

REVIEW ARTICLE

Hepatitis C: an Overview of Various Laboratory Assays with their Mode of Diagnostic Cooperation

Mohammad-Ayman A. Safi

Department of Medical Microbiology and Parasitology, Faculty of Medicine, King Abdul-Aziz University, Saudi Arabia

SUMMARY

Background: This review describes in chronological order the different assays for hepatitis C virus (HCV) antibodies, for the core antigen and for the HCV-RNA.

Methods: By ascending chronological order, the enzyme-linked immunosorbent assay (ELISA), rapid diagnostic tests (RDTs), HCV-Ab IgG avidity index (HCV AI), and Cy3-labeled microarray assay have been described for HCV antibodies in addition to ELISA for the total HCV core antigen (Ag).

Results: The recombinant immunoblot assay (RIBA) is a confirmatory test for HCV-Ab in blood, which is no longer needed due to the use of the sensitive third and fourth generation ELISA in addition to HCV-RNA detection by the Reverse Transcriptase-Polymerase Chain Reaction (RT-PCR). ELISA and RIBA measure current and prior exposure to HCV infection, but cannot discriminate between the two. RT-PCR is performed after ELISA for the diagnosis of HCV infection whether acute, chronic, false positive or false negative ELISA.

Conclusions: The cooperation between ELISA and RT-PCR in the diagnosis of HCV infection has been tabulated and discussed. HCV genotyping and subtyping testing is essential in pre-treatment evaluation of the patients for setting valuable treatment strategies and in understanding the epidemiology of the virus.

(Clin. Lab. 2017;63:xx-xx. DOI: 10.7754/Clin.Lab.2016.161113)

Correspondence:

Dr. Mohammad-Ayman A. Safi
Associate Professor of Immunology
Department of Medical Microbiology and Parasitology
Faculty of Medicine
King Abdul-Aziz University
PO Box 80205
21589 Jeddah
Kingdom of Saudi Arabia
Phone: +966 (2) 6400000 Ext 21116
Fax: +966 (2) 6403749
Email: aymansafi4@gmail.com

KEY WORDS

gastroenterology, HCV, ELISA, rapid diagnostic tests, RDT, RT-PCR, genotyping

INTRODUCTION

Hepatitis C virus (HCV) is an RNA virus belonging to genus Hepacivirus within the family Flaviviridae. It is estimated that about 3% of the world population is infected, with chronicity of about 80%, and is primarily parentally transmitted via blood transfusion, injection drug use, unsafe injection practices, and other health care related procedures [1].

The laboratory diagnostic tools of HCV infection are of two categories: indirect tests (serological assays) detecting specific antibody to HCV (anti-HCV) and direct assays that are directed toward the components of HCV viral particles, such as HCV RNA and core antigen. Both direct and indirect virological tests cooperate in the diagnosis of infection, therapeutic decision-making,

and assessment of virological response to therapy. In this review, various laboratory assays will be gathered and discussed in a chronological manner to show their concepts, developments, and principles and to demonstrate their applications and cooperation in the diagnosis of HCV infection.

I-Tests for HCV antibodies (HCV-Ab)

Enzyme-linked immunosorbent assays (ELISA)

Late in the last century, Hepatitis C Virus (HCV), was discovered as the etiologic agent for non-A non-B hepatitis [2-4], and testing began with the indirect enzyme-linked immunosorbent assays (indirect ELISA or indirect EIA) for detection of HCV-antibodies (HCV-Ab) appeared world-wide as 1st generation, then 2nd, 3rd, and 4th generation, with improvement in sensitivity and specificity [5-7]. In the indirect ELISA, the tested HCV-Abs interact with the recombinant HCV-proteins immobilized altogether on a plastic well, followed by washing, incubation in conjugate (peroxidase-labeled goat anti-human Ig), washing, and incubation with a substrate to yield a color. The intensity of the color is proportional to the concentration of the HCV-Abs. The RNA genome of HCV encodes at least nine proteins including six nonstructural (NS) proteins (NS-2, -3, -4A, -4B, -5A, and -5B) and three structural proteins (E1, E2, and Core) [5-7]. The original, 1st generation HCV-Ab EIA utilized a recombinant protein, c100-3, derived from the non-structural (NS-4) region of the viral genome as the solid-phase (immobilized) antigen [8]. The test was widely used, but found to possess low sensitivities for a high-prevalence population (approximately 80%) [9], inability to discover all HCV patients [10], long "window-phase" [11], and a high rate of false-positive reaction such as in primary biliary cirrhosis [12], rheumatoid arthritis [13], malaria [14], paraproteinemia [15], and in chronic active hepatitis [16], chronic liver disease [17] and autoimmune hepatitis [18].

Second generation ELISA (or EIA) included recombinant HCV antigens: c33c from the nonstructural NS-3 region, and structural (c22-3) protein from the core region of HCV [19,20]. They detected HCV-Ab at much higher rates [21]. The mean window-phase was reduced from 16 weeks (with first generation EIAs) to 10 weeks and the sensitivities were increased to about 95% [9]. In 1996, the U.S Food and Drug Administration (FDA) approved a third generation EIA that added a fourth antigen (NS-5) which allowed earlier detection of antibodies (26 days earlier) [22] and increased the sensitivity up to 97% [6]. Therefore, third generation EIA (and ELISA) for HCV-Ab was considered to be more sensitive and specific than those of the second generation [23,24].

Sensitive 4th generation ELISA has also been used, with increased sensitivity for NS (-3, -4A, -4B, -5A) in addition to the Core region, for which the antigens (NS-3 and NS-4) were derived from genotypes 1a, 1b, 2, and 3 [25]. The INNOTEST HCV-Ab IV showed a specificity of 99.8% and enhanced sensitivity [7].

ELISAs require sophisticated equipment, regular maintenance, a constant electricity supply, and skilled technicians. Thus, they are suitable for testing large numbers of samples per day, in blood banks or for surveillance studies such as screening of baby boomers. The generation born between 1945 and 1965 (baby boomers) account for approximately 3/4 of all chronic HCV infections in the United States. Thus, their screening for HCV infection was recommended by the Centers for Disease Control and Prevention (CDC) [26] and by the US Preventive Services Task Force [27-29].

Rapid diagnostic tests (RDTs)

As described above, ELISAs are sensitive and specific but sophisticated. Thus, they are called laboratory tests as they are not suitable for preliminary or emergency medical screening. Alternatively, Simple/Rapid tests are better for emergency testing, and in smaller laboratories with low numbers of tests per day. These are the tests that are quick (results in minutes) and easy to perform. They also allow point-of-care testing in primary care for different things that formerly only a laboratory test could measure [30,31]. They include agglutination, immuno-dot, immuno-chromatographic and/or immunofiltration techniques. In 2010, the FDA approved a Rapid Antibody Test (OraQuick HCV), in which HCV antigens (HCV-Ag) are fixed as a test strip on a cellulose membrane. HCV antibodies (if present) bind to HCV-Ag and then to colloidal gold particles (stored in the area) forming a red line indicating positivity. The test can be used for persons at risk for hepatitis or for those with signs or symptoms of hepatitis. The test strip can be used with a sample collected from a finger stick, venipuncture whole blood or oral fluids [32]. Screening for HCV in the emergency department (ED) has been found to be feasible, albeit costly [33,34].

HCV-Ab IgG Avidity Index (HCV AI) for HCV-Ab

HCV-Ab IgG Avidity Index (HCV AI) can be determined by ELISA by testing two aliquots of 200 μ L (for each serum sample), one diluted 1:10 with 1 M guanidine and the other with phosphate-buffered saline (PBS) and incubated for 10 minutes at room temperature. HCV AI is the ratio between the OD at 492 nm from the guanidine wells and the OD from the PBS wells [35]. HCV AI, within the first 8 days following the onset of clinical symptoms, may be useful in identifying actual acute HCV infection cases [36].

