NAFLD as a driver of chronic kidney disease

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Summary
Non-alcoholic fatty liver disease (NAFLD) and chronic kidney disease (CKD) are worldwide public health problems, affecting up to 25–30% (NAFLD), and up to 10–15% (CKD) of the general population. Recently, it has also been established that there is a strong association between NAFLD and CKD, regardless of the presence of potential confounding diseases such as obesity, hypertension and type 2 diabetes. Since NAFLD and CKD are both common diseases that often occur alongside other metabolic conditions, such as type 2 diabetes or metabolic syndrome, elucidating the relative impact of NAFLD on the risk of incident CKD presents a substantial challenge for investigators working in this research field. A growing body of epidemiological evidence suggests that NAFLD is an independent risk factor for CKD and recent evidence also suggests that associated factors such as metabolic syndrome, dysbiosis, unhealthy diets, platelet activation and processes associated with ageing could also contribute mechanisms linking NAFLD and CKD. This narrative review provides an overview of the literature on: a) the evidence for an association and causal link between NAFLD and CKD and b) the underlying mechanisms by which NAFLD (and factors strongly linked with NAFLD) may increase the risk of developing CKD.

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Introduction
Readers of the Journal will be very familiar with non-alcoholic fatty liver disease (NAFLD) and how to define it, but will perhaps be less familiar with chronic kidney disease (CKD) and its definition.

Both diseases are progressive chronic conditions that represent a spectrum of diseases extending from relatively mild disease, with only modest changes in function, to severe debilitating disease with end-stage organ damage, necessitating either chronic dialysis or organ transplantation in order to sustain life. NAFLD encompasses a histopathological spectrum of metabolic liver conditions encapsulating simple steatosis alone (non-alcoholic fatty liver, i.e. NAFL); steatosis, inflammation and ballooning of hepatocytes, with or without liver fibrosis (non-alcoholic steatohepatitis, i.e. NASH), and cirrhosis.1,2 When advanced fibrosis or cirrhosis occur, the risk of hepatocellular carcinoma also increases markedly.

CKD is a complex, progressive chronic condition that is defined by either abnormalities of kidney structure or function present for ≥3 months, with serious implications for health.1-4 Either markers of kidney damage or decreased glomerular filtration rate may be present.

The National Kidney Foundation has identified 5 stages of CKD from 1 to 5 (as shown in Fig. 1).5 In the presence of a urinary albumin-to-creatinine ratio (ACR) that is normal or very mildly increased (i.e. urinary ACR <30 mg/g) and an estimated glomerular filtration rate (eGFR) above 60 ml/min/1.73 m², the risk of progression to end-stage renal disease is very low and such patients usually do not undergo regular surveillance. Clinicians should therefore identify CKD stage 3 or above, because these stages of CKD are associated with a high or very high risk of disease progression. To define CKD stage ≥3, markers of kidney damage can include the presence of one of: abnormal albuminuria (ACR ≥30 mg/g) or overt proteinuria, urine sediment abnormalities and other abnormalities due to tubular disorders, abnormalities detected by kidney histology, structural abnormalities detected by imaging, or a history of renal transplantation. For decreased eGFR, CKD is defined by an eGFR value <60 ml/min/1.73 m².2,3,4 NAFLD and CKD are associated with poor outcomes and high costs; they have become major public health problems owing to their increasing prevalence and incidence. Indeed, NAFLD affects up to ~25–30%1,6 and CKD affects up to ~10–15% of the general adult population in many parts of the world.5,8 It is well established that CKD is also a major risk factor for cardiovascular disease (CVD) and all stages of CKD are associated with an increased risk of cardiovascular morbidity, premature mortality and decreased quality of life.9 Recently, it has also been shown that NAFLD is an independent risk factor for CVD, regardless of the coexistence of cardiometabolic risk factors, such as obesity, hypertension, type 2 diabetes mellitus (T2DM) or metabolic syndrome (MetS).10,11 Therefore, since NAFLD and CKD often occur with features of the MetS that adversely affect the kidney, elucidating the relative impact of NAFLD on the risk of incident CKD presents a
substantial challenge to investigators working in this field of research.

The main aims of this narrative review are to discuss: a) the evidence for an association and causal link between NAFLD and CKD and b) the putative mechanisms by which NAFLD (and factors strongly linked with NAFLD) may increase the risk of developing CKD.

Evidence of an association between NAFLD and CKD
Since patients with NAFLD exhibit multiple traditional and non-traditional risk factors for CKD (as summarised in Fig. 1), it is not surprising that these patients also have a higher prevalence and incidence of CKD compared with those who do not have steatosis.

Cross-sectional studies
Several hospital-based and community-based studies have documented that NAFLD, as assessed by imaging techniques or liver biopsy, is significantly associated with an increased prevalence of CKD (defined as eGFR <60 ml/min/1.73 m², abnormal albuminuria or overt proteinuria). As reviewed extensively elsewhere, in these studies the prevalence of CKD ranged from approximately 20% to 55% among patients with NAFLD compared to 5% to 30% among their counterparts without NAFLD. Notably, in most of these studies the significant association between NAFLD and increased prevalence of CKD persisted, both in patients with T2DM and in those without diabetes, even after adjustment for common risk factors for CKD. Some smaller case-control studies using liver biopsy to diagnose NAFLD have also shown a significant, graded association between the histologic severity of NAFLD (mainly the hepatic fibrosis stage) and the presence of either decreased eGFR or abnormal albuminuria. For example, in a previous case-control study, we found that patients with biopsy-confirmed NASH had a higher prevalence of both CKD and abnormal albuminuria than age-, sex- and body mass index (BMI)-matched controls, and that the histologic stage of liver fibrosis was associated with decreasing eGFR values, independently of age, sex, adiposity measures, hypertension, plasma triglyceride concentrations and homeostatic model assessment (HOMA)-estimated insulin resistance (Fig. 2).

Cohort studies
Although the current evidence, from cross-sectional studies, for the existence of an association between NAFLD and increased prevalence of CKD is robust and consistent across different ethnicities and patient populations, whether NAFLD is also a "driving force" for the development and progression of CKD remains uncertain. However, to date, an ever-increasing number of retrospective and prospective cohort studies, with a reasonably long duration of follow-up, have consistently documented that NAFLD (diagnosed

Key point
Both NAFLD and CKD are major public health problems whose prevalence and incidence is growing.

Fig. 1. Risk factors in patients with NAFLD and CKD. The stages of CKD are also reported in the figure. CKD is defined as either kidney damage or eGFR <60 ml/min/1.73 m² for at least 3 months or more. Kidney damage is defined as pathologic abnormalities or markers of damage, including abnormalities in blood or urine tests or in imaging methods. CKD, chronic kidney disease; eGFR, estimated glomerular filtration rate; NAFLD, non-alcoholic fatty liver disease.
either by abnormal levels of serum liver enzymes or by imaging techniques) is significantly associated with an increased incidence of CKD.\(^{44–46}\) (Table 1). Notably, in most of these studies the significant association between NAFLD and increased incidence of CKD persisted even after adjustment for age, sex, obesity, hypertension, T2DM and other potential confounding factors. For example, in the Valpolicella Heart Diabetes Study, including 1,760 outpatients with T2DM who had preserved kidney function at baseline, the presence of NAFLD on ultrasonography was associated with a nearly 50% increase in the risk of incident CKD (adjusted hazard ratio 1.49; 95% CI 1.1–2.2) over a follow-up period of 6.5 years, independent of age, sex, adiposity measures, blood pressure, smoking, duration of diabetes, haemoglobin A1c, plasma lipids, baseline eGFR, microalbuminuria, and the use of hypoglycaemic, lipid-lowering, antihypertensive or antiplatelet drugs.\(^{30}\) Similar results have also been found in adults with type 1 diabetes and in other large community-based cohort studies of different ethnicities (Table 1).

