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Evolution before our eyes

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Synopsis

Viruses can evolve fast and sometimes adapt quickly to a new host species. For example, an influenza virus that normally infects birds can become adapted to humans. The tobacco etch virus normally infects tobacco plants. Professor Santiago Elena from Valencia wants to find out what it takes to make the tobacco virus capable of infecting another plant: Arabidopsis. The movie shows how Santiago Elena does the evolutionary experiment and we see that after 30 rounds of experimental evolution the virus is indeed adapted to the new host plant! After the experiment, Elena looks at the genetic code of the adapted virus and finds that there are just three differences between the genetic code of the normal (tobacco loving) virus and the virus that is now adapted to rabidoptis. » Other Bookm

Winner at "Vedere la Scienza Festival 2010" (category: Educational videos for the web). Finalist at the "European Science TV and New Media Festival 2010".

Credits

Directors : Robert Sigl, Bemhard Rübe & Pleuni Pennings - Script & editing : Brant Backlund - Camera operator : Bemhard Rübe - Graphics : Simon Baker, James Doyle & Jacob Barrow - Equipment Manager : Robert Sigl -Narration : Francine Oliver - Narration Producer : Chris Kugelman - Music : Chris Tegg - Production Manager : Marion Hartl - Production Assistant : Kaitlin Hickey - Producer : Brant Backlund - Executive Producer : Pleuni Pennings - Production : LMU Munich (2009)

Scientific expert

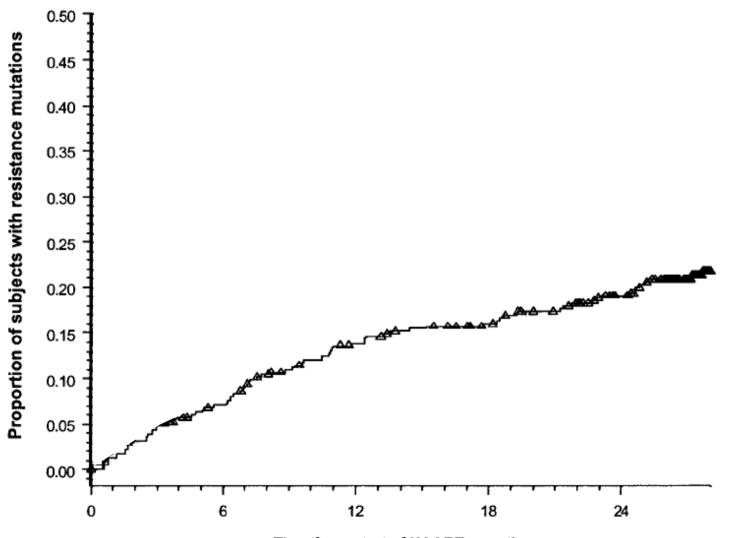
Santiago Elena

Instituto de Biologia Molecular y Celular de Plantas, Evolutionary Systems Virology Group, Spain Evolutionary Virology

•To design working antiviral strategies, we need to understand how new viruses appear and what principles govern their diversification and adaptation to hosts. Also, viruses allow us to witness evolution in real time and to test fundamental principles of the evolutionary theory.• What determines the rate of evolution of resistance of HIV?

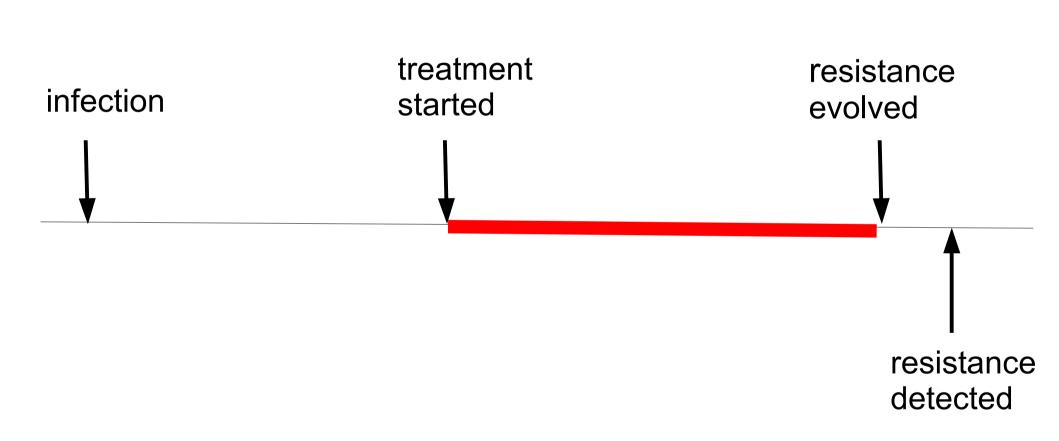
Pleuni Pennings March 2011 KITP, Santa Barbara

