

# CITATION REPORT

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Transmembrane 6 superfamily member 2 gene E167K variant impacts on steatosis and liver damage in chronic hepatitis C patients

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#	Paper	IF	Citations
50	TM6SF2 E167K Variant, a Novel Genetic Susceptibility Variant, Contributing to Nonalcoholic Fatty Liver Disease. <i>Journal of Clinical and Translational Hepatology</i> , <b>2015</b> , 3, 265-70	5.2	24
49	Insights from Genome-Wide Association Analyses of Nonalcoholic Fatty Liver Disease. <i>Seminars in Liver Disease</i> , <b>2015</b> , 35, 375-91	7.3	37
48	PNPLA3 p.I148M variant is associated with greater reduction of liver fat content after bariatric surgery. <i>Surgery for Obesity and Related Diseases</i> , <b>2016</b> , 12, 1838-1846	3	44
47	Age might confound the impact of the transmembrane 6 superfamily member 2 E167K polymorphism on hepatic fibrosis in hepatitis C virus-infected patients. <i>Hepatology</i> , <b>2016</b> , 63, 1052	11.2	
46	TM6SF2 rs58542926 is not associated with steatosis and fibrosis in large cohort of patients with genotype 1 chronic hepatitis C. <i>Liver International</i> , <b>2016</b> , 36, 198-204	7.9	12
45	PNPLA3 gene in liver diseases. <i>Journal of Hepatology</i> , <b>2016</b> , 65, 399-412	13.4	140
44	Host - hepatitis C viral interactions: The role of genetics. <i>Journal of Hepatology</i> , <b>2016</b> , 65, S22-S32	13.4	42
43	Meta-analysis of the influence of TM6SF2 E167K variant on Plasma Concentration of Aminotransferases across different Populations and Diverse Liver Phenotypes. <i>Scientific Reports</i> , <b>2016</b> , 6, 27718	4.9	16
42	Reply. <i>Hepatology</i> , <b>2016</b> , 63, 1052-3	11.2	
41	Diverse impacts of the rs58542926 E167K variant in TM6SF2 on viral and metabolic liver disease phenotypes. <i>Hepatology</i> , <b>2016</b> , 64, 34-46	11.2	65
40	Associations of TM6SF2 167K allele with liver enzymes and lipid profile in children: the PANIC Study. <i>Pediatric Research</i> , <b>2016</b> , 79, 684-8	3.2	11
39	Hepatic Transmembrane 6 Superfamily Member 2 Regulates Cholesterol Metabolism in Mice. <i>Gastroenterology</i> , <b>2016</b> , 150, 1208-1218	13.3	61
38	Genetics of nonalcoholic fatty liver disease. <i>Metabolism: Clinical and Experimental</i> , <b>2016</b> , 65, 1026-37	12.7	67
37	PNPLA3 rs738409 and TM6SF2 rs58542926 variants increase the risk of hepatocellular carcinoma in alcoholic cirrhosis. <i>Digestive and Liver Disease</i> , <b>2016</b> , 48, 69-75	3.3	60
36	The effect of the TM6SF2 E167K variant on liver steatosis and fibrosis in patients with chronic hepatitis C: a meta-analysis. <i>Scientific Reports</i> , <b>2017</b> , 7, 9273	4.9	14
35	Recent Advances in the Pathogenesis of Hepatitis C Virus-Related Non-Alcoholic Fatty Liver Disease and Its Impact on Patients Cured of Hepatitis C. <i>Current Hepatology Reports</i> , <b>2017</b> , 16, 317-325	1	1
34	MBOAT7 rs641738 variant and hepatocellular carcinoma in non-cirrhotic individuals. <i>Scientific Reports</i> , <b>2017</b> , 7, 4492	4.9	131

