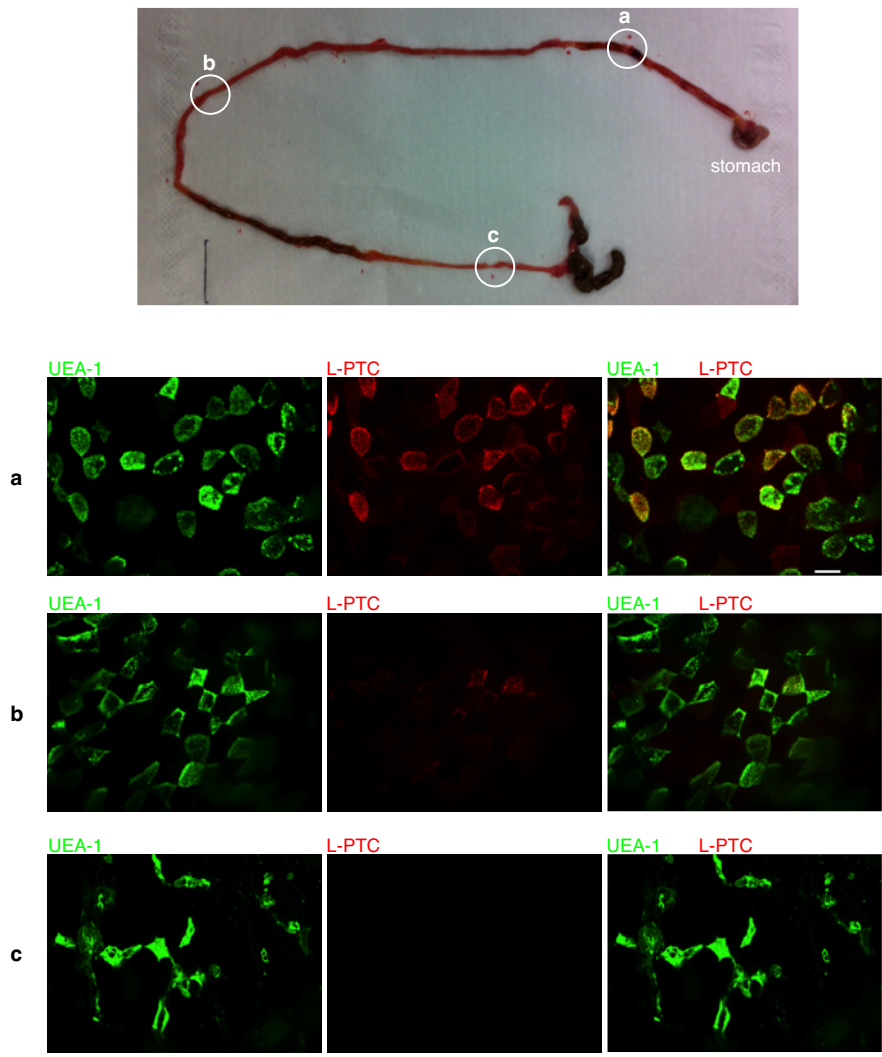


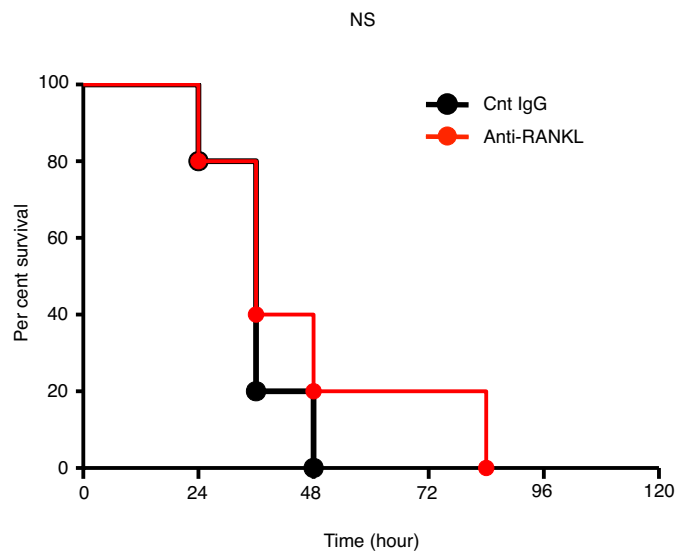
Supplementary Figure 1: Co-localization of reconstituted L-PTC and dendritic cells

