

Details from a special case: laboratory acquired HIV-1 infection

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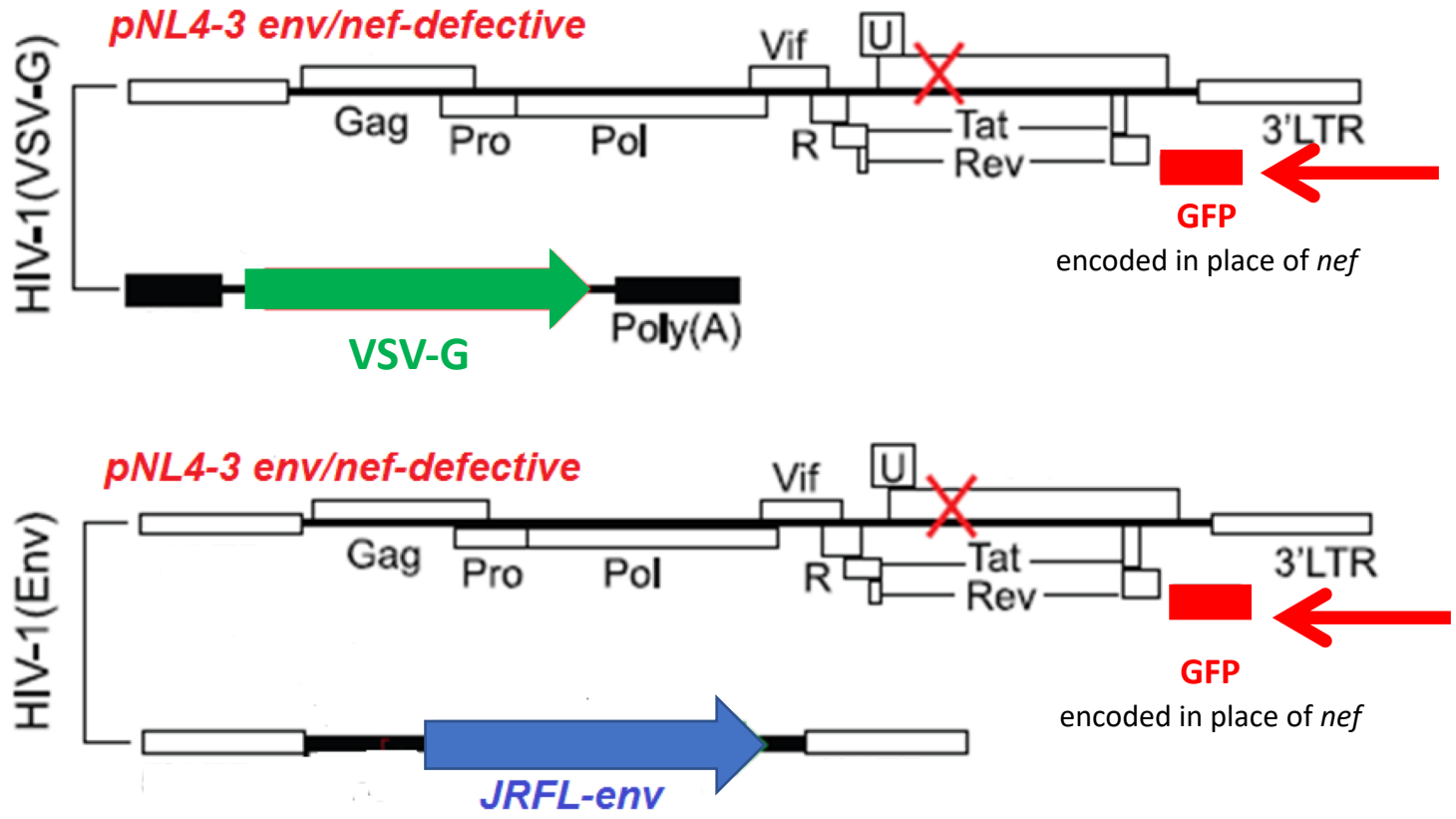
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Case report

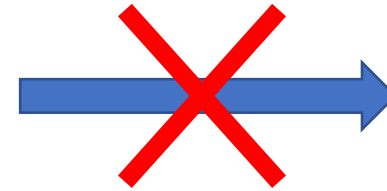
- A blood donor was found to be HIV-1 infected at routine screening
- Previous tests (the latest 7 months before) were negative
- No classical risk factors for HIV acquisition:
 - The unique partner was tested HIV-negative
 - No blood transfusions
 - No piercing or tattoos
 - No injecting drug use
 - No invasive medical procedures

However, during the 6 months preceding HIV diagnosis...

