Long-term prognosis of MAFLD patients according to non-invasive methods

ABSTRACT

LF-Fib is an alternative to LB for the (logrank p<0.0001). The TRF of TE and FIB4 estimation of the transition rate to fibrosis were also similar to LB however, less fit in [transition to stage F1 or more (TRF)] in T2D both T2D and noT2D groups for both TE [0.75 and noT2D, comparatively to other NITs [FIB- (0.63-0.89) vs 1.34 (1.12-1.60), p<0.001] and 4, liver stiffness measurement (LSM) by FIB4 [0.79 (0.63-0.99) vs 1.26 (1.01-1.59), Fibroscan]

Methods: TRF was evaluated using Cox-Mantel Hazard Ratio [HR(95%CI) and logrank comparison, p value<0,05] with a modeling of hazard from birth to age of LB or NIT in a prospectively collected NAFLD population evaluated for fibrosis with LB, and E concomitant NITs (LF-Fib, LSM, FIB4). NITs cut-offs with highest sensitivity for minimal fibrosis were used (0.28, 1.45 and 5.6kPa respectively).

Results: N=583 pts were included, 52% T2D, and LF, P<0.05 for male gender). 56% males, median (range) age 59.5 (18-85), HbA1c 6.6% (4.7-12), BMI 31.5 (20-54) kg/m2 (obesity 59%), mean (SE) time lapse between LB and NITs 1.7 (0.4) months. The estimation of TRF [HR (95%CI)] using LF-Fib was similar to that using LB in both T2D [0.67 (0.56-0.80) vs. 0.65 (0.54-0.79)], and noT2D [1.50 (1.26-1.78) vs. 1.54 (1.27-1.86)], respectively, with

Objective: The aim was to demonstrate that earlier TRF in noT2D compared to T2D p<0.05], respectively. In pts having ALT>30IU compared to those with ALT≤30IU, the TRF was faster in noT2D [2.13 (1.55-2.95) vs 0.47 (0.34-0.65), logrank p<0.001] and not significantly different in T2D [1.28 (0.93-1.75) vs 0.78 (0.57-1.07), p=ns]. In multivariate analysis, including NITs, arterial hypertension (AHT), HbA1c and BMI, only AHT, BMI≥35, male gender, FIB-4 and LF (Ste, Act and Fib) were significantly associated to TRF in T2D (all p < 0.001) and the same with the exception of BMI and FIB-4 in noT2D (P<0.0001 for AHT

> usion: Validated biomarkers such as IVERFASt should allow a powerful analysis of fibrosis progression in NAFLD, similar to LB and better screening strategies for stratifying

