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Metabolism of Non-Enzymatically Derived Oxysterols: Clues from sterol metabolic disorders^{⋄,⋄,⋄}

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ABSTRACT

Cholestane- 3β , 5α , 6β -triol (3β , 5α , 6β -triol) is formed from cholestan-5,6-epoxide (5,6-EC) in a reaction catalysed by cholesterol epoxide hydrolase, following formation of 5,6-EC through free radical oxidation of cholesterol. 7-Oxocholesterol (7-OC) and 7β -hydroxycholesterol (7β -HC) can also be formed by free radical oxidation of cholesterol. Here we investigate how 3β , 5α , 6β -triol, 7-OC and 7β -HC are metabolised to bile acids. We show, by monitoring oxysterol metabolites in plasma samples rich in 3β , 5α , 6β -triol, 7-OC and 7β -HC, that these three oxysterols fall into novel branches of the acidic pathway of bile acid biosynthesis becoming (25R)26-hydroxylated then carboxylated, 24-hydroxylated and side-chain shortened to give the final products 3β , 5α , 6β -trihydroxycholanoic, 3β -hydroxy-7-oxochol-5-enoic and 3β , 7β -dihydroxychol-5-enoic acids, respectively. The intermediates in these pathways may be causative of some phenotypical features of, and/or have diagnostic value for, the lysosomal storage diseases, Niemann Pick types C and B and lysosomal acid lipase deficiency. Free radical derived oxysterols are metabolised in human to unusual bile acids via novel branches of the acidic pathway, intermediates in these pathways are observed in plasma.

1. Introduction

The oxysterols 7-oxocholesterol (7-OC, for systematic names see Supplemental Table S1), 7β -hydroxycholesterol (7β -HC), 5α ,6-epoxycholesterol (5α ,6-EC), 5β ,6-epoxycholesterol (5β ,6-EC) and cholestane- 3β , 5α , 6β -triol (3β , 5α , 6β -triol) have for many years been thought of as autoxidation artefacts formed *ex vivo* from cholesterol during sample handling [1-3]. This view is now changing with the realisation that (i) 7-OC can be formed enzymatically from 7-dehydrocholesterol (7-DHC) [4], (ii) 7β -HC and 7-OC can be interconverted in enzyme catalysed reactions [5-9], while (iii) the 5α ,6-EC adduct dendrogenin A (DDA) is present in tissue [10], (iv) 5,6-EC is hydrolysed by the enzyme cholesterol epoxide hydrolase (ChEH) to 3β , 5α , 6β -triol which itself is

oxidised by hydroxysteroid dehydrogenase (HSD) 11B2 to $3\beta,5\alpha$ -dihydroxycholestan-6-one ($3\beta,5\alpha$ -diHC-6O, also called 6-oxocholestan- $3\beta,5\alpha$ -diol) found at elevated levels in breast cancer tissue [11], and (v) high levels of 7-OC and $3\beta,5\alpha,6\beta$ -triol are characteristic of the lysosomal storage diseases Niemann-Pick types A, B, C1 and C2 (NPA, NPB, NPC) and lysosomal acid lipase deficiency (LALD) [12-18]. In these inborn errors of metabolism, characterised by the accumulations of multiple lipid species including the build-up of lysosomal cholesterol, it is likely that 7-OC and 5,6-EC, the precursor of $3\beta,5\alpha,6\beta$ -triol, are formed via free radical reactions (Fig. 1). Importantly, 7-OC, 7 β -HC, $5\alpha,6$ -EC and $3\beta,5\alpha,6\beta$ -triol are precursors of biologically active molecules, including 25-hydroxy-7-oxocholesterol (25H,7O-C) and 26-hydroxy-7-oxocholesterol (26H,7O-C) which are ligands to the Hedgehog

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^{**} Swansea Innovations Ltd have licensed derivatisation technology described in this paper to Avanti Polar Lipids Inc. and Cayman Chemical Company.

