



## **5<sup>th</sup> Congress of the European Academy of Neurology**

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### **Teaching Course 13**

**Nervous system disorders due to retroviruses (Level3)**

**The possible role of retroviruses in  
neurological disorders of unknown aetiology**

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# The possible role of retroviruses in neurological disorders of unknown etiology



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## **Outline**

**Virus etiology in MS**

**Retrovirology**

**HERV in MS**

**HERV in ALS**

**Conclusions**



## *Viral etiology and MS*

An infectious origin of MS was suggested  
for the first time by Pierre Marie in  
1884

The topic is still relevant and unsettled

## *Viruses and MS: possible scenarios*

1. Single initial viral infection initiates chronic immune-mediated inflammatory reaction
2. Recurrent viral infections due to same virus or different viruses that trigger a relapse
3. Reactivation of latent or persistent virus or viruses within the CNS

## *Evidence for Viral etiology of MS*

1. Epidemiological Data
2. Laboratory Data in MS patients:
  - a) Serological data
  - b) Virus isolation
  - c) Viral histochemistry & molecular studies
  - d) Abnormalities of immune regulation
3. Animal models of Virus-Induced Demyelination

## *Epidemiological Data*



Faroe Islands Data

## *Serology: Higher antiviral antibodies in MS than in controls*

Serum	CSF
Measles	Measles
Parainfluenza 3	Parainfluenzas 1-3
Influenza C	Influenza A, B
Varicella	Varicella
Herpes simplex	Herpes simplex
Rubella	Rubella
Epstein-Barr	Epstein-Barr
HTLV-I (gag)	Mumps
HTLV-II	Respiratory syncytial
HHV-6	Coronaviruses Adenoviruses HTLV-I (gag) Simian virus 5

## *Possible culprits: isolated viruses*

- rabies virus,
- HSV,
- scrapie prion,
- parainfluenza virus 1,
- measles virus, simian virus 5,
- chimpanzee cytomegalovirus,
- coronavirus,
- EBV,
- tick-borne encephalitis virus,
- HTLV-1,
- VZV,
- HHV-6.

- Proposed viral mechanisms:
- direct brain or peripheral infection
  - activation of autoreactive T cells against nerve myelin
  - bystander activation
  - epitope spreading,
  - molecular mimicry,
  - virus-virus interactions.

***Leptomeningeal cell line from MS with reverse transcriptase activity and viral particles.***

H. Perron, C. Geny, A. Laurent, C. Mouriquand, J. Pellat, J. Perret, J.M. Seigneurin

***In Vitro transmission and antigenicity  
of a retrovirus isolated from an MS patient***

H. Perron , B. Gratacap, B. Lalande, O. Genoulaz, A. Laurent, C. Geny, M. Mallaret, P. Innocenti, E. Schuller, P. Stoebner, J.M. Seigneurin

***Molecular identification of a novel retrovirus repeatedly isolated from patients with MS***

H. Perron, J.A. Garson, F. Bedin, F. Beseme, G. Paranhos-Baccala, F. Komurian-Pradel, F. Mallet, P.W. Tuke, C. Voisset, J.L. Blond, B. Lalande, J.M. Seigneurin, B. Mandrand, and the Collaborative Research Group on Multiple Sclerosis.

***MS and HTLV retrovirus***

Hilary Koprowski, Elaine C. DeFreitas, Mary E. Harper,  
Magnhild Sandberg-Wollheim<sup>3</sup>, William A. Sheremata<sup>4</sup>,  
Marjorie Robert-Guroff, Carl W. Saxinger, Mark B. Feinberg,  
Fossie Wong-Staal, Robert C. Gallo

***Amplification and molecular cloning of HTLV-1 sequences from DNA of MS patients***

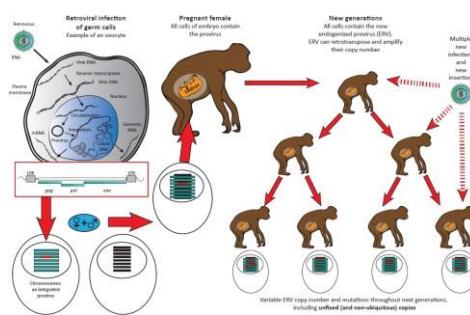
E. Premkumar Reddy, Magnhild Sandberg-Wollheim,  
Richard V. Mettus, Phillip E. Ray, Elaine DeFreitas,  
Hilary Koprowski

## Animal models: Some viruses capable of producing demyelination

- JMH strain of mouse hepatitis virus
- Theiler's infection of mice
- Visna virus of sheep
- Canine distemper
- Measles
- Mumps
- Influenza
- Papovaviruses
- Herpes simplex virus
- HIV
- Togaviruses

## Human endogenous retroviruses (HERVs)

- sequences belonging to human ERV families retaining structural features of retroviral genomes that have become integrated into the genome through repeated infections during evolution.



