



1 23 June 2016
2 EMEA/CHMP/EWP/30039/2008 Rev 1
3 Committee for Medicinal Products for Human Use (CHMP)
4

5 **Guideline on the clinical evaluation of direct acting**
6 **antivirals for the treatment of chronic hepatitis**
7 **Draft**

8

Draft agreed by Infectious Diseases Working Party (IDWP)	April 2016
Adopted by CHMP for release for consultation	23 June 2016
Start of public consultation	07 July 2016
End of consultation (deadline for comments)	31 December 2016

9
10

Comments should be provided using this [template](#). The completed comments form should be sent to IDWPsecretariat@ema.europa.eu

11
12

Keywords	<i>HCV, direct acting antivirals, Drug development, Guidance</i>
-----------------	---



13 **Guideline on the clinical evaluation of direct acting**
14 **antivirals for the treatment of chronic hepatitis**

15 **Table of contents**

16 **Executive summary 4**

17 **1. Introduction (background) 5**

18 **2. Scope..... 5**

19 **3. Legal basis and relevant guidelines 5**

20 **4. Pharmacodynamics and pharmacokinetics 7**

21 4.1. Nonclinical virology studies7

22 4.2. Clinical virology studies8

23 4.2.1. Viral drug resistance8

24 4.2.2. Determination of HCV genotype and subtype8

25 4.2.3. Determination of plasma HCV-RNA levels8

26 4.3. Clinical pharmacokinetics9

27 4.4. Drug-drug interactions9

28 **5. Assessment of efficacy 9**

29 5.1. General considerations for clinical trials.....9

30 5.2. Subject characteristics and the definition of patient populations 10

31 5.2.1. Viral genotypes 10

32 5.2.2. Host IL28B genotype 10

33 5.2.3. Treatment history 11

34 5.2.4. Assessment of liver fibrosis..... 11

35 5.3. Methods to evaluate efficacy 11

36 5.4. Dose finding studies 12

37 5.4.1. Monotherapy studies 12

38 5.4.2. Early combination dose ranging studies (phase 2a) 12

39 5.5. Phase IIb studies and confirmatory studies 12

40 5.5.1. Study populations..... 12

41 5.5.2. Selection of the study regimen 13

42 5.5.3. Add-on and substitution studies 13

43 5.5.4. Studies aiming at a shortened treatment duration 13

44 5.5.5. Fixed dose combinations..... 13

45 5.5.6. Follow-up after the primary endpoint 13

46 5.5.7. Combination of medicinal products and the demonstration of the contribution of each

47 component to regimen efficacy 14

48 5.5.8. The extrapolation of efficacy between viral genotypes 14

49 5.6. Studies in special patient populations..... 15

50 5.6.1. Treatment of patients with decompensated liver disease 15

51 5.6.2. Post-transplant treatment..... 16

52 5.6.3. HCV/HIV co-infected patients..... 16

53 5.6.4. Patients with prior DAA experience 16

54	5.6.5. Studies in paediatric patients	17
55	3.6.6 Studies in older patients	17
56	6. Safety aspects	18
57	7. Information in the Summary of the Product Characteristics	18
58	Definitions.....	19
59		

60 Executive summary

61 This draft guideline replaces the CHMP's *Guideline on the clinical evaluation of direct acting antiviral*
62 *agents intended for treatment of chronic hepatitis C (EMA/CHMP/EWP/30039/2008)*.

63 There have been considerable developments in the field of hepatitis C virus (HCV) therapy since the
64 adoption of *EMA/CHMP/EWP/30039/2008*. Since 2013 direct acting antivirals (DAAs) have been
65 approved for the treatment of chronic HCV infections within interferon-free combination regimens.
66 Therefore this revision of the prior guidance concerns the development of DAA-only regimens.

67 The mechanism of action of each new agent should be elucidated. In-vitro activity against different
68 HCV genotypes and subtypes should be characterised. The selection of resistance should be studied *in*
69 *vitro* for each genotype and the impact of mutations from wild-type on viral susceptibility should be
70 investigated. The viral drug target should be sequenced at baseline in clinical studies; furthermore,
71 genotypic resistance testing should be performed on samples from patients with virological failure and
72 phenotypic resistance testing should be performed if the impact of individual mutational events on
73 susceptibility remains uncharacterised or if no emerging mutations are detected.

74 The drug-drug interaction profile (DDI) of a new DAA or fixed dose combination (FDC) should be
75 adequately characterised, with focus on co-medications of crucial relevance for the target HIV infection
76 (e.g. including drugs used for the treatment of HIV, for management of liver transplantation and for
77 opiate substitution)

78 The primary endpoint in clinical trials aiming at viral clearance should be sustained virological response
79 defined as plasma HCV RNA below the lower limit of quantification of the assay (LLOQ) 12 weeks after
80 the planned end of therapy (SVR12). There should be further follow-up to confirm the durability of
81 response for novel drug regimens.

82 The sponsor should design the clinical development programme (pre- and post-initial licensure) so that
83 the efficacy and safety of the new DAA within one or more combination regimens is documented for
84 the full range of patients in whom beneficial effects and clinical use may be anticipated. The patient
85 and viral characteristics that should determine eligibility for each clinical trial will be selected
86 accordingly. As applicable, these characteristics may include viral genotype, level of liver damage
87 (degree of fibrosis, Child-Pugh classification category and any clinical features of decompensation) and
88 prior DAA regimen treatment history.

89 In general, randomized controlled trials with an active comparator, considered standard of care for the
90 study population, is the most informative study design for pivotal trials. This should be considered in
91 all cases. In case a DAA is developed as an add-on to an established combination (to increase efficacy
92 or to shorten treatment duration) or as a substitute for a component in such a combination,
93 randomized controlled trials against an active comparator are generally necessary to document efficacy.

