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The Ability of Hepascore to Predict Liver Fibrosis in Chronic Liver Disease: a Meta-analysis

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List of Abbreviations: HCV: chronic hepatitis C virus; HBV: chronic hepatitis B; ALD: alcoholic liver disease; NAFLD: non-alcoholic fatty liver disease; AUROC: area under ROC curve; CI: confidence interval; QUADAS: quality assessment of diagnostic accuracy studies; DANA: difference between the mean fibrosis stages in advanced fibrosis and non-advanced fibrosis groups; PPV: positive predictive value; NPV: negative predictive value; HIV: human immunodeficiency virus;

Conflict of interest: The University of Western Australia (employer of YH, LAA and GPJ) hold the patent for Hepascore and have a licencing agreement with Quest Diagnostics.

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ABSTRACT

Background & Aims: Hepascore is a serum model that was developed to assess the severity of liver fibrosis. It has been well validated in common causes of chronic liver disease. This study performed a meta-analysis to evaluate the pooled diagnostic performance of Hepascore and to compare it for different aetiologies of chronic liver disease.

Methods: Two reviewers searched electronic databases from October 2005 to September 2015 for studies that evaluated the diagnostic performance of Hepascore for liver fibrosis in chronic liver disease.

Results: 21 studies were included. The AUROC was adjusted according to the distribution of fibrosis stages. The mean adjusted AUROC was 0.83 (95%CI, 0.81-0.85) for significant fibrosis, 0.89 (95%CI, 0.85-0.92) for advance fibrosis and 0.93 (95%CI, 0.91-0.95) for

cirrhosis. A cut point of 0.50-0.55 achieved a summary sensitivity of 70% and a summary specificity of 79% to predict significant fibrosis. A cut point of 0.50-0.61 had a summary sensitivity of 81% and a summary specificity of 74% to predict advanced fibrosis. A cut point of 0.80-0.84 had a summary sensitivity of 72% and a summary specificity of 0.88% to predict cirrhosis. The accuracy of Hepascore was similar among all disease aetiologies for the prediction of cirrhosis. However, Hepascore had better diagnostic ability for significant and advanced fibrosis in chronic hepatitis C, chronic hepatitis B and alcoholic liver disease than for non-alcoholic fatty liver disease and HIV co-infected viral hepatitis.

Conclusions: Hepascore is a clinically useful measure of liver fibrosis in patients with common causes of chronic liver disease.

Key words: Hepascore; DANA; liver fibrosis; AUROC

KEY POINT BOX

- Hepascore has been validated in chronic hepatitis C, chronic hepatitis B, alcoholic liver disease and non-alcoholic fatty liver disease.
- Hepascore had an excellent accuracy to exclude cirrhosis in all four common causes of chronic liver disease.
- Hepascore had good diagnostic performance for significant fibrosis and advanced fibrosis in chronic hepatitis C, chronic hepatitis B and alcoholic liver disease.
- A cut point of 0.50-0.61 is predictive of significant fibrosis and advanced fibrosis and a cut point of 0.80-0.84 is predictive of cirrhosis.

INTRODUCTION

Chronic liver disease is a major global health problem. Chronic hepatitis C virus (HCV) infection, chronic hepatitis B virus (HBV) infection, alcoholic liver disease (ALD) and non-alcoholic fatty liver disease (NAFLD) are the main causes of chronic liver disease. Most chronic liver diseases have a similar clinical course with a prolonged asymptomatic early phase during which liver damage progresses silently and variable late clinical presentation of decompensated cirrhosis. The disease prognosis is closely associated with the severity of liver fibrosis and the majority of adverse outcomes occur after the development of liver cirrhosis [1]. As a result, liver fibrosis severity is currently the most reliable patient prognostic measure. Additionally, the measurement of liver fibrosis can help to guide important clinical management decisions that include the need for treatment and the initiation of hepatocellular carcinoma and liver decompensation surveillance.

Histopathological staging of liver biopsy has been used as the gold standard for liver fibrosis measurement. However liver biopsy is limited in its use due to its invasive nature, sampling error and risk of serious complications or death [2]. Furthermore, liver biopsy is inconvenient, expensive and not widely accessible to a large number of patients or physicians. Surrogate serum fibrosis models that can accurately predict the severity of liver fibrosis are of great clinical significance. Compared to liver biopsy, serum fibrosis models have advantages of low cost, wide availability, high reproducibility and non-invasive nature. During the last two decades, a number of serum fibrosis models have been developed and a few of them have been well validated and used in routine clinical practice.