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Abbreviations

ACOT acyl-CoA thioesterases ACOX2 acyl-coenzyme A oxidase 2 alpha-methylacyl-CoA-racemase AMACR **BACS** bile acyl CoA-synthetase ChEH cholesterol epoxide hydrolase cholesterol oxidase ChOx

cytochrome P450 CYP DBP D bifunctional protein DDA dendrogenin A Ηh Hedgehog

HSD hydroxysteroid dehydrogenase

GlcNAc N-acetylglucosamine GP Girard P reagent

LALD lysosomal acid lipase deficiency

LC-MS liquid chromatography mass spectrometry

multiple reaction monitoring MRM

 MS^n mass spectrometry with multistage fragmentation

NPA Niemann-Pick disease type A NPB Niemann-Pick disease type B NPC Niemann-Pick disease type C RICs reconstructed ion chromatograms SLOS Smith-Lemli-Opitz syndrome

Smo Smoothened

SPCx sterol carrier protein x

3β-НСА 3β-hydroxycholest-5-en-(25R)26-oic acid

3βH,7O-CA 3β-hydroxy-7-oxocholest-5-en-(25R)26-oic acid

 3β H,7O- Δ ⁵-BA 3β -hydroxy-7-oxochol-5-enoic acid 3β , 5α -diHC-6O 3β , 5α -dihydroxycholestan-6-one 3β,5α,6β-triHBA 3β,5α,6β-trihydroxycholanoic acid 3β,5α,6β-triHCa 3β,5α,6β-trihydroxycholestan-(25R)26-oic acid 3β,5α,6β-triol cholestane-3β,5α,6β-triol 3β , 5α , 6β , 24-tetra HCa 3β , 5α , 6β , 24-tetra hydroxycholestan-26-oic

 $3\beta,5\alpha,6\beta,26$ -tetrol cholestane- $3\beta,5\alpha,6\beta,(25R)$ 26-tetrol 3β,7α-diHCA 3β,7α-dihydroxycholest-5-en-(25R)26-oic acid 3β,7β-diHCA 3β,7β-dihydroxycholest-5-en-(25R)26-oic acid 3β , 7β -diH- Δ ⁵-BA 3β , 7β -dihydroxychol-5-enoic acid

3β,7β,24-triHCA 3β,7β,24-trihydroxycholest-5-en-26-oic acid

3β,24-diH,7O-CA 3β,24-dihydroxy-7-oxocholest-5-en-26-oic acid

 $5\alpha,6$ -EC $5\alpha,6$ -epoxycholesterol 5β,6-EC 5β,6-epoxycholesterol 7-DHC 7-dehydrocholesterol 7-OC 7-oxocholesterol

7α-HCO 7α-hydroxycholest-4-en-3-one

7α-hydroxycholesterol 7α-HC

7αH,3O-CA 7α-hydroxy-3-oxocholest-4-en-(25R)26-oic acid

7β-hydroxycholesterol 7β-НС 7β,26-diHC 7β,26-dihydroxycholesterol 24S-HC 24S-hydroxycholesterol 25-HC 25-hydroxycholesterol 25H,7O-C 25-hydroxy-7-oxocholesterol 26-HC (25R)26-hydroxycholesterol 26H,7O-C 26-hydroxy-7-oxocholesterol.

7α-НС LOH + ¹O₂ HSD3B7 HSD11B2 7α-HCO HSD11B2

Fig. 1. Free radical formation of 7α-HC, 7β-HC, 7-OC and 5,6-EC [34]. The reactions shown in blue boxes and by blue arrows are enzyme catalysed. CYP, cytochrome P450; HSD, hydroxysteroid dehydrogenase; ChEH, cholesterol epoxide hydrolase; LOO, lipidperoxy radical; LO, lipidalkoxy radical; LOH, lipidhydroxide; LOOH lipidhydroperoxide; RH, hydrogen donor.

