

FIGURE S1

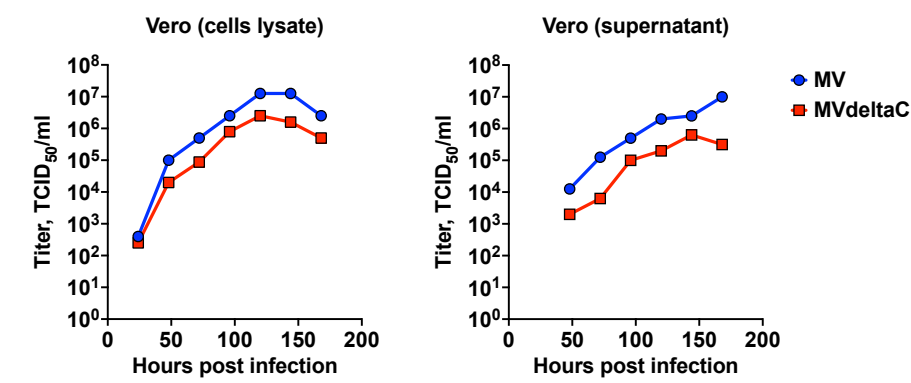


Figure S1. Growth kinetics of MV and MVdeltaC on Vero cells. Cells were infected at MOI 0.01 and incubated at 32°C. The data show reciprocal endpoint dilution titers of cell-associated (left) and cell-free (right) viruses.

MVdeltaC P5

Coverage depth (log X)

Genome position

iSNVs (%)

MVdeltaC P10

Coverage depth (log X)

Genome position

iSNVs (%)

Protein Domain Diagram:

N P/dC/V M F H L

1000 2000 3000 4000 5000 6000 7000 8000 9000 10000 11000 12000 13000 14000 15000

1830 (100% reads) acg ORF MVdC C: T

1863 (100% reads) taa stop

ORF MV Schwarz C: M S

original sequence atg tca

Figure S2. Sequencing analysis of MVdeltaC passages. Reads coverage and proportion of intrasample single nucleotide variants (iSNVs) obtained from next-generation sequencing of MVdeltaC at passages 5 (P5) and 10 (P10). A schematic of the measles virus genome is shown below using the same scale, with the two engineered mutations preventing C protein expression highlighted. No sequence changes were detected at either of these positions in P5 or P10 samples.

