

Lancet Seminar – Hepatitis C

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Search strategy and selection criteria

We searched MEDLINE and PUBMED databases using the search terms HCV and hepatitis C virus as well as epidemiology, clinical manifestation, virology, diagnosis, biopsy, treatment, drugs, immunology or vaccines. We selected publications mostly from the past 5 years, but did not exclude commonly referenced and highly regarded older publications.

Abstract

Hepatitis C virus (HCV) infection is a major global health problem that continues to grow with an estimated 170 million people infected. The consequences of chronic infection can include cirrhosis, end-stage liver disease and hepatocellular carcinoma. Due to shared routes of transmission, coinfection with HIV is a significant problem and individuals infected with both viruses have poorer outcomes. There is no effective vaccine though HCV is a potentially curable infection. For many years, the standard of care has been subcutaneous interferon-alpha and oral ribavirin for between 24 and 72 weeks of treatment. This treatment results in a sustained virological response in only about 50% of individuals and is complicated by significant adverse events. In recent years, advances in HCV cell culture has allowed a greater understanding of HCV virology and this has paved the way for the development of many new directly acting antiviral drugs that target key components of virus replication such as protease and polymerase inhibitors. We are now at a point of improved and simplified treatments for HCV that may be administered as oral regimes of short duration and with far greater tolerability than regimens of old. The remaining hurdles may be access to appropriate care and cost of treatment as the epidemic continues to grow.

Introduction

First discovered in 1989, hepatitis C virus (HCV) is a major global health problem affecting over 170 million people.¹ The problem continues to grow; globally the number of people who are seropositive for anti-HCV antibodies is estimated to have increased from 2.3% to 2.8% between 1990 and 2005.² Central and East Asia along with North Africa and the Middle East are estimated to have the highest prevalence (>3.5%) with moderate prevalence in Eastern and Western Europe (1.5%-3.5%).² The majority of subjects who become acutely infected (~80-85%) fail to clear the virus and progress to chronic infection. This figure may be higher in subjects who are coinfecting with HIV and lower in women and children.^{3,4} The consequences of chronic infection can be cirrhosis, portal hypertension, liver decompensation and the development of hepatocellular carcinoma (HCC) with HCV infection ultimately causing approximately 350,000 deaths per year.⁵ In regions of high endemicity, chronic viral hepatitis usually accounts for >50% of HCC and cirrhosis.⁶ Globally, 27% of cases of cirrhosis can be attributed to HCV and 25% of HCC is attributable to HCV infection. Aside from fatal consequences, individuals chronically infected with HCV have a decreased quality of life compared to the general population.⁷

For many years, treatment for chronic HCV has been inadequate with success rates of treatment estimated at around 50%, depending on genotype. The standard-of-care until 2011 was a combination of pegylated interferon-alpha (PEG-IFN), administered subcutaneously and ribavirin (RBV) taken orally. This combination could lead to a sustained virological response (SVR) and, based on long-term follow up results, SVR means cure and chronic HCV has become the first chronic viral infection to be cured by medical therapy. However, such treatment is associated with significant adverse events and furthermore, is poorly tolerated and less efficacious in subjects with advanced disease who are most at need.⁸ The introduction of direct acting antiviral drugs (DAAs), with two protease inhibitor drugs licensed in 2011, has improved treatment responses rates and heralded a new era of HCV treatment (figure 1).^{9,10} A pipeline of new DAAs are in various stages of pre-clinical and clinical development creating great optimism for the future of managing chronic HCV

infection with simple, short, interferon-free, all oral regimes.¹¹

Virology

HCV is a positive single-stranded RNA virus in the Flaviridae family, within the genus Hepacivirus. The positive-sense RNA genome is 9600 nucleotides in length from which a single HCV polyprotein of 3011 amino acids is translated. This is subsequently cleaved by cellular and viral proteases into three structural proteins (core, E1 and E2) and seven non-structural (NS) proteins (p7, NS2, NS3, NS4A, NS4B, NS5A and NS5B).¹² For some time since its discovery in 1989 there were no close relatives identified in other animal species. However, in recent years there has been an explosion of data on the subject of non-primate hepaciviruses (NPHVs). Initially a related virus – canine hepacivirus - was identified as the cause of respiratory infection in dogs,¹³ a discovery closely followed by detection of a related NPHV in horses.¹⁴ Related viral sequences have now also been identified in rodents and bats, extending the diversity of the family enormously.¹⁵

HCV infections within human populations also show extreme genetic diversity. This is partly explained by the long evolutionary association between the virus and humans, likely several centuries if not longer.¹⁶ There are currently 7 established genotypes, although most work has focused on genotypes 1-6, which although showing global spread, have different geographical origins. Genotypes 1, 2, 4 and 5 are found as endemic infections in Africa, whilst genotypes 3 and 6 have evolved in Asia.¹⁷ During the last century, a number of medical interventions such as schistosomiasis eradication campaigns in Egypt have amplified specific strains to epidemic proportions and subsequently many of these have spread internationally.¹⁸ In the UK, genotype 3a is co-dominant with genotype 1, a feature which has implications for both vaccines and therapy (see below).

