



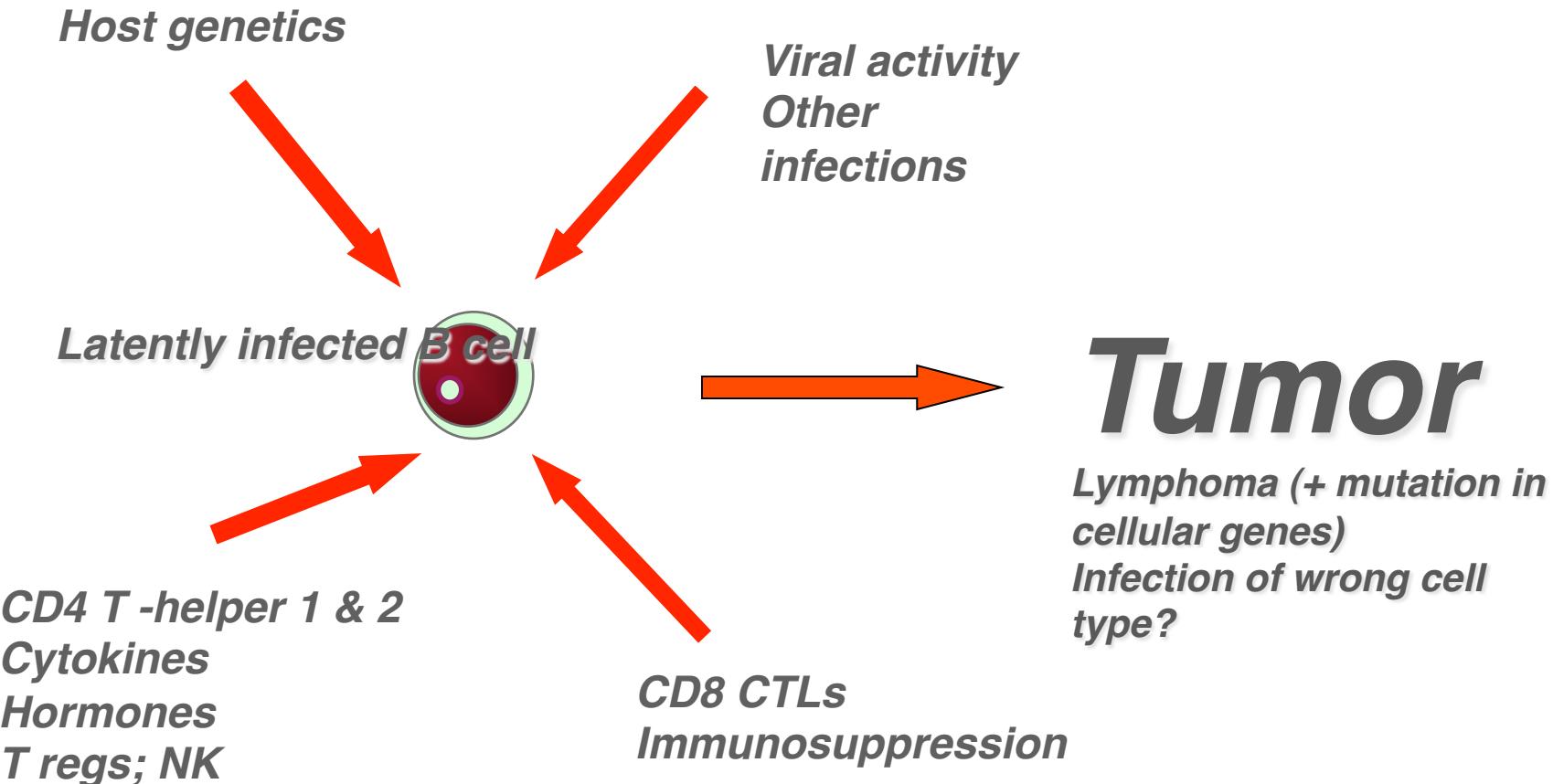
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# What is the most efficient immunotherapy to Epstein-Barr virus infection?

Ingemar Ernberg  
Riga  
May 2016

Dept of Microbiology, Tumor and  
Cell Biology (MTC)

# Control and disruption of EBV latent infection





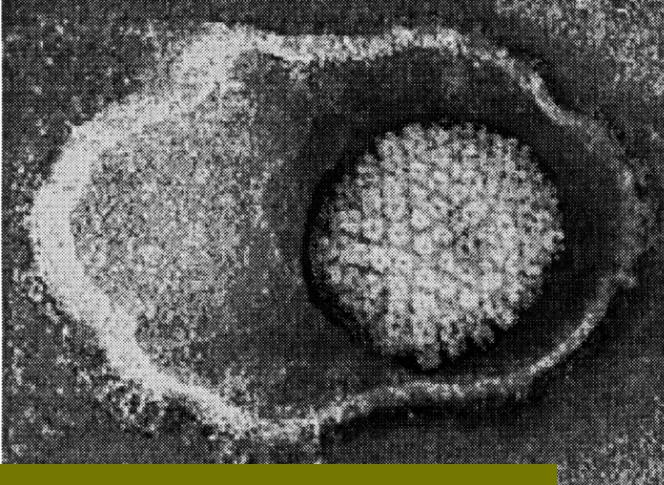
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# ***Prevention/vaccination or Immunotherapy***

Why? Who? When? How?

# *My points*

- ◆ EBV & infection *in vivo*
- ◆ Vaccine?
- ◆ Tumors
- ◆ High risk: transplant pat:s (& HIV-carriers)
- ◆ T-cell therapy



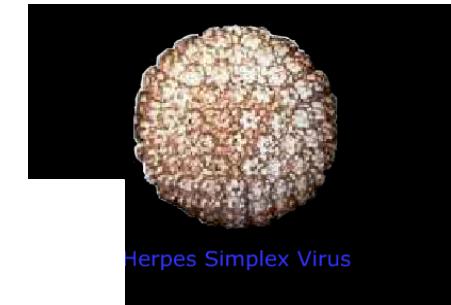
# Epstein-Barr virus

## *The paradoxes*

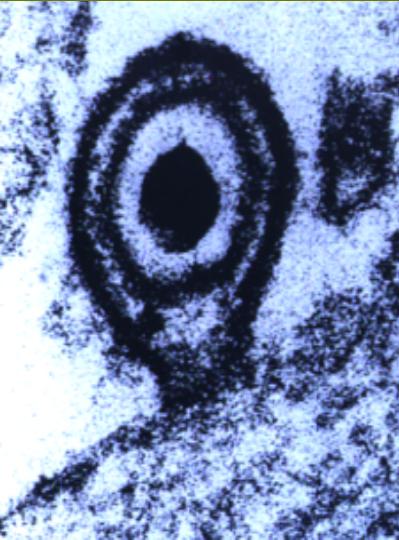
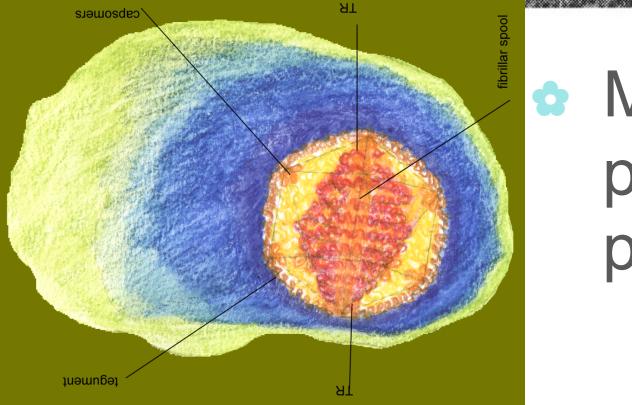
## *EveryBodies Virus*



