

Characteristics of patients with hepatitis C and distribution of viral genotypes in Lithuania

Čiupkevičienė Eglė, Šumskienė Jolanta, Kupčinskas Limas

Introduction

- In 2016, the World Health Assembly passed a resolution to eliminate hepatitis C as a public health threat by 2030.
- Effective drugs are now available; however, most patients remain unaware of the hepatitis C virus (HCV) infection, which may be recognized only in the late stages when the complications have occurred.
- Different virus genotypes may affect the pathobiology of disease.
- Thus, identifying patients at high risk of HCV infection and ensure proper treatment is a health care challenge.
- Aim of the study to evaluate the characteristics of patients with hepatitis C and HCV genotype distribution in Lithuania.





Methods

The study was carried out in three hospitals in Lithuania, which are the inpatient and outpatient centers consulting and treating patients with HCV infection. All patients who attended a routine visit at each hospital within 6 months and fulfilled the selection criteria were selected. Information was collected from the patient's medical documentation. Altogether 201 patients (51.7% men and 48.3% women) were involved in the study.

Selection criteria:

- positive HCV-RNR test;
- defined HCV genotype;
- age ≥ 18 .

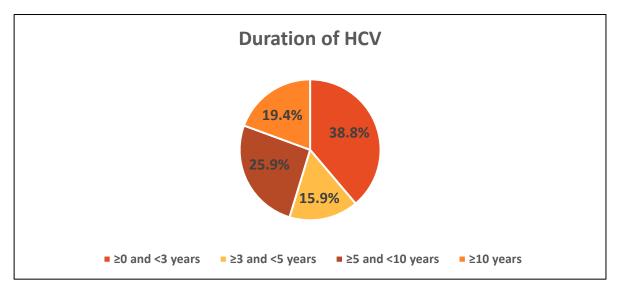
Collected information:

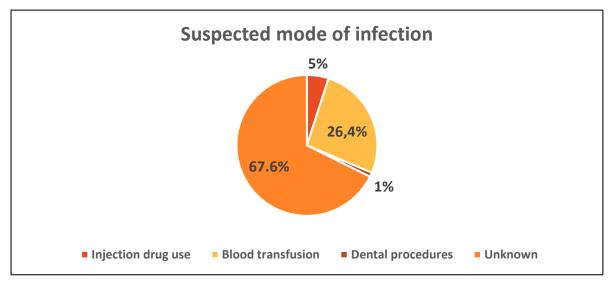
- demographic data;
- alcohol consumption and substance use;
- suspected mode of infection;
- duration of HCV infection;
- genotype;
- liver fibrosis stage;
- hepatocellular carcinoma status.

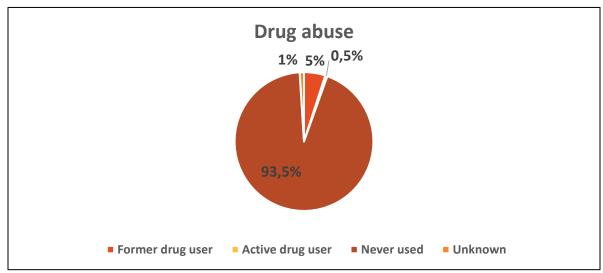




Results (1)



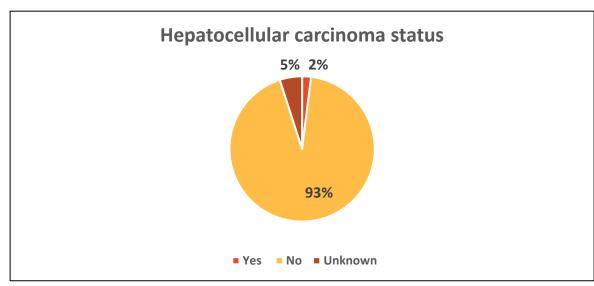


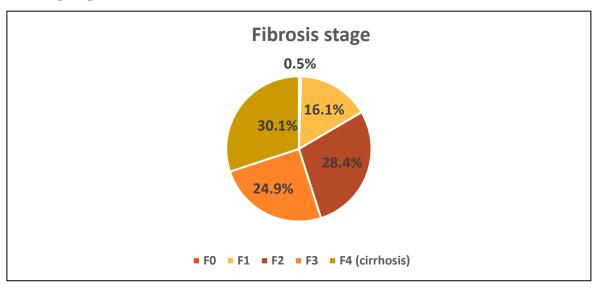


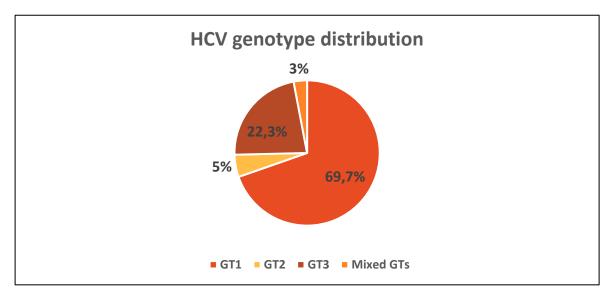


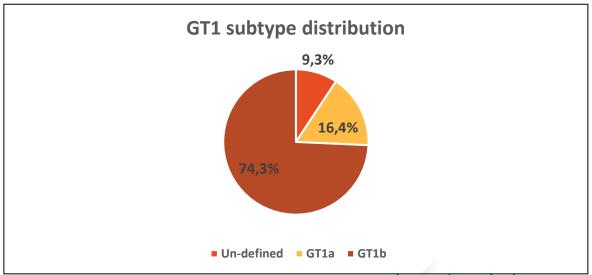


Results (2)











Conclusions

- Genotype 1 is the most common HCV genotype in Lithuania.
- Intravenous drug use was not the main mode of transmission of the infection.
- A high proportion of fibrosis grade 3 and 4 indicates the need for screening for this latent infection in the general population, especially in risk groups.