(Hh) pathway oncoprotein Smoothened (Smo) [19]; 7β,26-dihydroxycholesterol (7β,26-diHC), a ligand to the nuclear receptor RORγt [20]; DDA a tumour suppressor [21] and both DDA and dendrogenin B have potent proliferative effects in neural stem cells [10]; and the oncometabolite and glucocorticoid receptor ligand 3β,5α-diHC-6O [11]. Accepting that the 5,6-epoxides and 7-oxidised sterols are formed in vivo, interest now shifts to investigating how they are metabolised. With respect to 7-OC, we have recently reported that in patients with the recessive congenital disorder Smith-Lemli-Opitz syndrome (SLOS) 7-OC can be converted to 7β-hydroxy bile acids and ultimately conjugated with N-acetylglucosamine (GlcNAc) [22-25]. A similar pathway may also be operative in patients with NPC disease where 7-OC is also elevated and the same GlcNAc conjugated bile acids have been identified [26]. Patients with NPC, NPB or NPA show elevated 3β,5α,6β-triol in plasma, presumably derived from free radical oxidation of cholesterol to 5,6-EC followed by hydrolysis to the triol by ChEH (Fig. 1) [27]. Clayton and colleagues and Ory and colleagues have now both identified $3\beta,5\alpha,6\beta$ -trihydroxycholanoic acid $(3\beta,5\alpha,6\beta$ -triHBA) as the bile acid product of $3\beta,5\alpha,6\beta$ -triol metabolism [28,29]. An alternative route for metabolism of $3\beta,5\alpha,6\beta$ -triol is oxidation to the oncometabolite 3β ,5α-diHC-6O [11].

Although the end products of 7-OC and 5,6-EC/3β,5α,6β-triol

metabolism have been defined, the *in vivo* biochemical pathways generating these products have yet to be fully elucidated. To study further the metabolism of 7-OC and 5,6-EC/ $3\beta,5\alpha,6\beta$ -triol we took advantage of plasma samples from patients where levels of these substrates are particularly high. We find that 7-OC, 7β -HC and $3\beta,5\alpha,6\beta$ -triol fall into new branches of the acidic pathway of bile acid biosynthesis and in patient plasma where the concentration of these metabolites is high, almost all the necessary intermediates to bile acids are observed. Buildup of these intermediates may be responsible for some of the clinical features of diseases where free radical oxidation of cholesterol is prevalent and their measurement may have diagnostic value.

2. Materials and methods

2.1. Materials

Oxysterol standards were from Avanti Polar Lipids Inc (Alabaster, Al, USA), 3β , 5α , 6β -triHBA was prepared as in Ref. [29], other bile acid standards were kindly donated by Dr Jan Sjövall of Karolinska Institute, Stockholm, or as detailed in Griffiths et al. [30]. Materials for liquid chromatography – mass spectrometry (LC-MS) analysis were as in Refs. [31,32].

Fig. 2. Derivatisation of (i) 3β -hydroxy-5-ene, and (ii) 3β , 5α , 6β -triol, containing sterols via oxidation with bacterial cholesterol oxidase (ChOx) and reaction with $[^2H_5]$ GP, and of (iii) 7-oxo-sterols by reaction with $[^2H_0]$ GP in the absence of cholesterol oxidase. The reactions are illustrated for 7β -HC, 3β , 5α , 6β -triol and 7-OC, respectively.

