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Title
**N-terminal Propeptide of Type 3 Collagen-Based Sequential
Algorithm can identify high-risk steatohepatitis and fibrosis in
MAFLD**

Short Title: Sequential algorithm, steatohepatitis and fibrosis staging

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8

9 **Abbreviation list:**

10 ALT, alanine aminotransferase; AST, aspartate aminotransferase; AAR, aspartate

11 aminotransferase to alanine aminotransferase ratio; APRI, aspartate aminotransferase-

12 platelet ratio index; AUROC, area under the receiver operating characteristic curve;

13 BMI, body mass index; CI, 95% confidence interval; DCA, decision curve analysis;

14 FIB-4, fibrosis-4; FLIP, fatty liver inhibition of progression; MAFLD, metabolic

15 dysfunction-associated fatty liver disease; NFS, NAFLD fibrosis score; NPV, negative

16 predictive value; PPV, positive predictive value; PRO-C3, N-terminal propeptide of

17 type 3 collagen; VCTE, Vibration-Controlled Transient Elastography.

18

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10 All authors contributed to the manuscript for important intellectual content and
11 approved the final submission of the manuscript.

12

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14 Although the PRO-C3 ELISA test was carried out at Nordic Bioscience under a
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16 analysis, and manuscript drafting were carried out independently of the Nordic
17 Bioscience team. ADAPT has not been developed as a proprietary test. The PRO-C3
18 ELISA test is not currently commercially available, but can be obtained as a Nordic
19 Bioscience research test for research use only. Diana Julie Leeming and Morten
20 Karsdal are employed by, and own stock at Nordic Bioscience. Grace Lai-Hung Wong
21 and Vincent Wai-Sun Wong have served as speakers and/or consultants for Echosens.
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1 **Abstract**

2 **Background & Aims:** With metabolic dysfunction-associated fatty liver disease
3 (MAFLD) incidence and prevalence sharply increasing globally, there is an urgent
4 need for non-invasive diagnostic tests to accurately screen high-risk MAFLD patients
5 for liver inflammation and fibrosis. We aimed to develop a novel sequential algorithm
6 based on N-terminal propeptide of type 3 collagen (PRO-C3) for disease risk
7 stratification in patients with MAFLD.

8 **Methods:** A derivation and independent validation cohort of 327 and 142 patients
9 with biopsy-confirmed MAFLD were studied. We compared the diagnostic
10 performances of various non-invasive scores in different disease states, and a novel
11 sequential algorithm was constructed by combining the best performing non-invasive
12 scores.

13 **Results:** For patients with high-risk progressive steatohepatitis (i.e., steatohepatitis +
14 NAFLD activity score \geq 4 + F \geq 2), the AUROC of FAST score was 0.801 (95%
15 confidence interval (CI): 0.739-0.863), and the negative predictive value (NPV) was
16 0.951. For advanced fibrosis (\geq F3) and cirrhosis (F4), the AUROCs of ADAPT and
17 Agile 4 were 0.879 (95%CI: 0.825-0.933) and 0.943 (95%CI: 0.892-0.994), and the
18 NPV were 0.972 and 0.992. Sequential algorithm of ADAPT+Agile 4 combination
19 was better than other combinations for risk stratification of patients with severe
20 fibrosis (AUROC=0.88), with similar results in the validation cohort. Meanwhile, in
21 all subgroup analyses (stratifying by sex, age, diabetes, NAS, BMI and ALT),
22 ADAPT+Agile 4 had a good diagnostic performance.

1 **Conclusions:** The new sequential algorithm reliably identifies liver inflammation and
2 fibrosis in MAFLD, making it easier to exclude low-risk patients and recommending
3 high-risk MAFLD patients for clinical trials and emerging pharmacotherapies.

4

5 **Keywords:**

6 Metabolic dysfunction-associated fatty liver disease; Steatohepatitis; Sequential
7 algorithm; Fibrosis staging

8

1 **Introduction**

2 Metabolic dysfunction-associated fatty liver disease (MAFLD) affects up to nearly
3 30% of the adult population worldwide and its prevalence has reached more than 70%
4 in patients of severe obesity and type 2 diabetes [1-3]. The newly proposed definition
5 of MAFLD has been endorsed by global multi-stakeholders [4]. MAFLD has created
6 a huge clinical and economic burden and has become the most common global cause
7 of chronic liver diseases [5-7]. Although only a small fraction of patients with
8 MAFLD will progress to advanced fibrosis or cirrhosis [8, 9], the absolute number is
9 huge given the large number of MAFLD patients worldwide [10, 11]. Liver fibrosis,
10 in particular F3-4 fibrosis, has a strong association with adverse liver-related
11 prognosis [12, 13]. Therefore, it is important to accurately identify high-risk patients
12 with MAFLD to guide monitoring and treatment.