Hepascore is a serum model that was developed to predict the severity of liver fibrosis in patients with chronic hepatitis C [3]. Hepascore includes four biomarkers, namely: alpha2-macroglobulin, hyaluronic acid, bilirubin and gamma-glutamyl transpeptidase, as well as age and gender [3]. After the initial development and validation of Hepascore in chronic hepatitis

C, Hepascore has been more widely validated in chronic hepatitis B, ALD and NAFLD to detect significant liver fibrosis (Metavir F2, F3, F4), advanced fibrosis (Metavir F3, F4) and cirrhosis (Metavir F4). However, no meta-analysis of these studies has been performed. This study performed a meta-analysis of all Hepascore validation studies to evaluate the summary diagnostic performance of Hepascore and to compare it across different aetiologies of chronic liver disease.

MATERIALS AND METHODS

Literature search strategy

An electronic search was performed on PubMed and Cochrane library using the key word “Hepascore” from October 2005 to September 2015. Studies were included if they met the following inclusion criteria: 1) The whole population or a sub-group of patients who had chronic hepatitis C with or without HIV co-infection, chronic hepatitis B with or without HIV co-infection, ALD, NAFLD or mixed aetiology of chronic liver disease could be extracted from the study. 2) Both liver biopsy and Hepascore were performed for patients and the diagnostic performance of Hepascore was evaluated. 3) Studies provided the area under ROC curve (AUROC) (95% CI) of Hepascore for different fibrosis stages and/or the true positive, true negative, false positive and false negative of at least one cut point could be calculated from the data. Exclusion criteria included: 1) non-English literature. 2) reviews. 3) duplicated cohorts. 4) The target population were patients with liver disease other than chronic hepatitis C, chronic hepatitis B, ALD and NAFLD.

Study inclusion and quality assessment

The studies were assessed by two independent reviewers using predetermined inclusion and exclusion criteria. The quality of each study was determined by using the validated quality assessment of diagnostic accuracy studies (QUADAS) questionnaire [4]. The questionnaire included 14 questions that covered use of appropriate patient population and reference standard, disease progression bias, verification bias, review bias, clinical review bias, incorporation bias, test execution, study withdrawals and indeterminate bias [4]. For each question, yes, no or unclear was scored. Disagreements between the two reviewers were resolved through further review.

Statistical analysis

Two approaches were used to evaluate the summary diagnostic performance of Hepascore. Firstly, meta-analysis was performed using the random effects model for all included studies that provided both AUROC and 95% CI. Previous studies found that the AUROC was significantly influenced by the distribution of fibrosis stages [5]. Hence, the AUROC was standardized according to the difference between the mean fibrosis stages in advanced fibrosis and non-advanced fibrosis groups (DANA) using the formula: $adAUROC = obAUROC + (0.1056) * (2.5 - DANA)$ [5]. Meta-analysis was performed for both observed AUROC (obAUROC) and adjusted AUROC (adAUROC). ObAUROC and adAUROC of Hepascore was compared between different aetiologies of chronic liver disease. Meta-regression analysis and sensitivity analysis were used to evaluate the influence of seven characteristics of individual studies on AUROC, namely: patient inclusion methods (single centre vs multicentre), mean biopsy length (<20mm vs \geq 20mm), biopsy evaluation (blinded vs not blinded), interval time between biopsy and serum collection (within one month vs >1

month), serum collection (fasting vs non-fasting), Hyaluronic acid test kit (Corgenix vs other kit) and study quality (all question score yes vs one or more questions scored no or unclear).

Secondly, a summary ROC (SROC) model was calculated for all included studies from which at least one 2X2 table containing true positive, true negative, false positive and false negative could be created [6]. Summary sensitivity and specificity of validated cut points to predict significant fibrosis, advanced fibrosis and cirrhosis were calculated. The estimated positive predictive value (PPV) and negative predictive value (NPV) of each cut point was calculated using the observed prevalence.

RESULTS

Literature search results

A total of 55 articles were identified from the literature search. 45 were original studies and were published in English. 24 studies were excluded for the following reasons: 14 studies did not evaluate the diagnostic performance of Hepascore or did not provide sufficient data for the meta-analysis; six studies evaluated the utility of Hepascore in other types of liver disease; four studies had duplicated cohorts. 21 studies were included in the final analysis. The main characteristics of these studies were shown in table 1 [3, 7-27]. A total of 5686 patients were included: 3523 had HCV infection, 441 had HCV/HIV co-infection, 588 had HBV infection, 108 had HBV/HIV co-infection, 321 had ALD, 242 had NAFLD and 463 had mixed aetiology of chronic liver disease. 11% patients had Metavir F0 (range: 0-44%), 34% had Metavir F1 (range: 11-47%), 25% had Metavir F2 (range 17-40%), 15% had Metavir F3 (range: 3-25%), 15% had Metavir F4 (range: 6-32%). According to the QUADAS

questionnaire, the qualities of the final 21 studies were good to excellent. (table 2). Twenty of these used the Metavir staging system as the reference standard.