### Systematic reviews and meta-analyses

In a meta-analysis of 33 observational (20 cross-sectional and 13 longitudinal) studies published in 2014, Musso et al. examined the association between NAFLD and risk of prevalent and incident CKD (defined as eGFR <60 ml/min/1.73 m\(^2\), abnormal albuminuria or both).\(^{47}\) Meta-analysis of the data from 20 cross-sectional studies (involving nearly 30,000 individuals) showed that NAFLD was associated with a 2-fold increased prevalence of CKD (random-effects odds ratio 2.12, 95% CI 1.69–2.66). More interestingly, meta-analysis of data from the 13 longitudinal studies (involving a total of nearly 28,500 individuals) showed that NAFLD was associated with a nearly 80% increased risk of incident CKD (random-effects hazard ratio [HR] 1.79; 95% CI 1.65–1.95). Similarly, in a sub-group analysis of individual patient data from 5 small studies (involving a total of ~430 adults with biopsy-confirmed NAFLD with only 86 incident CKD cases), the authors also suggested that the presence of advanced hepatic fibrosis was associated with a higher prevalence (random-effects odds ratio 5.20; 95% CI 3.14–8.61) and incidence (random-effects HR 3.29; 95% CI 2.30–4.71) of CKD than either non-advanced fibrosis or simple steatosis, respectively.\(^{47}\) In all of the aforementioned analyses, the presence and severity of NAFLD were associated with a higher prevalence and incidence of CKD even after adjustment for pre-existing T2DM and other common risk factors for CKD, such as age, ethnicity, BMI and smoking history.\(^{47}\)

Recently, we have also performed a comprehensive systematic review and meta-analysis that involved a total of 9 observational cohort studies (published up to August 2017) with aggregate data on ~96,500 middle-aged individuals (34.1% with NAFLD) of predominantly Asian descent and ~5,000 new cases of incident CKD (stage ≥3, defined as occurrence of eGFR <60 ml/min/1.73 m\(^2\), with or without accompanying proteinuria) over a median follow-up period of 5.2 years.\(^{48}\) No studies with biopsy-proven NAFLD were available for the analysis. As shown in Fig. 3, this updated meta-analysis confirmed that NAFLD (detected by serum liver enzymes, fatty liver index or ultrasonography) was associated with a nearly 40% increase in the long-term risk of incident CKD (random-effects HR 1.37; 95% CI 1.20–1.53; \(I^2 = 33.5\%\)), a risk that appeared to increase in parallel with the severity of NAFLD, as assessed by the NAFLD fibrosis score or other non-invasive markers of advanced fibrosis (n = 2 studies; random-effects HR 1.50; 95% CI 1.25–1.74; \(I^2 = 0\%\)), and remained significant in those studies where analysis was adjusted for common risk factors and potential confounders (i.e., age, sex, BMI, hypertension, smoking, diabetes, baseline eGFR and use of certain medications). In addition, as also shown in Fig. 3, when the analysis was stratified by the type of study population, the association between NAFLD and risk of incident CKD was essentially consistent for both patients with diabetes, and those without diabetes at baseline.\(^{49}\) In the few studies involving patients with T2DM, the association between NAFLD and risk of CKD remained significant even after adjusting for duration of diabetes, glycaemic control, hypertension and other established risk factors for CKD.\(^{46}\) In addition, when the analysis was stratified either by study country (Asian vs. European countries), the association between

![Figure 2. Renal function parameters in patients with NASH and controls.](image)
Table 1. Principal observational cohort studies examining the association between NAFLD and risk of development or progression of CKD stage 3 (ordered by publication year).