HCV infection also shows enormous diversity within infected hosts, existing in blood as a 'swarm' of related sequences or quasispecies. This diversity is a consequence of the error-prone viral polymerase and the high viral replication rate and allows for rapid adaptation to host antibody responses, cellular immune responses and many antiviral drugs. Recent analyses suggest within-host viral diversity may be even greater than had previously been estimated, likely due to hidden populations within the liver.¹⁹

For many years, it was not possible to grow HCV in tissue culture, but this is now possible following the discovery of a specific strain of HCV genotype 2 which was able to infect specific hepatoma cell lines and allow a full replication cycle.²⁰ Culture of HCV has allowed study of entry, revealing a complex set of interactions with surface receptors, including CD81, SRB-1 (a scavenger receptor) and two tight junction proteins Occludin-1 and Claudin.²¹ Similarly these models have allowed critical insights into viral replication, and host-virus interactions.²² Tissue culture-derived virus has also recently facilitated imaging studies for analysis of mature virions by electron microscopy, revealing an unusually irregular structure.²³ Most importantly, the ability to analyse HCV replication in tissue culture, coupled with structural analysis of key proteins such as the NS3 protease and NS5b polymerase has driven the development of novel specific DAAs.²⁴⁻²⁶

Immunology

Immune responses to HCV play a critical role in determining the outcome of acute disease, and also play a more complex role in determining long-term disease progression. Acute responses to HCV include both innate and adaptive arms, with clear evidence for both. Genetic evidence has implicated polymorphisms in the region of the IL28B (IFNL3) gene to strongly impact on spontaneous resolution of infection.²⁷ IFNL3 is a lambda interferon, with sustained antiviral activity similar to that of interferon, but with a more restricted receptor

distribution. Whether the polymorphisms identified influence the regulation of IFNL3 itself or impact on a nearby gene, termed IFNL4, remains to be defined.²⁸ Similarly, genetic associations between genes in the KIR locus and acute resolution of infection point to a role for NK cell responses in viral control.²⁹

Adaptive responses mediated by CD8+ and CD4+ T cells are also involved in acute host defence and there are strong associations reported with HLA Class II alleles in many studies, including a well-powered genome-wide association study.²⁷ HLA Class I associations have been identified in single-source outbreaks, such as that seen in Irish women infected through a batch of Anti-D preparation. CD8+ T cell responses to peptides bound by HLA-A3 and HLA-B27 have been proposed as the mediators of such protection – these target regions of the virus where single mutations that allow immune escape also compromise viral fitness.³⁰ Studies from infected chimpanzees indicate that both CD4+ and CD8+ T cell responses are required for full protection.³¹ These studies have prompted the development of T-cell based preventive vaccines: a regimen based on two recombinant vectors expressing HCV nonstructural genes – a novel adenovirus construct followed by a modified vaccinia Ankara (MVA) construct) - is now in Phase II trials in the USA.³²

B cell responses to HCV leading to the generation of neutralising antibodies (NAbs), have also been intensely studied, although this is a complex area due to the extreme heterogeneity of key regions of the envelope genes – the major target for NAbs.³³ The variability of regions such as the hypervariable regions (HVRs) of HCV E2 within-hosts is a consequence of antibody-driven immune selection. Nevertheless, broadly cross-reactive Nabs have been described,³⁴ and further work to characterise these, especially in the context of a recently described HCV E2 crystal structure may lay the foundations for antibody-based vaccines.³⁵

Epidemiology

HCV has long been recognized as a parenterally-transmitted cause of viral hepatitis.³⁶ Transmission via blood transfusion was a major route before the implementation of universal screening of blood in the developed world though this route remains a problem elsewhere.³⁷ Where blood products and transplanted organs are now safe, intravenous drug use (IDU) has become the major route of HCV transmission.³⁸ Epidemiological studies highlight transmission sources other than the sharing of contaminated needles, including sharing of other drug paraphernalia such as foil and spoons.³⁹ These findings are of vital importance in targeted prevention interventions.

Mother to child transmission (MTCT) of HCV has been less intensively studied than other chronic viral infections. The natural history of HCV in pregnancy and in infected infants born to these mothers is poorly understood. Consequently effective methods for prevention of HCV vertical transmission have not been developed. MTCT rates are estimated to be 2 to 8% in HCV mono-infected mothers but may be two to four times higher in those coinfected with HIV.^{40, 41} There are currently no data from randomized controlled trials to support recommending caesarian section in this setting.⁴² The efficiency with which HCV is sexually transmitted has been controversial. However, in monogamous heterosexual couples where one partner has chronic HCV, the rate of transmission to a discordant partner is extremely low. Furthermore, HCV transmission is not associated with any particular sexual practice allowing for more unambiguous messages to be given to these couples.⁴³

In the context of HIV infection, the epidemiology of HCV infection has changed dramatically in the last decade. Since 2000 there has been an ongoing epidemic of acute HCV infections

in HIV positive men who have sex with men (MSM) that shows no sign of abating.⁴⁴ There is evidence that transmission is per mucosal rather than parenteral and is associated with certain sexual practices (fisting and group sex) and intranasal and intrarectal drug use.⁴⁵ Sophisticated molecular epidemiological and phylogenetic studies in several European countries have identified several transmission clusters within MSM networks where sexual risks are implicated.^{44, 46, 47} Of particular concern in this group is the rate of reinfection after successful treatment or spontaneous clearance. As many as 25% of individuals treated for HCV will become re-infected within 2 years and these data highlight the need for effective sexual health education and preventative interventions targeted at this group.⁴⁸ It further demonstrates that natural immunity is not adequate protection against a subsequent infection, even with the same genotype, and highlights the challenge in developing a prophylactic vaccine.