Diagnostic performance of Hepascore for significant fibrosis

19 cohorts reported AUROC and 95%CI for significant fibrosis. The mean obAUROC of Hepascore was 0.78 (95%CI, 0.76-0.80) and the mean adAUROC was 0.83 (95%CI, 0.81-0.85) (figure 1). Hepascore had an increased diagnostic accuracy in HCV, HBV, ALD and mixed aetiology with a mean obAUROC of 0.80 (95%CI, 0.77-0.82), 0.79 (95%CI, 0.75-0.83), 0.82 (95%CI, 0.76-0.87) and 0.80 (95%CI, 0.74-0.86) respectively. Less diagnostic accuracy of Hepascore was found for patients with HIV co-infection and NAFLD, with a mean obAUROC of 0.73 (95%CI, 0.67-0.79) and 0.73 (95%CI, 0.66-0.80) respectively. Compared to obAUROC, most adAUROC's for different causes of liver disease increased, with adAUROC of 0.85 (95%CI, 0.82-0.87) in HCV, 0.84 (95%CI, 0.80-0.88) in HBV, 0.83 (95%CI, 0.76-0.89) in ALD, 0.80 (95%CI, 0.76-0.84) in HIV co-infection and 0.81 (95%CI, 0.75-0.87) in mixed aetiologies (figure 1). The diagnostic accuracy for NAFLD remained less than other forms of liver disease with adAUROC of 0.74 (95%CI, 0.67-0.81). Significant heterogeneity was observed for obAUROC ($I^2=55.6\%$, $p=0.002$) and for adAUROC ($I^2=47\%$, $p=0.013$). Sub-group analysis of heterogeneity was performed according to the causes of chronic liver disease. Non-significant heterogeneity of obAUROC was found in chronic hepatitis B ($I^2=6.8\%$, $p=0.359$), HIV co-infection ($I^2=47\%$, $p=0.152$) and ALD ($I^2=7.4\%$, $p=0.299$), while significant heterogeneity was found in chronic hepatitis C ($I^2=60\%$, $p=0.015$). Non-significant heterogeneity of adAUROC was found in all analysed causes: chronic hepatitis C ($I^2=48.8\%$, $p=0.057$), chronic hepatitis B ($I^2=6.5\%$, $p=0.361$), HIV co-infection ($I^2=0.0\%$, $p=0.625$) and ALD ($I^2=27.9\%$, $p=0.239$). Sub-group analysis was not performed

for NAFLD and mixed aetiologies due to only one study of each cause was available for analysis.

The diagnostic performance of at least one cut point was reported in 18 cohorts and these were included in the SROC analysis (table 3). The average prevalence of significant fibrosis was 52% (ranged: 41% - 66%). The summary AUROC for all causes of liver disease was 0.79 (95%CI, 0.76-0.83) and this was similar to the AUROC calculated using the random effects model (figure 2A). The summary diagnostic odds ratio (DOR) was 7.3 (5.9-9.1). A cut point of 0.50-0.55 was validated in 10 cohorts with a summary sensitivity of 0.70 (95% CI, 0.60-0.78) and summary specificity of 0.79 (95% CI, 0.72-0.85). A lower cut point of 0.31-0.34 was validated in four cohorts with a summary sensitivity of 0.75 (95%CI, 0.60-0.86) and a summary specificity of 0.65 (95%CI, 0.57-0.73). Using the average observed prevalence of significant fibrosis (52%) in the included studies, the cut point of 0.5-0.55 had an estimated PPV of 0.78 and an estimated NPV of 0.71. The cut point of 0.31-0.34 had an estimated PPV of 0.70 and an estimated NPV of 0.71.

Diagnostic performance of Hepascore for advanced fibrosis

16 cohorts reported AUROC and 95%CI for advanced fibrosis. The mean obAUROC was 0.84 (95%CI, 0.81-0.87) and the mean adAUROC was 0.89 (95%CI, 0.85-0.92) (figure 3). Hepascore achieved a higher diagnostic accuracy in HCV and HBV with a mean obAUROC of 0.85 (95%CI, 0.80-0.90) and 0.86 (95%CI, 0.79-0.94) respectively. The obAUROC was 0.78 (95%CI, 0.72-0.85) for HIV co-infection, 0.83 (95%CI, 0.74-0.93) for ALD and 0.81 (95%CI, 0.73-0.90) for NAFLD. A similar pattern was found using adAUROC, with the mean adAUROC of 0.90 (95%CI, 0.86-0.95) in HCV, 0.91 (95%CI, 0.84-0.98) in HBV, 0.86 (95%CI, 0.82-0.90) in HIV co-infection, 0.84 (95%CI, 0.74-0.93) in ALD, 0.83 (95%CI, 0.74-0.91) in NAFLD and 0.73 (95%CI, 0.62-0.83) in mixed liver disease aetiologies.

