Glyphosate Excretion is Associated With Steatohepatitis and Advanced Liver Fibrosis in Patients With Fatty Liver Disease

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Introduction

Nonalcoholic fatty liver disease (NAFLD) is currently the most common chronic liver disease in developed countries ¹ Patients with non-alcoholic steatohepatitis (NASH) are considered to be at a higher risk of fibrosis progression and development to cirrhosis and hepatocellular carcinoma.

Among potential environmental contributors to the pathophysiology of NAFLD are exposure to pesticides and herbicides ². Glyphosate, the primary weed-killing ingredient in Roundup[®], is sprayed on genetically modified (GM) crops and on many non-GM grain crops and is found in these crops at harvest ³.

Rodents chronically fed with a low dosage of glyphosate exhibit signs of hepatotoxicity, liver congestion, necrosis, and DNA damage of the liver cells ⁴⁵⁶. This study examined excretion levels of glyphosate and its primary metabolite aminomethylphosphonic acid (AMPA) in a well-characterized and prospectively recruited cohort of patients with biopsyproven NAFLD.

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Author Contributions. Paul J. Mills: Study concept and design; acquisition of data; analysis and interpretation of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content; statistical analysis. Cyrielle Caussy: Study concept and design; acquisition of data; interpretation of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content. Rohit Loomba: Study concept and design; acquisition of data; interpretation of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content.

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COIs. The authors report no conflicts of interest.

Methods

Participants were originally recruited as part of a larger study between September 2012 and March 2018 at the University of California at San Diego (UCSD) NAFLD Research Center. As previously described ⁷, patients with suspected NAFLD with a clinical indication for liver biopsy underwent a careful evaluation for other causes of hepatic steatosis and liver disease through a standardized research visit including detailed medical and alcohol use history as well as anthropometric and physical examination. Histologic scoring was done using the Nonalcoholic Steatohepatitis Clinical Research Network Histologic Scoring System. Prior to conducting statistical analyses, cases were grouped as definite NASH or NAFLD not NASH. This study was approved by the UCSD Institutional Review Board. Informed written consent was obtained from each study participant.

Each patient provided a fasting urine sample which was stored at -80° C. Urine samples were analyzed for glyphosate and AMPA using HPLC coupled with mass spectrometry. Using the formula [(glyphosate + 1.5) × AMPA] we calculated the glyphosate residue, which provides an estimate of dietary intake and exposure to residues.

ANOVA, ANCOVA, Chi-square, and multivariate general linear models covarying for age, gender and BMI were used (SPSS Version 24.0 software package (IBM, Armonk, NY)). Dependent variables were glyphosate, AMPA and glyphosate residue. Results were considered statistically significant at the p 0.05 level. Prior to statistical analyses, data were tested for normality and homogeneity of variance.

Results

Patient characteristics are presented in Table 1. Neither age nor BMI were significantly related to glyphosate, AMPA or glyphosate residue. Similarly, neither diabetes status nor race / ethnicity were significantly related to glyphosate, AMPA or glyphosate residue. Glyphosate [women 0.373 μ g/L (SD=0.41)] vs [men 0.215 μ g/L (SD=0.17) (F=5.18; p=0.025)] and glyphosate residue [women 0.833 μ g/L (SD=0.67) vs men 0.594 μ g/L (SD=0.38) (F=4.09; p=0.046)] were elevated in women as compared to men.

In multivariate models adjusting for age, sex, and BMI, as compared to patients without NASH, AMPA (F=5.39; p=0.022) and glyphosate residue (F=7.43; p=0.008) were elevated in patients with definite NASH (Table 1). When compared to patients without advanced fibrosis (Stages 0 & 1), patients with advanced fibrosis (Stages 2, 3, & 4) had, respectively, elevated AMPA [0.196 μ g/L (SD=0.20) vs 0.365 μ g/L (SD=0.33) (F=9.44; p=0.003), glyphosate residue [0.525 μ g/L (SD=0.38) vs 0.938 μ g/L (SD=0.372) (F=11.9; p=0.001), and glyphosate [0.230 μ g/L (SD=0.19) vs 0.351 μ g/L (SD=0.45) (F=4.13; p=0.046).

Discussion

We report that glyphosate excretion is significantly higher in patients with NASH compared to patents without NASH. In addition, we also report a significant dose-dependent increase of glyphosate exposure with increase in fibrosis stages.

For individuals not working in the agricultural or horticultural industries, the primary route of glyphosate exposure is through ingestion of Roundup[®]-treated GM foods and/or non-GM crops such as wheat and oats ³. Glyphosate excretion was elevated in women, which presumably reflected an increased exposure to glyphosate.

