

# *New Vaccine Strategies using cytomegalovirus*

Klaus Früh



**Traditional vaccines work by mimicking the immune responses elicited by a given pathogen, using a safe alternative of the pathogen.**

**However, traditional vaccine approaches struggle to elicit protective immune responses for infectious diseases that do not elicit protective immunity upon natural exposure**

### **Prophylactic Vaccines Needed**

- AIDS
- Tuberculosis
- Malaria
- Hepatitis C
- Herpesviruses

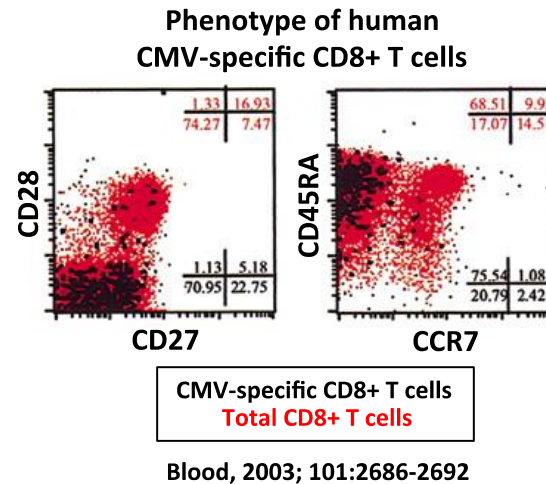
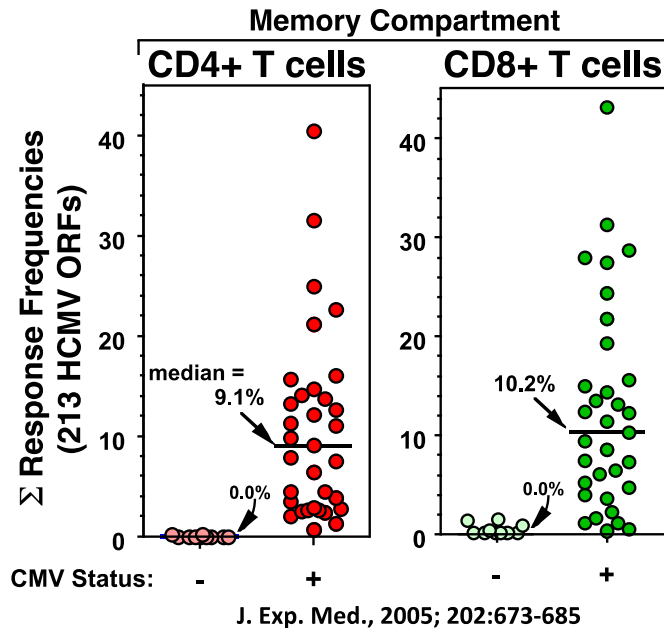
### **Therapeutic Vaccines Needed**

- Hepatitis B
- Human papillomavirus
- Herpesviruses
- Cancer

**Since we cannot mimic natural immunity, we need new approaches that elicit immune responses that are different and more efficacious than those induced by each pathogen**

# Using Cytomegalovirus (CMV) as a vaccine vector

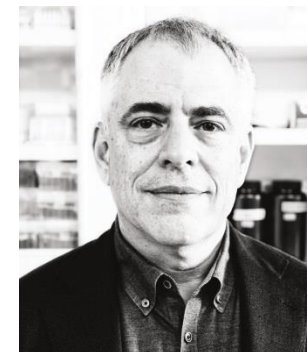
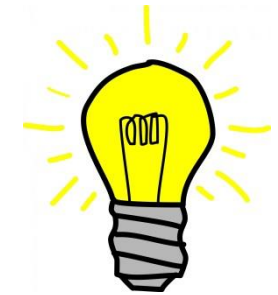
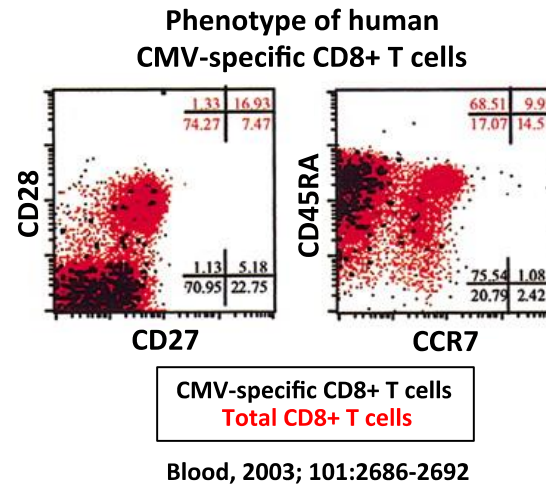
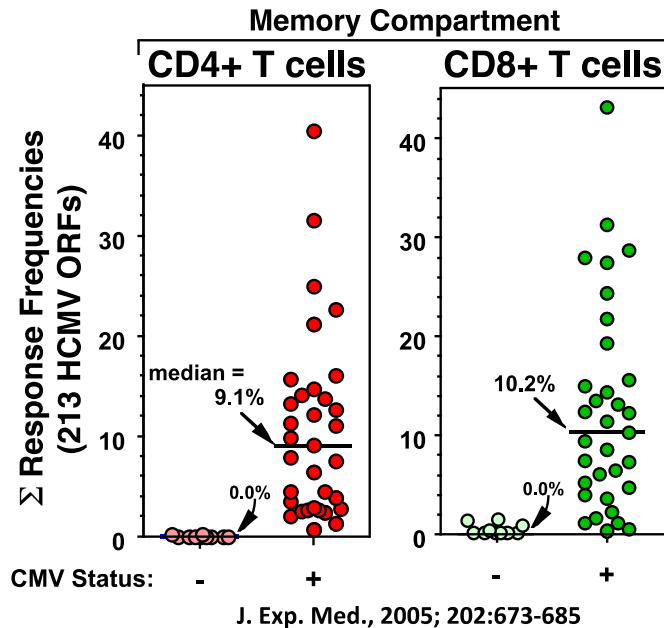
In the 1990s, it became evident that CMV is unique in its ability to elicit and maintain life-long, high frequency, effector (and tissue-resident) memory T cell responses



# Using Cytomegalovirus (CMV) as a vaccine vector

In the 1990s, it became evident that CMV is unique in its ability to elicit and maintain life-long, high frequency, effector (and tissue-resident) memory T cell responses

Can we use this unique immunological trait to target other pathogens?



Louis Picker

**Not only is the answer to this question “yes”, but the immunobiologic properties of CMV vectors turned out to be markedly different from any other known vector or non-vector vaccine modality . . .**

**In nonhuman primates, vectors based on rhesus cytomegalovirus:**

1. Elicit and indefinitely maintain high frequency, “effector memory” T cell responses\* in mucosal sites, lymphoid tissues and parenchymal organs
2. Efficiently re-infect & persist despite robust anti-CMV immunity
3. Manifest uniquely programmable CD8+ T cell immunogenicity with expanded breadth and conventional and/or novel epitope targeting.
4. Provide protection against SIV, TB and Malaria
5. Maintain immunogenicity even with profound attenuation (spread-deficiency and tropism restriction)

**\*Little to no antibodies**

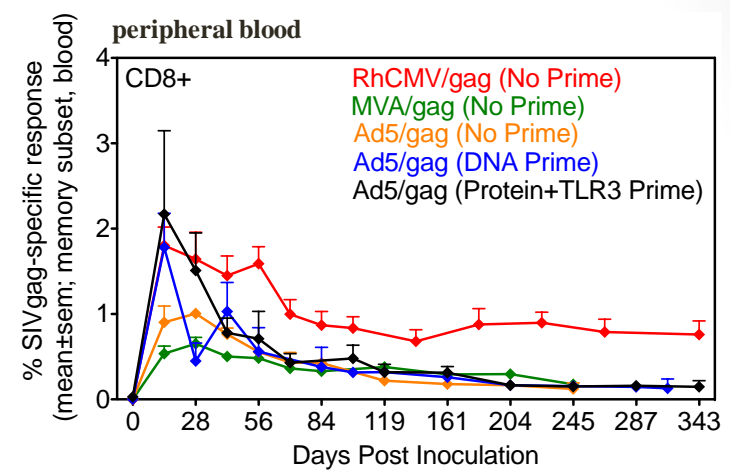
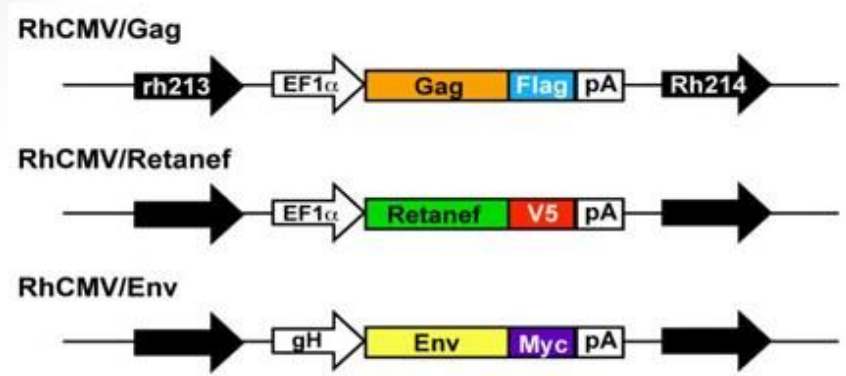
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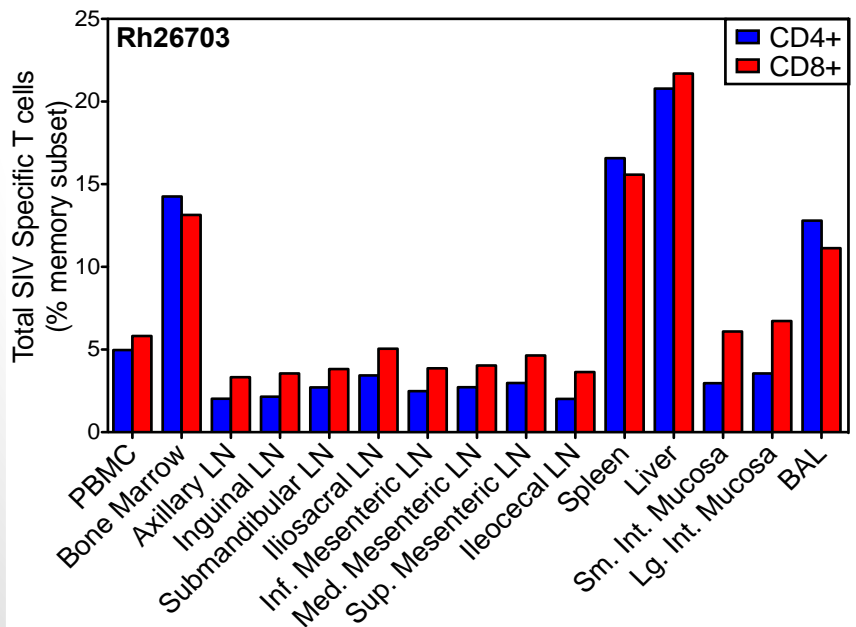
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# Inserting SIV antigens into RhCMV results in very high frequency SIV-specific T cells responses especially in tissues

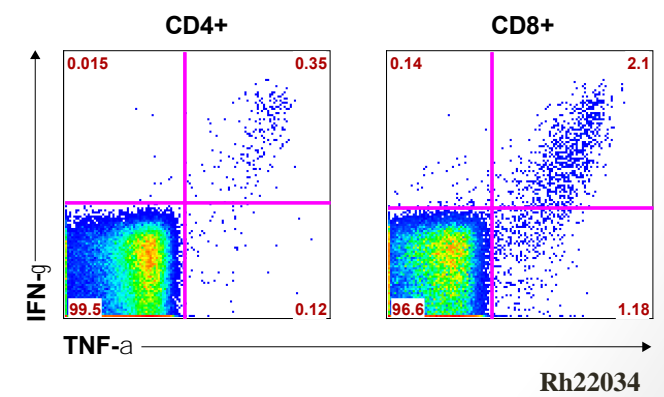


## Necropsy analysis 700 days post RhCMV/SIV vector vaccination



## T cell responses are maintained indefinitely

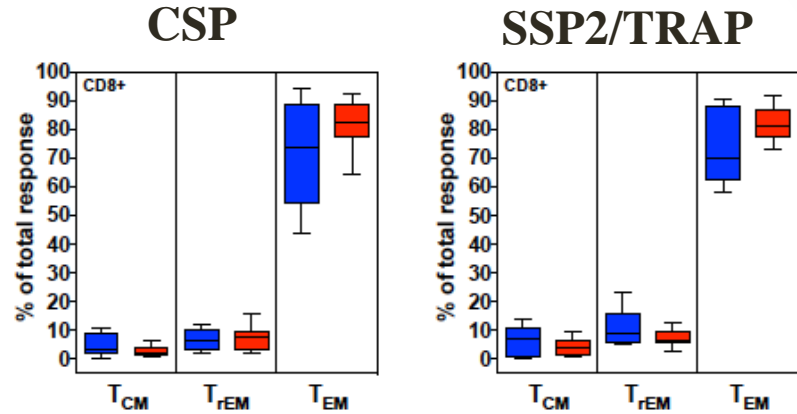
SIVgag-specific T cell responses in blood >8 years post RhCMV/gag vaccination





