

# Detection of carcinogenic etheno-DNA adducts in children and adolescents with non-alcoholic steatohepatitis (NASH)

Ulrike Teufel<sup>1\*</sup>, Teresa Peccerella<sup>2\*</sup>, Guido Engelmann<sup>1</sup>, Thomas Bruckner<sup>3</sup>, Christa Flechtenmacher<sup>4</sup>, Gunda Millonig<sup>2</sup>, Felix Stickel<sup>5</sup>, Georg F. Hoffmann<sup>1</sup>, Peter Schirmacher<sup>4</sup>, Sebastian Mueller<sup>2</sup>, Helmut Bartsch<sup>6</sup>, Helmut K. Seitz<sup>2</sup>

<sup>1</sup>Department of Paediatrics, University of Heidelberg, Heidelberg, Germany; <sup>2</sup>Alcohol Research Centre, University of Heidelberg and Department of Medicine (Gastroenterology & Hepatology), Salem Medical Centre, Heidelberg, Germany; <sup>3</sup>Institute of Medical Biometry and Informatics, <sup>4</sup>Department of Pathology, University of Heidelberg, Heidelberg, Germany; <sup>5</sup>Hepatology Unit, Clinic Beau-Site Hirslanden, Bern, Switzerland; <sup>6</sup>Erstwhile: Division of Toxicology and Cancer Risk Factors, German Cancer Research Centre (DKFZ), Heidelberg, Germany

**Contributions:** (I) Conception and design: HK Seitz, H Bartsch, S Mueller; (II) Administrative support: GF Hoffmann, P Schirmacher; (III) Provision of study materials or patients: GF Hoffmann, G Engelmann, U Teufel; (IV) Collection and assembly of data: U Teufel, G Engelmann; (V) Data analysis and interpretation: T Peccerella, T Bruckner, C Flechtenmacher, G Millonig, F Stickel, HK Seitz; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

\*These authors contributed equally to this work.

**Correspondence to:** Helmut K. Seitz, MD, AGAF. Professor of Medicine, Gastroenterology and Alcohol Research, University of Heidelberg, Department of Medicine, Salem Medical Centre, Heidelberg, Germany. Email: [helmut\\_karl.seitz@urz.uni-heidelberg.de](mailto:helmut_karl.seitz@urz.uni-heidelberg.de).

**Background:** Carcinogenic exocyclic-DNA adducts like 1,N<sup>6</sup>-etheno-2'-deoxyadenosine (εdA) are formed through reactive intermediates of 4-hydroxynonenal (4-HNE) or other lipid peroxidation (LPO) products with the DNA bases A, C, methyl-C and G. High levels of hepatic etheno-DNA adducts have been detected in cancer prone liver diseases including alcoholic liver disease (ALD). In ALD εdA levels correlated significantly with cytochrome P-450 2E1 (CYP2E1) expression which is also induced in non-alcoholic steatohepatitis (NASH). We investigated the occurrence of εdA adducts in children with NASH as a DNA damage marker.

**Methods:** Liver biopsies from 21 children/adolescents with histologically proven NASH were analysed for hepatic fat content, inflammation, and fibrosis. εdA levels in DNA, CYP2E1-expression and protein bound 4-hydroxynonenal (HNE) were semi-quantitatively evaluated by immunohistochemistry.

**Results:** Among 21 NASH children, εdA levels in the liver were high in 3, moderate in 5, weak in 9 and not elevated in 4 patients. There was a positive correlation between CYP2E1 and protein-bound 4-HNE ( $r=0.60$ ;  $P=0.008$ ) and a trend for a positive relationship for CYP2E1 vs. staining intensity of εdA ( $r=0.45$ ;  $P=0.06$ ). Inflammatory activity and fibrosis correlated significantly ( $r=0.49$ ,  $P=0.023$ ).

**Conclusions:** Our results demonstrate for the first time the presence of elevated carcinogenic etheno-DNA lesions (εdA) in the majority (17/21) of liver biopsies from young NASH patients. Our data suggest that LPO-derived etheno-adducts are implicated in NASH. Whether these adducts may serve as predictive risk markers in NASH children to develop hepatocellular cancer later in life remains to be investigated.

