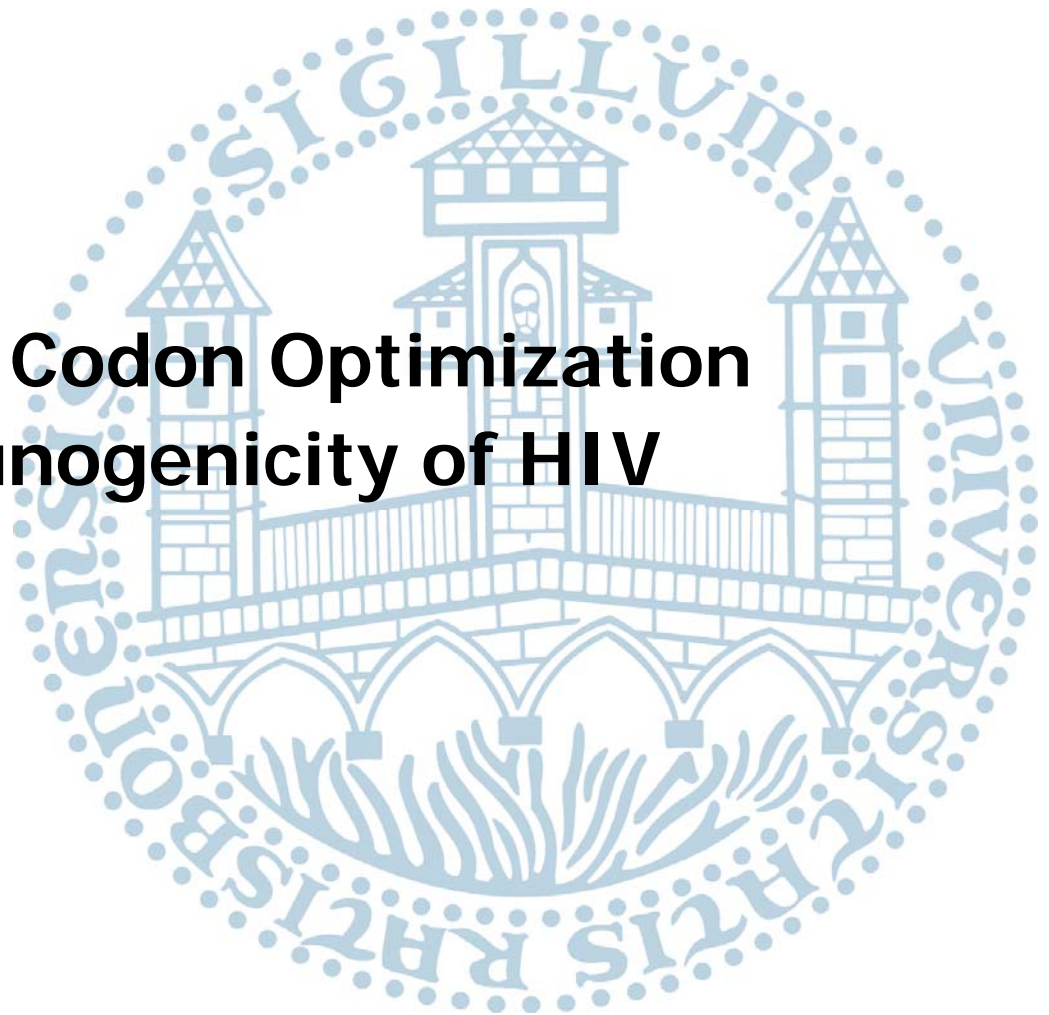


Impact of RNA- and Codon Optimization on Safety and Immunogenicity of HIV Candidate Vaccines

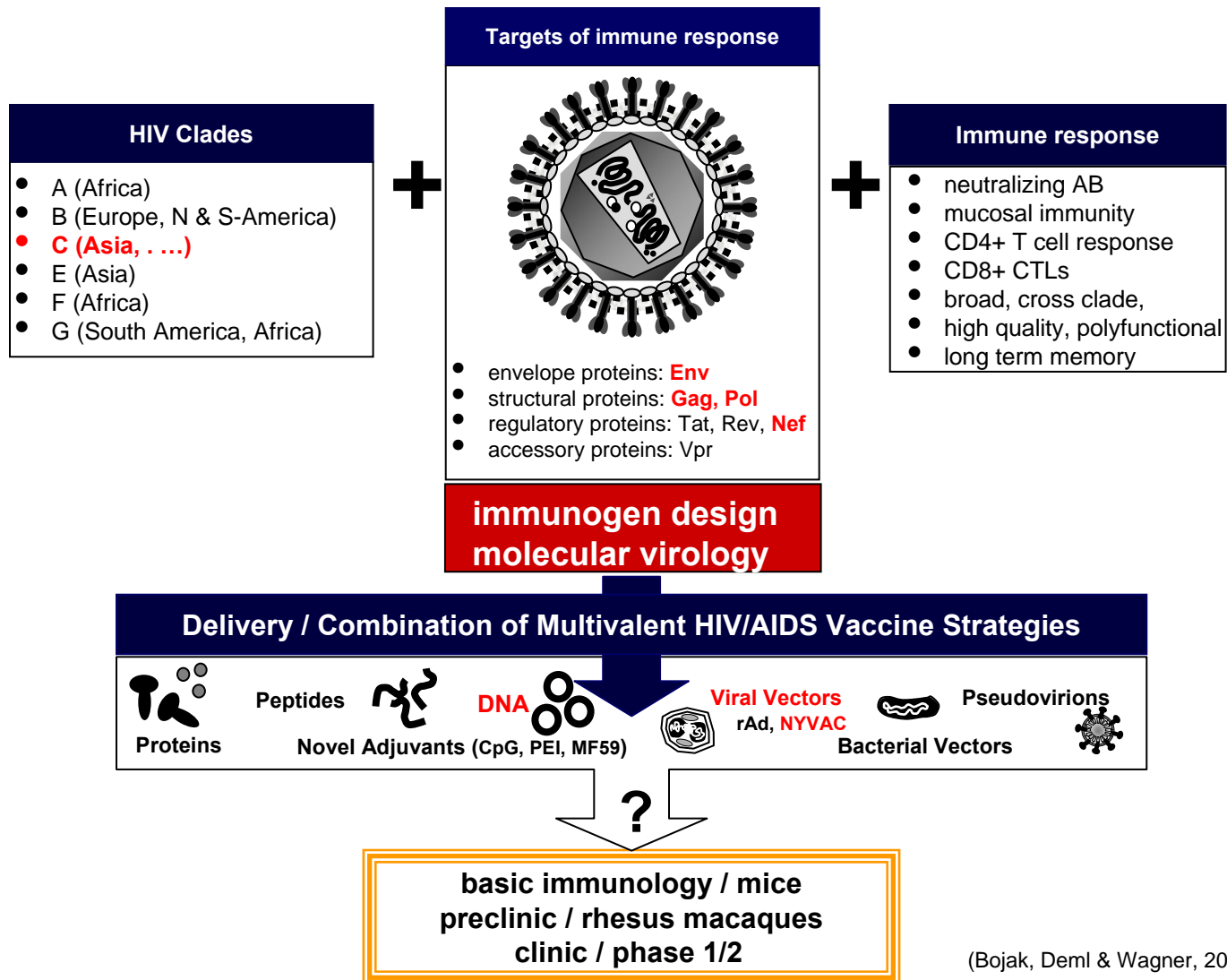
Ralf Wagner



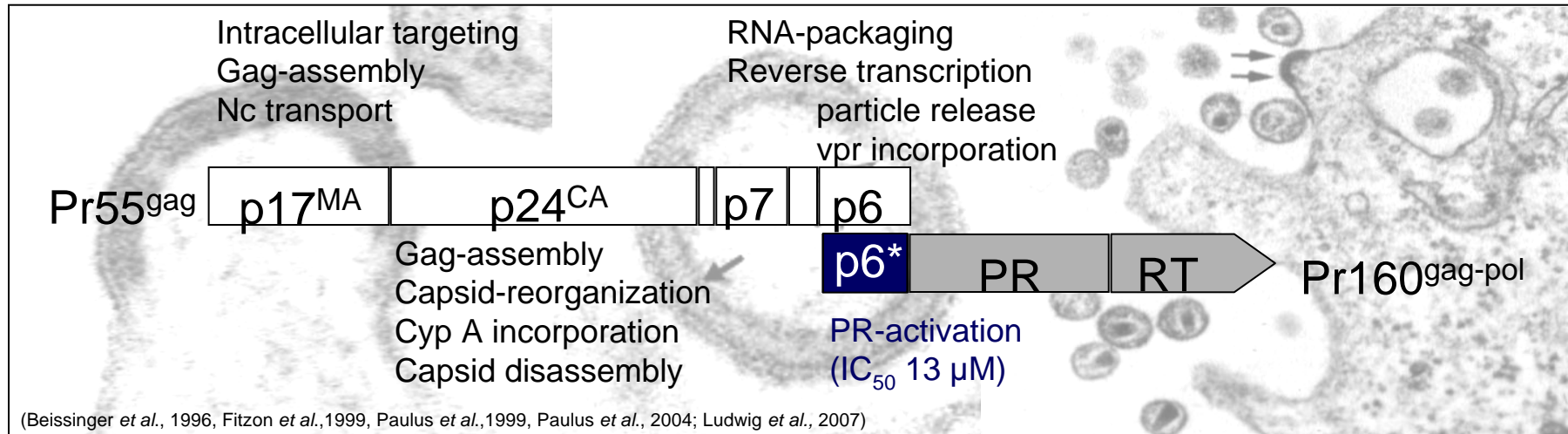
Molecular Microbiology and Gene Therapy Unit