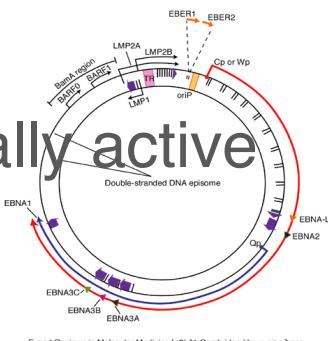
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Herpes Simplex Virus



- ✿ Most common virus in the human population (>90% of all adults, life long persistence)
- ✿ Tumor-associated virus
- ✿ 80% of carriers secrete biologically active virus (saliva)
- ✿ Very efficient in immortalizing B-cells



Expert Reviews in Molecular Medicine 12/31 © 2016 Cambridge University Press

Emberg 2016<sup>(2001)</sup>



# ***Stability***

## ***A suitable vaccine target***

- ◆ 50+ complete DNA-sequences
- ◆ Two major types: type I=type A and type II=type B (98 % homology)
- ◆ Subtypes ill-defined: a chinese?, deletions in critical genes (LMP 1)
- ◆ Variants: appr = familiar isolates (recombination between large repeats in latency genes)



EBV(s) has(ve) adapted over millions of years to a very successfull life within the immune system.

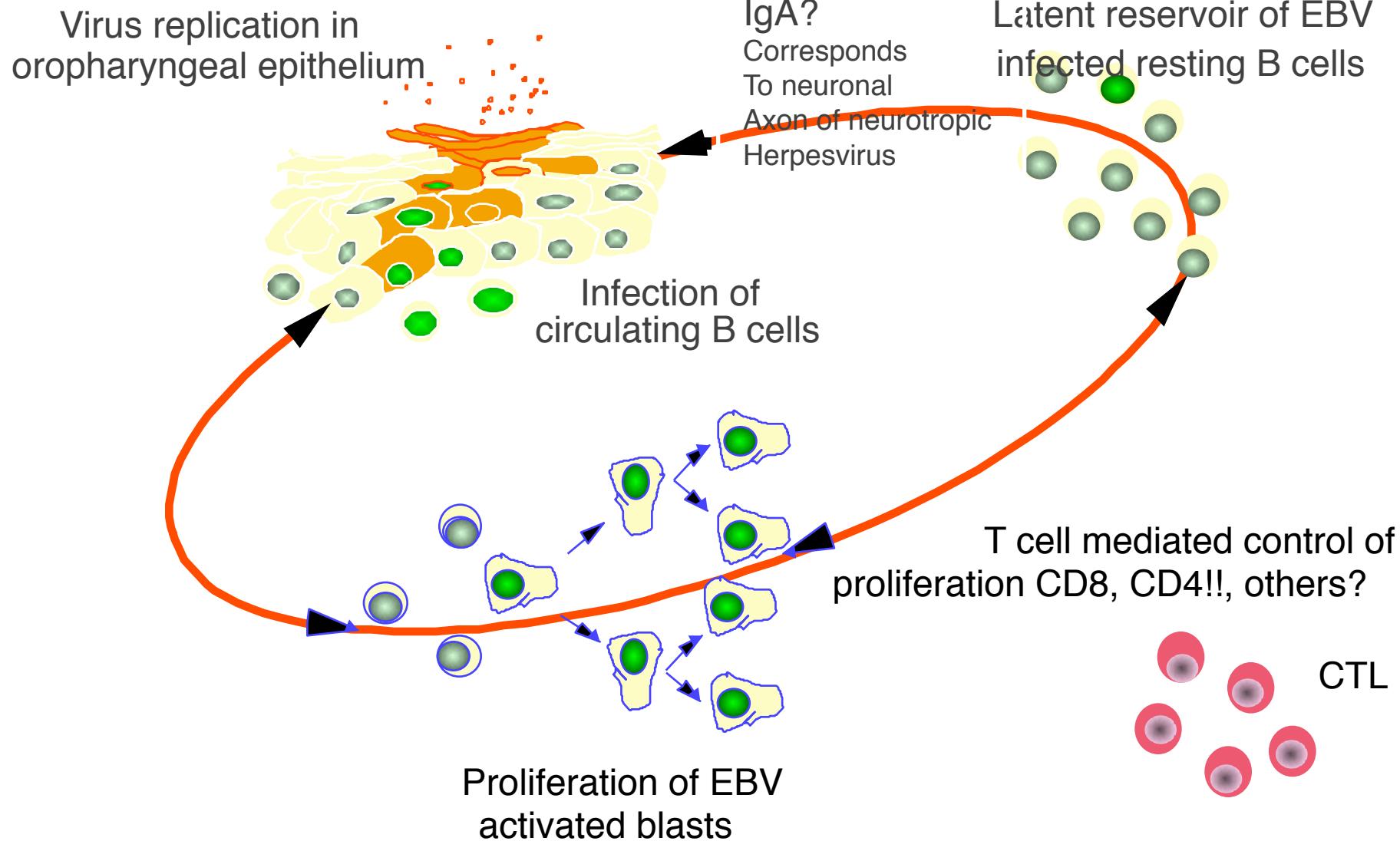
The viral strategy for survival, latency and spread in the population has evolved numerous mechanisms to master this challenging situation.

EBV-associated tumors are a rare trade off of this normally balanced virus-host interaction

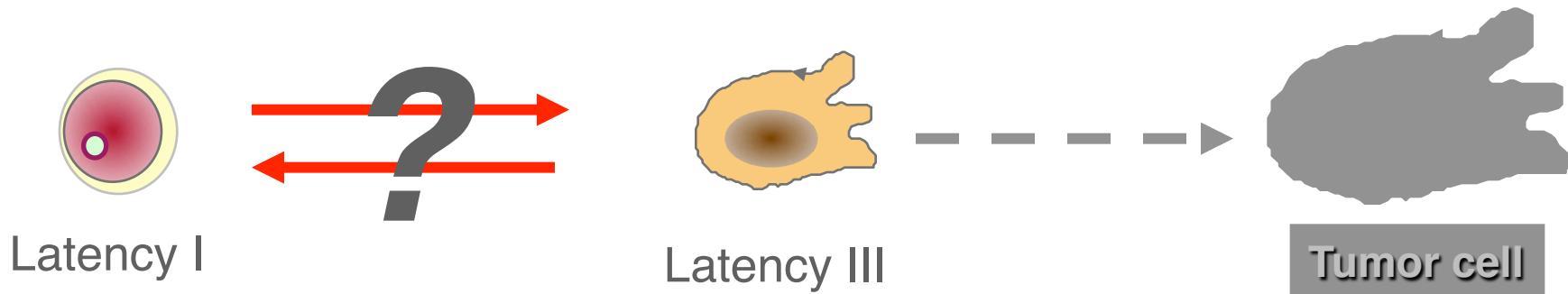
# EBV life cycle *in vivo*



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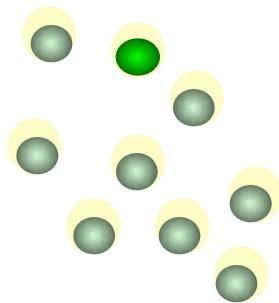
# *Control of the latency-switch, latency I to III and reverse*



# Model system for transformation in vitro

- HBV/HCV -
- HPV (+)
- EBV ++
- HHV 8/KSHV -
- HTLV +
  
- *H.Pylori* -
- Parasites (liver fluke,  
*Schistosomiasis*) -

## *Latency in vivo*



One infected B-lymphocyte/  
100ul-10 ml of blood  
(0,1-10/100.000 B-cells)

5000 infected cells in blood in a  
healthy carrier (+ X in lymphoid  
system)

Increases 100-10.000-fold in high  
risk groups

# *Host-virus interaction & diseases*



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- ◆ Childhood infection *subclinical*
- ◆ Primary infection > 5 yrs 35% risk of *infectious mononucleosis* (“an acute autoimmune syndrome”)
- ◆ In immunosuppressed: (Fatal/Chronic) *lymphoproliferative syndrome*
- ◆ *Lymphomas, epithelial cancers* as a result of accidents of the host-virus interaction

## To think about

Most biologically active EBV on earth is swallowed with the saliva.