**Keywords:** Cytochrome P-450 2E; etheno-DNA adducts; hepatocellular cancer; lipidperoxidation-induced DNA damage; non-alcoholic steatohepatitis in children (NASH in children)

Submitted Sep 29, 2015. Accepted for publication Nov 23, 2015.

doi: 10.3978/j.issn.2304-3881.2015.12.03

**View this article at:** <http://dx.doi.org/10.3978/j.issn.2304-3881.2015.12.03>

## Introduction

Nonalcoholic fatty liver disease (NAFLD) is frequently associated with obesity, insulin resistance and other features of the metabolic syndrome (1-3). While fatty liver *per se* is a benign condition, the occurrence of non-alcoholic steatohepatitis (NASH) is of major concern since it may progress to fibrosis and cirrhosis of the liver (4) and finally to hepatocellular carcinoma (HCC) (5). It is noteworthy that NASH not only occurs in adults but also in adolescents and children (6,7). When present at young age, it may be a risk factor for the occurrence of HCC later in life. Oxidative stress, excess generation of reactive oxygen species (ROS) and lipid peroxidation (LPO) products in the liver may be of particular importance for the disease onset (8).

In NASH, an induction of hepatic cytochrome P-450 2E1 (CYP2E1) has been described (9,10) which promotes oxidative stress, protein modifications, inflammation, and results in the generation of ROS causing LPO and subsequent cellular damage (11). Among the multiple LPO-products generated, 4-hydroxynonenal (4-HNE) a major product binds to proteins and to DNA *via* reactive intermediates forming carcinogenic etheno-DNA adducts such as 1,N<sup>6</sup>-etheno-2'-deoxyadenosine (εdA) (12-14). Both in cultured CYP2E1 overexpressing HepG2 cells and in liver biopsies from patients with alcoholic liver disease (ALD) a significant positive correlation between CYP2E1 induction and the generation of etheno-DNA adducts have been observed (15). Recently, we have detected εdA adducts in the livers of adult NASH patients (10). Since NASH also occurs in children, we investigated whether these carcinogenic DNA lesions also occur already at young age.

## Methods

### Patients

Twenty-one children/adolescents (13 males, 8 females) aged between 8 and 17 years were liver biopsied. They were recruited from the outpatient Clinic of the university Children Hospital Heidelberg which they visited because of elevated liver enzymes. A complete laboratory work up was performed and infectious, metabolic and autoimmune diseases were excluded. Clinical and metabolic data, including gamma-glutamyl transpeptidase (GGT), aspartate transaminase (AST), and alanine transaminase (ALT)-activities are given in *Table 1*. None of the patients had diabetes mellitus or consumed alcohol. Besides one patient (Asian) all were of Caucasian ethnicity.

**Table 1** Demographic, clinical and laboratory characteristics of the NASH patients studied

No.	Gender/age (years)	GGT NR <60 (U/l)	AST NR <50 (U/l)	ALT NR <50 (U/l)	BMI
1	f/8	73	89	185	15.05
2	m/11	66	37	53	19.96
3	f/14	79	90	201	21.45
4	f/9	16	35	68	17.67
5	m/16	N/A	N/A	N/A	N/A
6	m/17	48	50	126	29.08
7	m/15	28	58	161	34.60
8	m/17	48	117	297	32.37
9	f/12	40	78	132	30.23
10	m/12	136	257	430	26.59
11	f/11	68	61	163	30.21
12	m/13	19	78	191	29.67
13	m/14	19	30	51	24.18
14	m/10	118	55	96	24.09
15	m/12	140	243	537	26.26
16	m/11	33	44	78	26.26
17	m/14	32	26	47	23.73
18	f/11	N/A	89	89	26.20
19	f/17	106	166	90	32.20
20	f/11	N/A	66	68	22.48
21	m/15	37	85	147	25.34

All patients were Caucasians, except patient No. 2 who was of Asian origin. NASH, non-alcoholic steatohepatitis; GGT, gamma-glutamyl transpeptidase; AST, aspartate transaminase; ALT, alanine transaminase; NR, normal range.

Histologically normal liver sections from 3 healthy adult subjects were analyzed as controls for εdA background staining intensity. (These biopsies were originally taken to rule out hepatic tumours.) The study was approved in accordance with the declaration of Helsinki by the Ethical Committee of the University of Heidelberg, Germany (Ethic Approval Number: S-510-2009), and the parents of all subjects enrolled gave written informed consent for the participation in the present study.

All biopsies were assessed by histopathologists experienced in liver pathology (CF,PS) and diagnosed as NASH. Both investigators were blinded for immunohistopathology examinations related to the present study. Hepatic inflammation and fibrosis were assessed using the scoring system of Kleinert *et al.* (16). The

inflammation score identifies lobular inflammation with 0-3 (0, no inflammatory foci; 1, <2 foci/200×; 2, 2-4 foci/200×; and 3, >4 foci/200×).