94 If the sponsor is developing a wholly new combination regimen, and phase II data are indicative that
95 very high SVR rates are anticipated, it may not be essential to conduct randomised controlled studies
96 to describe efficacy. Since the spontaneous resolution rate of chronic HCV infection is negligible, and
97 key baseline demographic and disease factors that impact response are well described, it is possible to
98 assess the efficacy of a treatment regimen in uncontrolled trials in which the point estimate and its
99 precision (based on 95% confidence intervals) are documented. To document the safety profile, it is
100 recommended that at least one study in the program be of double-blind design vs. an active control or
101 placebo for the duration of the active treatment period(s), after which those assigned to placebo could
102 switch to open-label active treatment. Such a comparison is considered most valuable if performed in
103 patients with compensated cirrhosis.

104

105 For studies in patients with decompensated liver disease, an active standard-of-care comparator arm is
106 recommended.

107

108 **1. Introduction (background)**

109 Hepatitis C virus (HCV) is the most common infectious cause of chronic liver disease in Europe, and is
110 globally second only to Hepatitis B virus. Worldwide, approximately 3% of the population is estimated
111 to be infected, corresponding to around 200 million people at risk of developing serious liver related
112 morbidity. In Europe, where the vast majority of CHC cases are reported among patients with past
113 blood transfusion (before 1991) or with a history of intravenous drug use, the prevalence varies by
114 geographic region, from about 0.5% in the Northern countries to 2% and higher in the Mediterranean
115 countries and in Eastern Europe. HCV of genotype (GT) 1 is the predominant genotype globally as well
116 as in most European regions. In Europe and in the US, approximately 30% of HIV-infected patients are
117 co-infected with HCV, ranging up to 50% in some regions.

118 **2. Scope**

119 Guidance is provided on the design of clinical studies considered to be of relevance for the evaluation
120 of direct-acting anti-HCV compounds.

121 The scope of this guideline reflects the experience with DAA in the field of drug development for the
122 treatment of CHC. Sponsors planning modes of drug development that are not covered in this
123 guideline, are advised to consult with EU Regulators early in the clinical development programme, and
124 at least prior to initiating confirmatory studies.

125 **3. Legal basis and relevant guidelines**

126 This guideline has to be read in conjunction with the introduction and general principles (4) and parts I
127 and II of the Annex I to Directive 2001/83 as amended.

- 128 • Choice of a Non-Inferiority Margin - CPMP/EWP/2158/99
- 129 • Pharmacokinetic studies in man – CHMP/EWP/147013/04
- 130 • Investigation of drug interactions – CPMP/EWP/560/95
- 131 • Use of pharmacogenetic methodologies in the pharmacokinetic evaluation of medicinal products -
132 EMA/CHMP/37646/2009
- 133 • Evaluation of the pharmacokinetics of medicinal products in patients with impaired renal function -
134 CPMP/EWP/225/02
- 135 • Reporting the Results of Population Pharmacokinetic Analyses CHMP/EWP/185990/06
- 136 • Clinical investigation of medicinal products in the paediatric population – CPMP/ICH/2711/99
137 (ICH11)
- 138 • Role of Pharmacokinetics in the Development of Medicinal Products in the Paediatric Population
139 CHMP/EWP/147013/04

- 140 • Evaluation of the Pharmacokinetics of Medicinal Products in Patients with Impaired Hepatic
141 Function (CPMP/EWP/2339/02)
- 142 • Non-clinical Development of Fixed Combinations of Medicinal Products
143 (EMA/CHMP/SWP/258498/2005).
- 144 • Fixed Combination Medicinal Products CPMP/EWP/240/95
- 145 • Note for guidance on studies in support of special populations : Geriatrics (CPMP/ICH/379/95)
146

4. Pharmacodynamics and pharmacokinetics

4.1. Nonclinical virology studies

The preliminary in-vitro investigation of a new agent for the treatment of hepatitis C virus (HCV) should include the following:

1. A characterization of the mechanism of action of the new agent.
2. A determination of the antiviral activity (IC₅₀) in enzymatic assays (if such are available given the mechanism of action).
3. Determination of EC_{50/90} in cell based assays representing the different HCV genotypes and subtypes. Primarily, use of the sub-genomic replicon assay is anticipated to determine viral drug susceptibility. The choice of replicon representing each viral genotype/subtype (e.g., full length versus chimeric replicons) should be justified.
4. Determination of the impact of protein binding on EC_{50/90}.
5. Determination of the cytotoxicity and of the therapeutic index of the drug against the same cell line in which antiviral activity is determined.
6. For each viral genotype/subtype, an assessment of the in-vitro selection of resistant variants and characterisation of their phenotypic and genotypic properties. Selection experiments should be performed with a range of drug concentrations in relation to the EC₅₀, to characterize the concentration-dependency of the selection of resistant variants.
7. Characterization of the activity of the new agent against viruses/replicons (which may include clinical isolates or site directed mutants) harbouring a range of resistance associated mutations.
8. Studies of the activity of the new drug against other viruses (e.g. in particular HBV and HIV). If activity that might exert selective pressure against such viruses is detected, this should prompt further investigations to evaluate the potential for this to occur when using the agent to treat HCV in co-infected patients.
9. Studies of the potential for additive/synergistic or antagonistic effects to occur when the new agent is co-administered with other antiviral agents active against HCV. If the new agent is active against other viruses then further studies could be needed as appropriate to its spectrum.
10. If the new agent requires intracellular modification to form the active moiety (e.g. serial phosphorylation as for nucleoside/nucleotide analogues) it is important to assess the possible effects of co-incubation with other drugs that may compete for the activation pathway resulting in modification of antiviral activity.

