

Evolution of viruses antiviral defense

K. Moelling, University of Zurich, Zurich, CH and Institute of Advanced Studies, Berlin, Dtl

EVOLUTION OF VIRUSES AND ANTIVIRAL DEFENCE

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Organization of presently known viruses according to a hypothetical appearance during evolution (RNA to a DNA world).

Contemporary viruses can be organized in an evolutionary tree ranging from the RNA world to the DNA world, from ribozymes, via viroids, DNA-ribozymes, Influenza, retro- and para-retroviruses to DNA viruses – supporting a „virus-first“ hypothesis. Retroviruses have shaped or may have even built the human genome, where up to 50% are retrovirus-related sequences to which increasing and decreasing complexities contributed. Rudimentary reverse transcription from RNA to DNA is still ongoing today in telomeres during embryogenesis and cancer(1). Sequence analysis of the human genome witnesses our past, indicating how long HIV-like viruses, reverse transcriptase and RNases H have been around. Endogenization of retroviruses is actively ongoing in animal models and may allow a prediction on the future of HIV in people. Evolution of HIV takes place during antiretroviral therapies. An HIV suicide approach circumvents mutagenesis and escape mutants(2,3). Co-evolution or crossing arms, also known from phage and bacteria, can be deduced from structural and functional similarities of retroviral replication and the siRNA-mediated antiviral defence machineries (4). An evolutionary relationship between siRNA and interferon can be constructed by comparing their pathways. siRNA involving dicer as well as interferon are active antiviral defence mechanisms in mammalian cells, tested by dicer and interferon knock-down analyses (5;6). The systems differ in strength and sequence specificities.

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From the RNA world to the DNA world



Time, complexity, stability

Small interfering RNA is related to interferon as antiviral defence.

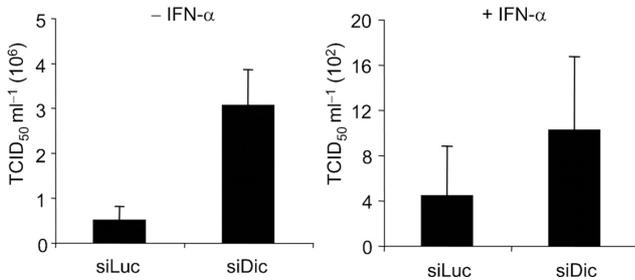
Relationship between Retroviruses and RISC

Retrovirus	siRNA
Reverse Transcriptase (RT)	PAZ
RNase H	PIWI
3'-OH, 5'-phosphate oligonucleotide	3'-OH, 5'-phosphate oligonucleotide
RT (primer-grip)	PAZ pocket
RT (RNA-dependent DNA polymerase/ DNA-dependent DNA polymerase)	RNA-dependent RNA polymerase
RT (DNA unwinding)	Helicase
Integrase (dinucleotides overhangs)	Dicer (dinucleotides overhangs)
Nucleocapsid (melting, RNA protection)	RNA Binding Proteins, TRBP, FMRP
TRBP	TRBP
Protease	Caspases, PPARP

Today's viruses as witness of evolution

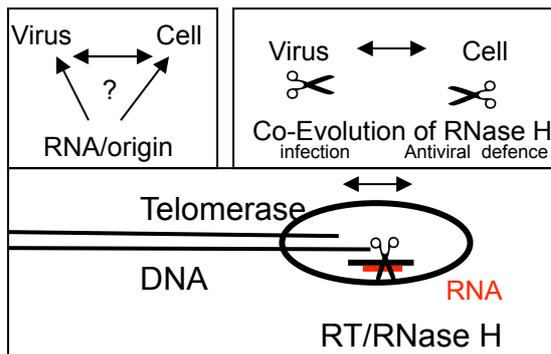
1.R-Rz --->2.R-RR--->3.R-RD-DD ---> 4.DD-R --->5.DD-DDR

- 1.RNA viroids, Ribozyme, catalytic, can open and close chemical bonds, bacterial viruses, plant viroids, Hepatitis Delta Virus
- 2.RNA Viruses, RNA into double-stranded RNA, then progeny RNA, bacterial viruses, Influenza, Hanta, SARS, single or segmented RNA
3. Retroviruses, RNA-DNA Hybrid intermediates, Double-stranded DNA proviruses, integrate, inheritable, stable for generations, transposons? Retroviruses (HIV, LV), und Retrotransposons, Telomerases(rudimentary RT with one RNA seq)
- 4.Para-Retroviruses, double-stranded DNA inside virions, single-stranded progenomic RNA, Hepatitis B, Caulimo Viruses, HBV developed Central Dogma and cellular nucleus
- 5.DNA into DNA via mRNA, all Organisms, Adeno, Papilloma, Herpes, Hepatitis C viruses, Transposons? Satellites of cellular origin?