300-500 ml/day, appr 500 infectious units entering the GI every day in 80% of humans.

“Part of the normal flora”

Role in infection, immune boosting, and pathogenesis?

# Infectious agents

## Number of new tumors annually & globally

- HBV/HCV 500,000
- HPV 450.000
- EBV 200.000
- HHV 8/KSHV 45.000
- HIV 10.000
- HTLV 2.200
  
- *H.Pylori* 460,000
- *Cl.psitacci, Borrelia burgdorfei,*  
*Campylobacter jejenum* 500?
- Parasites (liver fluke,  
*Shistsomiasis*) 11,000

# EBV-associated tumors in man

## *Lymphoid tissues*

Burkitt's lymphoma, endemic	98%
Burkitt's lymphoma, sporadic	30%
AIDS-immunoblastic lymphoma -in CNS	30% 100%
Post-transplant lymphoma	100%
Hodgkin's lymphoma	50%
T-cell lymphomas --lethal midline granuloma	10-30% >90%
Primary effusion lymphomas	>90%

## *Epithelial tissues*

Oral hairy leukoplakia	100%
Gastric adenocarcinoma	5-10%
Nasopharyngeal carcinoma, undifferentiated	100%
Salivary gland carcinomas	<100%

## *Smooth muscle tissue*

Leiomyosarcoma in immunosuppressed	100%
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# EBV-associated tumors in man

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# *Geographic distribution of EB virus-associated tumors*



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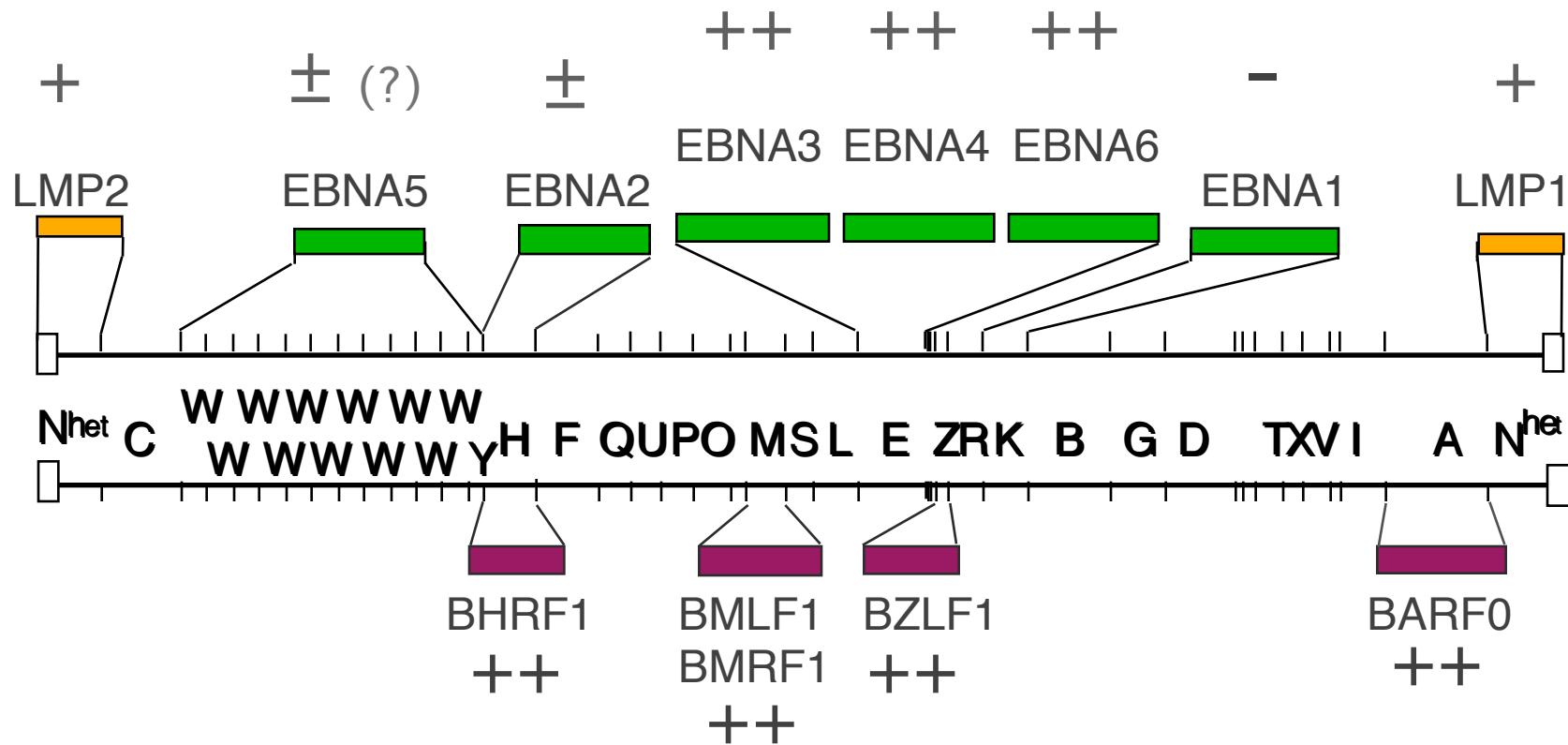
# EBV - therapeutic strategies

- ◆ Conventional cancertherapy
- ◆ Anti-viral drugs: spec DNA-polymerase inhibitors (acyclovir etc.) clears replication but not latent infection.  
Effect on lymphoproliferative syndrome
- ◆ Bone marrow transplantation can clear out the whole EBV-infection including latency
- ◆ Therapeutic vaccines
- ◆ Adoptive transfer of EBV-specific cytotoxic T-lymphocytes (CTL) reeducated in vitro  
(Check point-blockers PD1/CTLA4)

# *EBV specific cytotoxic T-cell (CTL) responses*



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- polyclonal responses preferentially directed against EBNA3,4,6 and antigens of the lytic cycle
- strongly focused on certain MHC:peptide combinations
- endogenously expressed EBNA1 is not recognised



# Components of immunity to EBV

- ◆ Antibodies are efficiently neutralizing infection (anti-gp 350; anti-MA)
- ◆ Antibodies + complement/LAK-cells eliminate virus replicating cells
- ◆ Earlier dogma: Cytotoxic T-lymphocytes are instrumental in controlling the life-long latency (Masucci, Rickinson, Moss)
- ◆ Recent discovery: CD4 cells producing specific cytokines (IL21) control the latent infection (Nagy, Kis, Klein; KI)



# *Receptors*

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B-cells: C3d (CR2)-receptor, CD 21  
MHC Class II (co-receptor)

Epithelial cells: MHC Class II  
Unorthodox mechanisms

Virus envelope receptor: gp 350/220

# EBV vaccine

## Primary preventive

- ◆ Recombinant gp350, major envelope glycoprotein, receptor binding, neutralizing epitope (prevents lymphomas in monkey models)
- ◆ To be used when, where and how? Risk groups for NPC, EBV-negative adolescents (IM)



# EBV vaccines....

## Therapeutic

- ◆ “Polytope”, several MHC Class I epitopes combined in a vaccinia-vector based vaccine (D.Moss et al)
- ◆ Peptide-loaded dendritic cells (NPC; A.Rickinson et al)
- ◆ Adoptive cytotoxic T-cell transfer (C.Rooney; M.Masucci et al)
- ◆ Recombinant gp350, major envelope glycoprotein, receptor binding, neutralizing epitope (therapeutic in post-transplants?; A.Morgan et al +CRC, UK)
- ◆ CTL modulation combined with demethylation (Rich Ambinder et al.)?

