



INVESTING IN YOUR FUTURE

DNA INTEGRITY AND EXPRESSION OF PROTEASOMAL GENES IN MULTIPLE SCLEROSIS

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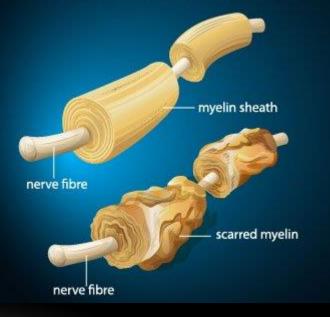
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MULTIPLE SCLEROSIS

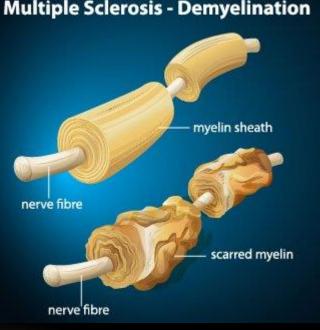
- Autoimmune disease
- Neurodegenerative disese
- Disease of the central nervous system brain, spinal cord, and optic nerves
- Sclerosis means scarring
- Consequences neurological disabilities

Multiple Sclerosis - Demyelination



MULTIPLE SCLEROSIS

- Around 2.5 million patients worldwide
- The chances of developing the condition are highest between ages 20 and 50
- Women have about 3 times increased likelihood of developing MS compared with men



Multiple Sclerosis - Demyelination

CAUSES

- The cause of MS is not clear
- Genetic susceptibility:
 - MS susceptible loci had been identified in the regions containing genes with immune, costimulatory, signal transduction functions and related to vitamin D function



Environment

Pathogens Chemicals Smoking Diet Sun exposure

Epigenetic, post-genomic and regulatory events

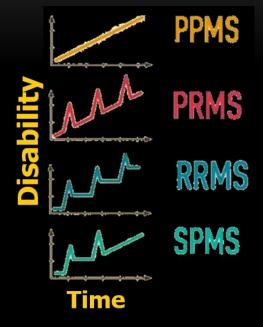
Gene rearrangements Messenger RNA splicing and/or editing Retroviral sequences Methylation MicroRNAs

Genome allelic variants

Single nucleotide polymorphisms Copy number variation Insertion/deletion polymorphisms Disease modifier genes Disease susceptibility genes

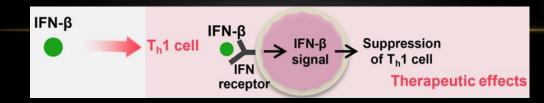
TYPES OF MS

- Primary-progressive (PPMS): 10-15%, MS slowly but steadily worsens.
- Progressive-relapsing (PRMS): 5%, the underlying disease steadily worsens. The patient has acute relapses, which may or may not remit.
- Relapsing-remitting (RRMS): 85%, symptoms flare during acute attacks, then improve nearly completely or "remit." This is the most common form of MS.
- Secondary-progressive (SPMS): Begins as relapsing-remitting type, then becomes progressive.
- In RRMS an inflammatory process predominates whereas in SPMS and PPMS a neurodegenerative process is more strongly expressed



INTERFERON BETA THERAPY

- T cells primary effectors of the autoimmune CNS inflammation
- Inside the CNS, activated T cells including T helper-1 (Th1) secrete proinflammatory cytokines
- Interferon β (IFN β) is the first class of disease modifying therapies for MS
 - directly increases expression and concentration of anti-inflammatory agents
 - downregulates the expression of proinflammatory cytokines
- IFNβ treatment may
 - reduce the trafficking of inflammatory cells across the blood-brain barrier
 - increase nerve growth factor production
 - lead to a potential increase in neuronal survival and repair



DNA LESIONS IN MULTIPLE SCLEROSIS

- Reactive oxygen species (ROS), if produced in excess, lead to oxidative stress
- ROS have been implicated as mediators of demyelination in MS
- The brain is particularly vulnerable to oxidative damage due to:
 - elevated use of oxygen
 - low antioxidant levels
 - high phospholipid levels
- The excessiveness of ROS causes oxidative damage to DNA
- ROS attack the nitrogenous bases and the sugar phosphate backbone
 → single- and double-stranded DNA breaks



DNA LESIONS IN MULTIPLE SCLEROSIS

- Blood sera of MS patients contain DNA-hydrolysing antibodies
- Antigen-activated lymphocytes bear an unusual genomic stress
 - DNA damage-response signalling pathways are activated
 - A similar situation is in the lymphocytes of the MS patients
- There was found increased level of γ-H2AX and poly(ADP) ribose in nucleated blood cells of the MS patients.
- Alkaline single-cell electrophoresis or comet assay a versatile and comparatively simple approach to detect DNA single-strand breaks



NITRIC OXIDE IN MULTIPLE SCLEROSIS

• Overproduction of nitric oxide (NO) by inducible NO synthase (iNOS) in glia

subsequent lesions of mitochondrial DNA and demyelination

↓ a crucial step in the pathogenesis of MS

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• Increased amount of nitrates and nitrites in the blood serum of MS patients

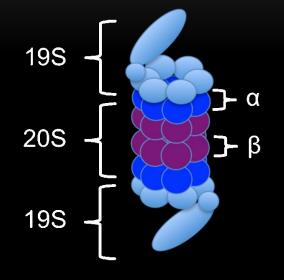
PROTEASOMES

- Accumulation of toxic protein aggregates can be observed in MS patients
- Ubiquitin-proteasome system (UPS) is responsible for degradation of intracellular proteins
- UPS is one of the key factors in regulation of the immune system
- MS \leftrightarrow dysfunction of UPS
- IFNβ therapy:
 - increased proteasome concentration
 - decreased proteasome activity
 - improved clinical course of MS