33	Combined effects of the PNPLA3 rs738409, TM6SF2 rs58542926, and MBOAT7 rs641738 variants on NAFLD severity: a multicenter biopsy-based study. <i>Journal of Lipid Research</i> , <b>2017</b> , 58, 247-255	6.3	108
32	The rs738409 polymorphism of the PNPLA3 gene is associated with hepatic steatosis and fibrosis in Brazilian patients with chronic hepatitis C. <i>BMC Infectious Diseases</i> , <b>2017</b> , 17, 780	4	11
31	PNPLA3 Variant p.I148M and Bariatric Surgery. <b>2017</b> , 389-396		
30	The rs4686434 variant in the fetuin B (FETUB) locus is associated with intrahepatic triglyceride content in obese Chinese adults. <i>Journal of Diabetes</i> , <b>2018</b> , 10, 916-925	3.8	1
29	Genetics and epigenetics of NAFLD and NASH: Clinical impact. <i>Journal of Hepatology</i> , <b>2018</b> , 68, 268-279	13.4	362
28	TM6SF2 Promotes Lipidation and Secretion of Hepatitis C Virus in Infected Hepatocytes. <i>Gastroenterology</i> , <b>2018</b> , 155, 1923-1935.e8	13.3	8
27	Non-alcoholic fatty liver disease: a narrative review of genetics. <i>Journal of Biomedical Research</i> , <b>2018</b> , 32, 389-400	1.5	24
26	Genetic and Epigenetic Modifiers of Alcoholic Liver Disease. <i>International Journal of Molecular Sciences</i> , <b>2018</b> , 19,	6.3	44
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23	The roles of transmembrane 6 superfamily member 2 rs58542926 polymorphism in chronic liver disease: A meta-analysis of 24,147 subjects. <i>Molecular Genetics &amp; Genomic Medicine</i> , <b>2019</b> , 7, e824	2.3	16
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21	and Gene Variants in Liver Fibrosis and Cirrhosis. <i>International Journal of Molecular Sciences</i> , <b>2019</b> , 20,	6.3	22
20	PNPLA3-A Potential Therapeutic Target for Personalized Treatment of Chronic Liver Disease. <i>Frontiers in Medicine</i> , <b>2019</b> , 6, 304	4.9	19
19	Genetic contributions to NAFLD: leveraging shared genetics to uncover systems biology. <i>Nature Reviews Gastroenterology and Hepatology</i> , <b>2020</b> , 17, 40-52	24.2	94
18	Donor PNPLA3 and TM6SF2 Variant Alleles Confer Additive Risks for Graft Steatosis After Liver Transplantation. <i>Transplantation</i> , <b>2020</b> , 104, 526-534	1.8	7
17	The influence of gene-chronic hepatitis C virus infection on hepatic fibrosis and steatosis. <i>Diagnostic Microbiology and Infectious Disease</i> , <b>2020</b> , 97, 115025	2.9	4
16	Comparison of performance, fatty acid composition, enzymes and gene expression between overfed Xupu geese with large and small liver. <i>Italian Journal of Animal Science</i> , <b>2021</b> , 20, 102-111	2.2	1

15	Study of CXCL9-11 gene polymorphisms in liver fibrosis among patients with chronic hepatitis C. <i>Pathogens and Disease</i> , <b>2021</b> , 79,	4.2	1
14	The impact of PNPLA3 and TM6SF2 in cirrhosis related complications.		
13	The role of PNPLA3 and TM6SF2 polymorphisms on liver fibrosis and metabolic abnormalities in Brazilian patients with chronic hepatitis C. <i>BMC Gastroenterology</i> , <b>2021</b> , 21, 81	3	1
12	Influence of Genetic Variants on Disease Regression and Outcomes in HCV-Related Advanced Chronic Liver Disease after SVR. <i>Journal of Personalized Medicine</i> , <b>2021</b> , 11,	3.6	2
11	Non-Alcoholic Fatty Liver Disease: Metabolic, Genetic, Epigenetic and Environmental Risk Factors. <i>International Journal of Environmental Research and Public Health</i> , <b>2021</b> , 18,	4.6	13
10	Novel association of rs58542926 genotype with increased serum tyrosine levels and decreased apoB-100 particles in Finns. <i>Journal of Lipid Research</i> , <b>2017</b> , 58, 1471-1481	6.3	35
9	The Genetics of Clinical Liver Diseases: Insight into the E167K Variant. <i>Journal of Clinical and Translational Hepatology</i> , <b>2018</b> , 6, 326-331	5.2	5
8	TM6SF2 E167K variant predicts severe liver fibrosis for human immunodeficiency/hepatitis C virus co-infected patients, and severe steatosis only for a non-3 hepatitis C virus genotype. <i>World Journal of Gastroenterology</i> , <b>2016</b> , 22, 8509-8518	5.6	9
7	TM6SF2 rs58542926 Polymorphism is not Associated With Risk of Steatosis or Fibrosis in Chilean Patients With Chronic Hepatitis C. <i>Hepatitis Monthly</i> , <b>2017</b> , 17,	1.8	0
6	Genetic Aspects of Non-Alcoholic Fatty Liver Disease. <i>Voprosy Sovremennoi Pediatrii - Current Pediatrics</i> , <b>2019</b> , 17, 442-448	0.5	1
5	PNPLA3 and TM6SF2, but Not MBOAT7, Are Associated with Steatosis and HBV Viral Persistence in Pakistani Population. <i>Jundishapur Journal of Microbiology</i> , <b>2020</b> , 13,	1.2	
4	Research progress on the relationship between TM6SF2 rs58542926 polymorphism and non-alcoholic fatty liver disease.. <i>Expert Review of Gastroenterology and Hepatology</i> , <b>2022</b> , 1-11	4.2	0
3	The influence of host genetics on liver microbiome composition in patients with NAFLD.. <i>EBioMedicine</i> , <b>2022</b> , 76, 103858	8.8	3
2	Evaluation of the association of a variant in PNPLA3 and TM6SF2 with fibrosis progression in patients with chronic hepatitis C infection after eradication: A retrospective study.. <i>Gene</i> , <b>2022</b> , 820, 146235	3.8	1
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