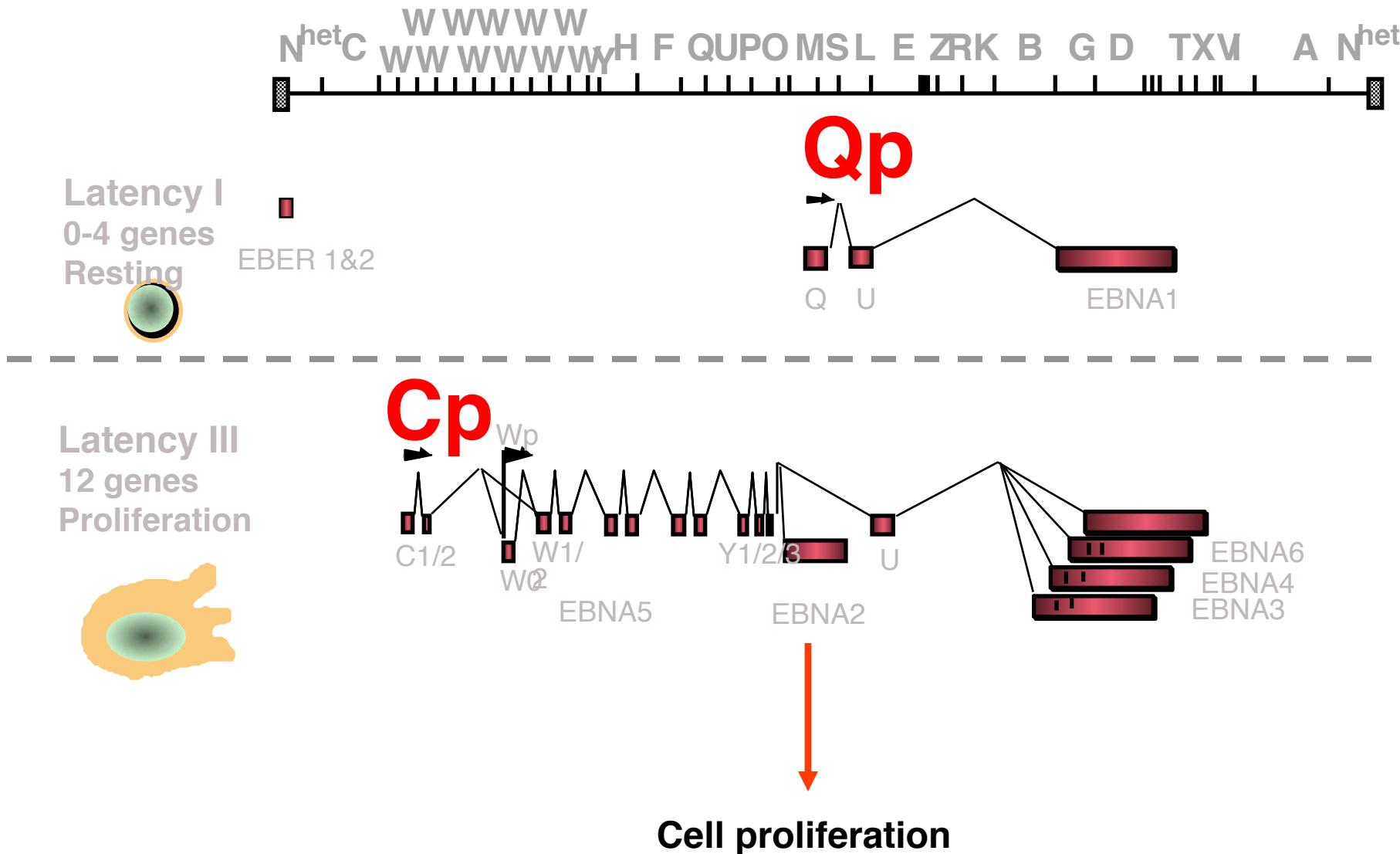
# Groups with high risk for EBV positive lymphomas

- Bone-marrow/HSC transplanted pat:s (with minor mismatch)
  - HIV-carriers (before HAART; 15-20 % CNS lymphomas)
- 
- Children with chronic malaria
  - Immunodisregulation+Chronic antigen stimulation

# Proliferation-rest is regulated by viral expression programs in latency



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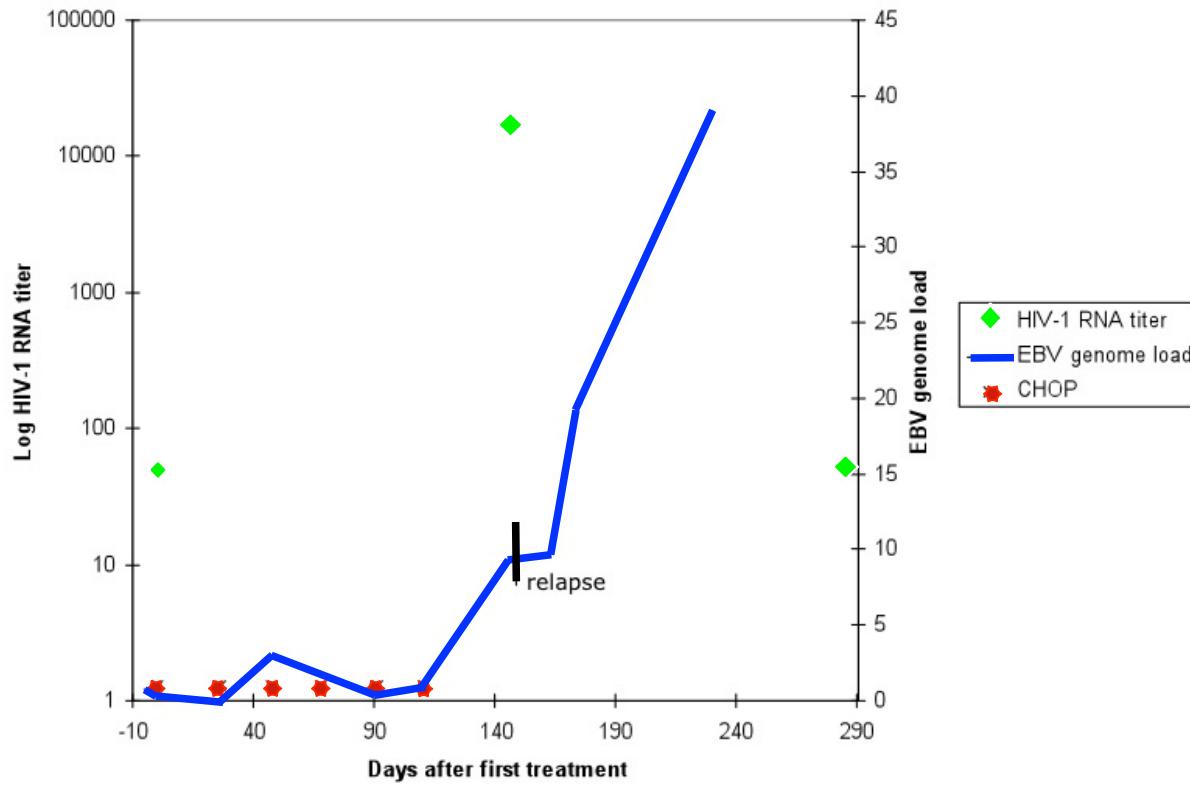
# Incidence of post-transplant lymphoproliferative disease (PTLD) depending on type of transplant