Significant heterogeneity was observed for both obAUROC ($I^2=83.5\%$, $p<0.001$) and adAUROC ($I^2=81.1\%$, $p<0.001$). Sub-group analysis showed that significant heterogeneity of both obAUROC and adAUROC was found in HCV (obAUROC: $I^2=89\%$, $p<0.001$, adAUROC: $I^2=85.7\%$, $p<0.001$) and HBV (obAUROC: $I^2=80.8\%$, $p=0.001$, adAUROC: $I^2=76.9\%$, $p=0.005$) but not for HIV co-infection (obAUROC: $I^2=43.7\%$, $p=0.183$, adAUROC: $I^2=0.0\%$, $p=0.646$). Sub-group analysis of heterogeneity was not performed for ALD, NAFLD and mixed aetiologies due to the presence of one study in each subgroup.

14 cohorts validated cut points to predict advanced fibrosis and these were included in the SROC analysis (table 3). The average prevalence of advanced fibrosis was 27% (ranged: 19% - 38%). The summary AUROC was 0.84 (95%CI, 0.81-0.87) and summary DOR was 11.9 (95%CI, 9.0 - 15.7) (figure 2B). A cut point of 0.50-0.61 was validated in seven cohorts with a summary sensitivity of 0.81 (95% CI, 0.71-0.87) and a summary specificity of 0.74 (95% CI, 0.72-0.77). Using the average observed prevalence of advanced fibrosis (27%), the same cut point had an estimated PPV of 0.54 and an estimated NPV of 0.91.

Diagnostic performance of Hepascore for cirrhosis

15 cohorts reported AUROC and 95%CI for cirrhosis. The mean obAUROC was 0.88 (95%CI, 0.86-0.90) and the mean adAUROC was 0.93 (95%CI, 0.91-0.95) (figure 4). Excellent accuracy was observed in all aetiologies of chronic liver disease with obAUROC of 0.89 (95%CI, 0.88-0.91) in HCV, 0.88 (95%CI, 0.83-0.92) in HBV, 0.87 (95%CI, 0.79-0.96) for HIV co-infection, 0.85 (95%CI, 0.70-1.00) for ALD, 0.91 (95%CI, 0.83-0.99) for NAFLD. The adAUROC was 0.95 (95%CI, 0.93-0.97) in HCV, 0.92 (95%CI, 0.87-0.98) in HBV, 0.95 (95%CI, 0.90-0.99) for HIV co-infection, 0.86 (95%CI, 0.70-1.00) for ALD and 0.92 (95%CI, 0.84-1.00) for NAFLD. Hepascore had less diagnostic accuracy in patients with

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mixed aetiologies with obAUROC of 0.81 (95%CI, 0.75-0.86) and adAUROC of 0.82 (95%CI, 0.76-0.88). Significant heterogeneity was observed for both obAUROC ($I^2=44.2\%$, $p=0.034$) and adAUROC ($I^2=62.2\%$, $p=0.001$). Sub-group analysis found significant heterogeneity of obAUROC in HIV co-infection ($I^2=82.2\%$, $p=0.018$) and ALD ($I^2=78.9\%$, $p=0.029$) but not in HCV ($I^2=0.0\%$, $p=0.671$) and HBV ($I^2=0.0\%$, $p=0.380$). Significant heterogeneity of adAUROC was only found in ALD ($I^2=81.2\%$, $p=0.021$) but not in chronic hepatitis C ($I^2=29.1\%$, $p=0.217$), chronic hepatitis B ($I^2=22.3\%$, $p=0.276$) and HIV co-infection ($I^2=26.1\%$, $p=0.245$). Sub-group analysis of heterogeneity was not performed for NAFLD and mixed aetiologies due to the presence of one study in each subgroup.

13 cohorts validated cut points to predict cirrhosis and were included in the SROC analysis (table 3). The average prevalence of advanced fibrosis was 13% (ranged: 6% - 31%). The summary AUROC was 0.90 (95%CI, 0.88-0.93) and the summary DOR was 25.3 (95%CI, 17.1-37.5) (figure 2C). A cut point of 0.80-0.84 was validated in seven cohorts with a summary sensitivity of 0.72 (95% CI, 0.64-0.79) and a summary specificity of 0.88 (95% CI, 0.85-0.91). Using the average observed prevalence of cirrhosis (13%), the same cut point had an estimated PPV of 0.47 and an estimated NPV of 0.95.

