#### Evolutionary origin on HIV

The closest relatives of HIV-1 and HIV-2 are simian immunodeficiency viruses (SIV).

- Here is evidence for multiple transmissions from SIV into humans.
- HIV-1 is very closely related to SIV from chimpanzees.

#### **Evolution of virulence**

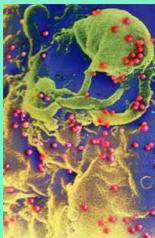
All SIVs appear to be apathogenic in their natural hosts.

SIV can be transferred to other species, where it induces AIDS.

**Short-sighted**' evolution of virulence.

#### HIV is a quasispecies

**Kiral replication is error prone**. **HIV reverse transcriptase and RNA** polymerase have error rates of about 1E-4. **#The virus population in any one patient is** extremely heterogeneous. **HIV can escape** from drug treatment. **HIV can escape from immune** responses.



#### **Evolution toward disease**

#Escape from immune responses
#Increasing viral diversity
#Faster replicating strains



#### Antigenic variation

virus mutant i

immune response against mutant i

$$\dot{v}_i = rv_i - px_iv_i$$
  

$$\dot{x}_i = cv_i - bx_i$$
  

$$i = 1, \dots, n$$

Each mutant goes to equilibrium:

$$v_i = \frac{br}{cp} \qquad x_i = \frac{r}{p}$$

Add new mutants over time.

#### Antigenic variation

Total virus load is proportional to antigenic diversity.

$$v \coloneqq \sum_{i} v_{i} = n \frac{br}{cp}$$

#### Antigenic variation

virus mutant i

specific immune response

cross reactive immune response

$$\dot{v}_i = v_i(r - px_i - qz)$$
  
$$\dot{x}_i = cv_i - bx_i \qquad i = 1,...,n$$
  
$$\dot{z} = kv - bz$$

Virus load:

$$v = \frac{brn}{cp + kqn}$$

#### Antigenic variation of HIV

virus mutant i

specific immune response

cross reactive immune response

$$\dot{v}_i = v_i(r - px_i - qz)$$
  

$$\dot{x}_i = cv_i - bx_i - uvx_i \quad i = 1,...,n$$
  

$$\dot{z} = kv - bz - uvz$$
  
Virus load: hrm

$$v = \frac{brn}{cp - (ru - kq)n}$$

#### Antigenic variation of HIV

Virus load:  

$$v = \frac{brn}{cp - (ru - kq)n}$$

#### **Diversity threshold:**

$$n_c = \frac{cp}{ru - kq}$$

# The 'diversity threshold' model has 3 possible outcomes

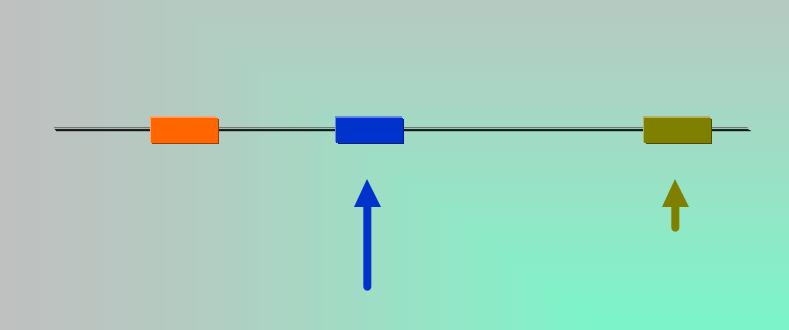
1. Disease after long asymptomatic period.

kq < ru < kq + cp



- 2. Indefinite virus control. ru < kq
- 3. Immediate disease. kq + cp < ru

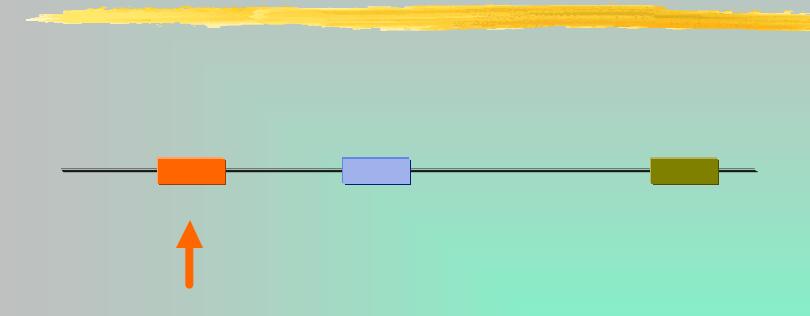
## Immune responses to multiple epitopes