PSMA3, PSMA6 AND PSMC6

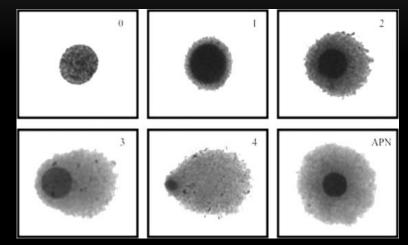
- Susceptibility of PSMA3, PSMA6 and PSMC6 genotypes involving rare alleles with autoimmune diseases
- Susceptibility of the *PSMA3* rs2348071 heterozygous genotype to multiple sclerosis
- Association of the *PSMA6* rs2277460 heterozygous genotype with MS in female group
- Association between *PSMC6* rs2295826 and rs2294827 loci and MS, for genotypes involving rare alleles

(*PSMC6* rs2295826 and rs2294827 loci are in complete linkage disequilibrium)



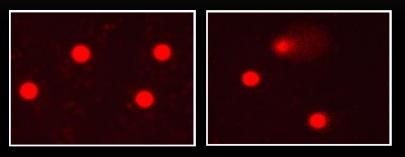
METHODS

- Comet assay
- Direct measurements of the NO radical by electron paramagnetic resonance (EPR) spectroscopy
- qPCR
 - PSMA3
 - PSMA6
 - PSMC6

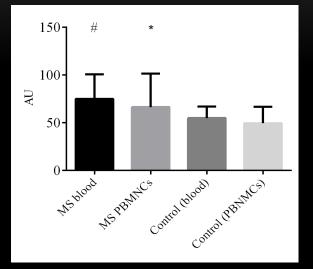


Comet types with the increasing damage. 0 – healthy cell and 4 extremely damaged cell. APN – apoptic nuclei (from: Collins et al. *Environ Mol Mutagen*. 1997;**30**, 139–146)

DNA INTEGRITY

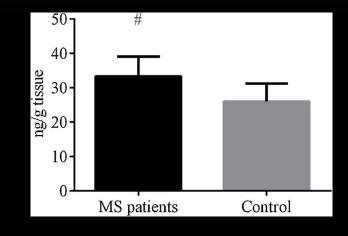


Left: Type 0 comets in the sample of a healthy patient. Right: Types 0, 1 and 3 comets in the sample of a patient with MS. Stained with EtBr. 40x magnification.



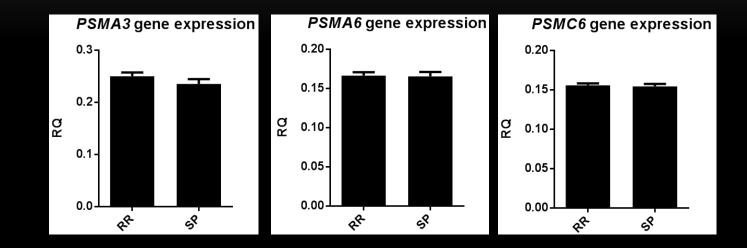
and *- p < 0.05 of whole blood and PBMNCs of MS group vs. control group

NO LEVEL IN BLOOD



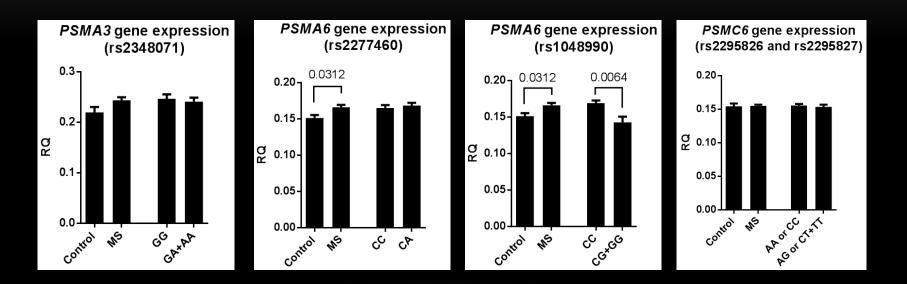
- p < 0.05 vs. control group

MS SUBTYPES

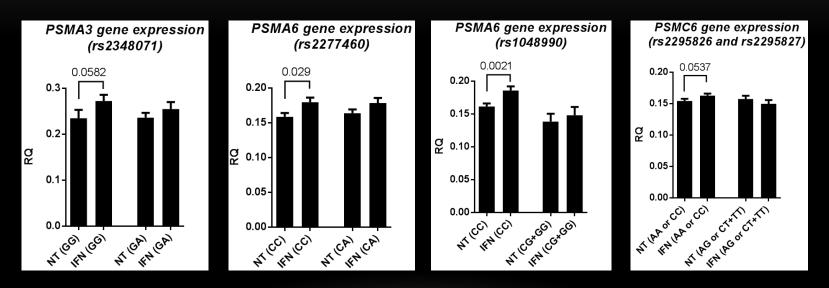


RR - Relapsing-remitting multiple sclerosis SP - Secondary-progressive multiple sclerosis

GENOTYPES



INTERFERON β THERAPY AND GENOTYPES



NT – no treatment; IFN - interferon β therapy

CONCLUSION

- Patients with MS have elevated level of DNA damage
- Patients with MS have significantly higher level of NO in the blood
- Overall group of MS patients have higher *PSMA6* gene expression
- IFNβ therapy increases PSMA3, PSMA6 and PSMC6 gene expression in patients with common genotypes compared to non-treated patients



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