- Kidney 1 %
- Liver 2 %
- Heart 3 %
- Lung 1 – 8 %
- Heart and lung 9 %
- Intestine 7 – 11 %
- Multivisceral 13 – 33 %
- BMT 1 % (up to 30 %)

# Risk factors for post-transplant lymphoproliferative disease (PTLD)

- Mismatched, unrelated stem cell grafts
- Primary immune deficiency
- T-cell depletion techniques which do not concomitantly reduce the B-cell content
- Use of anti-T-cell antibodies ( ATG, OKT-3 )
- Heavy immune suppression
- High levels of circulating EBV-infected cells \*
- EBV-seronegative recipient
- CMV infection / Young age ?/ GVHD treatment

# EBV-DNA load and relapse in EBV-associated lymphoma (HIV-infected patient)

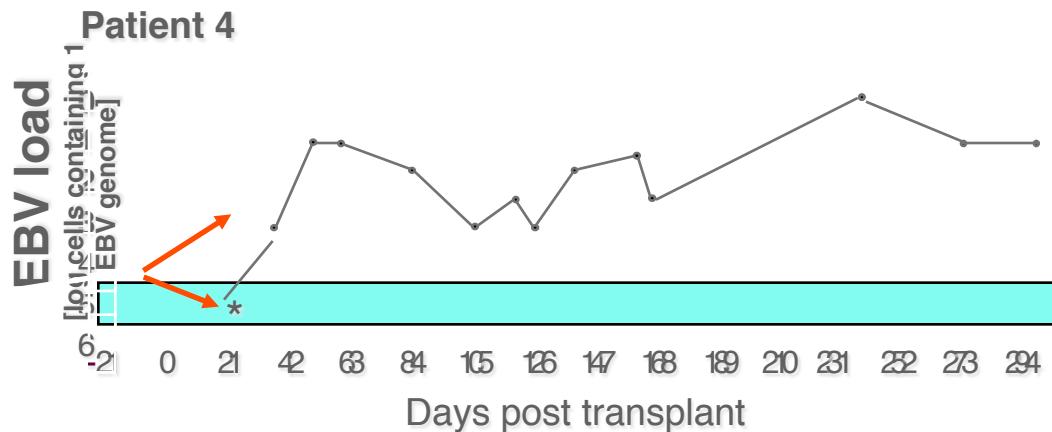
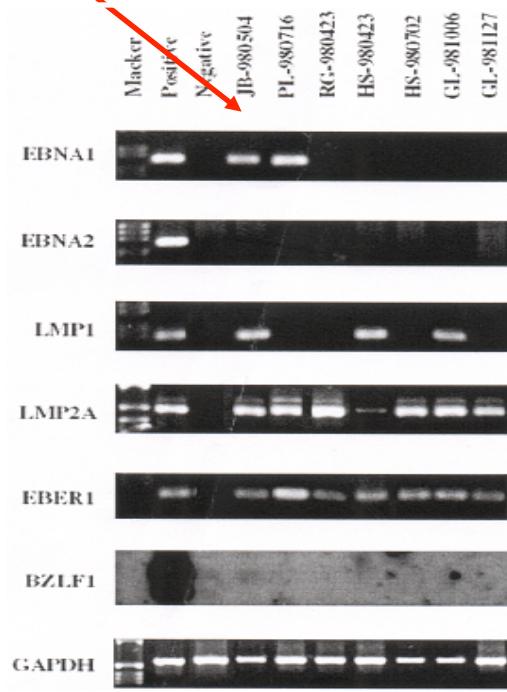


## Increased EBV load in patients receiving T-cell depleted BMT

Several log-fold increase of EBV-load post-BMT...

.... related to T-cell depletion of graft

....only latency 1 & 2 (restricted, non-proliferating in blood)



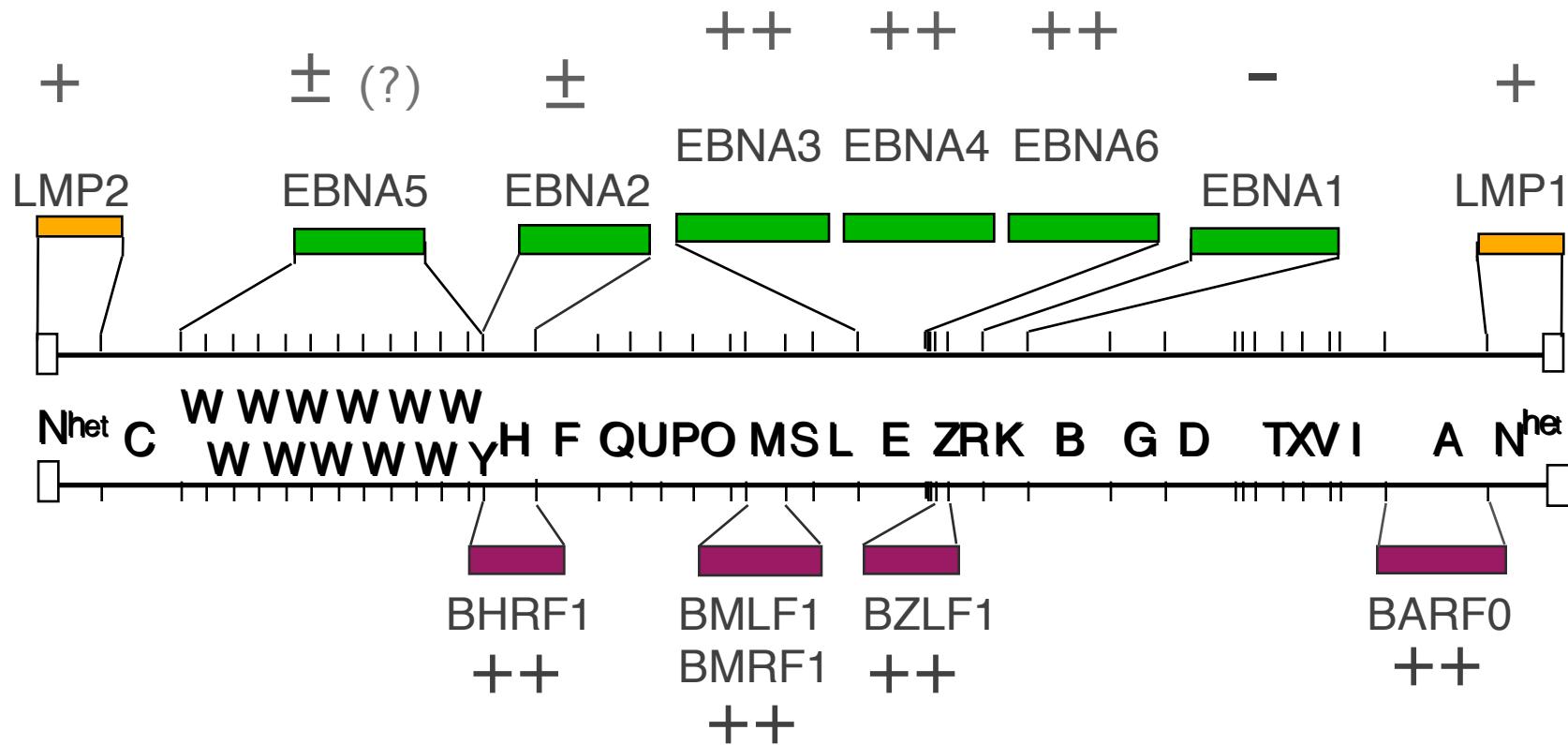
Maximal EBV load recorded in BMT recipients with high or low risk of PTLD