13
14 Presently, the diagnosis of active liver inflammation and progressive fibrosis mainly
15 relies on serum biomarkers, imaging methods, non-invasive scores and/or liver biopsy
16 [14]. The first three methodologies are the main screening tests that are widely used in
17 clinical practice [15]. Although these tests have a high negative predictive value, they
18 also have a high false positive rate, and they only target one disease state (such as the
19 degree of steatosis, inflammation, or the stage of fibrosis) [16-18], so that they cannot
20 assess the overall condition of the liver. Nevertheless, liver biopsy is the gold standard
21 for pathological evaluation of MAFLD, but its limitations mean that it is being used
22 less frequently in clinical practice [19, 20]. The current key challenge is how to

1 maximize the advantages of the first three aforementioned non-invasive diagnostic
2 methods, in order to assess liver inflammation and fibrosis in patients with MAFLD.
3
4 Based on the N-terminal propeptide of type 3 collagen (PRO-C3) and ADAPT score
5 that have good performance in fibrosis staging in Asian MAFLD patients [21, 22], we
6 performed a comparative analysis of several up-to-date non-invasive scores (i.e.
7 FAST, Agile 3+ and Agile 4) with the specific aim of developing a new combined
8 diagnostic model to screen high-risk patients with MAFLD [23, 24], in order to more
9 conveniently and accurately assess the overall liver status in MAFLD patients.
10 Moreover, the development of new sequential algorithms may optimize the positive
11 predictive value to increase the diagnosis rate, and reduce the number of patients with
12 indeterminate results who may need liver biopsy.

13

14 **Methods**

15 *The derivation and validation cohorts*

16 The derivation liver biopsy cohort included two Asian cohorts with a total of 327
17 adult patients aged 18-70 years with biopsy-proven MAFLD between January 2017
18 and December 2020 (i.e., 262 patients from Wenzhou and 65 patients from Hong
19 Kong). The Wenzhou and Hong Kong cohorts included patients from previously
20 published studies but also included additional patients with MAFLD [21, 22]. The
21 independent validation liver biopsy cohort included 142 MAFLD patients from
22 Wenzhou between June 2021 and December 2021. MAFLD was diagnosed by the

1 presence of hepatic steatosis on liver histology (defined as presence of more than 5%
2 of steatotic hepatocytes) with at least one of the following three coexisting metabolic
3 conditions, i.e., overweight/obesity, type 2 diabetes mellitus, or metabolic
4 dysregulation [25]. In both selected cohorts of the study, we included only patients
5 with a single aetiologic MAFLD, as we excluded those with excessive alcohol
6 consumption (> 10 g per day for women and > 20 g per day for men), autoimmune
7 liver diseases, chronic viral hepatitis (hepatitis B or C), secondary fatty liver, or prior
8 history of malignancy in the past two years [26]. Patients who have not undergone
9 vibration-controlled transient elastography (VCTE) measurement were also excluded
10 from the study.

11

12 We recorded clinical information within one day of liver biopsy examination. Fasting
13 venous blood samples for measurement of serum liver enzymes, total bilirubin,
14 albumin, lipids and other blood biochemical tests were obtained. Height and weight
15 were measured for the calculation of body mass index (BMI) = weight [kg]/height²
16 [m²]. Diabetes was defined as fasting glucose levels ≥ 7.0 mmol/L (≥ 126 mg/dL),
17 haemoglobin A1c $\geq 6.5\%$ (≥ 48 mmol/mol), a previous diagnosis of diabetes, or the
18 use of any anti-hyperglycaemic drugs [27]. The study was approved by the
19 institutional review boards of both centres and all participants gave their written
20 informed consent [21, 22].

21

22 ***Liver stiffness measurement***

1 Measurement of liver stiffness (LSM) and steatosis using VCTE (FibroScan equipped
2 with CAP software; Echosens, Paris, France) was undertaken within a week of the
3 diagnostic liver biopsy. The FibroScan M probe was used in the Wenzhou cohort, and
4 the M and XL probes were used in the Hong Kong cohort with BMI < 30 and ≥ 30
5 kg/m². Ten or more measurements and an interquartile range/median ratio of liver
6 stiffness ≤ 0.3 were required for valid LSMs.