Distribution of waiting times

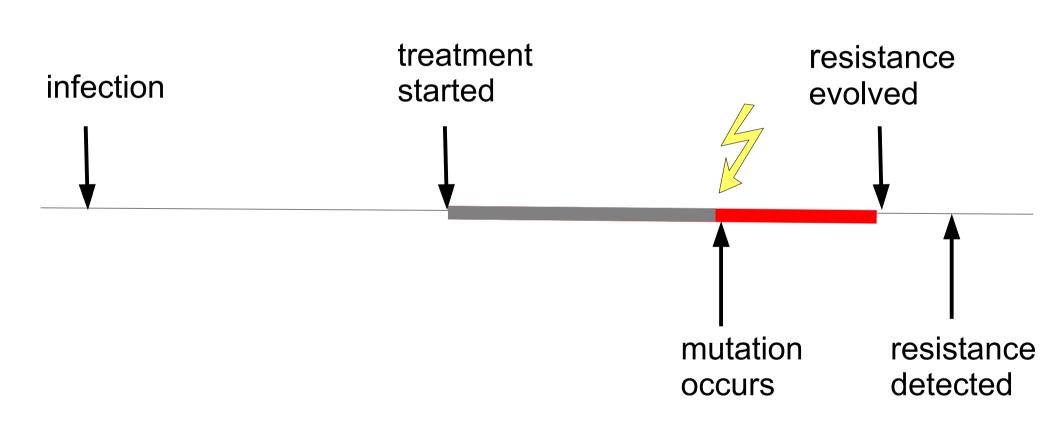


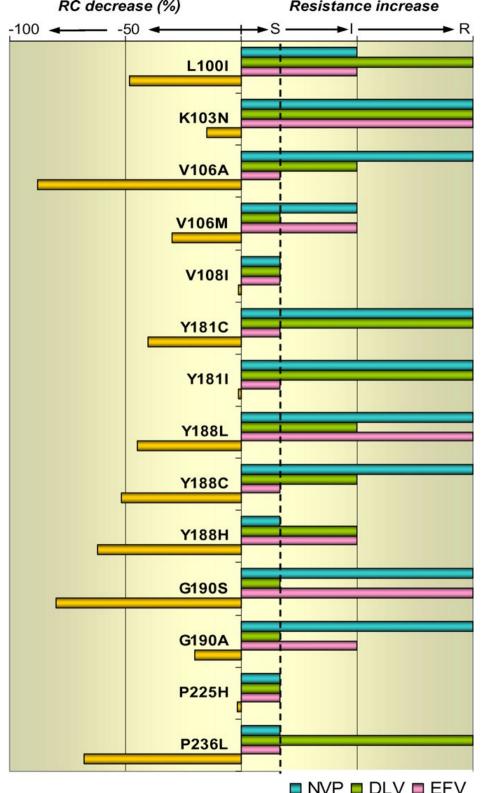
Time from start of HAART, months

Timeline



Timeline

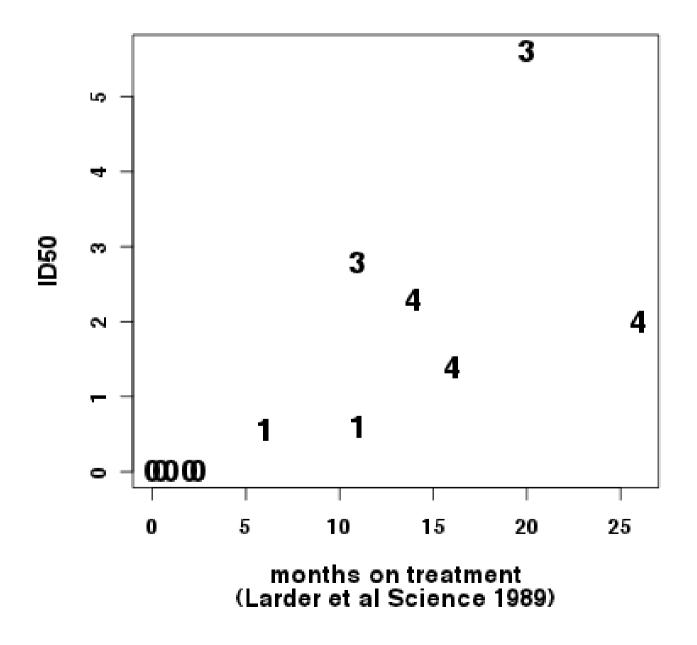




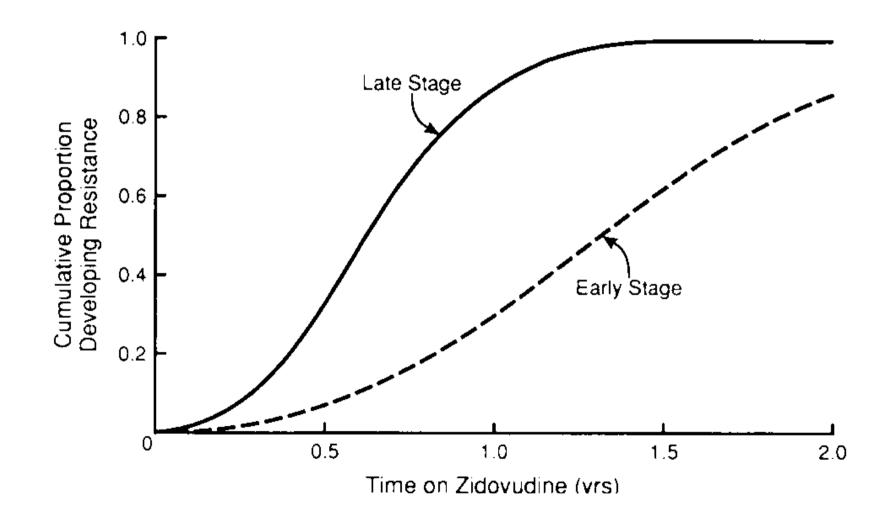
Mutations are well characterized

Martinez-Picado 2008 Virus Research

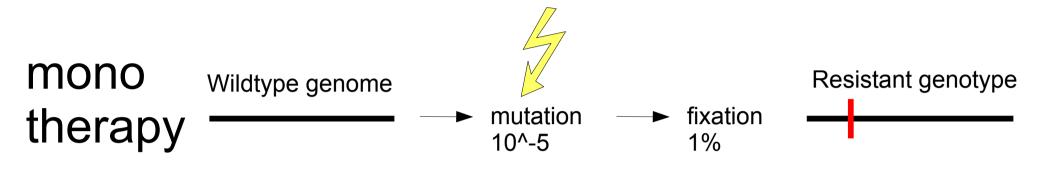
Resistance against AZT in 12 patients

