

# Systematic review

## Hepatitis C outcomes among HIV co-infected individuals in programmatic setting<sup>1</sup>

### OBJECTIVE

To determine the programmatic outcomes for the treatment of hepatitis C virus in chronically infected, treatment naïve individuals who are co-infected with HIV.

### METHODS

#### Search Strategy

- See annex

#### Databases

- MEDLINE
- EMBASE
- Bibliographies of relevant articles will be checked for further studies

#### Inclusions and exclusions

##### Types of studies

- Prospective and retrospective cohorts reporting outcomes for  $\geq 10$  patients
- All other study designs, including randomized trials, will be excluded.

##### Types of participants

###### *Inclusions:*

- Chronic HCV-infected adults co-infected with HIV who have completed treatment for HCV.

###### *Exclusions:*

- Cohorts of patients with other co-morbidities (eg HBV, haemophilia, liver transplant). Cohorts in which  $>10\%$  of patients have an additional co-morbidity will be excluded.
- Treatment-experienced cohorts. Cohorts in which  $<5\%$  of patients are treatment-experienced will be included.

##### Types of interventions

- Interferon (either pegylated or standard) with or without ribavirin.

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## **Types of outcomes**

### **Primary**

- Sustained virological response (SVR), defined as undetectable serum levels of viral RNA 24 weeks following the end of treatment

### **Secondary**

- Rapid virological response (RVR), defined as undetectable serum levels of viral RNA 24 weeks following the end of treatment
- Discontinuation of treatment due to adverse events
- Defaulting
- Mortality

## **DATA ANALYSIS**

Point estimates and 95% confidence intervals (95% CI) will be calculated for all primary and secondary outcomes. The primary outcome of SVR will be determined in an 'intent-to-treat' like analysis, with all patients initiating HCV therapy contributing to the denominator. Proportions will be root-transformed, and estimates pooled by random effects meta-analysis.

### **Heterogeneity**

The potential influence of clinical and programmatic covariates was explored through univariate subgroup analyses to assess the potential influence of:

- Genotype
- Type of HCV treatment
- Co-treatment with antiretroviral therapy
- Baseline liver damage

A P-value  $\leq 0.05$  considered as significant.

### **Statistical software**

All analyses were conducted in Stata version 12

Annex: Search strategy

#22	#13 AND #18 AND #21
#21	#19 or #20
#20	SVR
#19	Sustained virological response
#18	#14 OR #15 OR #16 OR #17
#17	Intervention
#16	Therap*
#15	Treat*
#14	Explode "Therapeutics"[MeSH terms]
#13	#4 AND #12
#12	#5 OR #6 OR #7 OR #8 OR #9 OR #10 OR #11
#11	Acquired Immune Deficiency Syndrome
#10	Acquired Immunodeficiency Syndrome
#9	AIDS
#8	Human Immunodeficiency
#7	HIV
#6	Explode "HIV Infections"[MeSH Terms]
#5	Explode "HIV"[MeSH Terms]
#4	#1 OR #2 OR #3
#3	HCV
#2	Hepatitis C
#1	Explode "Hepatitis C"[MeSH Terms]