#### ***Immunohistochemical detection of etheno-DNA adducts***

Staining was performed on liver tissue sections using the method developed in our laboratory (17-20). Paraffin-fixed slides were dipped in phosphate buffered saline (PBS) for 10 min and then placed in 0.3% H<sub>2</sub>O<sub>2</sub> in absolute methanol for 10 min to quench endogenous peroxidase. Slides were incubated with proteinase K (20 mg/mL) (Roche, Mannheim, Germany) in double distilled H<sub>2</sub>O at room temperature for 10 min to remove histone and non-histone proteins from DNA increasing antibody accessibility. After washing with PBS, slides were treated with 20 µg/mL RNase (Roche, Mannheim, Germany) (heated for 10 min at 80 °C to inactivate DNase) at 37 °C for 1 hour to prevent antibody binding to RNA adducts and then washed in PBS. To denature DNA, cells were treated with 4N HCl for 5 min at room temperature and subsequently rinsed in double distilled water and PBS. The pH was neutralized with 50 mM Trisbase buffer, pH 7.4, for 5 min at room temperature. Non-specific binding sites were blocked with 8% bovine serum albumin (BSA), 2% normal horse serum, 0.05% Tween 20 and 0.05% Triton X-100 for 20 min at 37 °C. Slides were incubated at 4 °C overnight with the primary monoclonal antibody EM-A-1 against εdA (provided by Drs. P. Lorenz and M. Rajewsky, University of Essen, Essen, Germany), at a dilution of 1:20 (20) and 2% normal horse serum to block nonspecific binding. After washing with PBS, the antibody detection was performed using the Vectastain Elite ABC kit (Vector Laboratories, Burlingame, CA, USA) according to the manufacturer's protocol (incubation with secondary antibody: horse anti-mouse IgG (H + L) 1:400 for 40 min at room temperature). Diaminobenzidine (DAB) was used as a chromogen to visualize the reaction. The reaction was stopped after 5 min with H<sub>2</sub>O. Slides were counterstained with 4',6'-diamidino-2-phenylindole and mounted with Roti-Histokitt II (Carl Roth, Karlsruhe, Germany). All slides were subjected to the same standard conditions. Negative controls were performed by omitting the primary antibody.

#### ***Immunohistochemical staining of CYP2E1 and protein bound 4-HNE***

Paraffin-embedded liver biopsy samples were cut into 6 mm

sections and placed on 3-aminopropyl-triethoxysilan-coated glass slides. Sections were treated with 0.5% H<sub>2</sub>O<sub>2</sub> in absolute methanol for 10 min to quench endogenous peroxidase activity. Thereafter, sections were incubated at room temperature for 2 hours with the primary antibody (rabbit anti-human CYP2E1, 1:400, Chemicon, Hofheim, Germany) or with rabbit anti-4-HNE (1:250, Alexis, Lörrach, Germany) and 5% normal goat serum to block nonspecific binding. Vectastain Elite ABC (Vector Laboratories, Burlingame, CA, USA) was used for detection according to the manufacturer's protocol. Staining was developed by incubating the sections for 5 min in DAB. Sections were counterstained with hematoxylin and mounted with Roti-Histokitt II (Carl Roth, Karlsruhe, Germany). Negative controls were performed by omitting the primary antibody.