**Institute of Medical Microbiology and Hygiene
University of Regensburg**

Integrated Vaccine Strategies



Pr55^{gag} – Necessary Component of HIV Vaccines



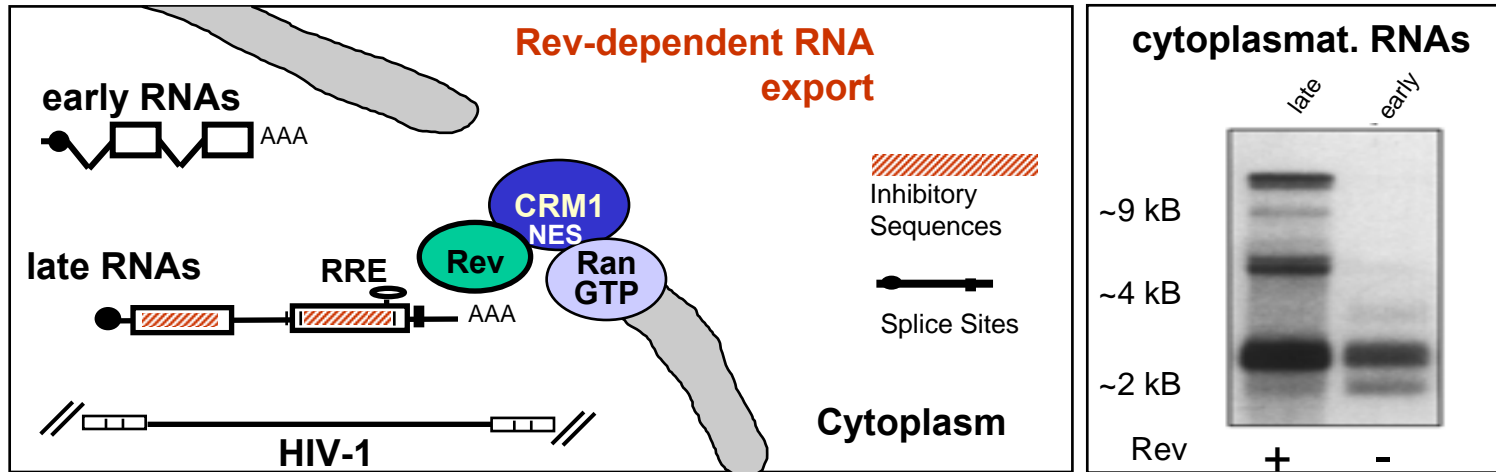
- Patients with strong Gag-Helper/CTL-Response control acute HIV infection better
(Rosenberg *et al.*, 1997; Ogg *et al.*, 1999, many others)
- LTNP HIV infected individuals control HIV replication by T cells recognizing highly conserved epitopes within Gag (Harrer *et al.*, 1998, Wagner *et al.*, 1999, many others)
- Recombinant and chimeric Gag VLPs induce Th1 / CTL responses in Balb/C mice;
Cross priming (Wagner *et al.*, 1993, 1994, 1996; Bojak *et al.*, 2002, 2003, 2004, Deml *et al.*, 1997a,b, 2005)
- Gag VLP partially protective in Rhesus monkey model (Wagner *et al.*, 1998; Notka *et al.*, 1999a, b, Ludwig & Wagner, 2007)

→ Gag specific T-cells have been linked to better control of HIV replication

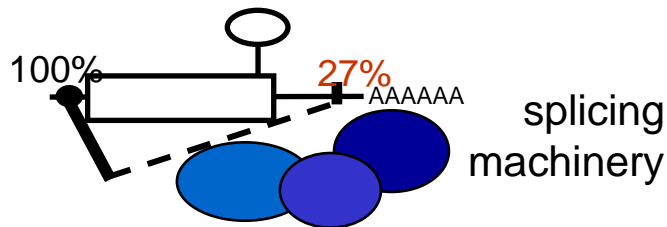
Rationals for Immunogen Selection



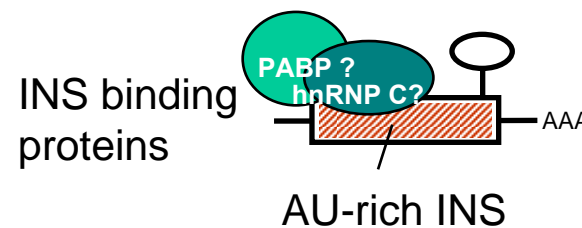
Late Lentiviral Gene Expression is Complex



strong splice donor and weak splice acceptors capture RNAs within splicing machinery



cis-located AU-rich repressor elements (INS) retain RNAs within the nucleus

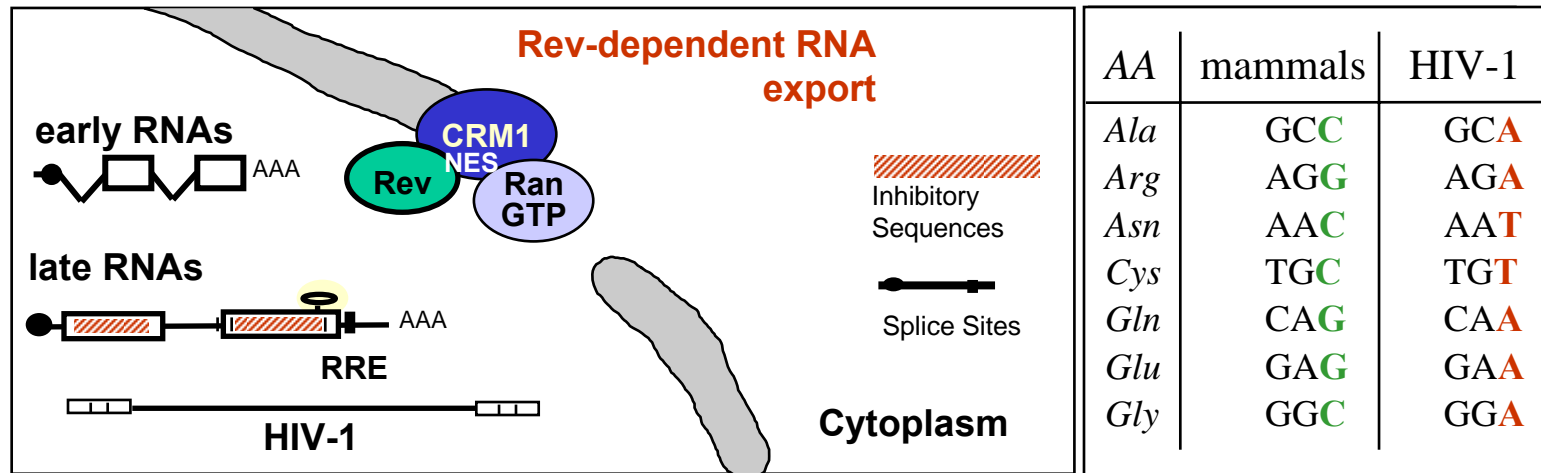


Intrinsic risks: Ψ -sequence, recombination, RRE/Rev, ...



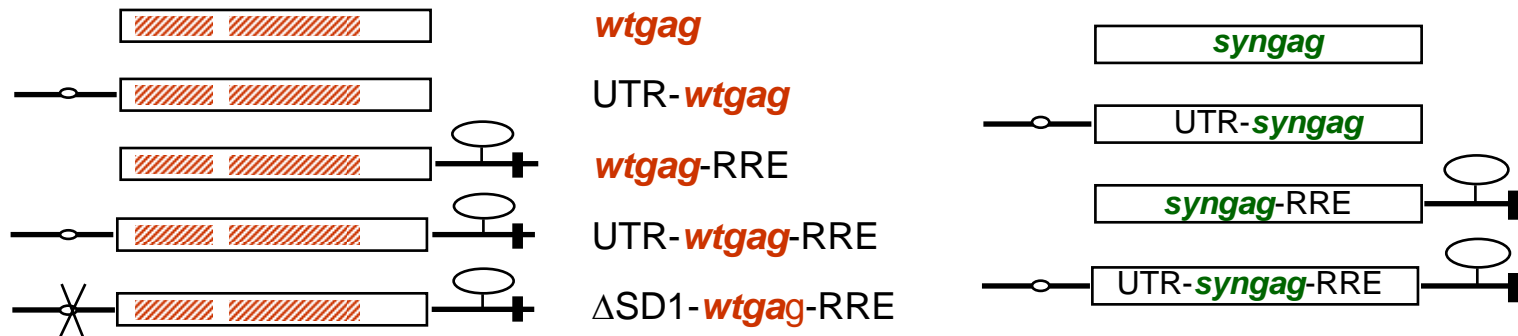
Bypassing Rev/RRE Regulation

Understanding Late Lentiviral Gene Expression



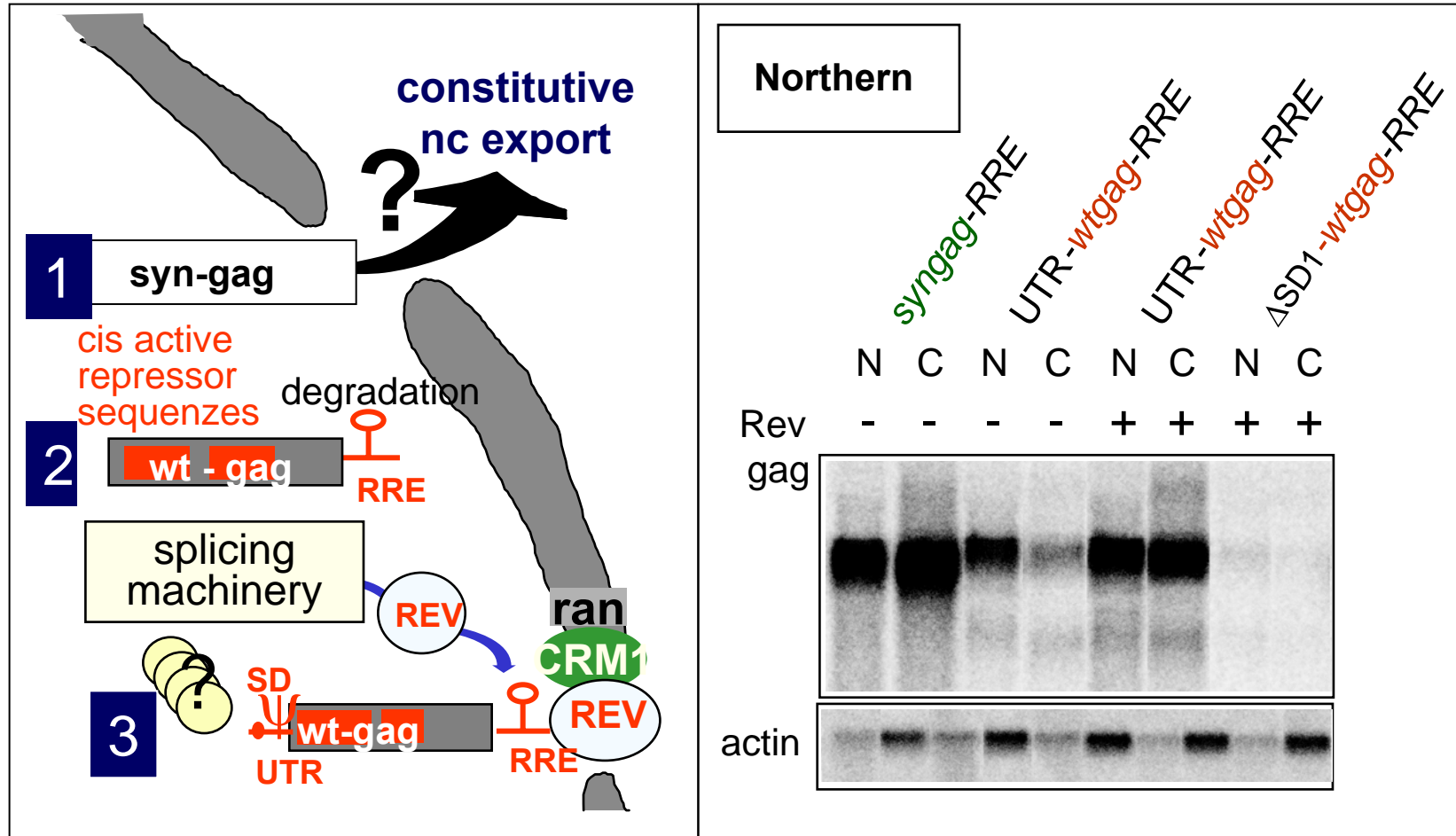
Influence of various cis-acting sequences [INS, UTR (SD), RRE (SA)]