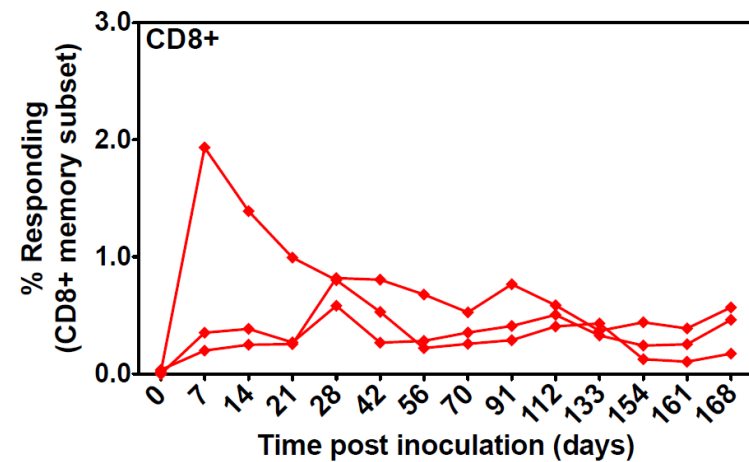
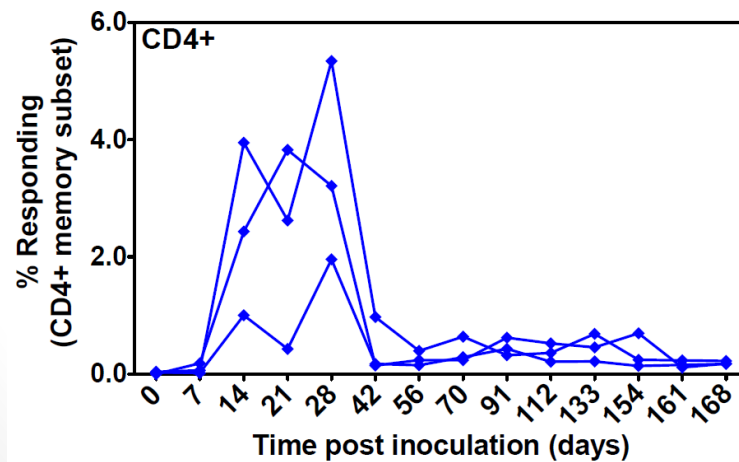
# CMV vectors are able to elicit effector memory responses to many different pathogens

- Mycobacterium Tuberculosis
- Plasmodium knowlesi
- Human Papillomavirus
- Ebola Virus
- Influenza Virus
- Hepatitis B Virus
- Herpes Simplex Virus



... even to cancer (=self) antigens

Peripheral blood T cell responses to Rhesus PAP induced in Rhesus monkeys inoculated with 68-1 RhCMV/RhPAP





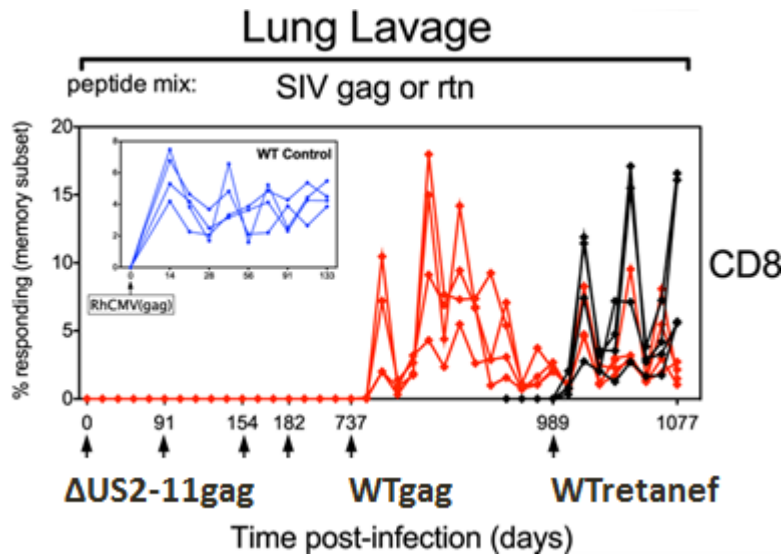
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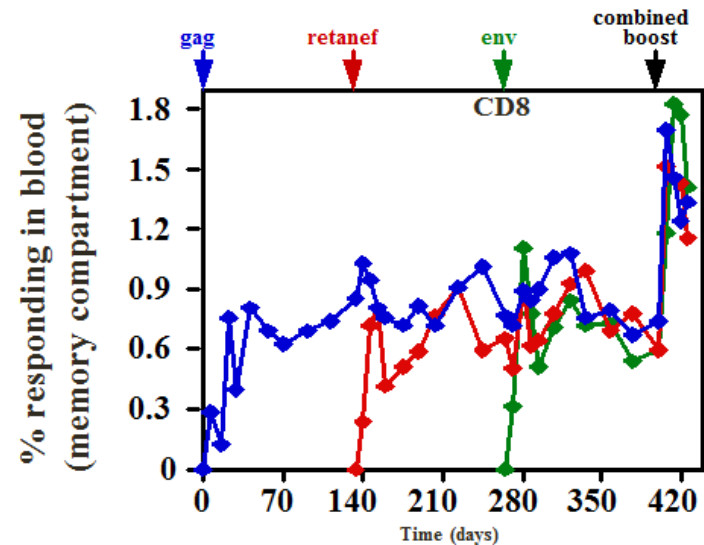
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# Viral inhibitors of antigen presentation (homologs of US2, US3, US6, US11) allow CMV-based vectors to be used repeatedly in CMV-immune recipients



*Science, 2010*



*Nature Medicine, 2009*

**10 PFU sufficient for subQ reinfection!**

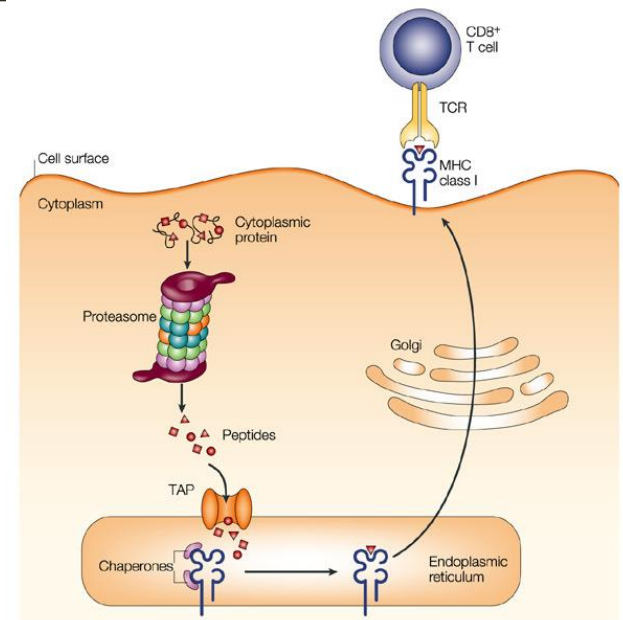
**This feature permits testing RhCMV-based vaccines in CMV-positive animals + humans**

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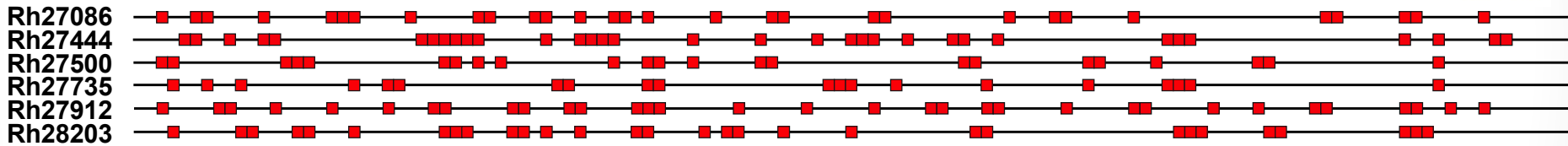
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# Viral infections and conventional vaccine vectors elicit CD8+ T cells that recognize peptides presented by polymorphic MHC-Ia molecules

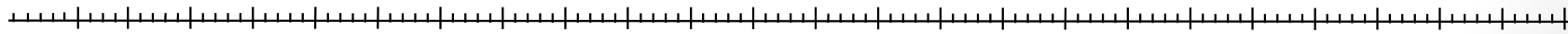
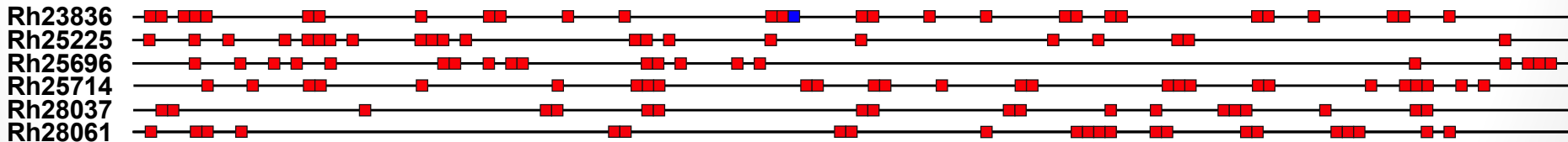


Nature Reviews | Immunology

## MVA/gag vector-vaccinated:



## SIVmac239-infected (plateau-phase plasma viral load <10,000 copies/mL):



Consecutive antigen insert 15mers (11 a.a. overlap)

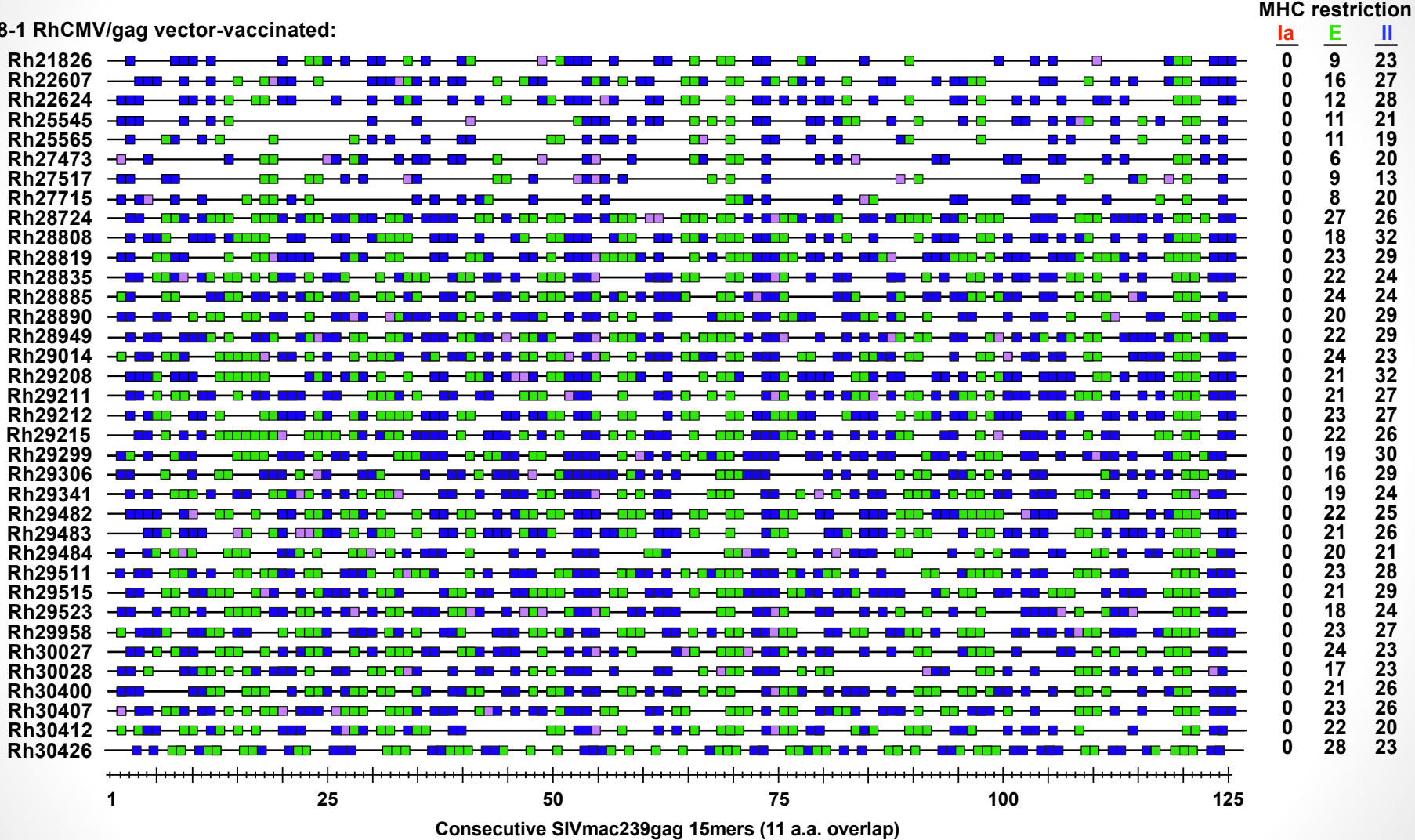
■ MHC-Ia restricted    ■ MHC-II restricted