FIGURE S3

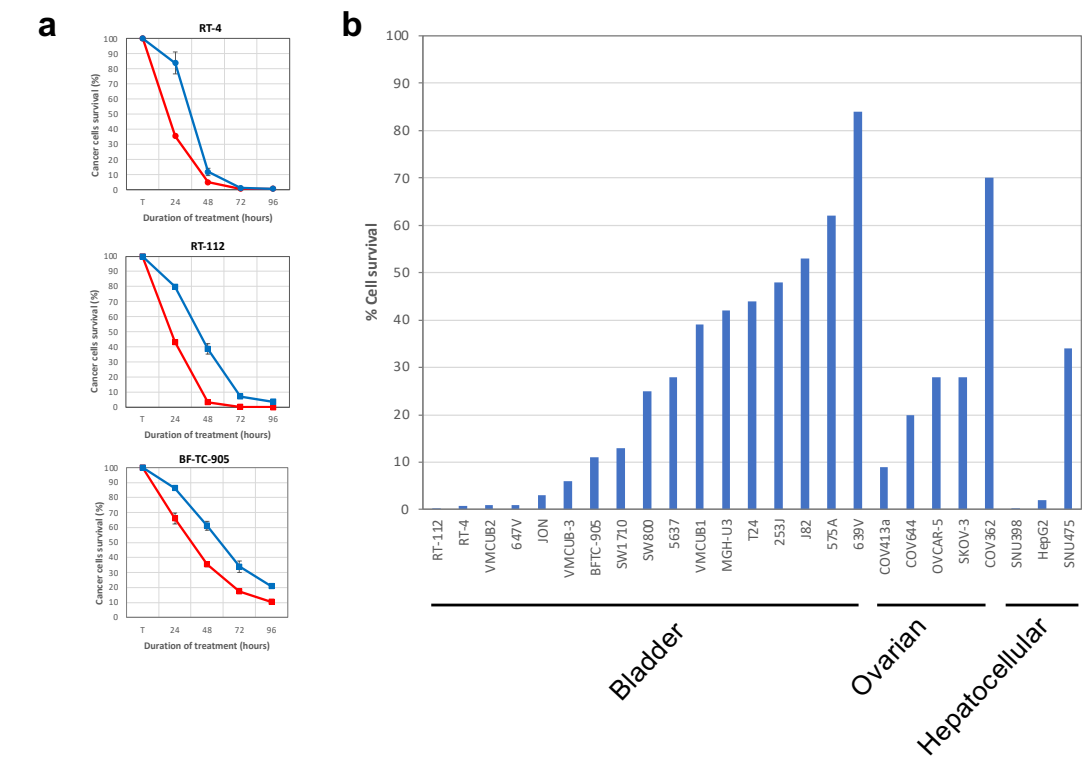


Figure S3. Infection and killing of human cancer cells by MVdeltaC. (a) Cell survival of three human bladder cancer cell lines after infection with MV (blue) or MVdeltaC (red) at MOI 5, measured by CellTiter-Glo assay; means \pm SD; measurements are in triplicate; data normalized to non-infected cells. (b) MVdeltaC killing of a series of human cancer cells of bladder, ovarian or hepatocellular origin as determined by CellTiterGlo assay (cancer cells survival after 72 hours of infection at MOI 5).

FIGURE S4

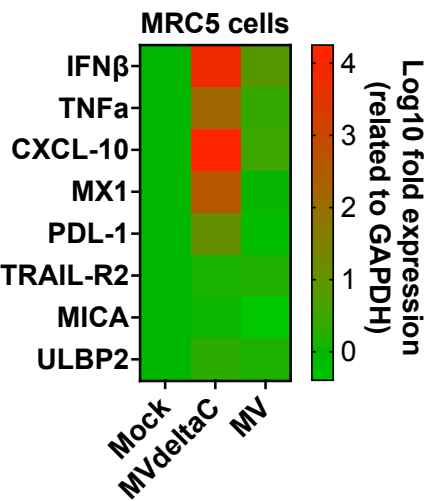


Figure S4. Gene expression analysis in infected MRC5 cells. Heatmap of differentially expressed genes in MRC5 cells after 24h incubation with MV, MVdeltaC (MOI 1) or PBS (Mock); experiment in triplicate, a gradient of colors from green to red represent log₁₀ fold change values from lowest to highest, respectively.