GENEART
THE GENE OF YOUR CHOICE

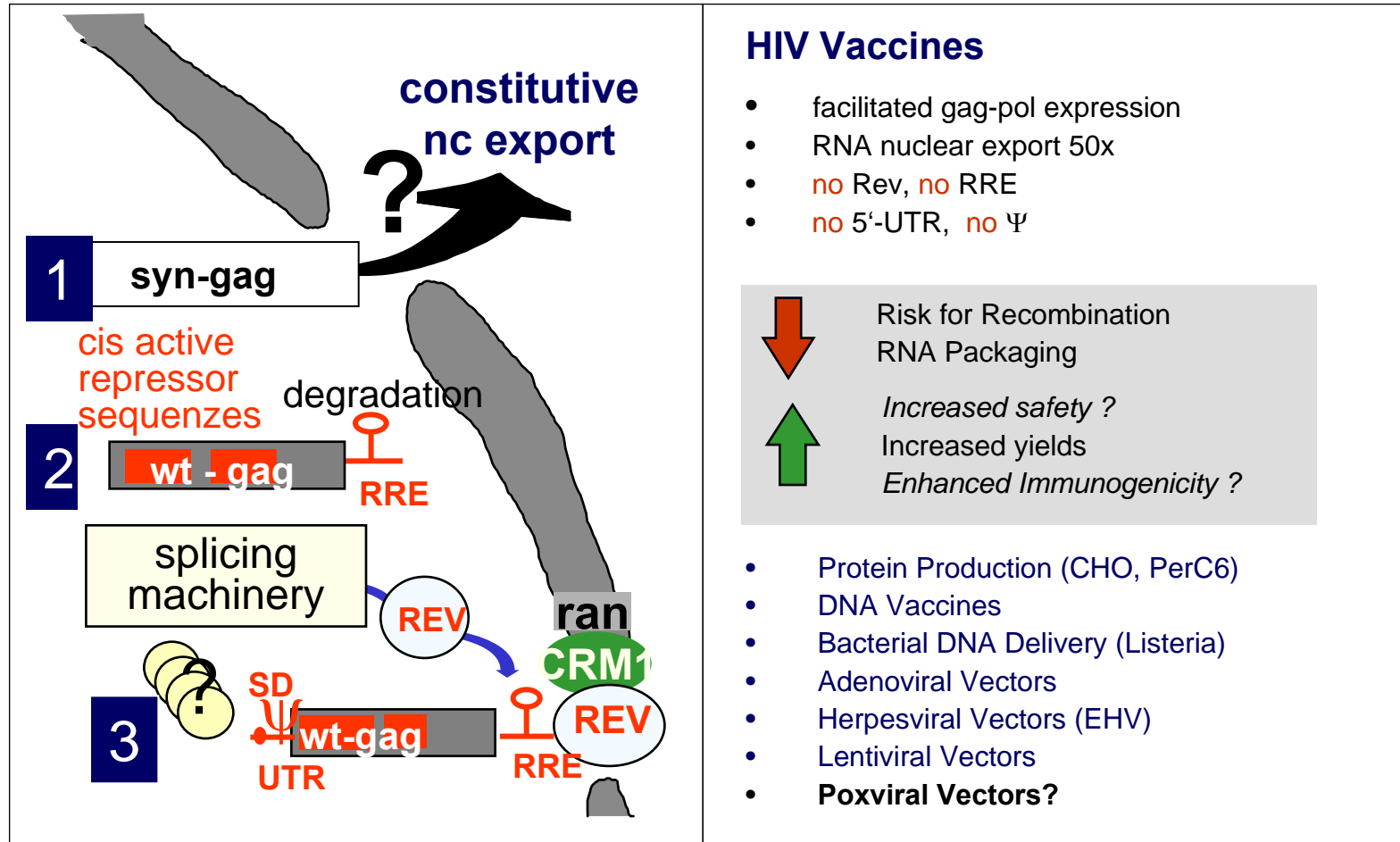


Bypassing Rev/RRE Regulation

Facilitating Late Lentiviral Gene Expression



Consequences of Modulated Late Lentiviral Gene Expression



HIV Vaccines

- facilitated gag-pol expression
- RNA nuclear export 50x
- no Rev, no RRE
- no 5'-UTR, no Ψ



Risk for Recombination
RNA Packaging



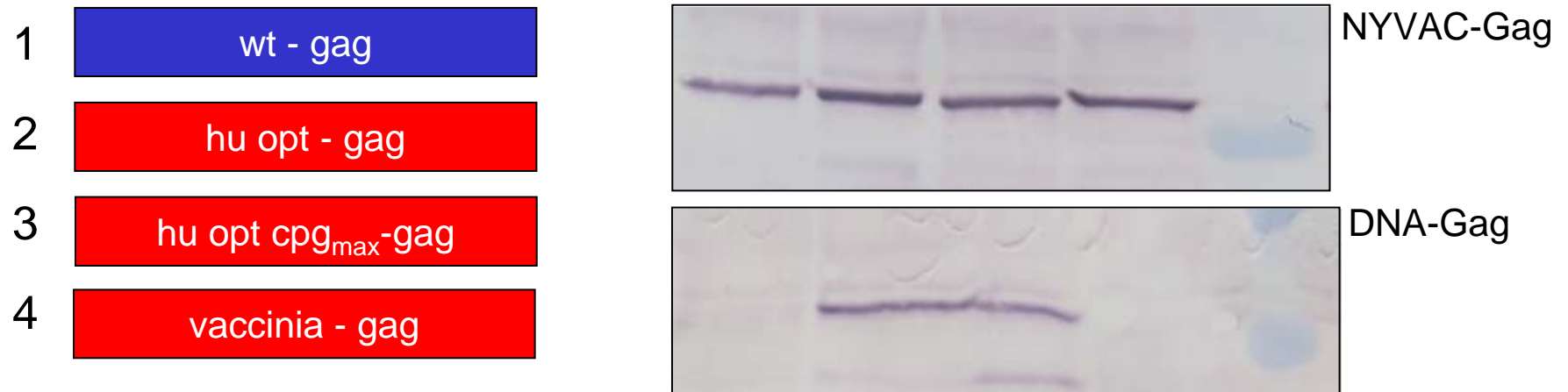
Increased safety ?
Increased yields
Enhanced Immunogenicity ?

- Protein Production (CHO, PerC6)
- DNA Vaccines
- Bacterial DNA Delivery (Listeria)
- Adenoviral Vectors
- Herpesviral Vectors (EHV)
- Lentiviral Vectors
- **Poxviral Vectors?**

Graf et al., 2000; Wagner et al., 2000; Deml et al., 2001; Bojak et al., 2003; Wild et al., 2004, 2007



Impact of RNA & Codon Optimization in Poxviral Vectors



WB - Antibody: CD-4/1 (anti p24 CA)

NYVAC: No major impact of RNA & codon optimization on transgene expression

DNA: Significant impact of RNA & codon optimization on transgene expression

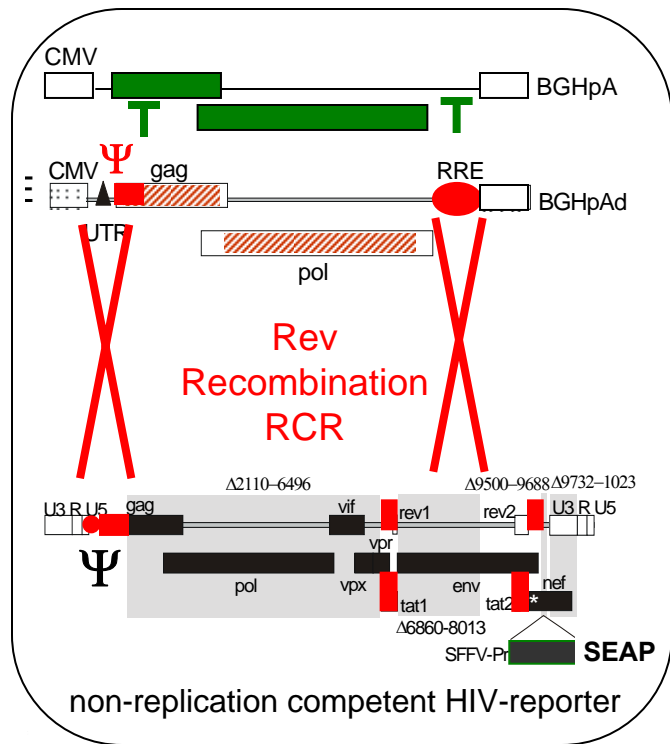
Conclusion: RNA- and codon-optimization recommended if various vector systems shall be compared / combined