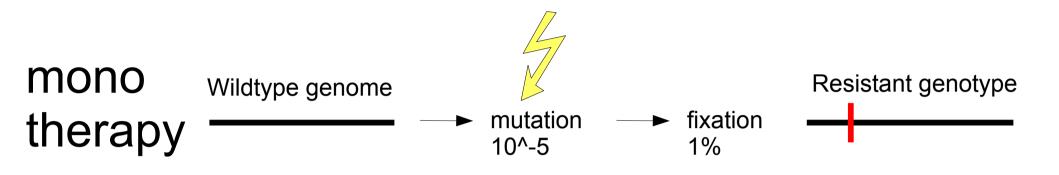


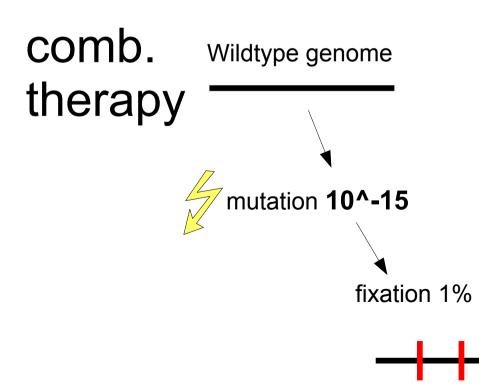
Evolution is mutation limited



Richman et al 1990 JAIDS

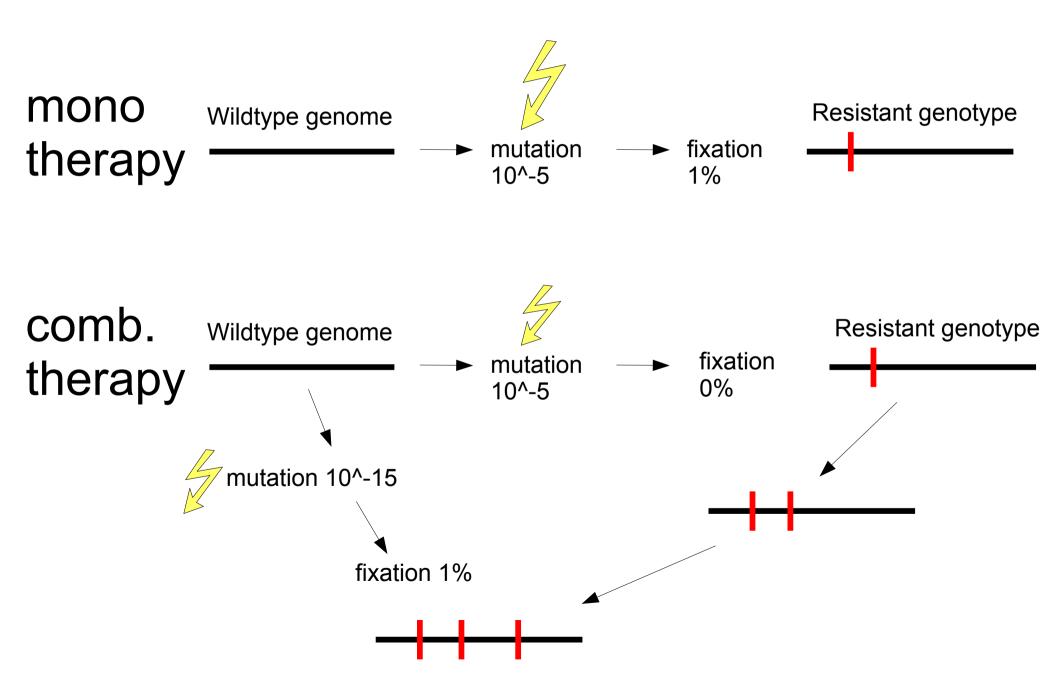


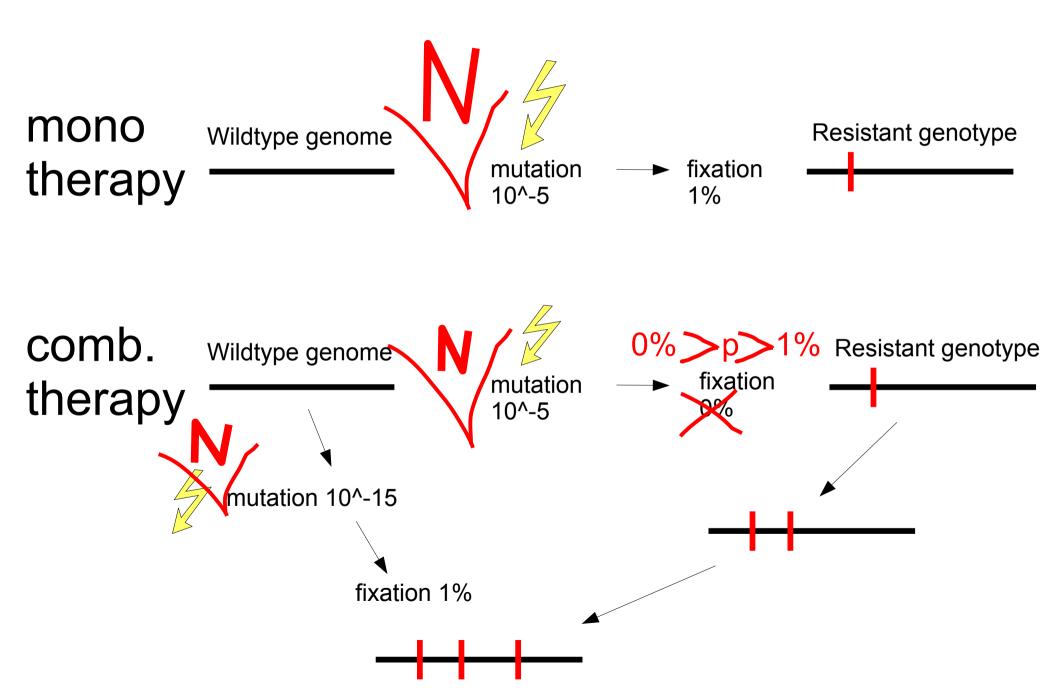


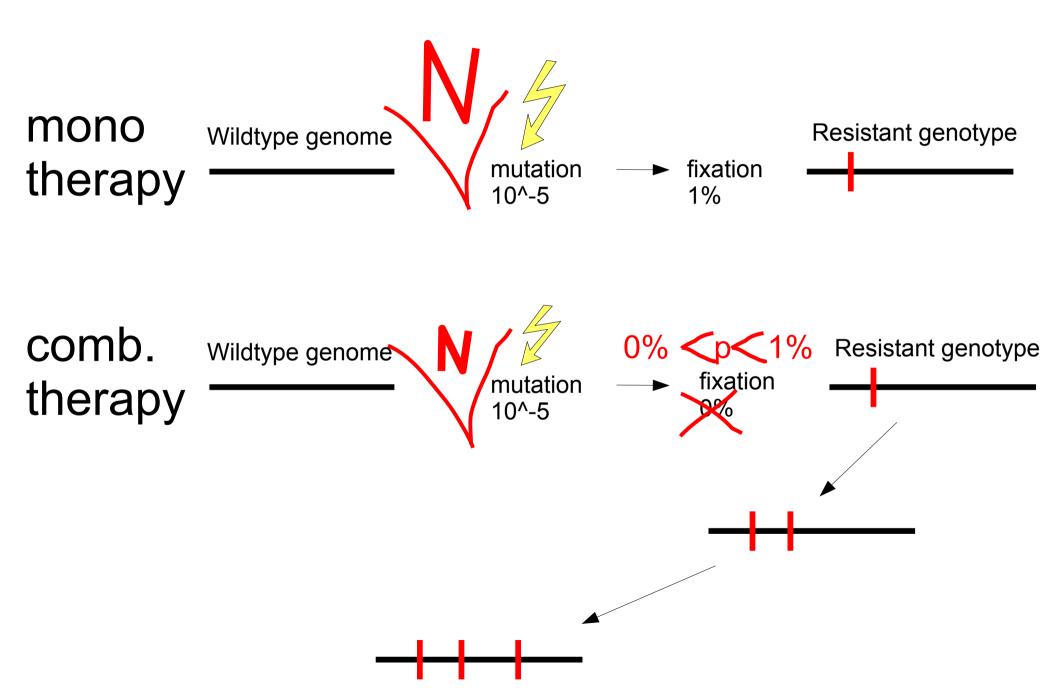


Common belief: "Combination therapy works

because it is unlikely that three mutations happen in one viral genome"