Donor	Type of BM graft	Maximal EBV load (genomes/10 <sup>6</sup> cells)	Days post BMT
<b>High risk patients</b>			
JB <sup>a</sup>	TCD	1,250	31
UH	TCD	6,250	129, 149, 366
PL	TCD	31,250	53, 63
OL	TCD	31,250	49, 125
RG	TCD	156,250	42, 52
GL	TCD	31,250	80
HS <sup>a</sup>	U	156,250	80
mean±SE		59107±25963	
Median		31250	
<b>Low risk patients</b>			
MU	U	50	34
NJ	U	250	51, 62
SG	U	1250	62, 79, 93
RT	U	50	95
EE	U	1250	55
PZ	U	1250	335
LS	U	250	13, 27, 53, 62, 73, 81
GS	U	1250	67
mean±SE		700±269	
Median		750	

# *EBV specific cytotoxic T-cell (CTL) responses*



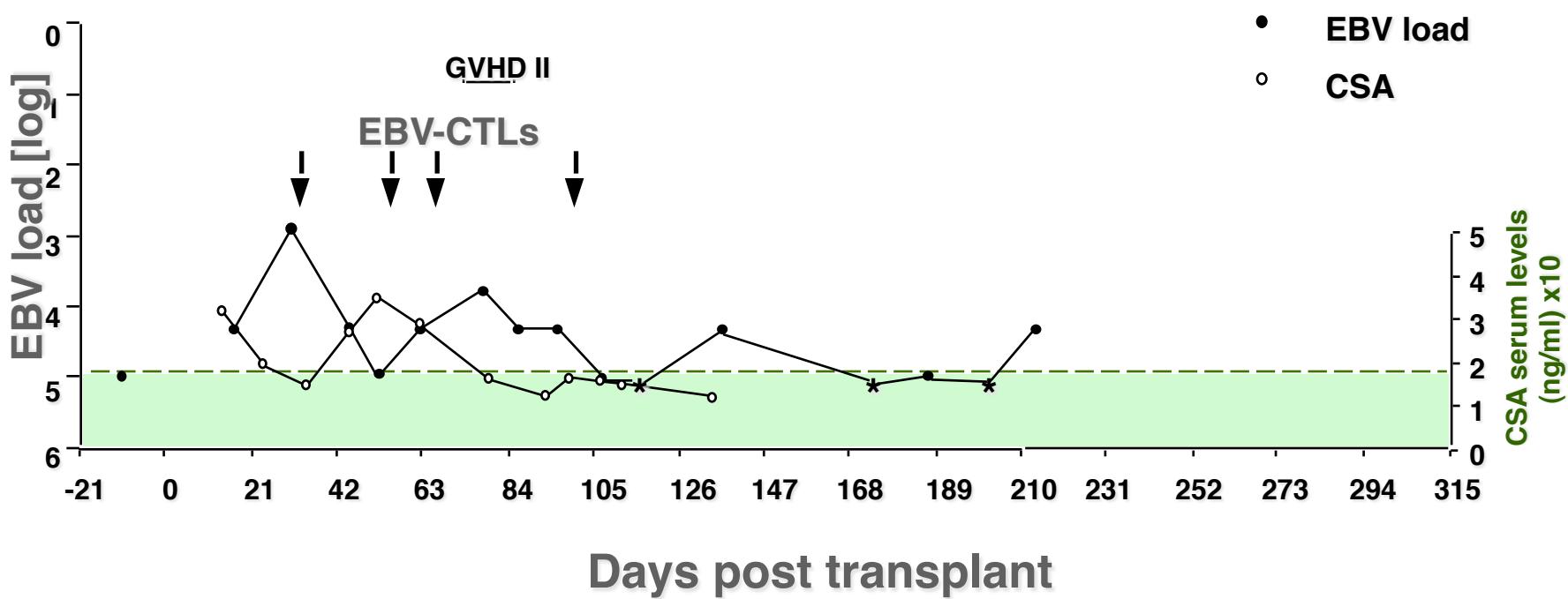
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- polyclonal responses preferentially directed against EBNA3,4,6 and antigens of the lytic cycle
- strongly focused on certain MHC:peptide combinations
- endogenously expressed EBNA1 is not recognised

## *Early administration of EBV-CTLs prevents the increase of EBV load*

Patient 5



# **Study of "EBV" at transplant unit of Karolinska University Hospital South (Huddinge)**



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## **Question**

Can we identify better tools for PTLD risk prediction at individual patients level?

T cell phenotyping on EBV-related immune reconstitution (tetramers, FCM)

## **Cohort**

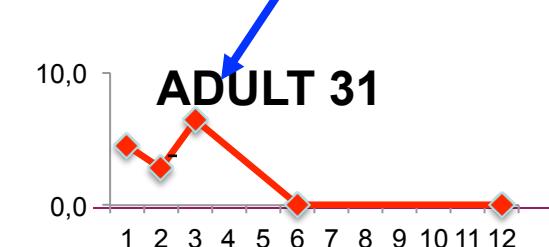
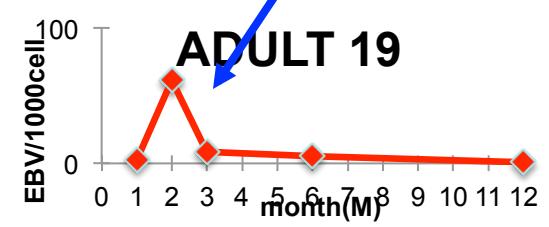
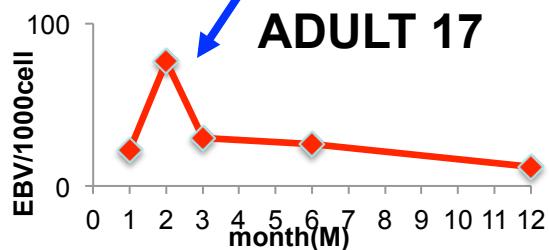
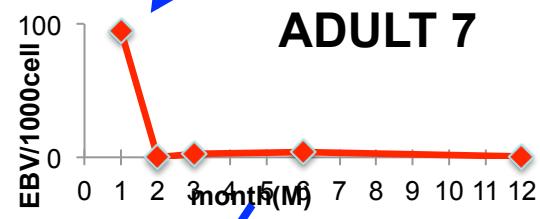
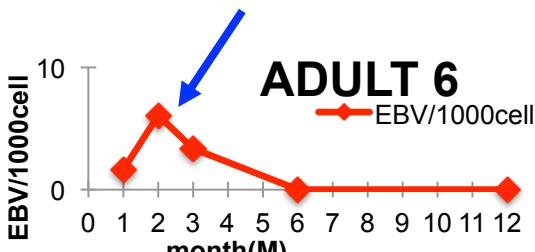
Entered so far: 350 adults & 50 children entered (open)