FIGURE S5

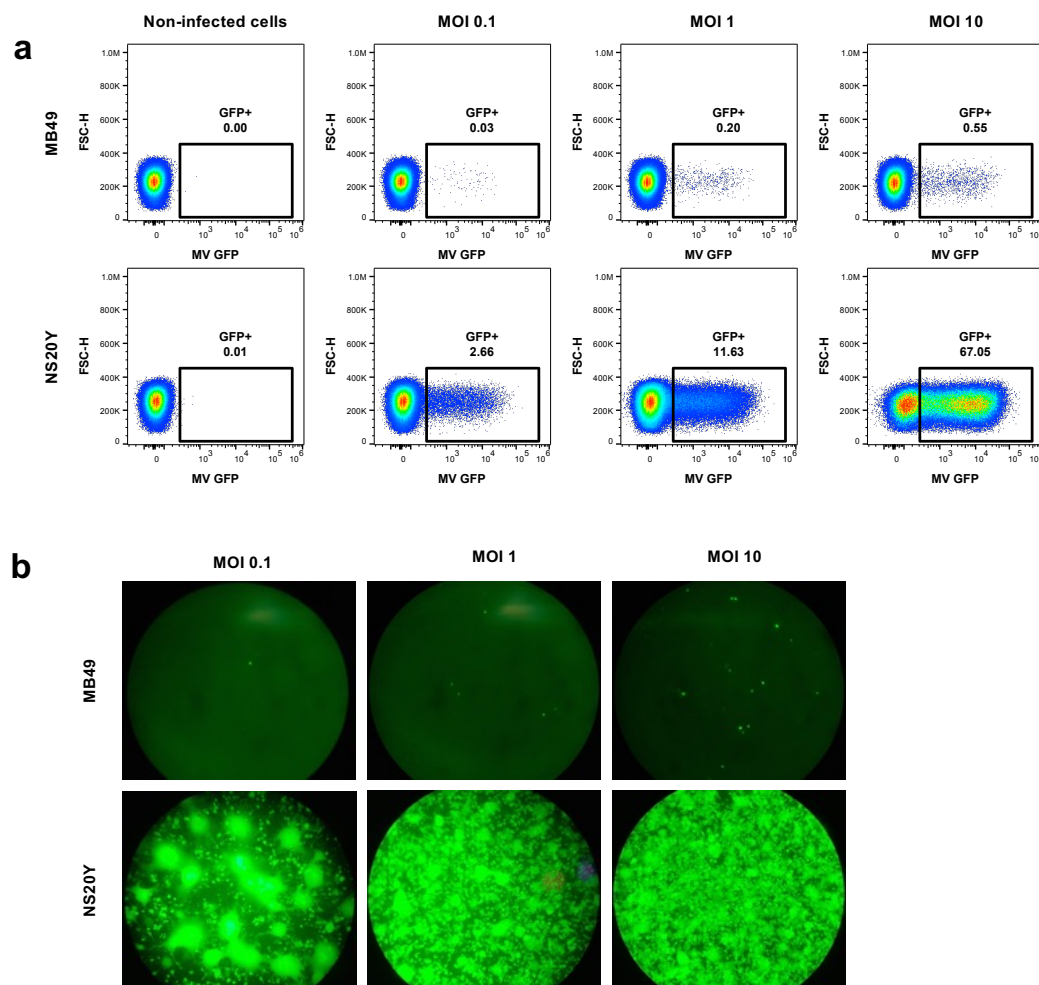


Figure S5. NS20Y cells are permissive to measles infection. Mouse neuroblastoma NS20Y cells and bladder carcinoma MB49 cells were infected with MV-eGFP at the indicated MOI and analyzed by (a) flow cytometry 24 hours post-infection, and (b) immunofluorescence microscopy 72 h hours post-infection.