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<th>Authors, Country, Year (Ref.)</th>
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<td>Ryu S et al., South Korea, 2007</td>
<td>Community-based cohort study: 10,337 non-diabetic and non-hypertensive male workers with normal kidney function and no overt proteinuria at baseline. Follow-up: 3 years</td>
<td>Serum liver enzymes (serum GGT concentrations)</td>
<td>eGFR &lt;60 ml/min/1.73 m² and/or overt proteinuria (urinary dipstick ≥1); 366 patients developed incident CKD during follow-up</td>
<td>Age, BMI, alcohol intake, smoking, baseline eGFR, triglycerides, HDL-cholesterol, C-reactive protein, HOMA-IR, and incident cases of hypertension and diabetes</td>
<td>NAFLD (i.e. top quartile of serum GGT concentrations) was independently associated with increased risk of incident CKD (AHR 1.87; 95% CI 1.31-2.67).</td>
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<tr>
<td>Chang Y et al., South Korea, 2008</td>
<td>Community-based cohort study: 8,329 non-diabetic and non-hypertensive men with normal kidney function and no overt proteinuria at baseline. Follow-up: 3.2 years</td>
<td>Ultrasonography; the prevalence of NAFLD was 30.2%</td>
<td>eGFR &lt;60 ml/min/1.73 m² and/or overt proteinuria (by urinary dipstick); 324 patients developed incident CKD during follow-up</td>
<td>Age, BMI, alcohol intake, hypertension, smoking, fasting glucose, baseline eGFR, triglycerides, HDL-cholesterol, LDL-cholesterol, HOMA-IR, C-reactive protein, incident cases of hypertension and diabetes</td>
<td>NAFLD was independently associated with increased risk of incident CKD (AHR 1.60; 95% CI 1.3-2.2)</td>
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<td>Targher G et al., Italy, 2008</td>
<td>Prospective cohort study (Valpolicella Heart Diabetes Study): 1,760 type 2 diabetic outpatients with preserved kidney function and no overt proteinuria, free of CVD and known chronic liver diseases at baseline. Follow-up: 6.5 years</td>
<td>Ultrasonography; prevalence of NAFLD was 73.2%</td>
<td>eGFR &lt;60 ml/min/1.73 m² and/or overt proteinuria; 547 patients developed incident CKD during follow-up (428 developed decreased eGFR alone, 112 developed proteinuria, irrespective of eGFR, and 7 developed kidney failure; no patients developed nephrotic syndrome)</td>
<td>Age, sex, BMI, waist circumference, blood pressure, smoking, diabetes duration, haemoglobin A1c, plasma lipids, baseline eGFR, use of antihypertensive, lipid-lowering, antiplatelet or hypoglycaemic agents</td>
<td>NAFLD was independently associated with increased risk of incident CKD (AHR 1.49; 95% CI 1.1-2.2)</td>
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<td>Arase Y et al., Japan, 2011</td>
<td>Retrospective cohort study: 5,561 middle-aged individuals with NAFLD and normal kidney function without overt proteinuria at baseline. Follow-up: 5.5 years</td>
<td>Ultrasonography and serum liver enzymes (serum GGT concentrations). Prevalence of NAFLD was 100%.</td>
<td>eGFR &lt;60 ml/min/1.73 m² and/or overt proteinuria (urinary dipstick); 263 patients developed incident CKD during follow-up</td>
<td>Age, sex, hypertension, diabetes, total cholesterol, triglycerides, HDL-cholesterol, liver enzymes, haemoglobin, white blood cell count, platelet count, baseline eGFR</td>
<td>Among patients with NAFLD, elevated serum GGT concentrations were independently associated with an increased risk of incident CKD (AHR 1.35; 95% CI 1.02-1.8). Measurement of NAFLD provided incremental risk reclassification beyond that of conventional CKD risk factors</td>
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<tr>
<td>Targher G et al., Italy, 2014</td>
<td>Prospective cohort study: 261 type 1 diabetic adult outpatients with normal kidney function, free of CVD and known chronic liver diseases at baseline. Follow-up: 5.2 years</td>
<td>Ultrasonography; prevalence of NAFLD was 50.2%</td>
<td>eGFR &lt;60 ml/min/1.73 m² and/or overt proteinuria; 61 patients developed incident CKD during follow-up (28 developed decreased eGFR with abnormal albuminuria, 21 developed reduced eGFR alone, and 12 developed macroalbuminuria alone; no patients developed kidney failure; no patients developed nephrotic syndrome)</td>
<td>Age, sex, diabetes duration, haemoglobin A1c, hypertension, baseline eGFR, presence of microalbuminuria</td>
<td>NAFLD was independently associated with an increased risk of incident CKD (AHR 1.85; 95% CI 1.03-3.3). Measurement of NAFLD provided incremental risk reclassification beyond that of conventional CKD risk factors</td>
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<td>Hub JH et al., South Korea, 2017</td>
<td>Prospective cohort study: 4,761 adults with normal kidney function and no overt proteinuria and free of CVD and known chronic liver diseases at baseline. Mean follow-up: 10 years</td>
<td>Fatty liver index (FLI); prevalence of NAFLD (defined as FLI ≥60) was 12.6%</td>
<td>eGFR &lt;60 ml/min/1.73 m²; 724 individuals developed incident CKD during follow-up</td>
<td>Age, sex, smoking, diabetes status, physical exercise, alcohol intake, protein intake, systolic blood pressure, total cholesterol, C-reactive protein, baseline eGFR</td>
<td>NAFLD (FLI ≥60) was independently associated with increased risk of incident CKD (AHR 1.46; 95% CI 1.19-1.79). FLI provided incremental risk reclassification beyond that of traditional renal risk factors</td>
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<td>Shen ZW et al., China, 2017</td>
<td>Prospective cohort study: 21,818 adults with normal kidney function and no overt proteinuria at baseline, who received routine health examination. Follow-up: 5 years</td>
<td>Serum liver enzymes (serum GGT concentrations)</td>
<td>eGFR &lt;60 ml/min/1.73 m² and/or overt proteinuria (urinary dipstick); 1,456 individuals developed incident CKD during follow-up</td>
<td>Age, sex, BMI, alcohol intake, serum creatinine, albumin, alanine aminotransferase, haemoglobin, white blood count, triglycerides, total cholesterol, hypertension, smoking, history of CVD, history of diabetes</td>
<td>NAFLD (i.e. top quartile of serum GGT levels) was independently associated with an increased risk of incident CKD (AHR 1.33; 95% CI 1.07-1.64)</td>
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<td>Kunutsor SK et al., Finland, 2017</td>
<td>Prospective cohort study (Kuopio Ischemic Heart Disease Study): 2,338 middle-aged men with normal kidney function at baseline. Median follow-up: 25.6 years</td>
<td>Serum liver enzymes (serum GGT concentrations)</td>
<td>eGFR &lt;60 ml/min/1.73 m²; 221 individuals developed incident CKD during follow-up</td>
<td>Age, BMI, systolic blood pressure, history of hypertension, smoking, history of coronary heart disease, diabetes, total cholesterol, HDL-cholesterol, alcohol intake, baseline eGFR</td>
<td>NAFLD (i.e. top quartile of serum GGT concentrations) was not independently associated with increased risk of incident CKD (aHR 0.97, 95% CI 0.64-1.47)</td>
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<td>Sinn DH et al., South Korea, 2017</td>
<td>Retrospective cohort study: 41,430 adults with normal kidney function and no overt proteinuria at baseline, free from known chronic liver diseases. Follow-up: 4.2 years</td>
<td>Ultrasonography; advanced NAFLD fibrosis assessed by the NFS (≥-1.455), FIB4 score (≥1.45) or APRI index (≥0.5); prevalence of NAFLD was 34.3%</td>
<td>eGFR &lt;60 ml/min/1.73 m²; 691 participants developed incident CKD during follow-up</td>
<td>Age, sex, BMI, smoking, alcohol intake, systolic blood pressure, haemoglobin A1C, LDL-cholesterol, use of hypoglycaemic and lipid-lowering medications, baseline eGFR, time-varying development of diabetes and hypertension over the follow-up</td>
<td>NAFLD was independently associated with increased risk of incident CKD (aHR 1.21, 95% CI 1.03-1.44). The association between NAFLD and CKD was consistent in all subgroups analysed. In addition, advanced NAFLD fibrosis (as detected by a NFS ≥-1.455) was associated with even a higher risk of incident CKD (aHR 1.59, 95% CI 1.31-1.93). When NAFLD participants were classified according to APRI index and FIB4 score, those with higher APRI index or FIB4 score also had an increasing risk of incident CKD</td>
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<td>Jang HR et al., South Korea, 2018</td>
<td>Retrospective cohort study: 1,525 adults with CKD (baseline mean eGFR 59 ml/min/1.73 m², 26% with overt proteinuria) free from known liver diseases, who underwent repeated health check-up examinations. Mean follow-up: 6.5 years</td>
<td>Ultrasonography; advanced NAFLD fibrosis assessed by the NFS (≥-1.455); prevalence of NAFLD was 40.9%</td>
<td>Annual percent decline in eGFR</td>
<td>Age, sex, year of visit, smoking, alcohol intake, BMI, hypertension, diabetes, dyslipidaemia, systolic blood pressure, haemoglobin A1c, LDL-cholesterol, triglycerides</td>
<td>NAFLD was independently associated with CKD progression. In multivariable-adjusted models, the average difference in annual percent change in eGFR decline comparing patients with NAFLD to those without NAFLD was -1.06% (-1.73%, -0.38%; p=0.002). The decline in eGFR associated with NAFLD was greater in patients with higher NFS, in those with proteinuria or with low eGFR (&lt;45 ml/min/1.73 m²) at baseline</td>
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<td>Wilechansky RM et al., United States, 2019&lt;sup&gt;44&lt;/sup&gt;</td>
<td>Prospective cohort study (Framingham Heart Study): 688 adults with normal kidney function and abnormal albuminuria at baseline, free from known liver diseases. Median follow-up: 12.5 years</td>
<td>Multidetector computed tomography</td>
<td>eGFR &lt;60 ml/min/1.73 m²; microalbuminuria; number of incident CKD cases was not reported</td>
<td>Age, sex, BMI, smoking, drinks per week, systolic/diastolic blood pressure, use of antihypertensive medications, HDL, total cholesterol, regular aspirin use, diabetes and follow-up interval</td>
<td>Liver fat (measured by the average liver attenuation on CT) was significantly associated with incident microalbuminuria and CKD in age- and sex-adjusted models. These relationships were not significant in multivariable-adjusted models. However, there was a discrepancy between the timing of baseline kidney function measurements (1998-2001) and CT assessment of liver fat (2002-2005) of a median of 4.1 years.</td>
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<td>Önnerhag K et al., Sweden, 2019&lt;sup&gt;45&lt;/sup&gt;</td>
<td>Retrospective cohort study (identified from a computerised register in Malmö): 144 adult patients with biopsy-proven NAFLD. Mean follow-up: 18.8 years</td>
<td>Non-invasive fibrosis scoring systems (i.e. FIB-4-index, NFS, APRI and BARD score)</td>
<td>eGFR &lt;60 ml/min/1.73 m²; 47 participants developed incident CKD during follow-up</td>
<td>Age, sex, overweight/obesity, prior CVD, hypertension, liver fibrosis stage</td>
<td>Both the intermediate and high-risk category of NFS and FIB-4 scores were independently associated with increased risk of incident CKD.</td>
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<tr>
<td>Park H et al., United States, 2019&lt;sup&gt;46&lt;/sup&gt;</td>
<td>Retrospective propensity-matched cohort analysis of the Truven Health MarketScan Database (2006-2015): 262,619 newly diagnosed patients with NAFLD and 769,878 propensity (1:3)-matched non-NAFLD patients. Follow-up: 9 years</td>
<td>International Classification of Diseases (ICD-9) codes</td>
<td>CKD stages 3-5 identified by the ICD-9-CM codes; There were 5,766 new CKD cases in the NAFLD cohort and 8,655 new CKD cases in non-NAFLD cohort</td>
<td>Age, sex, diabetes, hypertension, obesity, hyperlipidaemia, coronary artery disease, peripheral vascular disease, cerebrovascular disease, heart failure and chronic obstructive pulmonary disease, use of angiotensin-converting-enzyme inhibitors and angiotensin II receptor blockers, mean number of outpatient visits and mean number of inpatient visits, cirrhosis, decompensated cirrhosis and hepatocellular carcinoma</td>
<td>NAFLD was independently associated with increased risk of incident CKD (aHR 1.58, 95% CI 1.52-1.66). In the sensitivity analysis adjusting for time-varying covariates after NAFLD diagnosis, NAFLD persisted as a significant CKD risk factor (aHR 1.58, 95% CI 1.52-1.66).</td>
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eGFR was estimated by using either the 4-variable Modification of Diet in Renal Disease (MDRD) study equation or the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) study equation (that was used by the last 9 studies reported in this table).