## *Endogenous Retroviruses*

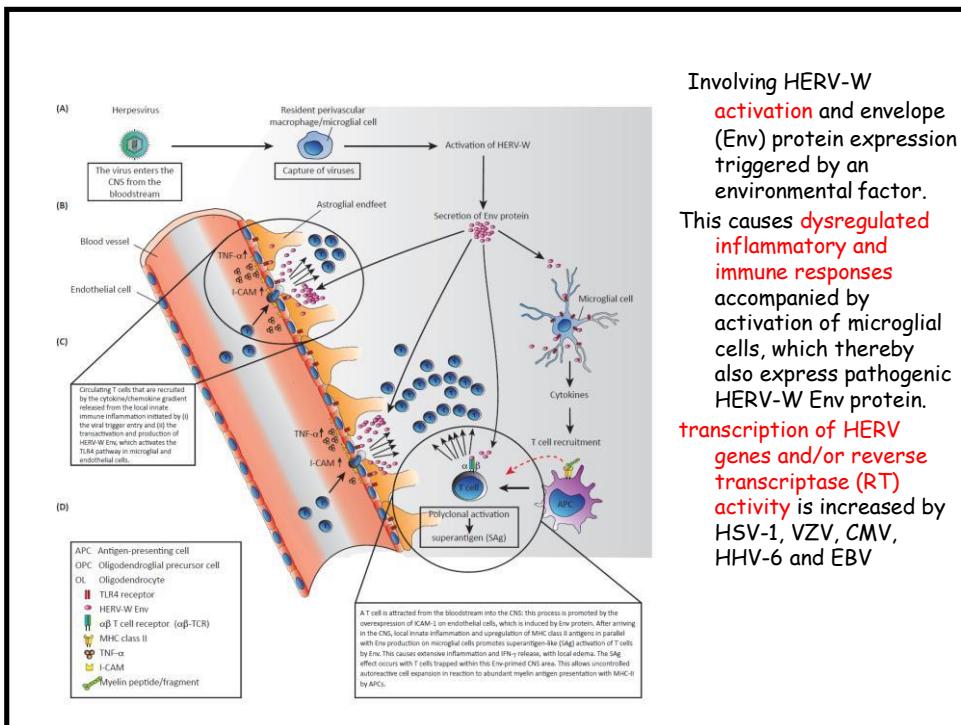
- The eukaryotic genome is composed of DNA sequences, many derive from mobile genetic elements estimated to account for about 50% of the entire human genome.
- There are 31 different **families of HERVs that together make up about 8% of the human genome**, four times more DNA than is devoted to protein coding genes
- HERV-W makes up about 1% of the human genome and is part of a superfamily of repetitive and transposable elements.

## *Silent HERVs can be activated by environmental riggers*

- DNA methylation & histone modification are essential to epigenetic control of human genes, HERV including.
- HERVs activation is linked to chromatin state.
- The baseline predisposition of a HERV copy to be activated can be tissue, cell, or maturation stage-specific.
- Inflammatory stimuli may activate HERVs via epigenetic dysregulation such as proinflammatory cytokines that act in cultured cells from MS patients

# EBV

- The most consistent and independently confirmed studies are for EBV by
  - history of infectious mononucleosis
  - high anti-EBNA-1 (EBV nuclear antigen 1) IgG titers before MS onset.
- EBV is known to activate HERV-W/MSRV *in vitro* and *in vivo*.
- Thus, EBV could be an initial trigger, while HERV-W/MSRV is a direct neuropathogenic contributor, before and during MS, in addition to its known contribution to promoting autoreactive T cells, immunoinflammation, and remyelination blockade.