Sensitivity analysis

A sensitivity analysis was performed to test the effect of study characteristics on the adAUROC of Hepascore. No significant effect on adAUROC was found when patient inclusion methods, mean biopsy length, biopsy evaluation, interval time between biopsy and serum collection, timing of serum collection, hyaluronic acid test kit and study quality were analysed (table 4).

DISCUSSION

This meta-analysis reviewed 21 studies that evaluated the diagnostic performance of Hepascore for measuring the severity of liver fibrosis in chronic liver disease. Duplicate cohorts were excluded. The strength of this meta-analysis was that these studies included worldwide populations of the most common aetiologies of chronic liver disease, namely chronic hepatitis C, chronic hepatitis B, ALD and NAFLD. Standardization of the AUROC according to the distribution of liver fibrosis stages amongst cohorts allowed a more accurate comparison of the diagnostic performance of Hepascore for different aetiologies of chronic liver disease.

Hepascore had an excellent accuracy to predict cirrhosis for all aetiologies of chronic liver disease with an adAUROC of 0.95 for chronic hepatitis C, 0.92 for chronic hepatitis B, 0.86 for ALD, 0.92 for NAFLD and 0.95 for HIV co-infected chronic hepatitis. Meta-analysis of chronic hepatitis C and chronic hepatitis B found that Hepascore also had an excellent adAUROC for significant fibrosis, 0.85 and 0.84 respectively and for advanced fibrosis, 0.90 and 0.91 respectively. In ALD, NAFLD and HIV co-infected chronic hepatitis the diagnostic performance of Hepascore was good for significant fibrosis and advanced fibrosis. The adAUROC for significant fibrosis in ALD, NAFLD and HIV co-infection hepatitis was 0.83, 0.74 and 0.80 respectively and for advanced fibrosis the adAUROC was 0.84, 0.83 and 0.86 respectively. Apart from NAFLD, all other chronic liver disease aetiologies had an increased adAUROC compared with obAUROC. The adAUROC according to DANA was developed in chronic hepatitis C, but we predict that the fibrosis distribution for other liver diseases has a similar effect on AUROC.

The diagnostic performance of validated cut points for significant fibrosis, advanced fibrosis and cirrhosis were also evaluated for all causes of chronic liver disease. The cut points proposed in the original study of Hepascore were those most commonly validated in subsequent studies. A cut point of 0.50-0.55 had a summary sensitivity of 70%, summary specificity of 79%, an estimated PPV of 0.78 and an estimated NPV of 0.71 to predict significant fibrosis. A cut point of 0.50-0.61 had a summary sensitivity of 81%, a summary specificity of 74%, an estimated PPV of 0.54 and an estimated NPV of 0.91 to predict advanced fibrosis. A cut point of 0.80-0.84 had a summary sensitivity of 72%, a summary specificity of 0.88%, an estimated PPV of 0.47 and an estimated NPV of 0.91 to predict cirrhosis. These results were similar to those reported in the original study. Hepascore had the most accurate ability to excluded advanced fibrosis and cirrhosis with an estimated NPV of 0.91 for both.

Significant heterogeneity of obAUROC was found between all studies. This was heterogeneity was reduced but still remained significant after adjustment of the AUROC for DANA. This suggested that heterogeneity might be partly caused by the different distribution of fibrosis stages between studies. Subgroup analysis of disease aetiology found that heterogeneity of adAUROC was no longer significant within: HCV, HBV, HIV co-infection and ALD to predict significant fibrosis; in HIV co-infection to predict advanced fibrosis and in HCV, HBV, HIV co-infection to predict cirrhosis. This suggests that in addition to fibrosis distribution different aetiologies of chronic liver disease was another cause of heterogeneity between studies. Meta-regression analysis and sensitivity analysis was performed using seven pre-defined study characteristics. However, none of these characteristics showed a significant effect on AUROC. Others had previously found that blinded biopsy reviewing and histological staging system had an effect on the AUROC of other serum models [28]. The small number of studies that included in each subgroup made further analysis of the source of

heterogeneity impossible. Another limitation of this study was the potential bias that may have been a result of only including published full length articles to ensure adequate and comprehensive assessment of study quality.

In summary, this study confirmed that Hepascore is a useful measure of the severity of liver fibrosis in patients with the common causes of chronic liver disease. Hepascore had an excellent accuracy to exclude cirrhosis in all four causes of chronic liver disease and had good diagnostic performance for significant fibrosis and advanced fibrosis in chronic hepatitis C, chronic hepatitis B and alcoholic liver disease.