HIV/HCV Coinfection

HIV and HCV share routes of transmission and therefore coinfection with both viruses is a common problem, affecting an estimated 20-30% of the world's 34 million HIV-infected individuals.⁴⁹ HCV-related liver disease has become a leading cause of morbidity and death in HIV patients in the era of HAART.⁴⁸ The impact of HIV on the natural history of HCV is well-established and significant, affecting every aspect of the disease.⁴⁹ Chronic HCV infection is more likely in HIV patients and this is associated with higher HCV viral loads.⁴⁹ Coinfected patients demonstrate a faster progression to cirrhosis and end-stage liver disease though this may be attenuated by early HAART where available,⁵⁰⁻⁵² and historically responded less well to interferon and ribavirin treatment regimens.⁵³ However, attempting treatment remains important as achieving a sustained virological response dramatically reduces the incidence of liver-related morbidity and mortality in this population.⁵⁴

Initiation of HAART in coinfecting patients is associated with a higher risk of hepatotoxicity. However, it is felt that this is outweighed by the potential benefits of immune restoration that might lessen disease progression and thus, initiation of HAART is generally recommended early in these patients.⁵⁵ Both HIV and HCV are infections associated with disorders of multiple systems. Alongside, the deleterious effect of HIV on HCV-related liver pathology, patients who have HIV-HCV coinfection have higher rates of HIV-related kidney disease,⁵⁶ more global neurocognitive dysfunction,⁵⁷ and the prevalence of cardiovascular disease and bone disease is higher.⁵⁸⁻⁶⁰

Diagnosis

The diagnosis of HCV relies on the detection of antibody to the virus and nucleic acid amplification tests to detect HCV RNA. Antibody tests were first approved by the US Food and Drug Administration (FDA) in 1990 and have evolved considerably since this time.⁶¹ HCV RNA is detected early in infection (~2 weeks) and will be followed by antibody seroconversion days to weeks later (~6 weeks), though the development of detectable antibody can be delayed or not occur at all in the immunocompromised such as HIV positive.⁶² Anti-HCV antibodies are detected by an enzyme immunoassay (EIA) or chemiluminescence immunoassay (CIA) and these are the first screening tests.⁶¹ The presence of anti-HCV antibody in the absence of detectable RNA indicates old spontaneously resolved or treated infection and, in the presence of RNA, indicates a current HCV infection. Therefore, all antibody tests that are repeatedly positive, should be assessed by a sensitive test for HCV RNA. In acute infection, RNA may be present without antibody and in immunocompromised patients (such as those with HIV) with abnormal liver function, tests to detect HCV RNA may be a more appropriate diagnostic test.

The different HCV genotypes respond with different success rates to treatment with interferon and ribavirin.^{9,10} Additionally, the genotype determines the duration of therapy necessary to achieve SVR.⁶³ Therefore, accurate genotyping of chronic infections is vital. Early genotyping was performed using assays that either determined the specificity of the antibody present or by hybridisation of amplified viral RNA from highly conserved regions of the virus. However, it is becoming apparent that these hybridisation assays may not be specific enough and that more accurate results may be obtained from molecular sequence data for example, sequencing both C/E1 and NS5B regions (or in future whole genomes using next generation sequencing approaches) and establishing phylogenetic relatedness.^{64,}

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On treatment, the measurement and quantification of HCV RNA at predetermined timepoints may allow for the shortening of treatment regimes (response-guided therapy) or determine that it is futile to continue with treatment due to a lack of response.⁶⁶ Assays have become more sensitive and the lower limits of detection and quantification of HCV RNA has decreased with inevitable consequences for response guided therapy and stopping rules. Resistance testing may become more important than previously. It is therefore important that treating clinicians remain familiar with updates in diagnostic technology and liaise closely with laboratory colleagues.⁶⁶

Acute HCV

Studies on the natural history of early HCV infection have been limited due to the asymptomatic nature of the majority of acute HCV infections.⁶⁷ Our understanding is improving following the study of acute cases in patients with HIV and in animal models of infections conducted in chimpanzees. 10 to 14 weeks after infection, an acute hepatitis with a corresponding increase in liver transaminase enzymes occurs.⁶⁸ A characteristic that has been observed in both human and chimpanzee infection is an early peak in HCV viral RNA load followed by a dip.⁶⁷ It is conservatively estimated that 15 to 20% of subjects will clear acute infection and in these, this downward trajectory in HCV viral load continues whereas chronic infection is associated with a recrudescence of viraemia.^{3, 67, 69} Multiple factors have been shown to be associated with spontaneous clearance of HCV infection including gender, IL28B polymorphisms, ALT levels or presence of jaundice, rate of decline in HCV-RNA, and blood IP-10 levels.⁷⁰ Attempts have been made to establish scoring systems that might reliably discriminate those patients with high potential for spontaneous clearance from those that should be treated early.⁷⁰ There is a balance between treating too early in patients who may go on to clear infection and delaying treatment which will result in reduced treatment efficacy. A positive HCV RNA 12 weeks into the course of acute HCV infection has been established as a helpful transition from acute into chronic infection and might help guide early treatment decisions.⁷¹

