

Supporting Information

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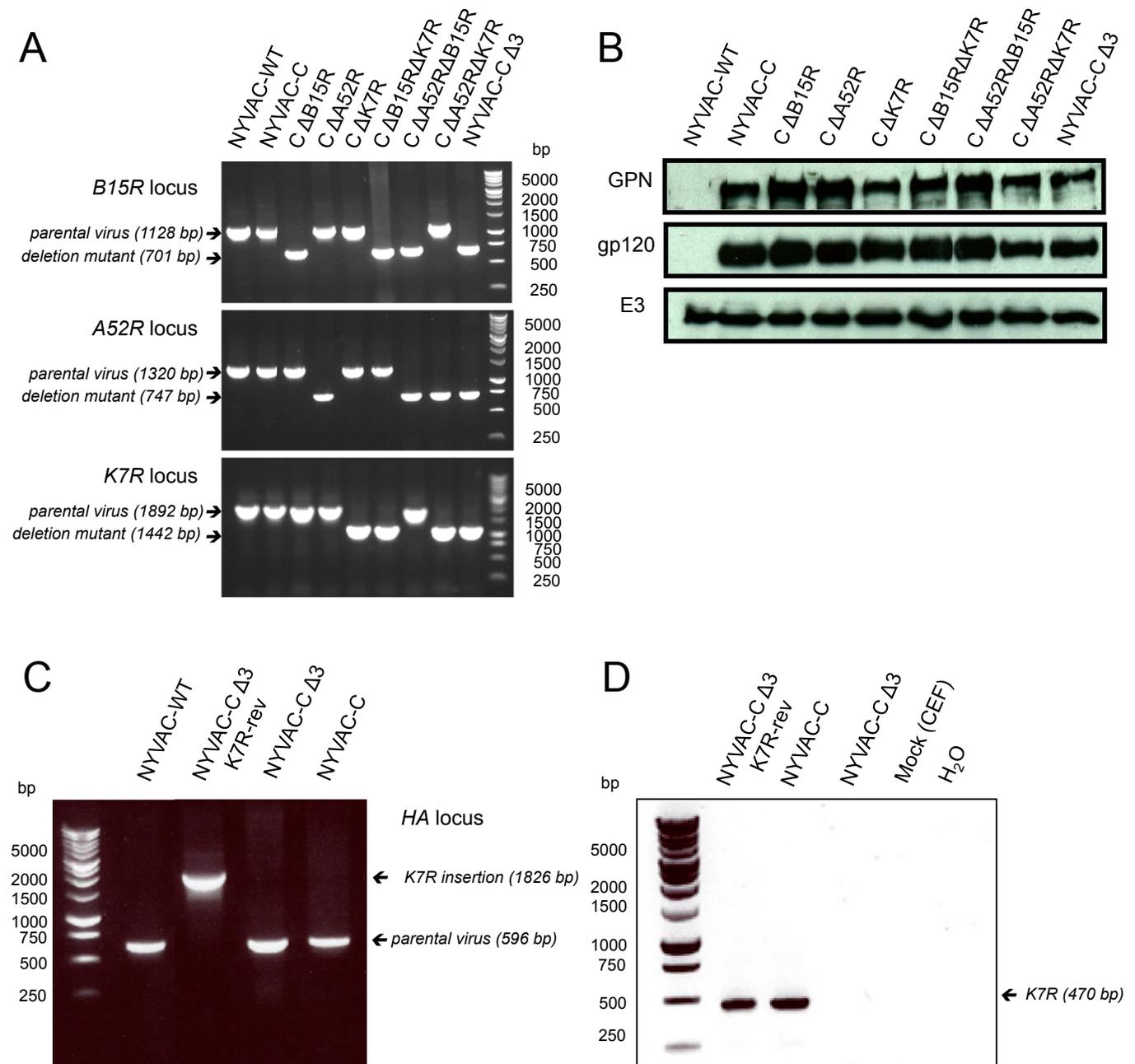


Fig. S1. Generation and characterization of NYVAC-C mutants. (A) PCR analysis to confirm A52R, B15R, and K7R gene deletion. Viral DNA was extracted from BSC-40 cells infected with the viruses as indicated (2 PFUs per cell, 24 h). (B) Western blot showing expression of HIV-1 antigens and E3 VACV protein in BSC-40 cells infected as in A. (C) PCR analysis to confirm K7R gene insertion. Viral DNA was extracted from BSC-40 cells infected with the viruses as indicated (1 PFU per cell, 24 h). (D) RT-PCR analysis to confirm K7R gene expression. RNA was extracted from CEF cells infected with the viruses as indicated (1 PFU per cell, 24 h).