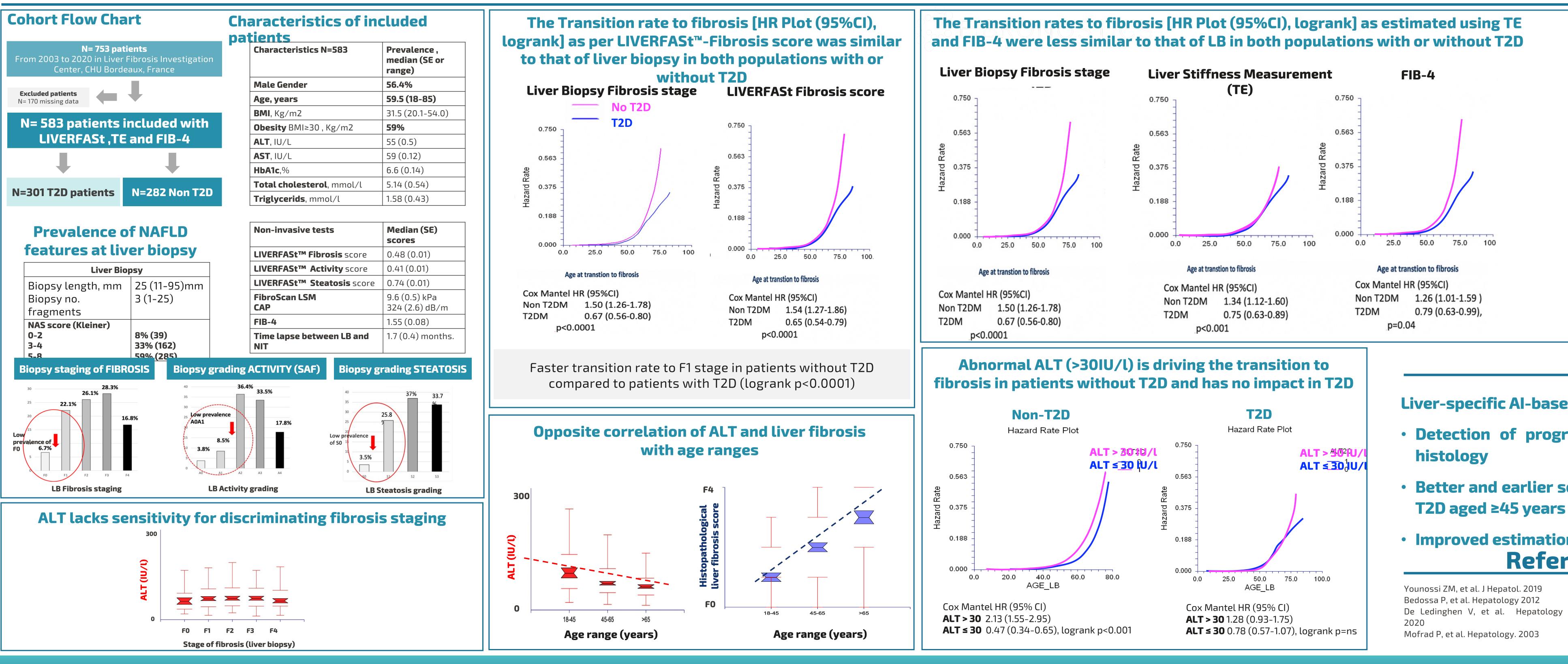
BACKGROUND

Liver biopsy is not adapted to routine diagnosis due to the high prevalence of NAFLD, 40% sample-related variability and poor acceptance

There is an urgent need for reliable non-invasive tools for differentiating NAFL from NASH and for disease staging.

Liver enzymes as are often normal despite advanced fibrosis in T2D and therefore, cannot be used to stage NASH fibrosis.

LIVERFASt[™] (LF) is a serum AI based algorithm (CPT 0166U)Q for assessing liver fibrosis along with steatohepatitis that demonstrated prognostic value to predict overall and liver-related morbimortality.



Hepatology unit, HôpitalHaut Lévêque, Bordeaux University Hospital, Bordeaux, France

INRIA, Talence, France INSERM U1053, Bordeaux University, Bordeaux, France

AIMS

To demonstrate that LIVERFASt[™] Fibrosis score (LF-Fib) is a surrogate of liver biopsy (LB) for the estimation of the transition rate to fibrosis F1 stage or more (TRF1), in type 2 diabetic (T2D) patients with better performances than liver stiffness measurement (LSM) by transient elastography and than FIB-4 index.

^oatients:

- Concomitant LB and LIVERFASt[™], TE, FIB-4.
- Transition rate to any fibrosis stage (TRF) was evaluated using modelling of hazard from birth to the age of the liver fibrosis estimator.
- Cut-offs for minimal fibrosis, F1 stage:
- LB SAF score: perisinusoidal zone 3 or portal fibrosis LIVERFASt[™] -Fibrosis: 0.28; TE: 5.6 kPa; FIB-4: 1.45
- Cox Mantel Hazard Ratios [HR (95%CI), logrank comparison p value between groups

Marie DECRAECKER1, Dan Dutartre2, Jean-Baptiste Hiriart1, Marie Irles-Depé1, Faiza Chermak1, Juliette Foucher1, Victor de Lédinghen1, 3