## *Additional triggers besides EBV*

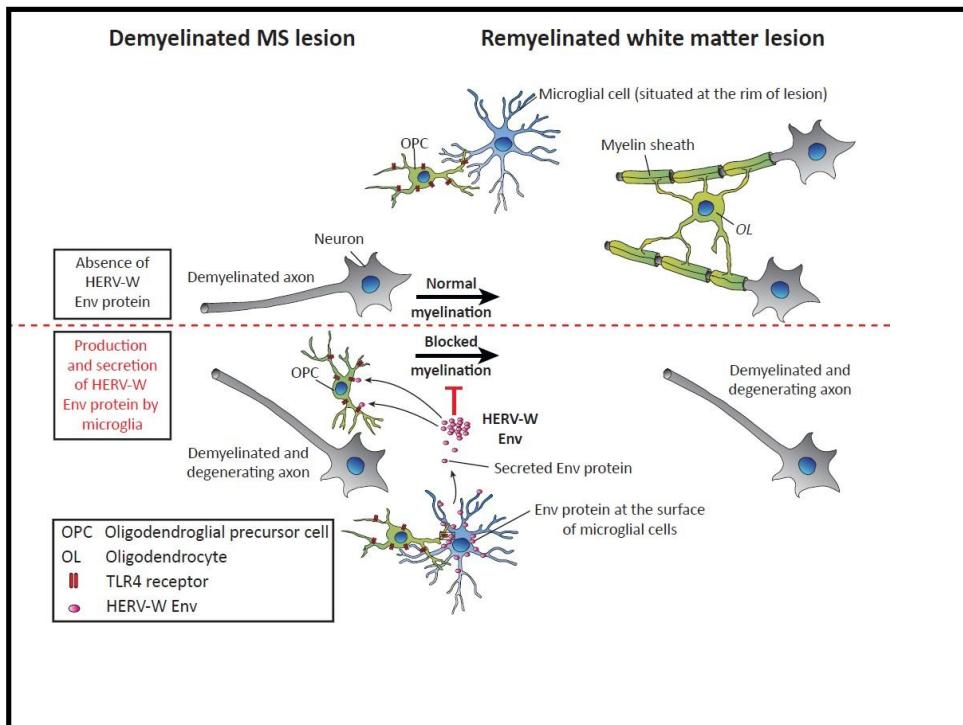
- **Herpesviridae** are now suggested to upregulate HERV-W expression, with its Env protein acting as a pathogenic effector in MS.
- HERVs express pathogenic proteins in disease, and the best evidence of an association and pathogenic involvement is for HERV-W/MSRV (detected in MS blood, spinal fluid, and brain, in parallel with MS stages, active/remission phases, and therapy outcome).
- **Other viruses** reported to transactivate HERVs are the exogenous retroviruses HTLV-1 and HIV-1

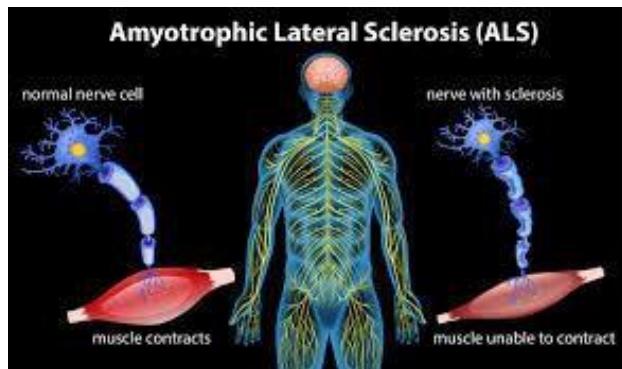
## *Mechanisms of activation*

- If HERV proteins are not expressed, HERV RNA expression (transcription alone) does not seem to have biological effects per se in humans.
- When produced, HERV proteins are not implicitly pathogenic.
- HERV H, K, W are abnormally represented in MS

## Therapy

- There is correlation between serum levels of HERV-W proteins (Env) and MS activity to suggest that it might be used as a therapeutic target in MS.
- A one year study phase IIb with a monoclonal antibody directed against Env in 260 RRMS patients is now on going





## HERV-K and ALS

- HERV-K is so-named because it uses lysine (K) tRNA as a primer
- 11 largely complete proviral sequences have been identified in human genomes
- Some endogenized HERV-K elements have complete ORFs and can form complete viral particles
- It is unknown if expression of HERV-K in ALS patients derives from a single complete retroviral sequence or represent trans-complementation between partially defective complementary copies.