When presenting in-vitro data, the assays and prototype strains used should be clearly defined and justified. It is preferable that the same methods should be used throughout the development programme to enable comparisons between studies. If methods are changed (e.g. due to modifications of or advances in assays over time) appropriate controls should be included to enable comparisons between studies.

183 **4.2. Clinical virology studies**

184 **4.2.1. Viral drug resistance**

185 The viral drug target gene should be sequenced at baseline for viruses obtained from all patients
186 entering clinical trials, unless otherwise justified. Naturally occurring polymorphisms associated with
187 differential drug efficacy should be identified. For example, the impact on drug susceptibility of
188 common polymorphisms should be analysed *in vitro* (see section 4.1) and trials should explore
189 correlations between baseline polymorphisms and viral response on-treatment and post-treatment.

190 Genotypic studies should be performed on samples obtained from patients at the time of documenting
191 lack of response, whether this is non-response or a loss of initial response. Any genotypic change that
192 has emerged since baseline should preliminarily be assumed to be due to the selective pressure of the
193 drug regimen, and should be explored for correlation with a phenotypic change if this has not
194 previously been established for the specific mutation(s) detected. If no genotypic change since baseline
195 is found then the isolate should undergo phenotypic analysis.

196 There are several different methods for the analysis of genotypic resistance. Population sequencing is
197 the standard method, but only detects variants with a frequency of about 20% (a figure that varies
198 depending on viral load). Clonal sequencing is more sensitive, and can provide additional information
199 about the linkage of mutations and the frequency of different quasispecies. Next generation
200 sequencing methods may provide a further understanding of on-treatment and post-treatment (in case
201 of failure to reach SVR) quasispecies dynamics. The sponsor should justify the methods used at each
202 stage of investigation, and should closely follow the scientific discussion and development of methods
203 within the field. Within clinical trials, samples should be stored to enable further analysis with different
204 methods, if required.

205 **4.2.2. Determination of HCV genotype and subtype**

206 The reference method for HCV genotype and subtype determination is direct sequencing and
207 phylogenetic analysis with either CE-marked or validated in-house techniques. Unless otherwise
208 justified, the target gene should be sequenced for all patients in the clinical investigation program (see
209 also above). Alternatively, one may use a CE-marked second generation line probe assay. Outside of
210 genotype 1, however, this is not sufficient for the determination of subtype; therefore, direct
211 sequencing is necessary. If other methods are used, this should be fully justified. Techniques based
212 solely on the analysis of the 5' non coding region are not recommended, as a too high incidence of
213 erroneous determination of the subtype has been reported.

214 **4.2.3. Determination of plasma HCV-RNA levels**

215 HCV RNA levels should be determined with a standardised, CE-marked quantitative assay based on
216 real-time PCR technology, with a lower limit of detection in the order of 10-15 IU/ml. Levels of viremia
217 below the lower limit of quantification (LLOQ), should be reported as "target detected" or "target not
218 detected" . The choice of assay should be appropriate for the genotypes in the study population, as
219 some assays have been reported to substantially underestimate HCV RNA levels in certain genotypes.

220 The same assay should be used for all samples from a single study and, whenever possible, throughout
221 the clinical development programme.

222 **4.3. Clinical pharmacokinetics**

223 The clinical pharmacokinetic study programme should follow the relevant CHMP guidelines
224 (Pharmacokinetic studies in man – CHMP/EWP/147013/04). In order to reduce the risk of selection of
225 drug resistant variants, the initial pharmacokinetic studies should be performed in healthy volunteers.
226 Studies of pharmacokinetics in patients with hepatic and renal impairment should be conducted in
227 accordance with the principles described in the relevant CHMP guidelines (CPMP/EWP/2339/02 and
228 CPMP/EWP/225/02). If it is known that the test agent has a high barrier to resistance, and selection of
229 resistance is unlikely, studies in patients with hepatic impairment may be performed in patients with
230 HCV infection.

231 **4.4. Drug-drug interactions**

232 The general principles described in CHMP guidance on the investigation of drug-drug interactions
233 should be followed (CPMP/EWP/560/95/Rev.1Corr*). In designing the mechanistically driven drug-drug
234 interaction programme, priority should be given to studies of oral contraceptives, as well as drugs used
235 in the management of HIV, liver transplantation, depression and substance abuse. Within these areas,
236 essential drugs (for which reasonable therapeutic alternatives are lacking) that have a foreseen
237 potential for interaction, should be prioritised for study.

238 Sufficient data to guide the safe use of the drug(s) in the target population is expected to be available
239 at the time of the initial marketing authorisation. If the possibility of a relevant interaction with an
240 important co-treating agent cannot be excluded *in vitro*, clinical studies should include an appropriate
241 design to allow for an assessment of the clinical significance of the putative interaction.

242 **5. Assessment of efficacy**

243 **5.1. General considerations for clinical trials**

244 Randomised, active-controlled studies with a standard-of-care regimen for the target population, is
245 generally considered the most informative design for confirmatory trials. In case such designs are not
246 used, a scientific justification is necessary. Further, unless specifically justified, randomised controlled
247 studies should be double-blind.

248 Due to the dynamics of the field, the appropriate design in terms of, e.g., genotypes and populations
249 to be studied, as well as in terms of appropriate comparator regimens, prior to commencing
250 confirmatory studies may change over time. A generally recommended standard of care regimen for
251 the particular target population would usually be considered the appropriate reference treatment in a
252 pivotal trial. However, spontaneous resolution of chronic HCV infection in the absence of therapy is a
253 very rare event. Therefore, studies without an active, prospective randomised control constituting an
254 approved and recommended regimen may be sufficiently informative if SVR12 rates are anticipated to
255 be very high (e.g., around 95%).

