

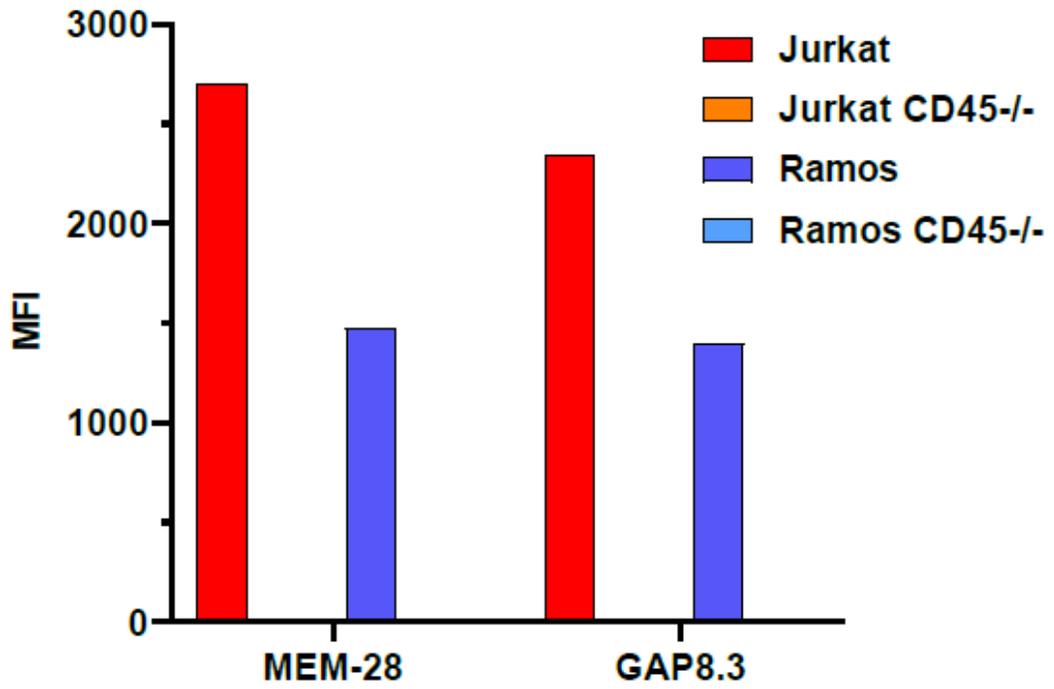
Supplementary Material

Inhibition of the B cell receptor signaling induced by the human adenovirus species D E3/49K protein

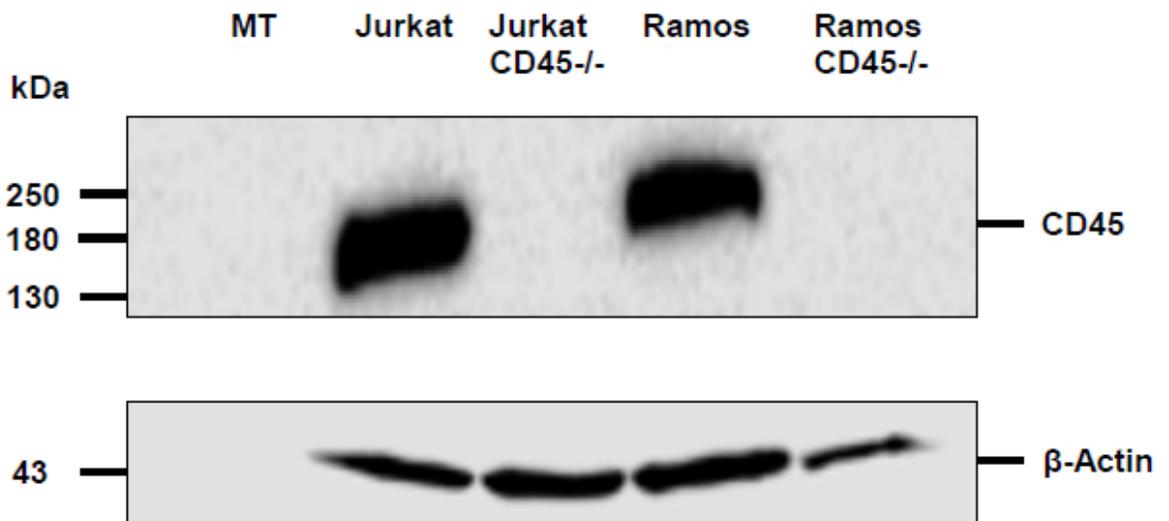
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A

