



Commentary

Extrahepatic manifestations and HEV, the genotype matters



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Extrahepatic manifestations and viral hepatitis: a topic attracting the attention of hepatologists for more than 30 years. Hepatitis B and C virus infections have been associated with various extrahepatic diseases and in some of these phenomena a causal relationship has been proven.

Within the last two decades there was growing interest and awareness for hepatitis E virus (HEV) infections worldwide [1,2]. HEV genotype (GT) 3, the predominant GT in Europe and the USA, presents as zoonosis, mainly transmitted by infectious swine meat [3]. In contrast GT 1 and 2 are mainly transmitted by contaminated drinking water in tropical developing countries. A special situation is the setting of GT 4. This is a tropical virus, but in contrast to GT 1 and 2 it is mainly transmitted by consumption of swine meat, as GT 3 [1,2].

In the last five years several assumed extrahepatic manifestations have been observed in the context of HEV infections [4]. In particular, pancreatitis has been associated with HEV GT 1 infections, while neurological diseases as neuralgic amyotrophy or Guillain-Barré syndrome have been observed in patients with GT 3 infections. One of the most impressive studies regarding this topic was the paper by Dalton et al. [5]. They studied the prevalence of acute HEV infections in a large cohort of neurological patients presenting with various non-traumatic acute neurological diseases. They reported evidence of current HEV infection in 11/464 patients (2.4%): HEV viremia was shown in seven patients and anti HEV IgM positivity only in four patients. Notably, none of these patients had jaundice or other clinically signs of acute hepatitis. The majority of these cases included Neuralgic amyotrophy and cerebral ischemia/infarction but also patients with seizures, encephalitis and acute facial and vestibular neuropathy. This observation affirmed an association of HEV and neurological diseases in general and especially Neuralgic amyotrophy.

In *EBioMedicine*, Wang et al. studied a similar approach from China, an area endemic for GT 4 [5]. They enrolled 1117 patients with neurological illness and 1475 healthy controls and tested them for HEV and anti-HEV. In contrast to the previous study from Europe, the rate of anti-HEV IgM antibodies did not differ significantly between groups, as authors found 6 positive cases in the neurological illness cohort (0.54%) and 10 positive cases in the healthy controls (0.68%). All detected individuals were HEV RNA positive. Patients with acute HEV infection included cases of viral encephalitis, posterior circulation ischemia, peripheral neuropathy as well as one case of Guillain-Barré

syndrome. Notably, rate of anti-HEV IgG antibodies did not show a significant difference between groups (neurological cases 39.51% vs. controls 35.63%; $p = \text{n.s.}$). Therefore, Wang et al. concluded that HEV genotype 4 seems not to contribute directly to acute neurological illness in China.

The study of Wang et al. raises several questions as their findings were unexpected and in absolute contrast to previous European and Japanese studies [6–9]. However, we have to be very careful in comparing such data. Whereas Wang et al. included a control group in their study, Dalton and van Eijk et al. reported cohort studies without controls [6,8,9]. Beside this, one could discuss that sample sizes and degree of heterogeneity of neurological entities of different etiologies vary a lot between these studies. Fukae et al. recently reported that 5% of Guillain-Barré patients in Japan were associated with acute HEV infection – in line with European findings by van den Berg [7,9]. However, what is true for Guillain-Barré syndrome does not need to be true in neuralgic amyotrophy. The main factor is the geographical distribution of HEV, as it can be assumed that the European studies reported solely HEV GT 3 associated cases, and the study by Wang et al. reported HEV GT 4 associated infections [5].

Irrespective from geographical factors, HEV GT 3 and GT 4 resemble each other regarding the majority of viral, epidemiological and clinical factors [2]. This is not really surprising if we look at the evolution of HEV: It has been supposed that lineages of the progenitors of HEV (which existed 536–1344 years ago) split into 2 main groups: the anthrotopropic GT 1/2 and the enzootic forms, GT 3/4 [10]. Here again, GT 3/4 behave similarly as an analysis of population dynamics of HEV proposed an expansion of both, GT 3 and GT 4, during the 20th century, followed by a clear decline in the 1990s. However, there are differences, e.g. GT 4 exhibits country specific variants, as in Chinese and Japanese claims different population dynamics have been observed [10].

The association of HEV and neurological disorders seems to be a further differentiation of GT 3 and GT 4 also depending on the neurological entity. Future studies are needed to further elucidate similarities and differences of these 2 GTs.

Disclosure

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