#### Immunodominance

breadth of the response is related to immune memory

### Immune responses to multiple epitopes



### Antigenic variation can lead to shifting immunodominance

### HIV disease progression according to this model

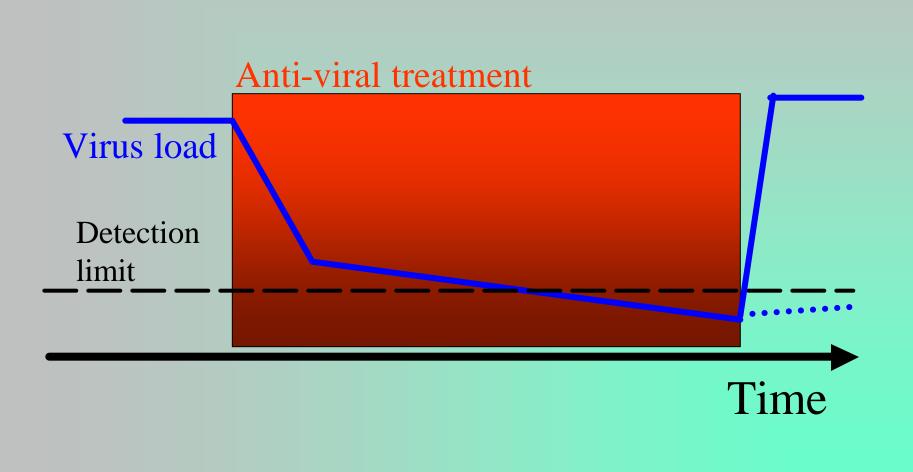
 There is a highly dynamic balance between the virus and the immune system with rapid virus turnover.
 The evolutionary adaptation of the v

The evolutionary adaptation of the virus in individual patients is the mechanism of disease progression.

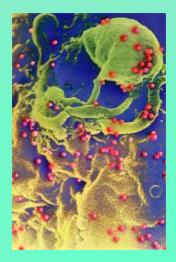
### Three possible mechanisms of HIV disease progression

#Evolution of the virus
#Slow break-down of the immune system
#Accumulation of opportunistic infections

## The virus will return if therapy is withdrawn



Is it possible to treat and help the patient's immune system to gain control of the virus?

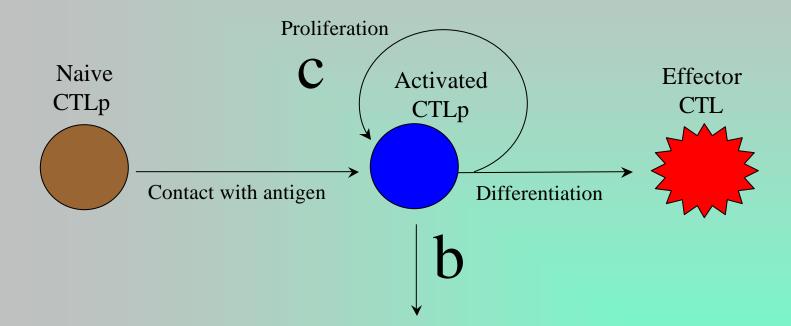


#### A new theory of CTL memory

The primary role of CTL memory is to eliminate virus infections or to reduce virus load to low levels.