(DCs). **(a)** Alexa Fluor 488–labeled L-PTC (green) was injected into ligated mouse intestinal loops and incubated for 3 h; FAE regions were stained with a anti-Na⁺K⁺ATPase antibody (basolateral marker), Cy3–labeled anti-rabbit IgG antibody (red), and Pacific Blue–labeled anti-CD11c antibody (DC marker, blue). X-Z images in lower panels correspond to the positions indicated by dotted lines in the X-Y images. Right panels show a higher-magnification image of the boxed region. L-PTC were captured by DCs in the SED (arrows). **(b and c)** For reconstitution of the L-PTC, Alexa Fluor 488–labeled BoNT (green) and Alexa Fluor 568–labeled NAPs (red) were mixed at a molar ratio of 5:1 (1000 nM : 200 nM) in PBS (pH 6.0), and then incubated for 3 h at 37°C. Reconstituted L-PTC was confirmed by pull-down using lactose gel beads (EY Laboratories, Inc.),

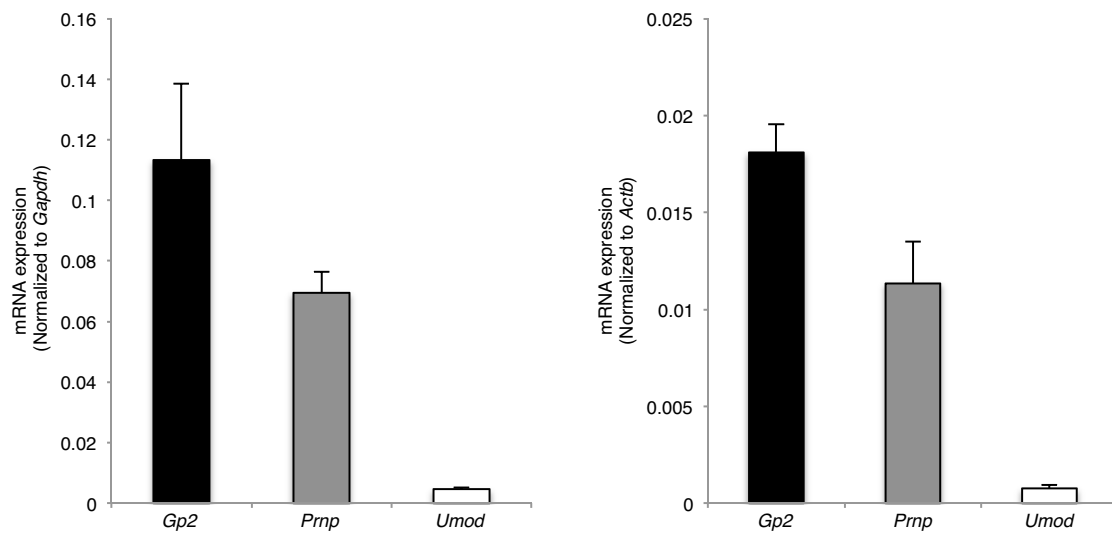
which bound to HA, followed by immunoblotting with anti-L-PTC antibody (data not shown). Reconstituted L-PTC was injected into ligated intestinal loops and incubated for 3 h. FAE was stained with a Pacific Blue–labeled anti-CD11c antibody (blue). **(b)** X-Y view. **(c)** Three-dimensional data, Top, X-Y view. Bottom, side (X-Z) view. X-Y images **(b)** correspond to the position indicated by dotted line in the X-Z view **(c)**. For three-dimensional imaging, see Supplementary Movie 1. Scale bars: 10 μm (**a**, **b**, and higher-magnification images in **a**); 5 μm (**c**). Data are representative of two independent experiments.



Supplementary Figure 2: Localization of orally administered L-PTC in small intestine. Alexa Fluor 568–labeled L-PTC (26.7 pmol: 20 μ g, red) was intragastrically administered to mice and incubated for 1 h. After incubation, the small intestine was removed, and PPs were excised from this tissue. PPs from duodenum (**a**), jejunum (**b**), or ileum (**c**) were stained with FITC–labeled UEA-1 (green), and analyzed on a confocal laser microscope. Scale bar, 10 μ m. Data are representative of two independent experiments.