The recombinant immunoblot assay (RIBA) as complementary for HCV-Ab

Initially, the main problem encountered with HCV testing was the lack of confirmatory tests. Recombinant immunoblot assays (RIBA) [13,37,38] had been developed as supplemental tests to discriminate between true- and false-positive results for samples that are repeatedly reactive by ELISAs [39]. RIBA is similar, in principle, to ELISA except that the recombinant and synthetic HCV-proteins are immobilized as bands on a nitrocellulose

strip rather than a plastic well in addition to a washing step after the final incubation with substrate to yield colored bands. Second generation RIBA (RIBA HCV 2.0) consisted of a nitrocellulose strip solid support on which four bands of recombinant HCV proteins are immobilized. The four bands contained four different antigens: the 5-1-1 antigen, the c100-3 recombinant antigen, the c33-c recombinant antigen, and the c22-3 antigen (respectively derived from the NS-4 region, the NS-4 region, the NS-3 region, and from the core region of the HCV genome). Since 1992, a greater specific and sensitive third-generation RIBA HCV strip immunoblot system (RIBA HCV 3.0 SIA) has been widely used, which had four significant differences from RIBA HCV 2.0: 1) it included a recombinant antigen from the NS-5 region, 2) the recombinant c22-3 antigen was replaced by a four-epitope core synthetic peptide c22(p), 3) the 5-1-1 and c100-3 recombinant antigens were replaced by a mixture of synthetic peptides c100 (p), and 4) more c33-c recombinant antigen was used [40-44]. However, 3rd generation ELISA tests for HCV-Ab are both highly sensitive and specific to the extent that only one single ELISA determination is necessary for the diagnosis of HCV infection, without the need for confirmation by RIBA. PCR for HCV-RNA detection resolves weakly positive or negative ELISA results in cases that are clinically compatible with hepatitis C [45].

Fluorescent cyanin3 (Cy3)-labeled microarray assay for HCV-Ab detection

In 2008, a new inexpensive, specific, and sensitive method, for the massive sample diagnosis, was described, in which the target HCV-proteins are immobilized altogether in gel in each well of a new 96-well chip-format plate; then the bound HCV-Abs are detected by cy3-labeled anti-human Ig [46]. The sol-gel contents included tetraethoxysilane, PEG8000, HCl, and sodium phosphate (pH 7.5) [47]. Each individual well was spotted with four different HCV antigens (Core, NS-5, NS-3, and E1/E2) that conferred 1000 times more sensitivity than that of the ELISA with high specificity (98.78%) yielding significantly fewer false-positive results than did the ELISA [46].

II-Tests for components of HCV particles (core antigen and RNA)

Tests for HCV core antigen detection

The availability of an anti-core antigen (Ag) monoclonal antibody allowed development of an ELISA detecting and quantifying total HCV core Ag in serum and plasma of HCV-infected patients, which required an initial step of immune complex dissociation to remove bound anti-core antibodies prior to the antigen detection with the monoclonal antibodies [48].

It was claimed to be accurate, precise, and specific but cannot be used for HCV RNA values below 20,000 IU/mL, making its clinical usefulness limited as HCV-RNA detection methods are more sensitive [48].

A new generation chemiluminescence immunoassay (CLIA) based quantitative test (Architect HCV Ag Test, Abbot, Germany) had been shown to possess a sensitivity comparable to that of end point PCR (~1,000 IU/mL) but less than that of real time PCR [49, 50].

The Reverse Transcriptase-Polymerase Chain Reaction (RT-PCR) for viral RNA

Polymerase Chain Reaction (PCR) [51-54] is a technology using molecular biology discovered in 1985 by Kary Mullis [53]. This method allows the amplification *in vitro* of a specific region of DNA many million-fold, in order to produce enough DNA to be adequately tested [53].

The Reverse Transcriptase-Polymerase Chain Reaction (RT-PCR) [55] is used for amplification of RNA (e.g., HCV-RNA) which has to be converted to cDNA by reverse transcription, using reverse transcriptase, and then PCR is performed [56,57]. RT-PCR is performed after extraction of the HCV-RNA from serum, which is carried out by using extraction kits (such as QIAGEN) according to the methods of Vogelstein and Gillespie (1979) [58] and Stull and Pisano (2001) [59].

Immunoserological tests (such as ELISA) measure current and/or prior exposure to HCV infection, but cannot discriminate between the two [60]; therefore, RT-PCR, which detects HCV-RNA, is performed for viremia detection within a few days after exposure to the virus and many weeks before elevation of transaminases and antibody titre [61-63]. This helps in monitoring anti-viral treatment for chronic HCV infection [64], which depends on the viral load [65], and in the diagnosis of vertical transmission of HCV from chronically-infected mothers to their offspring [66,67]. The cooperation between ELISA and RT-PCR in the diagnosis of HCV infection whether acute, chronic, false positive or false negative ELISA will be discussed later on, in Table 1. The PCR-amplified DNA can also be generated and identified by using the automated technology such as Cobas Amplicor RT-PCR qualitative assay with the limit of detection of less than 50 IU/mL and can be used to confirm viremia and assess ultimate treatment response [68] and SMART CYCLERII, a real-time PCR thermo-cycler, by which thermal and optical data can be monitored in real time, and graphs of temperature, growth, and melt curves are displayed during data collection [69].

HCV-RNA transcription-mediated amplification (HCV TMA) qualitative test

Based on transcription-mediated amplification (HCV TMA), a highly sensitive and reliable qualitative chemiluminescent assay for detection of HCV-RNA in human serum or plasma, was developed and manufactured [70, 71]. In HCV TMA, following a lysing step, target HCV-RNA is captured on magnetic particles coated with oligonucleotides complementary to the 5' untranslated region of the HCV genome [72] in separate poly-

Table 1. Diagnosis of HCV infection according to HCV-Ab and HCV- RNA ¹.

Test results		Case	Note
Ab ²	RNA ³		
- ⁴	- ⁴	False negative?	The first 1 - 3 weeks after accidental parenteral exposure
- ⁵	- ⁵	Unlikely acute hepatitis C	Patient whose liver disorders are due to another cause or “occult HCV infection”
+ ⁶	- ⁶	Unlikely acute hepatitis C	Healthy carriers, possible resolved infection, “occult HCV infection”, transient patients or PCR inhibition
+ ⁷	? ⁷	Chronic hepatitis C	Detection of HCV-RNA confirms chronic HCV infection
+ ⁷	+ ⁷	Chronic hepatitis C	Chronicity is certain in a patient with chronic liver disease
- ⁴	+ ⁴	Chronic hepatitis C with false negative serology	Occasionally, immunocompromised, hemodialysis, agammaglobulinemic or mixed cryoglobulinemic patients have false- negative serology results
+ ⁸	? ⁸	False positive Ab	If absence of HCV-RNA on at least twice at 6 months apart
- ⁹	+ ⁹	Acute hepatitis C (window phase)	Will be confirmed by subsequent seroconversion
+ ¹⁰	+ ¹⁰	Acute hepatitis C	HCV IgG avidity index and serial measurements of the HCV-Ab IgM titres may help

propylene tubes, followed by incubation, washing, then amplification with an isothermal TMA process [70,71]. Detection of the amplified product is based on hybrid protection and the dual kinetic assays [72,73], using HCV probe reagent (chemiluminescent oligonucleotide probe) which produces a chemiluminescent signal that is read as relative light units (RLU) [70,71]. Qualitative HCV TMA has a sensitivity of 5.3 IU/mL [74] and is able to equally detect different HCV genotypes [61,75, 76]. The conversion of the qualitative HCV TMA assay to a quantitative assay has been described [77].

Genotyping of the extracted RNA

HCV genotyping was described by sequence-based PCR products analysis techniques, via genotype-specific antibodies [78], restriction endonuclease digestion [79], reverse hybridization against genotype-specific probes [80], direct sequencing [81,82], and annealing genotype-specific capture probes (AGSCP) [83]. Six major genotypes (1-6) have been characterized [84], with subtypes 1a to 1c, 2a to 2d, 3a to 3f, 4a to 4k, 5a, and 6a, and new (sub) types are continually being discovered [85-87]. Although in one study [88] 12 genotypes were predicted, but currently, the existence of six major HCV types is generally accepted. Debates are ongoing about whether the extra types are real distinct types or subtypes of types 3 and 6 [89]. Different methods for HCV-genotyping have been reported, including detailed sequence homology studies [82], restriction fragment length polymorphisms of amplified PCR products [83,90], PCR amplification with subtype-specific primers [91-95], and line probe assay (LiPA) based on RT-PCR followed by hybridization with specific probes (for each genotype) immobilized

(as parallel lines) on nitrocellulose strips [96]. Genotyping and subtyping for HCV are clinically important in the understanding of HCV disease and therapy options [81,97-100].