FIGURE S6

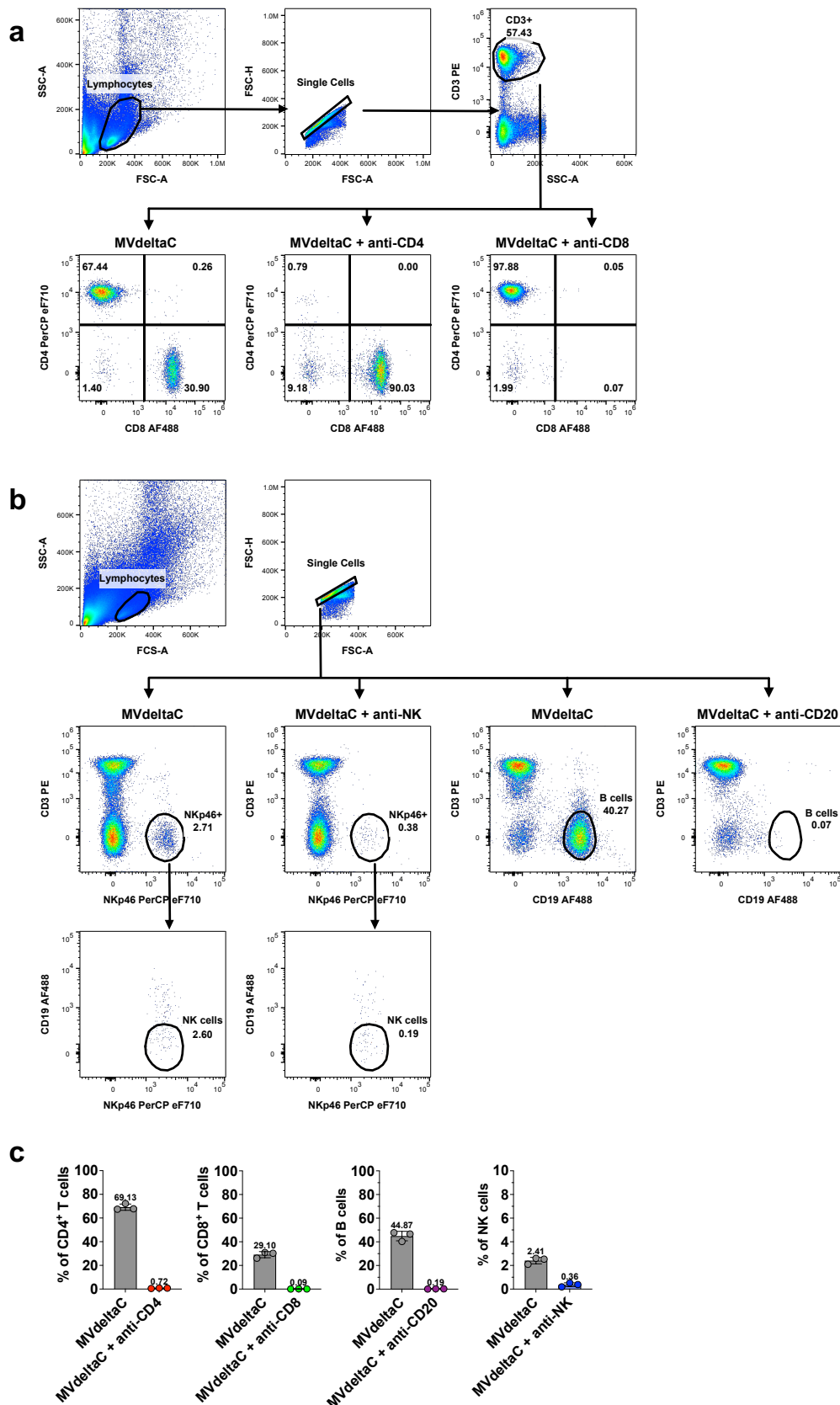


Figure S6. Flow cytometry analysis of immune cell populations in the blood of A/J mice depleted for CD4⁺, CD8⁺, B or NK cells. (a) Representative dot plots show the flow cytometry gating strategy to identify CD4⁺ (CD3⁺ CD4⁺ CD8⁻), CD8⁺ (CD3⁺ CD8⁺ CD4⁻) or (b) NK (CD3⁻ NKp46⁺ CD19⁻) and B (CD3⁻ CD19⁺) cell populations in the blood of treated mice. Mice were grafted with 5×10^6 NS20Y murine tumor cells on day 0. T cell depletion was achieved by i.p. injection of anti-CD4 or anti-CD8 mAbs (200 μ g in 100 μ L of PBS) on days 5, 6 and 7 after NS20Y cells injection. NK or B cell depletion was performed on days 5 and 7 using 50 μ g of anti-asialo GM1 or 200 μ g of anti-CD20 antibodies, respectively. Sustained immune cell depletion was achieved by weekly antibody administrations on days 14 and 21. Tumor therapy with MVdeltaC at 2×10^6 TCID₅₀ was performed on days 6, 9 and 12. Immune cell populations in the groups of A/J mice were analyzed on day 27. (c) Bar graphs show flow cytometry data for the three animals from each group, means \pm SD.

FIGURE S7

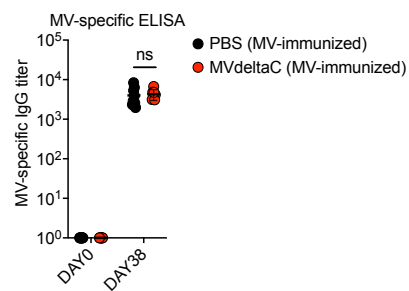
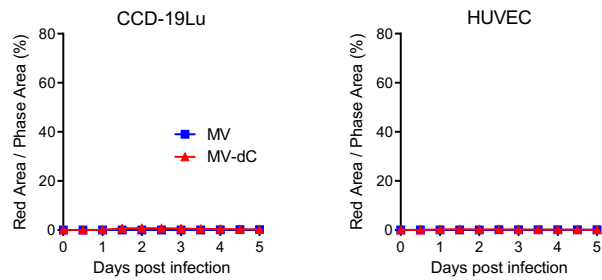


Figure S7. MV-specific IgG antibody responses in A/J mice following MV Schwarz vaccination. Mice were immunized i.p. with MV Schwarz (1×10^5 TCID₅₀) on day 0. Sera were collected at day 38 post-immunization and assessed for IgG antibody responses to MV antigens by ELISA. The data show the reciprocal endpoint dilution titers of MV-specific IgG antibodies in preimmune (day 0) and day 38 sera; each point represents an individual mouse; means \pm SD, n= 7-9 per group; unpaired two-tailed t test; ns, not significant.