In subjects not clearing the virus, early treatment has been shown to be more effective than delayed treatment in subjects both with and without HIV. In the context of HIV infection, treatment within 3 to 6 months is recommended, where possible, by European guidelines and this is associated with improved treatment outcomes.⁷² The optimal treatment regime for acute HCV infection with PEG-IFN in HIV has not been established but 24 or 48 weeks of PEG-IFN with RBV is recommended depending on the early viral kinetics.⁷¹ Treatment success rates ranging from 65 to 85% have been reported in this setting.^{73, 74} By contrast, in acute HCV in HIV-negative individuals, interferon-alpha, standard or pegylated, alone for up to 24 weeks may be sufficient to effect a cure in up to 98% of subjects.^{75, 76}

Natural history

HCV infection has a propensity to cause chronic hepatitis which may lead to cirrhosis, decompensated cirrhosis and HCC. The onset and accumulation of hepatic fibrosis is clinically silent in the early stages of disease, and it therefore remains difficult to accurately identify progression of the disease to cirrhosis in patient.^{77,78} Annual rates of progression of hepatic fibrosis from minimal disease to cirrhosis have been modelled and estimated. The prevalence of biopsy-proven cirrhosis after 20 years of infection has varied between 7% (in retrospective studies) to 18% (in clinical referred settings). The risk of cirrhosis is increased in individuals abusing alcohol, in those who acquire the disease at an older age, by concomitant obesity, in men, and in immunosuppressed HIV positive patients or in recurrent HCV following liver transplantation.^{79,80}

Patients with minimal fibrosis have a low risk of developing complications of liver disease over the ensuing two decades. Patients with bridging fibrosis or cirrhosis, conversely have a higher risk. It may be necessary to repeat liver biopsies in patients to determine progression. Alternatively and more practically, non-invasive blood tests, fibroelastography and hepatic imaging can be used to identify patients with advanced fibrosis to gauge indications for immediate or deferred treatment.⁸¹ Extrahepatic manifestations of HCV such as cryoglobulinaemia, or HCV-associated splenic lymphoma are also indications for antiviral therapy. Treatment will reduce infectivity and transmission in individuals using intravenous drugs.

Treatment

The primary goal of treatment for chronic HCV is cure, and thus prevention of progression of the disease. A sustained virological response (SVR), i.e HCV RNA < 15 iu/ml 12-24 weeks after completion of antiviral therapy is associated with an improvement in both all-cause and liver-related mortality from HCV.^{82,83} PEG-IFN and RBV have previously been the mainstay of treatment for all genotypes of HCV, given for up to 48 weeks, but are being superseded by the advent of direct acting antiviral agents (DAAs) (Fig. 2).

Indications for treatment

Patients with cirrhosis are at immediate threat of complications of liver disease and ideally treatment should be prioritized for patients with advanced disease. Unfortunately current IFN-based treatment response rates are lower in patients with cirrhosis, and thus suboptimal, albeit they are improved by the addition of first generation protease inhibitors. Furthermore the potential risks of adverse events are greater. Recent data suggest that response rates in patients with cirrhosis can be improved by DAA regimens without IFN. Treatment may also given to patients to prevent the development of advanced fibrosis, and cirrhosis, or for extrahepatic symptoms, and to prevent transmission of the infection. The high costs of current regimens may require stratification of patients for treatment.

Current treatment genotype 1

First generation protease inhibitors (PI) improve response rates in patients with genotype 1 infection. Telaprevir and boceprevir are inhibitors of the NS3/4a HCV protease. SVR rates are improved in both treatment-naïve and treatment-experienced patients treated with a combination of telaprevir or boceprevir plus PEG-IFN and RBV.^{9,10} Approximately 50- 60% of PI recipients also qualify for a reduced (six months) duration of treatment, based on achieving a rapid virological response. Lower response rates have been observed in patients with cirrhosis and those with a prior null response to PEG-IFN and RBV, i.e. those with a < 2 log₁₀ IU/ml decline in HCV RNA by treatment week 12. Also, patients with cirrhosis require a longer duration (48 weeks) of PEG-IFN and RBV.⁸⁴ Treatment with first generation PIs can

be complex and cause considerable adverse events. The safety profile of prolonged IFN and first generation DAA treatment in patients with cirrhosis is poor.⁸⁵

A single nucleotide polymorphism upstream of the IL28B gene influences response to PEG-IFN and RBV. Higher response rates have been reported in patients inheriting the IL28B rs12979860 CC genotype.⁸⁶ Several drug-drug interactions can occur.⁸⁷ Resistance associated viral variants, with substitutions located in the catalytic site of the NS3 protease, have been described following telaprevir and boceprevir treatment. Stopping rules to avoid the acquisition of more complex mutations are recommended.⁸⁸⁻⁹⁰