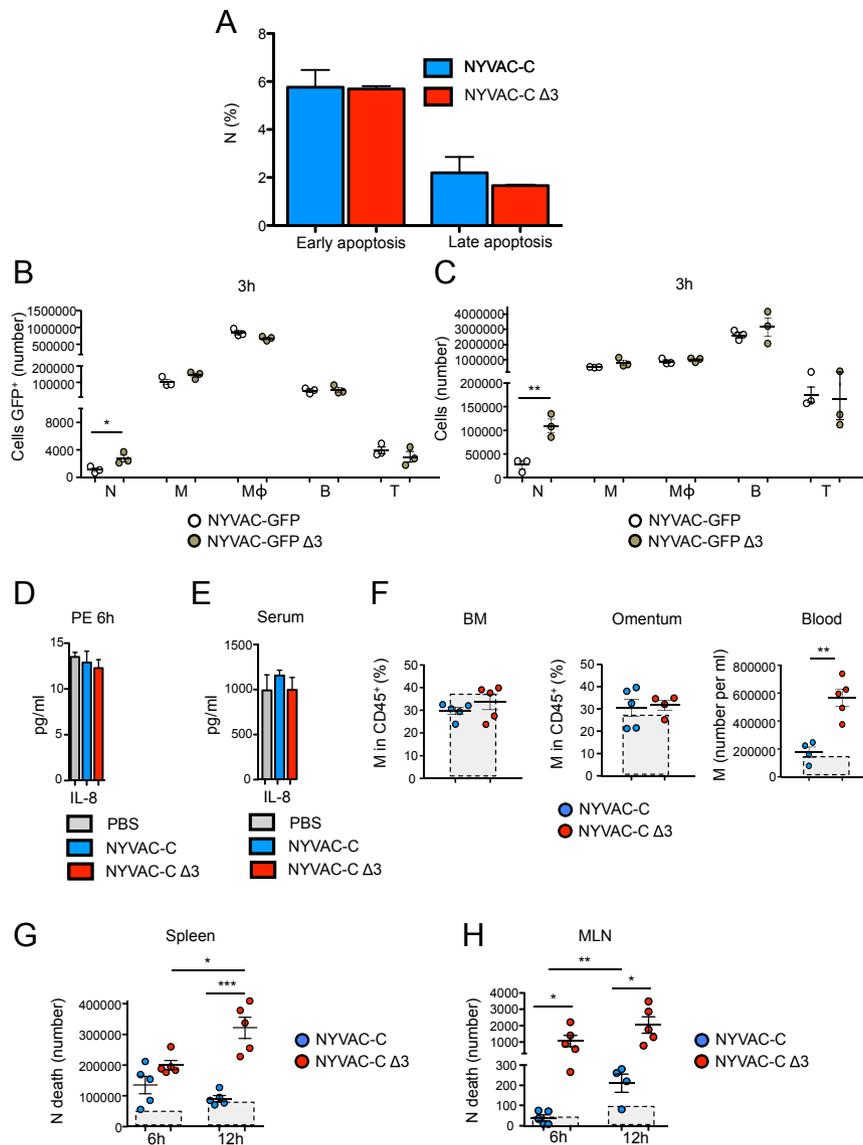


Fig. S3. In vitro and in vivo neutrophil cell death, PEC characterization after NYVAC-GFP or NYVAC-GFP $\Delta 3$ infection, and IL-8 secretion and monocyte trafficking after NYVAC-C or NYVAC-C $\Delta 3$ infection. (A) Percentages of early apoptotic (Annexin V⁺-PI⁺) and late apoptotic (Annexin V⁺-PI⁺) migrated neutrophils toward supernatants of NYVAC- or NYVAC-C $\Delta 3$ -infected (4 h) J774 macrophages. Values show mean \pm SEM of duplicates. (B) Absolute numbers of GFP⁺ neutrophils (N), monocytes (M), macrophages (M ϕ), B cells (B), and T cells (T) in the peritoneal cavity of NYVAC-GFP- or NYVAC-GFP $\Delta 3$ -injected (10^7 PFUs) mice at 3 h postinfection. Each point represents an individual mouse. (C) Absolute numbers of neutrophils (N), monocytes (M), macrophages (M ϕ), B cells (B), and T cells (T) in the peritoneal cavity of NYVAC-GFP- or NYVAC-GFP $\Delta 3$ -injected (10^7 PFUs) mice at 3 h postinfection. Each point represents an individual mouse. Shown are IL-8 levels at 6 h postinfection in peritoneal exudates (D) and serum (E) of PBS-, NYVAC-C-, or NYVAC-C $\Delta 3$ -injected mice ($n = 5$ per group). (F) Percentages of monocytes in BM and omentum, and total number of monocytes in blood at 6 h postinfection from NYVAC-C- or NYVAC-C $\Delta 3$ -injected mice. Boxes with dashed lines indicate absolute numbers of cells in PBS-injected mice. Each point represents an individual mouse. Total numbers of neutrophil cell death in spleen (G) and in MLNs (H) at 6 and 12 h postinfection from NYVAC-C- or NYVAC-C $\Delta 3$ -injected mice. Boxes with dashed lines indicate absolute numbers of cell deaths in PBS-injected mice. Each point represents an individual mouse. Graphs show mean \pm SEM. Data are representative of two independent experiments. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