FIGURE S8

a



b

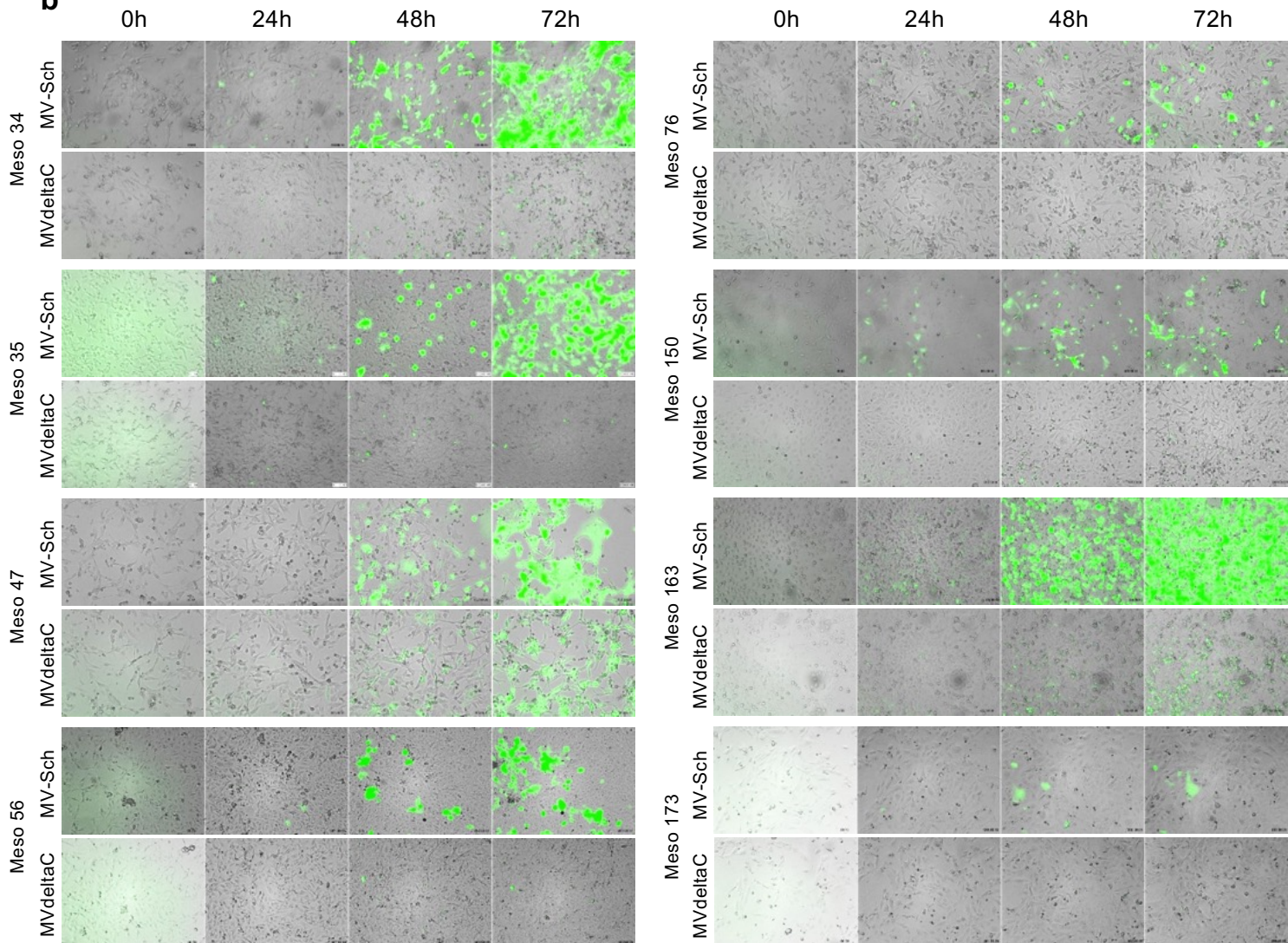


Figure S8. Infection of healthy and cancer human cells by MV or MVdeltaC. (a) Quantification of virus propagation in healthy lung fibroblasts (CCD19-Lu) or endothelial cells (HUVECs) infected with either MV-mCherry or MVdeltaC-mCherry. (b) Time-lapse microscopy analysis of mesothelioma cell lines infected with MV-eGFP or MVdeltaC-eGFP (MOI = 1).

