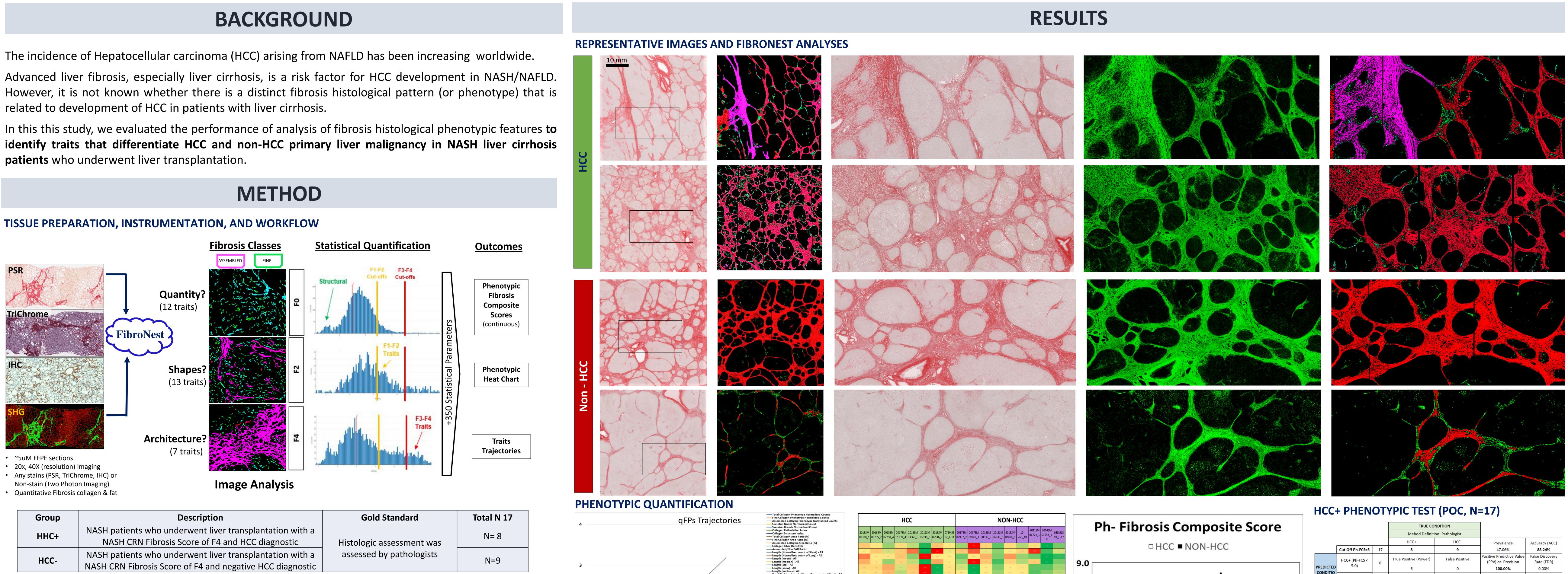


# Automated Fibrosis Phenotyping of NASH non-tumorous lesions digital images helps classify HCC and non-HCC NASH patients who underwent liver transplantation



Group	Description	Gold St
HHC+	NASH patients who underwent liver transplantation with a NASH CRN Fibrosis Score of F4 and HCC diagnostic	Histologic ass assessed by
HCC-	NASH patients who underwent liver transplantation with a NASH CRN Fibrosis Score of F4 and negative HCC diagnostic	

- FFPE sections (3-4 microns) of patient liver non-tumorous tissue were deparaffinized, stained for with Picro Sirus Red (no Hematoxylin bath) for Collagen and digitized at 40X (Aperio AT2)
- Using FibroNest<sup>™</sup> the fibrosis phenotype is described for its collagen content and structure (12 traits), the morphometric traits of the collagen fibers (13 traits), and fibrosis architecture / texture traits (7). In each image, each morphometric and texture trait is represented by a histogram distribution (e.g. Fiber Skeleton Length histogram).
- The histogram for each trait is described by up to seven quantitative fibrosis parameters (qFPs, 315 in total) to account for mean, variance, distortion and progression.
- Principal qFPs are automatically detected if their group distribution (N=8 or N=9) significantly different (p<0.001, T- TEST), and normalized group mean value ratio greater than 1.2
- Principal qFPS are used individually and collectively to describe the differences in phenotypes between the two groups. They ate combined into a normalized Phenotypic Composite Fibrosis Score (Ph-CFS), a continuous quantifier of the HCC fibrosis phenotype.
- Cut-off values are selected to optimize the performance of a test to classify the patients (HCC or non-HCC) based on their Ph-CFS. The performance of the test is evaluated using a confusion matrix and its related sensitivity and specificity values.