256 Possible alternative designs include a placebo control arm with delayed treatment, comparisons of
257 different regimens (doses, durations, number of drugs) including the new agent(s), or single arm
258 studies. If a pivotal study does not have a standard-of-care comparator arm, it is crucial that the
259 sponsor can justify that the demographic and disease characteristics of the patients included cover a

260 range that is relevant to the proposed recommended uses of the regimen. Enrichment of studies with
261 patients that have characteristics that may be associated with lower SVR12 rates, such as prior
262 treatment failure or advanced liver disease, may be considered in order to ascertain that SVR12 rates
263 are not driven by the selection of “easy to cure” patients.

264 It is notable that studies that do not randomise to a control arm may not be straightforward in their
265 interpretation if anticipated SVR rates turn out substantially lower than assumed at the planning stage;
266 from a scientific point randomised, active control trials remain the preferred option.

267 It is acknowledged that the pre-licensure clinical development programme may often include pivotal
268 trials with different study designs. In general, the applicant is encouraged to include at least one study
269 in which the test regimen is compared to placebo (deferred treatment), or to an active comparator, in
270 order to further the understanding of the safety profile of the regimen. Such comparative safety data
271 may be most informative in patients with cirrhosis.

272 **5.2. Subject characteristics and the definition of patient populations**

273 **5.2.1. Viral genotypes**

274 The patterns of activity (EC50 as well as barrier to resistance) of many DAAs are genotype- and
275 subtype dependent, with some agents showing *in vitro* and clinical activity only against certain
276 genotypes.

277 The range of genotypes for which clinical studies are relevant for a certain drug will be inferred initially
278 on the basis of *in-vitro* antiviral activity data. The results of early clinical studies (e.g. using
279 monotherapy against a range of genotypes) should be used to select the genotypes/sub-genotypes for
280 later studies.

281 The rationale for studying different genotypes and subtypes in separate studies or within the same
282 study should take into account which drug combinations, doses and treatment durations might be
283 optimal for each genotype. Such considerations may also include whether the same comparator
284 regimen is relevant for each genotype/subtype. If several genotypes/subtypes are studied within the
285 same trials in a development program, genotype or subtype may be an important stratification and/or
286 capping factor. The totality of evidence, from *in vitro* virological findings to clinical outcomes, must be
287 sufficient to enable a sound assessment of the benefit-risk relationship for each particular
288 genotype/subtype for which the use of a drug regimen is recommended. Concerning genotype/subtype
289 determination, see section 4.2.

290 **5.2.2. Host IL28B genotype**

291 Host IL28B genotype was first described as a major predictor of response to interferon-based regimens
292 in patients with genotype 1 (GT1) infection. It has subsequently emerged as a predictor of response
293 also to interferon-free regimens in GT1 when these are not optimized in terms of potency, barrier to
294 resistance and/or treatment duration. Furthermore, there are data to support the impact of IL28B
295 genotype on response to treatment of other viral genotypes too; however, this impact has tended to
296 be less consistent and smaller than in GT1. Therefore, categorisation of patients on the basis of a
297 favourable or non-favourable genotype (e.g., rs12979860 C/C vs C/T, T/T) is of potential importance
298 at several levels of drug development, and it is recommended that this parameter be recorded in all
299 patients participating in clinical trials within a drug development program for hepatitis C, regardless of
300 viral genotype/subtype. A sufficient number of patients with each IL28B genotype should be

301 investigated for inferences on the claimed treatment effect to be made for both C/C and non-C/C
302 genotypes.

303 **5.2.3. Treatment history**

304 It is recommended that peginterferon (pegIFN) +ribavirin treatment experience and prior response be
305 documented, as this is helpful in understanding the relationship of interferon response and response to
306 the interferon-free regimen. Furthermore, a targeted enrichment of treatment experienced patients
307 (particularly prior non/null responders) may be valuable in defining the optimal regimen (e.g.,
308 treatment duration) in those patients that have the lowest interferon response/host immunity to HCV.
309 The crucial issue is that the drug development program should provide the basis for the identification
310 of an appropriate regimen based on the known baseline characteristics of the individual patient.

311 For classifying prior response to pegIFN and ribavirin in genotype 1 infection, the following terms are
312 recommended:

- 313 • Null-response is defined as less than 2 log₁₀ decline in viral load at week 12.
- 314 • Partial-response is defined as at least 2 log₁₀ decline in viral load at week 12, but never achieving
315 an unquantifiable viral load
- 316 • Relapse is defined as unquantifiable virus at end of treatment but subsequent re-emergence of
317 quantifiable HCV-RNA.
- 318 • Breakthrough indicates the re-emergence of quantifiable virus while on treatment after previously
319 being unquantifiable or a confirmed increase of at least 1 log₁₀ in HCV-RNA during treatment.

320 Emerging categories of patients, in terms of treatment experience, include those that have failed
321 treatment with pegIFN+ribavirin in combination with a DAA, as well as patients that have failed
322 therapy with DAA only regimens. This issue is further discussed below, in section 5.7.4.

323 **5.2.4. Assessment of liver fibrosis**

324 The impact of cirrhosis on PK, efficacy and safety should be determined. The role of liver fibrosis
325 assessment within clinical trials may be to exclude patients with advanced fibrosis/cirrhosis from early
326 clinical trials, or, conversely, to correctly identify patients with cirrhosis, e.g., to enable stratification
327 and subgroup analysis of drug effect in such patients.