## ALS & Retroviruses

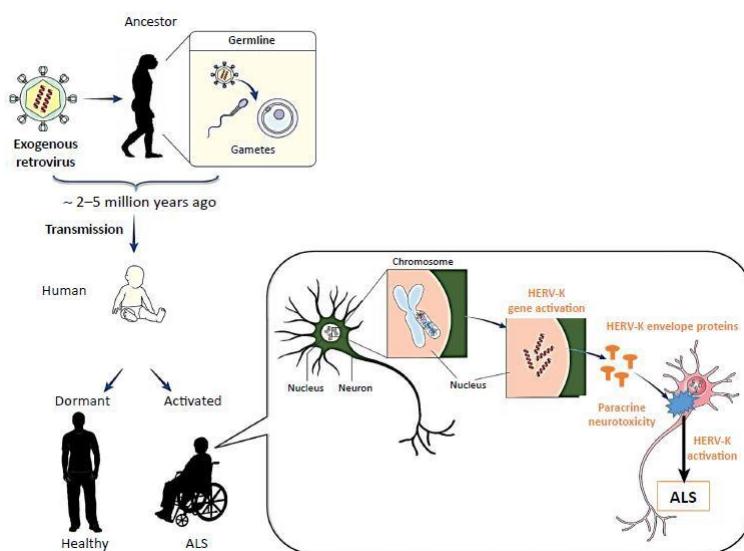
- The first evidence that retroviral elements might be activated in ALS came from a study where brain tissues from **two ALS patients in Guam** were found to have RNA-directed DNA polymerase activity.
- The **activity was RNase-sensitive suggesting RT activity**.
- Studies in ALS patients confirmed the presence of RT in serum and showed that **nearly 50% of the patients have detectable RT activity**.
- However, attempts to find an exogenous retrovirus in ALS patients were unsuccessful and no virus or transmissible agent was identified.

## ALS & Retroviruses

- The fact that HERVs in ALS arise from the genome and not from the environment might explain why RT was detected in ALS brain and blood samples, but no human-to-animal or human-to-human transmission of the disease was documented.
- RT encoded by the pol gene of HERV-K reported in the brain of sporadic ALS patients was specific for ALS
- Expression levels of HERV-K pol, env, and gag genes in brains of sporadic ALS patients correlated with each other, suggesting that an entire HERV-K genome is activated

## HERV-K & ALS

- If HERV-K expression is forced in neurons, it causes cellular degeneration mediated by its Env protein.
- Transgenic mice expressing HERV-K Env in neurons develop a clinical & pathological phenotype that resembles ALS, with upper and lower motor neuron degeneration
- What triggers HERV-K expression in adult neurons of ALS patients remains unknown.



## Antiretroviral therapy

- An approach similar to that taken for HIV could be considered
- A panel of antiretroviral drugs approved for treating HIV infection was screened, but elevated concentrations were found to be necessary to control HERV-K replication in HeLa cells *in vitro*
- A pilot clinical trial with indinavir, a protease inhibitor used for HIV, failed to show any efficacy in ALS

## Antiretroviral therapy

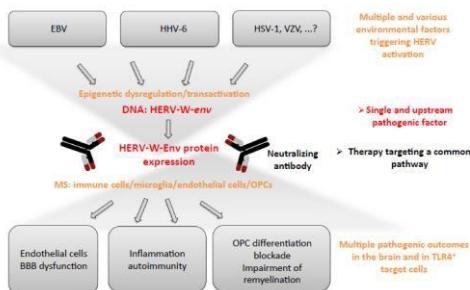
- Some patients with HIV infection can develop an ALS-like syndrome, which may show symptom regression and halted evolution under treatment with anti-HIV drugs
- These patients also have expression of HERV-K in blood, with levels that fell after antiretroviral drugs therapy
- It is possible that anti-HIV drugs might indeed inhibit HERV-K, but that their efficacy is not as good as for HIV

## Open trial

An open-label pilot study has been initiated in Australia and the UK. The trial is enrolling 40 patients with ALS, and will follow them for 3 months without treatment; then treat them for 6 months with triumeq - which includes two reverse transcriptase inhibitors (abacavir and amivudine) that have been shown to effectively inhibit HERV-K RTactivity *in vitro*, and an integrase inhibitor dolutegravir

## Therapeutic options for MS & ALS

- To test antiretroviral drugs
- To use a humanized neutralizing antibody targeting the toxic Env protein that mediates HERV-W pathogenicity



## Take home messages

- HERVs are evolutionarily acquired, mostly defective and inactive, and are epigenetically silenced genetic elements.
- They might have pathogenetic significance in neurological diseases.
- How can we neutralize HERV endogenous proteins is a major therapeutic challenge. Potential approaches include vaccination, antibody-mediated neutralization of pathogenic components, and antiretroviral compounds.