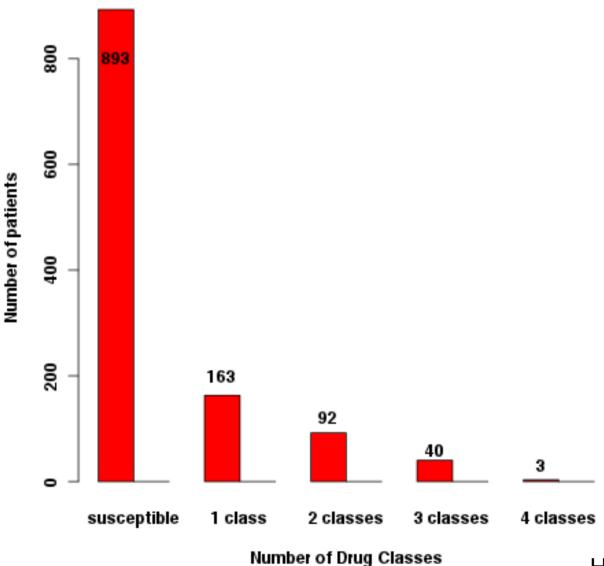






Resistance evolves one drug class at a time

Resistance after 2.5 years of HAART



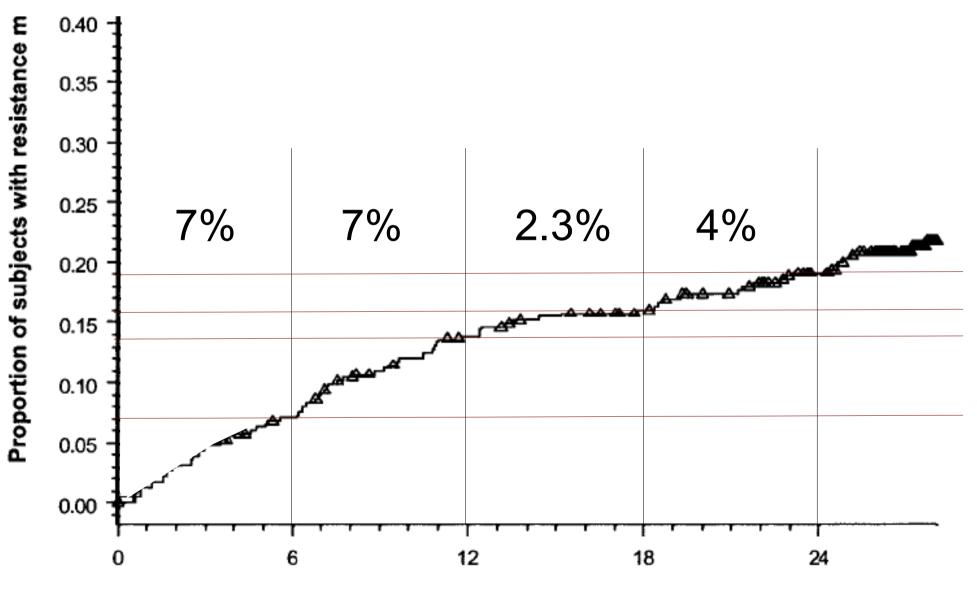
Harrigan et al 2005 JID

Summary of introduction

- 1. Waiting time distribution can be observed
- 2. Evolution is limited by mutations
- 3. First mutation most important
- 4. HIV evolves resistance one drug-class at a time