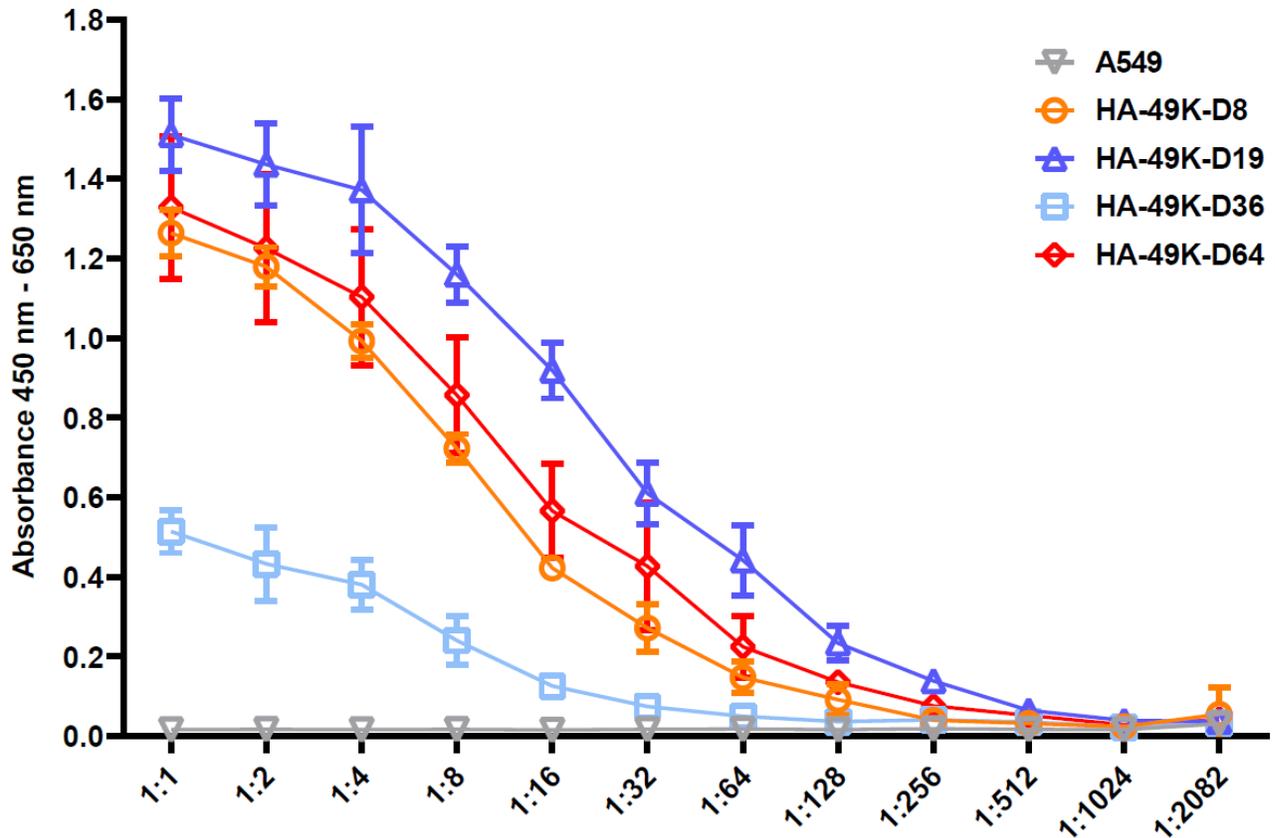


B



Supplementary Figure 1. Identification of CD45 levels of Jurkat and Ramos cell lines