III-Cooperation between HCV-Ab and HCV-RNA results in the diagnosis of HCV infection

Modes of cooperation between HCV-Ab and HCV-RNA results in the diagnosis of HCV infection can be classified into five categories: False negative (inquiry?), unlikely acute hepatitis C, chronic hepatitis C, False + veAb and acute hepatitis C (Table 1). The following discussion is a stepwise description of such cooperation according to the superscripts shown in Table 1.

Superscript 1

In the diagnosis of HCV infection; HCV-Ab, HCV-RNA, and serum liver chemistry values are commonly used, but the latter is not necessary from a virological point of view because up to 40% of viraemic patients can have a normal aminotransferase value on a single determination [101].

Only one single, third generation ELISA determination is necessary for the diagnosis of HCV infection without the need for confirmation of positive or weakly positive ELISAs with RIBA because HCV-RNA detection by RT-PCR helps to resolve weakly positive or negative ELISA results when clinically compatible with hepatitis C [45].

Cooperation between HCV-Ab and HCV-RNA results does not apply in case of offspring born to HCV-infected mothers, where the diagnosis depends on HCV-RNA detection, because maternal IgG is passively transferred *in utero* and remains detectable for several months after

delivery [102-104], while HCV-RNA can be detected within a few days after delivery and then persists or is cleared spontaneously [104]. Cord blood may get contaminated by maternal blood and should never be used for HCV-RNA testing in the diagnosis of prenatal HCV infection [105,106].

Routine testing of pregnant women for HCV-Ab is not recommended, because the number of pregnant women who would be positive is expected to be low, in addition to the lack of measures to prevent perinatal transmission of HCV, and no licensed therapy or guidelines for the treatment of HCV-infected infants or children exist [105-108].

Indeed, in patients who have no indication for therapy or have a contra-indication to the use of antiviral drugs, neither HCV-Ab nor the HCV-RNA are useful because these virological tests have no prognostic value as their loads do not correlate with the severity of liver inflammation or fibrosis, nor with their progression [109]. Alternatively, in untreated patients, the severity of liver inflammation and fibrosis must be evaluated every 3 - 5 years by other means such as liver biopsies or non-invasive serological or ultrasound-based testing [110].

Superscript 2

HCV-Ab should be tested by a sensitive and specific technique such as 3rd and 4th generation ELISA [6,7, 23,24], which mainly detect HCV-Ab IgG class, since HCV-Ab IgM class can be detected in 50 - 93% of patients with acute HCV infection and in 50 - 70% of chronic cases [111].

Superscript 3

HCV-RNA should be tested by a sensitive technique detecting ≤ 50 HCV-RNA IU/mL [112,113], such as qualitative Cobas Amplicor RT-PCR which detects < 50 IU/mL and can be used to confirm viremia [68,69], or by transcription-mediated amplification (HCV TMA) [70,72].

HCV-RNA detection (qualitative and quantification) helps to diagnose chronic HCV infection, identify patients who need antiviral therapy, and monitor the virological responses to antiviral therapy [108,114]. Qualitative HCV-RNA assays detect viral genomes confirming an active infection and demonstrate its presence 4 - 6 weeks before antibody seroconversion takes place [115].

Most qualitative HCV-RNA tests are 10- to 100-fold more sensitive than quantitative HCV-RNA tests, and up to 1,000-fold more sensitive than the HCV core antigen assay, making them the method of choice for confirming active infection and for assessing viral clearance in response to therapy [14,116].

Superscript 4

It was estimated that the rate of HCV seroconversion among health care workers after a needle stick injury (parenteral) exposure is 0 - 7% [106].

However, after an accidental parenteral exposure to the virus, negative HCV-RNA results should be retested a few weeks later, because HCV-RNA is detectable in serum within 1 - 3 weeks after accidental parenteral exposure to the virus and weeks before the onset of ALT elevations or the appearance of HCV-Ab [117,118]. Keeping in mind that occasionally immunocompromised patients, patients undergoing hemodialysis, and patients with agammaglobulinemia and with mixed cryoglobulinemia have false-negative serology results (detectable HCV replication in the absence of HCV-Ab), these patients may require HCV-RNA testing for diagnosis [119,122].

Superscript 5

Acute hepatitis C is very unlikely if both HCV-Ab and HCV-RNA are absent [109]; thus, patients with symptoms of acute hepatitis of uncertain origin that are negative HCV-Ab should undergo qualitative HCV-RNA testing [123]. However, negative HCV-Ab and negative serum HCV-RNA but positive liver HCV-RNA have been described in "occult HCV infection" in which the patients have abnormal liver function tests of unknown etiology since the patients lack markers of HBV infection and lack clinical or biochemical evidence of autoimmunity, genetic or metabolic disorders, alcohol intake or drug toxicity [124]. Most patients (70%) with intrahepatic HCV-RNA also have HCV-RNA in their peripheral blood mononuclear cells (PBMC) [124,125].

Superscript 6

Acute hepatitis C is also unlikely if HCV-Ab are present without HCV-RNA [109]. However, about 30% of HCV-Ab positive with HCV-RNA negative have been found to have significant liver disease on biopsy [126, 127]; thus, one should consider other possibilities for the negative HCV-RNA result such as "healthy" HCV-carriers, infection resolving, RNA transient drop, occult HCV-infection, and technical inhibition of PCR. Healthy HCV-carriers are those patients with positive HCV-Ab but negative for serum HCV-RNA and normal ALT levels, and considered to have encountered and cleared HCV at some time in the past [109]. It has been estimated that in 15 - 25% of HCV-infected patients, the infection resolves and the patients recover from hepatitis C [118,128].

A temporary (transient) drop in the level of viremia below the detection limit should be taken into consideration [118], which may occur due to partial control of viral replication (by the immune response) before replication escapes and chronic infection is established [129,130]. These patients should however be retested after a few weeks because HCV-RNA can be transient undetectable [129].

Patients, who are HCV-Ab positive with undetectable serum viral RNA and normal ALT levels, should be verified if they could have HCV-RNA in their liver and PBMC, which is another form of "Occult HCV infection" (see above, superscript 5) [125,131,132].

It is also important to rule out the possibility of technical inhibition of PCR due to the presence of heparin in the collected samples, which can be solved using an internal control incorporated in each test tube from the extraction step [133]. In addition to ruling out the possibility of losing a little of the RNA due to improper storing or as a result of repeated thawing and freezing [134] and in spite of other contrary results, the RNA level remained stable during repeated thawing and freezing for about 8 cycles [135].

Superscript 7

When an individual is found to be HCV-Ab positive during blood donation or screening of at-risk populations, then HCV-RNA should be tested since the PCR positive result, in such circumstances, will confirm chronic infection [119,136], keeping in mind that the persistence of HCV-RNA for more than 6 months defines chronic HCV infection [101], and that a continuing hepatitis disease without improvement for at least six months defines chronic hepatitis [137]. Chronicity is also certain when patients have clinical or biological signs of chronic liver disease [45,138].

Chronic infection is often not symptomatic [139] until evidence of liver failure becomes clinically apparent, in which the rate of progression to cirrhosis is usually slow, with 20 or more years elapsing between infection and the development of serious complications [140-142].

Superscript 8

If HCV-RNA is negative and HCV-Abs are positive, false positive EIA may occur, the exact prevalence of which is unknown and which cannot be differentiated from the pattern of patients who have recovered from a past HCV infection [109]. False positive EIA was seen in patients with autoimmune diseases [16,143], in neonates born to mothers with chronic HCV infection since passively transferred maternal HCV-Abs may be detected in the children of HCV-infected mothers for up to 1 year [106,144], a low viral load below the detection limit of PCR, inhibition of PCR due to the presence of heparin in the collected samples [133], and improper storing and/or repeated thawing and freezing which may lead to little loss of the RNA [134], despite other reports to the contrary, the RNA level remains stable during repeated thawing and freezing for about 8 cycles [135]. There is a possibility of a negative result by HCV-RNA testing during the convalescent period (the progression from the acute to the chronic phase) since the patient may lose HCV-RNA and remain negative in this period [130].