Each square represents the location of a peptide recognized by CD8+ T cells within SIVgag

The color represents the restriction element as determined by blocking antibodies or peptides

# RhCMV-based vectors elicit CD8+ T cells that recognize peptides in the context MHC-II or the non-polymorphic MHC-Ib molecule MHC-E

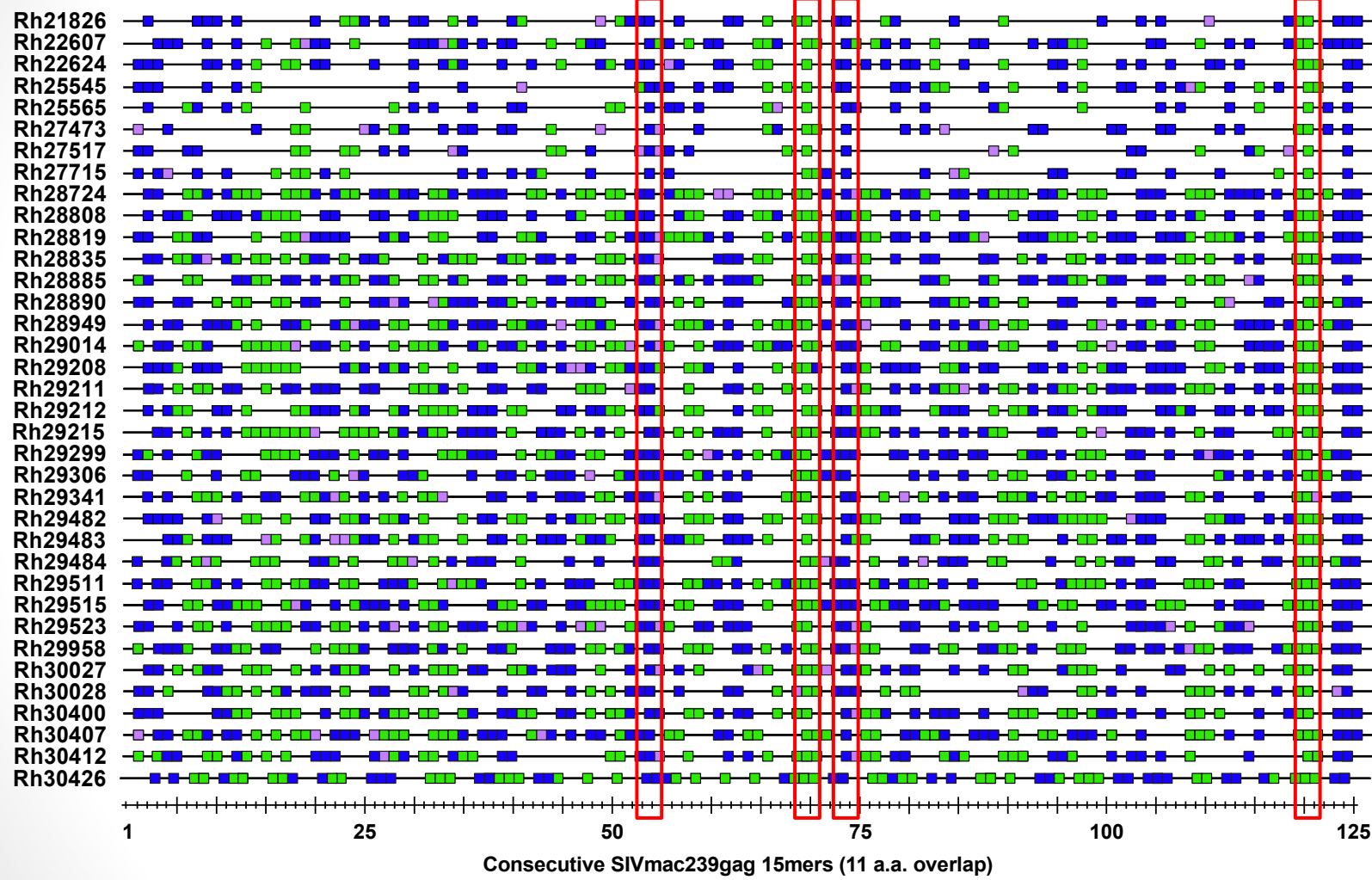
68-1 RhCMV/gag vector-vaccinated:



# RhCMV-based vectors elicit CD8+ T cells that recognize peptides in the context MHC-II or the non-polymorphic MHC-Ib molecule MHC-E

... and some peptides are recognized in every animal

68-1 RhCMV/gag vector-vaccinated:



MHC restriction

	Ia	E	II
Rh21826	0	9	23
Rh22607	0	16	27
Rh22624	0	12	28
Rh25545	0	11	21
Rh25565	0	11	19
Rh27473	0	6	20
Rh27517	0	9	13
Rh27715	0	8	20
Rh28724	0	27	26
Rh28808	0	18	32
Rh28819	0	23	29
Rh28835	0	22	24
Rh28885	0	24	24
Rh28890	0	20	29
Rh28949	0	22	29
Rh29014	0	24	23
Rh29208	0	21	32
Rh29211	0	21	27
Rh29212	0	23	27
Rh29215	0	22	26
Rh29299	0	19	30
Rh29306	0	16	29
Rh29341	0	19	24
Rh29482	0	22	25
Rh29483	0	21	26
Rh29484	0	20	21
Rh29511	0	23	28
Rh29515	0	21	29
Rh29523	0	18	24
Rh29958	0	23	27
Rh30027	0	24	23
Rh30028	0	17	23
Rh30400	0	21	26
Rh30407	0	23	26
Rh30412	0	22	20
Rh30426	0	28	23

■ MHC-Ia restricted

■ MHC-II restricted

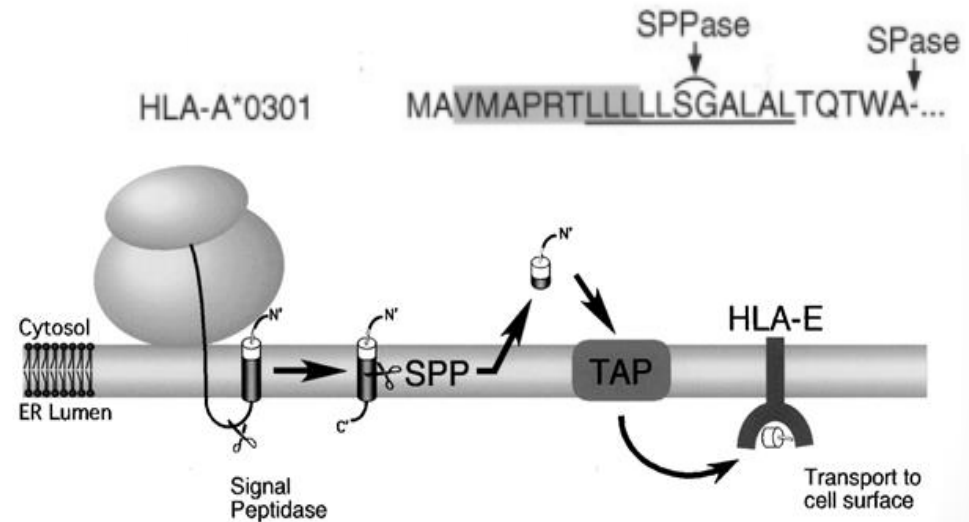
■ MHC-E restricted

■ Indeterminate

## Supertopes

# MHC-E is generally not involved in antigen presentation to T cells but presents peptide VL9 from MHC-I leader sequences as a “self” signal to NK cells

MHC-E is loaded with VL9  
TAP dependent



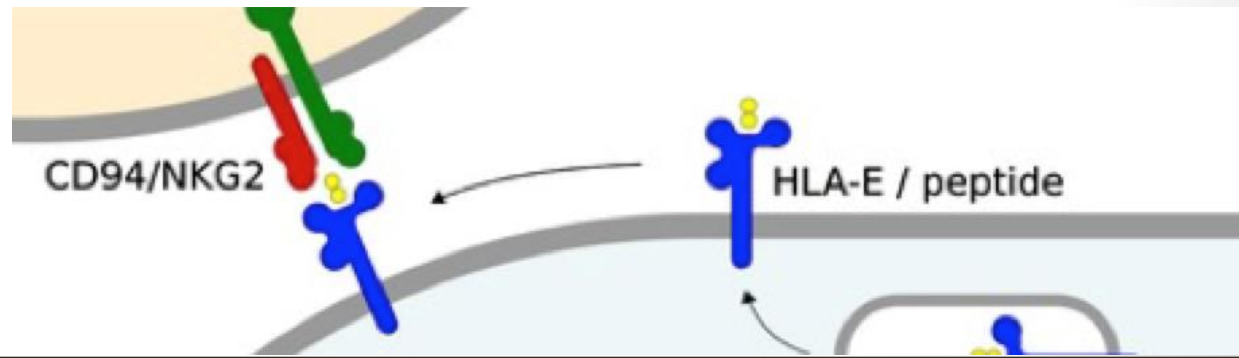
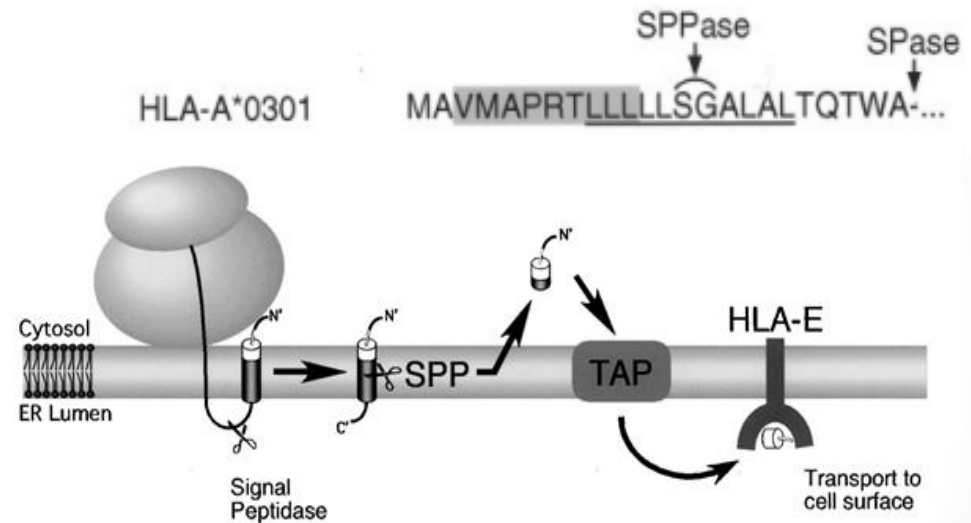


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MHC-E is loaded with VL9  
TAP dependent

MHC-E/VL9 complex represents  
a self-signal for NK cells

NKG2A: inhibitory receptor

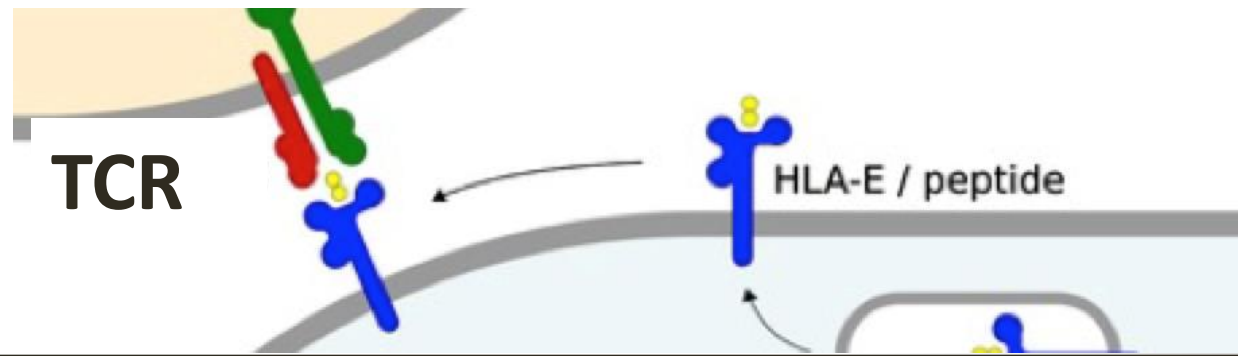


Importantly, although CMV is unique in its ability to elicit CD8+ T cells that recognize peptides in the context of MHC-E, other viruses (e.g. SIV, HIV, HBV) as well as cancer cells can be recognized by MHC-E-restricted CD8+ T cells.

