Special Report: 
Multiple Sclerosis—
The Truth Lies Within Us

Data presented at the recent ECTRIMS offer fresh insight and potentially new direction into the cause of multiple sclerosis.

BY DANIEL KANTOR, MD

Conferences are boring. Ever since press releases have mostly preceded and preempted scientific platform presentations at prestigious international conferences, the multiple sclerosis (MS) community has a lot less truly novel data to look forward to at major medical conferences, such as the annual Congress of the European Committee for Treatment and Research in MS (ECTRIMS). That was, until October 28, 2017, when Hans-Peter Hartung, MD, Chairman of the Department of Neurology of University of Düsseldorf in Germany, delivered a late-breaking lecture on the 24-week results from CHANGE-MS (Clinical trial assessing the pHERV-W Env Antagonist GNbAC1 for Efficacy in MS).1

Unlocking the Mysteries of MS

The pinnacle of the most important worldwide gathering of MS sub-specialists is the presentation of late-breaking abstracts that are embargoed until that time, which means that even the most influential MS specialist waits eagerly to learn what’s new. At 9:30am Paris time, when Dr. Hartung took to the stage for the penultimate time slot, few in the audience of several thousand doctors, scientists, and others, knew what to expect. What seemed like a fairly standard post hoc analysis of negative trial results was actually the most visible introduction to the key that may unlock the mysteries of MS.

GNbAC1 isn’t just another unnamed pharmaceutical compound, but rather the test case for a novel way of potentially understanding the actual cause of MS. Our current understanding of MS is that it is an autoimmune disease affecting the central nervous system (CNS). More than 200 genes have been associated with the risk for MS, and it is believed that an early (prepubescent) environmental factor triggers the neuro-inflammatory cascade that is characterized, in the vast majority of patients, by relapses of CNS dysfunction. Since MS was first described by Jean-Martin Charcot in the 19th century, there have been revolving theories concerning its underlying cause; the viral theory remains the most widely cited. Although everyone with MS has evidence of prior exposure to Epstein-Barr virus (EBV), based on immunoglobulin levels, and even actual viral particles identified in the meninges in some studies, the problem with the viral theory is that almost everyone in the general population has also been exposed to EBV.

Thus, if a common virus truly causes MS, then why does MS only affect a small fraction—1%—of the population? Also, if there isn’t a viral cause for MS, then why does it act like a postviral neurological syndrome?

The answers to these important questions lie in the insights gained by CHANGE-MS.

The Other 8%: Foreign DNA Lurking Inside Us

Instead of an early childhood viral exposure triggering MS in a genetically susceptible individual, CHANGE-MS suggests the possibility that the initial viral basis of MS is actually ancient viral DNA that lies dormant within us and is roused after having been incorporated into our genome for millions of years. For the past century, instead of looking for an external cause of MS, we should have been focused inward to the 8% of foreign DNA already incorporated into our genome. These DNA elements are termed pathogenic human endogenous retroviruses (pHERVs), and the expressed pHERV-W envelope (Env) protein is thought to be pathogenic in MS; pHERV-W Env is found in active MS lesions on monocytes and microglia.2–7 It still takes an environmental exposure, such as viral infection with EBV, to de-repress the pHERV-W Env gene
and allow transcription of mRNA (messenger RNA) and expression of the pHERV-W Env protein. pHERV-W Env is an agonist of toll-like receptor 4 (TLR4), leading to a pro-inflammatory cascade and inhibition of oligodendrocyte progenitor cell (OPC) maturation through nitrosative stress.\(^2\)\(^-\)\(^7\) GNbAC1 is a recombinant, humanized IgG4-kappa monoclonal antibody, which binds with high affinity to pHERV-W Env, thereby blocking pHERV-W activation of TLR4 and rescuing myelin expression in OPCs.

**New Hope**

Because GNbAC1 does not seem to have other mechanisms of action, any benefit would seem to argue that the HERV hypothesis in MS has some merit, and we may be considering a whole new understanding of the underlying cause of MS. Unfortunately, the August 28, 2017 press release by the companies developing GNbAC1 (GeNeuro SA and Servier) reported that while safe and well tolerated, GNbAC1 did not reach its primary six-month endpoint of reducing cumulative brain MRI gadolinium-enhancing lesions at 12, 16, 20, and 24 weeks as compared to baseline. Dr. Hartung’s presentation explained these primary results because GNbAC1, which is not an immunosuppressant, may simply take longer to help in MS, and in the post hoc analysis of patients with active MRIs, there is a statistically significant reduction in gadolinium-enhancing lesions when comparing week 24 to baseline data.\(^1\) Additionally, there was a statistical trend favoring remyelination, evidenced by improvement on magnetization transfer ratio in the normal-appearing white matter and cerebral cortex of people with MS treated with the highest dose of GNbAC1 studied in CHANGE-MS. These promising results highlight why GNbAC1 is the most interesting MS compound to watch next year; of note, 48-week results are expected in the first few months of 2018.

**Looking Beyond MS**

Since the FDA approval of the first disease-modifying therapy for MS in 1993, there has been an explosion of new approved therapies, with a current count of 13 separate branded products, one branded generic, and two unbranded generic disease-modifying therapies (with one to two new ones expected yearly for the next several years). The HERV hypothesis in MS is leading the way for other pHERV subclasses in other neurologic and autoimmune diseases. For example, there is ongoing interest and research into pHERV-K in amyotrophic lateral sclerosis and an enrolling Phase 2 trial of GNbAC1 in type 1 diabetes mellitus. The fate of our understanding of currently incurable neurologic and nonneurologic diseases lies in the results of GNbAC1 in MS. We have a lot to be excited for in 2018.

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