Superscript 9

The serological window phase, characterized by detectable HCV-RNA in the absence of HCV-Ab, has been estimated to be 60 days on average [145]. Some patients (50 - 70%) with acute infection who develop symptoms, will have detectable antibodies at that time, but 90%

will have measurable antibodies after 3 months [118]. Thus, using a PCR assay, viremia can be detected within a few days of exposure to the virus and many weeks before elevation of transaminases and antibody titre [61-63].

The presence of HCV-RNA in the absence of HCV-Abs is strongly indicative of acute HCV infection, which will be confirmed by seroconversion (i.e., the appearance of HCV-Abs) a few days to weeks later [109].

Superscript 10

Acutely infected patients can also have both HCV-RNA and HCV-Abs at the time of diagnosis [109]. It is difficult, in this case, to distinguish acute hepatitis C from an acute exacerbation of chronic hepatitis C or an acute hepatitis of another cause in a patient with chronic hepatitis C [109].

The incubation period for newly acquired (acute) HCV infection ranges from two weeks to six months, with an average incubation period of 6 - 10 weeks [146-149]. The course of acute hepatitis C is variable, although elevations in serum ALT levels (often in a fluctuating pattern) are its most characteristic feature [137]. Normalization of ALT levels might occur and suggests full recovery, but this is frequently followed by ALT elevations that indicate progression to chronic disease [127, 151]. However, 50 - 70% of patients with acute infection who develop symptoms will have detectable antibodies at that time, but 90% will have measurable antibodies after 3 months [118].

Tests are not available to distinguish acute from chronic HCV infection [137]. However, the HCV-Ab IgG avidity index within the first 8 days following the onset of clinical symptoms may be useful in identifying actual acute HCV infection cases [36].

HCV-Ab IgM were reported in 50 - 93% of patients with acute hepatitis C and 50 - 70% of patients with chronic hepatitis C [151-153]. Therefore, HCV-Ab IgM cannot be used as a reliable marker of acute HCV infection [140]. However, the serial measurements of the HCV-Ab IgM titres based on at least three determinations from the fifth to the 15th day from the onset of the symptoms may identify patients with acute hepatitis C [153].

About 80% of acute HCV infections are asymptomatic [141] or have mild symptoms (fatigue, nausea, anorexia, vague abdominal discomfort, and vomiting) progressing to jaundice in about 25% of patients, less frequently than hepatitis B [140,141].

Most (70 - 90%) of infected people fail to clear the virus during the acute phase and become chronic carriers [127,133,134], that progress at variable rates (15 - 20%) to cirrhosis [137] and may (1 - 4%) develop to hepatocellular carcinoma [153]. About 5% of infected persons may die from the consequences of long term infection (liver cancer or cirrhosis) [128]. HCV-associated end-stage liver disease is the leading indication for liver transplantation in American adults [150,154].

Declaration of Interest:

Author declared no conflict of interests.

References:

1. Forman MS, Valsamakis A. Hepatitis C virus. In: Versalovic J, Carrol KC, Funke G, Jorgensen JH, Landry ML, Warrock DW, editors. *Murray's Manual of Clinical Microbiology*. 10th ed. Washington: American Society of Microbiology Press 2011; pp. 143755.
2. Choo QL, Kuo G, Weiner AJ, Overby LR, Bradley D, Hought M. Isolation of a cDNA clone derived from a blood-borne non-A, non-B viral hepatitis genome. *Science* 1989;244:359-62 (PMID: 2523562).
3. Alter MJ, Hadler SC, Judson FN, et al. Risk factors for acute non-A, non-B hepatitis in the United States and association with hepatitis C virus infection. *JAMA* 1990;264:2231-5 (PMID: 217 0702).
4. Alter H. Discovery of non-A, non-B hepatitis and identification of its etiology. *Am J Med* 1999;107(Suppl. 6B):16S-20S (PMID: 10653450).
5. Kuo G, Choo Q L, Alter H J, et al. An assay for circulating antibodies to a major etiologic virus of human non-A, non-B hepatitis. *Science* 1989;244:362-64 (PMID: 2496467).
6. Kao JH, Lai MY, Hwang YT, et al. Chronic hepatitis without anti-hepatitis C antibodies by second-generation assay clinicopathologic study and demonstration of the usefulness of third-generation assay. *Dig Dis Sci* 1996;41:161-5 (PMID: 8565750).
7. [No authors listed]. ANTI-HCV (4th Generation). Fortress Diagnostics Limited. UK. Revision No.10 MAY/11 Accessed November 22, 2016 http://www.artekinmedikal.com.tr/upload/Pencere2/HCV_4th_Gen_96T_New.pdf
8. Maertens G, Stuyver L. Genotypes and genetic variation of hepatitis C virus. In: Harrison TJ, Zuckerman AJ, eds. *The Molecular Medicine of Viral Hepatitis*. England: Wiley 1997;183-233.
9. Gretch DR. Diagnostic tests for hepatitis C. *Hepatology* 1997;26 (3 Suppl. 1):43S-47S (PMID: 9305663).
10. Van der Poel C, Reesink H, Schaasberg W, et al. Infectivity of blood seropositive for hepatitis C virus antibodies. *Lancet* 1990; 335:558-60 (PMID: 1968574).
11. Tremolada F, Casarin C, Tagger A, et al. Antibody to hepatitis C virus in post-transfusion hepatitis. *Ann Intern Med* 1991;14:277-81 (PMID: 1846277).
12. Fusconi M, Lenzi M, Ballardini G, et al. Anti-HCV testing in autoimmune hepatitis and primary biliary cirrhosis. *Lancet* 1990; 336:823 (PMID: 1976190).
13. Theilmann L, Blazek M, Goeser T, Gmelin K, Kommerell B, Fiehn W. False-positive anti-HCV tests in rheumatoid arthritis. *Lancet* 1990;335:1346 (PMID: 1971406).
14. Aceti A, Taliani G, DeBac C, Sebastiani A. Anti-HCV false positivity in malaria. *Lancet* 1990;336:1442-3 (PMID: 1978891).
15. Boudart D, Lucas J-C, Muller J-Y, Le Carrer D, Planchon B, Harousseau JL. False-positive hepatitis C virus antibody tests in paraproteinemia. *Lancet* 1990;336:63 (PMID: 1973253).
16. McFarlane IG, Smith HM, Johnson PJ, Bray GP, Vergani D, Williams R. Hepatitis C virus antibodies in chronic active hepatitis: pathogenic factor or false-positive results? *Lancet* 1990;335:754-7 (PMID: 1969512).
17. Schruppf E, Elgjo K, Fausa O, Haukenes G, Kvale D, Rollag H. The significance of anti-hepatitis C virus antibodies measured in chronic liver disease. *Scand J Gastroenterol* 1990;25:1169-74 (PMID: 2177219).
18. Ikeda Y, Toda G, Hashimoto N, Kurokawa K. Antibody to superoxide dismutase, autoimmune hepatitis, and antibody tests for hepatitis C virus. *Lancet* 1990;335:1345-6 (PMID: 1971405).
19. Mattsson L, Gutierrez R, Dawson GH, Lesniewski RR, Mushahwar LK, Weiland O. Antibodies to recombinant and synthetic peptides derived from the hepatitis C virus genome in long-term-studied patients with post-transfusion hepatitis C. *Scand J Gastroenterol* 1991;26:1257-62 (PMID: 1722348).
20. Bakir TM, Ramia S. Importance of second generation and confirmatory testing in diagnosing hepatitis C virus (HCV) infection in patients with non-A, non-B (NANB) hepatitis. *Biomed Res (India)* 1994;5:61-6.
21. Bresters D, Cuypers HT, Reesink HW, et al. Enhanced sensitivity of a second generation ELISA for antibody to hepatitis C virus. *Vox Sang* 1992;62:213-7 (PMID: 1379394).
22. Barrera J, Francis M B, Ercilla G, et al. Improved detection of anti-HCV in post-transfusion hepatitis by a third-generation ELISA. *Vox Sang* 1995;68:15-8 (PMID: 7536987).
23. Lavanchy D, Mayerat D, Morel B, et al. Evaluation of a third generation assay for detection of anti-hepatitis C (HCV) antibodies and comparison with presence of HCV-RNA in blood donors reactive to c 100-3 antigen. *J Clin Microbiol* 1994;32:2272-5 (PMID: 7529253).
24. Vrieling H, Zaaijar HL, Reesink HW, van der Poel CL, Cuypers HTM, Leile PN. Sensitivity and specificity of three third generation anti-HCV ELISAs. *Vox Sang* 1995;69:14-7 (PMID: 7483 486).
25. Garcia FB, Pereira Gde A, Martins PR, Moraes-Souza H. Epidemiological profile of hepatitis C in blood donors at the Uberaba Regional Blood Center. *Rev Soc Bras Med Trop* 2009;42(1):1-4 (PMID: 19287926).
26. Smith BD, Morgan RL, Beckett GA, et al. Centers for Disease Control and Prevention. Recommendations for the identification of chronic hepatitis C virus infection among persons born during 1945-1965. *MMWR Recomm Rep* 2012 Aug 17;61:1-32 (PMID: 22895429).
27. Garcia J. Hepatitis C: USPSTF recommends all baby boomers be screened. *Medscape Medical News from WebMD*. Accessed November 22, 2016. <http://www.medscape.com/viewarticle/806836>
28. Moyer VA, U.S. Preventive Services Task Force. Screening for hepatitis C virus infection in adults: U.S. Preventive Services Task Force recommendation statement. *Ann Intern Med* 2013 Sep 3;159(5):349-57 (PMID: 23798026).
29. Ngo-Metzger Q, Ward JW, Valdiserri RO. Expanded hepatitis C virus screening recommendations promote opportunities for care and cure. *Ann Intern Med* 2013 Sep 3;159(5):364-5 (PMID: 237 97155).
30. [No authors listed]. WHO: Simple/Rapid tests. Accessed November 22, 2016. http://www.who.int/diagnostics_laboratory/faq/simple_rapid_tests/en