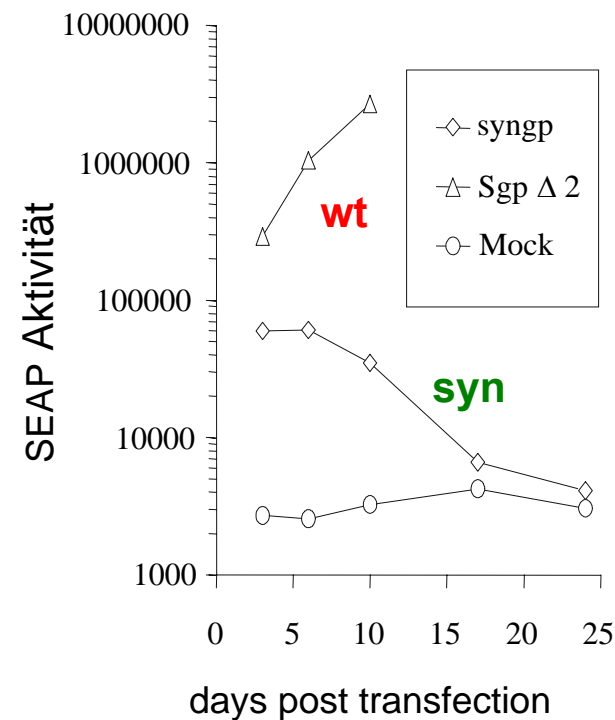
Reduced Risk for Generating Replication Competent Recombinants

...following RNA- and Codon-Optimization

- 2-Plasmid-system
- Transfervector gagpol/Env⁺
- packaging construct wt vs syn
- cotransfection and harvest particles

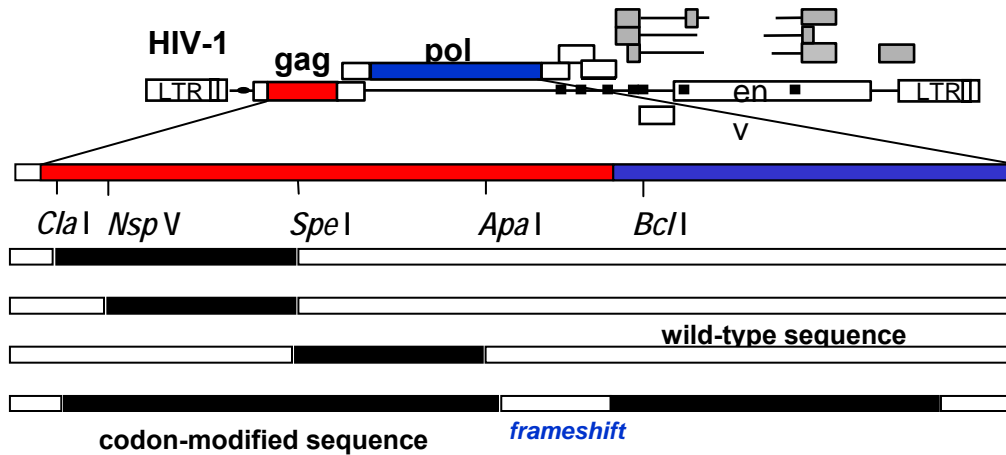


- Permissive CD4⁺ cell culture system
- Replication post recombination
- Tat driven SEAP expression



Reduced Replicative Capacity of Engineered Recombinant HIVs

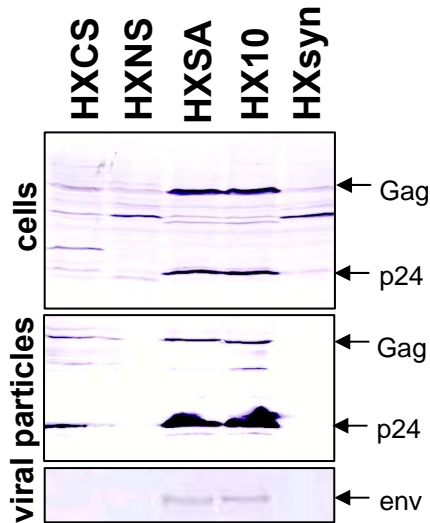
chimeric viruses



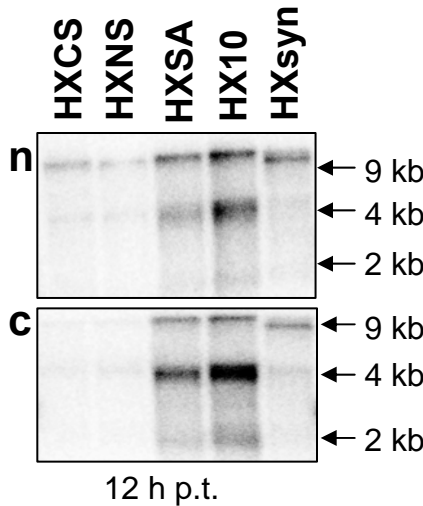
Mutant: Codon-optimized region (gag ATG=112):

<i>HXCS</i>	5' gag	(bp 154-830)
<i>HXNS</i>	5' gag	(bp 337-830)
<i>HXSA</i>	3' gag	(bp 830-1333)
<i>HX10</i>	-----	
<i>Hxsyn</i>	<i>gag/pol</i>	(bp 154-1333...)

Processing & maturation

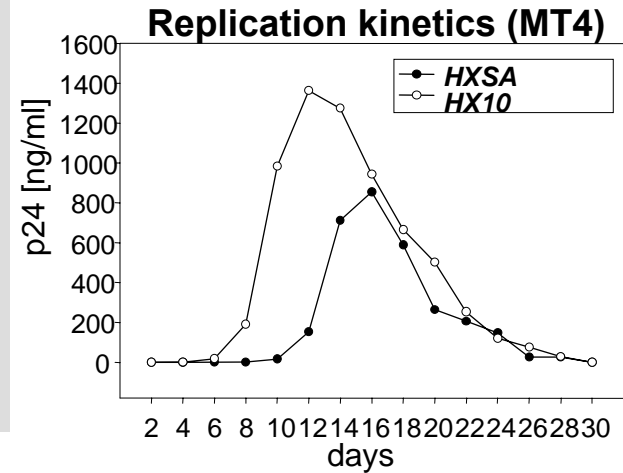


RNA splicing & export



(Wagner *et al.*, 2000; Steck *et al.*, in prep.)