7

8 ***Liver histological assessment of MAFLD***

9 Eligible patients with MAFLD for liver biopsy were aged 18 or older, able to give
10 written informed consent, and scheduled (independently from this study) to have a
11 liver biopsy within 2 weeks before or after VCTE measurements for investigation of
12 suspected MAFLD (usually due to elevated serum liver enzyme levels and hepatic
13 steatosis on ultrasound). Ultrasound-guided percutaneous liver biopsy was performed
14 using a 16G Tru Cut needle and tissue was stained with haematoxylin-eosin and
15 Masson trichrome. The reading and scoring of biopsy specimens were performed by
16 an experienced liver histopathologist in each centre, who was blinded to the clinical
17 and biochemical data of participants. The NAFLD activity score (NAS) was
18 calculated as the sum of hepatic steatosis, lobular inflammation and hepatocyte
19 ballooning [19, 28, 29]. Presence of steatohepatitis was diagnosed using the fatty liver
20 inhibition of progression (FLIP) definition (at least grade one for steatosis,
21 ballooning, and lobular inflammation) [30]. The histologic stages of liver fibrosis
22 were defined as: stage 0, no fibrosis; stage 1, peri-sinusoidal or portal venular fibrosis;

1 stage 2, peri-sinusoidal and portal vein/periportal fibrosis; stage 3, bridging fibrosis;
 2 and stage 4, cirrhosis [27]. Significant fibrosis was defined as fibrosis stage \geq F2, and
 3 advanced fibrosis was defined as fibrosis stage \geq F3.

5 *Non-invasive scoring tests*

6 The formation of N-terminal propeptide of type 3 collagen (PRO-C3) was assessed in
 7 serum using the PRO-C3 competitive ELISA kit from Nordic Bioscience, Herlev,
 8 Denmark, as previously described [31]. It has been established that the parameters
 9 that have a significant contribution to predicting the severity of liver fibrosis were
 10 liver stiffness measurement (LSM), aspartate aminotransferase-to-alanine
 11 aminotransferase ratio (AAR), platelet count (PLT), sex and diabetes status. After
 12 including the above-mentioned parameters, the **Agile 3+** and **Agile 4** algorithms
 13 included (diabetes status: yes = 1, no = 0 and sex: male = 1, female = 0) follow the
 14 two subsequent equations:

$$15 \quad \text{Agile 3+} = \frac{e^{\text{logit}(p_{F \geq 3})}}{1 + e^{\text{logit}(p_{F \geq 3})}}$$

16 with $e^{\text{logit}(p_{F \geq 3})} = -3.92368 + 2.29714 \times \ln(\text{LSM}) - 0.00902 \times \text{PLT} - 0.98633 \times$
 17 $\text{AAR}^{-1} + 1.08636 \times \text{Diabetes status} - 0.38581 \times \text{Sex} + 0.03018 \times \text{Age}.$

$$18 \quad \text{Agile 4} = \frac{e^{\text{logit}(p_{F=4})}}{1 + e^{\text{logit}(p_{F=4})}}$$

19 with $e^{\text{logit}(p_{F=4})} = 7.50139 - 15.42498 \times \frac{1}{\sqrt{\text{LSM}}} - 0.01378 \times \text{PLT} - 1.41149 \times \text{AAR}^{-1} -$
 20 $0.53281 \times \text{Sex} + 0.41741 \times \text{Diabetes status}.$

21
 22 Additionally, we used multivariate logistic regression to create a novel non-invasive

1 score to identify patients with advanced fibrosis in MAFLD, which including PRO-
2 C3, LSM, platelet levels and diabetes status.

$$3 \quad \text{MLR Model} = -3.89497 + 0.09313 \times \text{PRO} - \text{C3} + 0.20373 \times \text{LSM} \\ 4 \quad \quad \quad - 0.00603 \times \text{Platelets} + 1.29126 \times \text{Diabetes status}$$

5 Other widely used non-invasive scores of fibrosis, namely FibroScan-AST (FAST)
6 score [24], ADAPT score [31], APRI [32], FIB-4 [33], NAFLD fibrosis score (NFS)
7 [34], and BARD (BMI, AAR, diabetes) score [35] were calculated using available
8 clinical and laboratory variables.

9

10 ***Statistical analysis***

11 According to their normal or not normal distribution, continuous data were reported
12 as means and standard deviation or medians (interquartile ranges), respectively.