Sampled 1,2,3,6, 12 and 24 months > transplantation

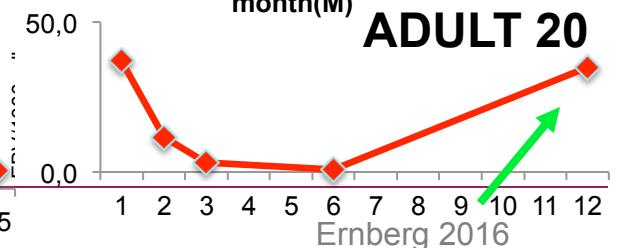
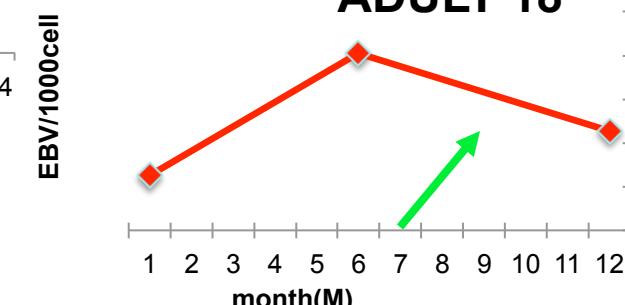
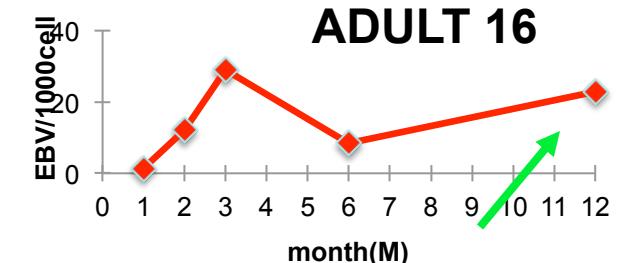
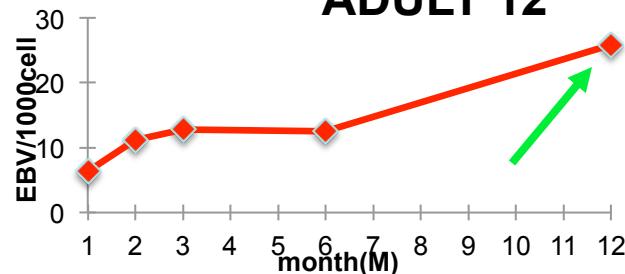
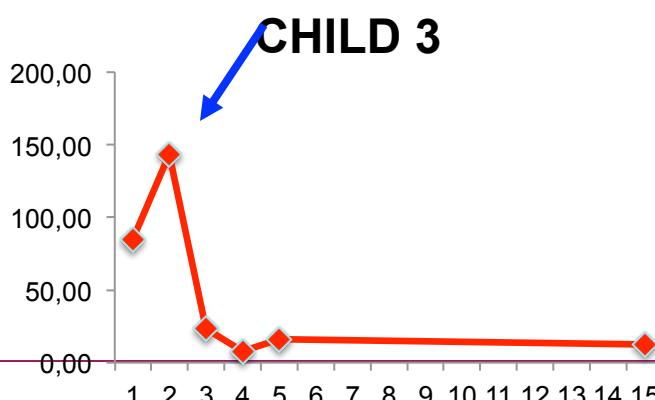
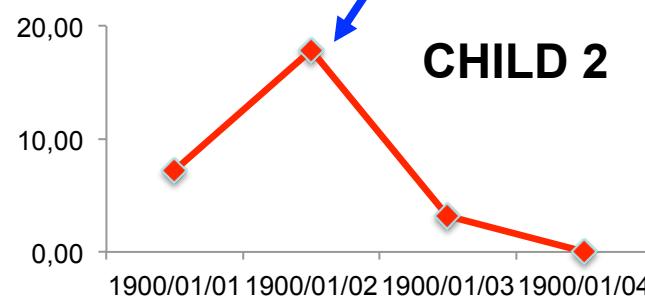
## **Colaborators**

Markus Maeurer, Olle Ringdén, Johan Matsson, Jacek Winiarski,

## EBV DNA load over time: TWO patterns



← Type I  
Type II →



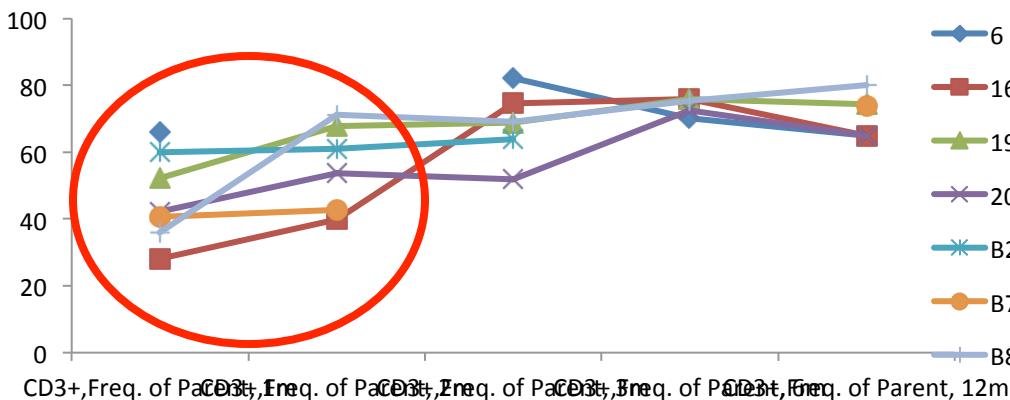
Ernberg 2016

# Fraction CD3+ cells

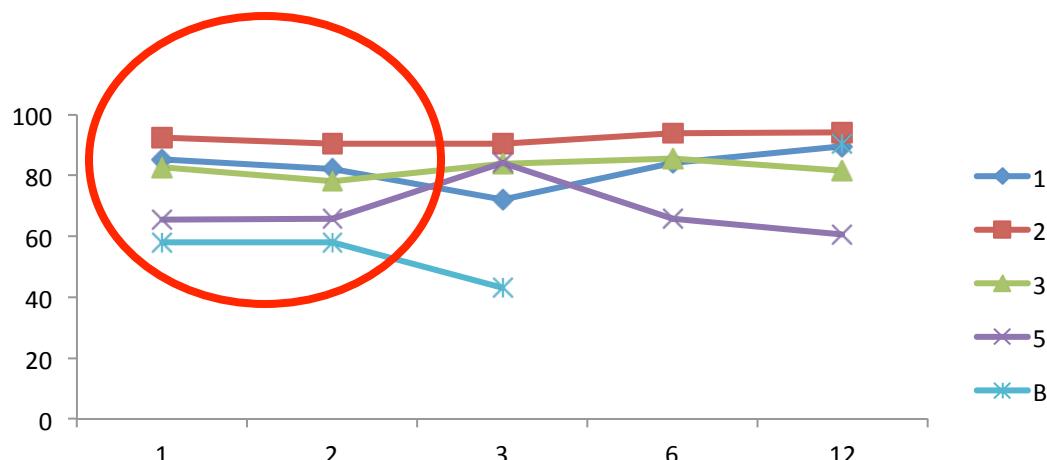


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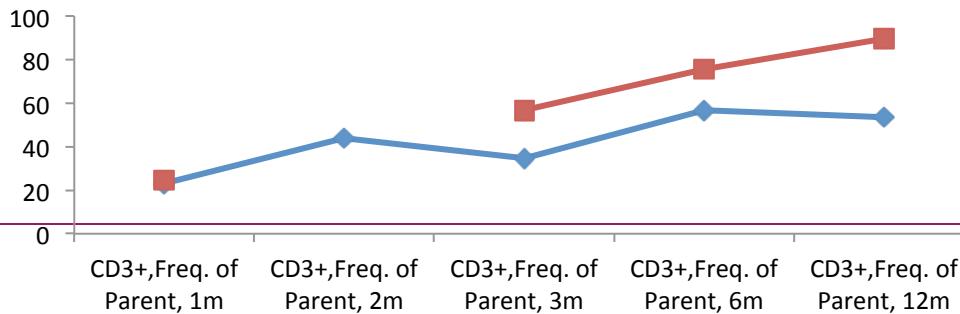
Intermediate



