

Supplementary Materials

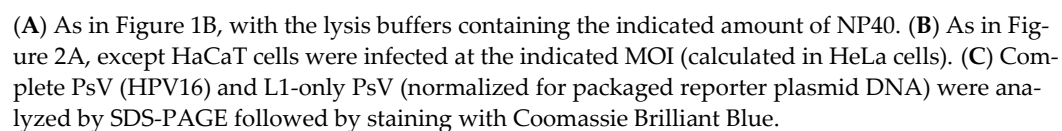
Human Papillomavirus L2 Capsid Protein Stabilizes γ -Secretase during Viral Infection

Table S1. qRT-PCR for packaged reporter genomes.

Virus	Genomes/ μ L
WT HPV16	7.11E+05
Null	1.08E+06
Notch	2.79E+05
WT HPV16	3.73E+07
BR6	2.00E+06
BR11	1.85E+07
WT HPV16	2.14E+07
L46A	1.01E+07
Q47L	2.43E+07
Y48L	1.49E+07
WT HPV16	1.28E+06
G49L	1.79E+06
S50L	2.54E+06
M51L	5.75E+06

Table S2. List of antibodies used in this study.

Antigen	Source	Catalogue Number	Dilution	Application
FLAG M2	Sigma	F3165	1:5000	IP
FLAG HRP	Sigma	A8592	1:1000	WB
HPV16 L1	BD Biosciences	554171	1:1000	WB
PS1	Cell Signaling	5643S	1:1000	WB
			1:500	IP
PEN2	Cell Signaling	8598	1:500	WB
APH1	ThermoFisher	PA1-2010	1:1000	WB
			1:500	IP
NCT	Santa Cruz	Sc-376513	1:1000	WB



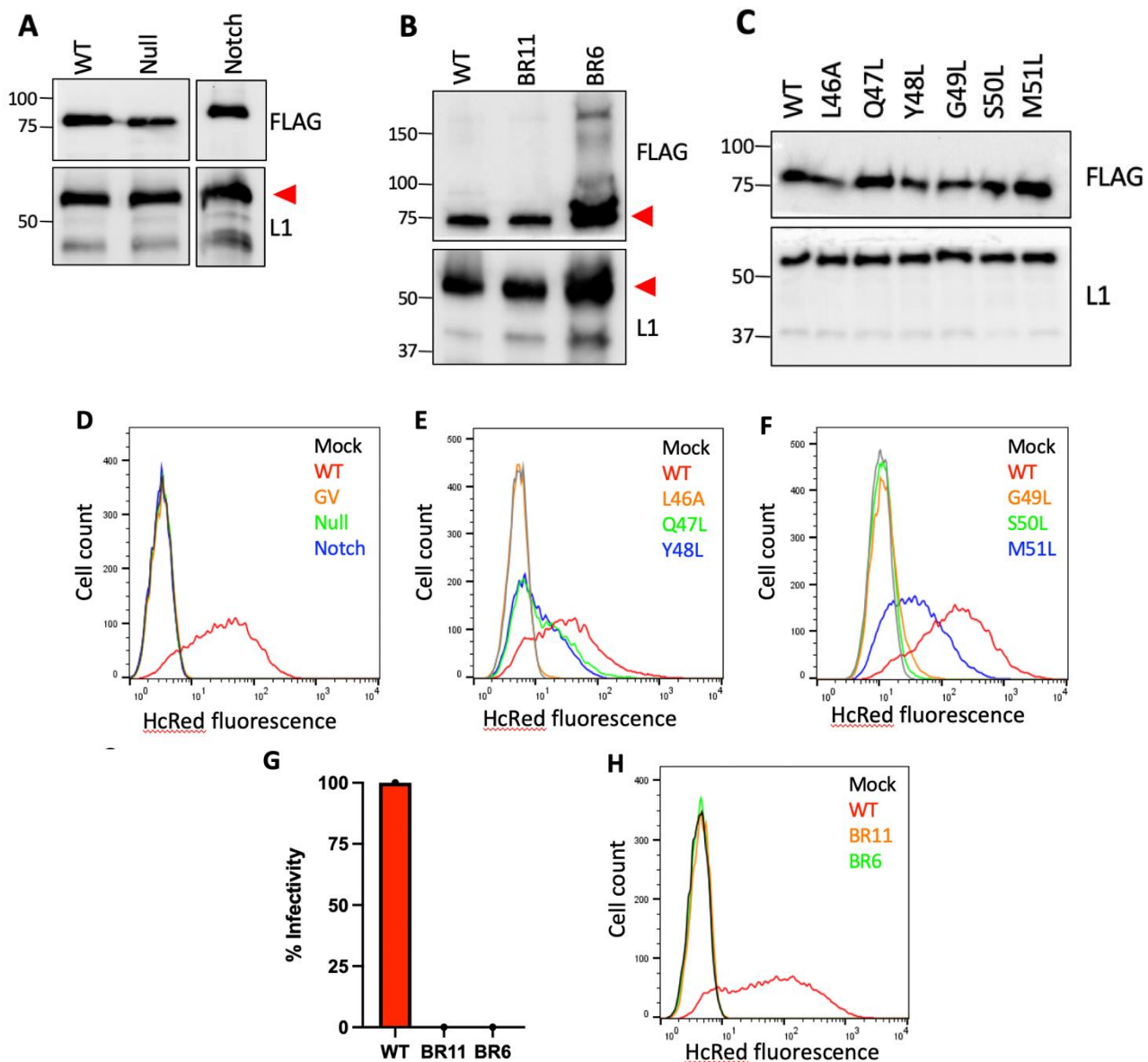


Figure S2. Related to figure 3.

(A) Normalized amounts of gradient-purified wild-type, L2 TM Null, and L2 TM Notch PsVs were analyzed by SDS-PAGE followed by western blotting for L1 or the FLAG tag on the C-terminus of HPV L2. Red arrowhead shows position of L1. All samples were analyzed on the same SDS-PAGE gel; an extraneous lane was removed between the Null and Notch mutants. (B) As in (A), except with the indicated TM mutants. Red arrowhead shows position of L1 and FLAG-tagged L2. (C) As in (A), except with the indicated TM point mutants. (D-F) Flow cytometry histograms of HcRed reporter gene expression used to generate the graphs shown in Figure 3C, 3D, and 3E, respectively. (G) Infection was analyzed as in Figure 3C for the indicated PsV. (H) Flow cytometry histograms are shown as in panel D-F.