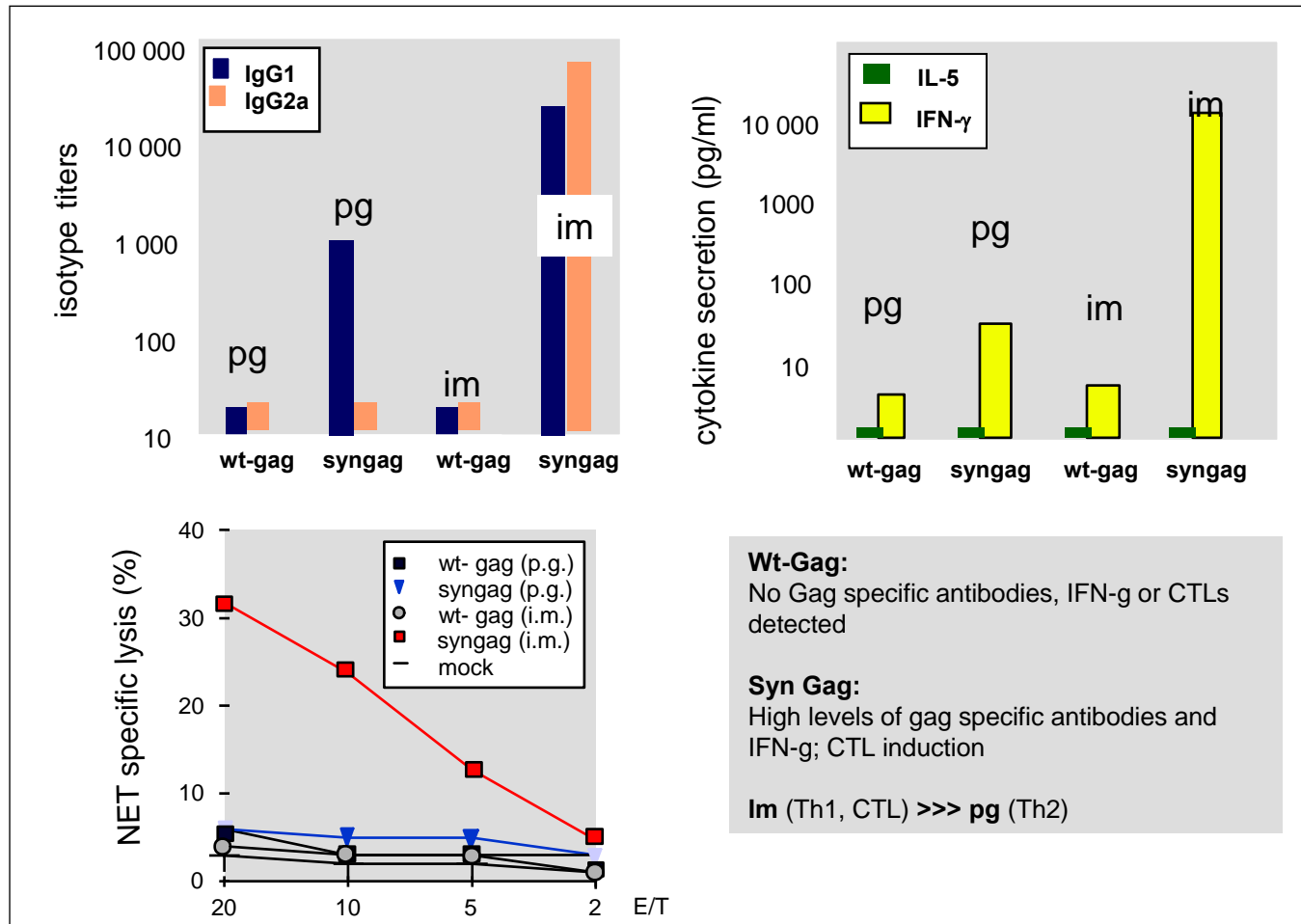
Viral infectivity



Safety Profiles

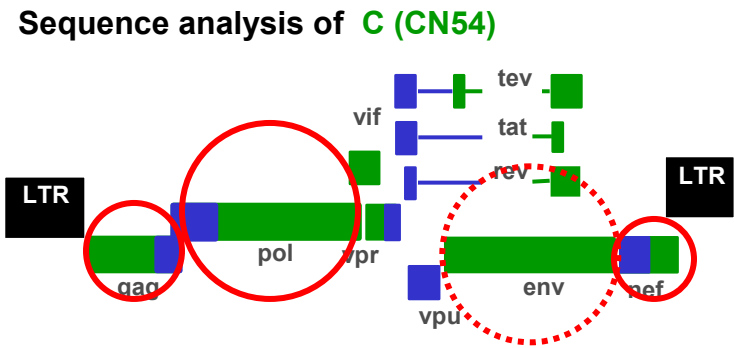
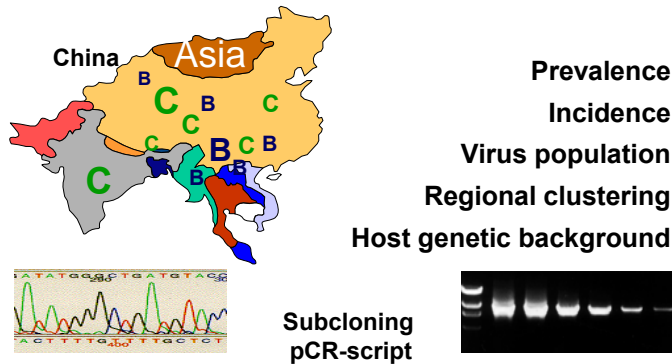
RNA and Codon Optimization DNA Vaccines

Increased immunogenicity in BALB/c mice



Immunogenicity

Molecular Epidemiology: Prerequisite for Immunogen Design



Graf et al., 1998, Su et al., 2000



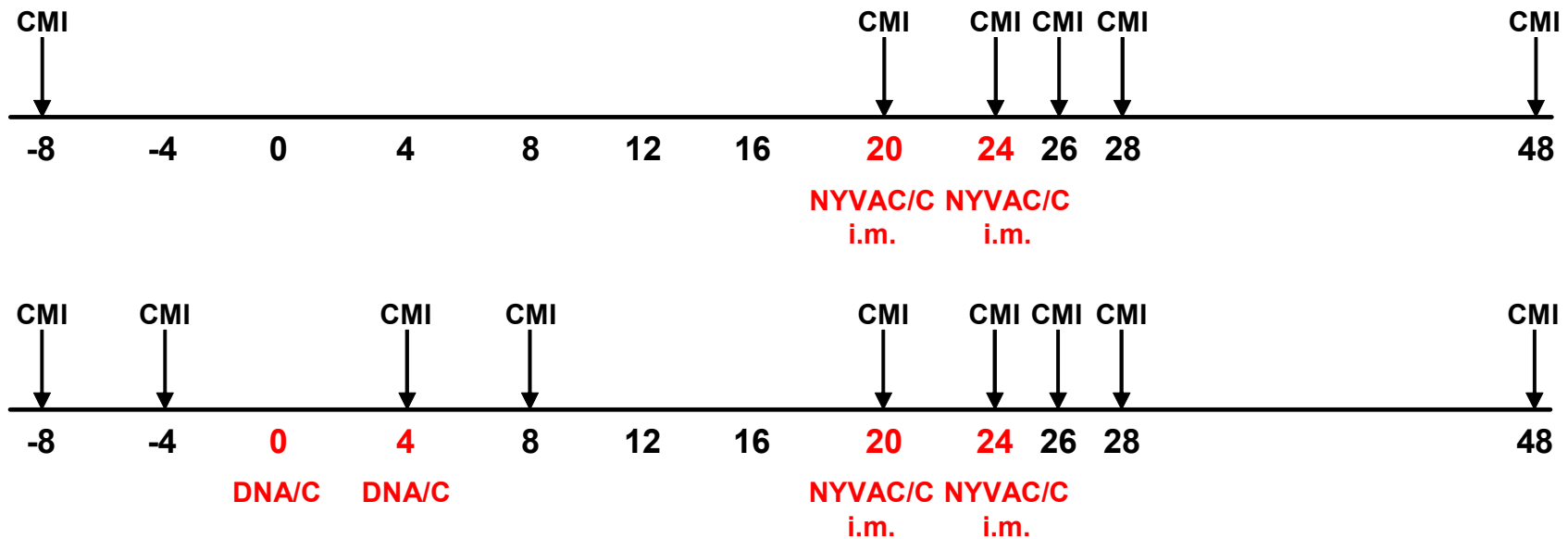
	Clade	Delivery	Status	Developer / Manufacturer
GagPolNef	C (CN54)	DNA-C	GMP, B, T, I, Fill.	Wagner / Wolf / Cobra
Env	C (CN54)	DNA-C	GMP, B, T, I, Fill.	Wagner / Wolf / Cobra
GagPolNef / Env	C (CN54)	NYVAC-C	GMP, B, T, I, Fill.	Sanofi Pasteur
GagPolNef / Env	C (CN54)	MVA-C	GMP in progress	Esteban
GagPolNef / Env	C (CN54)	vTT-C	in progress	Y. Shao / S-CDC
GagPolNef / Env	B	NYVAC-B	GMP, B, T, I, Fill.	Sanofi Pasteur
GagPolNef / Env	B	MVA-B	GMP, B, T, I, Fill.	Esteban



cGMP Immunogens

In vivo Studies in Rhesus Monkeys

Mimicking a Human Phase I Trial



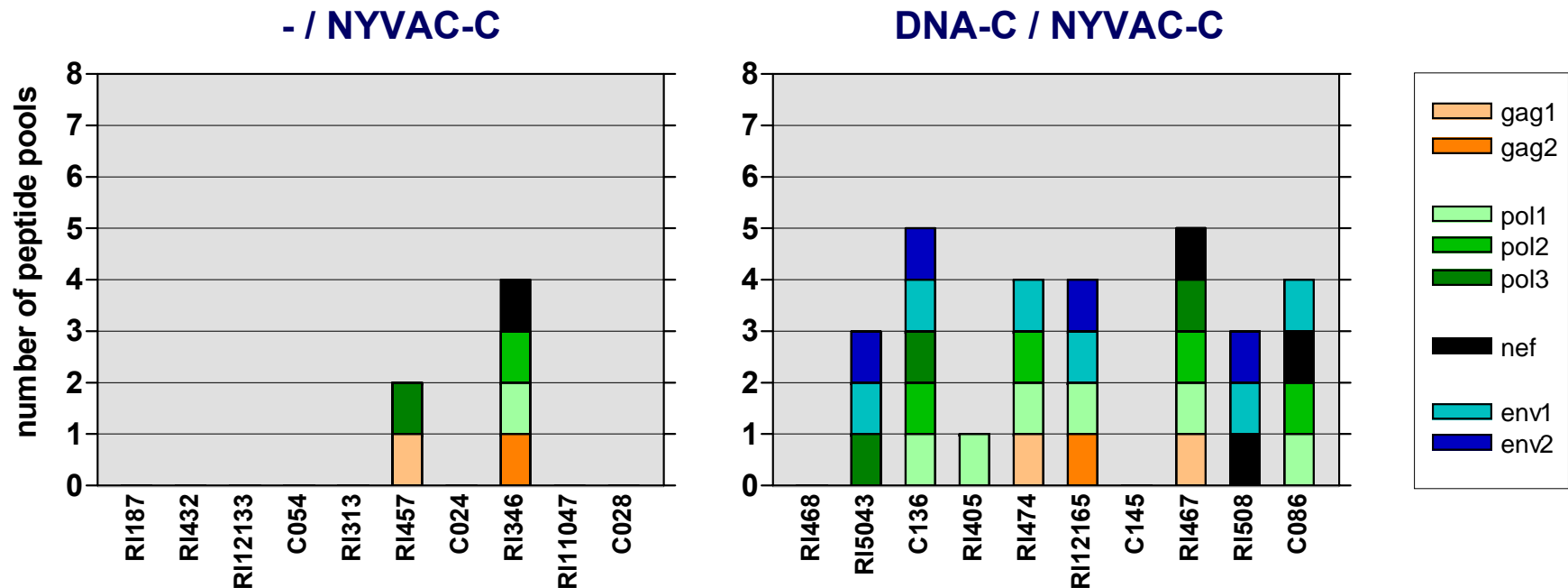
10 monkeys per group



In vivo Studies in Rhesus Monkeys

Mimicking a Human Phase I Trial

Antigen-specific **IFN γ release** at week 24 (4 weeks after 1st NYVAC)



2/10 respond

8/10 respond

several pools recognised

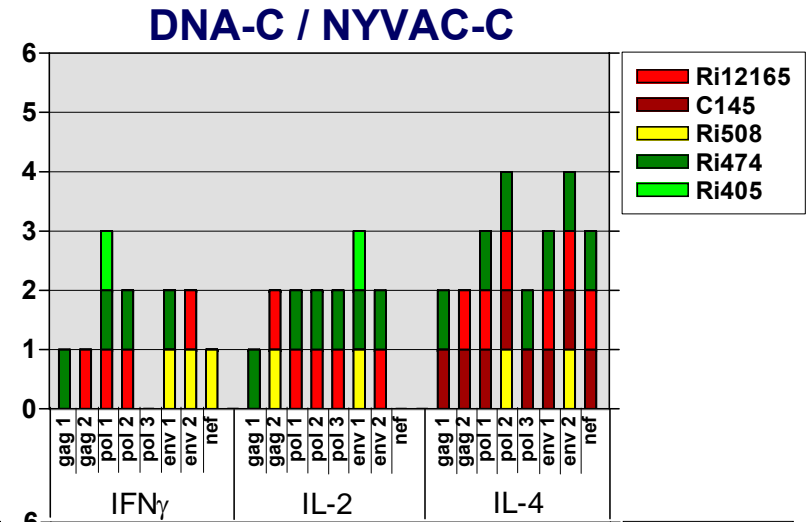
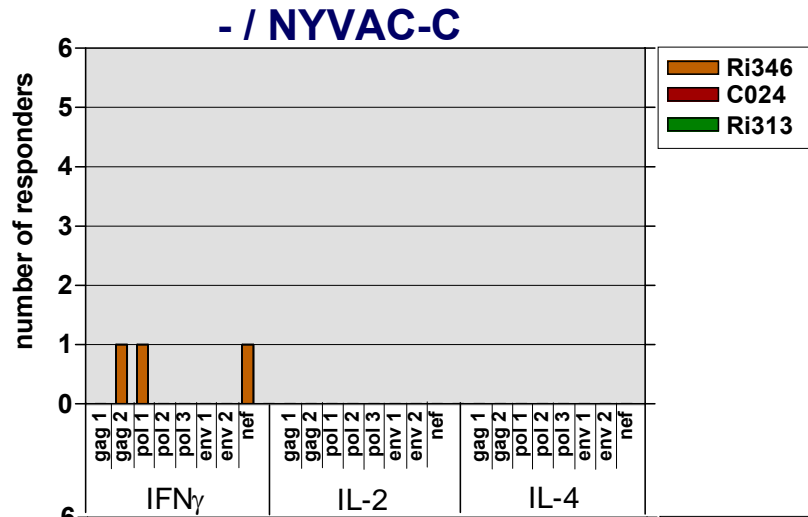