31. [No authors listed]. CDC: Rapid Diagnostic Tests: How They Work. Accessed November 22, 2016. http://www.cdc.gov/malaria/malaria_worldwide/reduction/dx_rdt.html
32. [No authors listed]. US Food and Drug Administration. FDA approves rapid test for antibodies to hepatitis C virus. FDA News Release. Accessed November 22, 2016. <http://www.fda.gov/NewsEvents/Newsroom/PressAnnouncement/s/ucm217318.htm>
33. Lowry F. Baby boomer Hep C screening practical in emergency department. Medscape Medical News from WebMD. 2014 Mar 12; Accessed November 22, 2016. <http://www.medscape.com/viewarticle/822310>
34. Galbraith JW, Morgan J, Rodgers J, et al. High yield and feasibility of baby boomer birth cohort HCV screening in two urban, academic emergency departments [abstract 59]. Presented at: International Conference on Viral Hepatitis (ICVH); March 17-18, 2014; New York, NY Accessed November 22, 2016. http://www.iapac.org/icvh/presentations/ICVH2014_OA59.pdf
35. Coppola N, Pisapia R, Martini S, et al. Anti-HCV IgG Avidity Index in acute hepatitis C. Clinical Microbiology and Infectious 2006, Volume 12, Supplement 4 – P774, Accessed March 17, 2017. http://www.blackwellpublishing.com/eccmid16/PDFs/clm_1428.pdf
36. Coppola N, Pisapia R, Marrocco C, et al. Anti-HCV IgG avidity index in acute hepatitis C. J Clin Virol 2007;40:110-5 (PMID: 17720621).
37. van der Poel CL, Cuyper HT, Reesink HW, et al. Confirmation of hepatitis C virus infection by new four-antigen recombinant immunoblot assay. Lancet 1991;337:317-9 (PMID: 1671231).
38. Dow BC, Coote I, Munro H, et al. Confirmation of hepatitis C virus antibody in blood donors. J Med Virol 1993;41:215-20 (PMID: 7505306).
39. Skidmore S. Recombinant immunoblot assay for hepatitis C antibody. Lancet 1990;335:1346 (PMID: 1971407).
40. Lott JA, Nolte FS, Gretch R, Koff RS, Seeff LB. Laboratory guidelines for screening, diagnosis, and monitoring hepatic injury. In: Dufour DR, editor. Laboratory medicine practice guidelines. Washington (DC): National Academy of Clinical Biochemistry; 2000. p. 21-3.
41. Pawlotsky JM, Fleury A, Choukroun V, et al. Significance of highly positive c22-3 "indeterminate" second-generation hepatitis C virus (HCV) recombinant immunoblot assay (RIBA) and resolution by third-generation HCV RIBA. J Clin Microbiol 1994;32(5):1357-9 (PMID: 7519631).
42. Buffet, C, Charmaux N, Laurent-Puig P, et al. Enhanced detection of antibodies to hepatitis C virus by use of a third-generation recombinant immunoblot assay. J Med Virol 1994;43:259-61 (PMID: 7523581).
43. Uyttendaele S, Claeys H, Mertens W, Verhaert H, Vermynen C. Evaluation of third-generation screening and confirmatory assays for HCV antibodies. Vox Sang 1994;66:122-9 (PMID: 7514324).
44. Zaaier HL, Vrieling H, van Exel-Oehlers PJ, Cuyper HT, Lelie PN. Confirmation of hepatitis C infection: a comparison of five immunoblot assays. Transfusion 1994;34:603-7 (PMID: 7519796).
45. Pawlotsky JM, Lonjon I, Hezode C, et al. What strategy should be used for diagnosis of hepatitis C virus infection in clinical laboratories? Hepatology 1998;27(6):1700-1702. Jun;27(6):1700-2 (PMID: 9620345).
46. Kwon JA, Lee H, Lee KN, et al. High diagnostic accuracy of antigen microarray for sensitive detection of hepatitis C virus infection. Clin Chem 2008;54:424-8 (PMID: 18223133).
47. Kim S, Kim Y, Kim P, et al. Improved sensitivity and physical properties of sol-gel protein chips using large-scale material screening and selection. Anal Chem 2006;78(21):7392-6 (PMID: 17073404).
48. Bouvier-Alias M, Patel K, Dahari H, et al. Clinical utility of total HCV core antigen quantification: a new indirect marker of HCV replication. Hepatology 2002 Jul;36(1):211-8 (PMID: 12085367).
49. Morota K, Fujinami R, Kinukawa H, et al. A new sensitive and automated chemiluminescent microparticle immunoassay for quantitative determination of hepatitis C virus core antigen. J Virol Methods 2009;157:8-14 (PMID: 19135481).
50. Kesli R, Polat H, Terzi Y, Kurtoglu MG, Uyar Y. Comparison of a newly developed automated and quantitative hepatitis C virus (HCV) core antigen test with the HCV RNA assay for clinical usefulness in confirming Anti-HCV results. J Clin Microbiol 2011;49(12):4089-93 (PMID: 21940466).
51. Mullis K B, Faloona F A, Scharf S, Saiki R K, Horn G, Erlich H A. Specific enzymatic amplification of DNA *in vitro*: the polymerase chain reaction. Cold Spring Harb Sympo Quan Biol 1986;51 Pt 1:263-73 (PMID: 3472723).
52. Mullis KB, Faloona FA. Specific synthesis of DNA *in vitro* via a polymerase-catalyzed chain reaction. Methods Enzymol 1987; 155:335-50 (PMID: 3431465).
53. Mullis KB. The Unusual Origin of the Polymerase Chain Reaction. Sci Am 1990;262(4):56-61, 64-5 (PMID: 2315679).
54. Saiki RK, Scharf S, Faloona F, et al. Enzymatic amplification of beta-globin genomic sequences and restriction site analysis for diagnosis of sickle cell anemia. Science 1985;230(4732):1350-4 (PMID: 2999980).
55. Dharmaraj S. The Basics: RT-PCR. Thermo Fisher Scientific. Accessed November 22, 2016. <https://www.thermofisher.com/sa/en/home/references/ambion-tech-support/rtpcr-analysis/general-articles/rt-pcr-the-basics.html>
56. Tang YW, Persing DH. Molecular detection and identification of microorganisms, p. 215-44. In P. R. Murray, E. J. Baron, M. A. Tenover, F. C. Tenover, and R. H. Tenover, ed., Manual of Clinical Microbiology 1999, 7th ed. ASM Press, Washington, D.C.
57. Blair GE, Blair Zajdel ME: Biochem Educ 20:87-90, 1992. In Murray PR, Rosenthal KS, Pfaller MA: Medical Microbiology, 6th ed. Philadelphia, Mosby 2009
58. Vogelstein B, Gillespie D. Preparative and analytical purification of DNA from agarose. Proc Natl Acad Sci USA 1979; 76:1203-6 (PMID: 768406).
59. Stull D, Pisano JM. Purely RNA: New innovations enhance the quality, speed, and efficiency of RNA isolation techniques. Scientist 2001;15(22):29-31. <http://www.the-scientist.com/?articles.view/articleNo/13691/title/purely-rna/>
60. Pawlotsky JM. Diagnostic tests for hepatitis C. J Hepatol 1993; 31:71-9 (PMID: 10622564).
61. Simizu YK, Weiner AJ, Rosenblatt J, et al. Early events in hepatitis C virus infection of chimpanzees. Proc Natl Acad Sci USA 1990;87:6441-4 (PMID: 2117282).