Low



High



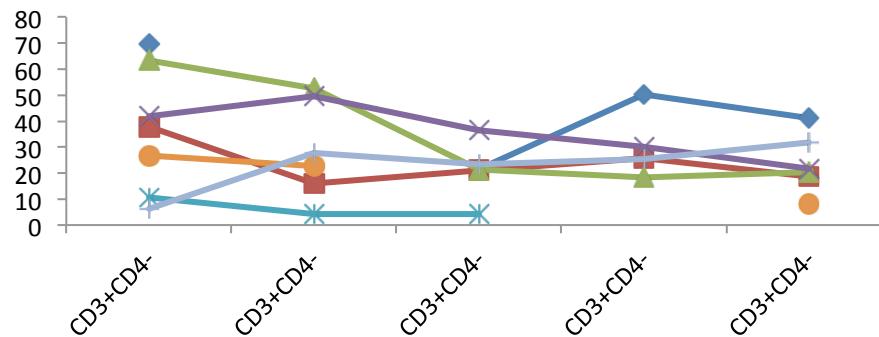
Ernberg 2016

# CD4-CD8- T cells

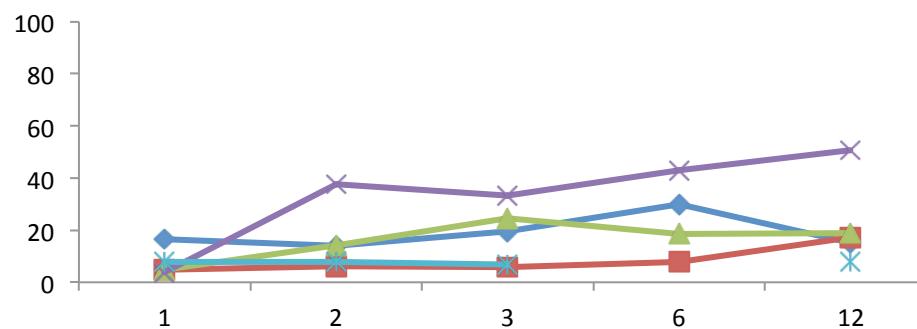


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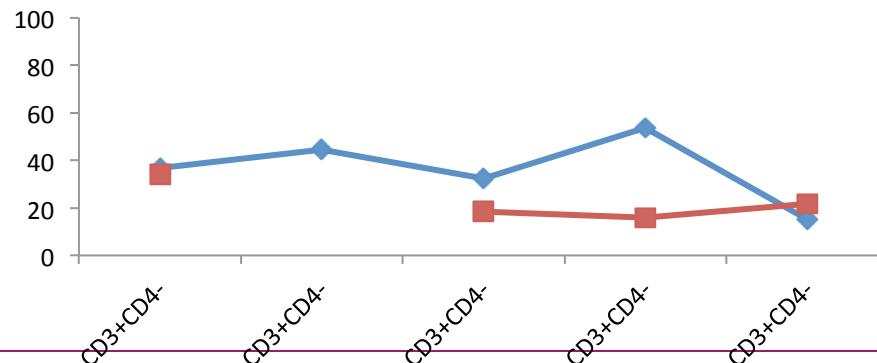
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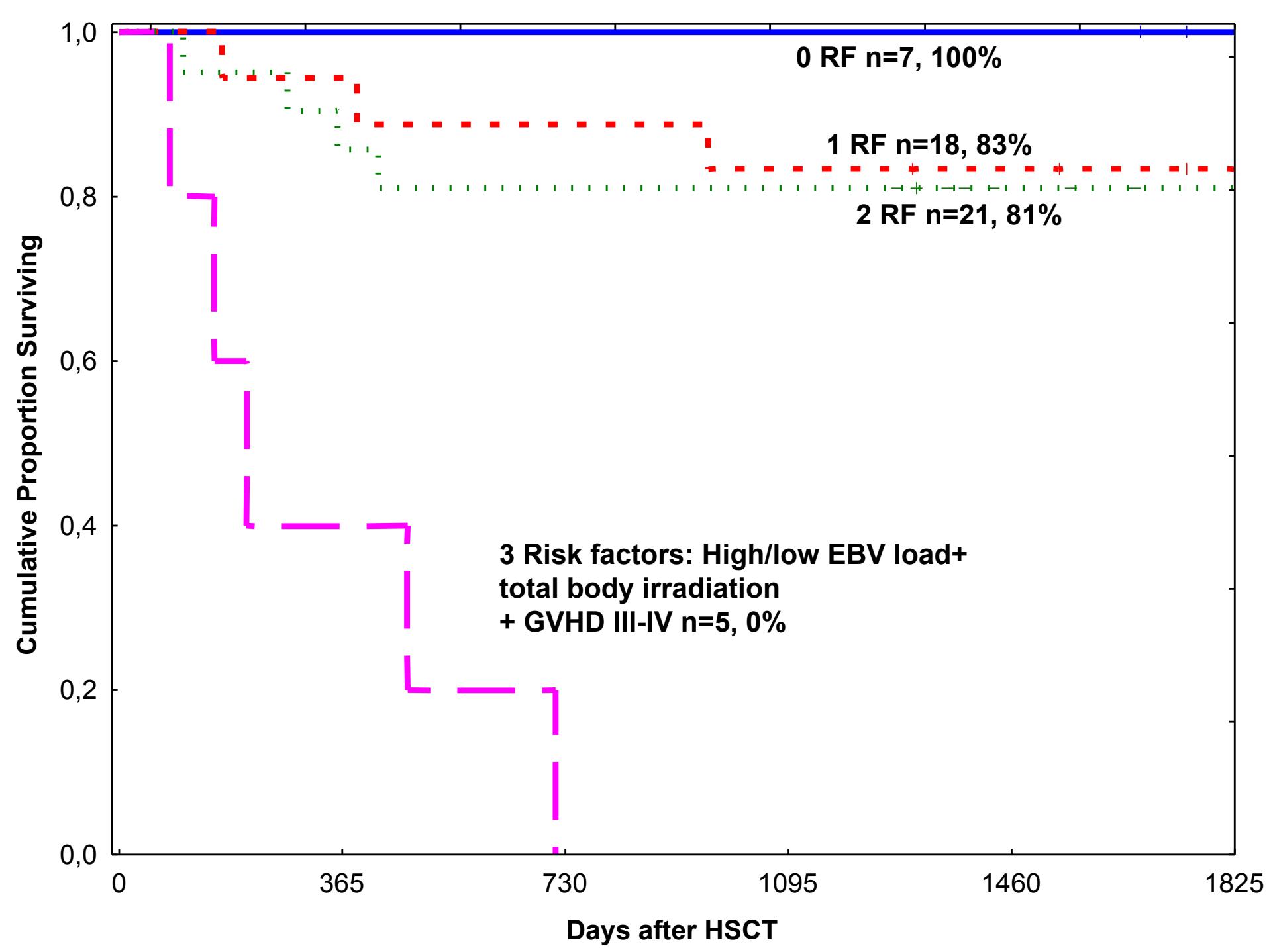


Low



High





## Concluding remarks



Target population for vaccination against EBV not identified; efficient vaccine exist (rec gp350); duration of protection not known

Normalization of EBV load in risk groups depends on functional immunereconstitution (cART; post-BMT)

Adoptive (EBV-specific) T-cell therapy prevents PTLD and cures some lymphomas (post HSCT)

New possibilities: check point inhibitors, CD4, interleukins (IL21?)

Extreme low or high EBV load in blood (B-cell bound) post-BMT/HSCT poor prognosis. With aGCVD & TBI predicts (100%) failure



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# Collaborators



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CTL work

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