**Dominik Wodarz** 

#### **CTL** dynamics



**CTL memory** is characterized by highly responsive and long-lived CTL precursors (high c and low b). **CTL memory** requires CD4 cell help

#### The basic model with CTL

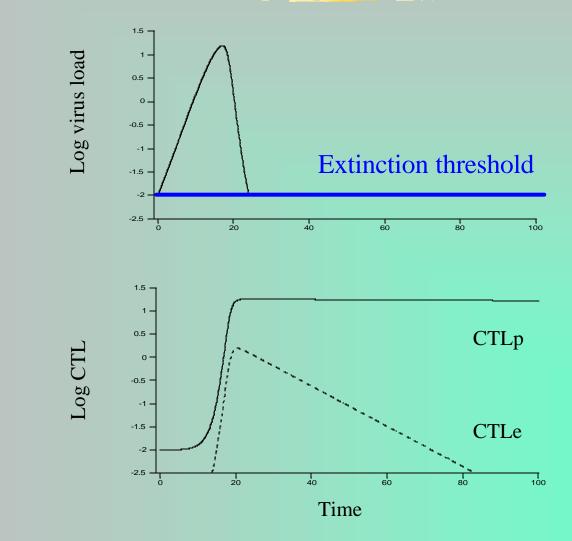
$$x = \mathbf{l} - dx - \mathbf{b} xy$$

$$y = \mathbf{b} xy - ay - pyz$$

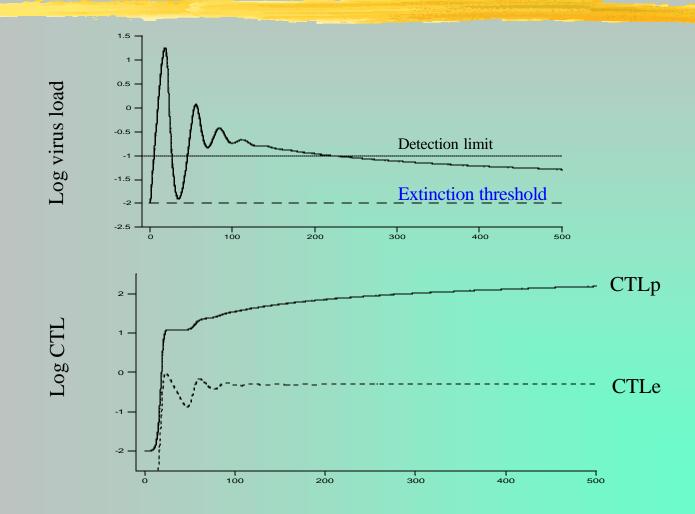
$$w = cyw (1 - q) - bw$$

$$z = cqyw - hz$$

### 2 possible outcomes: 1. Virus elimination



#### 2. Persistent infection



Time

### HIV specific model

$$\dot{x} = \mathbf{l} - dx - \mathbf{b} xy$$
$$\dot{y} = \mathbf{b} xy - ay - pyz$$
$$\dot{w} = cxyw - cqyw - bw$$
$$\dot{z} = cqyw - hz$$

#### HIV

#### HIV kills CD4 cells which are needed for CTL memory.

Failure to establish a CTL memory response leads to persistent infection, high virus load and rapid disease progression

A good CTL memory response leads to virus elimination (rare ?) or at least low virus load and slow disease progression