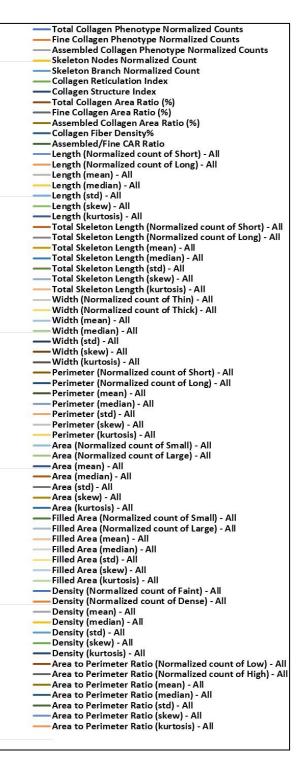
Hisamitsu Miyaaki<sup>1</sup>, Yuko Akazawa<sup>1</sup>, Li Chen<sup>2</sup>, Mathieu Petitjean<sup>2</sup>

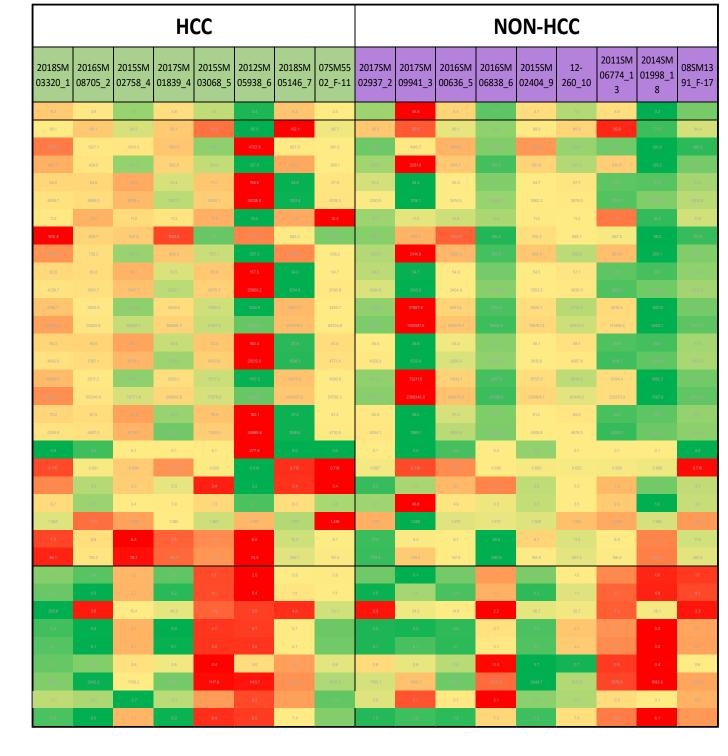
<sup>1</sup>Nagasaki University, Nagasaki, Japan <sup>2</sup>PharmaNest, Princeton, NJ, USA

qFP trajectories (normalized to HCC phenotype) describe the differences between HCC and non-HCC Phenotypes and highlight principal qFPs

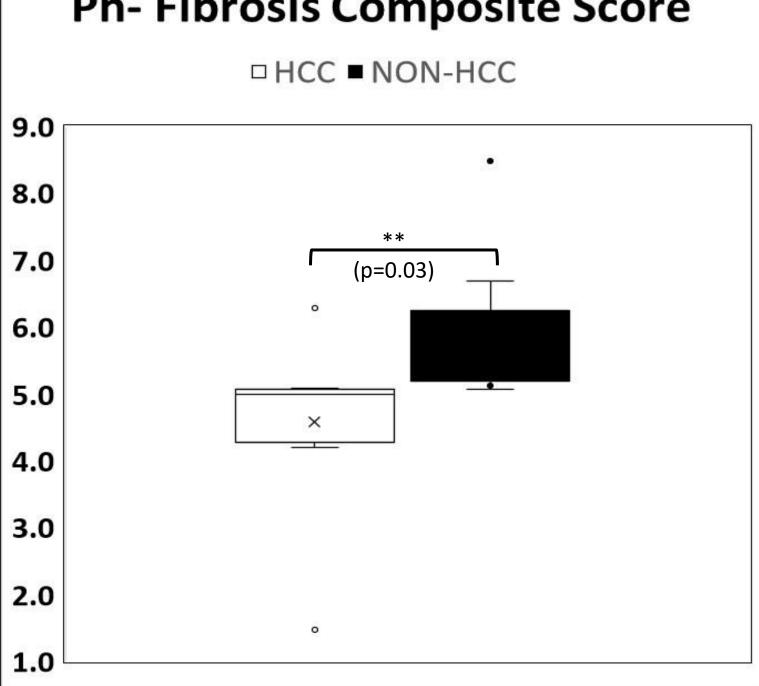
NON-HCC

HCC





Phenotypic Heat charts describe each Patient's phenotype (principal qFPs only)



The Phenotypic Composite score is assembled from Principal gFPs and helps classify the HCC and non-**HCC** Patients



## False Omission Rate Negative Predictive N False Negative HCC- (Ph- FCS > rue Negative Value (NPV) (FOR) 81.82% 18.18% rue Positive Rate (TPR) False Positive Rate (FPR) Positive Likelihood Ratio Diagnostic Odo or Recall, Sensitivity, or Fall-Out or Probability (LR+) 75% 0.00% Specificity (SPC) False Negative Rate Negative Likelihood Sselectivity or True FNR) or Miss Rate Ratio (LR-) Negative Rate (TNR) 0.25 25.00% 100%

100% Sensitivity and 100% Specificity (N=17)

## Conclusion

Fibrosis histological phenotypic quantification helps to distinguish between HCC and non-HCC in NASH liver cirrhosis patients who underwent liver transplant. These data show that phenotypic analysis of collagen with FibroNest<sup>™</sup> is an effective and automated method to classify HCC from non-HCC in histopathology imaging studies. This can be of clinical importance for appropriate guidance of treatment strategy