13 Categorical data were reported as a numbers (or percentages). The overall diagnostic
14 accuracy of the various non-invasive scores was evaluated by the receiver operator
15 characteristic (ROC) curve analysis and expressed as the area under the receiver
16 operator characteristic curve (AUROC). The diagnostic accuracies of the FAST score,
17 the ADAPT score, the Agile 3+ and Agile 4 algorithms, and other non-invasive
18 fibrosis scores were determined by calculating the sensitivity, specificity, positive
19 predictive value (PPV), negative predictive value (NPV), as well as positive and
20 negative likelihood ratios (LRs).

21

22 The DeLong test was performed on the AUROC curves by a bootstrap re-sampling

1 method with 500 repetitions. The dual cut-off value improves the diagnostic certainty
2 of screening high-risk groups of fibrosis. The curve with the greatest probability was
3 the best decision strategy to maximize the new advantage. We conducted a decision
4 curve analysis (DCA) to evaluate which combination of the two algorithms is more
5 advantageous, that is, whether the new decision tree curve was better at identifying the
6 severity of fibrosis rather than harmful (screening and deselecting the population that
7 are in most need of a liver biopsy, reducing the risk of an unnecessary invasive
8 inspection). All statistical analyses were performed by R version 3.6.1 ([https://www.r-
9 project.org/](https://www.r-project.org/)).

10

11 **Results**

12 *Patient Characteristics*

13 The clinical and biochemical characteristics of Asian patients with biopsy-proven
14 MAFLD in the derivation (n=327) and validation (n=142) cohorts are shown in **Table**
15 **1**. Most clinical and laboratory parameters, as well as serum PRO-C3 levels, FAST
16 score, ADAPT score, Agile 3+ score and Agile 4 score did not differ significantly
17 between the two cohorts of patients. Compared to those in the validation cohort,
18 patients in the derivation cohort had a lower prevalence of steatohepatitis (56%
19 vs.74%, $P < 0.001$), but a comparable prevalence of advanced fibrosis (13% vs.8%, $P =$
20 0.125) and $NAS \geq 4$ (51% vs.58%, $P = 0.158$) on liver histology.

21

22 *Diagnostic accuracy of the FAST score and other widely used non-invasive fibrosis*

1 *scores for steatohepatitis + NAS ≥ 4 + F ≥ 2*

2 In the derivation cohort, the FAST score was compared with the APRI, FIB-4, NFS

3 and BARD scores for the identification of patients with high-risk progressive

4 steatohepatitis, which was defined as presence of steatohepatitis + NAS ≥ 4 + F ≥ 2 .

5 **Supplementary Figure 1A** shows the diagnostic accuracy of the FAST and other

6 non-invasive scores. The AUROC of the FAST score was 0.801 (95% CI: 0.739-

7 0.863) that was higher than that of the other widely used non-invasive scores: 0.687

8 (95% CI: 0.604-0.770, P< 0.001) for APRI, 0.594 (95% CI: 0.506-0.683, P< 0.001)

9 for FIB-4, 0.625 (95% CI: 0.543-0.707, P< 0.001) for NFS, and 0.640 (95% CI:

10 0.565-0.714, P= 0.029) for BARD, respectively. In our cohort the cut-off value for

11 sensitivity ($\geq 90\%$) was 0.285 and for specificity ($\geq 90\%$) was 0.605, respectively.

12 Using a dual cut-off approach (**Supplementary Figure 2A**), we found that patients

13 with FAST score < 0.285 were low-risk groups (NPV= 0.951); patients with FAST

14 score > 0.605 were likely to have progressive steatohepatitis, while 122 (37.3%) of

15 327 patients were in the 'gray zone' between the two FAST cut-off values. The FAST

16 score showed a good diagnostic performance between the validation and derivation

17 cohort (AUROC= 0.742, 95% CI: 0.617-0.868, NPV= 0.964), and 58 (40.8%) patients

18 were in the 'gray zone' (**Supplementary Table 1**).