This suggests that not all of MHC-E is loaded with VL9 peptide and that non-VL9 loading (or VL9 exchange) is a common occurrence in MHC-E expressing cells.

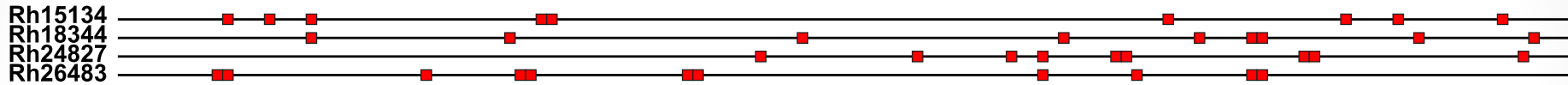
HLA-E is upregulated by chronic pathogens (e.g. HIV, HCMV, HCV) and on many cancers

The non-polymorphic nature of HLA-E offers an entirely new vaccine modality

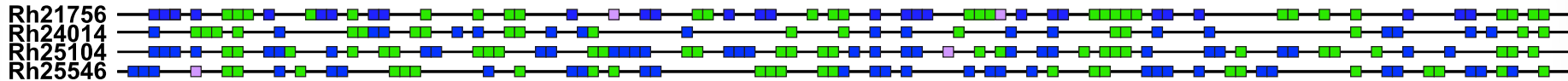


# Unconventionally targeted CD8+ T cell responses (to CMV proteins) are not found in naturally CMV infected monkeys (or humans)

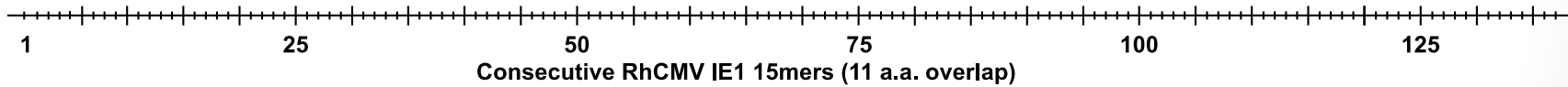
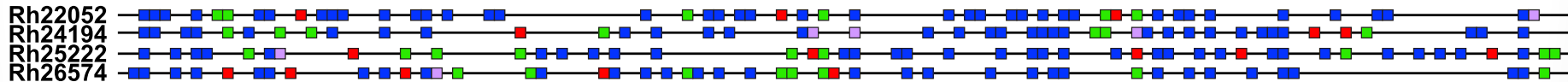
Naturally RhCMV-infected RM (infection with colony-circulating, true wildtype RhCMV; no vaccination):



Strain 68-1 RhCMV/gag vector-vaccinated RM that were RhCMV negative prior to vaccination:

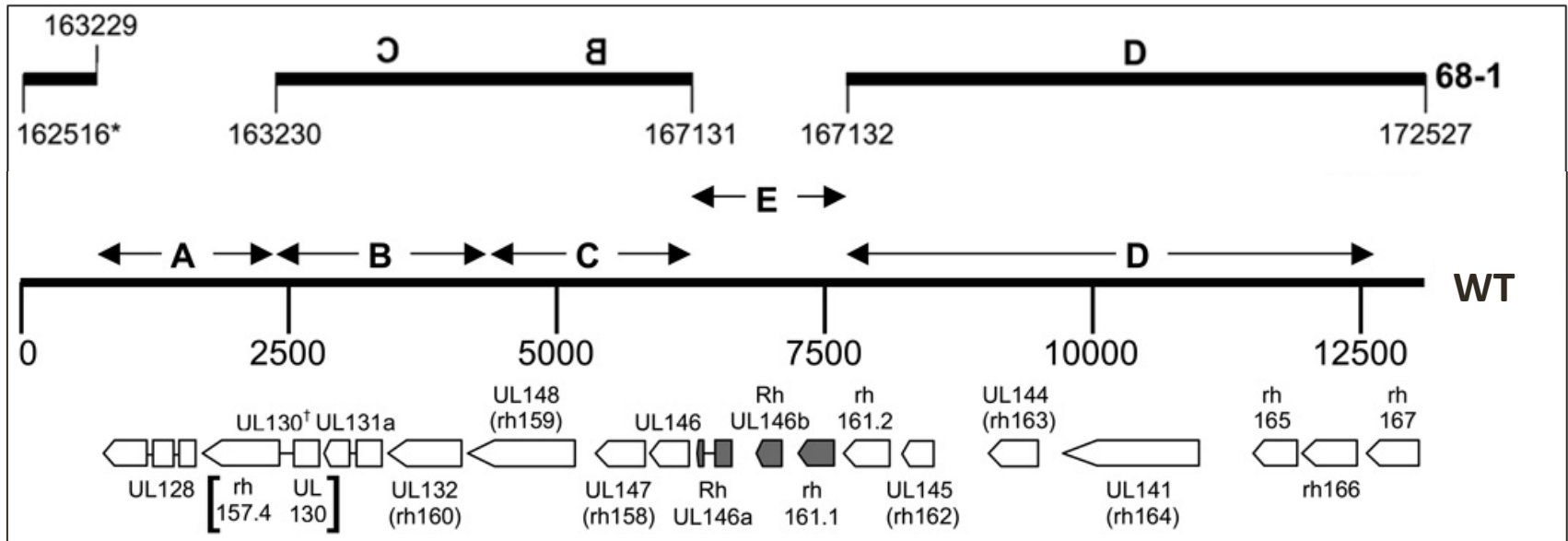


Strain 68-1 RhCMV/gag vector-vaccinated RM that were naturally RhCMV-infected (e.g. with wildtype RhCMV) prior to vaccination:



## Only in animals given RhCMV vectors derived from strain 68-1

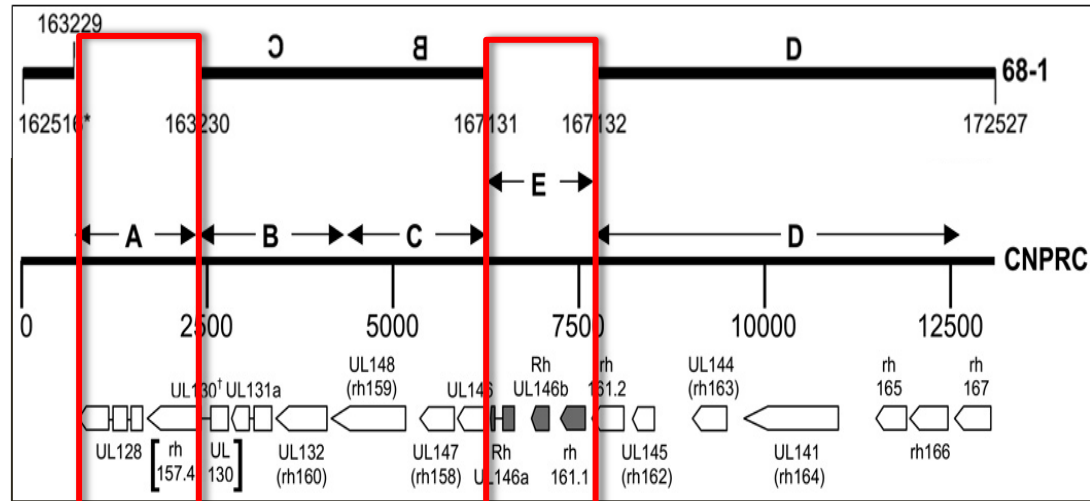
# RhCMV 68-1 is a fibroblast-adapted strain with gene deletions and inversions



Oxford Virology 2008

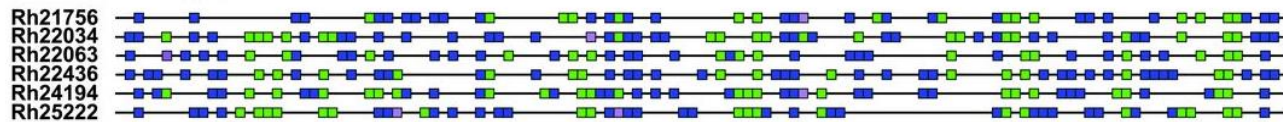
Which of these spontaneous genetic modifications enable RhCMV 68-1 to elicit unconventional CD8<sup>+</sup> T cells?

# Repair of deletions switches the CD8+ T cell responses from unconventional to conventional



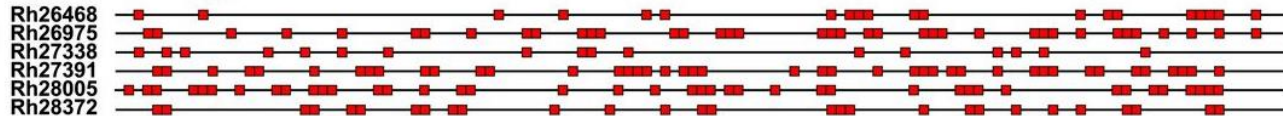
Oxford Virology 2008

Strain 68-1 RhCMV/gag vector-vaccinated:



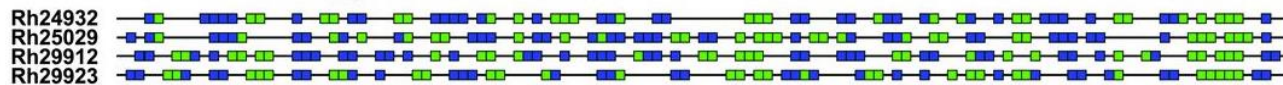
	Ia	E	II
Rh21756	0	13	21
Rh22034	0	15	24
Rh22063	0	14	24
Rh22436	0	14	26
Rh24194	0	17	21
Rh25222	0	14	21

Strain 68-1.2 RhCMV/gag vector-vaccinated:



Rh26468	15	0	0
Rh26975	27	0	0
Rh27338	16	0	0
Rh27391	28	0	0
Rh28005	27	0	0
Rh28372	18	0	0

ΔRh157.5/.4 Strain 68-1.2 RhCMV/gag vector-vaccinated:



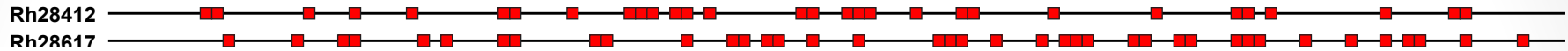
Rh24932	0	20	22
Rh25029	0	22	24
Rh29912	0	19	24
Rh29923	0	19	22

Consecutive SIVmac239gag 15mers (11 a.a. overlap)

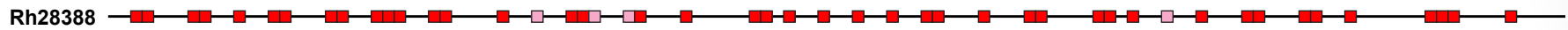
■ MHC-Ia restricted ■ MHC-E restricted ■ MHC-II restricted ■ Indeterminate