62. Garson JA, Tuke PW, Makris M, et al. Demonstration of viraemia patterns in haemophiliacs treated with hepatitis-C-virus-contaminated factor VIII concentrates. *Lancet* 1990;336:1022-5 (PMID: 1977017).
63. Garson JA, Ring C, Tuke R, Tedder RS. Enhanced detection by PCR of hepatitis C virus RNA. *Lancet* 1990;336:878-9 (PMID: 1976900).
64. Davis GL. Interferon treatment of chronic hepatitis C. *Am J Med* 1994;96:41S-46S (PMID: 7509123).
65. Gretch DR, dela Rosa C, Carithers RL Jr, Willson RA, Williams B, Corey L. Assessment of hepatitis C viremia using molecular amplification technologies: correlations and clinical implications. *Ann Intern Med* 1995;123:381-4 (PMID: 7542853).
66. Thaler MM, Park CK, Landers DV, et al. Vertical transmission of hepatitis C virus. *Lancet* 1991;338:17-8. PMID: 1676085).
67. Lin HH, Hsu HY, Chang MH, et al. Low prevalence of hepatitis C virus and infrequent perinatal or spouse infections in pregnant women in Taiwan. *J Med Virol* 1991;35(4):237-40 (PMID: 1724982).
68. Lee SC, Antony A, Lee N, et al. Improved version 2.0 qualitative and quantitative AMPLICOR reverse transcription-PCR tests for hepatitis C virus RNA: Calibration to international units, enhanced genotype reactivity, and performance characteristics. *J Clin Microbiol* 2000;38(11):4171-9 (PMID: 11060086).
69. [No authors listed]. SmartCycler. Accessed November 22, 2016 http://www.cepheid.com/administrator/components/com_product_catalog/library-files/9843d40ce3fb6eef0e1e4521b18b12ba-SmartCycler-Brochure.pdf
70. Hendricks DA, Friesenhahn M, Tanimoto L, Goergen B, Dodge D, Comanor L. Multicenter evaluation of the VERSANT HCV RNA qualitative assay for detection of Hepatitis C Virus RNA. *J Clin Microbiol* 2003;41(2):651-6. PMID: 12574262).
71. Krajden M, Ziermann R, Khan A, et al. Qualitative detection of hepatitis C virus RNA: comparison of analytical sensitivity, clinical performance, and workflow of the Cobas Amplicor HCV test version 2.0 and the HCV RNA transcription-mediated amplification qualitative assay. *J Clin Microbiol* 2002;40(8):2903-7 (PMID: 12149349).
72. Sarrazin C, Teuber G, Kokka R, Rabenau H, Zeuzem S. Detection of residual hepatitis C virus RNA by transcription-mediated amplification in patients with complete virologic response according to polymerase chain reaction-based assays. *Hepatology* 2000;32(4 Pt 1):818-23 (PMID: 11003628).
73. Sawyer L, Leung K, Friesenhahn M, Duey D, McMorrow M, Eguchi B. Clinical laboratory evaluation of a new sensitive and specific assay for qualitative detection of hepatitis C virus in clinical specimens. *J Hepatol* 2000;32(Suppl. 2):116. Accessed November 22, 2016. [http://www.journal-of-hepatology.eu/article/S0168-8278\(00\)80780-1/pdf](http://www.journal-of-hepatology.eu/article/S0168-8278(00)80780-1/pdf)
74. Gorrin GM, Friesenhahn P, Lin M, et al. Performance evaluation of the VERSANT HCV RNA qualitative assay by using transcription-mediated amplification. *J Clin Microbiol* 2003;41:310-7 (PMID: 12517866).
75. Ross RS, Viazov SO, Hoffmann S, Roggendorf M. Performance characteristics of a transcription-mediated nucleic acid amplification assay for qualitative detection of hepatitis C virus RNA. *J Clin Lab Anal* 2001;15(6):308-13 (PMID: 11793430).
76. Comanor L, Elkin C, Leung K, et al. Successful HCV genotyping of previously failed and low viral load specimens using an HCV RNA qualitative assay based on transcription-mediated amplification in conjunction with the line probe assay. *J Clin Virol* 2003 Sep;28(1):14-26 (PMID: 12927747).
77. Covill S, Mathis J, Hodge P, Cass M, Bott M, Bodrug S. Quantitative HCV TMA Assay with Variable Dynamic Range. Abstracts of the Interscience Conference on Antimicrobial Agents and Chemotherapy 41:294. (Chicago, IL 2001 Dec 16-19; Abstract no. H-755.). accessed November 22, 2016. <https://eurekama.com/research/035/595/035595963.php>
78. Ansaldi F, Torre F, Bruzzone BM, Picciotto A, Crovari P, Icardi G. Evaluation of a new hepatitis C virus sequencing assay as a routine method for genotyping. *J Med Virol* 2001;63(1):17-21 (PMID: 11130882).
79. Arens M. Clinically relevant sequence-based genotyping of HBV, HCV, CMV, and HIV. *J Clin Virol* 2001;22(1):11-29 (PMID: 11418349).
80. Le Pogam S, Dubois F, Christen R, Raby C, Cavicchini A, Goudeau A. Comparison of DNA enzyme immunoassay and line probe assays (Inno-LiPA HCV I and II) for hepatitis C virus genotyping. *J Clin Microbiol* 1998;36(5):1461-3 (PMID: 9574733).
81. Mellor J, Walsh EA, Prescott LE, et al. Survey of type 6 group variants of hepatitis C virus in Southeast Asia by using a core-based. genotyping assay. *J Clin Microbiol* 1996;34(2):417-423 (PMID: 8789027).
82. Zhang ZX, Yun ZB, Chen M, Sönnnerborg A, Sällberg M. Evaluation of a multiple peptide assay for typing of antibodies to the hepatitis C virus: relation to genomic typing by the polymerase chain reaction. *J Med Virol* 1995;45(1):50-5 (PMID: 7536231).
83. Rho J, Ryu JS, Hur W, et al. Hepatitis C virus (HCV) genotyping by annealing reverse transcription-PCR products with genotype-specific capture probes. *J Microbiol* 2008;46(1):81-7 (PMID: 18337698).
84. Nouroz F, Shaheen S, Mujtaba G, Noreen S. An overview on hepatitis C virus genotypes and its control. *Egyptian Journal of Medical Human Genetics* 2015;16 (4):291-8. <http://www.sciencedirect.com/science/article/pii/S1110863015000506>
85. Reyes GR, Baroudy BM. Molecular biology of non-A, non-B hepatitis agents: hepatitis C and hepatitis E viruses. *Adv Virus Res* 1991;40:57-102 (PMID: 1659777).
86. Germer JJ, Zein NN. Advances in the molecular diagnosis of hepatitis C and their clinical implications. *Mayo Clin Proc* 2001;76(9):911-20 (PMID: 11560302).
87. Robertson B, Myers G, Howard C, et al. Classification, nomenclature, and database development for hepatitis C virus (HCV) and related viruses: proposals for standardization. International Committee on Virus Taxonomy. *Arch Virol* 1998;143(12):2493-503 (PMID: 9930205).
88. Bukh J, Purcell RH, Miller RH. At least 12 genotypes of hepatitis C virus predicted by sequence analysis of the putative E1 gene of isolates collected worldwide. *Proc Natl Acad Sci USA* 1993;90(17):8234-8 (PMID: 8396266).
89. Stuyver L, Wuseur A, Van Arnhem W, et al. Hepatitis C virus genotyping by means of 5'-UR/core line probe assays and molecular analysis of untypeable samples. *Virus Res* 1995;38:137-57 (PMID: 8578855).