19

20 *Diagnostic performances of novel scores and other widely used non-invasive*

21 *fibrosis scores for advanced fibrosis and cirrhosis*

22 Based on the previously observed strong association between serum PRO-C3 levels

1 and the histological severity of liver fibrosis [21], we tested the diagnostic
2 performance of the ADAPT score for identifying advanced fibrosis in the derivation
3 cohort and compared it with Agile 3+ score and other widely used serum-based non-
4 invasive scores (i.e., APRI, FIB-4, BARD and NFS scores). As shown in the
5 **Supplementary Table 2**, the AUROC of the ADAPT score was 0.879 (95% CI:
6 0.825-0.933, NPV= 0.972) for advanced fibrosis, better than Agile 3+ score
7 (AUROC=0.805, 95% CI: 0.725-0.886, P= 0.015, NPV= 0.952) and Agile 4 score
8 (AUROC=0.826, 95% CI: 0.747-0.904, P= 0.019, NPV= 0.97), also higher than that
9 of the other widely used non-invasive scores: 0.642 (95% CI: 0.550-0.734, P< 0.001)
10 for APRI, 0.732 (95% CI: 0.643-0.820, P< 0.001) for FIB-4, 0.800 (95% CI: 0.723-
11 0.877, P< 0.001) for NFS, and 0.709 (95% CI: 0.632-0.786, P< 0.001) for BARD,
12 respectively (**Supplementary Figure 1B**). The ADAPT score also performed best in
13 the validation cohort (AUROC=0.885, 95% CI: 0.772-0.998, NPV= 0.988) compared
14 to other non-invasive scores (see the **Supplementary Table 2**). We also used the dual
15 cut-offs approach for ruling in and ruling out MAFLD patients with advanced fibrosis
16 (**Supplementary Figure 2B**).

17
18 Furthermore, the AUROC for the Agile 4 score was 0.943 (95% CI: 0.892-0.994) for
19 cirrhosis (n= 18), 0.909 (95% CI: 0.843-0.974, P= 0.169) for the Agile 3+ score,
20 0.582 (95% CI: 0.438-0.727, P< 0.001) for the APRI score, 0.794 (95% CI: 0.685-
21 0.903, P= 0.006) for the FIB-4 score, 0.905 (95% CI: 0.849-0.961, P= 0.113) for NFS,
22 and was 0.801 (95% CI: 0.720-0.882, P= 0.134) for BARD in the derivation cohort

1 (Supplementary Figure 1C). In view of the excellent diagnostic performance of the
2 Agile 4 score in patients with cirrhosis, we used the dual cut-offs (Supplementary
3 Figure 2C) to identify patients with advanced fibrosis. Patients with Agile 4 values <
4 0.318 ($\geq 90\%$ sensitivity) did not have cirrhosis, and patients with Agile 4 values >
5 0.586 ($\geq 90\%$ specificity) were likely to have cirrhosis. Due to the small number of
6 patients with cirrhosis (F4) in the validation group (n= 4), the comparison of
7 diagnostic performance lacks confidence, but on the whole cohort, Agile 4 score still
8 performed well (Supplementary Table 3).

9

10 ***Combination of Agile 4 and other scores in a sequential algorithm improves the***
11 ***diagnostic accuracy of advanced fibrosis***

12 We compared combinations of multiple sequential algorithms, including MLR Model,
13 PRO-C3, VCTE, and other clinical and hematological parameters. The combination of
14 ADAPT + Agile 4 showed the best diagnostic performance, with an AUROC of 0.880
15 (95% CI: 0.824-0.935) and accuracy of 88.1% (NPV= 0.955) that was better than
16 MLR Model (AUROC=0.876, 95% CI: 0.817-0.935, accuracy of 85.3%) and ADAPT
17 + LSM (AUROC=0.879, 95% CI: 0.821-0.936, accuracy of 84.4%), and also higher
18 than that of the other widely used non-invasive scores. At the same time, a
19 requirement for liver biopsy in only 31.4% of patients (gray zone) and the specificity
20 was the highest (90.5%) (Supplementary Table 4). Taking advantages of these
21 parameters (VCTE can be performed on an outpatient basis and hematological
22 parameters can be obtained on the day of admission at no additional cost), sequential

1 combinations of non-invasive scores were used to assess the overall liver status of
2 patients with MAFLD.

3

4 Based on the performance of the FIB-4 and NFS scores for diagnosing advanced

5 fibrosis, we further evaluated the performance of the sequential combination

6 algorithm of ADAPT, Agile 3+, FIB-4, NFS scores and Agile 4 score for the diagnosis

7 of advanced fibrosis. The best performance was obtained by using ADAPT + Agile 4

8 (AUROC= 0.880, 95% CI: 0.824-0.935), which was better than combinations of the

9 other sequential predictive algorithms: 0.823 (95% CI: 0.745-0.901, P= 0.009) for

10 (Agile 3+) + Agile 4, 0.785 (95% CI: 0.703-0.866, P= 0.002) for FIB-4 + Agile 4, and

11 0.812 (95% CI: 0.736-0.889, P= 0.002) for NFS + Agile 4, respectively (**Figure 1A**).