VIDEOS MESO13

Figure S9. Infection of Meso 13 mesothelioma cells by MV or MVdeltaC.

Time-lapse microscopy analysis of mesothelioma cell lines infected with MV-eGFP (a) or MVdeltaC-eGFP (b) (MOI = 1).

FIGURE S10

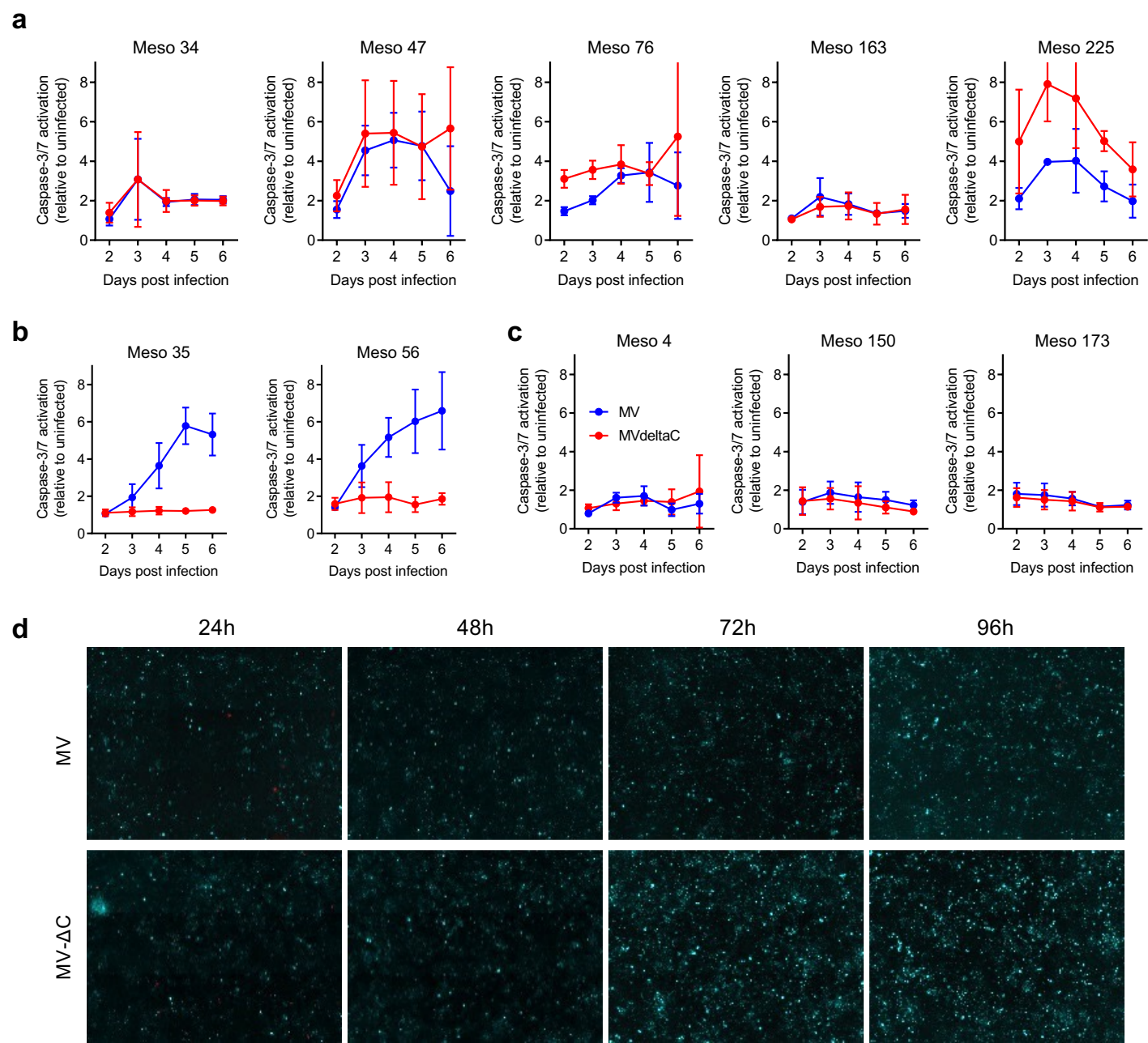


Figure S10. Activation of Caspases-3/7 in mesothelioma cells infected by MVdeltaC. (a-c) Luminescence-based quantification of caspase-3/7 activity in human mesothelioma cell lines. (a) MV-permissive, (b) MVdeltaC-resistant, and (c) MV-resistant cells. (d) Fluorescence microscopy of Meso 4 cells showing caspase-3/7 activation (cyan) after infection with MV-mCherry or MVdeltaC-mCherry (red).

FIGURE S11