90. Cha T, Brall E, Irvine B, et al. At least five related, but distinct, hepatitis C viral genotypes exist. *Proc Natl Acad Sci USA*. 1992; 89:7144-8 (PMID: 1323128).
91. McOmish F, Chan SW, Dow BC, et al. Detection of three types of hepatitis C virus in blood donors: investigation of type specific differences in serologic reactivity and rate of alanine aminotransferase abnormalities. *Transfusion* 1993;33(1):7-13 (PMID: 7678709).
92. Nakao T, Enomoto N, Takada N, Takada A, Date T. Typing of hepatitis C virus genomes by restriction fragment length polymorphism. *J Gen Virol* 1991;72 (Pt 9):2105-12 (PMID: 1716652).
93. Andonov A, Chaudhary RK. Genotyping of Canadian hepatitis C virus isolates by PCR. *J Clin Microbiol*. 1994;32(8):2031-4 (PMID: 7989565).
94. Chayama K, Tsubota A, Arase Y, et al. Genotypic subtyping of hepatitis C virus. *J Gastroenterol Hepatol*. 1993;8(2):150-6 (PMID: 8386022).
95. Okamoto H, Sugiyama Y, Okada S, et al. Typing hepatitis C virus by polymerase chain reaction with type-specific primers: application to clinical surveys and tracing infectious sources. *J Gen Virol* 1992;73(Pt 3):673-9 (PMID: 1312125).
96. Schröter M, Zöllner B, Schäfer P, et al. Genotyping of hepatitis C virus types 1, 2, 3, and 4 by a one-step LightCycler method using three different pairs of hybridization probes. *J Clin Microbiol* 2002;40(6):2046-50 (PMID: 12037062).
97. Hoofnagle JH. A Step Forward in Therapy for Hepatitis C. *N Engl J Med* 2009;360(18):1899-901 (PMID: 19403908).
98. Wilck MB, Hamel MB, Baden LR. Management of Incidental Hepatitis C Virus Infection - Polling Results. *N Engl J Med* 2009; 360(23):e30 (PMID: 19494211).
99. Tamori A, Enomoto M, Kawada N. Recent Advances in Antiviral Therapy for Chronic Hepatitis C. *Mediators Inflamm* 2016;6841628 (PMID: 27022210).
100. Chevaliez S. Virological tools to diagnose and monitor hepatitis C virus infection. *Clin Microbiol Infect*. 2011 Feb;17(2):116-21 (PMID: 21054664).
101. Chevaliez S, Pawlotsky JM. How to use virological tools for optimal management of chronic hepatitis C. *Liver Int* 2009;29(s1):9-14 (PMID: 19207960).
102. Pembrey L, Newell ML, Tovo PA; EPHN Collaborators. The management of HCV infected pregnant women and their children. European Pediatric HCV Network. *J Hepatol* 2005;43:515-25 (PMID: 16144064).
103. Resti M, Bortolotti F, Vajro P, Maggiore G; Committee of Hepatology of the Italian Society of Pediatric Gastroenterology and Hepatology. Guidelines for the screening and follow-up of infants born to anti-HCV positive mothers. *Dig Liver Dis* 2003;35(7):453-7 (PMID: 12870728).
104. Ketzinel-Gilad M, Colodner SL, Hadary R, Granot E, Shouval D, Galun E. Transient transmission of hepatitis C virus from mothers to newborns. *Eur J Clin Microbiol Infect Dis* 2000;19(4):267-74 (PMID: 10834815).
105. American Academy of Pediatrics, Committee on Infectious Diseases. 1997 Red book: report of the committee on infectious diseases. 24th ed. Elk Village, IL.: American Academy of Pediatrics 1997:263.
106. [No authors listed]. Recommendations for prevention and control of hepatitis C virus (HCV) infection and HCV-related chronic disease. Centers for Disease Control and Prevention. *MMWR Recomm Rep* 1998 Oct 16;47(RR-19):1-39 (PMID: 9790221).
107. Dienstag JL. Sexual and perinatal transmission of hepatitis C. *Hepatology* 1997;26(3 Suppl 1):66S-70S (PMID: 9305667).
108. Mast EE, Alter MJ. Hepatitis C. *Semin Pediatr Infect Dis* 1997; 8:17-22. Accessed November 22, 2016. <http://www.sciencedirect.com/science/article/pii/S1045187097800052>
109. Chevaliez S, Pawlotsky JM. Hepatitis C Virus Serologic and Virologic Tests and Clinical Diagnosis of HCV-Related Liver Disease. *Int J Med Sci* 2006;3:35-40 (PMID: 16614740).
110. NIH Consensus Statement on Management of Hepatitis C: 2002. *NIH Consens State Sci Statements* 2002;19(3):1-46 (PMID: 14768714).
111. Pawlotsky JM. Diagnostic tests for hepatitis C. *J Hepatol*. 1999; 31 Suppl 1:71-9 (PMID: 10622564).
112. Pawlotsky JM. Use and interpretation of virological tests for hepatitis C. *Hepatology* 2002;36 Suppl 1: S65-73 (PMID: 12407578).
113. Brink ATP, Snijders PJF and Meijer CJKM. Target Amplification -Bases Techniques, in: *Nucleic Acid Testing for Human Disease*. Attila L (Ed) 2006. by CRC Press, Taylor & Francis Group. Pages 3-18.
114. [No authors listed]. Global surveillance and control of hepatitis C. Report of a WHO consultation organized in collaboration with the Viral Hepatitis Prevention board, Antwerp Belgium. *J Viral Hepat* 1999;6:35-47 (PMID: 10847128).
115. Carithers RL Jr, Marquardt A, Gretch DR. Diagnostic testing for hepatitis C. *Semin Liver Dis* 2000;20:159-71 (PMID: 10946421).
116. Krajden M. Hepatitis C virus diagnosis and testing. *Can J Public Health* 2000;91 Suppl 1:S34-S39 (PMID: 11059131).
117. Young KK, Archer JJ, Yokosuka O, Omata M, Resnick RM. Detection of hepatitis C virus RNA by a combined reverse transcription PCR assay: comparison with nested amplification and antibody testing. *J Clin Microbiol* 1995;33(3):654-7 (PMID: 7751372).
118. [No authors listed]. National Institutes of Health Consensus Development Conference Panel statement: management of hepatitis C. *Hepatology* 1997;26(3 Suppl 1):2S-10S21 (PMID: 9305656).
119. Fabrizi F, Poordad FF, Martin P. Hepatitis C infection and the patient with end-stage renal disease. *Hepatology* 2002;36(1):3-10 (PMID:12085342).
120. Lauer GM, Walker BD. Hepatitis C virus infection. *N Engl J Med* 2001;345(1):41-52 (PMID: 11439948).
121. Thio CL, Nolt KR, Astemborski J, Vlahov D, Nelson KE, Thomas DL. Screening for hepatitis C virus in human immunodeficiency virus-infected individuals. *J Clin Microbiol* 2000;38(2):575-7 (PMID: 10655348).
122. Lok AS, Chien D, Choo QL, et al. Antibody response to core, envelope and nonstructural hepatitis C virus antigens: comparison of immunocompetent and immunosuppressed patients. *Hepatology* 1993;18(3):497-502 (PMID: 7689528).
123. [No authors listed]. EASL International Consensus Conference on Hepatitis C. Paris, 26-28, February 1999, Consensus Statement. European Association for the Study of the Liver. *J Hepatol* 1999;30(5):956-61 (PMID: 10365827).