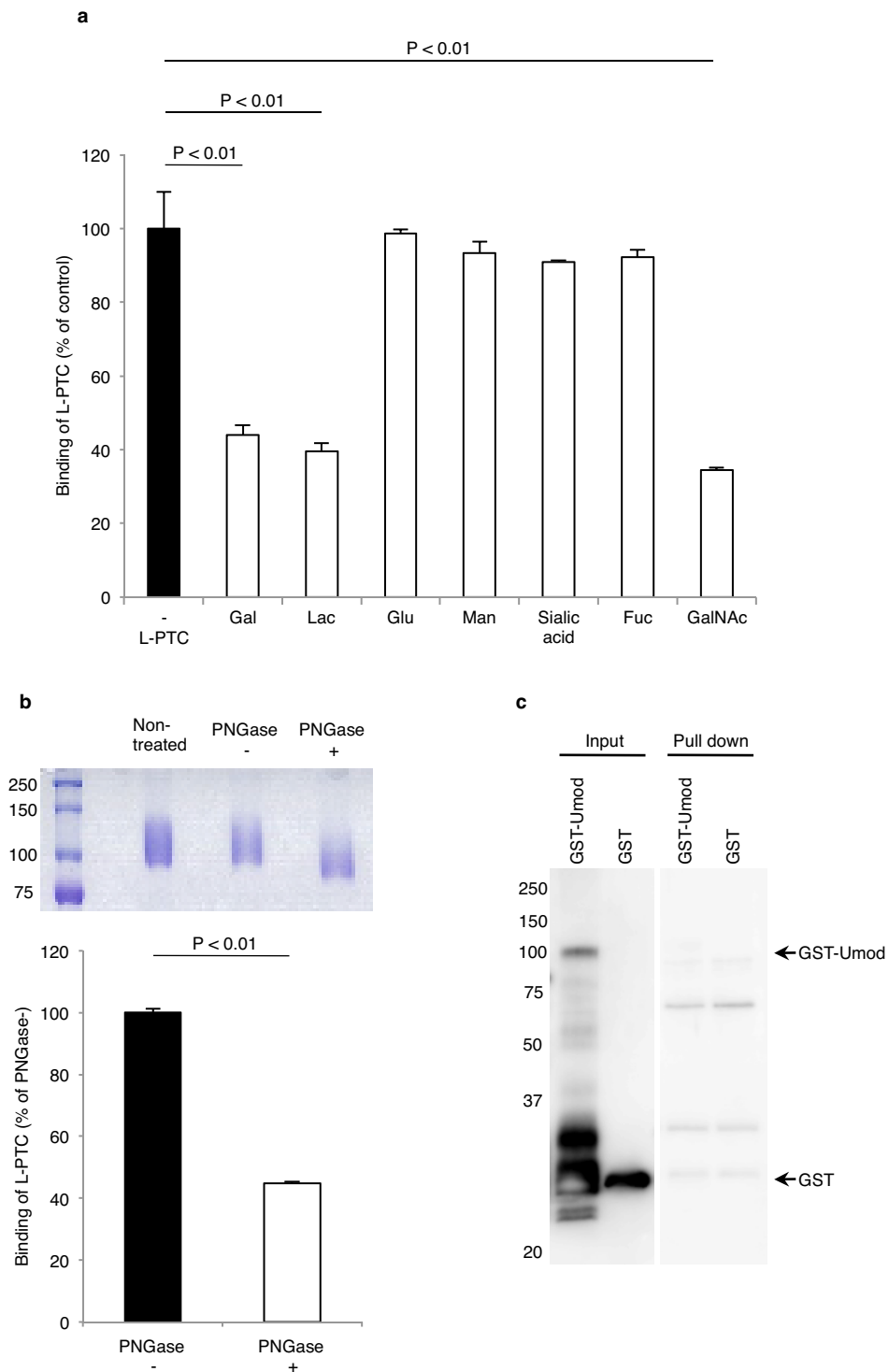


Supplementary Figure 3: M-cell depletion does not influence lethality upon oral administration of M-PTC. Mice were treated intraperitoneally with anti-RANKL antibody or control rat IgG. M-PTC were intragastrically (174.4 pmol: 50 μ g) administered to mice treated with anti-RANKL antibody or rat IgG ($n = 5$ per group). Statistical analyses were performed with the log-rank test. NS, not significant.



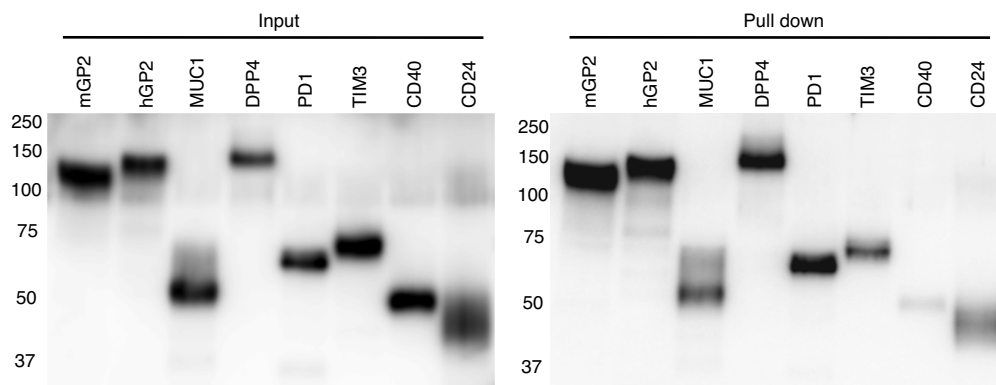
Supplementary Figure 4: qPCR analysis of mouse GP2, PrP^C, and uromodulin