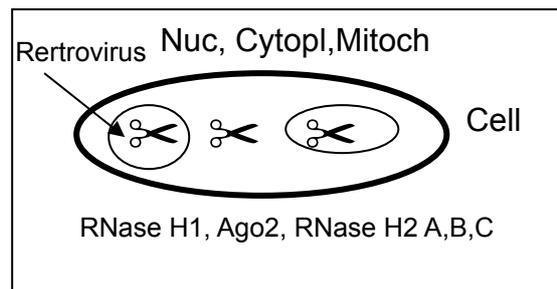
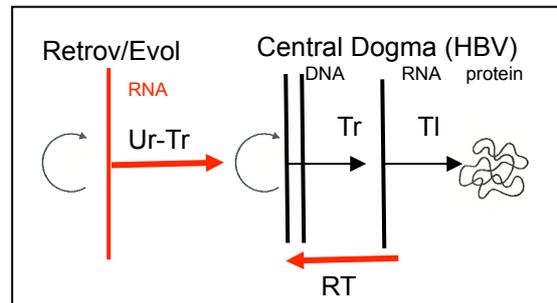


Structural/enzymatic relationship - evolutionary ?

AGO2	N-term	PAZ/PIWI
HIV	NC?	RT/RNaseH



Dicer protects against Influenza A



50% of human genome defective retroviruses, transposable elements, Junk? Benefit?

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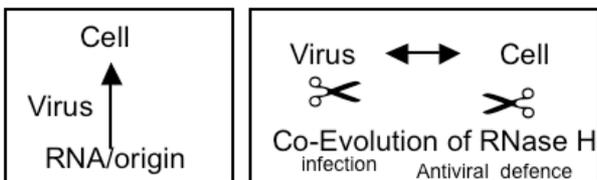
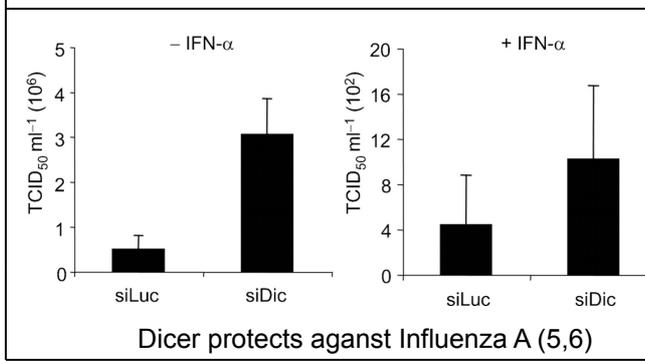
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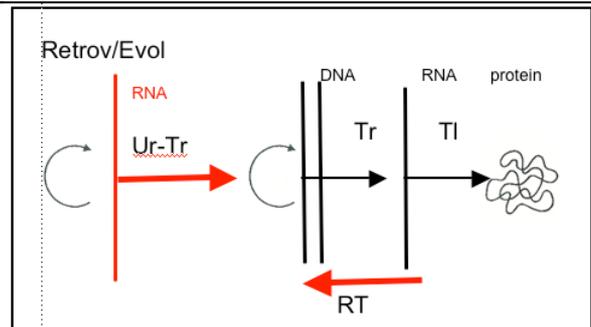
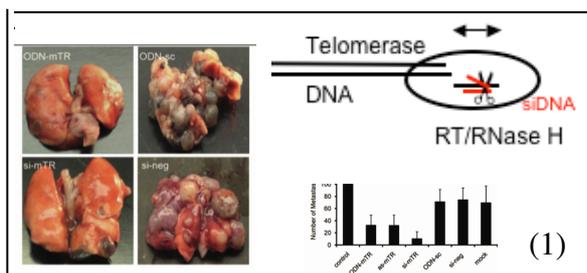
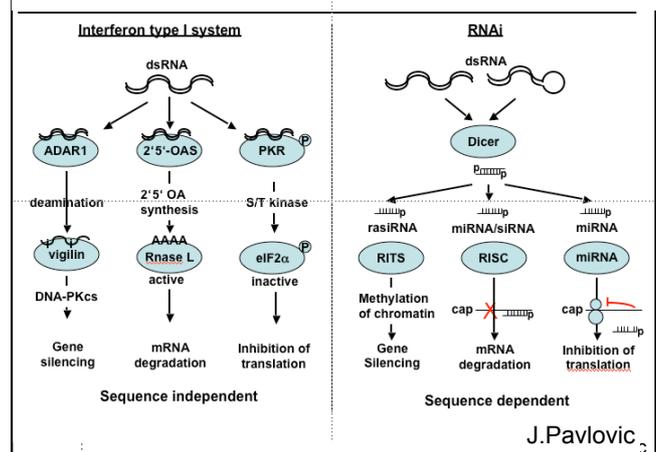


