A BRAZILIAN HEPATITIS A VIRUS ISOLATED AND ADAPTED IN PRIMATE AND PRIMATE CELL LINE AS A CHANCE FOR THE DEVELOPMENT OF A VACCINE

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Hepatitis A is caused by a member of the genus Hepatovirus within the family Picornaviridae (R. I. B. Francki, C. M. Fauquet, D. L. Knudson, F. Brown (eds) 1991, Arch. of Virol., Suppl., 2: 320-6). The virus has a great importance for Brazilian Public Health because it is widespread in the environment. On one hand infection in children develops often as a subclinical or mild disease. On the other hand, overt hepatitis develops in the majority of adults, and the incidence of subclinical infection is very low (F. Sutmoller et al., 1982, Mem. Inst. Oswaldo Cruz, 77: 9-17). Hepatitis A virus (HAV) is responsible for 50% of acute hepatitis diagnosed in the National Reference Center for Viral Hepatitis, Virology Department, FIOCRUZ. Epidemiological data show that approximately 90% of the adult Brazilian population has antibodies anti-HAV IgG (A. R. N. Abuzwaide et al., 1987, Rev. Inst. Med. Trop. São Paulo, 29: 219-23) in opposition to developed countries, in which the prevalence of susceptible adults is very high with risk of huge outbreaks.


A Brazilian hepatitis A virus (HAF-203) was isolated in fetal rhesus kidney cells-FRhK-4 from faeces collected from an infected child in the beginning of the acute phase, (Gaspar et al., 1992, Braz. J. Med. Biol. Res., in press). A great tool for this study was the availability of tests, standardized by our group, for the detection of viral antigen (HAVAg) and RNA. Enzyme immunoassay (EIA-HAVAg) was applied for viral protein detection; specific human anti-HAV IgG, from serum of individuals recovering from hepatitis A infection, was purified by a combination of ammonium sulfate precipitation followed by anion exchange chromatography and used to coat the surface of the solid phase and conjugated with horseradish peroxidase. RNA was detected by cDNA-RNA hybridization using a 32p labelled nick translated cDNA probe kindly supplied by J. R. Ticehurst (NIH, Bethesda, USA).

After the adaptation of HAF-203 in cell culture, we selected fast-growing virus particles with a higher antigenic titer. Through the follow-up of the viral growth curve in the third passage, after the adaptation phase to the cellular system, we could observe maximum levels of RNA on the 21st day and of viral protein on the 49th day. Based on these data, sequential passages were performed always using as inoculum the cellular lysate from the 21st day. Now, we have reduced the time for maximum protein release from 49 days p. i. (1st passage) to 7 days p. i. (8th passage).

The infectivity of HAF-203 was demonstrated in South-American marmosets Callithrix jacchus (C. L. Vitral, 1991, Infecção intragás-
trica em Callithrix jacchus com uma cepa brasileira do vírus da hepatite A – HAF-203, Thesis, Instituto Oswaldo Cruz, 131p.). In the acute phase, accentuated hepatic lesions were observed together with the elevation of aminotransferases. HAVAg was also demonstrated in hepatic and renal tissues using the technique of immunoperoxidase. We could observe that the course of experimental infection in these animals showed significant correlation to natural infection in man, as described by Teixeira et al., 1982 (Liver, 2: 53-60). These data were previously demonstrated with other species of marmosets (P. Karayiannis et al., 1986, J. Med. Virol., 18: 261-76) and chimpanzees (A. N. Schulman et al., 1976, J. Infect. Dis., 134: 80-4). C. jacchus, which was shown to be susceptible to hepatitis A virus, can now be used as an animal model for hepatitis A vaccine control. Currently, studies are being developed with the aim of cloning and sequencing the genome of HAF-203.