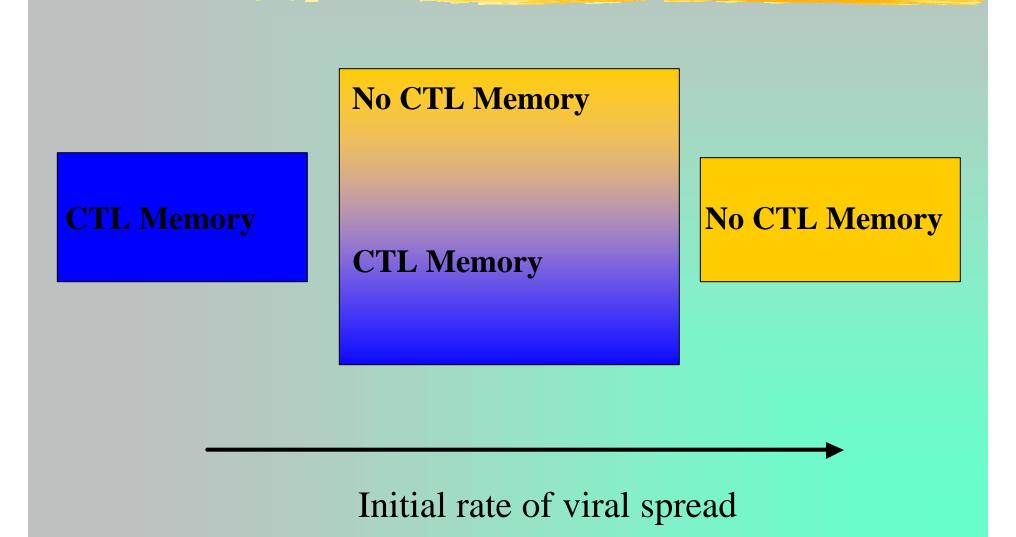
#### HIV: rate of disease progression

#### **Fast progressors: high virus load**

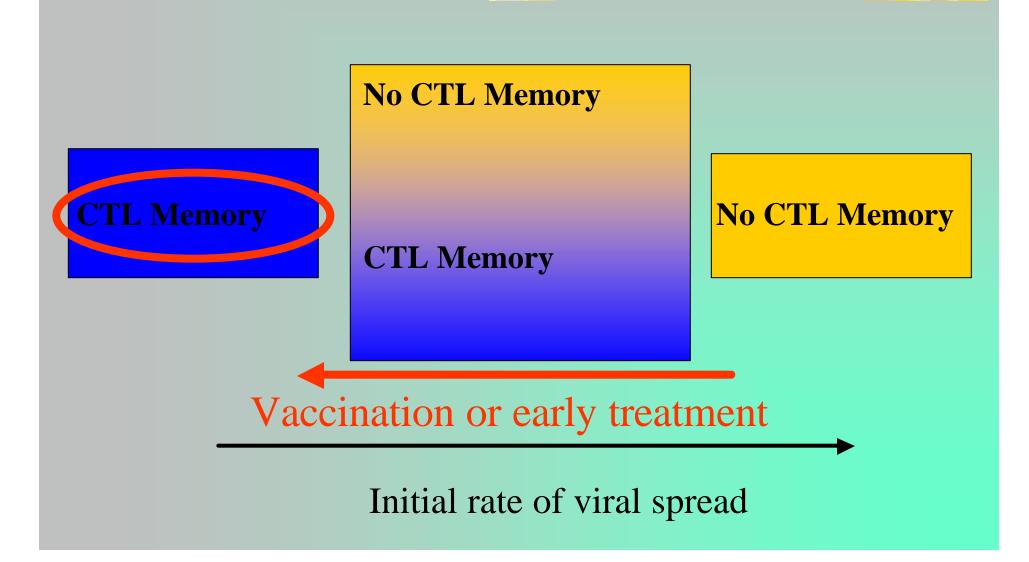
#### CTL memory makes the difference.

**Slow progressors: low virus load** 

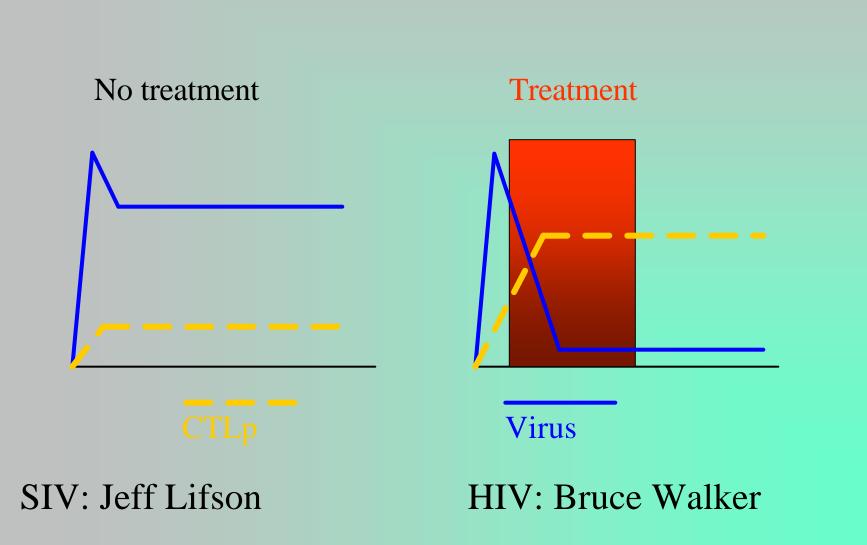
# HIV replication and establishment of memory