mRNAs. Total RNA was prepared from FAE. Expression of each gene was analyzed by real-time qPCR. The expression levels of mRNAs encoding GP2 (*Gp2*), PrP^C (*Prnp*), and uromodulin (*Umod*) were normalized to the level of the mRNA encoding glyceraldehyde-3-phosphate dehydrogenase gene (*Gapdh*, **a**) or β -actin (*Actb*, **b**). Error bars indicate s.d. ($n=3$).

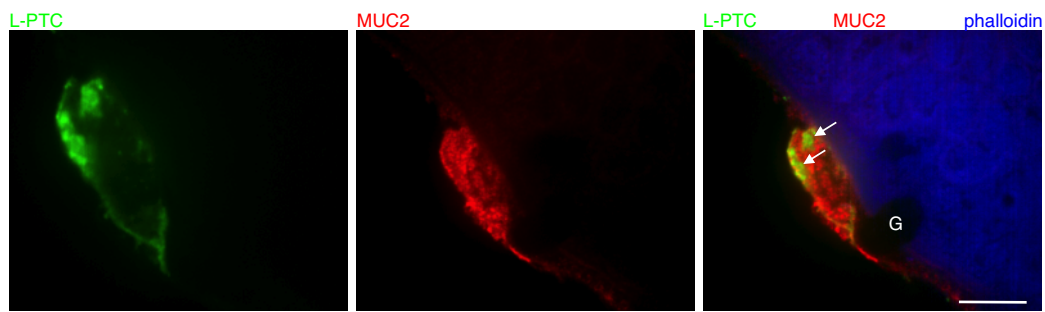


Supplementary Figure 5: Binding of L-PTC to GP2 and uromodulin is mediated by sugar chains. (a) L-PTC (100 nM) was pre-incubated with galactose (Gal), lactose (Lac),

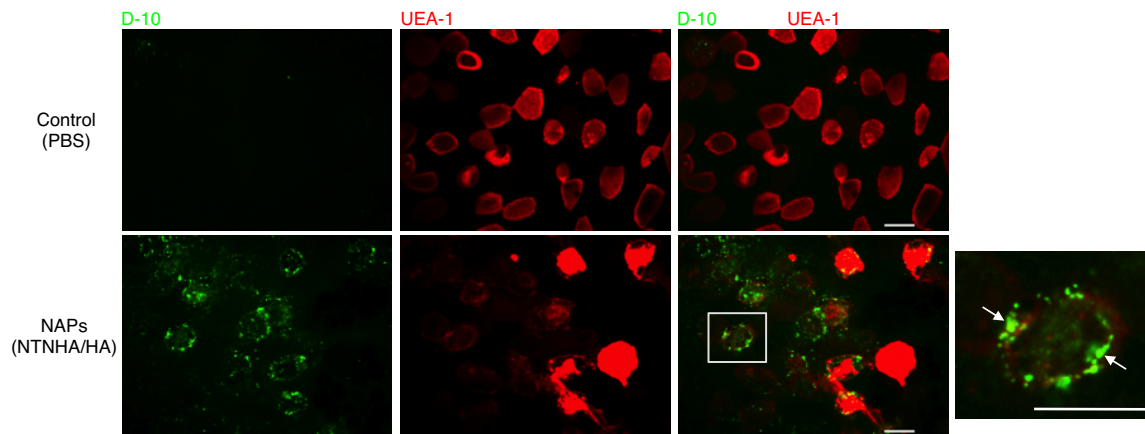
glucose (Glu), mannose (Man), sialic acid, fucose (Fuc), or N-acetylgalactosamine (GalNAc) (10 mM), and then added to plates coated with mGP2-Fc proteins. Bound L-PTC was detected using anti-BoNT and HRP-labeled anti-rabbit IgG antibodies. The data are expressed as a percentage of the binding of L-PTC in the absence of added carbohydrates. Error bars indicate s.d. ($n = 3$). Statistical analyses were performed with the Student's *t*-test. **(b)** Deglycosylation of mGP2-Fc was confirmed by SDS-PAGE. Binding of L-PTC to deglycosylated mGP2-Fc was analyzed by ELISA. Bound L-PTC was detected using anti-BoNT and HRP-labeled anti-rabbit IgG antibodies. Error bars indicate s.d. ($n = 3$). Statistical analyses were performed with the Student's *t*-test. **(c)** GST-mouse uromodulin or GST was incubated with Strep-Tactin Superflow agarose pre-bound to HA. HA-bound proteins were analyzed by immunoblotting with anti-GST antibody and HRP-labeled anti-rabbit IgG antibody.