The subject worked in a research laboratory, with the exclusive task of producing pseudoviruses by co-transfecting 293T cells with HIV-1 **NL4-3 env/nef-defective**, green fluorescent protein–labeled vector, and **JRFL Env-** or **VSV-G** encoding plasmid in **BSL-2**



No access to BSL-3 was allowed or reported



No major or minor laboratory **accidents** (percutaneous injury, medium splash, breaking of gloves, eye or skin contact with cell-derived medium, etc) were recalled

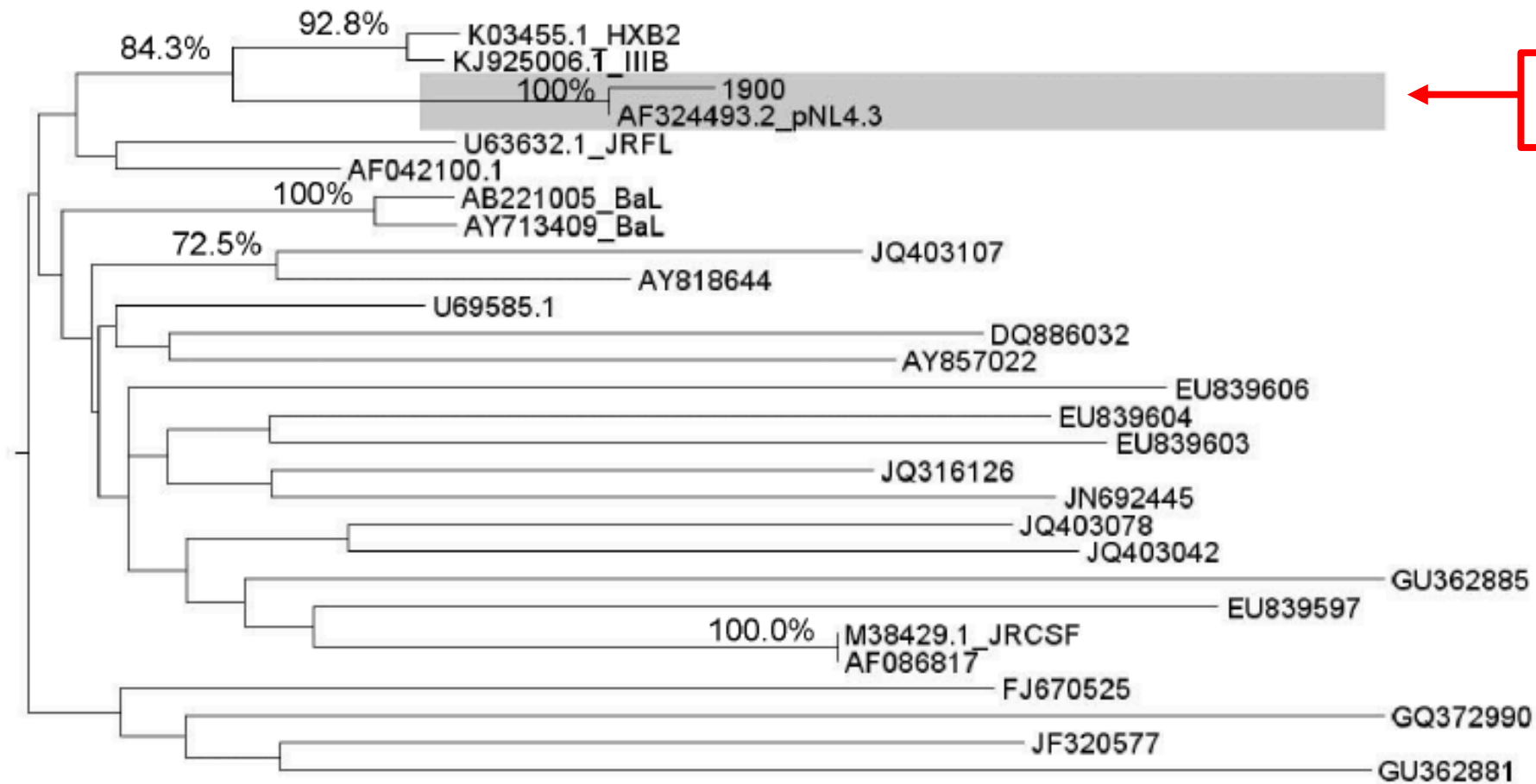
Methods

- Whole genome HIV-1 sequencing
- Phylogenetic analysis
- Viral isolation
- Phenotypic determination
- Neutralizing activity assessment
- Viral evolution analysis

Phylogenetic tree for gag-pol



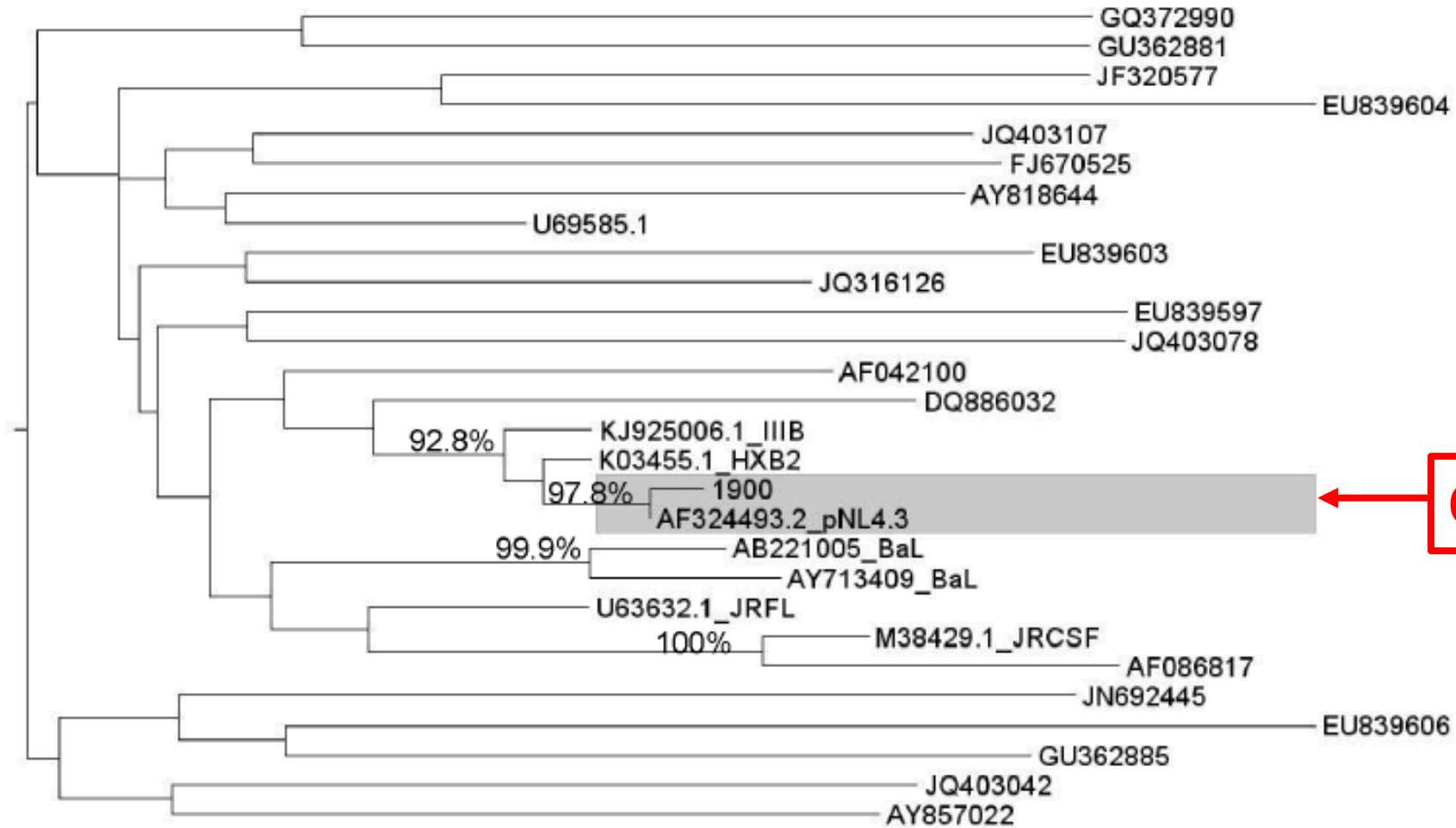
Phylogenetic tree for **Vif**, **Vpr**, **Vpu**, **Tat/Rev**