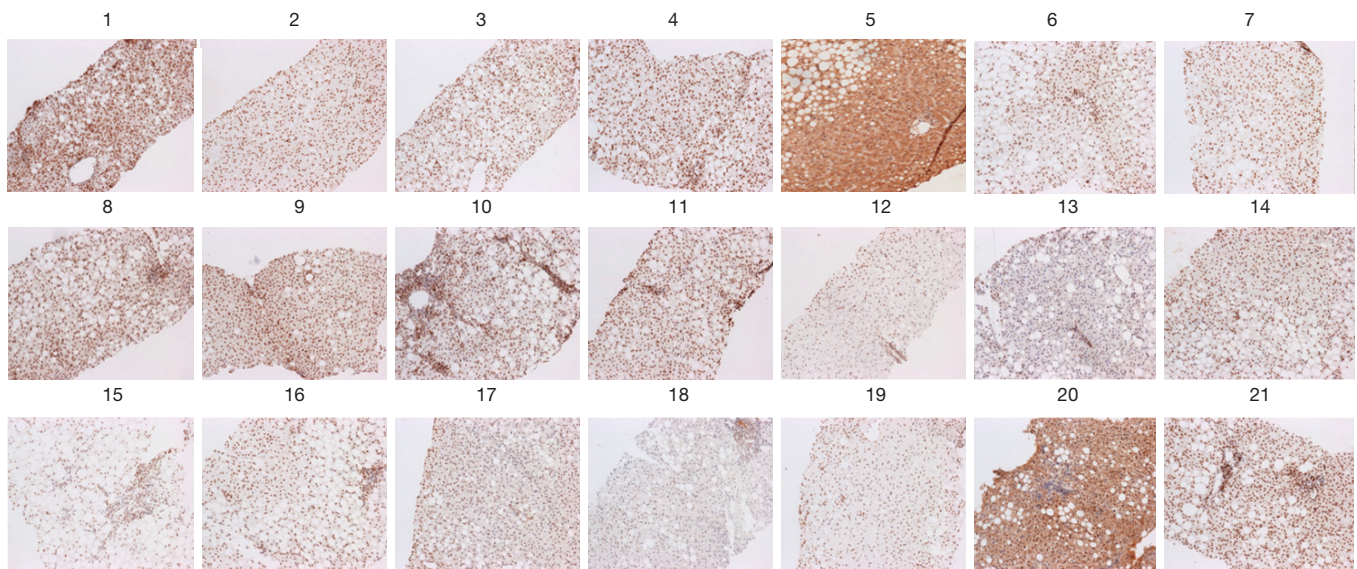
#### ***Immunohistochemical detection of alkyladenine DNA glycosylase (AAG) expression***

AAG expression was determined in 6 biopsies. Paraffin-embedded tissue was dewaxed in xylene, rehydrated in a decreasing ethanol series and finally washed in PBS. Antigen retrieval was performed by heating the slides in citrate buffer (10 mM, pH 6 with 0.05% Tween-20) for 10 minutes. After cooling down to room temperature, the slides were treated with 0.3% H<sub>2</sub>O<sub>2</sub> in absolute methanol for 20 minutes to quench endogenous peroxidase activity. Thereafter, sections were incubated overnight at 4 °C with the primary antibody (mouse antihuman APNG, 1:50, Santa Cruz, CA, USA) and 2% normal horse serum to block nonspecific binding. VECTASTAIN Elite ABC kit was used for detection according to the manufacturer's protocol (secondary antibody: anti-mouse species 1:200 and 2% normal horse serum in PBS). Staining was developed by incubating the sections for 5 minutes with DAB. After dehydrating the sections in an increasing alcohol series, they were counterstained with hematoxylin and mounted in Roti-Histokitt II (Carl Roth GmbH, Karlsruhe, Germany) mounting medium. Negative controls were performed by omitting the primary antibody.

#### ***Imaging and semi-quantitative analyses of etheno-DNA adducts, CYP2E1 expression and protein-bound 4-HNE***

After immunohistochemical staining for εdA, CYP2E1 and 4-HNE, the blinded slides were independently scored





**Figure 1** Immunohistochemistry of  $\epsilon$ dA in 21 children/adolescents with NASH (100 $\times$ ). Immunohistochemical staining of an exocyclic etheno-DNA adduct (etheno-deoxyadenosine,  $\epsilon$ dA) in hepatic nuclei of liver biopsies from 21 children (aged 8-17 years) using a highly specific monoclonal antibody (22). The staining intensity varied from strong (#1,5,20), moderate (#8,9,10,11,21), mild (#2,3,4,6,7,13,14,16,17) and non-detectable background levels only in (#12,15,18,19).  $\epsilon$ dA, 1,N<sup>6</sup>-etheno-2'-deoxyadenosine; NASH, non-alcoholic steatohepatitis.

by a second investigator. Representative pictures were taken at a magnification of 100 $\times$ /400 $\times$  with a SPOT FLEX system (Model 15.2 64 Mp, shifting pixel, DIAGNOSTIC Instruments, Inc., Sterling Heights, Michigan, USA, SPOT VERSION 4.6.4.69) and analyzed using Image J software (U. S. National Institutes of Health, Bethesda, Maryland, USA, <http://imagej.nih.gov/ij/>).

Staining intensity for CYP2E1 and 4-HNE was assessed according to the scale devised by Tsutsumi *et al.* (21), whereby 3+, 2+, 1+ and 0 denote intense, moderate, slight and no specific immunostaining, respectively. Staining intensity of  $\epsilon$ dA was estimated and recorded as 1 to 4. These 4 graded levels were used as there is always a very low background staining for  $\epsilon$ dA which was assigned as grade 1. In addition; the frequency of nuclei positively stained for  $\epsilon$ dA was calculated as % of stained cell nuclei over a total number of cells counted.