Supplementary Figure 6: HA binds to various glycoproteins. Recombinant Fc proteins were incubated with Strep-Tactin Superflow agarose pre-bound to HA. HA-bound proteins were analyzed by immunoblotting using HRP-labeled anti-human IgG antibody. Data are representative of two independent experiments.



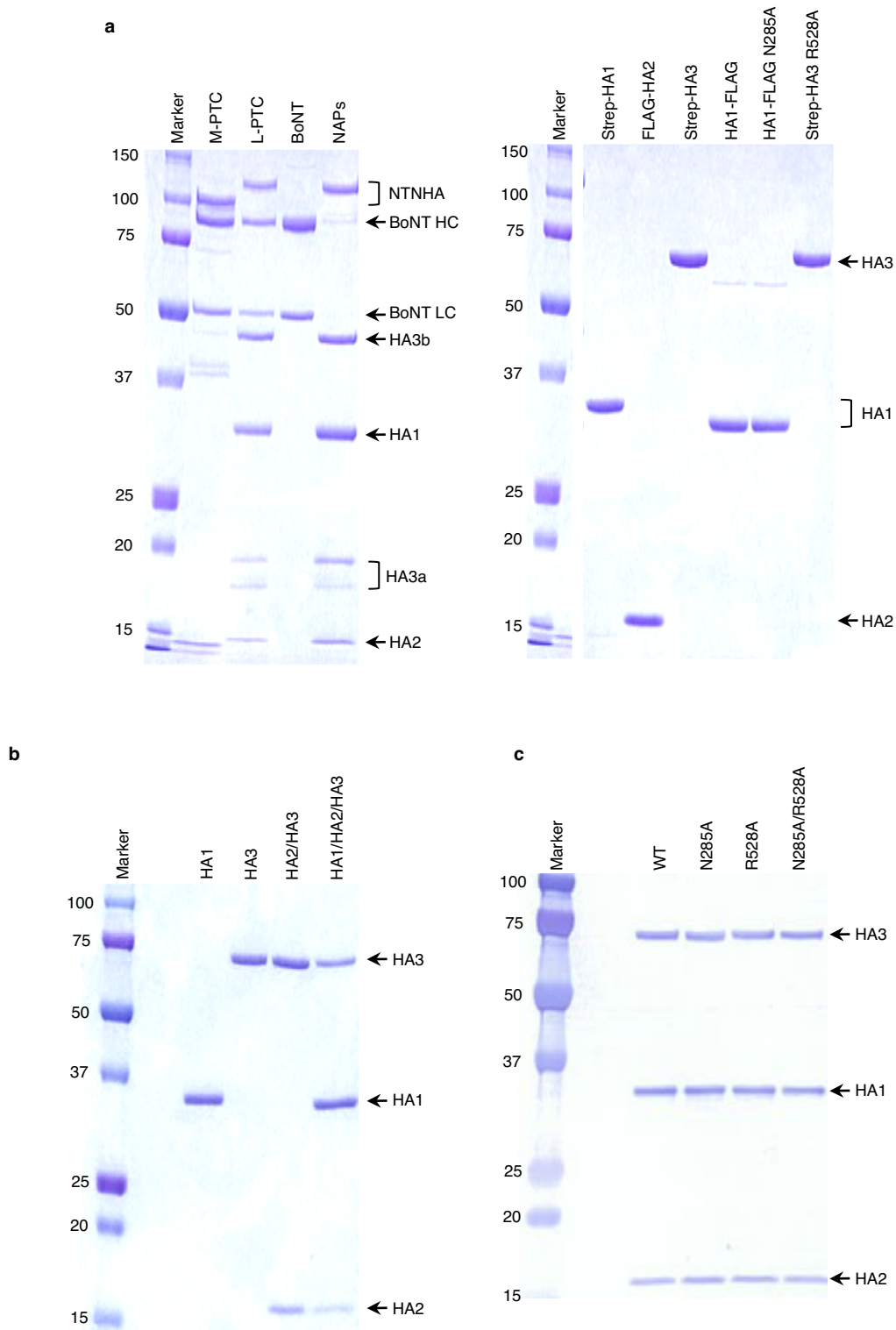
Supplementary Figure 7: L-PTC is trapped into the mucus layer. Alexa Fluor 488–labeled L-PTC (green) were injected into ligated mouse intestinal loops and incubated for 1 h. Whole-mount specimens were fixed with Carnoy’s solution and stained with anti-MUC2 antibody and Cy3–labeled anti-mouse IgG antibody (red) and iFluor 405–labeled phalloidin (blue). L-PTC was trapped in the mucus layer (arrows). G: goblet cell. Scale bar, 10 μ m. Data are representative of two independent experiments.