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Table 1. Characteristics of studies included in the analysis.

| Cohorts | year | aetiology | Country | centre | No. | mean age | male (%) | Staging system | F0 (%) | F1 (%) | F2 (%) | F3 (%) | F4 (%) | Mean biopsy size (mm) | Serum collection | Interval days | DANA |
|---------------------------|--------|-----------|-----------|---------------|------|----------|----------|----------------|--------|--------|--------|--------|--------|-----------------------|------------------|---------------|------|
| Adams -training [3] | 2005 | HCV | Australia | Single centre | 117 | 40 | 68 | Metavir | 20 | 37 | 25 | 13 | 6 | 13 | no fasting | - | 1.92 |
| Adams-validation [3] | 2005 | HCV | Australia | Multi centre | 104 | 41 | 73 | Metavir | 16 | 27 | 34 | 7 | 16 | 13 | no fasting | - | 2.07 |
| Bourliere [26] | 2008 | HCV | France | Multi centre | 467 | 47 | 59 | Metavir | 15 | 36 | 22 | 20 | 7 | 20 | no fasting | 0 | 2.00 |
| Cales [22], Boursier [17] | 2008-9 | HCV | France | Multi centre | 1056 | 46 | 60 | Metavir | 4 | 43 | 27 | 14 | 11 | 21 | no fasting | - | 1.79 |
| Becker [14] | 2009 | HCV | US | Single centre | 391 | 50 | 70 | Metavir | 16 | 34 | 15 | 16 | 19 | 16 | no fasting | <90 | 2.41 |
| Guéchet [13] | 2010 | HCV | France | Multi centre | 512 | 50 | 60 | Metavir | 7 | 45 | 18 | 15 | 15 | 25 | fasting | <60 | 2.06 |
| Lee [10] | 2010 | HCV | Australia | Single centre | 95 | 45 | 63 | Scheuer | - | - | - | - | - | 15 | no fasting | < 210 | - |
| Kalantari [25] | 2011 | HCV | Iran | Single centre | 80 | 35 | 85 | Metavir | 15 | 31 | 25 | 9 | 20 | - | no fasting | - | 2.23 |
| Crisan [21] | 2012 | HCV | Romania | Single centre | 446 | 49 | 38 | Metavir | 7 | 30 | 36 | 18 | 9 | 11 | fasting | 0 | 1.76 |
| Leroy [24] | 2014 | HCV | France | Single centre | 255 | 47 | 57 | Metavir | 15 | 38 | 26 | 11 | 11 | 24 | fasting | 0 | 1.97 |
| Cacoub [7] | 2008 | HCV/HIV | France | Multi centre | 272 | 40 | 72 | Metavir | 0 | 25 | 40 | 25 | 10 | 19 | fasting | - | 1.60 |
| Cales [9] | 2010 | HCV/HIV | France | Multi centre | 169 | 41 | 65 | Metavir | 8 | 26 | 33 | 13 | 20 | 25 | fasting | <90 | 2.03 |
| Wu [20] | 2010 | HBV | China | Single centre | 78 | 33 | 84 | Metavir | 17 | 42 | 17 | 13 | 12 | 18 | no fasting | - | 2.16 |
| Raftopoulos [27] | 2011 | HBV | Australia | Multi centre | 179 | 42 | 71 | Metavir | 15 | 43 | 20 | 14 | 8 | 21 | no fasting | <180 | 1.99 |
| Basar [19] | 2013 | HBV | Turkey | Single centre | 76 | 45 | 45 | Metavir | 12 | 21 | 29 | 21 | 17 | - | no fasting | 0 | 2.18 |
| Leroy [24] | 2014 | HBV | France | Single centre | 255 | 40 | 72 | Metavir | 15 | 38 | 26 | 11 | 11 | 25 | fasting | 0 | 1.97 |
| Bottero [18] | 2009 | HBV/HIV | France | Multi centre | 108 | 42 | 90 | Metavir | 10 | 33 | 26 | 16 | 15 | 17 | no fasting | <180 | 2.04 |
| Naveau [15] | 2008 | ALD | France | Single centre | 218 | 47 | 78 | Metavir | 7 | 30 | 22 | 10 | 31 | 15 | no fasting | <30 | 2.34 |
| Nguyen-Khac [12] | 2008 | ALD | France | Single centre | 103 | 53 | 74 | Metavir | 8 | 17 | 23 | 19 | 32 | 12 | no fasting | 0 | 2.42 |
| Adams [16] | 2011 | NAFLD | Australia | Multi centre | 242 | 47 | 60 | Metavir | 36 | 24 | 18 | 12 | 10 | - | no fasting | 0 | 2.38 |
| Boursier [8] | 2009 | Mixed | France | Multi centre | 390 | 52 | 68 | Metavir | 7 | 18 | 23 | 20 | 31 | - | no fasting | - | 2.39 |
| Costelloe [11] | 2015 | Mixed | UK | Single centre | 73 | 51 | 62 | Metavir | 44 | 11 | 12 | 3 | 30 | - | no fasting | 0 | 3.19 |