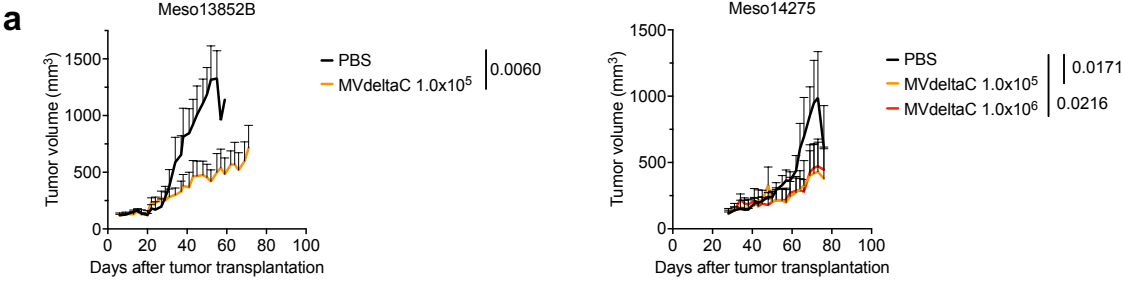
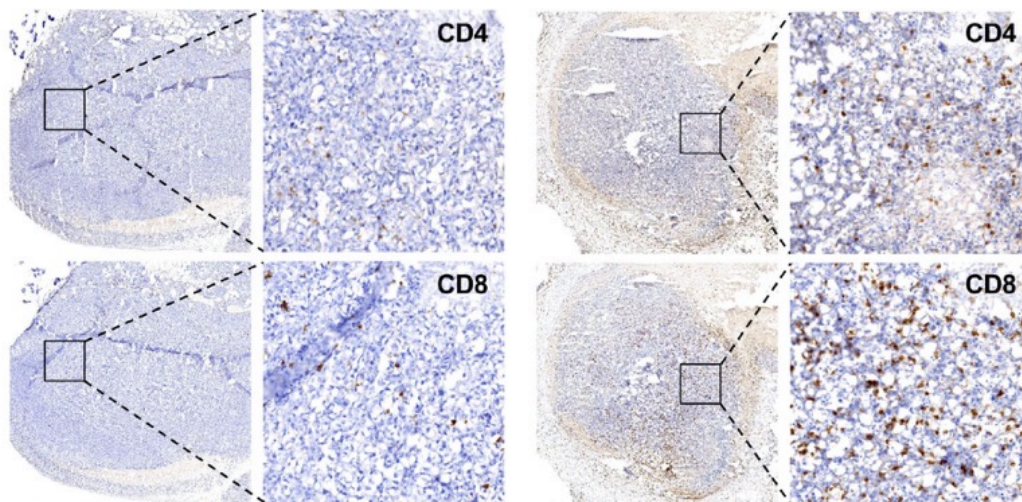


Figure S11. MVdeltaC inhibits human PDX growth in immunodeficient mice. Growth of mesothelioma PDX tumors in NMRI nu/nu mice after weekly MVdeltaC therapy at two doses; means \pm SEM, $n = 3$ mice per group; each group was compared to PBS group (unpaired one-tailed t test).



PBS

MVdeltaC

Cold Tumor

Hot Tumor

- Exclusion of CD8+ T cells and NK cells from the tumor
- Immunosuppressive immune cells in tumor (ie. Tregs)
- Poor prognosis and response to immunotherapy

- CD8+ T cells and NK cells are present in tumor
- Suppression of immunosuppressive cell types
- Improved prognosis