328 A number of different techniques for non-invasive assessment of liver histology are available. The
329 choice of method should be justified on the basis of the operating characteristics of the methods, in
330 view of the predictive value to include or exclude advanced fibrosis/cirrhosis, as relevant for the
331 particular purpose.

332 For patients in whom baseline histology is available through routine clinical care (liver biopsy
333 performed within 2 years prior to study entry), biopsy data should be collected and the relation
334 between baseline histology and efficacy and safety reported.

335 **5.3. Methods to evaluate efficacy**

336 The recommended primary endpoint for studies aiming at defining cure rate is sustained virological
337 response (SVR), defined as HCV-RNA < LLOQ 12 weeks after the *planned* completion of therapy
338 (SVR12), regardless of the actual duration of treatment. Patients with missing data should be

339 accounted as failures; the exception being that SVR12 may be imputed in patients for whom SVR has
340 been shown to be reached at a later date (e.g., SVR24).

341 SVR24 data should also be collected, and all available SVR24 data should be submitted at the time of
342 licensure, followed by submission of the remaining data as they emerge. Preferably the main study
343 protocols should follow patients up to one year after the planned end of treatment (EOT). Concerning
344 the long term follow up of patients, see section 5.5.6.

345 Apart from SVR, the kinetics of on-treatment viral response should be fully investigated and reported
346 in the drug development program,

347 Due to the approximate 90% predictive value of SVR4 for SVR12, it is reasonable to make decisions
348 within a clinical development program (e.g., going from phase II to phase III) on the basis of such
349 data.

350 **5.4. Dose finding studies**

351 **5.4.1. Monotherapy studies**

352 An adequate range of doses should be studied, based on protein binding-adjusted EC50 values *in vitro*
353 and on available dose-related drug exposure data from healthy volunteers. EC50 values of both wild-
354 type virus and viruses with mutations (single and in combination) derived during drug pressure *in vitro*
355 should be taken into account, so that selected doses for combination studies will be likely to provide
356 sufficient exposure for activity also against pre-existing variants with reduced drug susceptibility, if this
357 is feasible.

358 It is expected that monotherapy studies will initially be performed in chronic HCV-infected patients
359 without advanced fibrosis. Currently, 3 days of monotherapy, covering the first phase of viral decay, is
360 considered sufficient to assess the antiviral effect of a dose regimen in the general case. If *in vitro* data
361 and available knowledge of the drug class are strongly suggestive of a high barrier to resistance,
362 longer term monotherapy studies could be considered.

363 **5.4.2. Early combination dose ranging studies (phase 2a)**

364 As combination therapy is generally anticipated, such studies should be performed with the aim of
365 characterising appropriate doses, regimens and treatment durations for further investigation in phase 3.
366 It is anticipated that such studies will initially be performed in patients without advanced liver disease,
367 and subsequently in patients with more advanced disease. When including patients with a more urgent
368 need of treatment in experimental protocols, remaining options for treatment aiming at viral clearance
369 in case of failure should be considered. In particular, allocating cirrhotic patients to regimens of short
370 duration for which efficacy has not yet been established in patients with less advanced disease should
371 be avoided unless a likely effective salvage regimen would be available in case of virological failure
372 with the selection of drug resistant virus.

373 **5.5. Phase IIIb studies and confirmatory studies**

374 **5.5.1. Study populations**

375 Sponsors are generally encouraged to study the widest relevant range of patients in confirmatory
376 phase III studies, and particularly patients with advanced fibrosis. Unless there are specific

377 pharmacokinetic or safety concerns, it is expected that patients with compensated cirrhosis be included
378 in phase IIb/III studies.

379 Which subpopulations in terms of, e.g., viral (sub)genotype, IL28B genotype, cirrhosis/non-cirrhosis
380 and treatment experience are appropriate to study under the same protocol or under different
381 protocols may vary from case to case. This may depend on the known qualities of the regimen (e.g.,
382 the anticipated required potency and treatment duration), as well as on the availability of licensed and
383 recommended comparator regimens for the particular population. A specific concern is patients with
384 advanced fibrosis, who may require longer treatment duration for maximizing SVR rates.

385 **5.5.2. Selection of the study regimen**

386 Presently all clinically useful regimens for the treatment of HCV are combination regimens. An
387 investigational agent may be added to one or more previously approved drugs, or a test agent may be
388 substituted for a component of a recommended regimen, or the test regimen may exclusively consist
389 of two or more investigational drugs. As an increasing number of DAAs are approved, the sponsor
390 should carefully consider the respective value of add-on or substitution studies based on previously
391 approved drugs and regimens, versus the investigation of an entirely novel drug combination.

392 **5.5.3. Add-on and substitution studies**

393 In some cases, an active comparator arm is generally necessary. If the investigational drug is used as
394 an add-on or substitution to an approved regimen, that regimen should primarily be considered for
395 comparison, unless other designs can be justified. In the case of a substitution study, or an add-on
396 trial where the aim is to shorten treatment duration, a non-inferiority design would be relevant. If the
397 intent of the add-on study is to increase efficacy, a superiority design is required.

398 **5.5.4. Studies aiming at a shortened treatment duration**

399 Drug development may aim at documenting the efficacy of regimens shorter than those presently
400 generally recommended (i.e. <12 weeks). When including patients in trials with a shortened treatment
401 duration, patients in relatively urgent need of therapy (e.g., cirrhotic patients) should only be included
402 if there is a clear interferon-free treatment option in case of failure, taking anticipated cross-resistance
403 with approved agents into account. These considerations apply also to situations where the
404 recommended standard of care in a target population has a longer duration than the maximal duration
405 studied in the development program of the test agent.