Cluster with **NL4.3**

0.005

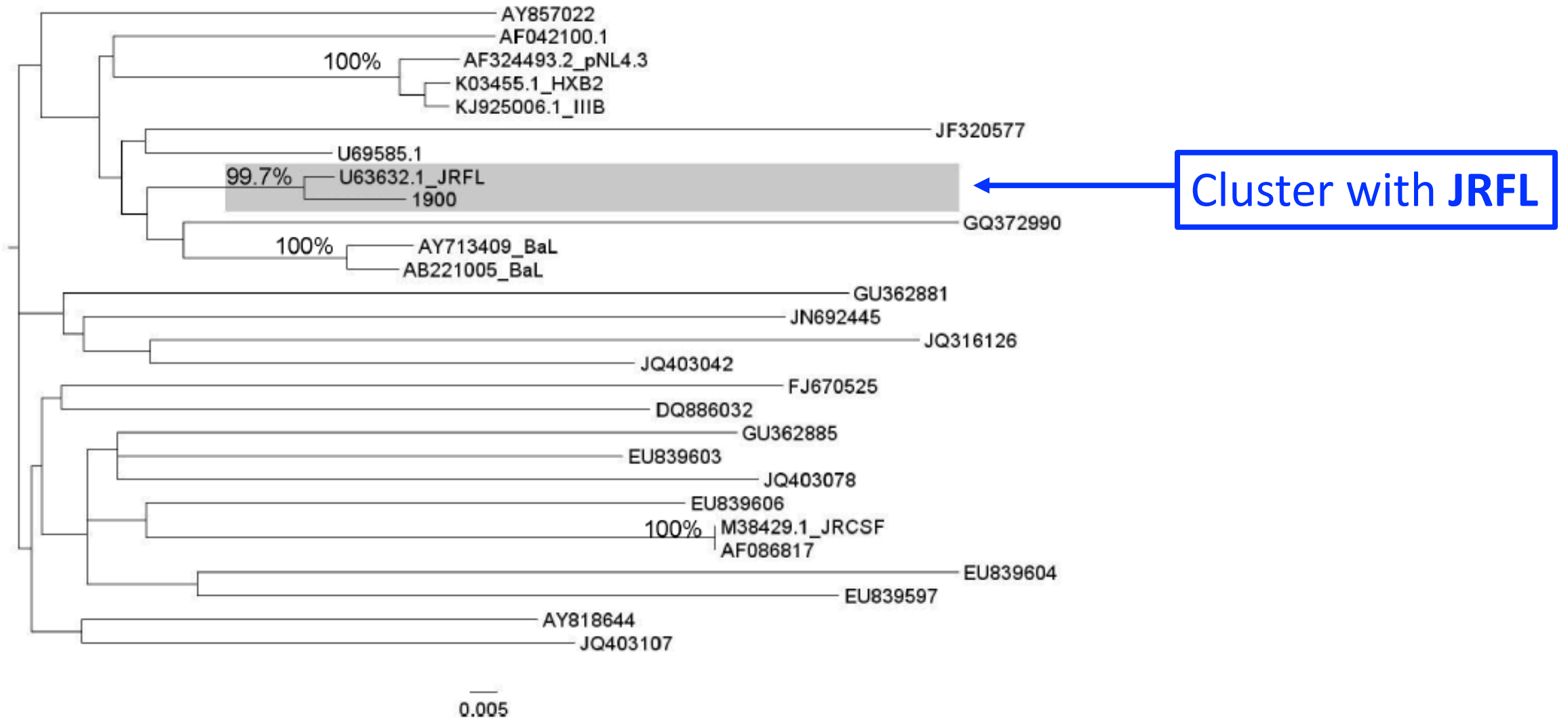
Phylogenetic tree for **nef**



Cluster with **NL4.3**

0.005

Phylogenetic tree for Env