Preclinical Trials

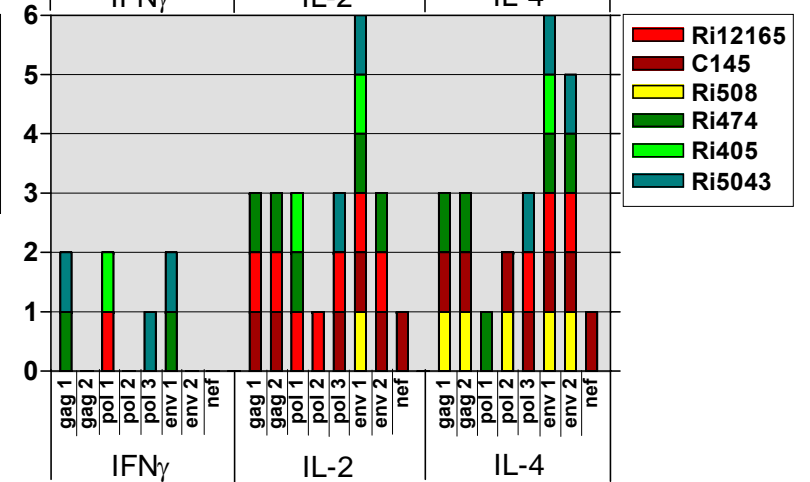
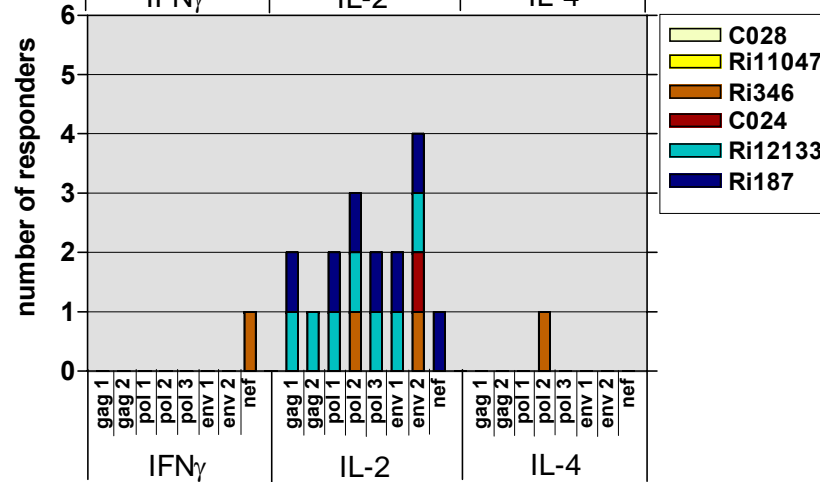
P. Mooij, J. Heeney *et al.*

In vivo Studies in Rhesus Monkeys

week 24



week 26



EV02: Study Design – HIV Negative Volunteers

Objective: Determine the impact of DNA-C versus NYVAC-C alone on priming T-cell responses

Study design:

2 x 20 HIV negative volunteers

A. DNA-C prime (W0, W4; 4mg im), NYVAC-C boost (W20, W24; 2x10E7 im)

B. NYVAC-C (W20, W24; 2x10E7 im)

London (MRC; J.Weber, S. McCormack), Lausanne (CHUV; G. Pantaleo)

Readout (Lausanne):

Several time points

ELISPOT analysis (γ IFN, IL2)

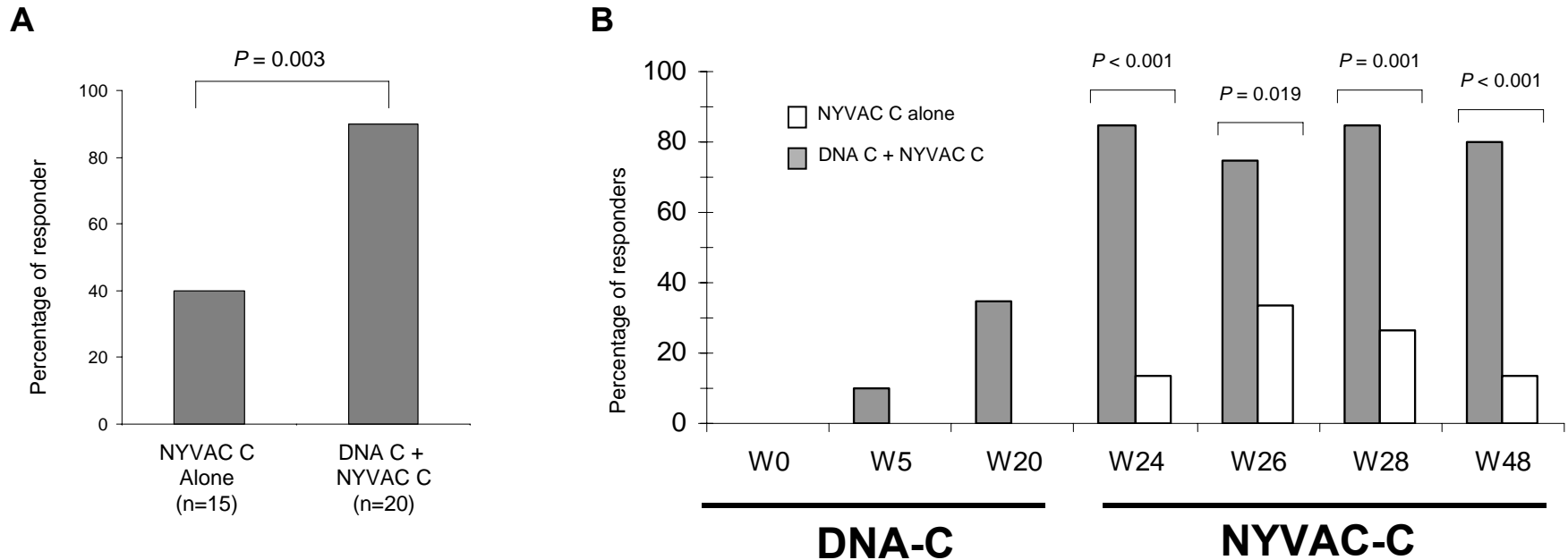
FACS analysis (CD4, CD8, CCR7, IL2, γ IFN)

T cell phenotype / function

Epitope mapping



EV02: Percentage of Responders (γ IFN+ T cells)



Higher percentage of responders in DNA-C prime / NYVAC-C boost group (>90%) compared to NYVAC-C group (<40%)