406 **5.5.5. Fixed dose combinations**

407 Sponsors may develop single drugs or drugs formulated in FDCs. The latter may combine previously
408 approved drug(s) with new compounds, or only contain new compounds. The present guideline
409 concerns all these scenarios.

410 The specific guidelines for the development of fixed dose combination medicinal products should be
411 consulted and applied as relevant (EMA/CHMP/SWP/258498/2005).

412 **5.5.6. Follow-up after the primary endpoint**

413 The primary endpoint in confirmatory trials should be SVR (for further details, see above, section 5.3.).
414 A representative subset of patients achieving, as well as not achieving, SVR should be monitored after
415 determination of SVR12. For those that achieve SVR12, a total of one year follow up post EOT for

416 durability of response is requested (though not necessary at the time of the MAA). For patients not
417 reaching SVR12, a total of 3 year follow up post EOT with assessment of genotypic resistance is
418 requested. The aim of the latter is to understand the kinetics of reversion to wild-type and/or long-
419 term persistence of drug-resistant variants after the cessation of the selective pressure of the
420 treatment regimen. These follow-up data do not need to be available at the time of a market
421 authorisation application submission, but should be reported subsequently. If relevant, patients in a
422 long term follow up programme could be recruited for a re-treatment study

423 **5.5.7. Combination of medicinal products and the demonstration of the** 424 **contribution of each component to regimen efficacy**

425 The likely need for combination therapy from Phase 2a onwards is recognised. Given available
426 knowledge of general virological principles, as well as preclinical virology data relevant to the particular
427 regimen, trials that have a full factorial design to directly demonstrate the contribution of each agent
428 to efficacy, are not generally expected. The drug development programme should be designed to
429 provide a reasonable rationale for the need for each drug, given the totality of evidence (see also
430 section 5.5.3 concerning add-on and substitution studies).

431 **5.5.8. The extrapolation of efficacy between viral genotypes**

432 The different HCV genotypes show a different geographic distribution. Genotypes 1 and 3 dominate in
433 the EU, followed by genotypes 2 and 4. Genotypes 5 and 6 remain uncommon in areas where clinical
434 trials are generally performed. From a drug efficacy perspective, the genotypes differ in several
435 respects. First, it is well-known that the difficulty of achieving viral clearance with interferon-based
436 immune therapy differs between genotypes, e.g., with SVR rates despite longer treatment duration
437 and higher ribavirin dose in genotype 1 compared to genotypes 3 and -2. This may reflect intrinsic
438 differences in the host's ability to clear the different genotypes. Further, the activity of a particular
439 direct acting antiviral may differ between genotypes or subtypes for reasons that may be more or less
440 understood. This difference in activity may be due to different EC50s of the most common variant(s),
441 but may also be due to different barriers to resistance in different (sub)genotypes, due to the
442 frequency of resistant quasispecies. Moreover, the frequency of detectable, polymorphic variants may
443 differ between genotypes or subtypes (e.g., the NS3/4A Q80K polymorphism or the NS5A L31M
444 polymorphism). Furthermore, available evidence indicates that genotype 3 infections may intrinsically
445 be somewhat more difficult to cure with DAA therapy compared to other genotypes, even though viral
446 susceptibility may be similar. The reason for this is not fully understood.

447 Subject to the in-vitro virological data, it may be possible to use clinical efficacy data obtained against
448 one genotype to support a conclusion of efficacy against another genotype for which clinical data are
449 relatively limited. For example, efficacy against genotype 1 may support a conclusion on efficacy
450 against genotypes 4, 5 and 6. This approach may make it possible to give dose regimen
451 recommendations in section 4.2. of the SmPC for less commonly encountered genotypes (see section
452 7). In such a bridging exercise, available data are used to address relevant aspects concerning the sum
453 antiviral efficacy of the drug/regimen against the dominant quasispecies or most common
454 subtypes/variants and against detectable minor quasispecies. In order to support bridging of efficacy,
455 the following elements need to be taken into account.

456 First, there should be clear indications that the genotype to which the bridge is created, is not
457 intrinsically more difficult to clear than the genotype from which the bridge is built (e.g., a bridge from
458 genotype 2 to genotype 3 would not be accepted). It is anticipated that clinical efficacy data from
459 genotype 1 would generally be used for bridging.

460 Second, all available clinical and virological data must be taken into account when considering the
461 appropriateness of the bridging exercise. For example, there may be clinical efficacy data for individual
462 components of a regimen against the genotype(s) for which bridging is proposed. If there are no or
463 very few such the bridging exercise must be adequately supported by other evidence such as on-
464 treatment viral kinetics, including any available monotherapy data.

465 Third, the presumed similarity of on-treatment antiviral potency between genotypes must be supported
466 by similar replicon EC50s.

467 Fourth, the sponsor must provide an analysis of the genetic heterogeneity of the genotype to which
468 efficacy is bridged, with particular focus on the frequency of potentially relevant polymorphisms in the
469 gene coding for the molecular target. The case must be made that resistant variants or quasispecies
470 are not more common in the genotype(s) to which efficacy assumptions are bridged, than in the
471 genotype(s) from which assumptions are bridged.

472 **5.6. Studies in special patient populations**

473 **5.6.1. Treatment of patients with decompensated liver disease**

474 While the term “decompensated liver disease” often denotes those with present or past clinical
475 decompensation events such as variceal haemorrhage, ascites, serious bacterial infections or
476 encephalopathy, and the term “hepatic impairment” usually refers to a functional classification as
477 Child-Pugh B or C, these terms are here used interchangeably to denote either or both of these states.