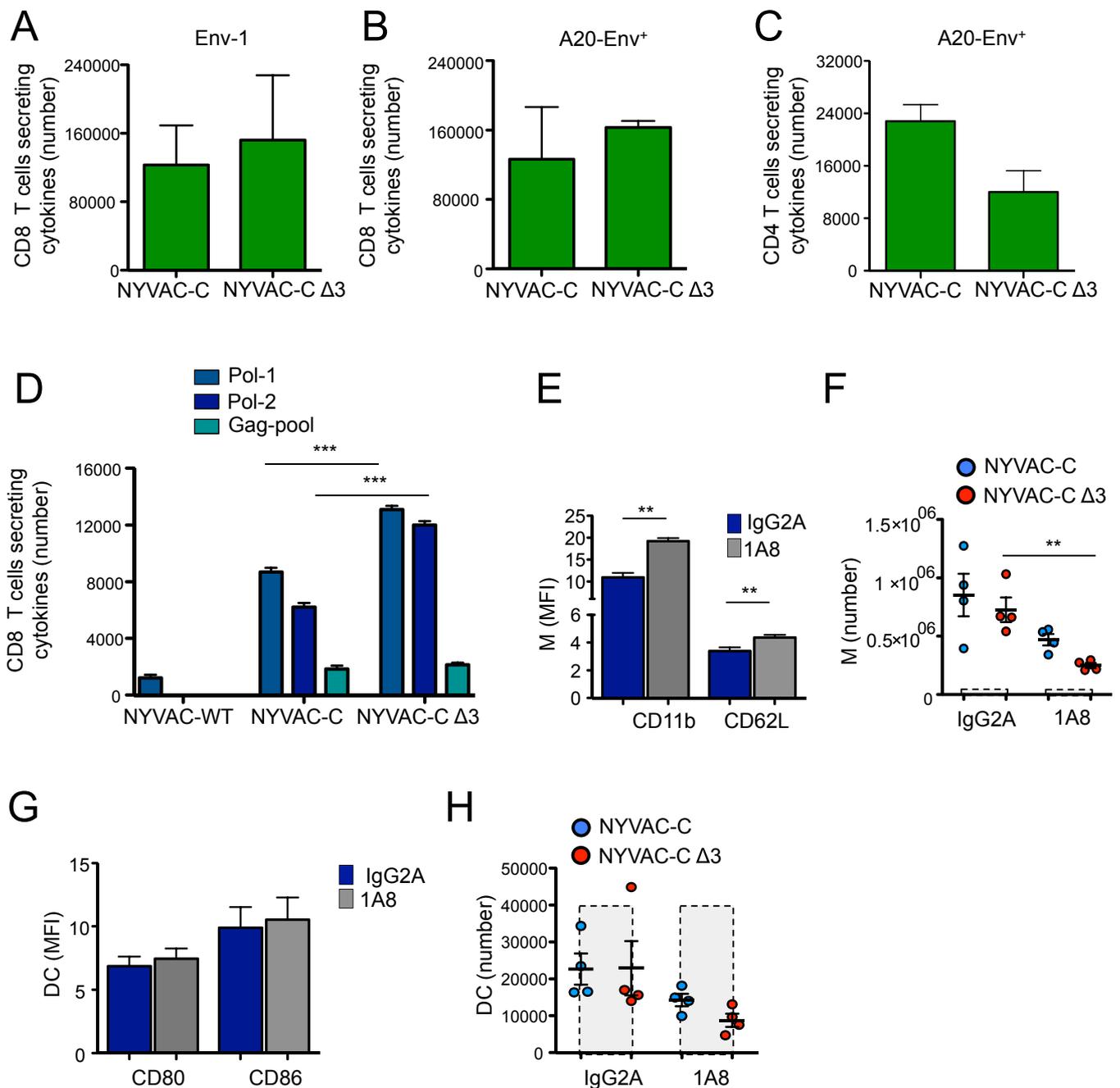


Fig. 55. Magnitude of the specific CD8 and CD4 T-cell response to HIV-1 antigens, and M or DC activation and absolute numbers in 1A8-pretreated/NYVAC-infected mice. Shown is the vaccine-induced HIV-1-specific T-cell response in mice ($n = 4$ per group) infected with 10^7 PFUs of NYVAC-C or NYVAC-C $\Delta 3$. The response was measured 11 d after the last i.p. immunization, after stimulation of splenocytes with Env-1 peptide (A) or with A20 Env⁺ (B and C), or after the last intramuscular immunization and after stimulation of splenocytes with HIV-1 peptides/pools (D). Total value (magnitude) is the sum of percentages of CD8 or CD4 T cells per spleen that secrete IFN- γ and/or TNF- α and/or IL-2 and/or CD107a. Values show mean \pm SEM of two independent experiments. Shown are the MFIs of indicated markers in monocytes (M) (E) and in DCs (G) at 6 h postinfection in NYVAC-injected and IgG2A- or 1A8-pretreated mice. Also shown are M (F) and DC (H) absolute numbers at 6 h postinfection in NYVAC-injected and IgG2A- or 1A8-pretreated mice. Boxes with dashed lines indicate numbers in PBS-injected mice. Graphs show mean \pm SEM; each point represents an individual mouse. ** $P < 0.01$, *** $P < 0.001$.