Supplementary Figure 8: NAPs (NTNHA/HA) disrupt the paracellular barrier

around the M cells. NAPs (NTNHA/HA) and paracellular tracer (fixable FITC–dextran 10K; D-10, green) or vehicle control (PBS) and the tracer were injected into ligated mouse intestinal loops and incubated for 2 h, and FAE regions were stained with rhodamine-labeled UEA-1 (red). Right panel shows a higher-magnification image of the boxed region. Paracellular tracer penetrated between M cells and enterocytes (arrows).

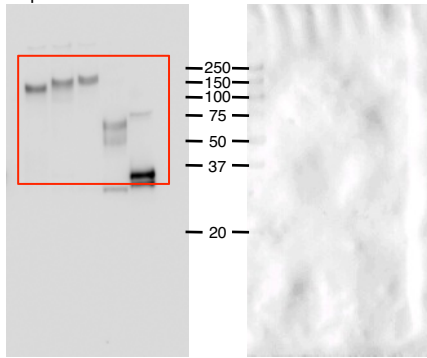
Scale bars: 10 μ m. Data are representative of two independent experiments.



Supplementary Figure 9: Toxins, NAPs, and recombinant HA proteins used in this

study. (a) SDS-PAGE of M-PTC, L-PTC, BoNT, NAPs (2.0 μg), and recombinant HA protein (1.0 μg). HA3 protein is cleaved to yield two fragments, designated HA3a and HA3b (alternatively termed HA20 and HA52, respectively), in L-PTC and NAPs⁸. (b and c) Recombinant HA1, HA2, and HA3 proteins were mixed at a molar ratio of 4:4:1 (20 μM :20 μM :5 μM) in PBS (pH 7.4), and then incubated for 3 h at 37°C. Reconstituted HA (complex of Strep-HA1, FLAG-HA2, and Strep-HA3, b) and reconstituted HA harboring mutant HA1 or HA3 (complex of HA1-FLAG [N285A], FLAG-HA2, and Strep-HA3 [R528A], c) were confirmed by pull-down using Strep-Tactin Superflow agarose and SDS-PAGE.