The most common side effects of telaprevir are anaemia, pruritis, nausea, diarrhoea and anorectal discomfort. About 4% of patients develop a severe dermatitis, necessitating cessation of treatment. Drug Reaction with Eosinophilia and Systemic Symptoms (DRESS syndrome) or Stevens Johnson syndrome are rare, but reported. Boceprevir causes dysgeusia and anaemia. The use of PIs is constrained in patients with decompensated cirrhosis and post-transplant HCV infection because of the risk of severe adverse effects of the IFN backbone.⁹¹

Interferon-sparing regimens with new agents for genotype 1

Treatment for genotype 1 HCV is evolving rapidly. Key viral replication targets have been identified, namely the HCV protease, NS5a protein and the NS5b RNA polymerase. In 2014, several potent antiviral inhibitors with once daily dosing and, because of improved potency, a shorter duration of treatment with PEG-IFN and RBV have been licenced. These IFN-“sparing” regimens include the addition of simeprevir (second generation PI), daclatasvir (a NS5a inhibitor) and sofosbuvir (a uridine nucleotide prodrug NS5b polymerase inhibitor) in combination with PEG-IFN and RBV for 12 to 24 weeks.⁹²⁻⁹⁵ High response rates have been observed in naïve genotype 1 patients: up to 90%.

Simeprevir 150 mg daily for 12 weeks with PEG-IFN and RBV for 24 weeks resulted in SVR rates in treatment naïve genotype 1 patients of 80% and 81% in the QUEST-1 and QUEST-2 studies respectively.^{96, 97} A rapid virological response occurs in 85% of whom 91% subsequently achieve an SVR at 12 weeks. Lower response rates were observed in patients with F3-F4 fibrosis. A Q80K mutation detectable at baseline in the NS3 sequence impairs the response to simeprevir. Simeprevir with PEG-IFN and RBV resulted in SVR rates in 79% of prior relapsers compared to 36% of PEG-IFN retreated controls.⁹⁸

Sofosbuvir together with PEG-IFN and RBV has been given for 12 weeks in treatment-naïve patients with genotypes 1,4,5 and 6 in the ATOMIC and NEUTRINO studies. SVR rates of 89% (genotype 1) and 82% (genotype 4) were reported in ATOMIC and 89% and 100% for genotype 1 and 4 respectively in NEUTRINO.^{95, 99} Smaller numbers of patients with genotype 5 and 6 were treated. Prior non-responders were not included in these clinical trials and only a limited number of patients with cirrhosis were included.

In a dose finding study 332 patients were treated MK-5172 (100, 200, 400, or 800 mg) once daily for 12 weeks together with PEG IFN and RBV. A control group received boceprevir plus PEG IFN and ribavirin. SVR rates of 89% to 91% were observed in patients given MK-5172 versus 61% of controls. Transient increases in serum aminotransferase concentrations occurred in the MK-5172 groups given 400 and 800 mg, suggesting that 100 mg is the safe dose of the protease inhibitor for utilisation in interferon free regimens.¹⁰⁰

Interferon-free regimens for genotype 1

It is likely that IFN-sparing regimens will be relatively quickly displaced in 2015 by the introduction of IFN-free regimens with improved efficacy and tolerability. These regimens will comprise a protease inhibitor or a NS5a inhibitor plus a nucleoside NS5b inhibitor \pm RBV; a protease inhibitor plus a NS5a inhibitor plus a NS5b non nucleoside inhibitor \pm RBV; or a protease inhibitor plus a NS5a inhibitor \pm RBV.¹⁰¹ Updated guidelines have been published by the European Association for the Study of the Liver (EASL). Sofosbuvir can also be used in combination with RBV in patients with genotype 1 infection who are intolerant to IFN, particularly in non-cirrhotic patients, and up to 72% response rates can be achieved.¹⁰²

Both sofosbuvir and daclatasvir,¹⁰³ or sofosbuvir plus ledipasvir (without RBV) are highly effective regimens for genotype 1 naïve and prior IFN non-responders including telaprevir and boceprevir non-responder patients.¹⁰¹ SVR rates of 97% in treatment naïve patients, (ION-1),¹⁰⁴ 93% of treatment experienced patients (ION-2) treated with sofosbuvir and ledipasvir without ribavirin for 12 weeks and 94% of genotype 1 naïve patients treated for 8 weeks have been reported,¹⁰⁵ the first two studies have included 15-20% of patients with cirrhosis. Anaemia was rare in the ribavirin-free arms. Clinical resistance is extremely rare, although a S282T mutation in the replicon model confers resistance to sofosbuvir.¹⁰⁶ However, relapses, when they occur, occur with wild type HCV. Although the numbers of patients are relatively small, a subgroup analysis indicated a SVR rate of 22/22 patients treated for 24 weeks versus 86% treated for 12 weeks (19/22). Further studies are examining the efficacy of sofosbuvir in combination with either ledipasvir and a non-nucleoside HCV NS5B inhibitor, (GS 9669) 500 mg per day or a protease inhibitor (GS 9451) 80 mg per day in naïve non-cirrhotic patients and further shortening of treatment to as little as four or six weeks.¹⁰⁷ Phase 3 studies examining the efficacy of sofosbuvir and GS-5881- a next generation NS5a inhibitor are in progress based on the pangenotypic efficacy of this combination in preliminary studies.¹⁰⁸