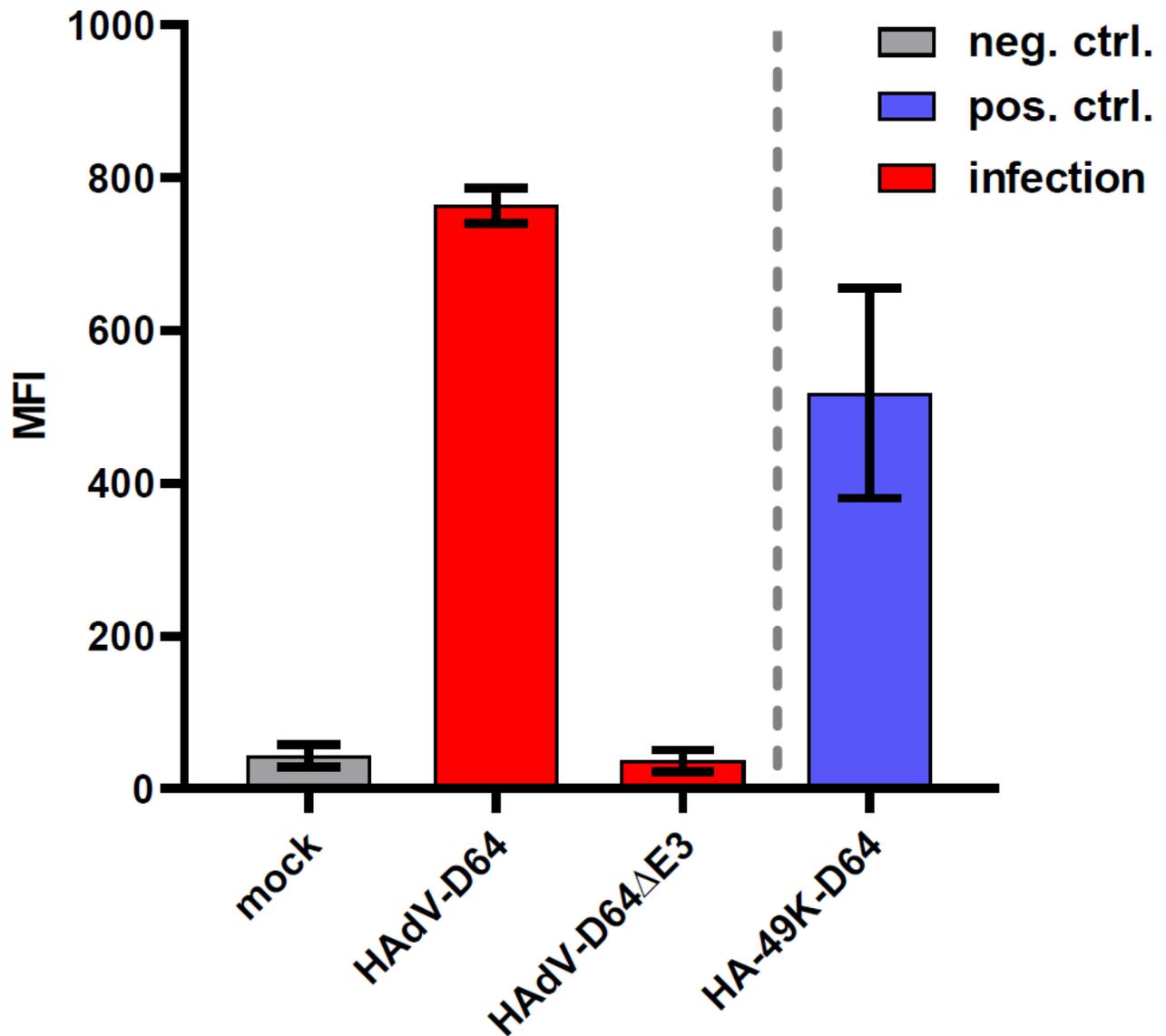
Flow cytometry measurements of CD45 cell surface expression levels for wild-type Jurkat (red) and Ramos cells (blue), and in CD45-deficient Jurkat (orange) and Ramos cells (light blue). CD45 levels were determined by staining with the two pan-specific α -human CD45 mAbs MEM-28 and GAP8.3. One representative experiment out of two is displayed (A). Cell lines were lysed and lysates were analyzed for CD45 expression using α -human/monkey CD45 (D9M8I) XP (Cell Signaling Technology) Ab. Detection of β -actin levels was used as loading control (B).