### Statistics

The empirical distribution of continuous parameter was reported with mean, minimum, and maximum, with absolute and relative frequency by categorical variables.

Spearman rank correlation was calculated to examine possible correlation between characteristics. A value of  $\alpha=0.05$  was used as level of significance.

### Results

Figure 1 shows the various grades of immunohistochemical staining for  $\epsilon$ dA in cell nuclei of hepatocytes in liver biopsies from NASH children. Among 21 biopsies,  $\epsilon$ dA levels in the liver were high in 3 (14.3%), moderate in 5 (23.8%), weak in 9 (42.9%), and not elevated in 4 (19%) individuals showing only a low background level. As biopsies from healthy children cannot be taken for ethical reasons, we used liver sections from healthy adult subjects (biopsy was taken to rule out hepatic tumours) as the only available controls for  $\epsilon$ dA background staining. A very low background level of etheno adducts (range, 0-27 adducts per 10<sup>9</sup> parent bases) has been detected in liver DNA from asymptomatic humans and untreated rodents (22) (data not shown).

Table 2 summarizes disease grade (inflammatory activity) and stage (degree of fibrosis) as classified by histopathological evaluation (fat, inflammation, fibrosis).

**Table 2** Etheno-DNA adducts, CYP2E1, 4-HNE and histological characteristics of the NASH patients studied

No.	$\epsilon$ dA (% pos. nuclei)	$\epsilon$ dA (ISc)	CYP2E1 (ISc)	4-HNE (ISc)	Grade	Stage
1	100	4	3	3	1	3
2	100	2	2	3	1	2
3	99.4	2	2	3	2	3
4	99.7	2	2	2	1	3
5	99.8	4	3	3	1	0
6	99.4	2	2	2	1	2
7	97.3	2	3	3	1	3
8	98.9	3	3	3	1	2
9	99.5	3	2	3	1	0
10	97.1	3	2	2	2	4
11	99.6	3	1	1	2	4
12	68.0	1	1	1	1	2
13	79.1	2	3	3	1	0
14	98.9	2	3	2	1	2
15	94.5	1	2	2	1	3
16	99.6	2	2	2	1	3
17	54.7	2	Nd	Nd	1	2
18	34.5	1	Nd	Nd	1	0
19	65.2	1	Nd	Nd	1	2
20	97.6	4	3	3	1	4
21	98.9	3	3	2	1	3

Grade of inflammation: 0, no inflammatory foci; 1, <2 foci/200 $\times$ ; 2, 2-4 foci/200 $\times$ ; 3, >4 foci/200 $\times$ . Stage of fibrosis: 0, none; 1, perisinusoidal or periportal fibrosis; 2, persinusoidal and periportal fibrosis; 3, bridging fibrosis; 4, cirrhosis.  $\epsilon$ dA, 1,N<sup>6</sup>-etheno-2'-deoxyadenosine; ISc, intensity score; CYP2E1, cytochrome P-450 2E1; 4-HNE, 4-hydroxynonenal; Nd, not determined.

Staining intensity of CYP2E1, protein bound 4-HNE and  $\epsilon$ dA in nuclear DNA is shown.

A significant correlation between CYP2E1 and protein bound 4-HNE ( $r=0.60$ ;  $P=0.008$ ) was found. A trend for a positive relationship between the staining intensity of  $\epsilon$ dA and CYP2E1 ( $r=0.45$ ;  $P=0.06$ ) was apparent but not seen for  $\epsilon$ dA and protein bound 4-HNE ( $r=0.39$ ;  $P=0.11$ ). Neither  $\epsilon$ dA, protein bound 4-HNE nor CYP2E1 correlated significantly with serum enzyme activities.

A comparison of  $\epsilon$ dA, CYP2E1, and 4-HNE staining in representative biopsies of 6 children is depicted in *Figure 2*.

In general, strong staining intensity paralleled each other for CYP2E1 and protein-bound 4-HNE, but was less apparent for CYP2E1 *vs.*  $\epsilon$ dA, as reflected by the Spearman rank correlation ( $r=0.45$ ;  $P=0.061$ ).

*Figure 3* shows AAG expression *vs.*  $\epsilon$ dA in 6 biopsies. In 5 out of 6 biopsies the level of etheno adducts seems reversely related to AAG expression.

Inflammatory activity (grading) and degree of fibrosis (staging) correlated significantly in these biopsies ( $r=0.49$ ;  $P=0.02$ ). However, a significant correlation between histology,  $\epsilon$ dA, CYP2E1 or 4-HNE was not observed.