Table 2. Quality assessment of included studies.

| Author, year | Q1 | Q2 | Q3 | Q4 | Q5 | Q6 | Q7 | Q8 | Q9 | Q10 | Q11 | Q12 | Q13 | Q14 |
|------------------------|---------------|--------------------|--------------------------------|--------------------------|---------------------------|--------------------------------|--------------------|------------------------|-----------------------------|------------------|------------------------|----------------------|----------------------|-------------|
| | Spectrum bias | Selection criteria | Appropriate reference standard | Disease progression bias | Partial verification bias | Differential verification bias | Incorporation bias | Test execution details | Reference execution details | Test review bias | Diagnostic review bias | Clinical review bias | Intermediate results | withdrawals |
| Adams, 2005[3] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Cales, 2008 [22] | yes | yes | yes | unclear | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes |
| Bourliere, 2008 [26] | yes | yes | unclear | yes | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes |
| Boursier, 2009 [17] | yes | yes | yes | unclear | yes | yes | yes | unclear | yes | yes | unclear | yes | yes | yes |
| Becker, 2009 [14] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Guéchet, 2010 [13] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Lee, 2010 [10] | yes | unclear | unclear | yes | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes |
| Kalantari, 2011 [25] | yes | no | yes | unclear | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes |
| Crisan, 2012 [21] | yes | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes | yes | yes | yes |
| Leroy, 2014 [24] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Cacoub, 2008 [7] | yes | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes | yes | yes | yes |
| Cales, 2010 [9] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Wu, 2010 [20] | yes | yes | unclear | unclear | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes |
| Raftopoulos, 2011 [27] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Basar, 2013 [19] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Bottero, 2009 [18] | yes | unclear | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes | yes | yes |
| Naveau, 2008 | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Nguyen-Khac, 2008 [12] | no | yes | yes | yes | yes | yes | yes | unclear | yes | yes | yes | yes | yes | yes |
| Adams, 2011 [16] | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |
| Boursier, 2009 [8] | no | unclear | yes | yes | yes | yes | yes | unclear | yes | yes | unclear | yes | yes | yes |
| Costelloe, 2015 [11] | no | no | unclear | unclear | yes | yes | yes | yes | yes | yes | yes | yes | yes | yes |

Table 3. Validated cut points of Hepascore by included studies.

| Cohorts | Year | Aetiology | Significant fibrosis | | | Advanced fibrosis | | | Cirrhosis | | |
|---------------------------|--------|-----------|----------------------|------|------|-------------------|------|------|-----------|------|------|
| | | | Cut point | Sen | Spe | Cut point | Sen | Spe | Cut point | Sen | Spe |
| Adams -training [3] | 2005 | HCV | 0.5 | 0.67 | 0.92 | 0.50 | 0.95 | 0.81 | 0.84 | 0.71 | 0.84 |
| Adams-validation [3] | 2005 | HCV | 0.5 | 0.63 | 0.89 | 0.50 | 0.88 | 0.74 | 0.84 | 0.71 | 0.89 |
| Bourliere [26] | 2008 | HCV | 0.5 | 0.63 | 0.86 | - | - | - | 0.84 | 0.71 | 0.88 |
| Cales [22], Boursier [17] | 2008-9 | HCV | 0.47 | 0.66 | 0.79 | 0.50 | 0.82 | 0.71 | 0.80 | 0.80 | 0.83 |
| Becker [14] | 2009 | HCV | 0.55 | 0.82 | 0.65 | 0.80 | 0.72 | 0.77 | - | - | - |
| Guéchet [13] | 2010 | HCV | 0.5 | 0.77 | 0.70 | 0.60 | 0.80 | 0.70 | 0.75 | 0.86 | 0.74 |
| Lee [10] | 2010 | HCV | 0.5 | 0.77 | 0.79 | - | - | - | - | - | - |
| Kalantari [25] | 2011 | HCV | 0.34 | 0.67 | 0.56 | 0.61 | 0.82 | 0.86 | 0.84 | 1.00 | 0.97 |
| Crisan [21] | 2012 | HCV | 0.34 | 0.57 | 0.72 | 0.61 | 0.61 | 0.73 | - | - | - |
| Leroy [24] | 2014 | HCV | 0.5 | 0.52 | 0.85 | 0.47 | 0.79 | 0.85 | 0.84 | 0.60 | 0.89 |
| Cales [9] | 2010 | HCV/HIV | 0.31 | 0.90 | 0.59 | - | - | - | - | - | - |
| Wu [20] | 2010 | HBV | 0.5 | 0.88 | 0.50 | - | - | - | - | - | - |
| Raftopoulos [27] | 2011 | HBV | 0.52 | 0.77 | 0.76 | 0.72 | 0.68 | 0.85 | 0.88 | 0.87 | 0.86 |
| Basar [19] | 2013 | HBV | 0.32 | 0.78 | 0.68 | 0.50 | 0.71 | 0.79 | 0.52 | 0.85 | 0.75 |
| Leroy [24] | 2014 | HBV | 0.5 | 0.42 | 0.84 | 0.42 | 0.75 | 0.71 | 0.84 | 0.56 | 0.92 |
| Bottero [18] | 2009 | HBV/HIV | 0.48 | 0.67 | 0.68 | 0.76 | 0.72 | 0.87 | 0.9 | 0.80 | 0.89 |
| Naveau [15] | 2008 | ALD | 0.25 | 0.90 | 0.37 | - | - | - | 0.97 | 0.90 | 0.87 |
| Adams [16] | 2011 | NAFLD | 0.44 | 0.51 | 0.88 | 0.37 | 0.76 | 0.84 | 0.7 | 0.87 | 0.89 |
| Costelloe [11] | 2015 | Mixed | - | - | - | 0.99 | 0.79 | 0.74 | - | - | - |