High homology between patient's virus and constructs he/she had handled

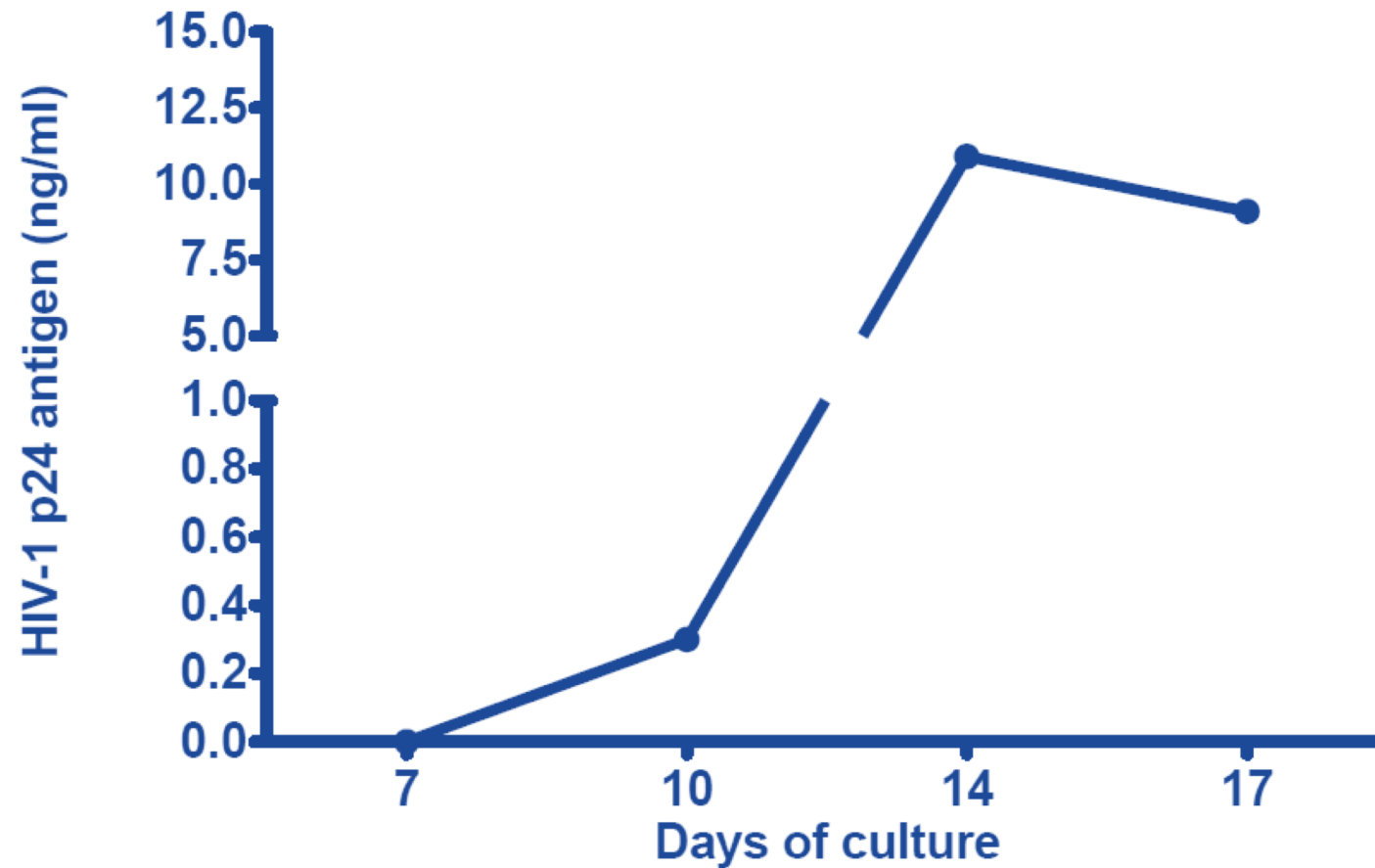
However, **nef** gene was present in patient's virus