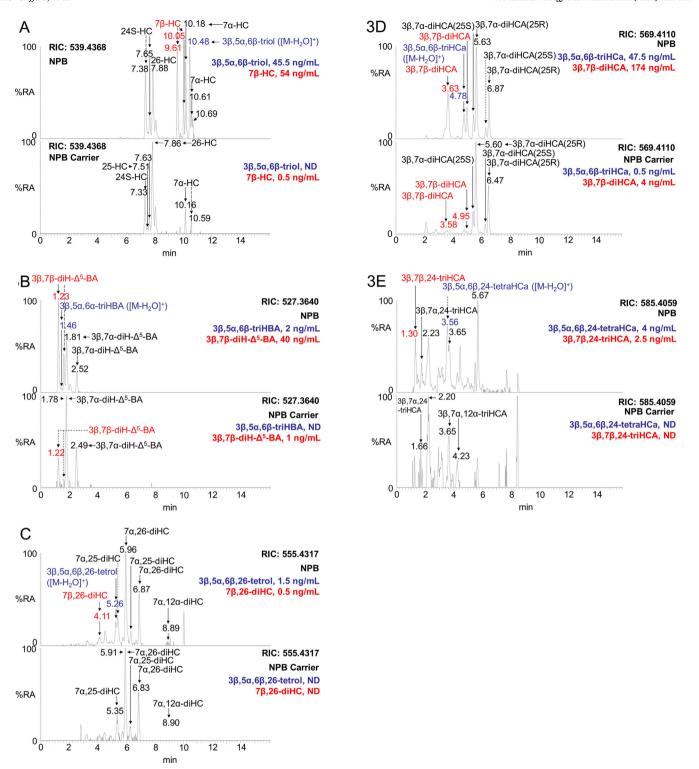


Fig. 3. LC-MS reconstructed ion chromatograms (RICs) of oxidised/ $[^2H_5]$ GP derivatised oxysterols and bile acids in plasma from an NPB patient and NPB carrier. RICs for (A) m/z 539.4368 corresponding to $[M]^+$ ions of hydroxycholesterols (HC) and $[M-H_2O]^+$ ions of 3β ,5 α ,6 β -triol, (B) m/z 527.3640 corresponding to $[M]^+$ ions of dihydroxychol-5-enoic acids (diH- Δ^5 -BA) and $[M-H_2O]^+$ ions of 3β ,5 α ,6 β -triHBA, (C) m/z 555.4317 corresponding to $[M]^+$ ions of dihydroxycholesterols (diHC) and $[M-H_2O]^+$ ions of 3β ,5 α ,6 β ,26-tetrol, (D) m/z 569.4110 corresponding to $[M]^+$ ions of dihydroxycholest-5-en-26-oic acids (diHCA) and $[M-H_2O]^+$ ions of 3β ,5 α ,6 β -triHCa, (E) m/z 585.4059 corresponding to trihydroxycholest-5-en-26-oic acids (triHCA) and $[M-H_2O]^+$ ions of 3β ,5 α ,6 β ,24-tetraHCa. The upper panels show RICs of NPB patient and lower panels NPB carrier plasma. Oxysterols with a the 3β ,5 α ,6 β -triol function are labelled in blue, those with a 3β ,7 β -dihydroxy function in red. Concentrations are given in the righthand corner of chromatograms. GP derivatised oxysterols give syn and anti conformers, which may or may not be chromatographically separated. For quantification of 3β ,5 α ,6 β ,26-tetrol and 3β ,5 α ,6 β ,24-tetraHCa extended chromatographic gradients were exploited as described in Griffiths et al. [30].

2.2. Patient samples

All participants or their parents/guardians provided informed consent and the study was performed with institutional review board

approval (REC08/H1010/63) and adhered to the principles of the Declaration of Helsinki.

Fig. 4. Formation and metabolism of 5,6-EC, 7β-HC and 7-OC. The enzymes catalysing the reactions are cholesterol epoxide hydrolase (ChEH), hydroxysteroid dehydrogenase 11B1 (HSD11B1), HSD11B2, cytochrome P450 27A1 (CYP27A1), bile acyl CoA-synthetase (BACS), alpha-methylacyl-CoA-racemase (AMACR), acyl-coenzyme A oxidase 2 (ACOX2), D bifunctional protein (DBP) and sterol carrier protein x (SPCx). For simplicity, the individual reactions catalysed by BACS, AMACR, ACOX2, DBP and SPCx are omitted and carboxylic acids rather than CoA-thioesters are shown. The peroxisomal enzyme bile acid CoA thioesterase (ACOT) will hydrolyse the CoA-thioesters to free acids [38,39].