While there are strengths to this study, including the use of a well-characterized cohort using liver biopsy for the diagnosis of NASH and stage of liver fibrosis, we acknowledge limitations, including no information on dietary intake or occupation and no patients without NAFLD. We did not find an association between glyphosate excretion and BMI, suggesting that glyphosate intake was independent of total caloric intake.

As far as potential mechanisms of glyphosate on the liver, Mesnage et al. showed that rats fed glyphosate have disrupted liver mitochondrial oxidative phosphorylation leading to proteome disturbances reflecting peroxisomal proliferation, steatosis and necrosis, a profile consistent with NAFLD and its progression to NASH ⁴. Other studies show that glyphosate inhibits fatty acid oxidation and increases fat and cholesteryl ester levels in mice livers, leading to increased lipid mass per gram of liver ⁸.

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Abbreviations.

NAFLD nonalcoholic fatty liver disease

NASH Non-alcoholic steatohepatitis

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Table 1.

Patient Characteristics (Mean \pm SD)

	All (n=97)	Not NASH (n=34)	Definite NASH (n=63)	p-value*
Demographics				
Age (years)	50.5 (13.2) (range 19 – 74)	47.3 (12.5) (range 21 – 72)	51.9 (13.8) (range 19 – 74)	0.249
Male (%)	41.2	53.3	36.5	0.124
White (%)	42.2	39.6	47.1	0.27
Hispanic or Latino (%)	35.0	35.2	34.9	0.81
BMI (kg/m ²)	31.8 (7.0)	29.2 (3.5)	33.0 (7.8)	0.014
Clinical				
Type 2 Diabetes (%)	38.10	17.6	44.4	0.008
Biological data				
AST (U/L)	47.0 (31.7)	35.4 (13.2)	52.5 (36.2)	0.014
ALT (U/L)	65.1 (43.2)	56.3 (24.8)	69.3 (48.3)	0.179
Hemoglobin A1c (%)	6.19 (1.2)	5.71 (0.9)	6.45 (1.2)	0.002
Triglycerides (mg/dL)	148.4 (66.3)	168.0 (78.5)	143.3 (58.8)	0.813
Total cholesterol (mg/dL)	185.0 (46.8)	195.3 (31.8)	180.2 (51.9)	0.153
HDL-cholesterol (mg/dL)	44.8 (12.9)	44.2 (10.4)	45.1 913.9)	0.753
LDL-cholesterol (mg/dL)	106.7 (29.7)	117.7 (25.0)	101.8 (30.4)	0.020
Platelet count (10 ³ μL)	238880 (72375)	238566 (83977)	239032 (66794)	0.977
Histology				
Fibrosis (%)	(n=74)	(n=27)	(n=47)	<0.001
Stage 0	39.0	88.9	10.6	
Stage 1	5.40	3.70	6.4	
Stage 2	27.1	3.70	40.4	
Stage 3	23.1	3.70	34.1	
Stage 4	5.40	0	8.5	
Steatosis (%)	(n=97)	(n=34)	(n=63)	0.017
Stage 0	0.1	0	1.5	
Stage 1	37.7	60.1	26.6	
Stage 2	39.3	23.3	47.4	
Stage 3	22.9	16.6	24.5	
Lobular inflammation (%)	(n=90)	(n=32)	(n=58)	0.003
Stage 0	4.5	13.3	0	
Stage 1	47.7	60.0	41.4	
Stage 2	45.5	23.4	56.9	
Stage 3	2.3	3.3	1.7	
Ballooning (%)	(n=97)	(n=34)	(n=63)	<0.001
Stage 0	38.7	84.7	13.3	

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All (n=97) Not NASH **Definite NASH** p-value* (n=34)(n=63) 46.3 15.3 63.4 Stage 1 Stage 2 15.0 0 23.3 **Glyphosate Excretion** Glyphosate (µ/L) 0.308 (0.34) 0.241 (0.18) 0.344 (0.40) 0.164 AMPA (μ/L) 0.284 (0.28) 0.197 (0.18) 0.331 (0.31) 0.022 Glyphosate Residue (µ/L) 0.735 (0.58) 0.538 (0.35) 0.841 (0.65) 0.008

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Data are provided as mean values \pm standard deviation or %. BMI: body mass index; ALT: alanine aminotransferase; AST: aspartate aminotransferase; HDL: High Density Lipoprotein; LDL: Low Density Lipoprotein. Not NASH group defined as patients with NAFL (n=24) and borderline NASH (n=10) as opposed to patients with definite NASH.

^{*} P-values determined by comparing characteristics of definite NASH as compared to not NASH, using ANOVA and ANCOVA or Chi-square test when appropriate to compare categorical variables.