Significant correlations also occurred between serum GGT activity and the activities of AST ( $r=0.67$ ;  $P=0.002$ ) as well as of ALT ( $r=0.49$ ;  $P=0.037$ ) and between the serum activities of both transaminases ( $r=0.78$ ;  $P=0.0001$ ).

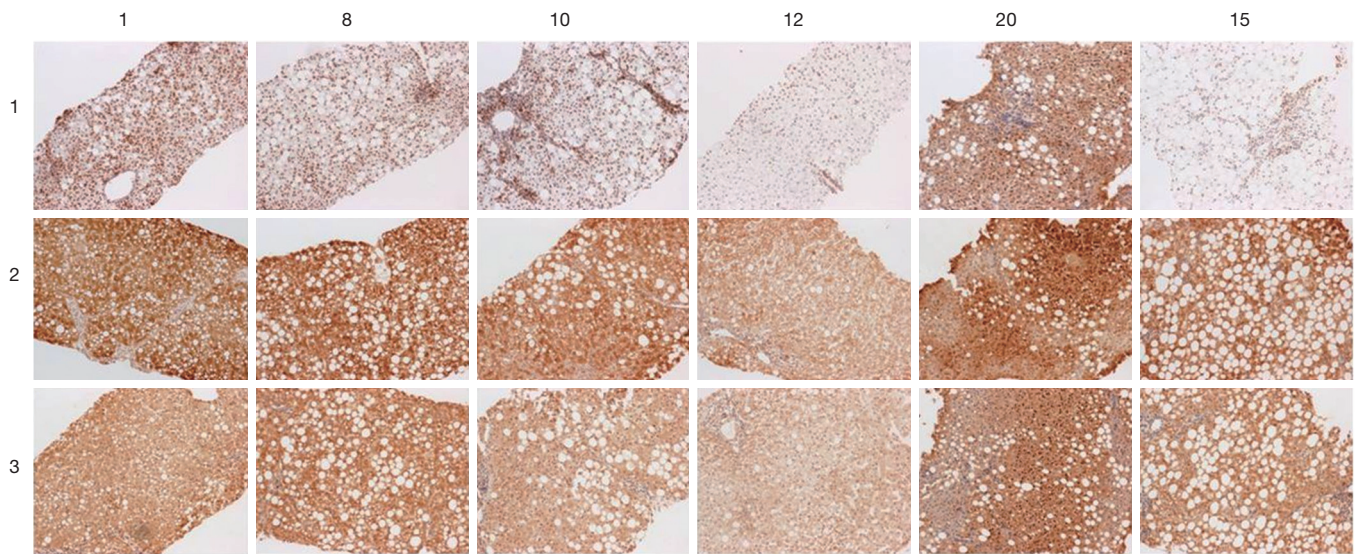
## Discussion

Our results presented here demonstrate for the first time the occurrence of carcinogenic, LPO-derived DNA adducts in the majority of liver biopsies of young NASH patients.  $\epsilon$ dA was analyzed, using a highly specific monoclonal antibody (17) for immunohistochemical detection. Etheno-DNA adducts are highly mutagenic lesions producing specific point mutations and other types of genomic lesions in all organisms tested so far (23). Exocyclic DNA adducts are derived from reaction of DNA bases A, C, 5methyl-C, and G with LPO-products that are formed from inflammation-driven oxidative stress reactions (24-26). One of the major LPO-product, 4-HNE induces DNA damage through its reactive intermediates (27), one of which is 2,3-epoxy-4-hydroxynonenal (28). Complex LPO reactions *in vivo* lead to both substituted (with fatty acid chains attached) such as heptanone-etheno adducts (27,29,30) and to the unsubstituted adducts  $\epsilon$ dA, 3,N<sup>4</sup>-etheno-2'-deoxycytosine ( $\epsilon$ dC), as well as two isomeric etheno-2'-deoxyguanosine adducts (26,31,32). A hitherto unknown LPO-derived DNA adduct formed *in vivo*, 3,N<sup>4</sup>-etheno-5-methyl-2'-deoxycytidine was recently detected in human liver that may perturb genome methylation (33).