It is truly a simple problem!

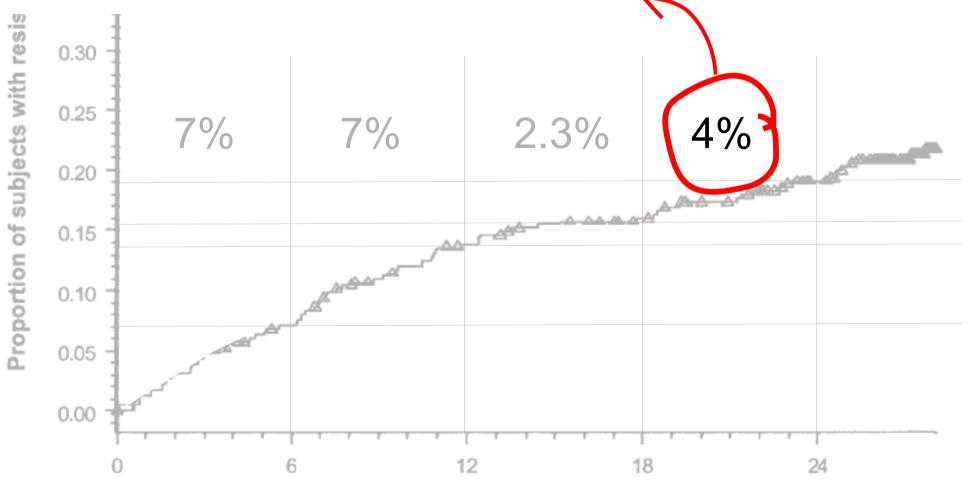
Example calculation



Time from start of HAART, months

Example calculation

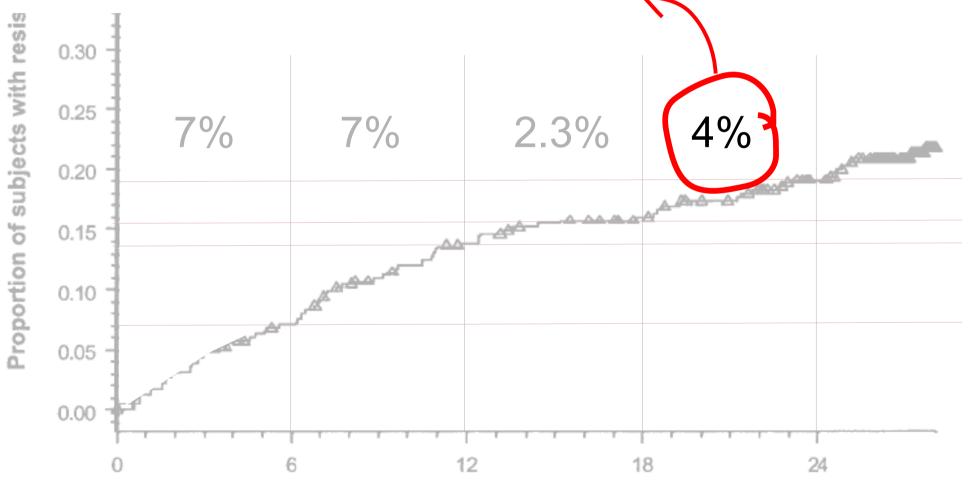
100 generations x 10⁻⁵mut/gen x 40 possible mutations x 100 pop size x 1% fixation probability



Time from start of HAART, months

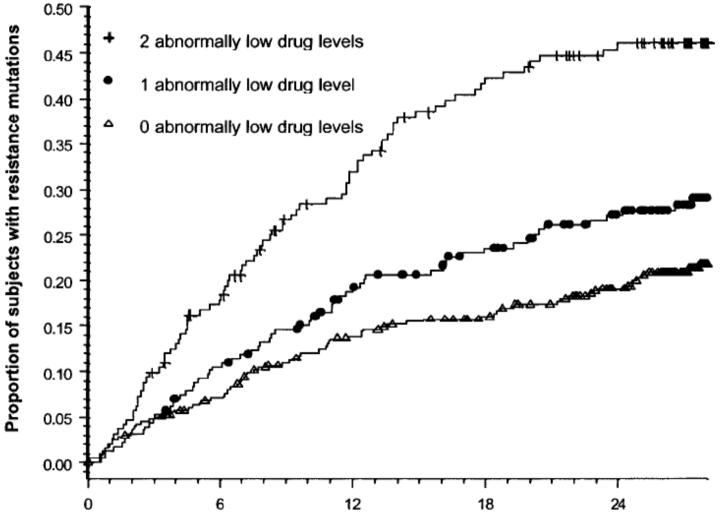
Example calculation

100 generations x 10⁻⁵mut/gen x 40 possible mutations x 100 pop size x 1% fixation probability