2.3. LC-MS methods

The LC-MS methods have been described in detail elsewhere [30,31]. In brief, a charge-tagging method was adopted [31,32], where sterols, including oxysterols and bile acids, with a 3 β -hydroxy group were oxidised with bacterial cholesterol oxidase (ChOx) to 3-oxo analogues and derivatised with [2H_5]-labelled Girard P (GP) reagent (Fig. 2), then analysed by LC-MS at high mass-resolution (120,000 at m/z 400, full-width at half-maximum height definition) with parallel multistage fragmentation (MS^n). Sterols with a natural oxo group were derivatised with [2H_0]GP reagent in the absence of cholesterol oxidase and analysed together with the [2H_5]GP-derivatised sterols in a single LC-MS(MS^n) run. Quantification was by the isotope dilution method.

3. Results

3.1. GP derivatisation of 3β-hydroxy-5-enesterols, 3β,5α,6β-trihydroxysterols and 7-oxosterols

Cholesterol oxidase from *Streptomyces* sp converts sterols with a 3 β -hydroxy-5-ene function to analogues with a 3-oxo-4-ene structure, which are substrates for derivatisation with [2H_5]GP (Fig. 2). The derivatisation products give [M] $^+$ ions in LC-MS analysis. Upon oxidation/derivatisation of 3 β ,5 α ,6 β -trihydroxysterols the 5 α -hydroxy group is labile with the result that these oxysterols becomes converted to [2H_5]GP-derivatives of a 6 β -hydroxy-3-oxo-4-ene structure, i.e. the [M - H $_2$ O] $^+$ ion of the oxidised/derivatised triol [33]. 7-Oxosterols do not require cholesterol oxidase treatment prior to [2H_0]GP derivatisation and are hence evident as [M] $^+$ ions in preparations in the absence of enzyme (Fig. 2). GP-derivatised sterols containing a 3 β ,5 α ,6 β -triol function give characteristic MS 3 spectra this allows their presumptive identification by LC-MS(MS n) in the absence of authentic standards. This is also true for 7-oxo- and 3 β ,7 β -dihydroxysterols [30].

3.2. Metabolism of 5,6-EC

Both 5α,6-EC and 5β,6-EC are converted in vivo by ChEH to 3β,5α,6β-triol [34,35] (Fig. 1) and both Mazzacuva et al. [28] and Jiang et al. [29] have found 3β,5α,6β-triHBA to be a bile acid produced by NPC patients in whom the $3\beta,5\alpha,6\beta$ -triol is abundant. We thus sought to confirm these finding by analysis of NPC patient plasma (n = 3) and to identify the intermediates in the biochemical pathway from $3\beta,5\alpha,6\beta$ -triol to $3\beta,5\alpha,6\beta$ -triHBA. $3\beta,5\alpha,6\beta$ -triol is also reported to be abundant in plasma samples from NPB and LALD patients [15-18], so we similarly analysed plasma samples from patients with these disorders (NPB, n = 3; LALD, n = 2). We also analysed plasma samples from carriers of these disorders (see Supplemental Table S1). To minimise problems of ex vivo oxidation of cholesterol producing 5,6-EC artefacts, cholesterol was removed from samples by reversed-phase solid phase extraction as a first step in sample preparation, consequently 5,6-EC and 3β ,5 α ,6 β -triol, the latter of which may be formed by ex vivo acid catalysed hydrolysis of 5,6-EC, are minimised in plasma samples from healthy donors, including disease carriers (5,6-EC < 0.5 ng/mL; 3β , 5α , 6β -triol < 3.5 ng/mL), see also Supplemental Table S1 which embraces data from the standard reference material NIST SRM 1950, prepared from plasma samples from 100 individuals between 40 and 50 years of age, whose ethnicity was representative of the US population and which included an equal number of men and women [36]. However, upon analysis of patient plasma for 3β,5α,6β-triol, elevated levels were found in NPC (range 31-53 ng/mL) and NPB (23.5-45.5 ng/ mL) samples (Supplemental Table S1, Fig. 3A) confirming the results of earlier studies [13,15-18], although the triol was only increased in one of the LALD samples (3.5–7.5 ng/mL). The precursor of 3β , 5α , 6β -triol, 5,6-EC, was similarly elevated in NPC (1-2.5 ng/mL) plasma, although in only one of the NPB (0.5-2 ng/mL) and LALD (0.5-1 ng/mL) plasma samples. The product of $3\beta,5\alpha,6\beta$ -triol metabolism, $3\beta,5\alpha,6\beta$ -triHBA,