Other demonstrably effective, short duration, oral combinations include ritonavir-boosted ABT-450 (a PI) (150 mg/100 mg) co-formulated together with ABT-267 25 mg once daily (an NS5a inhibitor) (ombitasvir) and ABT-333 250 mg twice daily (dasabuvir, a non-nucleoside NS5b polymerase inhibitor) with RBV (weight-based). (SAPPHIRE I). Ninety-six percent of patients responded.¹⁰⁹ Virological failure was thus rare, but in this and other multiple drug regimens, the emergence of NS3, NS5a and NS5b resistance-associated variants will require an appropriate salvage therapy. In the SAPPHIRE-2 study, 96% of non-cirrhotic prior non-responders to PEG-IFN and RBV, including 49% who were prior null responders achieved an SVR.¹¹⁰ Ninety-one and 98% of patients with compensated cirrhosis responded to the same regimen given for 12 or 24 weeks respectively.¹¹¹ Response rates were improved in 1a patients with a prior null response given for 24 weeks versus 12 weeks (92% versus 80%) These studies have also demonstrated that ribavirin need only be used if required, and is not advantageous in naïve non-cirrhotic 1b patients. In the (PEARL-III) study 419 patients with genotype 1b infection, and 305 patients with genotype 1a infection (PEARL-IV) were given 12 weeks of ABT-450/r-ombitasvir, dasabuvir (250 mg twice daily), and ribavirin or placebo for ribavirin. SVR rates in 1b infection were 99.5% with ribavirin and 99.0% without ribavirin; in patients with 1a the SVR rates were 97.0% and 90.2%, respectively.¹¹² A longer duration of treatment (24 weeks) with ribavirin is advisable for 1a patients with cirrhosis and a prior non response or other adverse factors. PEARL-II evaluated the efficacy of 12 weeks of treatment with the same regimen with or without RBV in non-cirrhotic pegIFN RBV treatment-experienced HCV genotype 1b patients. The response rates with or without RBV were 96% and 100%.¹¹³

Sofosbuvir 400 mg qd plus simeprevir 150 mg qd have been assessed in cirrhotic and non-cirrhotic naïve and prior non-responder patients; two cohorts have been studied; F0-F2 null-responders and F3-F4 naïve or null-responders (COSMOS).¹¹⁴ In the cohort of prior null-responders with early stage fibrosis, 92% of patients treated without RBV had an SVR. In cohort 2 (naïve and prior null-responders with metavir F3-F4 fibrosis), 96% of patients responded.¹¹⁴ There is no apparent need for RBV with this combination of a PI and NS5b polymerase inhibitor. Prior Q80K mutations in patients with subtype 1a had little impact on treatment. Anaemia and bilirubin elevations were more common in the RBV arms. The combination of daclatasvir 30 mg bd, asunaprevir 200 mg bd and BMS-791325 (a non-nucleoside NS5b polymerase inhibitor, 75 or 150 mg bd) for 12 weeks in treatment-naïve patients with or without cirrhosis,¹¹⁵ are similarly encouraging.¹¹⁶⁻¹¹⁸ Daclatasvir and asunaprevir (a PI) is effective in patients with subtype 1b infection (HALLMARK DUAL).¹¹⁹

The combination of MK5172 (a PI) 100 mg qd and MK8742, (a NS5a inhibitor) 20mg ± RBV in naïve or experienced genotype 1 patients with or without cirrhosis treated for 8-18 weeks gave high rates of response 90-97% (C-WORTHY).¹²⁰

Sofosbuvir can be utilised in patients with decompensated cirrhosis. The dose of sofosbuvir in patients with eGFR < 30 ml /minute/1.73m² is not yet established. No dose adjustment of daclatasvir is required for renal or hepatic impairment. The results obtained in phase 3 studies portend the advent of IFN-free regimens in 2014 for naïve and prior non-responder patients, including those with resistance to telaprevir and boceprevir and patients with cirrhosis. Refinements of treatment will be ascertained in further phase 3 trials. RBV is dispensable (in the “correct” DAA regimen). Ultrashort regimens of four to six weeks are being studied with several combinations of next generation protease inhibitors, NS5a inhibitors and polymerase inhibitors. Interestingly, appropriate IFN-free regimens may overcome the biological ineffectiveness of IFN in advanced liver disease.

These interferon free regimens are remarkably effective in patients with genotype 1 infection. Further analysis is required, but baseline factors including baseline RAVs, unfavourable IL28b genotype, viral load, subtype and cirrhosis can influence response with potent multiple DAA regimens. Treatment response rates could be reduced in prior null-responder patients with cirrhosis; longer durations of treatment could be required to optimise response rates in patients with more advanced liver disease. Other factors acting in concert such as baseline viral load or baseline NS5a mutations could affect response to NS5a inhibitors although their detection does not preclude a response. In a small percentage of patients multiply drug resistant viruses will be encountered after treatment failure or a relapse.¹²¹

Treatment for genotypes 2 to 6.

