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LETTERS TO THE EDITOR

# **PNPLA3 I148M variant affects non-alcoholic fatty liver disease in liver transplant recipients**

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## Abstract

De novo non-alcoholic fatty liver disease (NAFLD) is a common late complication for long-term survivors after liver transplantation. Genomic studies confirmed that PNPLA3 I148M and TM6SF2 E167K polymorphisms affected NAFLD susceptibility in the general population. However, this association was not validated in survivors after liver transplantation (LT). We performed a crosssectional survey to investigate this relationship. A comprehensive survey, including anthropometric measurements, fasting venous blood sampling, ultrasound, and questionnaires was performed in the shortterm. The clinical indications and patient's steatosis status before LT were collected from inpatient medical records. Sixty-five long-term recipients with a survival exceeding 10 years were enrolled in the final analysis. De novo NAFLD was more frequent in PNPLA3 GG carriers (0.33 vs 0.10 for GG vs CC + CG carriers, P = 0.018), while the genetic impact on NAFLD susceptibility was insignificant when categorized by the TM6SF2 polymorphism (0.19 in CC vs 0.14 in CT + TT carriers, P = 0.883). Multi-covariate analysis revealed that PNPLA3 exerted a significant genetic effect on de novo NAFLD following a recessive model (GG vs CC + CG, OR = 14.2, 95%CI: 1.78-113, P = 0.012). Compared to recipients with only the PNPLA3 GG allele or obesity (defined as body mass index > 25 kg/m<sup>2</sup>), steatosis was highly prevalent (71.4%) in PNPLA3 GG carriers



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with obesity. In conclusion, PNPLA3 I148M, but not TM6SF2 E167K, affects *de novo* NAFLD occurrence with a prominent interaction with obesity. Weight control might be a meaningful method to reduce the genetic susceptibility to NAFLD exerted by PNPLA3 variants.

Key words: PNPLA3; TM6SF2; Non-alcoholic fatty liver disease; Liver transplantation; Recipient

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**Core tip:** Previous genomic studies identified PNPLA3 I148M and TM6SF2 E167K polymorphisms as the most prominent genetic variations associated with nonalcoholic fatty liver disease (NAFLD) susceptibility in general populations. However, these impacts have never been evaluated in long-term liver transplant recipients. In a collection of survivors 10 years after liver transplantation, we found that the PNPLA3 I148M, but not TM6SF2 E167K polymorphism, affected *de novo* NAFLD predisposition and interacted with obesity. Our results revealed that liver transplant recipients might benefit from weight control to limit the deleterious effect exerted by genetic factors.

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### TO THE EDITOR

*De novo* non-alcoholic fatty liver disease (NAFLD) is a common late complication for long-term survivors after liver transplantation (LT)<sup>[1]</sup>. *De novo* NAFLD affects allograft survival indirectly by increasing cardiovascular and infectious disease occurrence<sup>[2]</sup>. Previous genomic studies identified the PNPLA3 I148M and TM6SF2 E167K polymorphisms as the most likely single nucleotide polymorphisms to influence NAFLD susceptibility in the general population<sup>[3]</sup>. However, this relationship was not confirmed in long-term survivors after LT as a specific population. Therefore, we performed a cross-sectional survey to investigate the impact of genetic and environmental risk factors for *de novo* NAFLD in adult long-term survivors after receiving LT.

After obtaining written informed consent, a comprehensive survey, including anthropometric measurements (for body weight and height), fasting venous blood sampling (for liver function, lipid, glucose, viral biomarker testing, genotyping, *etc.*), ultrasound examination, and questionnaires (for alcohol intake, smoking, exercise, and immunosuppression) were