aHR, adjusted hazard ratio; AST, aspartate aminotransferase; APRI, AST to platelet ratio index; BMI, body mass index; CKD, chronic kidney disease; CVD, cardiovascular disease; eGFR, estimated glomerular filtration rate; FIB4, fibrosis-4 score; GGT, gamma-glutamyltransferase; HOMA-IR, homeostasis model assessment-insulin resistance; NFS, NAFLD fibrosis score.
NAFLD and the risk of incident CKD appeared to be stronger in studies performed in Asian populations (n = 5 studies; random-effects HR 1.40; 95% CI 1.22–1.58; I² = 36.5%) than in European populations (n = 3 studies; random-effects HR 1.29; 95% CI 0.82–1.76; I² = 33.5%).

Taken together, the findings of these 2 meta-analyses7,48 clearly support the assertion that NAFLD identifies a group of individuals who are at increased risk of CKD, and who need more careful surveillance and treatment to reduce their risk of developing CKD. The results of these 2 meta-analyses also suggest that it is advanced NAFLD that carries a greater risk of incident CKD. This finding is in line with the results of a comprehensive meta-analysis supporting a strong link between the severity of NAFLD and increased risk of fatal and non-fatal cardiovascular outcomes.10

However, this question remains largely unsolved, and further prospective studies in larger cohorts of both Asian and non-Asian patients with biopsy-confirmed NAFLD are needed, in order to definitively prove whether the severity of NAFLD adversely affects the risk of developing CKD. That said, we believe that the evidence from these 2 meta-analyses and other more recent follow-up studies published in 2018 and 2019 (as listed in Table 1) calls for a more active and systematic search for CKD in patients with NAFLD with a view to implementing earlier and more aggressive surveillance and treatment to reduce their risk of developing CKD. The results of these 2 meta-analyses also suggest that it is advanced NAFLD that carries a greater risk of incident CKD. This finding is in line with the results of a comprehensive meta-analysis supporting a strong link between the severity of NAFLD and increased risk of fatal and non-fatal cardiovascular outcomes.10

In line with these observations, Vilar-Gomez et al. found that the histologic resolution of NASH and improvement in liver fibrosis stage were independently associated with an increase in eGFR values in a post hoc analysis of a published clinical trial that included 261 patients with biopsy-confirmed NASH, who were treated with lifestyle modifications during a period of 52 weeks.49 Recently, Önnerhag et al. examined the risk of overall mortality in patients with biopsy-proven NAFLD with the aim of investigating whether any increase in all-cause mortality was due to the presence of CKD.50 The authors measured eGFR values both at baseline and at the end of follow-up in a cohort of 120 middle-aged Swedish patients with biopsy-proven NAFLD, who were followed for a mean period of 19.5 years. The authors found that although patients with NAFLD and CKD had a significantly higher crude overall mortality rate than patients with NAFLD without CKD, the increased mortality risk was more strongly explained by an increased prevalence of metabolic comorbidities (including T2DM) rather than CKD.

![Fig. 3. Risk of CKD in patients with NAFLD.](image)

**Study limitations**

It should be noted that the observational design of the available studies (Table 1) does not allow us to establish a causal association between NAFLD and risk of CKD stage ≥3, and that it is currently uncertain whether NASH or NAFLD with advanced fibrosis carry a higher risk of incident CKD than simple steatosis. Most of the available studies used ultrasonography, which is the recommended first-line imaging method for detecting NAFLD in clinical practice. No studies used liver biopsy, which is considered the ‘gold standard’ for diagnosing and grading NAFLD.