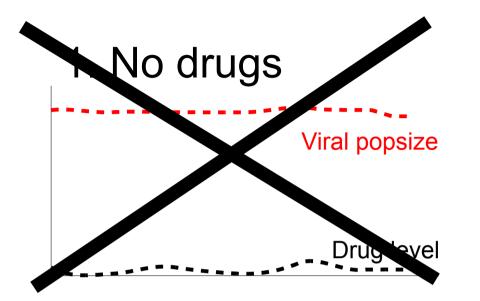
Time from start of HAART, months

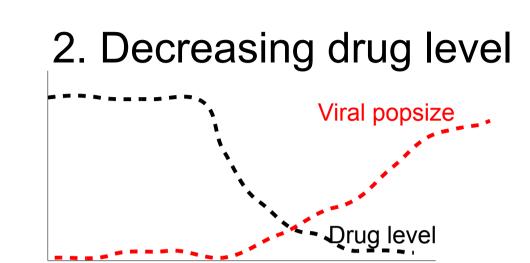
Non-adherence increases rate of evolution

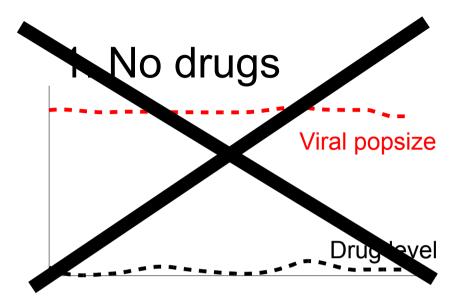


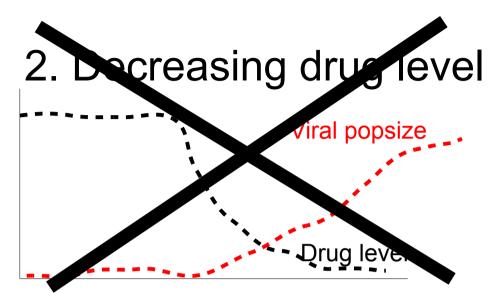
Time from start of HAART, months

1. No drugs Viral popsize Drug level

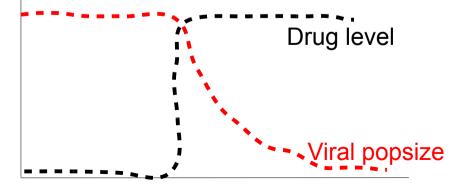


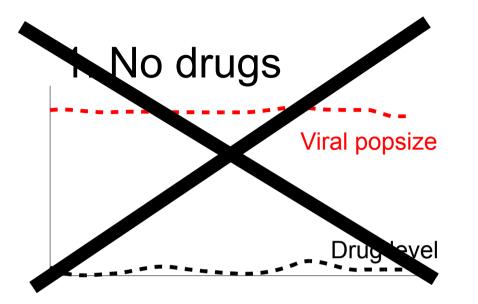


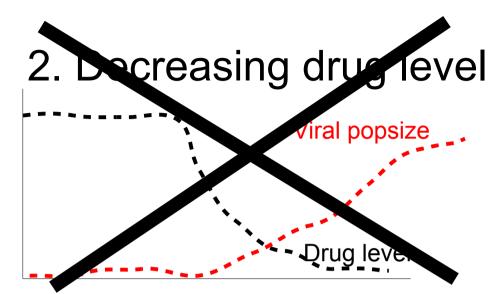




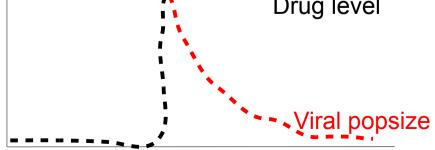
3. Increasing drug level

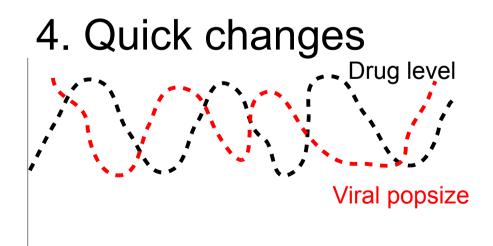


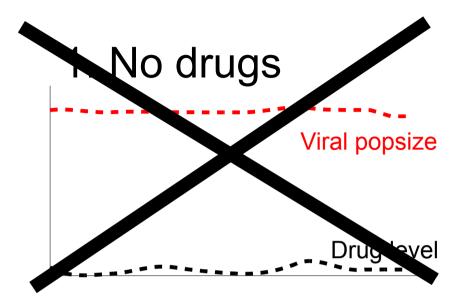


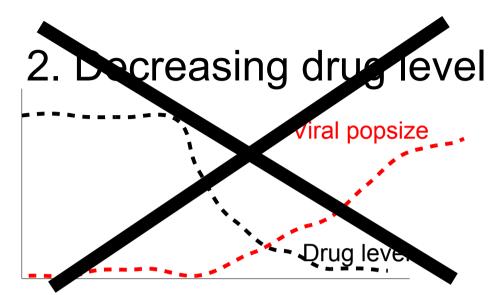


3. Increasing drug level

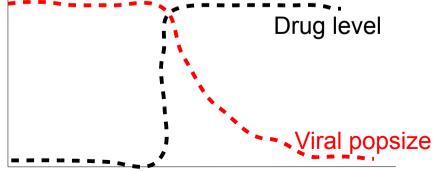


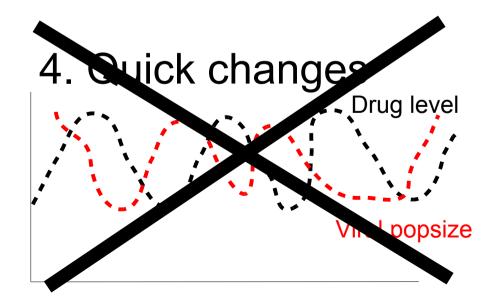


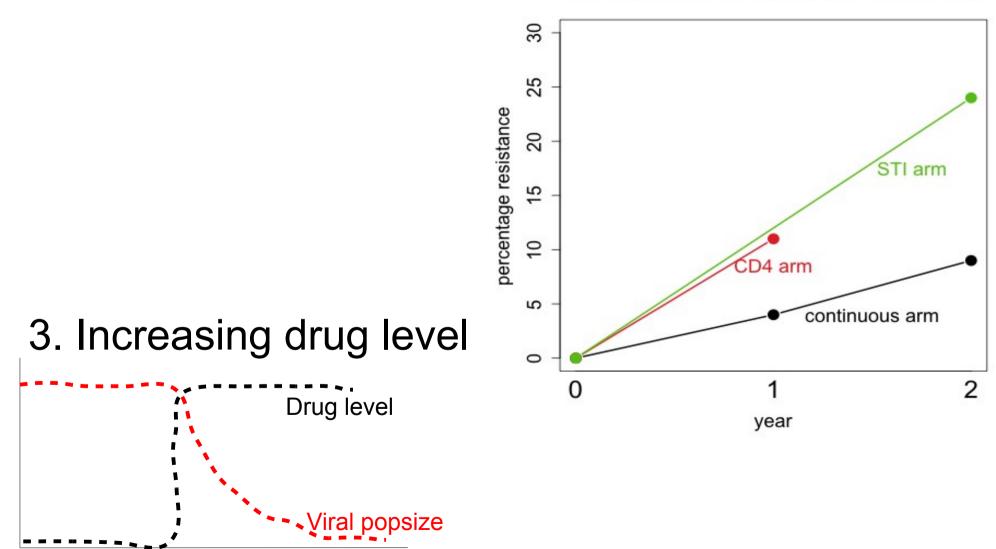




3. Increasing drug level