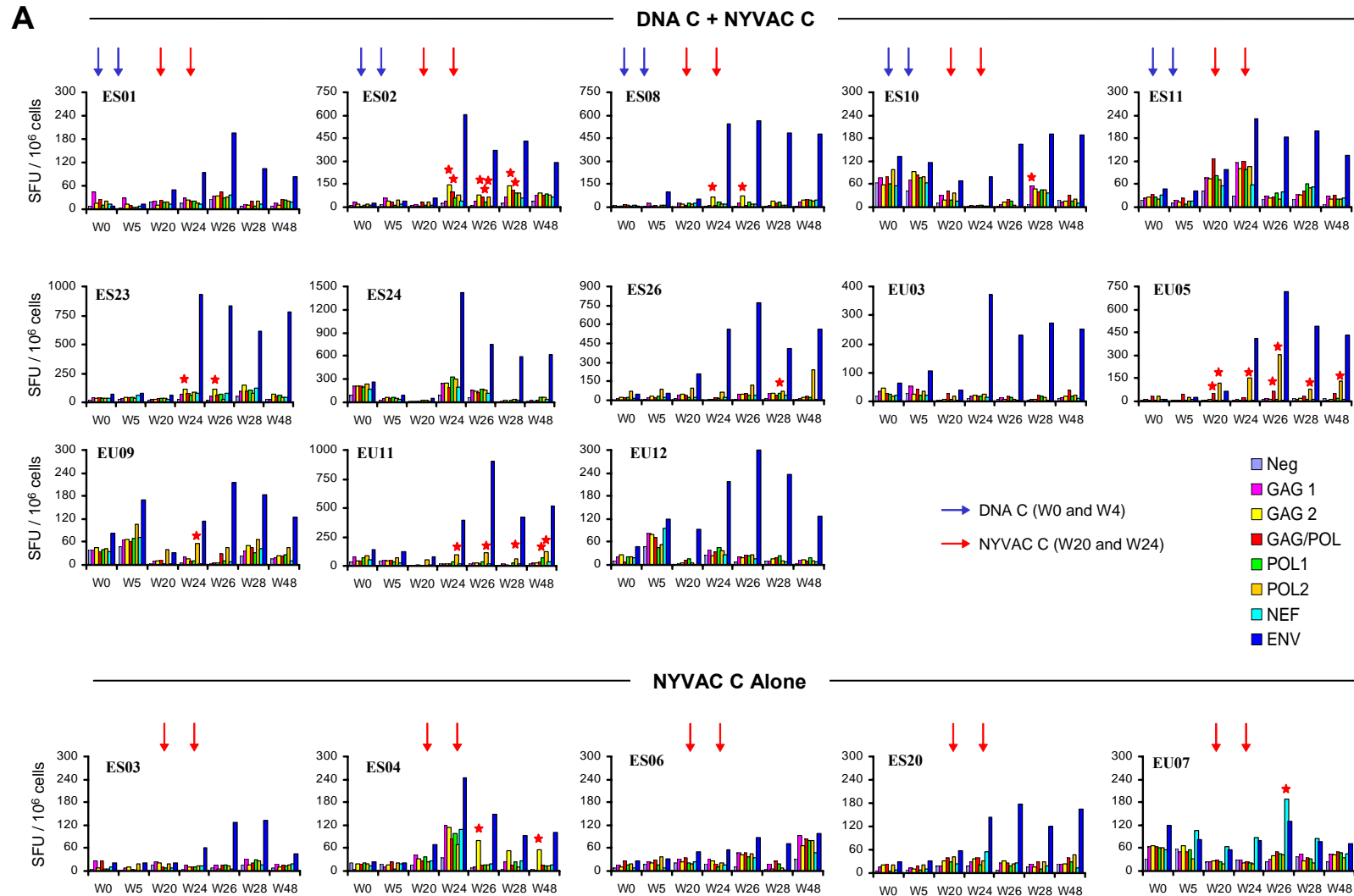
Durability DNA-C / NYVAC-C >> NYVAC-C



Phase I Clinical Trial

A. Harrari, G.Pantaleo *et al.*

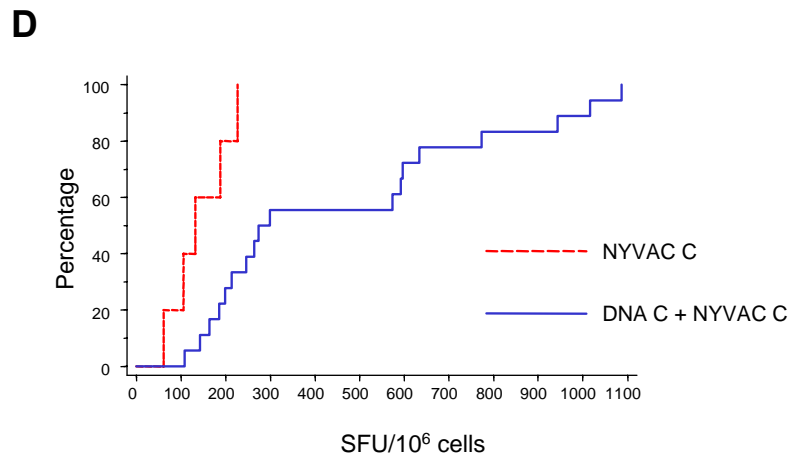
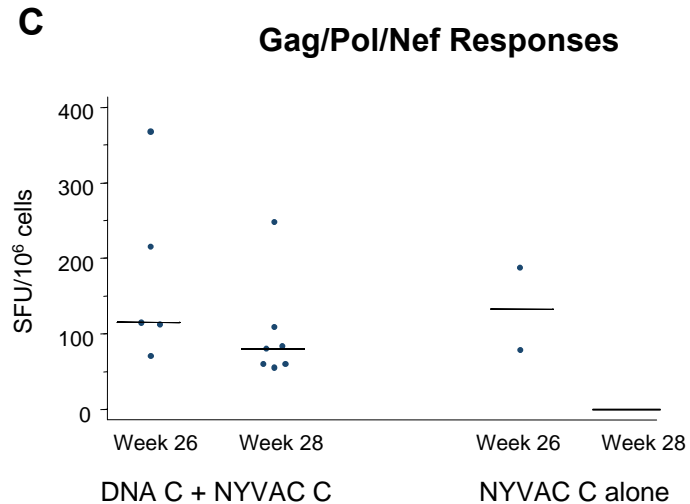
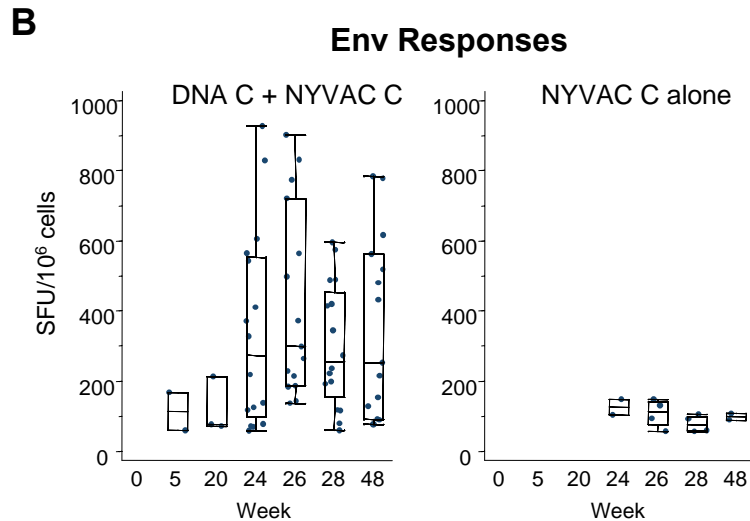
EV02: Magnitude of T cell Responses (γ IFN)



Phase I Clinical Trial

A. Harrari, G.Pantaleo *et al.*

EV02: Magnitude of T cell Responses (γ IFN)



Higher magnitude of T cell responses in DNA-C prime / NYVAC-C boost group (>90%) compared to NYVAC-C group (<40%)

Env >> GagPolNef

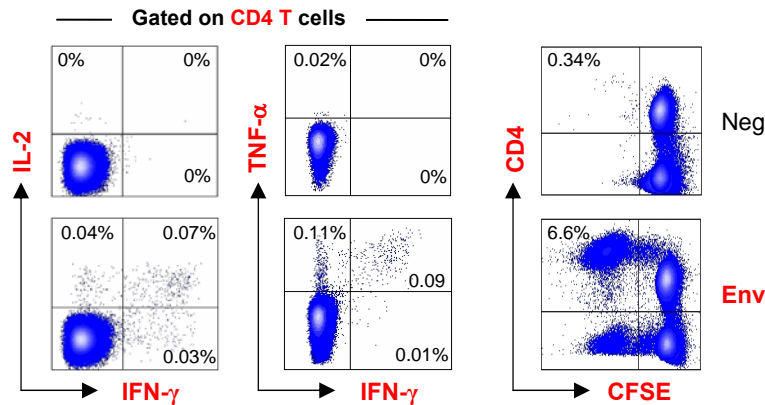


Phase I Clinical Trial

A. Harrari, G.Pantaleo *et al.*

EV02: Functional Analysis of T cell Responses

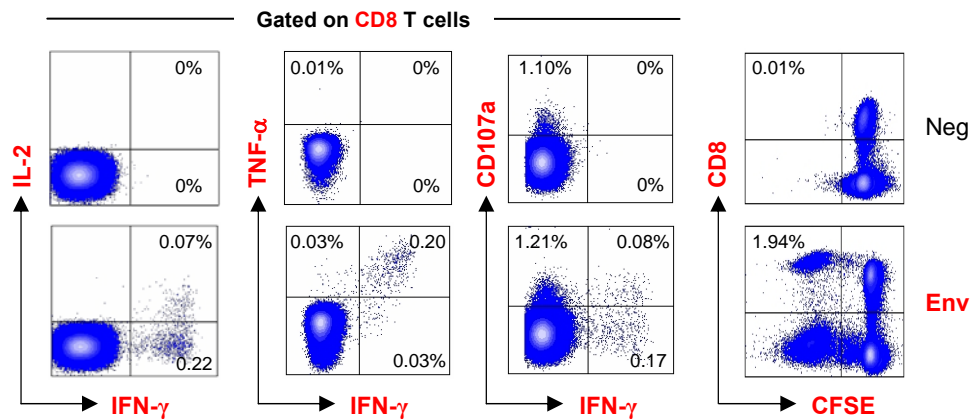
A Volunteer # EU11, DNA C + NYVAC C



CD4 and CD8 responses are polyfunctional

(IL-2, γ IFN, TNF α)