Table 4. Sensitivity analysis of study characteristics.

| Characteristic | Significant fibrosis | | Advanced fibrosis | | Cirrhosis | |
|--------------------|----------------------|---------------------|-------------------|---------------------|------------------|---------------------|
| | No. | AdAUROC (95% CI) | No. | AdAUROC (95% CI) | No. | AdAUROC (95% CI) |
| | All | 19 | 0.83 (0.81-0.85) | 16 | 0.89 (0.85-0.92) | 15 |
| Patients inclusion | | | | | | |
| single centre | 9 | 0.82 (0.80-0.85) | 8 | 0.87 (0.81-0.94) | 6 | 0.92 (0.87-0.97) |
| multicentre | 10 | 0.83 (0.81-0.86) | 8 | 0.89 (0.87-0.91) | 9 | 0.93 (0.91-0.96) |
| Mean biopsy length | | | | | | |
| <20 mm | 9 | 0.82 (0.79-0.85) | 7 | 0.90 (0.84-0.97) | 6 | 0.94 (0.90-0.98) |
| ≥20 mm | 7 | 0.85 (0.84-0.87) | 6 | 0.93 (0.88-0.89) | 6 | 0.95 (0.93-0.97) |
| Biopsy evaluation | | | | | | |
| blinded | 15 | 0.83 (0.81-0.84) | 13 | 0.87 (0.83-0.91) | 12 | 0.93 (0.91-0.96) |
| not blinded | 4 | 0.86 (0.84-0.88) | 3 | 0.93 (0.88-0.98) | 3 | 0.92 (0.85-0.99) |
| Interval time | | | | | | |
| ≤1 month | 9 | 0.81 (0.78-0.84) | 8 | 0.86 (0.83-0.89) | 7 | 0.93 (0.90-0.95) |
| > 1 month | 10 | 0.85 (0.83-0.87) | 8 | 0.91 (0.87-0.96) | 7 | 0.94 (0.90-0.98) |
| Serum collection | | | | | | |
| fasting | 6 | 0.82 (0.79-0.85) | 5 | 0.86 (0.82-0.90) | 4 | 0.92 (0.90-0.95) |
| non-fasting | 13 | 0.84 (0.82-0.86) | 11 | 0.90 (0.86-0.94) | 11 | 0.93 (0.90-0.96) |
| HA test kit | | | | | | |
| Corgenix | 11 | 0.84 (0.81-0.86) | 9 | 0.91 (0.88-0.94) | 9 | 0.96 (0.94-0.97) |
| other kits | 4 | 0.85 (0.82-0.88) | 4 | 0.86 (0.76-0.96) | 3 | 0.92 (0.90-0.95) |
| QUADAS score | | | | | | |
| all scored yes | 10 | 0.84 (0.81-0.86) | 7 | 0.90 (0.85-0.95) | 8 | 0.94 (0.91-0.96) |
| not all scored yes | 9 | 0.82 (0.80-0.85) | 9 | 0.87 (0.83-0.92) | 7 | 0.92 (0.88-0.96) |

FIGURE LEGENDS:

Figure 1. ObAUROC and adAUROC of Hepascore to predict significant fibrosis.

Figure 2. SROC curves of Hepascore. (A): SROC curve for significant fibrosis. (B): SROC curve for advanced fibrosis. (C): SROC curve for cirrhosis.

Figure 3. ObAUROC and adAUROC of Hepascore to predict advanced fibrosis.

Figure 4. ObAUROC and adAUROC of Hepascore to predict cirrhosis.