Hepatitis C: an Overview of Various Laboratory Assays with their Mode of Diagnostic Cooperation

124. Castillo I, Pardo M, Bartolome J, et al. Occult hepatitis C virus infection in patients in whom the etiology of persistently abnormal results of liver-function tests is unknown. *J Infect Dis* 2004; 189(1):7-14 (PMID: 14702147).
125. Carreño V. Occult hepatitis C virus infection: A new form of hepatitis C. *World J Gastroenterol* 2006 November 21;12(43) 6922-5 (PMID: 17109511).
126. McGuinness P, Bishop GA, Lien A, Wiley B, Parsons C, McCaughan GW. Detection of serum hepatitis C virus RNA in HCV antibody-seropositive volunteer blood donors. *Hepatology* 1993; 18(3):485-90 (PMID: 7689526).
127. Bresters D, Zaaier HL, Cuypers HIM, et al. Recombinant immunoblot assay reaction patterns and hepatitis C virus RNA in blood donors and non-A, non-B hepatitis patients. *Transfusion* 1993;33 (8):634-63 (PMID: 7688158).
128. [No authors listed]. Recommendations for Prevention and Control of Hepatitis C Virus (HCV) Infection and HCV-Related Chronic Disease: US Department of Health and Human Services, Centers for Disease Control and Prevention. *MMWR* 1998;47(RR19):1-39. Accessed November 22, 2016. <https://www.cdc.gov/mmwr/preview/mmwrhtml/00055154.htm>
129. Lavallette D, Morice Y, Germanidis G, et al. Human serum facilitates hepatitis C virus infection, and neutralizing responses inversely correlate with viral replication kinetics at the acute phase of hepatitis C virus infection. *J Virol* 2005;79(10):6023-34 (PMID: 15857988).
130. Al-Quaiz MN, Madani TA, Karwai MA. The natural history of hepatitis c virus infection. *Saudi Med J* 2003;24 suppl 2:S67-S70 (PMID: 12897903).
131. Carreno V, Pardo M, Lopez-Alcorocho JM, Rodriguez-Inigo E, Bartolome J, Castillo I. Detection of hepatitis C virus (HCV) RNA in the liver of healthy, anti-HCV antibody-positive, serum HCV RNA-negative patients with normal alanine aminotransferase levels. *J Infect Dis* 2006;194(1):53-60 (PMID: 16741882).
- a. Falcón V, Acosta-Rivero N, Shibayama M, et al. Evidences of hepatitis C virus replication in hepatocytes and peripheral blood mononuclear cells from patients negative for viral RNA in serum. *Am J Infect Dis* 2005;1:34-42. Full text accessed November 22, 2016. <http://thescpub.com/PDF/ajidsp.2005.34.42.pdf>
132. Poljak M, Seme K, Koren S. Evaluation of the automated COBAS AMPLICOR hepatitis C virus PCR system. *J Clin Microbiol* 1997;35 (11):2983-4 (PMID: 9350774).
133. Krajden M, Minor JM, Rifkin O, Comanor L. Effect of multiple freeze-thaw cycles on hepatitis B virus DNA and hepatitis C virus RNA quantification as measured with branched-DNA technology. *J Clin Microbiol* 1999;37(6):1683-6 (PMID: 10325307).
134. Kessler HH, Stelzl HE, Raggam RB, et al. Effects of storage and type of blood collection tubes on hepatitis C virus level in whole blood samples. *J Clin Microbiol* 2001;39(5):1788-90 (PMID: 11325991).
135. Pawlotsky JM, Bouvier-Alias M, Hezode C, Darthuy F, Rerr J, Dhumeaux D. Standardization of hepatitis C virus RNA quantification. *Hepatology* 2000;32(3):654-9 (PMID: 10960464).
136. [No authors listed]. WHO: Hepatitis C. Accessed November 22, 2016. <http://www.who.int/mediacentre/factsheets/fs164/en/>
137. [No authors listed]. EASL International Consensus Conference on Hepatitis C. Paris, 26-28, February 1999, Consensus Statement. European Association for the Study of the Liver. *J Hepatol* 1999;30:956-61 (PMID: 10365827).
138. Mast EE, Alter MJ, Margolis HS. Strategies to prevent and control hepatitis B and C virus infections: a global perspective. *Vaccine* 1999;17:1730-3 (PMID: 10194830).
139. Hsu HH, Greenberg HB. Hepatitis C. In: Hoeprich PD, Jordan MC, Ronald AR, eds. *Infectious Diseases. A treatise of infectious processes*, 5th ed. JB Lippincott Co., Philadelphia 1994:820-825.
140. [No authors listed]. EASL International Consensus Conference on Hepatitis C. Paris, 26-27 February 1999. Consensus Statement. *J Hepato* 1999;31(Suppl 1):3-8 (PMID:10622553).
141. Lemon SM, Brown EA. Hepatitis C virus. In: Mandell GL, Bennett JE, Dolin R, eds. *Principle and Practice of Infectious Disease*, Fourth. New York, Churchill Livingstone 1995:1474-86.
142. Czaja AJ, Magrin S, Fabiano C, et al. Hepatitis C virus infection as a determinant of behavior in type I autoimmune hepatitis. *Dig Dis Sci* 1995;40(1):33-40 (PMID: 7529673).
143. Zanetti AR, Tanzi E, Newell MI. Mother-to-infant transmission of hepatitis C virus. *J Hepatol* 1999;31 Suppl 1:96-100 (PMID: 10622569).
144. Busch MP, Glynn SA, Stramer SL, et al. A new strategy for estimating risks of transfusion-transmitted viral infections based on rates of detection of recently infected donors. *Transfusion* 2005; 45(2):254-64 (PMID: 15660836).
145. Koretz RL, Brezina M, Polito AJ, et al. Non-A, non-B posttransfusion hepatitis: comparing C and non-C hepatitis. *Hepatology* 1993;17(3):361-5 (PMID: 8383087).
146. Marranconi F, Mecenero V, Pellizzer GP, et al. HCV infection after accidental needlestick injury in health-care workers. *Infection* 1992;20(2):111 (PMID: 1582682).
147. Seeff LB. Hepatitis C from a needlestick injury. *Ann Intern Med* 1991;115(5):411 (PMID: 1907442).
148. Hsu HH, Greenberg HB. Hepatitis C. In: Hoeprich PD, Jordan MC, Ronald AR, eds. *Infectious Diseases. A treatise of infectious processes*, 5th ed. JB Lippincott Co., Philadelphia, 1994:820-5.
149. Houghton M. Hepatitis C viruses. In: Fields BN, Knipe DM, Howley PM, eds. *Fields Virology*, 3rd ed. Philadelphia, Lippincott - Raven, 1996:1035-58.
150. Hellstrom UB, Sylvan SP, Decker RH, Sonnerborg A. Immunoglobulin M reactivity towards the immunologically active region sp75 of the core protein of hepatitis C virus(HCV) in chronic HCV infection. *J Med Virol* 1993;39(4):325-32 (PMID: 8388029).
151. Negro F, Troonen H, Michel G, et al. Lack of monomeric IgM anti-hepatitis C virus (HCV) core antibodies in patients with chronic HCV infection. *J Virol Methods* 1996;60(2):179-82 (PMID: 8844624).
152. Sagnelli E, Coppola N, Marrocco C, et al. Diagnosis of hepatitis C virus related acute hepatitis by serial determination of IgM anti-HCV titres. *J Hepatol* 2005;42(5):646-51 (PMID: 15826712).
153. Kim WR, Brown RS Jr, Terrault NA, El-Serag H. Burden of liver disease in the United States: summary of a workshop. *Hepatology* 2002;36(1):227-42 (PMID: 12085369).