# Deletion of US11 “programs” RhCMV to additionally induce canonical MHC-I-restricted CD8+ T cells

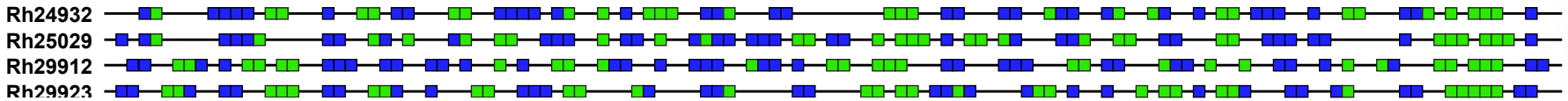
## RhCMV SIVgag



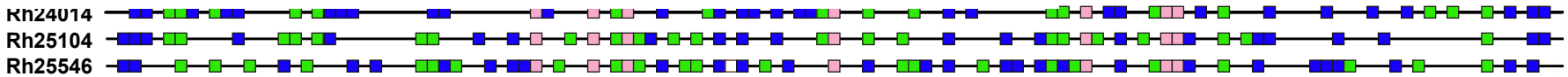
## ΔUS11 RhCMV SIVgag



## 68-1 RhCMV SIVgag



## ΔUS11 68-1 RhCMV SIVgag



1 25 50 75 100 125  
Consecutive SIVmac<sub>239</sub> gag 15mers (11 a.a. overlap)

- MHC-Ia restricted
- MHC-II restricted
- MHC-Ia restricted (canonical)
- Indeterminate
- MHC-E restricted
- Deconvolution/restriction pending

**Non-canonical**  
**Canonical**

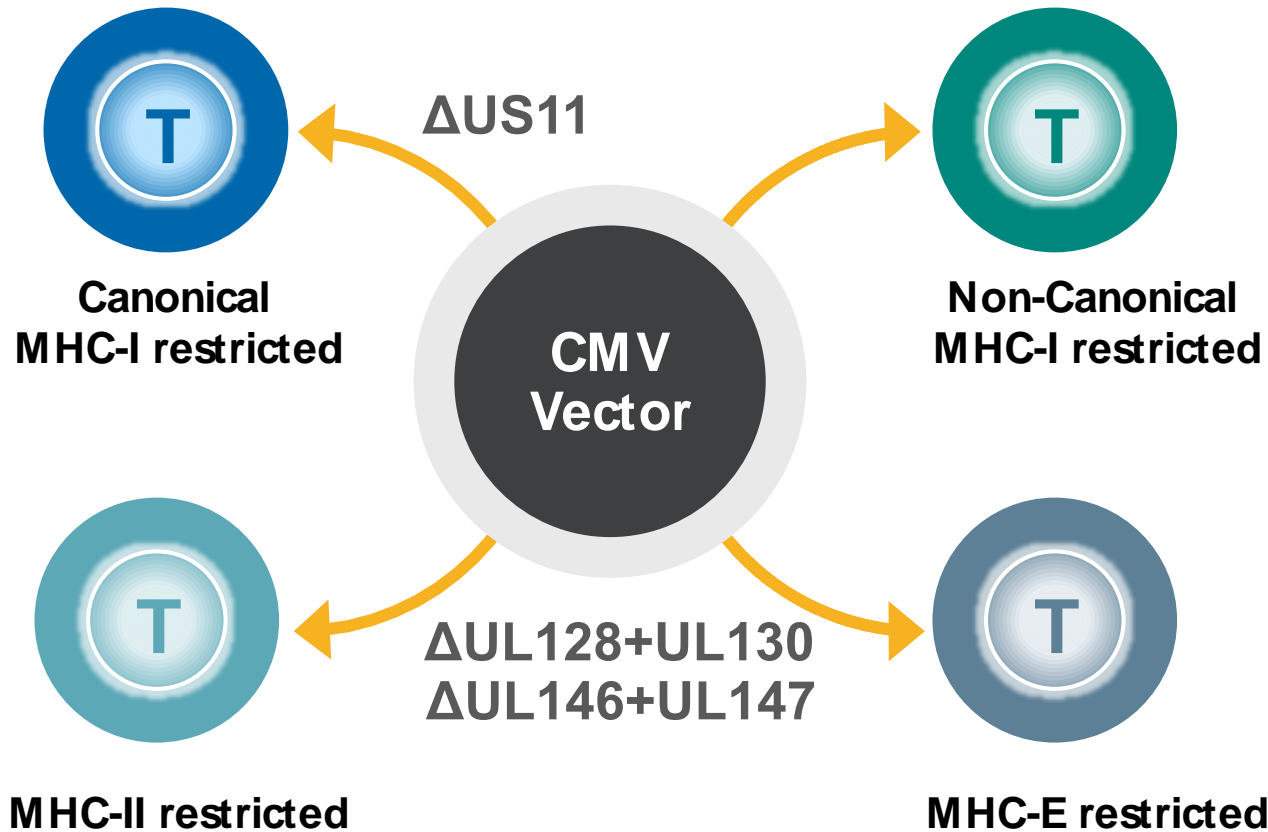
**in either a conventional or unconventional response background**

# We can program four non-overlapping CD8+ T cell responses!

## *Conventional T cell epitopes*

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Non-CMV  
Vectors  
mainly elicit  
canonical  
CD8+ T cells

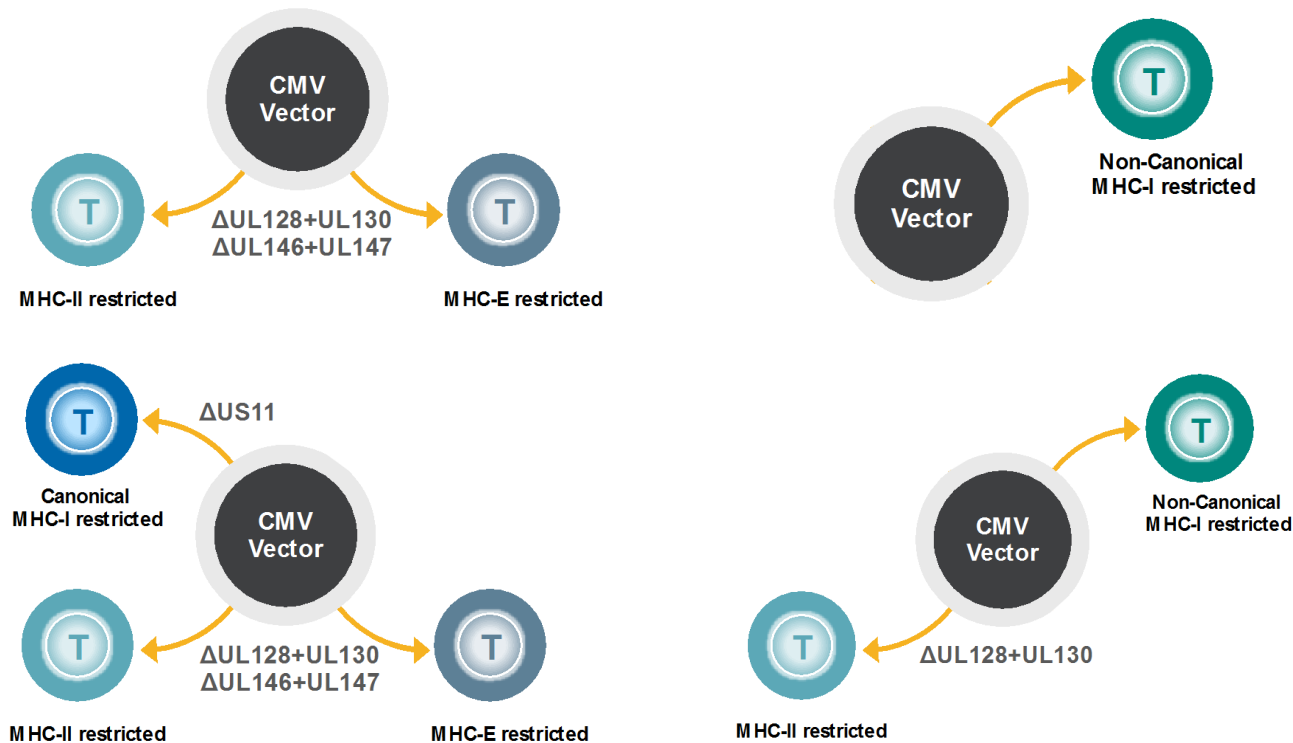


## *Unconventional T cell epitopes*

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# Using different vector backbones we can now determine the role of conventional (canonical/non-canonical) and unconventional (MHC-II, MHC-E) CD8+ T cell responses in protection against pathogen challenge





**SIV**

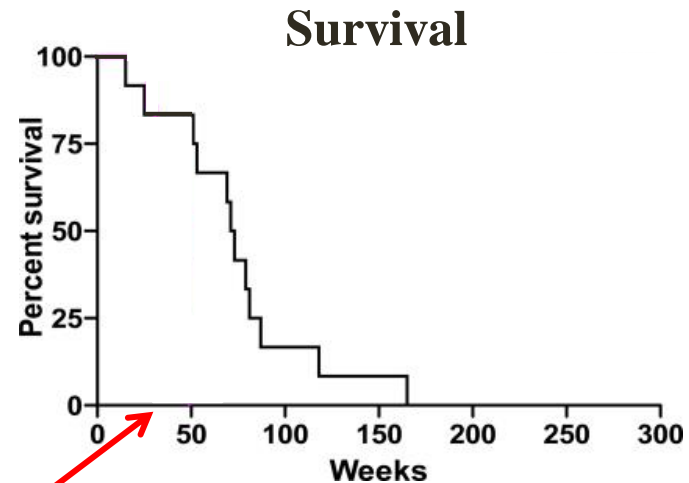
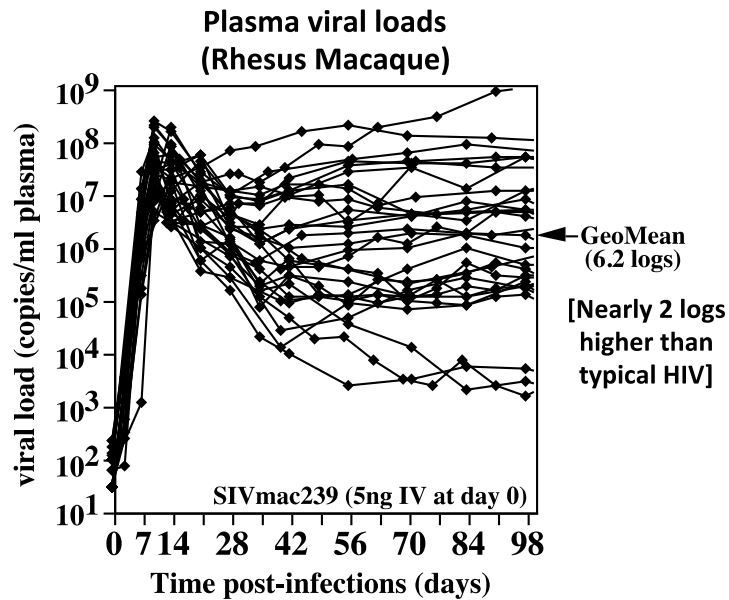
# SIVmac239: non-human primate AIDS model

- SIVmac239
  - Highly pathogenic in Indian origin Rhesus macaques
  - 10-100 times more virulent than HIV in humans
- Challenge model: repeated low dose intra-rectal or intra-vaginal inoculation to recapitulate natural infection with HIV