478 Once there is sufficient evidence of an appropriate dosing regimen capable of delivering high rates of
479 SVR, as well as PK data in patients with hepatic impairment and a reasonable and acceptable safety
480 database in patients with less advanced disease, trials in patients with very advanced liver disease
481 may commence. Trials in this population are particularly encouraged for genotypes where there is
482 limited evidence for available treatment options or where the efficacy of these may be suboptimal.
483 Available general evidence concerning required treatment duration and the need for ribavirin to
484 optimize outcomes in patients with decompensated liver disease should be taken into account when
485 selecting regimens for study.

486 SVR is considered an appropriate primary endpoint also in studies of patients with decompensated liver
487 disease, along with prevention of graft infection in case of transplantation. In order to describe the
488 clinical benefit of SVR 12 in this population, it is recommended that patients be further followed up to
489 capture data on mortality, need for transplantation, hepatic function (e.g., MELD score), incidence of
490 hepatocellular carcinoma and reversal of fibrosis.

491 Prior to initiating clinical trials in patients with decompensated liver disease, pharmacokinetics and
492 short term safety should be investigated in patients over the relevant functional range (e.g., Child-
493 Pugh B and C). If the drug(s) do not have a high barrier to resistance, pharmacokinetic studies should
494 be performed in patients that are not infected with HCV. It is recommended that an established
495 treatment regimen for the target population (in terms of the viral genotypes included for study) is used
496 as an active comparator in order to appropriately characterise the safety and efficacy of the new drug
497 or regimen relative to the existing standard of care. An immediate versus deferred (placebo-controlled)
498 design may be less feasible in these patients with an urgent medical need.

499 It is crucial that the safety of study participants is appropriately monitored when testing new
500 compounds in the population with decompensated liver disease.

501 **5.6.2. Post-transplant treatment**

502 Reinfection of the liver graft is inevitable in patients with detectable HCV-RNA prior to transplantation.
503 Progress to cirrhosis is rapid, and the prognosis of patients transplanted due to HCV is worse than
504 when transplanted for other indications. The tolerability of ribavirin is compromised in this group, and
505 several studies of interferon-free combinations have initiated patients on lower than standard doses of
506 ribavirin. Furthermore, ensuring that potential drug interactions with immunosuppressive agents can
507 be appropriately managed is an important goal of studies in this population. It is recognised that
508 formal drug interaction studies with some immunosuppressive agents may not readily be conducted in
509 healthy volunteers, except on a single dose basis, and that close monitoring of pharmacokinetics may
510 be required during trials. It is presently not entirely clear whether post-transplant status, including the
511 impact of immunosuppression, impacts response to DAA therapy independently of other factors such
512 as fibrosis status; e.g., most available data are on regimens containing ribavirin, and it has not been
513 clarified whether this is needed in the general case. Therefore, clinical efficacy studies in this
514 population are encouraged.

515 **5.6.3. HCV/HIV co-infected patients**

516 The progression of liver disease may be more rapid in patients co-infected with HIV, at least in those
517 with low CD4+ cell counts. Response rates to pegIFN+ribavirin has historically been lower than in
518 mono-infected patients; this however, has generally not been the case when direct acting antivirals are
519 used. Furthermore, based on emerging data and the DDI profile of a given regimen, the inclusion of
520 HCV/HIV co-infected patients in general confirmatory trials may be considered, provided that similar
521 treatment regimens are studied regardless of co-infection status. In such a case, stratification and/or
522 capping for co-infected patients may be relevant. It is of particular importance that a majority of the
523 patients studied are receiving antiretroviral therapy, to confirm that recommendations concerning the
524 management of drug interactions provided in section 4.5 of the SmPC, are in fact useful in providing
525 efficacious and safe co-therapy against HIV and hepatitis C. Population pharmacokinetic studies should
526 be part of these trials, to confirm that the expected exposures are yielded (for new agents and
527 antiretrovirals with proven/potential interactions).

528 **5.6.4. Patients with prior DAA experience**

529 This patient population is of considerable heterogeneity. For instance:

- 530 • The prior DAA class and compound(s) tried differ(s).
- 531 • The reason for unsuccessful treatment with a DAA regimen may be virological failure or lack of
532 tolerance including adherence issues.
- 533 • Patients may or may not have evidence of persistent viral resistance.

534 The most important *scientific* question pertaining to patients with prior virologic failure and/or selection
535 of variants resistant to DAAs, may be to understand its impact on the contribution of the same agent
536 or a cross resistant agent as a component in a more potent regimen (e.g., including more drugs, a
537 longer treatment duration and/or higher doses). However, the *clinically* most relevant retreatment
538 scenario in most cases may be with a potent combination of drugs of classes to which the patient has
539 not been exposed or to which cross-resistance is not anticipated, with or without ribavirin.

540 Much remains unknown concerning the impact of emergent drug resistance on subsequent therapy
541 with a partially or potentially cross-resistant compound. It is clear, however, that virtually all patients
542 that fail virologically when treated with DAAs while adhering to therapy are intrinsically “difficult to

543 cure". This should be taken into account when designing studies for patients that have experienced
544 virological failure on DAA-containing regimens. The virological rationale for regimens used in studies of
545 retreatment of patients with prior failure on DAA regimens should be carefully considered (e.g., the
546 anticipated potency and barrier to resistance of the experimental regimen), and emerging data should
547 be taken into account. It is anticipated that drug pressure (sum potency, treatment duration) will need
548 to be increased compared to the previous treatment attempt, in order to optimise responses in
549 patients with prior virological treatment failure.