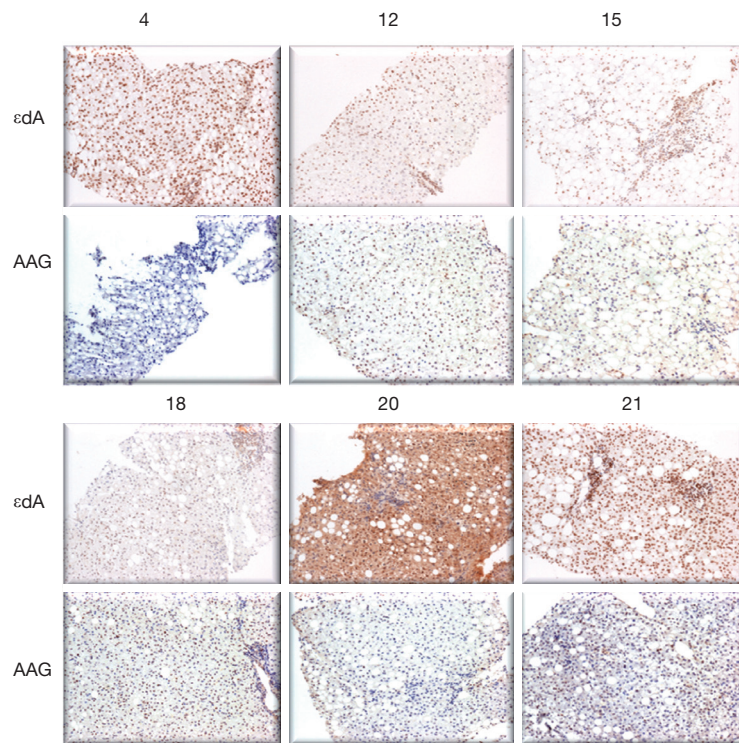
$\epsilon$ dA, a marker lesion for exocyclic adducts is formed by reaction of LPO-products with DNA bases in human (31) and rodent tissues (14). After DNA repair in internal organs, etheno adducts are eliminated and excreted in urine (20).

Strong support for the pathological relevance of etheno adducts is provided by previous studies which found up to 1-2 orders of magnitude elevated adduct levels in





**Figure 2** Immunohistochemistry of  $\epsilon$ dA, CYP2E1, and 4-HNE in 6 children/adolescents with NASH (100 $\times$ ). Immunohistochemical detection of  $\epsilon$ dA (upper panel, 1), CYP2E1 (mid panel, 2) and 4-HNE (lower panel, 3) in liver biopsies of six NASH children. The staining intensity between CYP2E1 expression and protein bound 4-HNE correlated significantly ( $r=0.60$ ;  $P=0.008$ ); correlation between  $\epsilon$ dA and CYP2E1 in the 21 biopsies showed a trend ( $r=0.45$ ;  $P=0.06$ ).  $\epsilon$ dA, 1, $N^6$ -etheno-2'-deoxyadenosine; CYP2E1, cytochrome P-450 2E1; 4-HNE, 4-hydroxynonenal; NASH, non-alcoholic steatohepatitis.



**Figure 3** Immunohistochemical staining of  $\epsilon$ dA and alkyladenine DNA glycosylase (AAG) in liver biopsies of six NASH children (100 $\times$ ). While in 1 biopsy (#4) high  $\epsilon$ dA staining is associated with high AAG staining, in all other biopsies  $\epsilon$ dA staining and AAG staining seem to be inversely related to each other.  $\epsilon$ dA, 1, $N^6$ -etheno-2'-deoxyadenosine.

human liver of patients with cancer prone inflammatory diseases (31,34).

In liver biopsies from ALD patients, the induction of CYP2E1 by chronic ethanol consumption resulted in excess generation of LPO products measured as protein bound 4-HNE, and etheno-DNA adducts (15). A prominent role of CYP2E1 in hepatocarcinogenesis has been further supported in CYP2E1 overexpressing human cells, and in a rodent NASH model using Zucker rats with and without ethanol ingestion (15,35,36). Thus, in both human and experimental studies, a significant positive correlation between CYP2E1 expression and  $\epsilon$ DA levels could be consistently demonstrated; moreover, the inhibition of CYP2E1 by clomethiazole resulted in a significant reduction of  $\epsilon$ DA levels in rat liver (15). CYP2E1 has also been found to be responsible for ethanol-induced oxidative DNA damage in the rodent liver (37) and for the occurrence of hepatic adenomas in chemically induced hepatocarcinogenesis following chronic alcohol ingestion in rats (35). Hepatic CYP2E1 was also induced in adult NASH patients (9,10), regardless whether they have developed diabetes or not (38). It has been shown in ob/ob mice, an experimental model for NASH, that CYP2E1 promotes liver injury (39).

It was demonstrated by Fujita *et al.* (40) that oxidative stress also plays an important role in disease pathogenesis of NASH. 8-oxodG, a DNA base-modified product generated by hydroxyl radicals, was significantly higher in NASH liver biopsy samples compared to subjects with simple steatosis.

