Supplementary material

N-glycosylation of rotavirus NSP4 protein affects viral replication and pathogenesis

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Supplementary Figure S1. N-glycosylation patterns of recombinant NSP4 in MA104 cells.

MA104 cells were infected with rsSA11, rsSA11-Y9P, rsSA11-N18A, or rsSA11-Y9P-N18A at an MOI of 10 in the absence of trypsin. Cell lysates were collected at 10 h post-infection, followed by treatment with Endo-H or distilled water.



Supplementary Figure S2. Co-localization of NSP4 and an ER marker. Pearson's correlation coefficient and Mander's overlap were calculated. Data are expressed as the mean \pm SD from 7–10 cells. Statistical significance was determined using a t-test, with p < 0.05 considered significant (*p < 0.05).



Supplementary Figure S3. Validation of co-immunoprecipitation results using HT29. HT29 cells were infected with rsSA11 or rsSA11-Y9P-N18A at an MOI of 5. Cell lysates were collected at 18 h post-infection and subjected to co-immunoprecipitation using anti-calnexin and anti-NSP4 antibodies. Molecular weight (kDa) is shown on the right. (IP = immunoprecipitation; WCL = whole cell lysate; CNX = calnexin).



Supplementary Figure S4. Co-immunoprecipitation assay using MA104. MA104 cells were infected with rsSA11 or rsSA11-Y9P-N18A at an MOI of 5. Cell lysates were collected at 10 h post-infection and subjected to co-immunoprecipitation using anti-calnexin and anti-NSP4 antibodies. Molecular weight (kDa) is shown on the right. (IP = immunoprecipitation; IB = immunoblotting; WCL = whole cell lysate; CNX = calnexin).