12 The diagnostic accuracy of this sequential algorithm that utilized the ADAPT + Agile

13 4 scores for the detection of advanced fibrosis was also superior to replacing ADAPT

14 by any of Agile 3+, FIB-4, or NFS, with a diagnostic accuracy of 88.1%, compared

15 with 86.9%, 85.9%, and 80.7%, respectively. The same results were also confirmed in

16 the validation group (AUROC= 0.905, 95% CI: 0.788-1.000, NPV= 0.984), ADAPT

17 + Agile 4 could best rule out advanced fibrosis in MAFLD (see the **Table 2**).

18

19 The decision curve analysis (DCA) used for assessing the sequential algorithms is

20 presented in **Figure 1B**. This figure analyzes the clinical utility of the ADAPT + Agile

21 4 sequential algorithm compared with (Agile 3+) + Agile 4, FIB-4 + Agile 4, and NFS

22 + Agile 4 in identifying patients with advanced fibrosis ($F \geq 3$). DCA showed that

1 from a threshold probability > 10%, we could obtain more net benefit by using a
2 variance graph. In particular, if a patient's threshold probability was > 10% and <
3 70%, the use of the ADAPT + Agile 4 score for predicting the risk of advanced
4 fibrosis had more benefits than the reference strategy (using Agile 3+, FIB-4, or NFS
5 scores combined with Agile 4 score).

6

7 We combined the algorithms of ADAPT (lower value of 3.705 and upper value of
8 4.93) and Agile 4 (lower value of 0.318 and upper value of 0.586), and then compared
9 it with other combinations. This new combination algorithm approach could more
10 accurately deselect patients with high-risk steatohepatitis (i.e., the number of true
11 positive (TP) groups has increased from 18%, 21% and 31% to 32%), while reducing
12 the number of patients requiring liver biopsy (including false positive (FP)), false
13 negative (FN) and indeterminate groups from 34%, 32% and 26% to 20%) (**Figure 2**).

14

15 *Diagnostic performance of ADAPT + Agile 4 in patient subgroups*

16 Circulating levels of hyaluronic acid, type III procollagen, type IV collagen, and
17 laminin were also used as routine liver fibrosis tests in the Wenzhou cohort. The best
18 performance was still obtained by combining ADAPT + Agile 4 (AUROC= 0.761,
19 95% CI: 0.657-0.864), which was better than established fibrosis markers and
20 combinations of other sequential algorithms: 0.508 (95% CI: 0.335-0.682) for
21 hyaluronic acid, 0.661 (95% CI: 0.506-0.815) for type III procollagen, 0.635 (95% CI:
22 0.472-0.798) for type IV collagen, 0.570 (95% CI: 0.437-0.703) for laminin, 0.638

1 (95% CI: 0.468-0.808) for (Agile 3+) + Agile 4, 0.642 (95% CI: 0.480-0.804) for
2 FIB-4 + Agile 4, and 0.663 (95% CI: 0.499-0.825) for NFS + Agile 4, respectively. In
3 particularly, the ADAPT + Agile 4 combination algorithm had 100% sensitivity and
4 100% NPV for excluding advanced fibrosis in the Wenzhou cohort (see the
5 **Supplementary Table 5**).

6

7 The ADAPT + Agile 4 combination algorithm had a good diagnostic performance for
8 advanced fibrosis among different subgroups of MAFLD patients. In men, the
9 AUROC of this sequential algorithm was 0.845 (95% CI: 0.771-0.919), and among
10 women the AUROC was 0.953 (95% CI: 0.907-0.999). Similarly, in different patient
11 subgroups, such as those with age < 44 years (AUROC=0.822, 95% CI: 0.726-0.917),
12 age ≥ 44 years (AUROC=0.894, 95% CI: 0.829-0.958), NAS < 4 (AUROC=0.889,
13 95% CI: 0.794-0.984), NAS ≥ 4 (AUROC=0.880, 95% CI: 0.814-0.945), BMI < 25
14 kg/m² (AUROC=0.914, 95% CI: 0.804-1.000), BMI ≥ 25 kg/m² (AUROC=0.865,
15 95% CI: 0.798-0.931), ALT < 40 U/L (AUROC=0.821, 95% CI: 0.701-0.942), ALT ≥
16 40 U/L (AUROC=0.915, 95% CI: 0.858-0.972), presence of diabetes
17 (AUROC=0.871, 95% CI: 0.796-0.946) or absence of diabetes (AUROC=0.769, 95%
18 CI: 0.660-0.878), the ADAPT + Agile 4 combination algorithm showed comparable
19 good performances for predicting advanced fibrosis in patients with MAFLD (**Figure**
20 **3**).