Supplementary Figure 2. Quantification of HA-49K levels in cell supernatants derived from the A549-based expression system

Cell culture supernatants from the A549-based expression system was generated as described above. HA-49K levels from supernatants of the individual producer cell clone for HA-49K of HA Δ V-D8 (orange), -D19 (blue), -D36 (light blue) and -D64 (red) were quantified via an α -HA based Sandwich ELISA. Supernatant from untransfected A549 were used as negative control (grey). Detection was based on its binding activity to its target receptor human CD45. 50 μ l of supernatants were used per sample and diluted as indicated on the X-axis. For the ELISA Nunc MaxiSorp™ flat-bottom plates were coated with 0.25 μ g/ml anti-HA Abs (both from Thermo Fisher Scientific) overnight at 4 $^{\circ}$ C, followed by blocking with 10 % goat serum (Merck AG) overnight. HA-49 supernatant dilutions were added to the plate followed by 1 μ g/ml hCD45-Fc in PBS solution (50 μ l per well) and incubated at least for 2 h at RT, followed by 30 min incubation with 0.16 μ g/ml of a horseradish peroxidase coupled α -human IgG. Detection was performed via a colorimetric using TMB substrate (Thermos Fisher

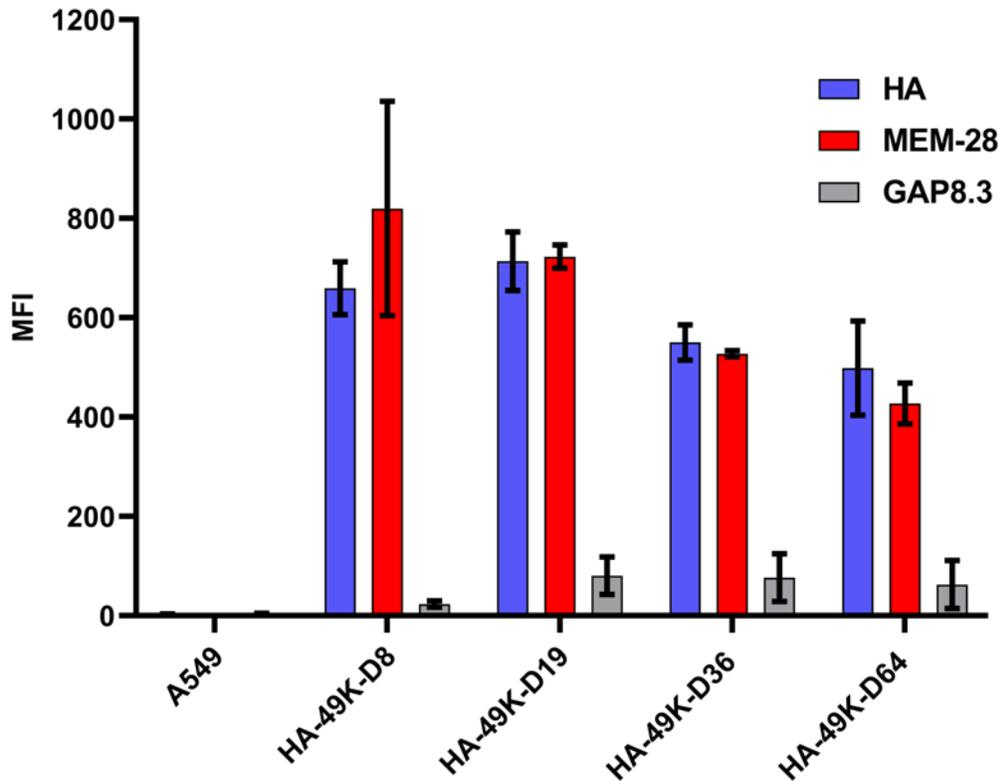
Scientific). Reaction was stopped at 20 min with 1 M sulfuric acid. Colorimetric absorbance measurement was executed at 450 nm and a reference wavelength of 620 nm, using the ELISA reader “Infinite® M PLEX” and the “software i-control™, 2.0” from the Tecan Trading AG.



Supplementary Figure 3. Detection of cell surfaced expressed E3/49K on HAdV-D64 infected cells