Although in ALD patients CYP2E1 expression and activity did significantly correlate with  $\epsilon$ DA (15), this was not found in hepatic biopsies from NASH children, where only a trend with borderline significance ( $r=0.45$ ;  $P=0.06$ ;  $N=21$ ) was found. Several explanations for this weak relationship may exist:

- (I) Our sample size is too small;
- (II) Only  $\epsilon$ DA was determined while correlations may exist with other types of etheno adducts that have been found in human tissues (24,31);
- (III) Most of ROS formation was inflammation driven rather than generated through CYP2E1, since it has been reported that the high prevalence of oxidative stress in children with NAFLD is associated with increased severity of steatohepatitis (41);
- (IV) The large inter-individual variations of hepatic  $\epsilon$ DA adduct levels could be caused by varying rates of their formation (affected by detoxifying enzymes,

antioxidants and levels of inflammatory mediators) and by their removal rates by DNA repair enzymes. These modifying factors have not been measured.

To support the latter assumption, a study on the expression of a DNA repair enzyme, AAG in liver biopsies from NASH children was attempted (Figure 3). Unsubstituted etheno-DNA adducts are eliminated mainly by base-excision repair. AAG and the mismatch-specific thymidine-DNA glycosylase remove  $\epsilon$ DA and  $\epsilon$ DC respectively (42-44).

Indeed, low AAG staining seemed to parallel a high etheno-DNA adduct formation, suggesting that low (or impaired) base excision repair activity may lead to increased DNA adduct generation. Confirmation of this hypothesis with a larger sample size is mandatory but of great interest in the context of a reduced DNA damage repair in NASH (44). Indeed, in steatotic livers of obese subjects with NASH, a decrease in nucleotide excision repair (NER) was found for  $M_1$ dG, another major LPO-derived DNA adduct, formed by reaction of malondialdehyde with deoxyguanine and by other pathways. Importantly the observed reduction in NER capacity upon hepatic inflammation in NASH patients was associated with and may be a consequence of reduced damage recognition (45). For an HNE-derived substituted etheno-dG adduct (termed HNE-dG) possibly also generated in NASH liver, NER was shown to be the major repair pathway in human cells (46). Thus, studies whether the hepatic DNA repair enzyme capacities of AAG and NER are reduced, leading to increased DNA damage are highly warranted.

Finally, it is noteworthy that DNA damage is particular detrimental during foetal development. Results from a cohort study of white blood cells (WBC) from mother-newborn child pairs lend support that NASH related oxidative stress is a self perpetuating process, and LPO-derived DNA damage could already be triggered in utero. As shown in 77 WBC-DNA samples from mother-newborn child pairs, highly variable levels of etheno-DNA adducts were detected (47). These results confirm that this DNA damage arises *in vivo* from LPO-derived reactive aldehydes such as 4-HNE and indicate that a similar signature of DNA damage is found in foetus and mother.

In conclusion, in children and young adults with NASH, one of the mutagenic and carcinogenic etheno-DNA adducts  $\epsilon$ DA was unequivocally detected in 17 out of 21 liver biopsies. Semi-quantitative analysis of  $\epsilon$ DA by



immunohistochemistry revealed large inter-individual variations. CYP2E1 expression and  $\epsilon$ DA levels showed only a trend for a positive relationship, possibly because  $\epsilon$ DA may additionally be formed through ROS generation due to inflammation. Whether hepatic etheno adducts may serve as predictive risk markers in NASH children to develop HCC later in life remains to be investigated.

## Acknowledgements

This study was supported by grants of the Dietmar Hopp and Manfred Lautenschläger Foundations to H. K. Seitz. We thank Ms. Heike Grönebaum for typing this manuscript.

## Footnote

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

*Ethical statement:* The study was approved in accordance with the declaration of Helsinki by the Ethical Committee of the University of Heidelberg, Germany (Ethic Approval Number: S-510-2009), and the parents of all subjects enrolled gave written informed consent for the participation in the present study.

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**Cite this article as:** Teufel U, Peccerella T, Engelmann G, Bruckner T, Flechtenmacher C, Millionig G, Stickel F, Hoffmann GF, Schirmacher P, Mueller S, Bartsch H, Seitz HK. Detection of carcinogenic etheno-DNA adducts in children and adolescents with non-alcoholic steatohepatitis (NASH). *HepatoBiliary Surg Nutr* 2015;4(6):426-435. doi: 10.3978/j.issn.2304-3881.2015.12.03