PEG-IFN and RBV are effective against genotypes 2-6. Forty-eight weeks of PEG-IFN and RBV are generally required for genotypes 4, 5 and 6, although rapid virological responders may be successfully treated for 24 weeks. Twenty-four weeks of PEG-IFN/RBV are given to patients with genotype 2 and 3. SVR rates are highest in patients with genotype 2 (85-90%). SVR rates of between 43 and 70% have been recorded in patients with genotype 4 treated with PEG-IFN and ribavirin. Genotype 4 patients with low baseline viral concentrations and a rapid virological response can be treated for 24 weeks. SVR rates of 60-85% occur in genotype 6.¹²² It has become apparent that patients with genotype 3 infection and cirrhosis have higher relapse and thus lower response rates.

Interferon-sparing regimens

Daclatasvir has been administered with PEG-IFN and RBV to naïve patients with genotype 2 or 3 infection. SVR rates of approximately 83% in genotype 2 and 70% in genotype 3 have been reported.¹²³ Similar encouraging results were reported in genotype 4 patients.¹²⁴ Simeprevir, a second wave protease inhibitor, is active against genotype 4, particularly in treatment-naïve and relapsed patients.¹²⁵ As noted above, sofosbuvir administered with 12 weeks of PEG-IFN and RBV is active against all genotypes. In the LONESTAR-2 study, sofosbuvir given in combination with PEG-IFN and RBV for 12 weeks to treatment-experienced genotype 2 or 3 patients resulted in SVR rates in 96% in genotype 2 and 83% in genotype 3 patients. Although the numbers were small, cirrhosis did not affect the response.¹²⁶ Presently, genotype 3 treatment-experienced patients with cirrhosis may require 12 weeks PEG-IFN and RBV plus sofosbuvir to attain the highest response rates.

Interferon-free regimens for genotype 2 and 3

A 12-week combination of sofosbuvir and ribavirin is highly efficacious (97%) in genotypes 2, but a longer treatment period of 24 weeks is required for patients with genotype 3. In the POSITRON and FUSION studies a lower proportion of patients with genotype 3 responded to 12 weeks of therapy, and in these genotype 3 patients, responses were lower among those with cirrhosis than among those without cirrhosis.¹²⁷ Response rates are similar in patients with genotype 2 irrespective of the presence of cirrhosis, but are reduced to 62% in experienced patients with genotype 3 and cirrhosis treated with sofosbuvir and ribavirin for 24 weeks.¹²⁸ However, 94% SVR rates occurred in naïve, non-cirrhotic genotype 3 patients treated for 24 weeks.¹²⁸ Virological failures are usually due to relapse with wild-type HCV, and discontinuations for drug-related adverse events have been rare. SVR rates of 94-100% were reported with the combination of daclatasvir plus sofosbuvir in treatment-naïve patients infected with genotypes 2 or 3 for 12 or 24 weeks, with or without ribavirin.¹⁰² Preliminary results of a 12 week combination of sofosbuvir and ledipasvir for patients with genotype 3 showed a lower SVR rate (16/25, 64%) versus 26/26 of those treated with sofosbuvir, ledispavir and RBV.¹²⁹ Sofosbuvir plus GS-5816 and other combinations including MK5172 plus MK874 (or next generation NS5a inhibitors) plus next generation polymerase inhibitors are being studied. SVR rates of > 90% have been reported genotype 4 patients with sofosbuvir and ribavirin treated for 24 weeks. By extrapolation the combination of sofosbuvir and simeprevir or daclatasvir should be active against genotype 4.¹³⁰ A 12-week dual regimen of ABT450/r + ombitasvir, ± RBV in treatment-naïve patients (treatment-experienced patients all received RBV) showed excellent response rates [ref Hezode C, et al. EASL 2014, London, O58](#)), as did all-oral therapy with daclatasvir plus asunaprevir + BMS-791325 for treatment-naïve patients with chronic HCV G4 infection.¹³¹

HIV/HCV coinfecting patients.

Telaprevir and boceprevir have both been assessed in chronic coinfecting patients.¹³²⁻¹³⁵ Although promising SVR rates have been obtained, drug-drug interactions can be problematic. Telaprevir can be given for 12 weeks in combination with PEG-IFN and RBV to treat acute genotype 1 co-infection.¹³⁶ Several promising IFN-sparing as well as IFN-free regimens have been tested in HIV/HCV coinfecting patients.¹³² These include the combination of simeprevir together with PEG IFN and RBV, sofosbuvir plus RBV, sofosbuvir plus ledipasvir, for 12 wks in HIV/HCV-coinfecting patients,¹³⁷ or MK5172 plus MK8742 with or without RBV in the C-WORTHY study.¹³⁸ In the PHOTON study, coinfecting genotype 1, 2 and 3 treatment-naïve patients were treated with sofosbuvir 400 mg daily and ribavirin for 24 weeks (genotype 1) and 12 weeks (genotype 2 and 3).¹³⁹ Multiple antiretroviral therapies were permitted as sofosbuvir is cleared by renal elimination. Seventy-six percent of 114 patients with genotype 1 (treated for 24 weeks), 88% of 26 patients with genotype 2