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**Key point**

A number of observational studies have provided robust evidence of the association between NAFLD and CKD and the relevance of their coexistence on clinical outcomes.
staging NAFLD. Moreover, as shown in Table 1, the published cohort studies employed varying degrees of baseline adjustments for risk factors of CKD. In particular, almost all studies adjusted their results for BMI, but only a few of these studies additionally adjusted their results for body fat distribution, which plays a key role in the pathogenesis of NAFLD and CKD. An accurate assessment of abdominal visceral fat accumulation would be particularly important to better understand whether the association between CKD and NAFLD is affected by this metabolic risk factor. Other limitations include the use of the Modification of Diet in Renal Disease or the CKD-Epidemiology Collaboration study equations to calculate eGFR, neither of which are reliable in the presence of severe obesity or cirrhosis.51 Furthermore, most of the available cohort studies have been conducted in Asian countries, where large populations under regular health check-up programmes, including liver ultrasonography. Since Asian and non-Asian populations have different genetic/cultural backgrounds, dietary factors and adipose tissue distributions, we believe that additional studies should be conducted in non-Asian populations. Another potential limitation is that no large prospective studies are available that have examined the rates of CKD progression to kidney failure (stage 5 CKD) in cohorts of patients with NAFLD, nor in cohorts of patients with advanced CKD. Finally, none of these studies have used renal biopsy to examine the specific renal pathology associated with NAFLD. So, it is currently uncertain if NAFLD is associated with a specific type of kidney disease, although we suggest that it is reasonable to assume that NAFLD may promote kidney injury, mostly through accelerated atherothrombosis.

Nevertheless, it is important to consider that in a recent analysis of the Third National Health and Nutrition Survey database that included a total of ~11,700 Americans, Paik et al. showed that amongst patients with ultrasound-detected NAFLD, the presence of moderate to advanced stages of CKD was independently associated with increased all-cause mortality over a mean follow-up period of 19 years.52 These findings point out that identification of CKD in patients with NAFLD has important prognostic implications. As also suggested by the authors, these data should inform clinicians and policy makers to identify those at the highest risk of adverse outcomes so that appropriate management strategies can be implemented.

Putative mechanisms linking NAFLD with CKD

T2DM and metabolic syndrome

Although not conclusive, as discussed above, the current epidemiological evidence suggests that NAFLD is an independent risk factor for CKD and that the presence of NAFLD and associated features of the MetS53 may be causally involved, at least in part, in the development and progression of CKD.

More than one-third of patients with NAFLD have impaired renal function and impaired renal function in patients with NAFLD is also associated with the severity of liver disease and presence of T2DM.54,55 When common diseases co-exist and share common risk factors, it can be difficult to disentangle causal relationships and understand the role of potential confounders. T2DM or MetS could be examples of confounding conditions linking NAFLD and CKD. In centrally obese individuals with T2DM, insulin resistance frequently occurs alongside other cardiometabolic risk factors that increase the risk of both NAFLD and CKD. The clustering of cardiometabolic risk factors that occur with visceral obesity and insulin resistance are encapsulated within the features of MetS, such as atherogenic dyslipidaemia, increased blood pressure and dysglycaemia.51

As mentioned in the Introduction, CKD stage ≥3 is defined by either abnormal albuminuria or decreased eGFR values. Whereas abnormal albuminuria (or overt proteinuria) is strongly associated with microvascular damage in renal glomeruli in diabetes and is a classical microvascular complication of diabetes, decreased eGFR values are more strongly associated with macrovascular damage. Macrovascular disease is strongly associated with CVD risk factors that occur alongside MetS, such as hypertension and atherogenic dyslipidaemia, as well as other risk factors not related to MetS, such as increased low-density lipoprotein-cholesterol concentrations. Thus, the clustering of cardiometabolic risk factors, occurring in patients with NAFLD and commonly referred to as the MetS, has the potential to cause both microvascular and macrovascular damage, giving rise to CKD.

CKD also commonly occurs alongside the MetS.56,57 Consequently, it can be very difficult to disentangle the differential effects of insulin resistance, visceral obesity and the linked macrovascular and microvascular risk factors on the kidneys, from the consequences of liver disease perse. That said, this may be a moot point, given the close inter-relationships of insulin resistance, visceral adiposity and other MetS features with liver disease in NAFLD. Given that all these MetS risk factors are closely inter-related in NAFLD, even in the presence of known genetic modifiers of NAFLD severity, such as the common genetic variants in patatin-like phospholipase domain-containing 3 (PNPLA3) and transmembrane 6 superfamily member 2 (TM6SF2),58,59 it may be an arcane point to try and dissect out the relative contributions of individual NAFLD-associated hepatic or cardiometabolic risk factors on kidney function. In fact, since these risk factors cluster tightly together and NAFLD may also promote the development of these cardiometabolic risk
As mentioned, visceral obesity is a classical feature of the MetS. Many potential mechanisms by which accumulation of visceral fat causes chronic inflammation are well described\(^{65-68}\) and a detailed discussion of these is beyond the scope of this review. However, briefly, alongside increases in visceral obesity and ectopic fat accumulation, there is an increase in plasma concentrations of non-esterified fatty acids (NEFAs) and a failure to adequately suppress NEFA concentrations during hyperinsulinaemia.\(^{69-72}\) With the increase in NEFA supply to the liver, a link has been proposed between NEFA overflow from the expanded and dysfunctional visceral adipose tissue and activation of hepatic macrophages that is independent of BMI.\(^{73}\) Activation of hepatic macrophages and hepatic inflammation is associated with an increase in proinflammatory cytokines\(^{74}\) and hepatic/systemic insulin resistance.\(^{75,76}\) Increased activity of the renin-angiotensin-aldosterone system\(^ {77}\) and oxidative stress mediated by proinflammatory and profibrotic mediators.\(^ {78}\) The liver-kidney crosstalk in NAFLD also includes the role of the energy sensor 5′-AMP-activated protein kinase (AMPK) and its regulation of fetuin-A and adiponectin. In the liver and kidney, AMPK is pivotal to directing hepatocytes and renal podocytes to compensatory and potentially deleterious pathways, leading to inflammatory and profibrotic cascades culminating in end-organ damage.\(^ {79}\) Collectively, NAFLD and CKD share common proinflammatory and profibrotic mechanisms of disease progression.\(^ {80,81}\) Experimental evidence also supports a role of the inflammasome and innate immune system in CKD.\(^ {82,83}\) Therefore, all of these factors and pathways could indicate a causal link between NAFLD and CKD, whereby NAFLD increases the risk of incident CKD.

Not only are there many traditional risk factors shared between NAFLD, CKD and T2DM/MetS, but it has now become clear that newer and emerging risk factors are also frequently present with each of these conditions. These newer risk factors include perturbation of the intestinal microbiota (dysbiosis) with associated inflammation, intestinal dysfunction and platelet activation. The role of these newer risk factors in NAFLD and CKD will be discussed in the following sections.