Table S1. Primers used for the deletion of *A52R*, *B15R*, and *K7R* genes and for the insertion of *K7R* gene

Genes	Flanks of genes	Name of primers	Sequence of primers
<i>A52R</i>	Left	LFA52R-Apa	TATATTGGGCCCTACTACGATTAA
<i>A52R</i>	Left	LFA52R-Sph	CTTTATGCATGCGGTGATCACACA
<i>A52R</i>	Repeated left	LF'A52R-Eco	ATTAGAGAATTCTACGATTAACGA
<i>A52R</i>	Repeated left	LF'A52R-Cla	CTTTATATCGATGGTATCACACA
<i>A52R</i>	Right	RFA52R-Cla	GAAATATCGATGAAAGTTAAAAG
<i>A52R</i>	Right	RFA52R-Bam	TCTGCCGGATCCAATGTAGTAATG
<i>B15R</i>	Left	LFB15R-Kpn-F	TTTATAGGTACCTTCTTTGAGGACT
<i>B15R</i>	Left	LFB15R-Kpn-R	GTTGGCGGTACCCTTACCAATTGCA
<i>B15R</i>	Repeated left	LF'B15R-Aat	TTCTTTGACGCTCTGTTTTCTGAAG
<i>B15R</i>	Repeated left	LF'B15R-Xba	GTTGGCTCTAGACTTACCAATTGCA
<i>B15R</i>	Right	RFB15R-Kpn	CGTATGGGTACCTGAGTTGTACATC
<i>B15R</i>	Right	RFB15R-Bam	GTGTCGGGATCCGAATTAGCATATT
<i>K7R</i>	Left	LFK7R-Aat	CAAAGGACGTCATCATCATTTTTTCACC
<i>K7R</i>	Left	LFK7R-Xba	GCTCTAGAGACTATCTCACAAAAG
<i>K7R</i>	Repeated left	LF'K7R-Eco	CGGAATTCATCATCATTTTTTCACC
<i>K7R</i>	Repeated left	LF'K7R-Cla	CCATCGATGACTATCTCACAAAAG
<i>K7R</i>	Right	RFK7R-Cla	CCATCGATTCTAGAAAAAATTGAATTG
<i>K7R</i>	Right	RFK7R-Bam	CGGGATCCAACAAGGGTTGG
<i>K7F</i>		K7RinternalFwd-Bam	CGCGGATCCATGGCGACTAAATTAG
<i>K7R</i>		K7RinternalRev-Not	AAGGAAAAAAGCGGCCGCTCAATTCAATTTTTTTC

Restriction enzyme cleavage sites are underlined.