1, LRP6, or any two-protein combination of LRP6, DKK1 and Kremen2 may cause dominant-negative effect, depending on the basal level of these three proteins. Only the complete form of the ternary complex is fully functional to internalize PA-ANTXR1/2-LFnDTA, and this ternary complex could undergo internalization by itself [26].

antagonism of Wnt signaling, as has been reported previously. However, the effects of such increased production on toxin sensitivity differed among the cell lines tested and were shown to be dependent on the extent of overproduction in at least two cell types. Similarly, whereas increased antagonism of LRP6-mediated Wnt signaling by up-regulation of DKK1 increased toxin lethality in HeLa, HEK293T and BHK cells, the reverse was true in M2182 and RAW267.4 macrophages. Collectively, these findings indicate that the effects of DKK1 antagonism of LRP6 are not congruent with the effects on Wnt signaling—which can also be bidirectional for a given DKK1 perturbation.

It has been reported that propeller domain 3 of LRP6 plays a dominant role in binding to DKK1 [26,46]. However, DKK1 has also been shown to bind to the first propeller domain of LRP6, and monoclonal antibody targeting on this region completely protected LRP6 from DKK1-mediated internalization [47]. Study of crystal structure revealed that the C-terminal domain of DKK1 binds to the top surface of the third propeller domain of LRP6, as well as its first propeller domain [48]. LRP6 has been shown to interact with both PA receptors, ATRs [20]. Similarly, DKK1 was found to be a component of complexes of PA with ANTXR1/TEM8 and ANTXR2/MG2. This finding is not surprising given known ability of LRP6 to interact with DKK1 [49]. However, whether DKK1 interacts with ATRs through LRP6 or conversely, whether DKK1 serves as a bridge between LRP6 and PA receptors, is not currently known.

Anthrax toxin triggers endocytosis of its receptors via a lipid raft-mediated process [43], and PA treatment was found to be able to recruit LRP6 protein from detergent soluble fractions to detergent-resistant membrane (DRM), mimicking the process of ATRs upon PA addition [21]. Interestingly, LRP6 was found present in lipid raft microdomains [50], and DKK1 treatment is able to remove LRP6 from the lipid raft [51]. Moreover, the entry of anthrax toxin proceeds mainly through a clathrin-dependent pathway because PA internalization is dependent on the actions of dynamin and Eps15 [43]. Intriguingly, DKK1-induced internalization of LRP6 is also dependent on clathrin, and the knockdown of clathrin was able to suppress the DKK1-dependent inhibition of Wnt signaling [52]. DKK1 belongs to a protein family that contains four members. Among them, both DKK2 and DKK4 can also bind to LRP6, although with lower affinity than DKK1, and inhibit Wnt signaling [41,53,54]. Whether the DKK2 and DKK4 proteins also participate in anthrax toxin endocytosis, and whether there are other secreted factors beyond DKK family proteins that can affect anthrax toxin uptake is unknown.

Consistent with earlier evidence that LRP6 promotes PA uptake [20,21], antagonism of LRP6 function by overexpression of DKK1 reduced toxin sensitivity in M2182 and RAW267.4 cells. However, such enhanced antagonism of

LRP6 function, as measured by a decrease in Wnt signaling, sensitized certain cell lines to PA/FP59, including HeLa, HEK293T and BHK cells (this report), suggesting that a decrease in LRP6 functionality in cells undergoing constitutive Wnt signaling did not significantly decrease availability of LRP6 for internalization of toxin. While providing additional evidence that Wnt signaling pathway components have important roles in toxin endocytosis, our findings also show that anthrax toxicity is independent of Wnt signaling per se.