**Dysbiosis and perturbed intestinal function affecting NAFLD and CKD**

Emerging experimental evidence suggests a role of the intestinal microbiota in the pathogenesis of both CKD\(^ {32,83}\) and NAFLD.\(^ {54-87}\) Fig. 4 shows mechanisms and factors potentially linking intestinal dysbiosis, visceral adipose tissue dysfunction, NAFLD and CKD. With perturbation of the gut microbiota (dysbiosis), there is an increase in gram negative organisms, lipopolysaccharide, gut permeability, secondary bile acids (BAs) and renal toxins that may increase the risk of development and progression of both NAFLD\(^ {88,89}\) and CKD.\(^ {32,84}\) Intestinal microbiota-generated production of uraemic toxins (e.g., trimethylamine, cresol and indole)\(^ {83,90-93}\) has the potential to further damage renal, hepatic and cardiovascular function through inflammatory, oxidative and fibrotic pathways. The metabolism of the amino acids tyrosine and phenylalanine by a variety of obligate or facultative anaerobes, including the genera *Bacteroides*, *Lactobacillus*, *Enterobacter*, *Bifidobacterium*, and especially *Clostridium difficile*, results in the increased production of para-cresyl and the conjugate para-cresyl sulfate.\(^ {94,95}\) *Escherichia coli* has been shown to metabolise tryptophan, resulting in the production of indole that is metabolised in the liver to the uraemic toxin indoxyl sulfate.\(^ {94,95}\) A variety of other potentially nephrotoxic metabolites are also produced, such as ammonia, thiols and phenols.\(^ {83}\)

A complex interaction also exists between the gut microbiota and BA metabolism in NAFLD,\(^ {89}\) wherein a diverse range of BAs can be detected in the plasma and have the potential to influence development and progression of the disease. Secondary BAs are generated from the 1–5% of primary BAs that are not re-absorbed in the jejunum. These BAs enter the ileum and colon and are modified by the gut microbiota hydrolases and dehydroxylases to create secondary BAs, such as deoxycholic acid and lithocholic acid. Further bacterial enzymes that include epimerases, oxidases and esterases, are capable of further modifying BAs before they are excreted in the stool. There is now evidence that the dynamic interaction existing between the microbiota and the BA pool can be modified by certain microbiota species to change the BA profile.\(^ {96}\) Specifically, NASH is associated with changes in the intestinal microbiota...
Composition and metabolome, an intestinal and systemic inflammatory response, and BA profiles, and it has also been suggested that the composition of the gut microbiome associated with dysregulation of BA biosynthetic pathways may contribute to the persistence of NAFLD. Modification of the BA profile may be important in the treatment of NAFLD but mechanistic studies are required to elucidate causal links between intestinal dysbiosis, NAFLD and CKD.

Dietary changes mediating a link between NAFLD, dysbiosis and CKD
Increased consumption of sugar-sweetened beverages is linked with the development of NAFLD, hypertension, MetS and T2DM in both laboratory animals and humans, although the association may be confounded by excess calorie intake or by unhealthy lifestyles. Today the most commonly consumed sugar is high fructose corn syrup. However, a causal role of excessive fructose consumption in the development of these metabolic diseases is still debated and the molecular mechanisms by which fructose elicits effects on dysregulated liver metabolism remain incompletely understood. Increased dietary fructose intake is associated with NASH and increased dietary fructose intake is also associated with increased serum uric acid concentrations in children and adolescents. Emerging experimental data suggest that increased dietary fructose might induce NAFLD, at least in part, due to the generation of uric acid during fructose metabolism that results in mitochondrial oxidative stress and impairment in ATP production. Although it has been thought that most fructose in the body is derived from dietary fructose intake (principally sugar/corn syrup sweetened drinks), it has recently been shown that endogenous fructose can also be generated in the liver with activation of the poloy pathway. In this pathway, glucose is converted to sorbitol by aldose reductase and sorbitol is converted to fructose by sorbitol dehydrogenase. Aldose reductase is a NADPH-dependent aldo-
Platelet activation as a mediator of the link between NAFLD and CKD

The hepatic microenvironment plays a crucial role in liver disease development, as hepatic stellate cells, resident liver macrophages (Kupffer cells), endothelial cells, extracellular matrix and a variety of immune cells or platelets may interact in complex and intertwined signalling pathways. As mentioned, dyslipidaemia is a key feature of MetS, NAFLD and CKD and the specific dyslipidaemia involves an increase in hepatic-derived triglyceride-rich lipoproteins. Oxidative stress is a key feature of NAFLD and CKD: both oxidative stress and an increase in triglyceride-rich lipoproteins (such as very low-density lipoprotein and remnant lipoproteins) are key regulators of platelet activation. With oxidative stress and kidney dysfunction, there is a reduction in anti-oxidant protective factors produced by the kidneys, such as the Klotho protein, and generation of metabolites such as plasma F2-isoprostanes, 8-oxo-7,8-dihydro-2′-deoxyguanosine, malonyldialdoxyde, advanced oxidation protein products, carbamylated proteins, asymmetric dimethylarginine and oxidised lipoproteins. When platelets are activated, alpha granules and dense granules are released containing multiple proinflammatory cytokines, chemokines and growth factors. These include chemokine (CXC motif) ligand 4 (CXCL4), endothelial growth factor (EGF), interleukin-6 (IL-6), platelet-derived growth factor (PDGF), serotonin, insulin-like growth factor 1 (IGF-1), transforming growth factor (TGF)-beta, tumour necrosis factor (TNF)-alpha, vascular endothelial growth factor A (VEGF-A), hepatocyte growth factor (HGF) and fibroblast growth factor (FGF). The release of TGF-beta, PDGF, serotonin and CXCL4 can cause progression of liver disease by activating stellate cells with a consequent increase in extracellular matrix production. With CKD, there is also increased platelet activation, and an attenuated response to dual antiplatelet therapy, compared to patients without CKD. Dysbiosis may also act to promote increased platelet activation since indoxyl sulfate activates platelets. Decreased urinary Klotho protein levels have been identified as one of the earliest biomarkers of CKD progression, and the Klotho protein is also able to modulate the effect of indoxyl sulfate on platelet hyperactivity and thrombus formation, protecting against indoxyl sulfate-induced atherosclerosis in mice with CKD. Thus, increased oxidative stress, intestinal dysbiosis, an increase in hepatic-derived triglyceride-rich lipoproteins and platelet activation are all closely inter-related with the potential for a “vicious” spiral of worsening liver and kidney disease in NAFLD and CKD. Recently, Malehmir et al. showed that platelet number, platelet activation and platelet aggregation are increased in NASH but not in simple steatosis or insulin resistance. Antiplatelet therapy (aspirin/clopidogrel, ticagrelor) but not non-steroidal anti-inflammatory drug treatment with sulindac also prevented NASH and subsequent development of hepatocellular carcinoma in a murine hIL4r-alpha-/GP1b-alpha transgenic mouse model of NASH. In addition, intra-vital microscopy also showed that antiplatelet therapy reduced intrahepatic platelet accumulation and the frequency of platelet-immune cell interaction, thereby limiting hepatic immune cell trafficking. Taken together, these experimental results suggest that blocking platelet activation might ameliorate NASH and subsequently decrease the risk of developing hepatocellular carcinoma.