# Table 1Current status of long-term recipients surviving<br/>more than 10 years

	Univariate			Multivariate	
	$\begin{array}{l} NAFLD \\ (n = 12) \end{array}$	Control $(n = 53)$	<i>P</i> value	OR	<i>P</i> value
Age (yr)	$56.5 \pm 8.4$	$53.6 \pm 10.1$	0.356	1.04	0.528
				(0.92-1.18)	
Gender (M/F)	10/2	47/6	0.611	1.40	0.427
				(0.14-14.2)	
Indication for LT					
Hepatitis/cirrhosis/	1/8/	7/35/	0.889		
cancer/others	2/1	9/2			
Survival time (yr)	$11.2 \pm 0.9$	$11.5 \pm 1.4$	0.541		
BMI $(kg/m^2)$	$25.1 \pm 3.0$	$22.5 \pm 2.6$	0.003	1.47	0.032
				(1.03-2.08)	
TG (mmol/L)	$1.6 \pm 1.1$	$1.1 \pm 0.6$	0.038	1.34	0.652
,				(0.38-4.71)	
HDL-C (mmol/L)	1.2 (1.0-1.4)	1.3 (1.0-1.7)	0.267		
FBG (mmol/L)	7.6 ± 3.4	5.7 ± 1.9	0.013	1.49	0.095
· · · /				(0.93-2.37)	
Hypertension	3/9	21/32	0.343		
(Yes/no)	,	,			
SUA (µmol/L)	381.6 ± 75.6	342.6 ± 76.4	0.116		
MetS (Yes/no)	4/8	9/44	0.201		
ALT (U/L)	36.7 ± 7.0	$38.8 \pm 6.7$	0.882		
Alcohol intake	$11.6 \pm 7.3$	$21.0 \pm 8.2$	0.766		
(g/wk)					
Smoking (cigar/d)	$4.2 \pm 3.4$	$4.3 \pm 1.2$	0.969		
Exercise (min/d)	$18.5 \pm 6.0$	22.9 ± 2.9	0.513		
Immunosuppression					
Tacrolimus/	11/1	36/12	0.575		
cyclosporine/	/0/0/0	/1/2/2			
MMF/sirolimus/	, , ,	, , ,			
none					
PNPLA3	1/3/8	16/21/16	0.018	14.2	0.012
(CC/CG/GG)				(1.78-113)	
TM6SF2	11/1/0	47/5/1	0.883	2.68	0.413
(CC/CT/TT)				(0.25-28.5)	
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Continuous variables with equal variance are presented as the mean ± SD; Continuous variables with unequal variance are presented as the median (interquartile range); Categorical variables are presented as the number of subjects. One-way ANOVA was used for the comparison between continuous variables with equal variance, Mann-Whitney U test was used for the comparison between continuous variables with unequal variance, chi-square test was used for the comparison between categorical variables in univariate analysis, and logistic regression analysis was used in the multivariate analysis. The effect of the PNPLA3 I148M polymorphism was evaluated by a recessive genetic model (GG vs CG + CC); the effect of the TM6SF2 E167K polymorphism was evaluated by a dominant genetic model (CC + CT vs TT) for decreased prevalence of TT carrier. ALT: Alanine aminotransferase; BMI: Body mass index; F: Female; FBG: Fasting blood glucose; HBsAg: Hepatitis B surface antigen; HDL-C: Highdensity lipoprotein cholesterol; LT: Liver transplantation; M: Male; MetS: Metabolic syndrome; MMF: Mycophenolatemofetil; SUA: Serum uric acid; TG: Triglyceride.

performed over the short-term (December 13<sup>th</sup>-14<sup>th</sup>, 2014). Indications for LT and patients' steatosis status before LT were collected from inpatient medical records. The study was approved by the Institutional Review Board of our hospital.

NAFLD and metabolic syndrome were defined according to previous criteria<sup>[4]</sup>. Participants with recurrent liver steatosis were excluded. Accordingly, 65



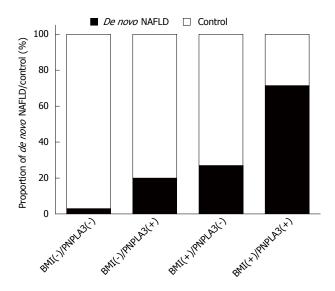


Figure 1 Proportion of the *de novo* non-alcoholic fatty liver disease/ control categorized by the body mass index-PNPLA3 I148M polymorphism. BMI (+) represents BMI  $\leqslant 25$  kg/m<sup>2</sup>; BMI (-) represents BMI > 25 kg/m<sup>2</sup>; PNPLA3 (+) indicates PNPLA3 I148M GG carriers; PNPLA3 (-) indicates the PNPLA3 I148M CC + CG carriers. BMI: Body mass index; NAFLD: Nonalcoholic fatty liver disease.

subjects (57 males and 8 females) receiving LT (from September, 1999 to November, 2004) in our hospital with a survival exceeding 10 years were enrolled into the final analysis.

All survivors were Han Chinese and negative for hepatitis B virus DNA/hepatitis C virus RNA assay. As shown in Table 1, twelve of the patients were diagnosed with *de novo* NAFLD. By univariate analysis, the NAFLD subgroup hada significantly higher body mass index (BMI), triglyceride (TG) levels, and fasting blood glucose (FBG) levels. *De novo* NAFLD was more frequent in PNPLA3 GG carriers than in CC + CG carriers (0.33 vs 0.10, P = 0.018), while the genetic impact on NAFLD susceptibility was insignificant when categorized by the TM6SF2 polymorphism (0.19 in CC vs 0.14 in CT + TT carriers, P = 0.883). Multicovariate analysis revealed that PNPLA3 exerted a significant genetic effect on *de novo* NAFLD following a recessive model (GG vs CC + CG, OR = 14.2, 95%CI: 1.78-113, P = 0.012). Compared to recipients only carrying the PNPLA3 GG allele or being obese (defined as BMI > 25 kg/m<sup>2</sup>), the prevalence of steatosis was disproportionally higher (71.4%) in PNPLA3 GG carriers who were obese (Figure 1).

This is the first report on the risk factors associated with *de novo* steatosis in Chinese long-term survivors after LT. PNPLA3, but TM6SF2, affects *de novo* NAFLD occurrence and has a prominent interaction with obesity. Weight control in recipients might be a potential method to reduce the genetic susceptibility of NAFLD exerted by the PNPLA3 variant.

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