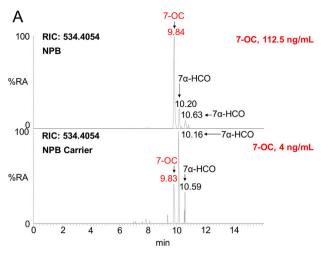
was found in NPC (1–3.5 ng/mL), NPB (1–2 ng/mL) and one of the two LALD (0–1.5 ng/mL) plasma samples (Fig. 3B), but was absent from controls and carriers (\leq 0.1 ng/mL). It is likely that 3β ,5 α ,6 β -triol is converted to 3β ,5 α ,6 β -triHBA in a series of enzyme catalysed reactions, like those operative in the acidic pathway of bile acid biosynthesis [37].

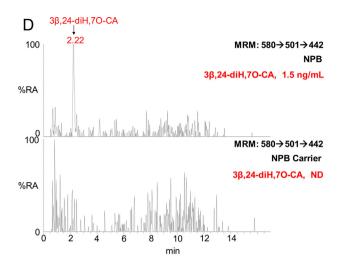
3.2.1. Acidic pathway

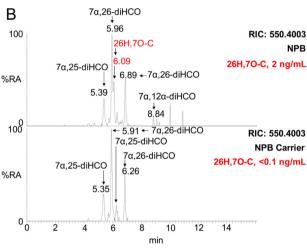
If $3\beta,5\alpha,6\beta$ -triol falls into a branch of the acidic pathway of bile acid biosynthesis the next metabolite, generated by the enzyme cytochrome P450 (CYP) 27A1, should be cholestane-3β,5α,6β,(25R)26-tetrol $(3\beta,5\alpha,6\beta,26$ -tetrol, Fig. 4). Although we do not have an authentic standard of the tetrol, it was presumptively identified in plasma samples from patients with NPC (0.5–1.5 ng/mL) and NPB (1–1.5 ng/mL) based on LC retention time, exact mass and MS3 spectrum, but was absent from control and carrier plasma (Fig. 3C, see Figure 6G in Ref. [30] for MS³ spectrum). Further, metabolism catalysed by CYP27A1 should lead to 3β,5α,6β-trihydroxycholestan-(25R)26-oic $(3\beta,5\alpha,6\beta$ -triHCa). Evidence for the presence of $3\beta,5\alpha,6\beta$ -triHCa in plasma samples from patients with NPC (6-16 ng/mL), NPB (31.5-47.5 ng/mL) and LALD (2-21 ng/mL) was again provided by LC retention time, exact mass and MS³ spectra (see Figure 6D in Ref. [30] for MS^3 spectrum). $3\beta,5\alpha,6\beta$ -triHCa was present at low levels in control and carrier plasma ($\leq 0.5 \text{ ng/mL}$, Fig. 3D). It is noteworthy that in the separate patient groups the concentrations of this acid appears to reflect those of the triol. However, the highest concentration of 3β , 5α , 6β triHCa in NPC plasma is only about half that of the lowest concentration in NPB plasma. Interestingly, the concentration of 3β -hydroxycholest-5en(25R)26-oic (3β-HCA), 3β,7α-dihydroxycholest-5-en-(25R)26-oic $(3\beta,7\alpha-diHCA)$ and 7α -hydroxy-3-oxocholest-4-en-(25R)26-oic (7αH,3O-CA) acids, three cholestenoic acids of the conventional acidic pathway, are also higher in the NPB plasma samples (3β-HCA, 300-648 ng/mL; 3β , 7α -diHCA, 132–336 ng/mL; 7αH.3O-CA. 120.5–316 ng/mL) than in NPC (3β-HCA, 54.5–111 ng/mL; 3β,7αdiHCA, 13.5-42 ng/mL; 7αH,30-CA, 29.5-76 ng/mL) or controls and carriers (3 β -HCA, 53–143.5 ng/mL; 3 β ,7 α -diHCA 10.5–35.5 ng/mL; 7αH,3O-CA, 43.5–69 ng/mL). Importantly, the initial metabolite of the conventional acidic pathway, (25R)26-hydroxycholesterol (26-HC), formed by CYP27A1 metabolism of cholesterol, was higher in NPB plasma samples (40-68 ng/mL) samples than NPC (12-16 ng/mL), LALD (7.5-11.5 ng/mL) or controls and carriers (16-28 ng/mL). These trends were not observed for initial members of the neutral pathway of bile acid biosynthesis i.e. 7α -hydroxycholesterol (7α -HC, NPB, 10-12.5 ng/mL; NPC, 12-35 ng/mL; LALD, 2.5-3 ng/mL; control/carrier, 1.5-13.5 ng/mL), the 24-hydroxylase pathway i.e. 24S-hydroxycholesterol (24S-HC, NPB, 9-34.5 ng/mL; NPC, 11.5-15.5 ng/mL; LALD, 28.5-34.5 ng/mL; control/carriers, 7-13.5 ng/mL) or the 25hydroxylase pathway i.e. 25-hydroxycholesterol (25-HC, NPB, 1.5–2 ng/mL; NPC, 1–1.5 ng/mL; LALD < 1 ng/mL, control/carriers, 0.5-1 ng/mL).