21

22 ***The combined hierarchical screening using FAST, combined with ADAPT and***

1 *Agile 4 significantly improves the diagnostic accuracy of fibrosis staging*

2 We utilized three non-invasive scores that had good performance in different stages of
3 liver fibrosis, and developed a combined diagnostic method for staging the severity of
4 fibrosis in MAFLD patients. Firstly, calculating the FAST score with FibroScan, we
5 propose that patients with a FAST score < 0.605 could undergo routine follow-up
6 (NPV= 0.9). Then, for patients with higher FAST scores who may have progressive
7 steatohepatitis, those with an ADAPT value < 4.93 or in the gray area need to confirm
8 by liver biopsy whether it is in the stage of severe inflammation and fibrosis. Finally,
9 the Agile 4 score was used to assess the risk of cirrhosis in patients with advanced
10 fibrosis with an ADAPT value > 4.93 (NPV= 0.955), and patients with an Agile 4
11 value < 0.586 were excluded from cirrhosis and were referred for tertiary care referral
12 (NPV= 0.99). The performance of using this combined diagnostic method is better
13 than using non-invasive algorithms or serum markers alone, and may reduce the
14 number of patients requiring liver biopsy (see **Figure 4**).

15

16 **Discussion**

17 In this study we combined three non-invasive diagnostic scores (i.e. the FAST,
18 ADAPT and Agile 4 scores) for identifying MAFLD progressive steatohepatitis and
19 advanced fibrosis to classify severity of liver disease. The diagnostic performance and
20 accuracy of this sequential prediction algorithm is better than other non-invasive
21 scores or sequential algorithms (including the MLR Model, the combined ADAPT +
22 LSM or the combined (Agile 3+) + Agile 4), and its superiority has been further

1 demonstrated in the independent validation cohort. This sequential prediction
2 algorithm effectively reduces the need for liver biopsy, and provides a more
3 comprehensive classification of MAFLD patients, so that patients with different risks
4 could be better targeted for clinical management.

5

6 Although the degree of liver fibrosis determines the poor prognosis of MAFLD
7 patients [36], increased necrotizing inflammatory activity causes progressive damage
8 and may influence treatment response [37, 38]. The high-risk MAFLD patients with
9 progressive steatohepatitis (defined as steatohepatitis + NAS ≥ 4 + F ≥ 2) could be
10 selected and treated as a secondary or tertiary care referral. The diagnostic
11 performance of the FAST score in the derivation cohort (AUROC= 0.801) could
12 better exclude high-risk patients with MAFLD (NPV= 0.951), and better predict
13 progressive steatohepatitis patients who may have a poor prognosis (NPV= 0.9). At
14 the same time, two thresholds were used to reduce the number of patients in the gray
15 or intermediate zones.

16

17 As the level of fibrosis is the most critical indicator that affects the prognosis of
18 MAFLD patients [39], it is particularly important to assess the severity of the disease
19 as accurately as possible. The ADAPT score based on the new serum marker PRO-C3
20 is good at diagnosing advanced fibrosis (AUROC = 0.879). We used the ADAPT
21 score for further risk assessment of patients with progressive liver inflammation and
22 significant fibrosis screened in the first step (i.e. those with

1 steatohepatitis + $NAS \geq 4 + F \geq 2$). Our results showed that an ADAPT score < 3.705
2 could basically eliminate the risk of advanced fibrosis (sensitivity = 89.7%, NPV =
3 0.972), but it is necessary to confirm by the liver biopsy with patients in the gray
4 zone, through pathological examination to confirm the pathological conditions of the
5 liver, because it is not ruled out that individual patients are still at risk of severe
6 fibrosis. For patients with a high ADAPT value, we used the Agile 4 score for final
7 screening (AUROC = 0.943). After excluding cirrhosis (sensitivity = 88.9%, NPV =
8 0.992), we will advise patients to make tertiary referral as soon as possible for better
9 medical management.