Fig. 3a
Input



Pull down

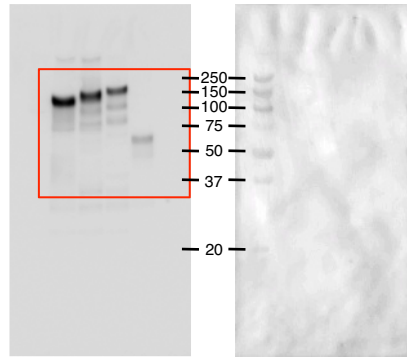


Fig. 5c

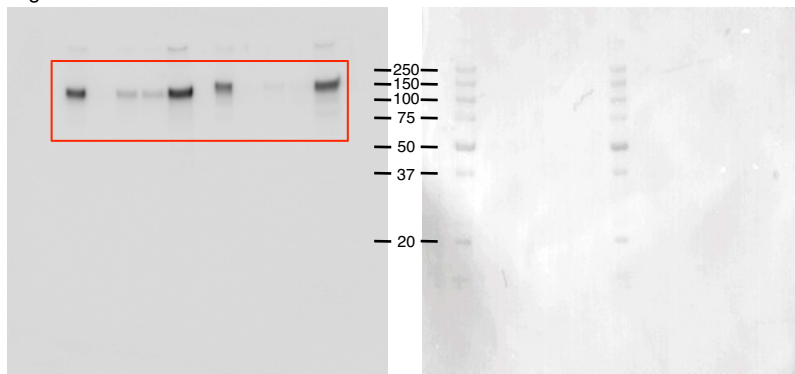


Fig. 5e

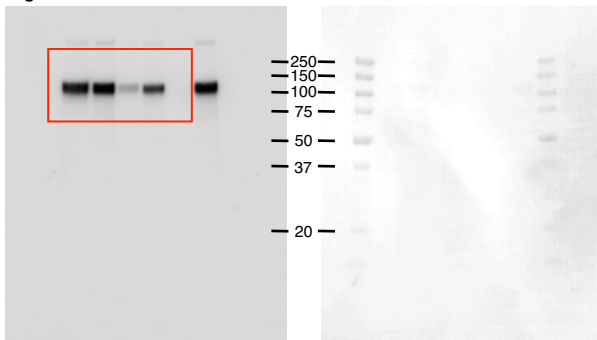
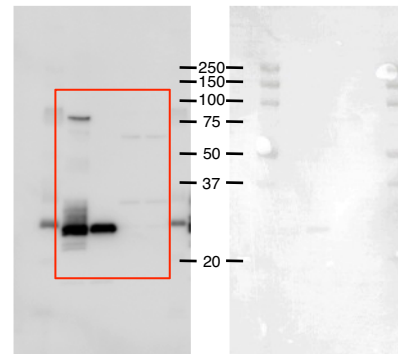
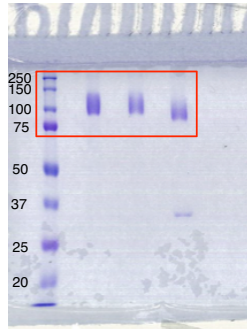