# SIVmac239: non-human primate AIDS model

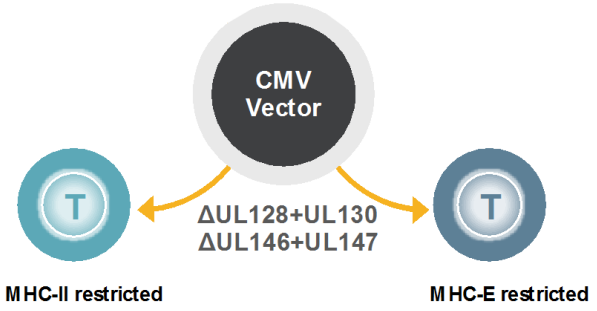
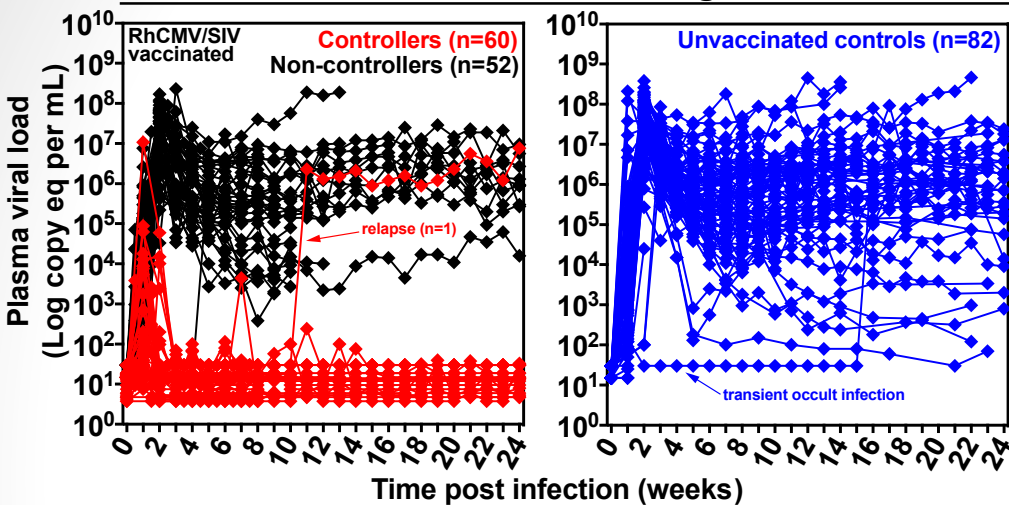
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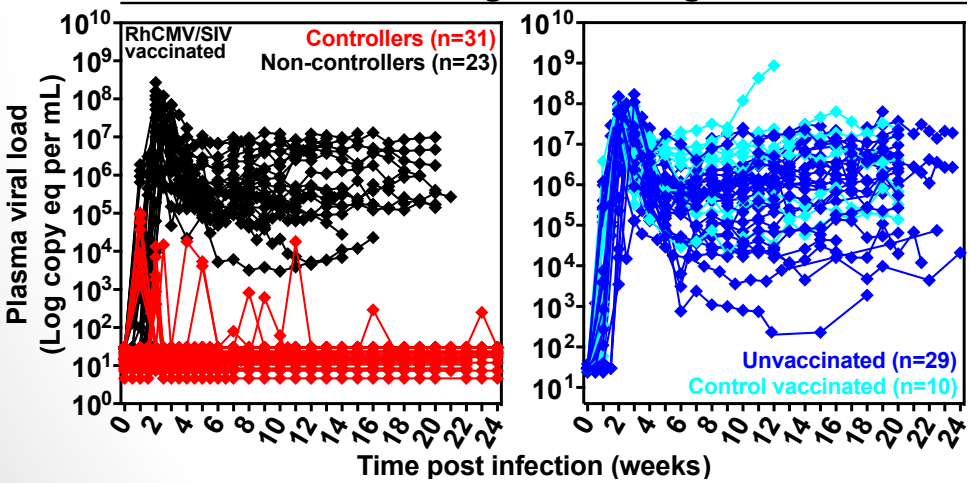
**Ave. time to AIDS in SIVmac239-infected RM is ~8-fold faster than for HIV infection!**

# RhCMV/SIV vaccine are the first to control highly virulent SIV

**Intra-rectal challenge**



**Intra-vaginal challenge**



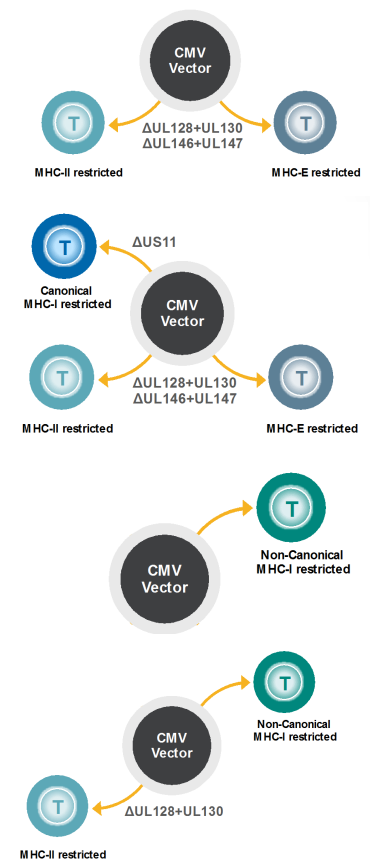
**Vaccine: 68-1 RhCMV**  
**Multiple vectors**  
**Expressing gag, rev, tat,**  
**Nef, pol +/- env**

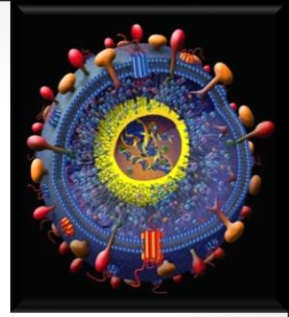
**All or nothing control!**

**Overall efficacy = 55% (91/166 animals protected)**

# Testing different CD8+ T cell programs against SIV

- 68-1 (MHC-E + MHC-II)
  - 55% protection (91/166 animals challenged)
- 68-1 $\Delta$ US11 (MHC-I<sub>canonical</sub> + MHC-E + MHC-II)
  - 21 % protection (3/14 animals challenged)
- 68-1.2 (MHC-I<sub>non-canonical</sub>)
  - 0% protection (0/10 animals challenged)
- 68-1.2 $\Delta$ UL128 (MHC-I<sub>non-canonical</sub> + MHC-II *non-supertopes*)
  - 0% protection (0/14 protected)
- **Current Status: Protection against SIV requires MHC-E and/or MHC-II<sub>supertope</sub> targeting CD8+ T cells**
  - Ongoing experiments: testing protection by CD8+ T cells that are MHC-E only, MHC-II only, MHC-I<sub>a</sub> only (not 68-1.2), MHC-E supertopes only, MHC-II supertopes only

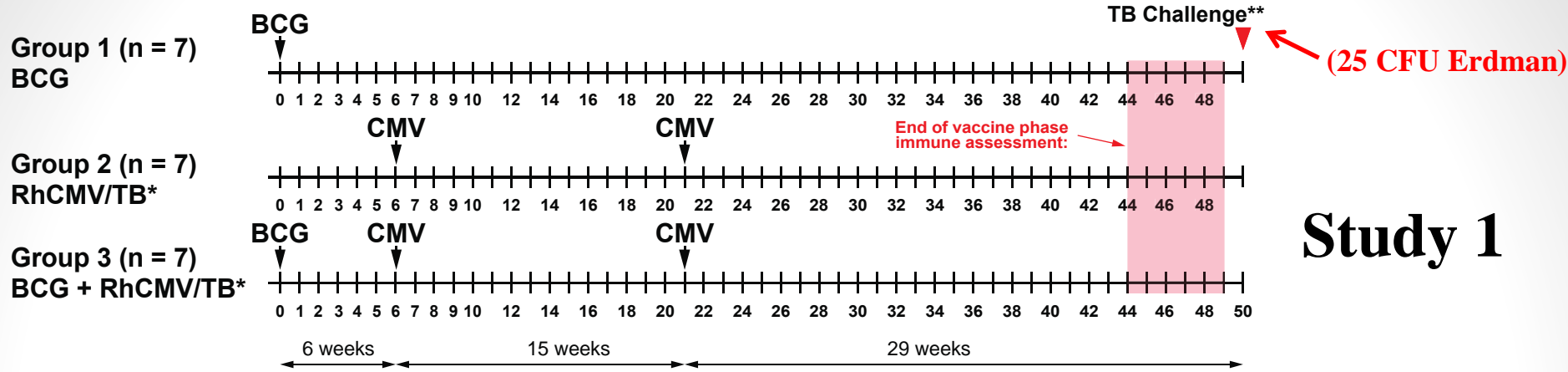




**TB**



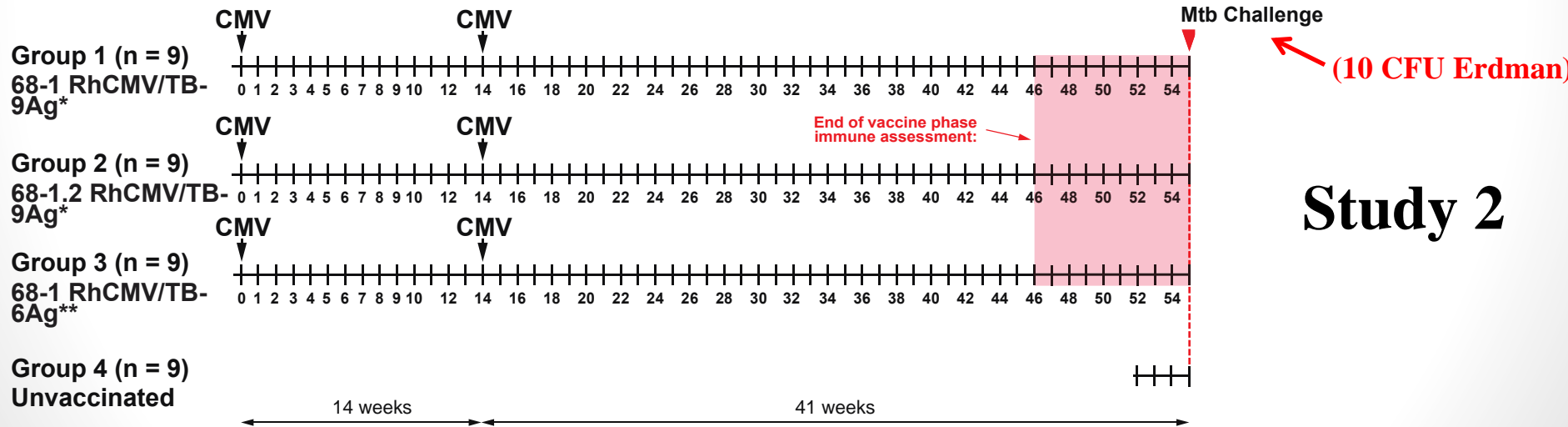
# Rhesus macaques are highly susceptible to Mtb