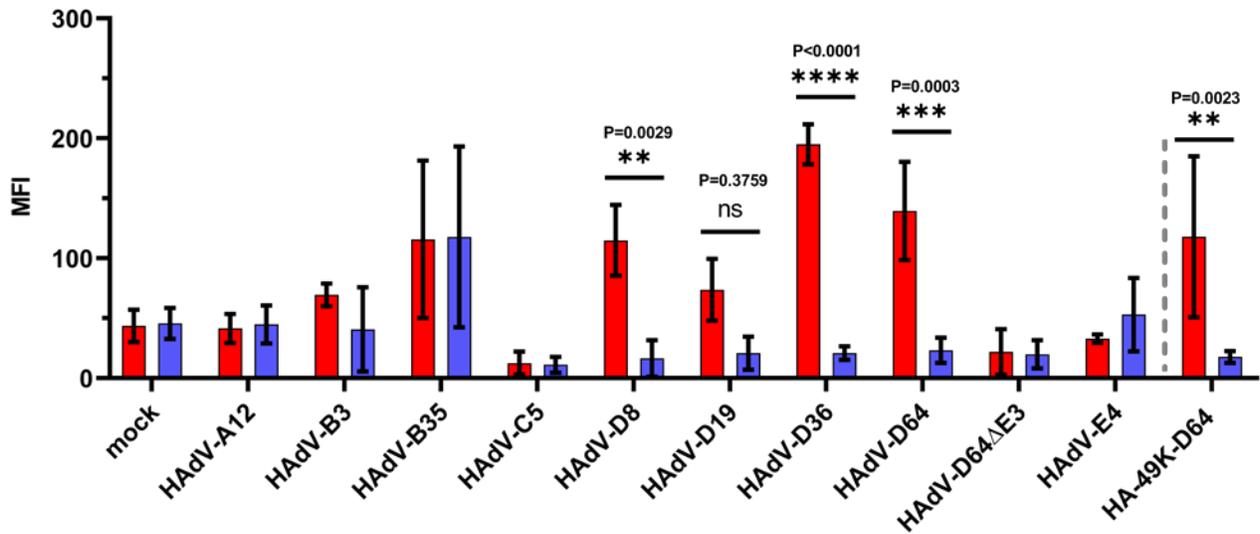
A549 cells were infected with an MOI of 5 with HAdV-D64 or HAdV-D64 Δ E3 (red). Cell surface resident E3/49K was detected with mAb 4D1 Ab by flow cytometric analysis. The mock infection

(grey) was performed as negative control, and a cell clone expressing stably HA-49K of HAdV-D64 was utilized as positive control (blue) in this infection assay (HA-49K of HAdV-D64 clone, separated from data derived from infection with grey dashed line). As expected, the presence of E3/49K was measured for HAdV-D64 infected cells as well as for the HA-49K stable transfected cell clone. The mean of 3 individual experiments including standard deviation is displayed.



Supplementary Figure 4. Measurement of the binding activity of soluble CD45 molecules to HA-49K transfectants

Stable transfectants expressing HA-49K of HAdV-D8, -D19, -D36 and -D64, and untransfected A549 cells, were treated with 0.5 µg per sample hCD45 Fc. Bound CD45 molecules were detected with α-human pan-CD45 MEM-28 (red) and GAP8.3 (grey) Abs. For comparison, HA-49K surface expression levels of the individual cell clone was measured by α-HA staining (blue). Untransfected A549 cells, being negative in all cases, were utilized as a negative controls. Bound CD45 to the different HA-49K types with MEM-28 was possible to similar levels compared to the HA-49K surface levels. In contrast to that, only low levels of bound CD45 molecules were detectable because of the competition of HA-49Ks with the GAP8.3 Ab for a potential similar binding site on CD4524. MEM-28 allows a precise measurement of a CD45 ECD binding activity. The mean of 3 individual experiments including standard deviation, presented as error bars, is shown.



Supplementary Figure 5. CD45-E3/49K interactions are blocked with GAP8.3 anti-CD45 Abs

A549 cells were infected with equal MOIs of 5 HAAdV-A12, -B7, -B35, -C5, -E4, -D8, -D19, -D36, -D64 and -D64ΔE3 viruses (MOI of 5). Infected cells were treated with +/- (red/blue) 0.5 μg hCD45-Fc per sample. Flow cytometric detection of bound CD45 molecules with usage of blocking α-CD45 GAP8.3 Abs was inhibited, because of the competition between the blocking Ab and E3/49K for the CD45-binding site²⁴. Mock infected cells as well as infection with HAAdV-D64 ΔE3 virus was performed as negative controls. The cell clone expressing stably HA 49K of HAAdV-D64 was applied as positive control (HA-49K -D64 clone, indicated with grey dashed line). The mean of 3 individual experiments including standard deviation, presented as error bars, is shown. Significant differences between +/- hCD45-Fc treatment were determined using two-way ANOVA test and indicated in the figure.