Premature ageing and age-related changes

Older age is a risk factor for NAFLD, CKD and T2DM. Decreased urinary Klotho protein occurs with ageing, and with decreased Klotho protein excretion, there is an associated vascular phenotype of medial calcification, intima hyperplasia, endothelial dysfunction, arterial stiffening, hypertension and impaired angiogenesis. Decreased urinary Klotho protein has been also identified as one of the earliest biomarkers of CKD progression and the Klotho gene was identified first as a putative ageing-suppressor gene that extended life span when overexpressed, and accelerated ageing-like phenotypes when disrupted in mice. As mentioned, the Klotho protein modulates the effect of indoxyl sulfate on platelet hyperactivity, and thus there is the potential for low levels of Klotho...
protein to mediate the increase in platelet reactivity that occurs with ageing.131 Older patients with NAFLD are at a higher risk of CKD as a function of their increased age, but with advancing age, obesity and increased serum uric acid concentrations there is also increased risk of developing CKD.132 Age-related changes in the liver may also occur with alterations in hepatic sinusoidal endothelial cells,133 increases in the hepatokine fetuin-A and decreases in adiponectin, potentially linking MetS, NAFLD and CKD.79

In summary, these new mechanistic data suggest plausible mechanisms and new pathways linked to MetS, intestinal dysbiosis, excessive fructose consumption, platelet activation and ageing that might, at least in part, mediate links between NAFLD and the risk of CKD. However, more research is needed to better understand if experimental models of NAFLD/NASH, initiated by primary changes in lipid storage in the liver, ultimately lead to CKD.

Effect of PNPLA3 polymorphism on renal function

Several susceptibility gene variants predisposing to NAFLD have been consistently identified in different populations.58,59 Among the genetic factors that may influence the onset and progression of NAFLD, the minor allele G of rs738409, i.e. a non-synonymous single nucleotide polymorphism in the PNPLA3 gene encoding an Ile148Met change, has been recognised as a major common genetic variant associated with a greater predisposition to NASH and progressive liver fibrosis in both paediatric and adult populations.58,59

Emerging evidence is now suggesting that the G allele of rs738409 is significantly associated with decreased eGFR values, irrespective of established renal risk factors and presence of NAFLD, across different ethnicities and patient populations. Indeed, as summarised in Table 2, there are now half a dozen studies that have examined whether, and to what extent, the PNPLA3 rs738409 polymorphism is associated with decreasing kidney function in both adults and children or adolescents.134-140 For instance, our group recently showed for the first time that the presence of the risk allele (G) of rs738409 was strongly associated with both decreasing eGFR and increasing 24-hour urinary protein excretion in a sample of 142 overweight Italian children/adolescents with biopsy-proven NAFLD.138 Notably, these associations were independent of sex, age, measures of adiposity, blood pressure, HOMA-estimated insulin resistance and also the histologic severity of NAFLD (i.e. NASH and liver fibrosis stage).138 Similarly, Sun et al. reported that the PNPLA3 GG genotype was significantly associated with a higher risk of prevalent CKD, abnormal albuminuria or increased levels of urinary neutrophil gelatinase-associated lipocalin (i.e. a reliable marker of renal tubular injury) in 217 Chinese adults with biopsy-confirmed NAFLD.137 Also in this study, PNPLA3 GG genotype remained significantly associated with renal glomerular and tubular injury after adjusting for age, sex, BMI, waist circumference, hypertension, diabetes, HOMA-estimated insulin resistance, hyperuricemia, and histologic severity of NAFLD.137 In another study involving 740 elderly Japanese individuals, Oniki et al. found that the PNPLA3 GG genotype was associated with lower levels of eGFR, independently of common renal risk factors and presence of ultrasound-detected NAFLD, especially in those with normal body weight.134 Notably, in a subgroup of these individuals, the authors also showed that the PNPLA3 GG genotype was associated with a significant decline in eGFR over a mean follow-up of 5.5 years.134

To date, the putative mechanisms underlying the association between the G allele of rs738409 and decreasing kidney function are not entirely understood. The published studies134-140 show that the association between the PNPLA3 GG genotype and kidney dysfunction was largely independent of the shared renal/metabolic risk factors. It is reasonable to hypothesise that the G allele of rs738409, which is highly expressed in liver sinusoidal pericytes,141 might also exert direct adverse effects on the kidneys. These nephrotoxic effects may occur via activation of renal pericytes, as renal pericytes are stromal cells that play a key role in angiogenesis and in regulating renal medullary and cortical blood flow, promoting renal fibrogenesis and glomerulosclerosis.142,143 However, further research is needed to better understand the role of the PNPLA3 rs738409 polymorphism on the development of glomerular and interstitial fibrosis.

If confirmed in future larger studies, we believe that the results of these studies may have important clinical implications, because they support the notion that PNPLA3 genotyping might be useful not only to identify individuals with greater susceptibility to NAFLD development and progression, but also for those patients with NAFLD who may be at higher risk of CKD, thus promoting the implementation of specific prevention programmes and treatment strategies for CKD among carriers of the PNPLA3 rs738409 GG genotype.

Conclusions

This review article outlines the existence of a strong association between the presence and severity of NAFLD and the increased prevalence and incidence of CKD, independent of obesity, hypertension, T2DM and other common cardio-renal risk factors. The data also suggest that PNPLA3 genotyping might be useful not only to identify those with greater susceptibility to NAFLD development and progression, but also to identify a subgroup of patients with NAFLD who are at higher

Key point

The PNPLA3 rs738409 polymorphism which is associated with a predisposition to NASH has recently been shown to associate with worse kidney function.
Table 2. Association between PNPLA3 rs738409 polymorphism and kidney dysfunction both in adults and in children or adolescents (ordered by publication year).