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Small interfering RNA related to interferon antiviral

Analogies between Interferon and RNAi pathways



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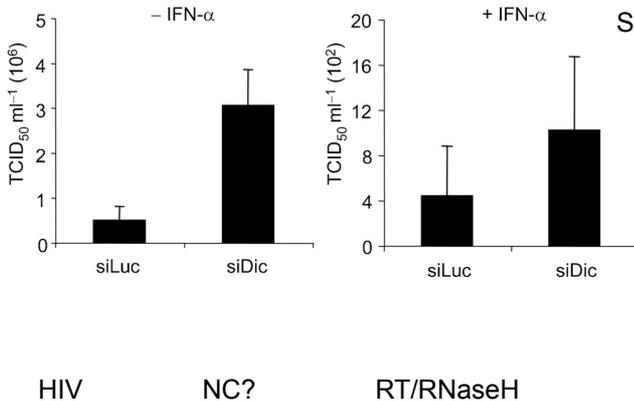
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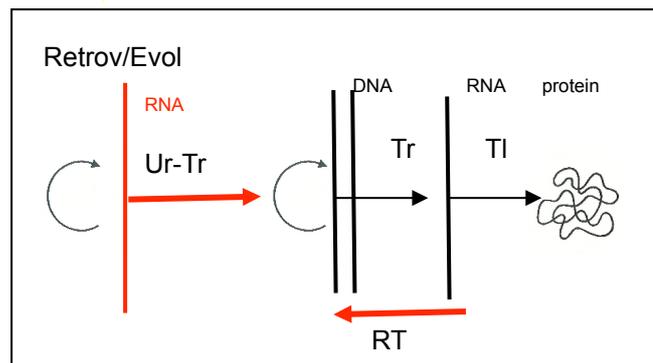
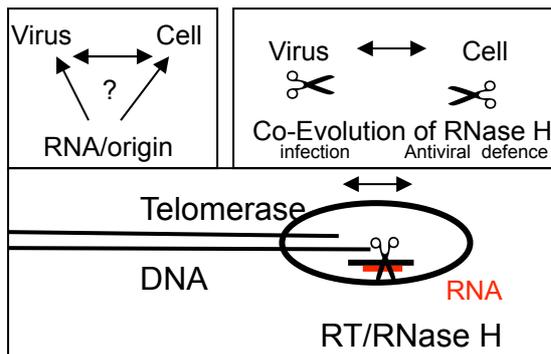
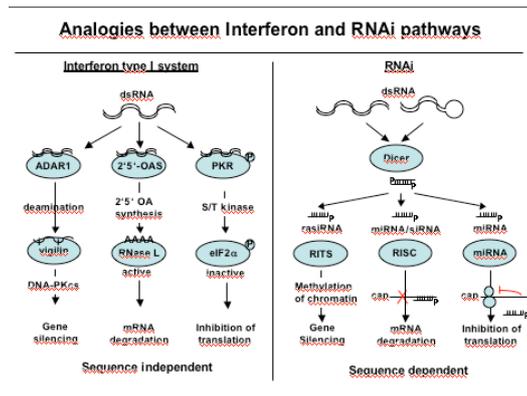


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