Recombination between **NL4.3** env/nef defective vector and **JRFL** Env-encoding plasmid



Unaware use of a full-length **infective clone**

Viral isolation



Determination of viral tropism

Phenotyping

	HIV p24-Ag (ng/ml day 7)
U87.CD4.CCR5	34 (++) Syncytia)
U87.CD4.CXCR4	4.9 (no Syncytia)

Virus sensitivity to antibodies (autologous)

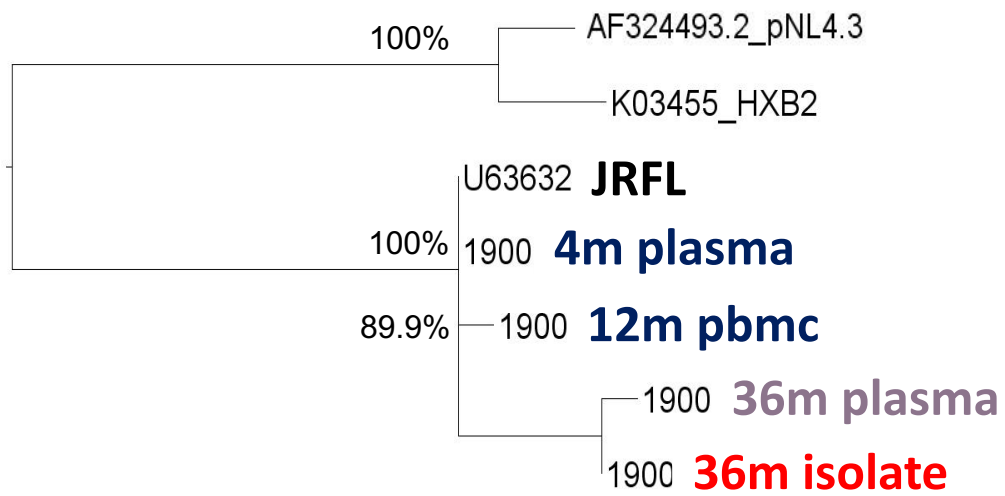
HIV-1 Viruses	Serum neutralization (IC50) in Tzm-bl assay (ug/ml)
Primary virus #1900	<20
JRFL (B)	495
<u>Tier 1</u>	
SF162 (B)	916
93MW965 (C)	1,397
GS015.EC12 (C)	304
TH023 (CRF01)	40
<u>Tier 2</u>	
DU151 (C)	<20
CM235 (CRF01)	<20
<u>Control</u>	
VSV-G	<20

Virus sensitivity to monoclonal antibodies

	Neutralization (IC50) with MoAbs in Tzm-bl assay (ug/ml)					
Virus	PG9	PG16	b12	2G12	4E10	2F5
Primary virus #1900	0.0049	0.0059	3.37	>5	>5	>5
JRFL (B)	>5	>5	0.014	1.21	3.77	0.78

Viral evolution

A.