<table>
<thead>
<tr>
<th>Authors, Country, Year (Ref.)</th>
<th>Study design and population</th>
<th>Kidney outcome measures</th>
<th>Covariate adjustments</th>
<th>Main results</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Adults</strong></td>
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<tr>
<td>Oniki K et al., Japan, 2015[134]</td>
<td>Cross-sectional of 591 elderly individuals without known liver diseases (472 without NAFLD and 119 with NAFLD on ultrasonography) recruited during a health screening program and selected from an initial cohort of 740 individuals. Among these individuals, a retrospective longitudinal analysis with a median follow-up of 5.0 (range: 1.0–6.0 years) was also performed in 341 subjects (51 with NAFLD and 290 without NAFLD)</td>
<td>eGFR&lt;sub&gt;CKD-EPI&lt;/sub&gt; (considered as continuous measure)</td>
<td>Age, sex, BMI, diabetes, hypertension, dyslipidaemia and presence of NAFLD on ultrasonography</td>
<td>PNPLA3 GG genotype was independently associated with lower eGFR values compared with carriers of the CC genotype, only in the subgroup of individuals with normal body weight (but not in those with overweight or obesity) in cross-sectional analyses. This association was also replicated in the longitudinal analyses</td>
</tr>
<tr>
<td>Musso G et al., Italy, 2015[135]</td>
<td>Cross-sectional study of 202 nonobese, nondiabetic adults (61 non-cirrhotic biopsy-proven NAFLD and 81 controls)</td>
<td>CKD (i.e. eGFR&lt;sub&gt;CKD-EPI&lt;/sub&gt; &lt;60 ml/min/1.73 m&lt;sup&gt;2&lt;/sup&gt; and/or microalbuminuria); microalbuminuria (30 - 300 mg/g)</td>
<td>Not specified</td>
<td>PNPLA3 GG or CG genotype (combined) was significantly associated with a higher risk of prevalent CKD, lower eGFR or microalbuminuria both in NAFLD patients and in controls</td>
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<tr>
<td>Mantovani A et al., Italy, 2019[136]</td>
<td>Cross-sectional study of 101 Caucasian post-menopausal women with non-insulin treated type 2 diabetes mellitus without known liver diseases</td>
<td>CKD (i.e. eGFR&lt;sub&gt;CKD-EPI&lt;/sub&gt; &lt;60 ml/min/1.73 m&lt;sup&gt;2&lt;/sup&gt; and/or abnormal albuminuria ≥30 mg/g)</td>
<td>Age, duration of diabetes, haemoglobin A1c, HOMA-IR, systolic blood pressure, hypertension treatment and presence of NAFLD on ultrasonography</td>
<td>PNPLA3 GG genotype was independently associated with lower eGFR values and higher prevalence of CKD compared with the CC or GC genotype</td>
</tr>
<tr>
<td>Sun DQ et al., China, 2019[137]</td>
<td>Cross-sectional study of 217 adults with non-cirrhotic biopsy-proven NAFLD</td>
<td>CKD (i.e. eGFR&lt;sub&gt;CKD-EPI&lt;/sub&gt; &lt;60 ml/min/1.73 m&lt;sup&gt;2&lt;/sup&gt; and/or abnormal albuminuria ≥30 mg/g); abnormal albuminuria ≥30 mg/g, urinary NGAL levels ≥31.2 ng/ml</td>
<td>Age, sex, BMI, waist circumference, hypertension, diabetes, HOMA-IR, hyperuricemia, presence of NASH (i.e. defined as a NAS ≥5) and histologic stage of fibrosis</td>
<td>PNPLA3 GG genotype was independently associated with a higher risk of prevalent CKD, abnormal albuminuria or increased NGAL levels, especially in patients with persistently normal serum alanine aminotransferase levels</td>
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<td><strong>Overweight/obese children or adolescents</strong></td>
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<tr>
<td>Targher G et al., Italy, 2019[138]</td>
<td>Cross-sectional study of 142 Caucasian children/adolescents with biopsy-proven NAFLD</td>
<td>eGFR (using the Bedside Schwartz equation); 24-hour proteinuria (both considered as continuous measures)</td>
<td>Age, sex, systolic blood pressure, waist circumference, presence of NASH (i.e. defined as a NAS ≥5) and histologic stage of liver fibrosis</td>
<td>PNPLA3 GG genotype was independently associated with lower eGFR values and increasing 24-hour proteinuria</td>
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<tr>
<td>Marzuillo P et al., Italy, 2019[139]</td>
<td>Cross-sectional study of 591 Caucasian children/adolescents with obesity</td>
<td>eGFR (using the Schwartz equation) considered as continuous measure</td>
<td>Sex, duration of obesity, HOMA-IR score, BMI, LDL-cholesterol, triglycerides</td>
<td>PNPLA3 GG genotype was independently associated with lower eGFR values compared with the CC or GC genotype, only in the subgroup of children with NAFLD (defined by ultrasonography and/or serum ALT &gt;40 IU/L)</td>
</tr>
<tr>
<td>Di Costanzo et al., Italy, 2019[140]</td>
<td>Cross-sectional study of 230 overweight/obese children (105 with NAFLD defined as liver fat fraction ≥5% by magnetic resonance imaging)</td>
<td>eGFR (using the Schwartz equation); abnormal albuminuria ≥30 mg/g</td>
<td>Age, sex, pubertal status, diastolic blood pressure, waist circumference and presence of NAFLD on magnetic resonance imaging</td>
<td>PNPLA3 G risk allele was not independently associated with eGFR &lt;90 ml/min/1.73 m&lt;sup&gt;2&lt;/sup&gt; and/or abnormal albuminuria</td>
</tr>
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</table>

NB: In all aforementioned studies, the PNPLA3 genotyping was determined on blood samples. BMI, body mass index; CKD, chronic kidney disease; eGFR<sub>CKD-EPI</sub>, estimated glomerular filtration rate (estimated by the CKD-EPI equation); HOMA-IR, homeostasis model assessment-insulin resistance; NAFLD, non-alcoholic fatty liver disease; NAS, NAFLD activity score; NASH, non-alcoholic steatohepatitis; NGAL, neutrophil gelatinase-associated lipocalin; PNPLA3, patatin-like phospholipase domain-containing protein-3.
risk of developing CKD. Despite the convincing evidence linking NAFLD with a higher risk of CKD, it remains to be definitively established whether a causal association also exists. Moreover, it should also be noted that none of the available studies have used renal biopsy to examine the specific renal morphology/pathology associated with NAFLD, and therefore it is uncertain if NAFLD is associated with a specific type of kidney disease, although it is reasonable to assume that NAFLD may promote kidney damage, mostly through accelerated atherothrombosis. We suggest that future prospective and interventional studies of well-characterised cohorts of patients with biopsy-proven NAFLD are required to try and better elucidate whether it is the presence and severity of NAFLD, or whether it is the presence of co-existing risk factors that increases risk of incident CKD. In the meantime, however, given the close link between NAFLD and CKD, more careful surveillance of these patients is warranted.

Abbreviations

ACR, albumin-to-creatinine ratio; aHR, adjusted hazard ratio; AMPK, AMP-activated protein kinase; APRI, AST to platelet ratio index; AST, aspartate aminotransferase; BAs, bile acids; BMI, body mass index; CKD, chronic kidney disease; CVD, cardiovascular disease; CXCL4, chemokine (CXC motif) ligand 4; DAGs, diacylglycerols; EGF, endothelial growth factor; eGFR, estimated glomerular filtration rate; FGF, fibroblast growth factor; FLI, fatty liver index; GGT, gamma-glutamyltransferase; HGF, hepatocyte growth factor; HOMA, homeostatic model assessment; HR, hazard ratio; IGF-1, insulin-like growth factor 1; IL-6, interleukin 6; LCFAs, long-chain fatty acids; LPS, lipopolysaccharide; MDRD, Modification of Diet in Renal Disease; MetS, metabolic syndrome; NAFLD, non-alcoholic fatty liver disease; NAS, NAFLD activity score; NASH, non-alcoholic steatohepatitis; NEFA, non-esterified fatty acids; NGS, NAFLD fibrosis score; NGAL, neutrophil gelatinase-associated lipocalin; PAI-1, plasminogen activator inhibitor 1; PDGF, platelet-derived growth factor; PNP, proprotein convertase subtilisin/kexin type 2; PON-1, paraoxonase 1; SCFAs, short-chain fatty acids; TGF, transforming growth factor; TM6SF2, transmembrane 6 superfamily member 2; TMA, trimethylamine; TMAO, trimethylamine oxide; TNF, tumour necrosis factor; VEGF, vascular endothelial growth factor; VLDL, very low-density lipoprotein.

Conflict of interest

The authors declare no conflicts of interest that pertain to this work.

Please refer to the accompanying ICMJE disclosure forms for further details.

Authors’ contributions

The authors contributed equally to this manuscript.

Supplementary data

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References