**Study 1**

\*68-1 RhCMV: Ag85A/Ag85B/Rv3407; Rpf A/Rpf C/Rpf D; Rv1733/Rv2626; Ag85B/ESAT-6

+ Group 4 (n = 8): unvaccinated controls

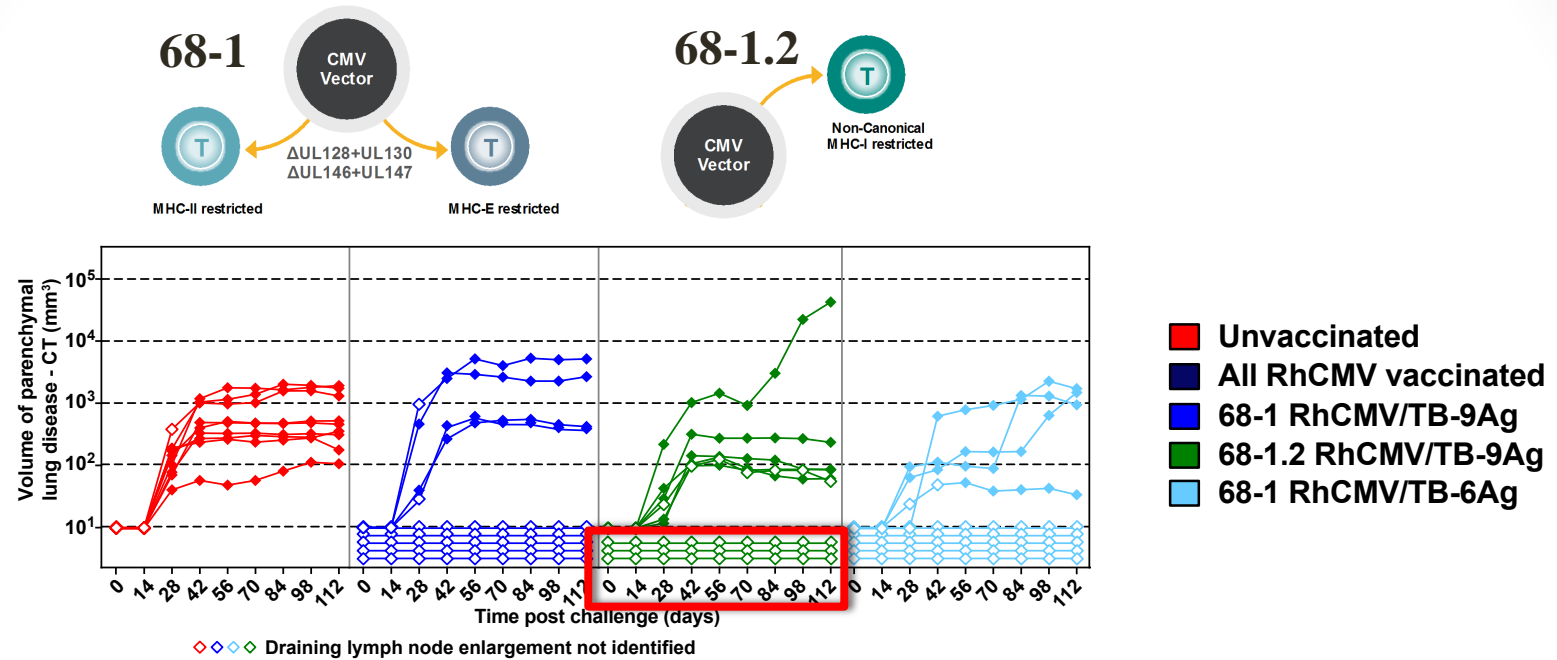


**Study 2**

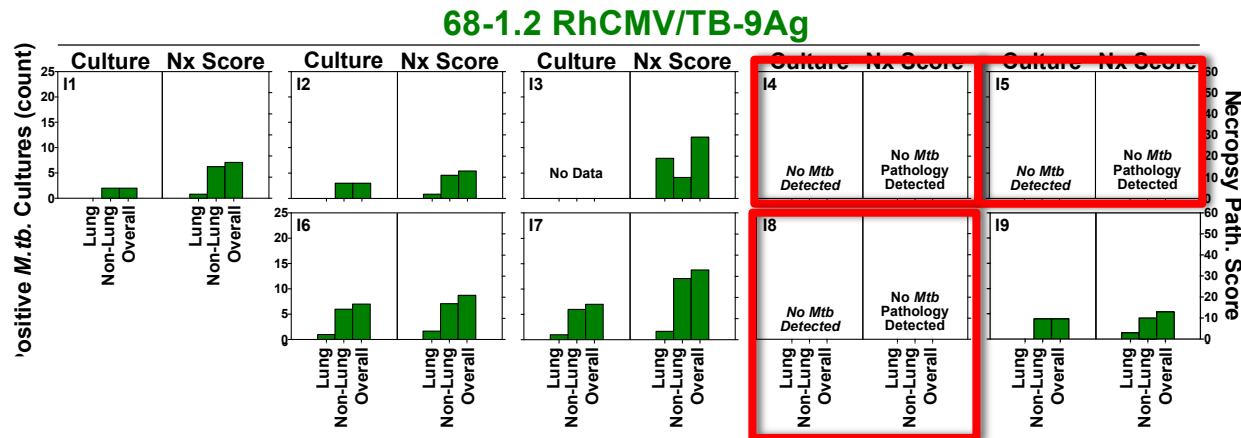
\*RhCMV(Ag85A/Ag85B/Rv3407); RhCMV(Rpf A/Rpf C/Rpf D); RhCMV(Rv1733/Rv2626); RhCMV(Ag85B/ESAT-6)

\*\*RhCMV(Ag85A/ESAT-6/Rpf A/Rpf D/Rv2626/Rv3407)

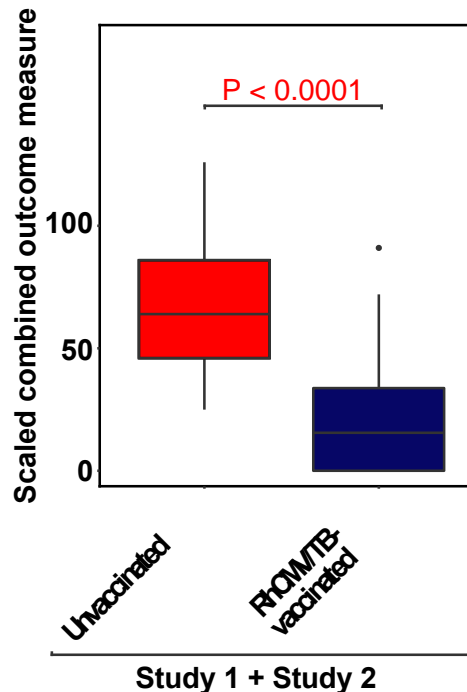
# Both conventional and unconventionally targeted CD8+ T cells protect against TB



**Disease at necropsy**



# Combined Outcome Analysis of RhCMV/TB Efficacy Studies #1 and #2



Using a scaled, combined (culture + path score) outcome measure, the extent of overall disease significantly reduced in RM vaccinated with RhCMV/TB alone compared to unvaccinated controls.

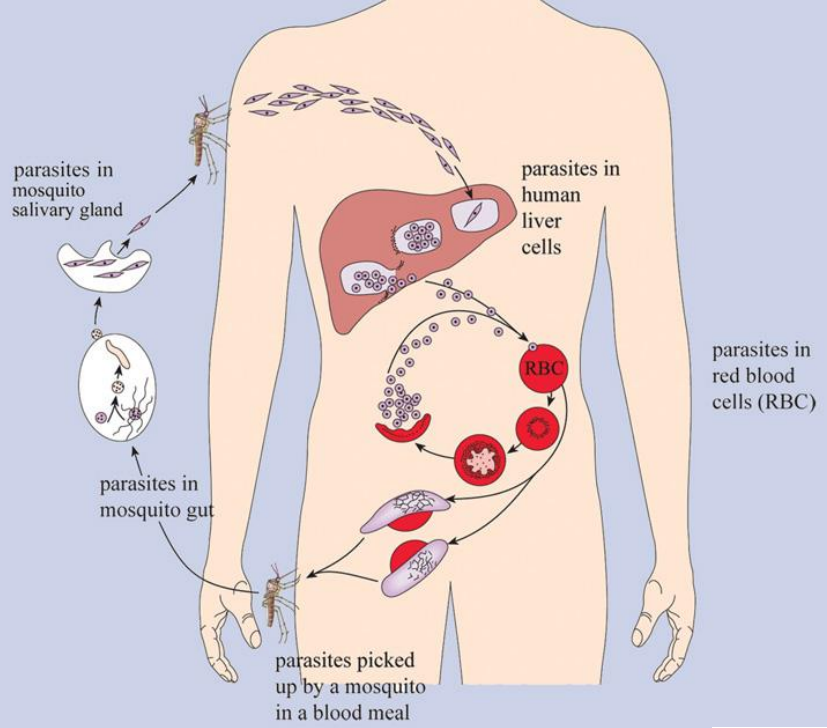
By Poisson modeling, the overall reduction in this combined outcome score over both studies was 68% (95% CI: 35.9-85.1%;  $P=0.0019$ ).

Note: no significant efficacy was observed with standard BCG vaccination

Currently comparing 68-1 and 68-1.2 6Ag in larger cohorts



# Malaria



## *Plasmodium knowlesi*

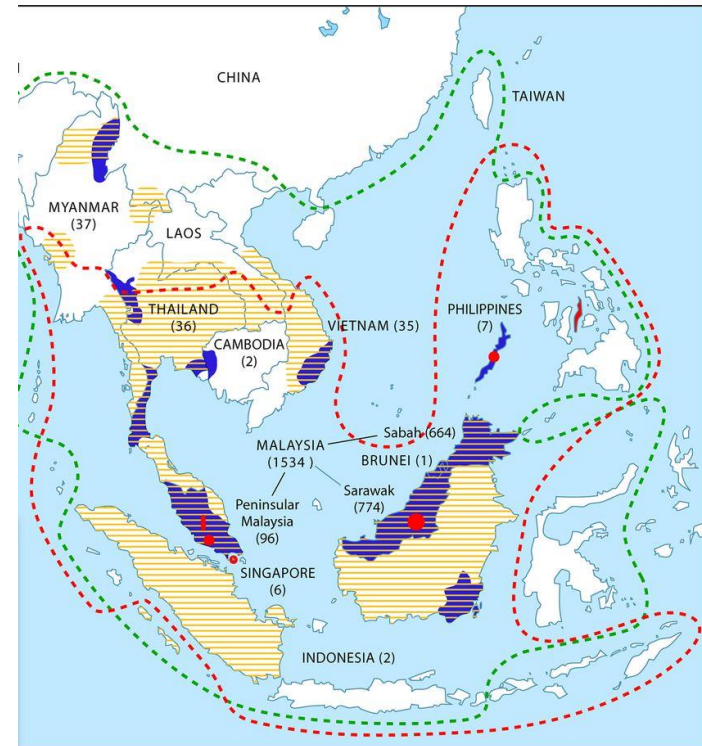
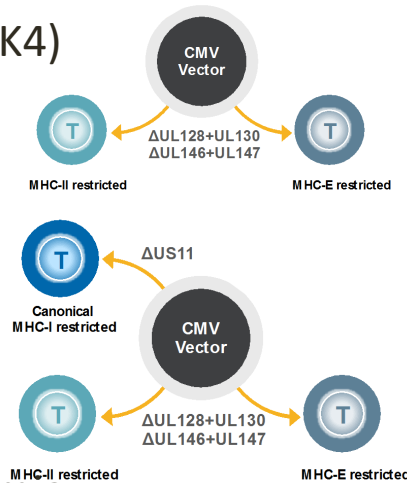
- wide-spread in SE Asia
- Zoonotic infection in humans
- Rhesus macaques not natural hosts (highly susceptible)

### Vaccine: CSP/MSP1/TRAP/AMA1 (PK4)

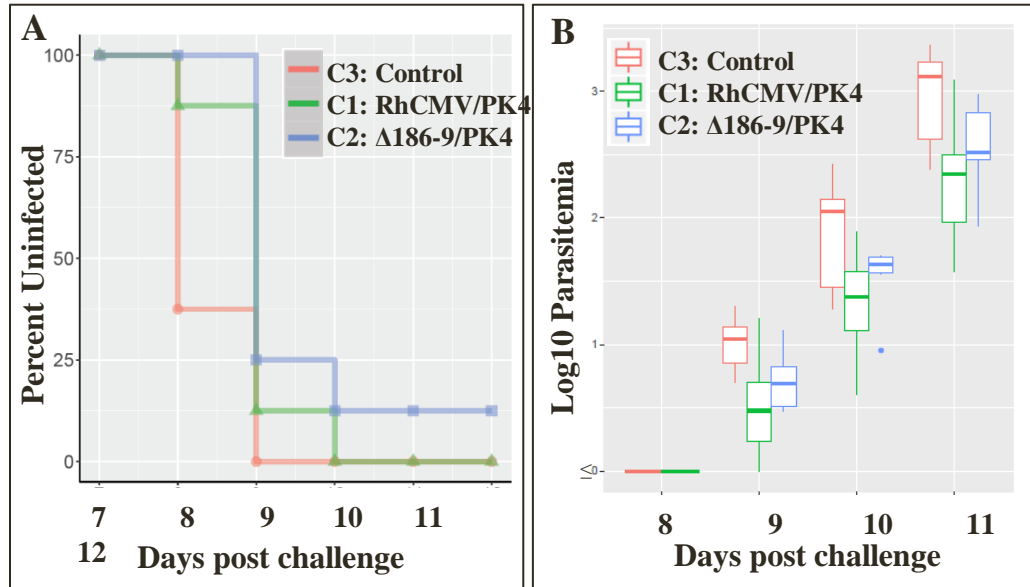
- 68-1 RhCMV/PK4 (n=8)
- 68-1 RhCMV $\Delta$ US11PK4 (n=8)
- Control cohort (n=8)

### Challenge

- 100 Sporozoites inoculated intravenously
- After day 6: monitor blood viremia by thin smear



# Challenge of RhCMV/PK4 immunized RM with *P. knowlesi* sporozoites



Delay in appearance of blood stage parasites in both vaccine cohorts

~ 80% reduction of liver stage

No impact on blood stage

One sterile animal in  $\Delta$ US11 68-1 group

Currently testing additional antigen and vectors eliciting conventional CD8+ T cells

## Vaccine Efficacy (% reduction at day 8)

Cohort	VE	2.5%	97.5%	P value
<b>C3 vs. C1</b>	<b>75%</b>	<b>49%</b>	<b>88%</b>	<b>0.0002</b>
<b>C3 vs. C2</b>	<b>80%</b>	<b>60%</b>	<b>90%</b>	<b>&lt;0.0001</b>

Not only is the answer to this question “yes”, but the immunobiologic properties of CMV vectors turned out to be markedly different from any other known vector or non-vector vaccine modality . . .