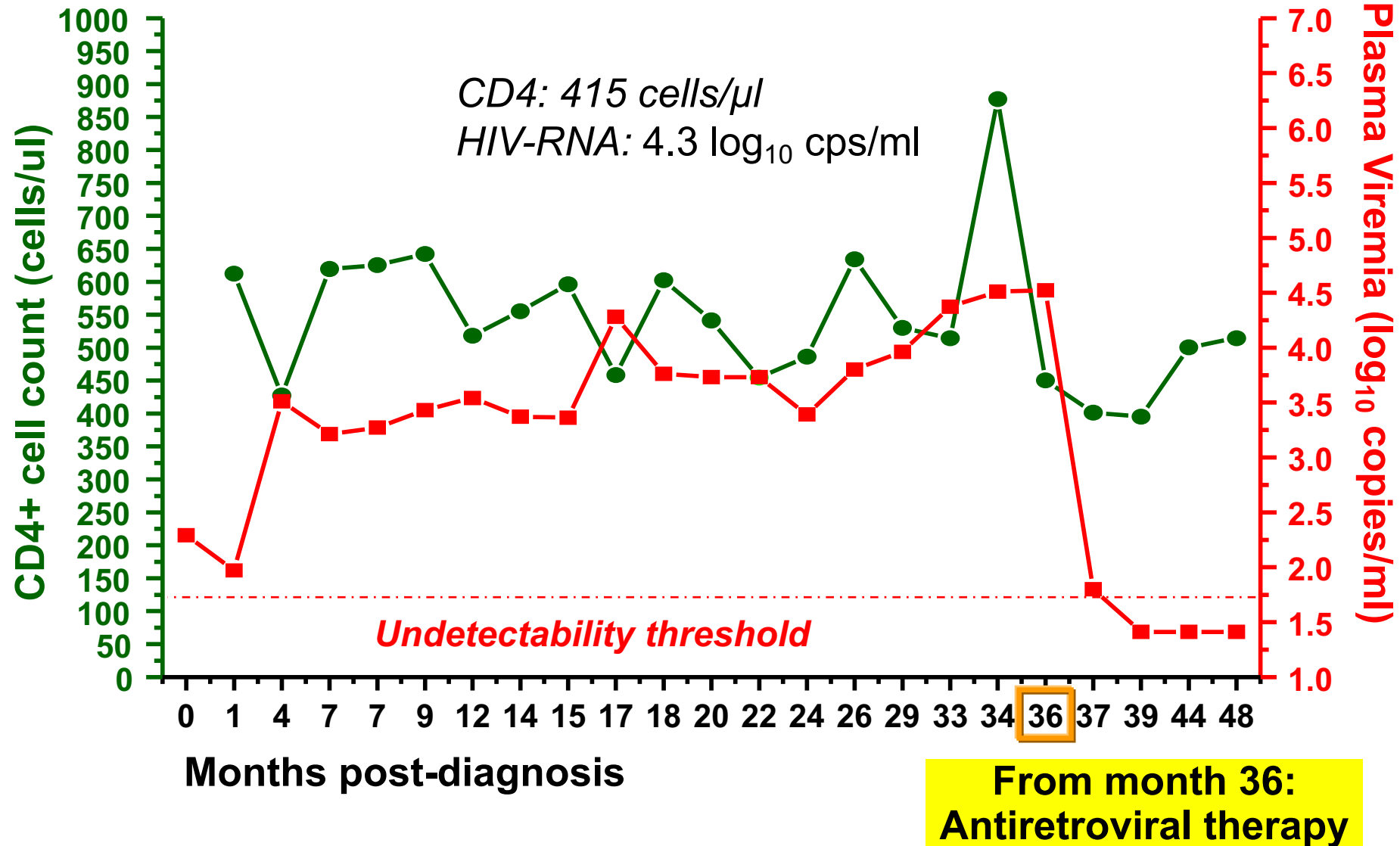


0.005

B.

	<i>Env</i>		
dN/dS	N-glycosylation sites	FPR (%)	
-	28 ^{TOT}	24.7	
0	<i>na</i>	24.7	
1	+1 (C2) 29 ^{TOT}	27.1	
2.2	+1 (C2) 29 ^{TOT}	17.0	
2	<i>na</i>	8.1	

Clinical follow-up



Finally: what's really happened?

We do not know yet. We can only make some hypotheses:

1. Labelling error



2. Unaware handling of infective clone in co-transfection experiments with VSV-G plasmid



3. Unaware contact with mucous membrane of a high-titer replicating HIV-1 culture

Lesson learned

- Trust the patient!
- Implement procedures to avoid contamination between BSL-2 and BSL-3
- Strict procedures of clone labelling
- ...

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For further details see publication: *Soria A, Alteri C, et al. Occupational HIV infection in a research laboratory with unknown mode of transmission: a case report. Clin Infect Dis 2017;64(6):810-3*