B Volunteer # EU11, DNA C + NYVAC C

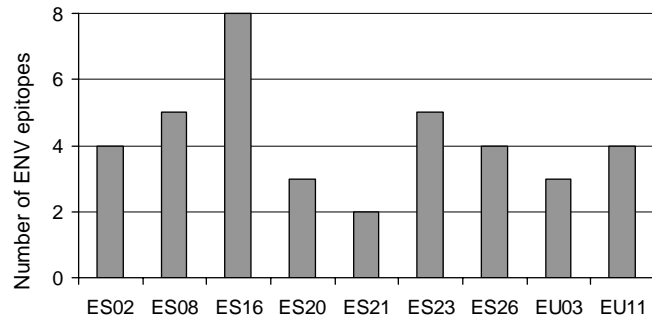


Phase I Clinical Trial

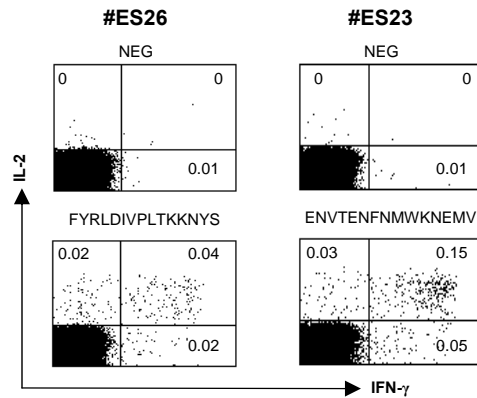
A. Harrari, G.Pantaleo *et al.*

EV02: Epitope specific T cell Responses

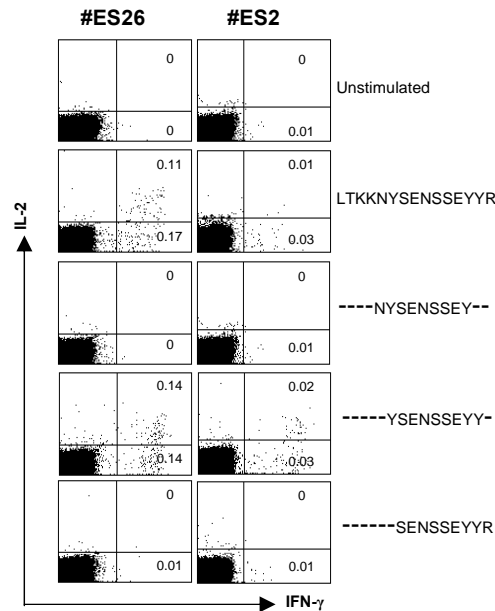
A number of epitopes



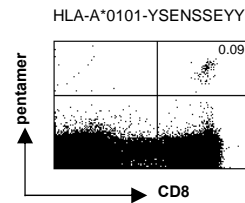
B Gated on CD4 T cells



C Gated on CD8 T cells



D #EU16



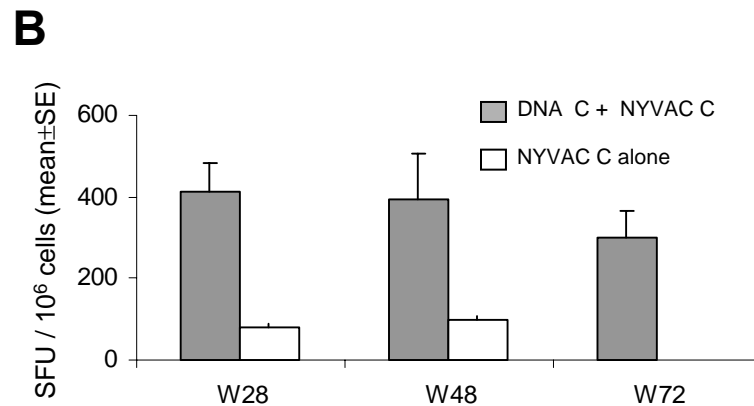
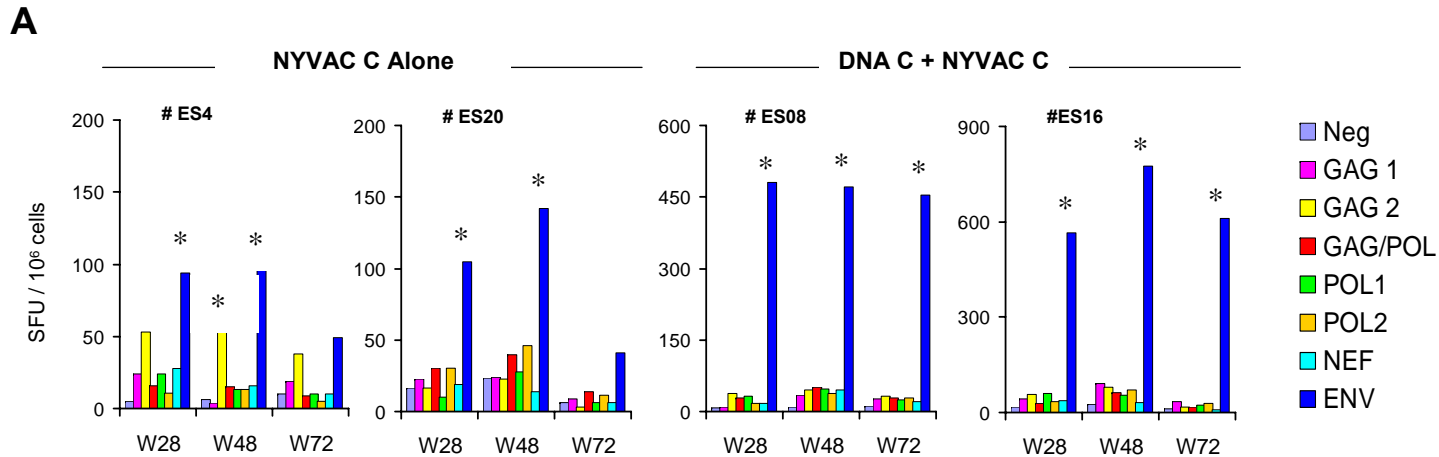
T cell responses are broad (mean 4,2 epitopes) and polyfunctional



Phase I Clinical Trial

A. Harrari, G.Pantaleo *et al.*

EV02: Longevity of T cell Responses (γ IFN, IL2)



Responses are durable

**70% positivity at W72
(DNA-C / NYVAC-C)**

~300 SFU / 10E6 cells at W72



Phase I Clinical Trial

A. Harrari, G.Pantaleo *et al.*

Dominant Env- / Mitigated Gag Responses

Possible explanations:

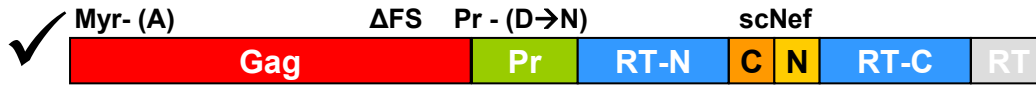
- Intracellular trafficking / localisation: gp120 secreted, GagPolNef not
- Gp120: 55 kDa + glycosylation; GPN: 160 kDa non glycosylated
- Gag expression >>> GagPolNef; SIV-Gag >> HIV-Gag
- Env vs GagPolNef: competition for HLA epitope presentation

Modifications:

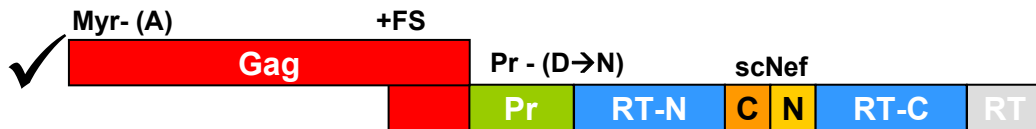
- Altered immunization regimen: 3 x DNA-C (mixture DNA-GPN and DNA-Env) followed by 1 x NYVAC-C (EV03) instead 2 x DNA-C followed 2 x NYVAC-C (EV02)
- At least 2 different NYVAC viruses: Env + GagPolNef instead Env / GagPolNef; different inoculation sites
- Increase immunogen processing and epitope presentation → DRIP
- Improve cross-presentation: VLP / secreted Gag derivatives
- Targeting to DCs (e.g. Dec205 targeting, FcR targeting)
- Incorporate molecular adjuvants (CD40L, CTB...)



Outlook: BMGF-PTVDC Discovery Group



reference construct



allow ribosomal frameshifting
(Gag 95 / PolNef 5)



allow ribosomal frameshifting
(Gag 95 / PolNef 5), VLP budding



allow ribosomal frameshifting
No NEF, (Gag 95 / Pol 5)



allow ribosomal frameshifting
No NEF, VLP budding



2 distinct ORFs



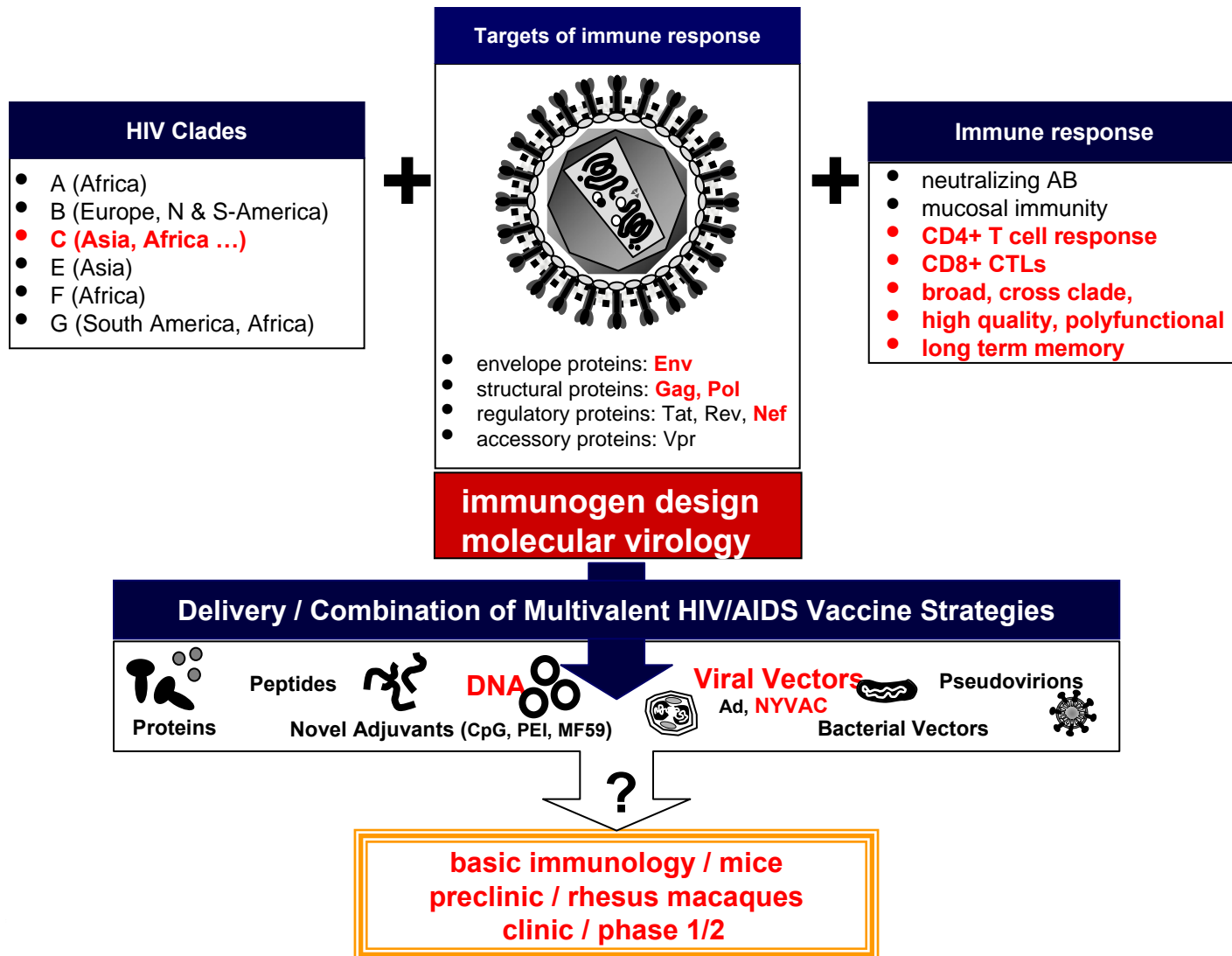
secretion, DC targeting

➔ best in class: increase expression & improve access to DRIP pathway



Immunogen design

What we have achieved – Where we need to go



Acknowledgements

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