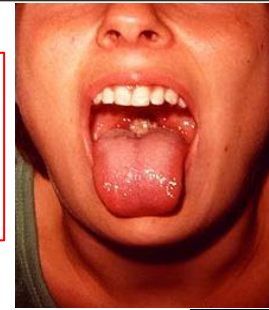
**In nonhuman primates, vectors based on rhesus cytomegalovirus:**

1. Elicit and indefinitely maintain high frequency, “effector memory” T cell responses\* in mucosal sites, lymphoid tissues and parenchymal organs
2. Efficiently re-infect & persist despite robust anti-CMV immunity
3. Manifest uniquely programmable CD8+ T cell immunogenicity with expanded breadth and conventional and/or novel epitope targeting.
4. Provide protection against SIV, TB and Malaria
5. Maintain immunogenicity even with profound attenuation (spread-deficiency and tropism restriction)

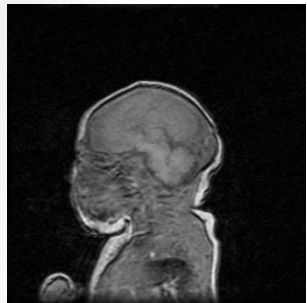
\*Little to no antibodies



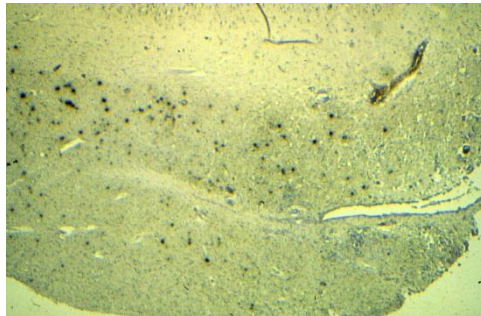
**Since HCMV can be a pathogen, vector safety is a top priority issue . . .**



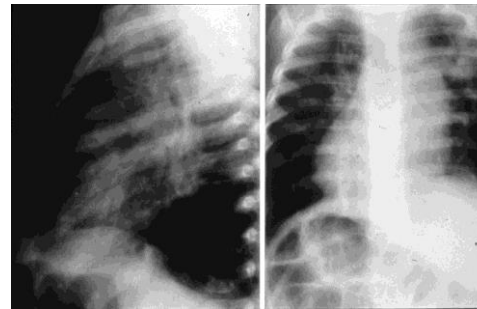
**mononucleosis**



**Congenital disease**



**encephalitis**



**pneumonitis**



**retinitis**

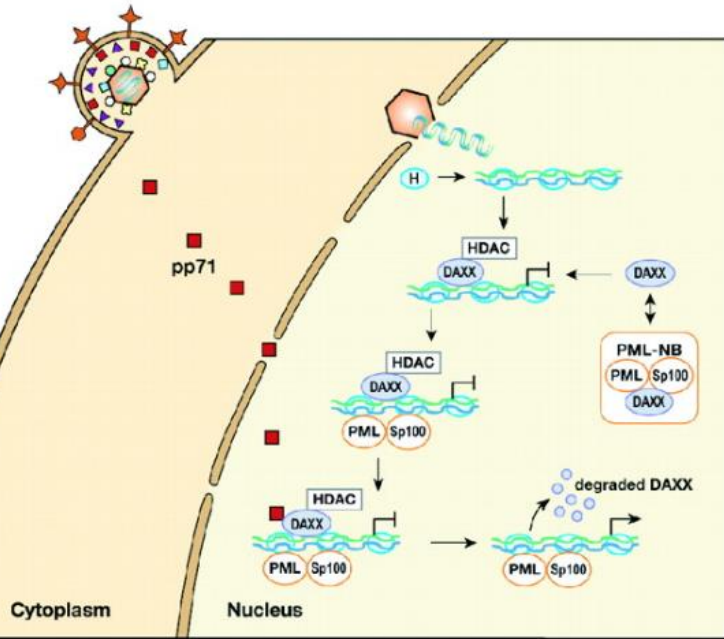
**Thus, for clinical translation, CMV vectors must be be engineered to be have:**

- **greatly reduced or no ability to cause disease,**
- **greatly reduced or no ability for person-to-person transmission,**
- **stability *in vitro* and *in vivo*, and**
- **the ability to be GMP manufactured at large scale,**

**While retaining the ability to super-infect, persist, and elicit/maintain all the immune responses needed for efficacy . . .**

# Employing intrinsic and innate immunity to attenuate CMV

The CMV tegument protein pp71 counters DAXX, a host intrinsic immune protein that blocks the viral gene expression required for lytic infection. pp71 has also been shown to counteract the innate immune signaling molecule STING



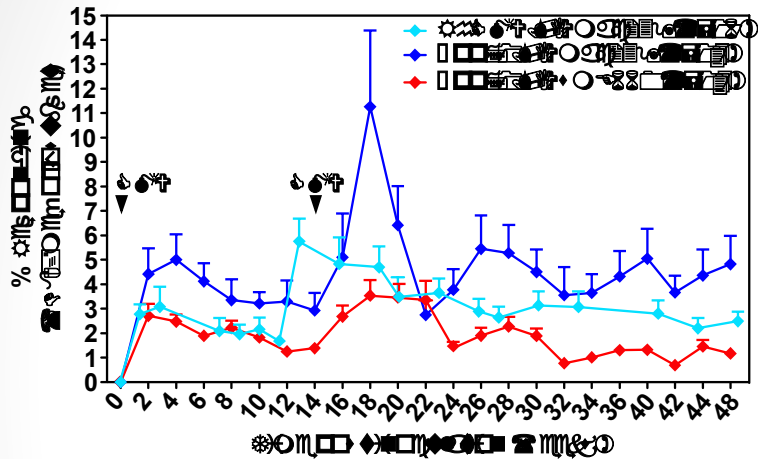
Kalejta R F Microbiol. Mol. Biol. Rev. 2008;72:249-265

## RhCMV lacking pp71 is highly attenuated in vivo:

- **Dissemination:** pp71-deleted RhCMV is limited to site of inoculation, very little dissemination
- **Shedding:** pp71-deleted RhCMV cannot be detected in urine by co-culture at any time post-infection
- **Transmission:** pp71-deleted RhCMV is not transmitted upon leukocyte transfusion, pp71-deleted RhCMV is not transmitted via breast milk from mother to infant
- **Fetal inoculation:** reduced spreading and fetal loss compared to wildtype RhCMV upon direct inoculation of high doses ( $10^6$ ) of pp71-deleted RhCMV

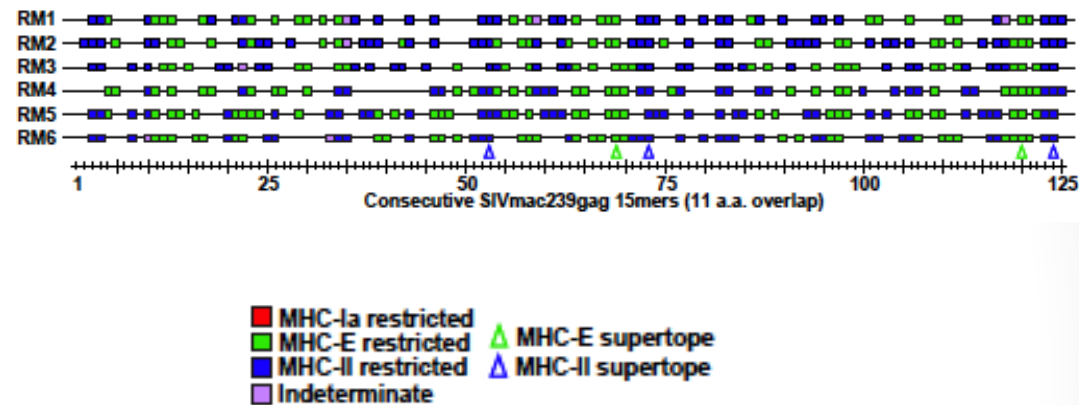
# Yet, $\Delta$ pp71 68-1 RhCMV/SIV vectors are similarly immunogenic and protective as WT 68-1 vectors, including against heterologous challenge

RhCMV/SIV vector-elicited CD8+ T cell responses to SIVmac239 sequence peptides (total)



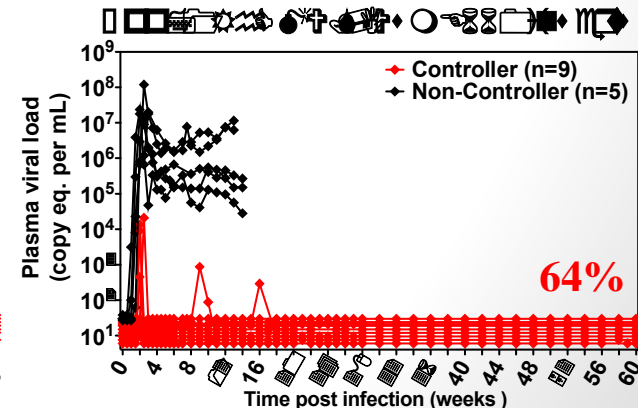
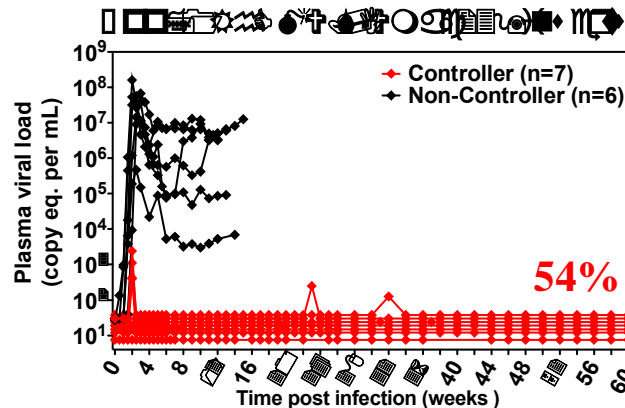
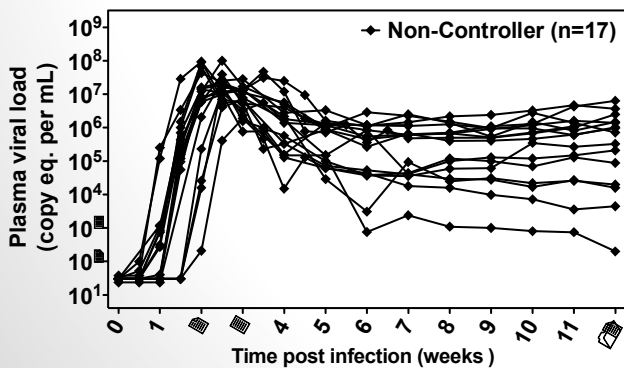
[RhCMV inserts: Gag, Pol, Rev/Nef/Tat, Env]

Epitope Targeting of pp71-deleted 68-1 RhCMV

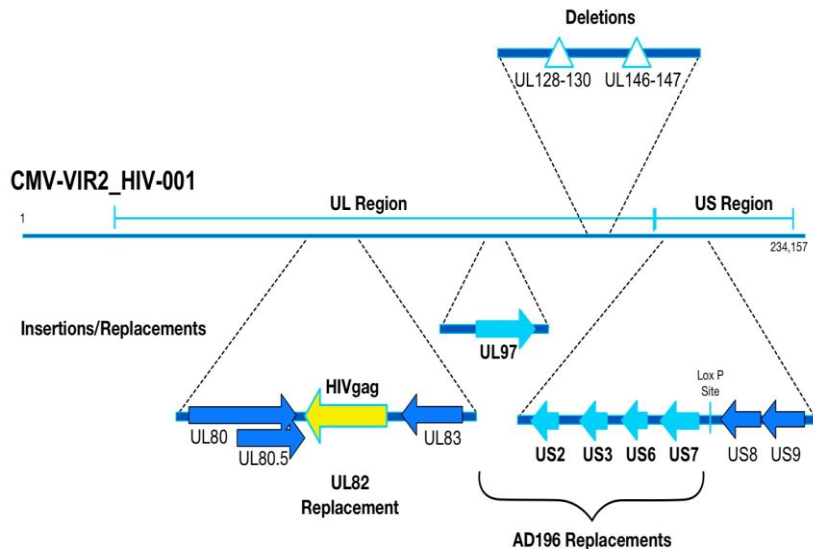


## Outcome of SIVmac239 challenge:

Unvaccinated controls



# Based on RhCMV results we now generated a pp71-deleted “68-1” like HCMV/HIV vector for clinical testing



Vector = Vaccine Prototype designed to determine

- Safety
- Reinfection capability
- Immunogenicity (CMV, HIV, T cell phenotype)
- Immune Programming

## Characteristics of pp71-deleted HCMV:

- highly attenuated in vitro (4log at low MOI in fibroblasts)
- reduced ability to infect non-fibroblast cells (due to UL128-130 deletion)
- maintains ability to establish latency in humanized mouse model
- reduced ability to reactivate in humanized mouse model
- immunogenic in monkeys  $>10e5$  FFU

**This vector is now in GMP manufacturing with clinical testing planned for 2019**

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Global HIV Vaccine  
Enterprise

OHSU has licensed CMV technology, of which Dr. Frueh is an inventor, to VIR Biotechnology, a company in which both OHSU and Dr. Frueh have significant financial interest. Potential individual and institutional conflicts of interest have been reviewed and are actively managed by OHSU.



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