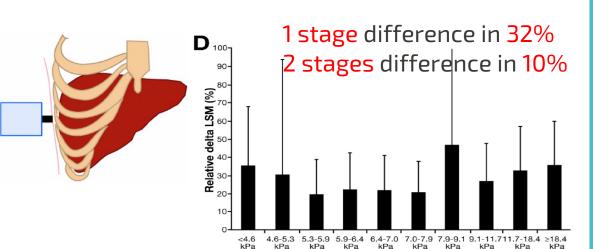
 Prospectively collected NAFLD patients Center tertiary Liver from а (Bordeaux, France) (NCT01241227)

Statistics:



Vibration Controlled Transient

- **Elastography (TE) by Fibroscan** (Echosens, Paris, France) **Ouality criteria:** IQR/median, Success rate, 10 valid LSM
- **Variability** in 531 NAFLD patients paired measurements



Decile group of LSM

Overestimation: Cytolysis with ALT > 3x ULN non-fasting, MetS: T2D, BMI>30, high-blood pressure

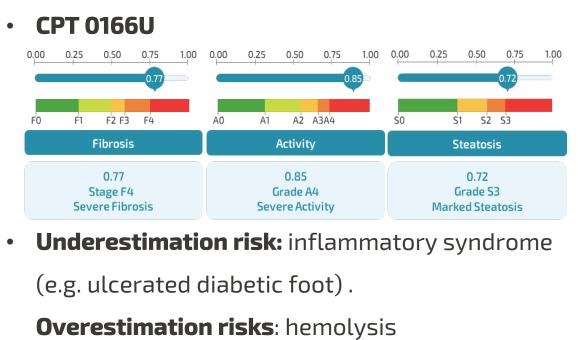
www.echosens.com

Ratziu V, et al. Gastroenterology. 2005 Nascimbeni F, et al. Clin Gastroenterol Hepatol 2014 Castera L, et al. Hepatology 2010; McPherson S, Am J Gastroenterol.

PATIENTS & METHODS

LIVERFASt[™] (Fibronostics, Orlando, Florida) Al computer aided biomarkers constructed using

- **SAF histological scoring** for assessing noninvasively fibrosis, activity and steatosis
- **Combines** 10 biomarkers including liver-specific fibrosis markers, lipid panel, liver enzymes, BMI age, and gender



www.fibronostics.com

FIB-4 Index

Algorithm : platelet count, age, AST, and ALT age(years)×AS(IU/L)

Platelet count $(10^9/L) \times ALT(IU/L)^{\overline{2}}$

- **Dual cut-off** for advanced fibrosis (<1.45, >3.25) **Over or underestimation** : age range, cytolysis,
- normal ALT and AST (T2D) Lower diagnostic performance for cirrhosis in T2D

Multivariate analysis

In T2D NAFLD patients, LIVERFASt[™] Fibrosis, Activity and Steatosis, high blood pressure and male gender were independently associated to the histological transition to fibrosis

Parameter	T2D patients	Non- T2D patients
LIVERFASt™ Fibrosis	P<0.0001	P<0.0001
LIVERFASt [™] Activity	P<0.0001	P<0.0001
LIVERFASt™ Steatosis	P<0.0001	P<0.0001
LSM (TE by Fibroscan)	ns	ns
FIB-4	P<0.0001	ns
Blood Pressure (=high)	P<0.0001	P<0.0001
HbA1c	ns	ns
BMI≥35 Kg/m²	P<0.0001	ns
Gender (=male)	P<0.01	P<0.05

CONCLUSION

Liver-specific AI-based blood biomarkers, such as LIVERFASt[™], allow:

• Detection of progression from simple NAFL to NASH fibrosis, similar to liver

Better and earlier screening strategy for stratifying high-risk patients for NASH, as T2D aged \geq 45 years or having co-morbidities as obesity or arterial hypertension

Improved estimation of elementary liver lesions with noninvasive standard-of-care References Disclosures

Fibronostics: RQ, MM, IA