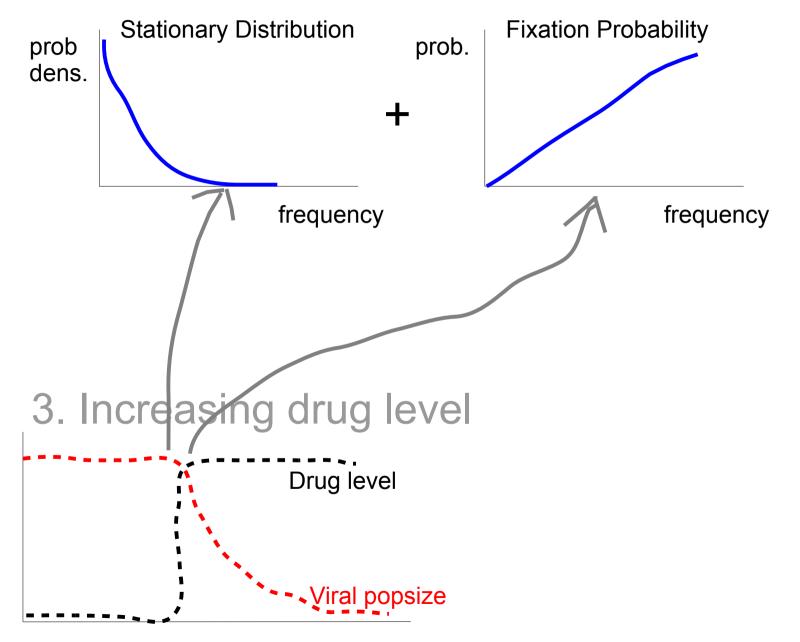




Rate of evolution of resistance Trivacan trial

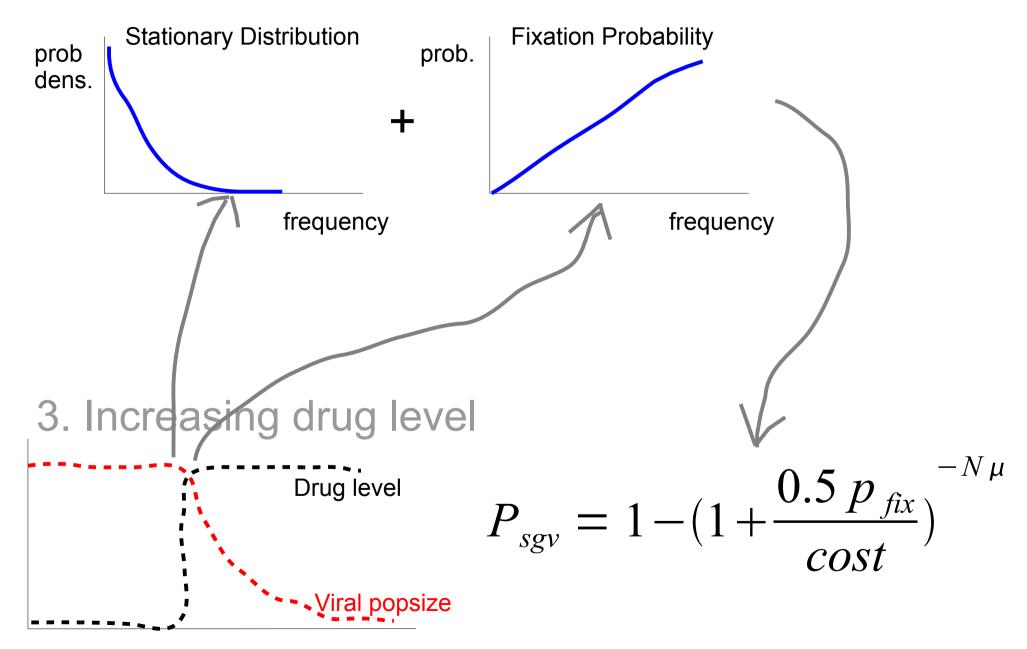
Danel et al 2006, 2009

Adaptation from standing genetic variation



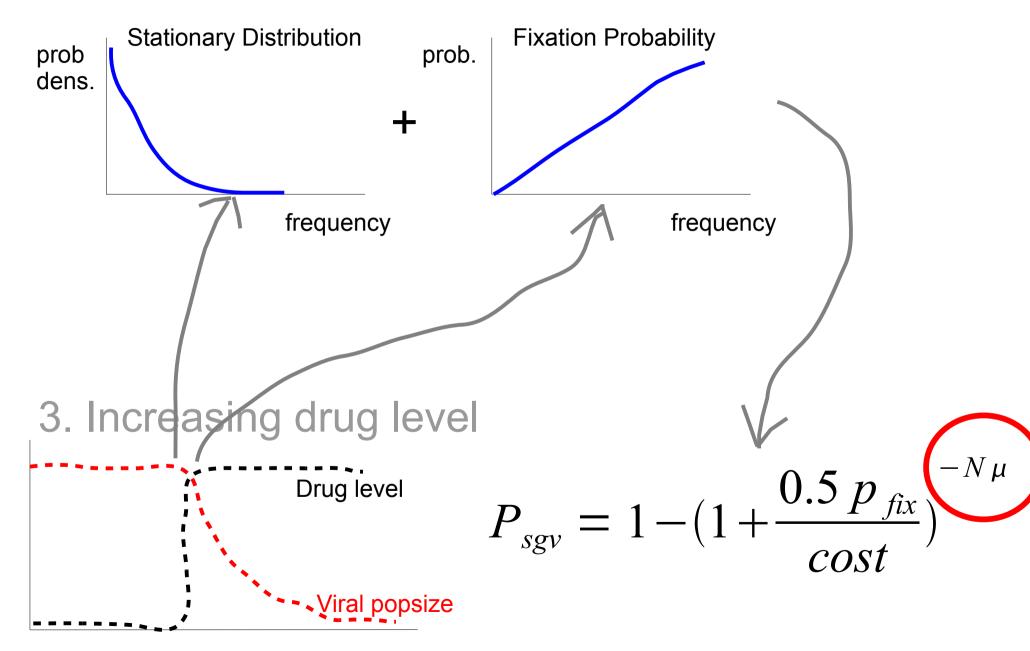
Hermisson & Pennings, Genetics 2005

Adaptation from standing genetic variation



Hermisson & Pennings, Genetics 2005

Adaptation from standing genetic variation



Hermisson & Pennings, Genetics 2005

Data

7 trials of "Structured Treatment Interruptions"

Example: Swiss-Spanish Intermittent Treatment Trial

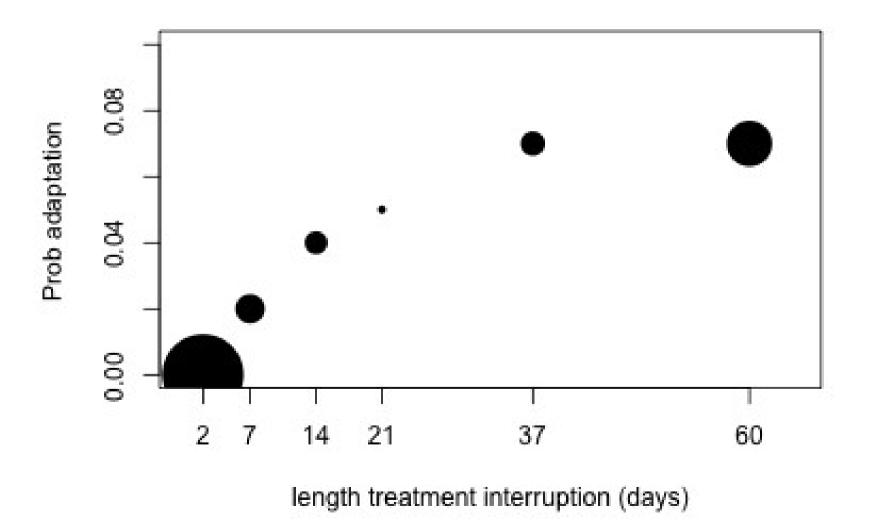
- 87 patients (-7)
- 4 interruptions of 14 days
- 11 patients had resistance

$$\frac{69}{80} = (1-x)^4$$

$$x = 0.04$$

Data

Prob of evolution resistance per interruption



$$P_{sgv} = 1 - (1 + \frac{0.5 p_{fix}}{cost})^{-N\mu}$$

$$7\% = 1 - \left(1 + \frac{0.5 \ 1\%}{5\%}\right)^{(-N4010^{-5})}$$

10^-5mut/gen x 40 possible mutations

1% fixation probability

5% cost

N = 1875

First year bonus?

	YEAR 1	YEAR 2	YEAR 3	P_SGV
Margot 2006	10.5	3.4	4.0	6.8
Harrigan 2005	14	6		8

Conclusions

- Very simple models work well - predictions fit with data.

- Population size can explain why treatment interruptions lead to faster evolution.

- HIV uses both standing variation and new mutations to adapt.

- Idea: adaptation from standing variation can be avoided with different treatment protocol.

- Unclear: how come Ne is so low?
- How can single-drug resistance evolve?

Thanks



John Wakeley (Harvard)

Joachim Hermisson (Vienna)





Dan Kuritzkes (Harvard)

Jonathan Li (Harvard)

Many people here and elsewhere for discussions