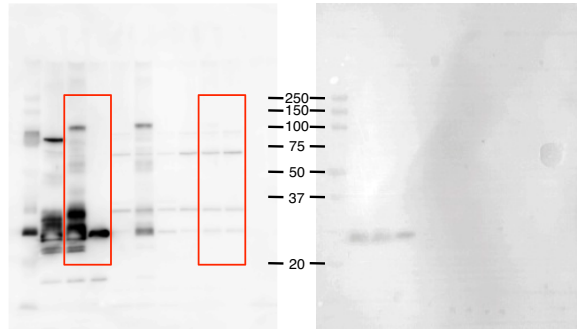
Fig. 5f



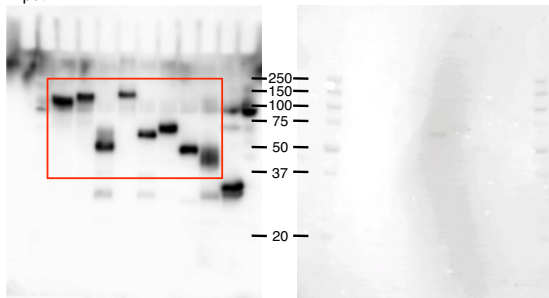
Supplementary Fig. 5b



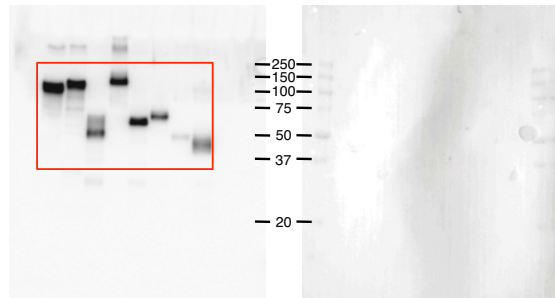
Supplementary Fig. 5c



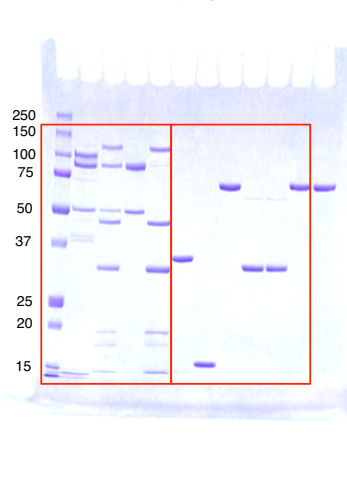
Supplementary Fig. 6
Input



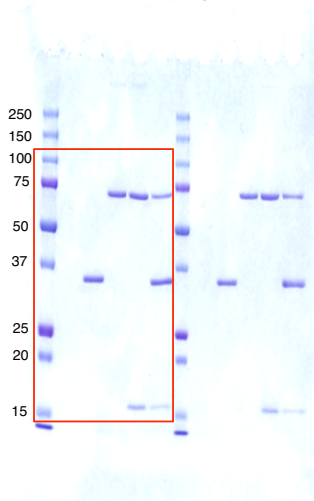
Pull down



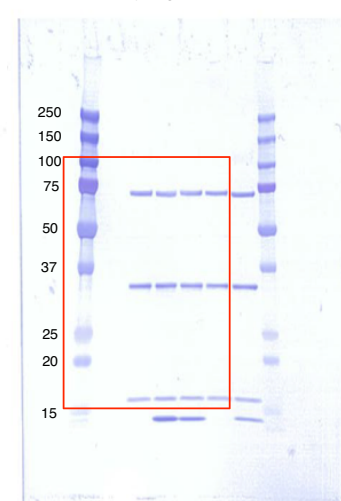
Supplementary Fig. 9a



Supplementary Fig. 9b



Supplementary Fig. 9c



Supplementary Figure 10: Uncropped scan images of immunoblots and SDS-PAGE gels (CBB staining). The red boxed regions in the uncropped scan images were used in the manuscript. The right side of the uncropped chemiluminescence image is a bright-field image of the blot membrane, showing molecular markers (kDa).

Supplementary Table 1: Primers sequences used in this study.

Primer	sequence (5'-3')
Strep-HA1-F	GGAACGGTACCCAGTAATCCAAAATTCATTAATG
Strep-HA1-R	ATAGTCGACTTATGGGTTACGAATATTCCA
FLAG-HA2-F	TGAATAAGCTTTCAGTTGAAAGAACTTTTCTAC
FLAG-HA2-R	CTTTGGTACCTTATATTTTTTCAAGTTTGAAC
Strep-HA3-F	AAAGTTAGGTACCCTAGTGATACTATTGATTTAG
Strep-HA3-R	CGTGTCGACTTAATTAGTAATATCTATATGC
HA1-FLAG-F	CATGCCATGGTAATCCAAAATTCATTAATA
HA1-FLAG-R	CGGGATCCTTACTTGTTCATCGTCATCCTTGTAGTCTGGGTTACGAATATTCCATTTTC
HA1-FLAG N285A-F	GATGATGCTCAGAAATGGAATATTCGT
HA1-FLAG N285A-R	TTTCTGAGCATCATCTCCATGATAATT
Strep-HA3 R528A-F	TATACAGCTCAAAGCCCTGATGTCCATG
Strep-HA3 R528A-R	GCTTTGAGCTGTATAGTAATGTGCACC
mGP2-Fc-F	GAAGATCTACCATGAAAAGGATGGTGGGTT
mGP2-Fc-R	CCGCTCGAGTGTAGTGTGGGGAGTCCCC
hGP2-Fc-F	CGCGGATCCACCATGGAAAGGATGGTGGGC
hGP2-Fc-R	CCGCTCGAGTCCATTCATGACACCGGGAGA
Umod-Fc-F	CGCAGATCTACCATGGGGATCCCTTTGACC
Umod-Fc-R	CGCGTCGACCTTGGACACTGAGGCCTGG
PrP ^C -Fc-F	CGGGATCCACCATGGCGAACCTTGGCTACT

Supplementary Table 1: Primers sequences used in this study (continued).

Primer	sequence (5'-3')
PrP ^C -Fc-R	CGCTCGAGGGATCTTCTCCCCTCGTAATAG
GST-mGP2-F	TCCAGGGGCCCCTGGGATCTACAATACATCAAGGTTATGGGAG
GST-mGP2-R	ATGCGGCCGCTCGAGTCATGTAGTGTGGGAGTCCCCTTC
GST-Umod-F	TCCAGGGGCCCCTGGGAAGTAACTCAACAGAAGCGAGAC
GST-Umod-R	ATGCGGCCGCTCGAGTCACTTGGACACTGAGGCCTGGAC
<u>Used for qPCR</u>	
<i>Actb</i> -F	GAGCGCAAGTACTCTGTGTG
<i>Actb</i> -R	AACGCAGCTCAGTAACAGTCC
<i>Gapdh</i> -F	TGTGTGCGTCGTGGATCTGA
<i>Gapdh</i> -R	TTGCTGTTGAAGTCGCAGGAG
<i>Gp2</i> -F	AACTGCTATGCCACCCCTTC
<i>Gp2</i> -R	TCGGCTTCTGAGGACACAC
<i>Umod</i> -F	CAGGTGTTTCCAGGACAGAGG
<i>Umod</i> -R	CATTCAGAACACCGTCTCGC
<i>Prnp</i> -F	GGTCCCTTTGATGGAGTCTGTC
<i>Prnp</i> -R	TGTGGATGCTCTAGCTATCCCA
<u>Used for mGP2 transfection</u>	
mGP2-F	GAAGATCTACCATGAAAAGGATGGTGGGTT
mGP2-R	CCGCTCGAGGAACAGTAGAGCCAGGAAGAC