550 If the investigational regimen includes a DAA to which the patients have been exposed, or a potentially
551 cross-resistant agent, baseline drug resistance should be thoroughly investigated so that firm
552 conclusions can be drawn about its impact on treatment response. Retreatment studies of patients with
553 DAA experience that have reverted to wild-type after the selection of resistance during therapy are
554 considered of particular importance for understanding the impact of acquired drug resistance.

555 Patients that have failed DAA based regimens due to lack of tolerability, and that do not have evidence
556 of drug resistance, should be evaluated on a case to case basis as regards re-treatment, and are not
557 considered a well-defined target population for clinical trials.

558 **5.6.5. Studies in paediatric patients**

559 It is currently not generally anticipated that clinical efficacy and safety studies in children will be
560 performed until after completion of Phase 3 studies in adults. However, PK studies in adolescents
561 anticipated to require the adult dose regimen may begin earlier and these patients may be included in
562 adult confirmatory trials.

563 Suitable age-appropriate formulations should be developed, palatability being of particular concern.

564 Similar to the case with HIV, it is considered that efficacy data may be bridged from adults to children,
565 provided that similar drug exposure is reached in plasma at the recommended doses. Studies primarily
566 aiming at characterising PK and selecting appropriate doses should cover an appropriate range of ages
567 (generally from 3 years and upwards), and should aim at achieving adult plasma drug exposures.
568 Treatment should be continued for a duration that is sufficient to reach SVR to provide clinical benefit
569 for study participants and to generate some safety and efficacy data. Such studies could include the
570 full range of patients (e.g., in terms of viral genotypes and other disease characteristics) for whom the
571 use of the drug/regimen is recommended in adults. It is recognised that the number of children and
572 adolescents with chronic hepatitis C eligible for clinical trials is limited. If there are no specific safety
573 concerns relevant to the paediatric population, pre-authorisation studies could be limited in size to 30-
574 40 patients distributed across the age range from 3 to less than 18 years old. As stated above, these
575 studies could primarily focus on the determination of PK, but would also collect, albeit in a rather
576 limited fashion, data on safety and efficacy. After authorisation, additional safety data would need to
577 be collected, possibly in form of a registry.

578 **3.6.6 Studies in older patients**

579 Hitherto pivotal studies have included relatively few elderly people. While the elderly are not
580 considered a special population in the sense of the abovementioned categories, the inclusion of elderly
581 subjects in clinical trials is generally encouraged.

582 **6. Safety aspects**

583 Specific safety concerns related to the treatment of chronic hepatitis C that are of relevance for the
584 development of new DAAs include impaired liver function at baseline, the known toxicity of currently
585 licensed drugs such as ribavirin, the potential for additive or synergistic toxicities of co-treating agents,
586 PK interactions and development of drug resistance. It is expected that mechanism-related toxicities
587 (such as mitochondrial toxicity for nucleoside analogues) will have been well characterised in non-
588 clinical and clinical studies. Any signals that emerge from the non-clinical studies should be followed in
589 the clinical development programme.

590 A particular problem concerns the investigation of the safety profile might arise when two or more
591 DAAs are investigated in combination, without either agent having previously characterised as to its
592 individual safety profile. Sponsors studying combinations of novel drugs are urged to consider this
593 problem. One way to address this issue is to also investigate one or both DAAs in combination with
594 agents with a previously described safety profile, where the safety profile of the individual
595 investigational agent can be characterised.

596 If the drug is subject to an expanded access program in patients outside criteria of clinical trial
597 population, safety data should be collected, as appropriate.

598 **7. Information in the Summary of the Product** 599 **Characteristics**

600 In the general case, the indication (section 4.1. of the SmPC) for DAAs against HCV infection should be
601 as follows:

602 *"[TRADENAME] is indicated in combination with other agents for the treatment of chronic hepatitis C*
603 *(CHC) in adults (see sections 4.2., 4.4. and 5.1.)*

604 *for genotype specific activity, see sections 4.4 and 5.1."*

605 For fixed dose combinations that may constitute a full regimen, a similar indication, excluding the
606 statement "in combination with other agents" is appropriate in the general case.

607 Section 4.4. should contain information on lack of data in clinically relevant subpopulations, and thus
608 reflect the potential absence of data to underlie a regimen recommendation, as well as any relevant
609 uncertainty concerning the optimal regimen in different clinical situations. This section may contain
610 recommendations for non-use in case of certain viral genotypes, viral polymorphisms, clinical
611 situations or certain prior DAA experience.

612 The efficacy data underlying regimen recommendations should be cited in section 5.1., as well as other
613 efficacy data considered of relevance to the prescriber and clinically relevant information on drug
614 resistance. Furthermore, this section should contain a summary of the in vitro potency against each
615 genotype, resistance pathways on in vitro selection and short term monotherapy activity against each
616 genotype. Any molecular understanding of genotype specific activity, such as conserved baseline viral
617 polymorphisms that might impact the activity of the drug, should be highlighted.

618

619 **Definitions**

620	CE	European Conformity
621	CHC	Chronic Hepatitis C
622	DAA	Direct acting antiviral
623	DDI	Drug-drug interactions
624	EC50	Median Effective Concentration to induce a 50% effect
625	EOT	End of treatment
626	FDC	Fixed dose combination
627	GT	Genotype
628	HCV	Hepatitis C virus
629	HIV	Human Immunodeficiency Virus
630	IL28	Interleukin 28B
631	LLOQ	Lower limit of quantification
632	MELD	Model End Stage Liver Disease
633	pegIFN	Peginterferon alfa
634	RNA	Ribonucleic acid
635	SVR	Sustained virological response