10

11 Our newly proposed three-step sequential algorithm only requires patients to carry out
12 a simple blood test for measurement of serum PRO-C3 level and undergo the
13 FibroScan-measured CAP and LSM measurements. Combined with some basic
14 patient data the algorithm allows evaluation of disease status of patients accurately
15 and quickly. In the face of such a large MAFLD population, our proposed three-step
16 sequential algorithm could effectively screen out low-risk patients and high-risk
17 patients who need different management, so as to reduce the unnecessary social and
18 economic burden of the liver disease.

19

20 The applicability of the ADAPT+ Agile 4 sequential algorithm was also verified in
21 various subgroups of patients with MAFLD. In different patient subgroups, the
22 diagnostic performance of the ADAPT+ Agile 4 sequential algorithm for identifying

1 advanced fibrosis was better than other non-invasive diagnostic score combinations
2 (AUROC = 0.88, accuracy = 88.1%, NPV = 0.955). The combined sequential
3 algorithm only utilized a few simple serological markers and the results of FibroScan
4 to perform a hierarchical screening of MAFLD patients from mild to severe, so as to
5 more accurately implement suitable treatment measures and lifestyle interventions.

6

7 Our study has some limitations that should be mentioned. Firstly, liver biopsy may be
8 affected by sampling variability, intra-observer and inter-observer variability.

9 Secondly, we conducted the study only in Asian patients. In addition, although we

10 compared the diagnostic performance of several non-invasive fibrosis scores and

11 included three models in developing the new combined sequential algorithm, we were

12 unable to compare our newly proposed combined sequential algorithm with magnetic

13 resonance elastography (MRE) [40-42] and other newer serological markers for

14 fibrosis diagnosis (such as CK-18 M30, CK-18 M65, CHI3L1, and M2BPGi) [43].

15 Thirdly, analysing data from larger subsets in each of the MAFLD sub-groups to

16 examine for the performance of x, y and z, would be an important area for future

17 research. Finally, our patients were from tertiary hospital-based centres, so that the

18 effectiveness of the test in primary care needs to be tested.

19

20 In conclusion, our study shows that the combined model of FAST+ ADAPT+ Agile 4

21 scores can be effectively used for risk stratification of liver disease in MAFLD and

22 can reduce the need for unnecessary liver biopsy. Further studies are certainly needed

- 1 to corroborate these findings in other cohorts of patients with MAFLD of different
- 2 countries.
- 3

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Figure legends

Figure 1. (A) The area under the receiver operating characteristic curves (AUROC) for predicting $F \geq 3$ fibrosis. (B) Decision curve analysis (DCA) for the combined diagnosis of non-invasive fibrosis scores. The y-axis represents net benefits, calculated by subtracting the relative harms (true positives) from the benefits (false positives). The x-axis measures the threshold probability. A screening strategy is superior if it has the highest value compared with other models, including two simple strategies, such as all patients (sloping solid line) or no patients (horizontal solid line).

Figure 2. The Sankey diagrams showed the distribution of patients in the true positive (TP), true negative (TN), false positive (FP), false negative (FN), and indeterminate groups. Different thresholds are used for each test layer through the ADAPT score and the Agile 4 score. When applying the two tests, the lower threshold was used to rule out patients without advanced fibrosis and cirrhosis, and the upper threshold was used to rule in patients with advanced fibrosis and cirrhosis (A). In the other three models, we compared the combination of Agile 3+, FIB-4 and NFS scores with Agile 4 scores respectively, and also used lower thresholds and upper thresholds to screen patients with advanced fibrosis and cirrhosis (B, C, D). Two pairs of different thresholds were selected for this hybrid strategy: the lower threshold corresponds to 90% sensitivity; the upper threshold corresponds to 90% specificity. The table next to each panel contains the number and proportion of patients in each TP, TN, FP, and FN groups.

Figure 3. Forest plot for the AUROC of ADAPT + Agile 4 in different subgroups by

gender, age (years), diabetes status, ALT (U/L), BMI (kg/m²), and NAS.

Figure 4. Flow chart for the 3-step serial combination of tests in the sequential algorithm. The FAST score is used as the first test, the ADAPT score is used as the second test, and the Agile 4 score is used as the third test to evaluate and judge the degree of liver inflammation and fibrosis in the biopsy cohort of MAFLD patients.

Supplementary Figure 1. The AUROC of FAST, ADAPT and Agile 4 scores in the derivation cohort.

Supplementary Figure 2. The dual cut-off approach of FAST, ADAPT and Agile 4 scores in the derivation cohort.