and 67% of 42 with genotype 3 achieved SVR12 (after treatment for 12 weeks in the latter groups). Amongst the prior non-responder patients, 92% with genotype 2 and 94% of 17 genotype 3 achieved SVR12 after 24 weeks of treatment. It appears fortunately that patients who are coinfecting with HIV and HCV have similar outcomes as those with HCV mono-infection.¹⁴⁰ However, appropriate dose modifications will be required in coinfecting patients. Efavirenz, etravirine and nevirapine are not recommended with daclatasvir, simeprevir or sofosbuvir. The dose of daclatasvir should be reduced to 30 mg from 60 mg if atazanavir/r is used but increased to 90 mg if dosed together with efavirenz, nevirapine or etravirine. Co-administration of atazanavir/r, lopinavir/r or darunavir/r with simeprevir is not recommended.

Patients with decompensated cirrhosis and pre- and post-liver transplant patients

Patients with decompensated cirrhosis are not candidates for interferon therapy. A preliminary report of the use of pre-transplant sofosbuvir and RBV for up to 48 weeks, stopping on day of transplant for patients transplanted for HCV and HCC (within Milan criteria), has resulted in a 64% post-transplant SVR rate. The duration of undetectable HCV RNA pre-transplant was the best predictor of response.¹⁴¹ Several controlled clinical trials of for example sofosbuvir and ledipasvir, sofosbuvir and daclatasvir and sofosbuvir and simeprevir are in progress to determine the safety and efficacy of interferon free DAA regimens in patients with advanced liver disease. Definitive guidance on the optimal duration of therapy in this group is critical; less tolerant of a relapse because of the risk of acute on chronic liver injury.

Telaprevir and boceprevir have been utilised to treat post-transplant recurrent HCV.¹⁴² Preliminary reports indicated that higher SVR rates than with PEG-IFN and RBV occur. However, toxicity, particularly anaemia and sepsis, and drug-drug interactions with the calcineurin inhibitors complicate treatment.¹⁴³ Discontinuation rates of up to 40 percent have been encountered. Fortunately, these treatments are likely to be displaced by better tolerated and more effective DAA therapies. Sofosbuvir and ribavirin have been used for the treatment of recurrent post-transplant hepatitis C (all genotypes). Virological response rates of 77% after 24 weeks of treatment have been reported.¹⁴⁴ These landmark treatments, which will be improved upon with time, greatly reduce the risk and complications of post-transplant HCV. Several pre and post transplant regimens are under investigation, including sofosbuvir plus daclatasvir, and sofosbuvir plus ledipasvir and sofosbuvir plus simeprevir ± RBV. ABT 450/r, ombitasvir and dasabuvir plus ribavirin give a 7 fold and 3 fold increase in tacrolimus and cyclosporin half life but the immunosuppressive therapy is manageable although requiring dosing of tacrolimus 0.5 to 1.0 mg at one to two week intervals. Excellent post-transplant SVR results were observed in a small group of genotype 1 patients without advanced fibrosis.¹⁴⁵

Conclusion

Improved, efficacious and simplified, interferon free, and for most, ribavirin free treatments for hepatitis C are now available. Detailed guidelines which will be updated at frequent intervals have been published for genotypes 1-6, and subcategories of patients. Simple all-oral regimes of short duration have now become a reality. Treatment could be expanded into groups for whom interferon was not tolerated. Emerging evidence that patients on stable opioid replacement therapy are good candidates for DAA regimens. Treatment algorithms may still be necessary with sofosbuvir and ledipasvir, or the AbbVie three DAA regimen, or sofosbuvir and daclatasvir for example to attain the highest chance of an SVR; (a slower primary response, may perhaps indicate a need to extend the duration of therapy

and may be necessary particularly in treatment experienced patients with advanced cirrhosis).¹⁴⁶

However the next generation of DAA treatments are costly drugs. Meeting the demand for therapy of a numerically common disease with these breakthrough therapies is concerning for policy makers because of the immediate budgetary impact. The cost may limit access, thus limiting societal benefit. Stratification and prioritisation of patients based on cost-effectiveness, stage of disease, and the potential gain from treatment, may be required. Prices may decrease as several effective drugs offering a high cure are licenced. However, the epidemic continues to grow. A major hurdle currently is the identification and appropriate referral of people in need of treatment and widespread delivery in primary care. Treatment will form part of the control of the disease; however, successful treatment of an infection has never led to its eradication. The search for an effective prophylactic vaccine must continue and advances in molecular vaccinology are paving the way for progress in this era.

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Figure 1. The evolution of the standard of care for HCV and improvements in sustained virological response rates.^{9, 10, 147, 148}

Figure 2. The treatment of HCV in 2015 (including approved or imminently approved protease, NS5b and NS5a inhibitors).

¥ protease inhibitor *NS5a inhibitor # NS5b inhibitor