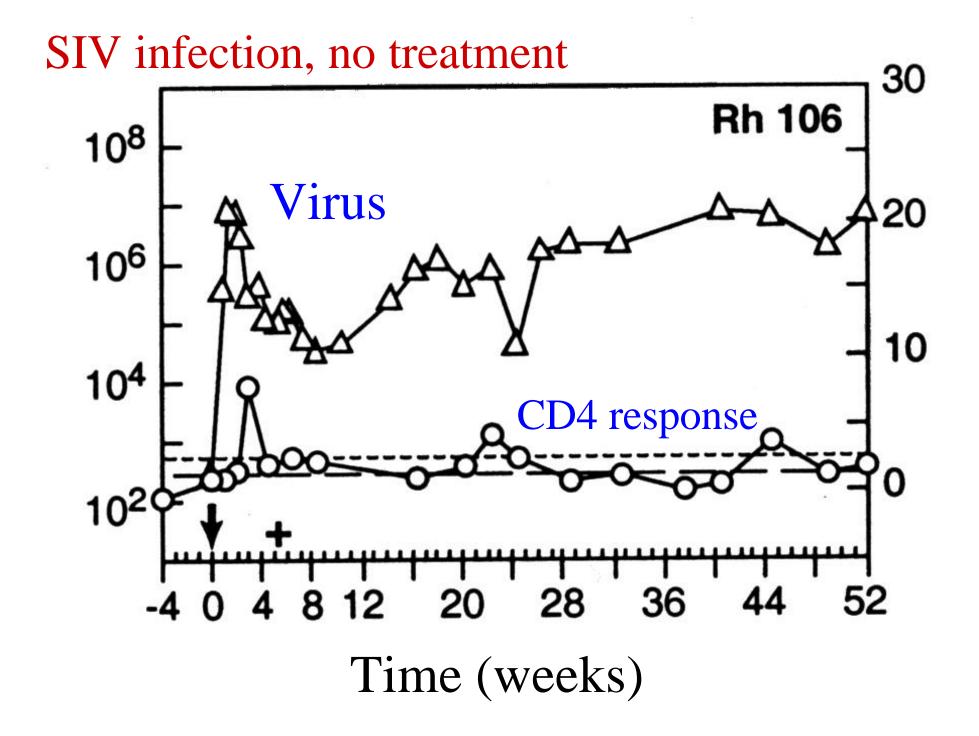


# HIV replication and establishment of memory

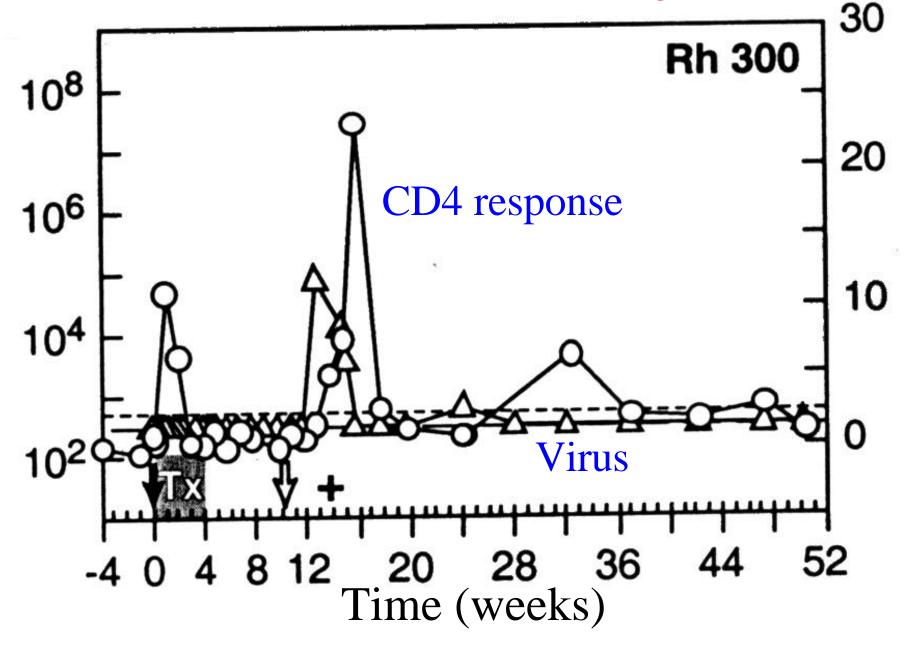


## Treatment during primary infection

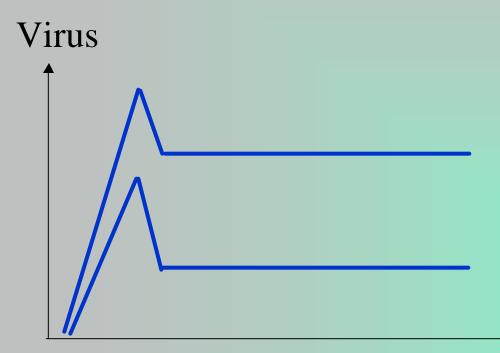




#### 4 weeks of treatment ; re-challenge



### SIV primary infection without treatment

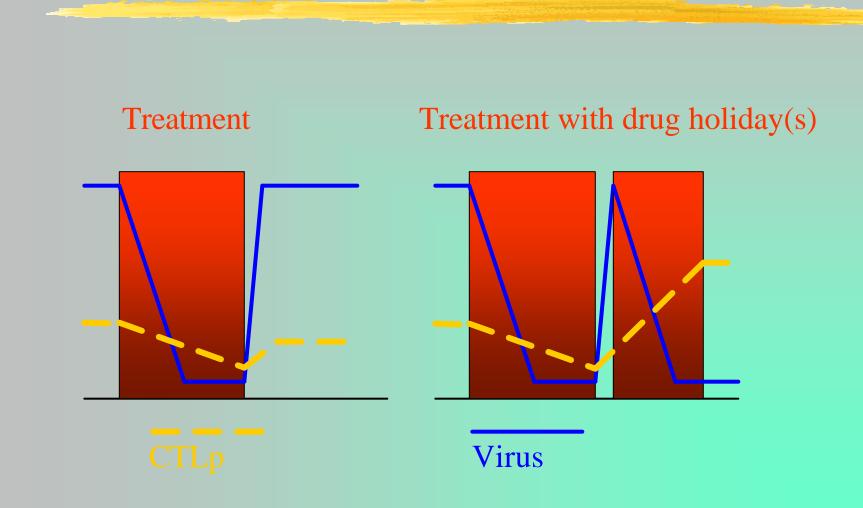


Virus load in the first week of infection is correlated with set-point is correlated with survival.

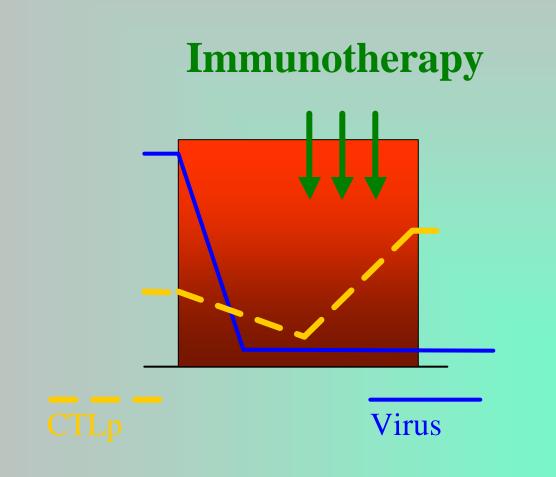
Time

Jeff Lifson: 12 monkeys, 12 authors

### Treatment during chronic HIV infection



# Anti-viral treatment and immunotherapy



### A new approach for HIV therapy

**For primary infection:** Use vaccination and early treatment to reduce the initial viral growth rate and bring patients into a state of long term non-progression.

For chronic infection: Use treatment and immunotherapy to switch patients into a state of long term non-progression.