The effects of perturbation of DKK1 production and consequently on its antagonism of LRP6 function in different cell types may reflect possible differences of DKK1/LRP6 ratio, which may in turn be influenced by cell type-specific differences in the extent of use of LRP6 for autocrine Wnt signaling or the production of other proteins that affect toxin entry (e.g., ARAP3; Lu et al.). The bidirectional effect of DKK1 overproduction observed in HEK293T cells and M2182/DKK^{-/-} is reminiscent of the role of DKK2 in LRP-mediated signaling, where the LRP6 ligand can act as either an inhibitor or an activator of the Wnt pathway, depending on the cellular context (i.e., the presence or absence of Kremen2) [55]. Kremen2 enhance Wnt signaling by maintaining LRP5/6 at the plasma membrane in the absence of DKK1, while they inhibit Wnt signaling by promoting the endocytosis of LRP5/6 in the presence of DKK1 [24].

Importantly, using TALEN technology, we were able to knock out *DKK1* gene expression in both HEK293T and M2182 cells. The complete loss of function of DKK1 confirmed its important role in anthrax toxicity. In addition, from the results of expressing DKK1, LRP6 or Kremen2, either individually or in combination, in high- or low-DKK1-expression M2182 cells, we demonstrated that (i) Kremen2 significantly increased cell sensitivity in high DKK1/LRP6 background; (ii) LRP6 significantly increased cell sensitivity in high DKK1/Kremen2 background; (iii) overexpression of LRP6 plus DKK1 greatly decreased cell sensitivity to PA-mediated toxicity, a dominant-negative effect in comparison with the result of overexpression of LRP6/DKK1/Kremen2; and, (iv) effects of LRP6 and/or Kremen2 on PA-mediated toxicity were dependent on DKK1. We therefore propose a working model in which the ternary structure, consisting of DKK1-LRP6-Kremen2, promotes the internalization of PA-ANTXR1/2-LF/EF to exert their toxic effect. It was reported that the association of DKK1, LRP6 and Kremen2 triggers rapid internalization of this ternary complex [26]. Overexpression of DKK1, LRP6, or any two-protein combination of LRP6, DKK1 and Kremen2 may cause dominant-negative effect by competing the ternary complex to bind to the anthrax toxin complex without mediating the endocytosis of the latter (Figure 8C), possibly explaining why both up- and down-regulation of DKK1 could reduce cell's sensitivity to the toxin.

Worthy of special mention is the finding that even the

complete loss of DKK1 in TALEN-mediated knockout cells did not make cells completely resistant to PA-mediated toxicity. Similarly, the effects of decreased LRP6 function on anthrax toxicity have been controversial [20,21,56,57], and it has been suggested that differences in results obtained in different laboratories may reflect variation in the type of cells tested [56,57]. We conclude from this finding that there are multiple mechanisms for toxin entry, as suggested by evidence that at least three cell surface proteins capable of interacting with PA have been identified [8–10]. Although a central role of ANTXR2/CMG2, which binds to PA more tightly than other receptors [58,59], has been shown for anthrax toxin lethality in mice [11], receptor expression is known to differ in different cell types [8,9,50,59]. Additionally, knockout of ANTXR1/TEM8 in HeLa cells abolished susceptibility to PA/LFnDTA, while the knockout of ANTXR2/CMG2 in HeLa cells had no effect at all [60]. On the contrary, knockout of ANTXR2/CMG2 in HEK293T cells completely abolished susceptibility to PA/LFnDTA, while the knockout of ANTXR1/TEM8 in HEK293T cells had no effect [60]. Interestingly, a recent publication by Leppla and colleagues [61] showed that LT and ET induced lethality are different on distinct cell types.

We thank S. Leppla (National Institute of Allergy and Infectious Diseases, NIH) for kind gifts of reagents, R.J. Collier (Harvard Medical School) for the plasmid pET-15b-LFN_{DTA}, P. Chen (Peking University) for LF-EGFP protein, and Christine Tan, J. Michael Elliot, and Yan Gong (Genentech) for DKK1 antibody production and initial characterization. These studies were supported in part by the National Natural Science Foundation of China (30770465, 31070115), the National Basic Research Program of China (2010CB911800), and the Peking-Tsinghua Center for Life Sciences (to WenSheng Wei), and by an award (HDTRA1-06-C-0039) from the US Defense Threat Reduction Agency (to Stanley N. Cohen). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

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