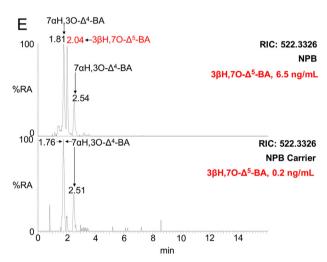
3.3. Shortening C_{27} to C_{24} bile acids

Side-chain shortening of 3β , 5α , 6β -triHCa is likely to proceed in the peroxisome. This would involve formation of a CoA-thioester via the enzyme bile acyl CoA-synthetase (BACS), its epimerisation at C-25 by alpha-methylacyl-CoA-racemase (AMACR), dehydrogenation of the side-chain between C-24 and C-25 by acyl-coenzyme A oxidase 2 (ACOX2), then hydration of the resultant double bond by peroxisomal multifunctional enzyme type 2, also known as D bifunctional protein (DBP), to give 3β , 5α , 6β ,24-tetrahydroxycholestan-26-oic acid (3β , 5α , 6β ,24-tetraHCa) as the CoA thioester (in Fig. 4, for simplicity the individual steps between 3β , 5α , 6β -triHCa and 3β , 5α , 6β ,24-tetraHCa have been omitted and the free acids are shown rather than the CoA thioesters). Using our oxidation/derivatisation method the free acids are preferentially observed and, in the absence of an authentic standard, 3β , 5α , 6β ,24-tetraHCa was presumptively identified by LC









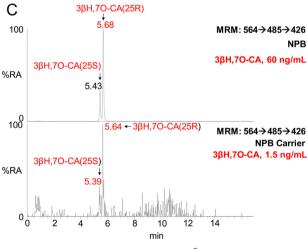


Fig. 5. LC-MS RICs or MRM chromatograms of [2 H₀]GP derivatised oxysterols and bile acids from NPB patient and NPB carrier plasma. RICs of (A) m/z 534.4054 corresponding to [M]⁺ ions of 7α-hydroxycholest-4-en-3-one (7α-HCO) and 7-OC, (B) m/z 550.4003 corresponding to [M]⁺ ions of dihydroxycholest-4-en-3-ones (diHCO) and 26H,7O-C, and (E) m/z 522.3326 corresponding to [M]⁺ ions of hydroxy-3-oxochol-4-enoic acids (H,3O- 4 -BA) and 3βH,7O- 5 -BA. MRM chromatograms of (C) m/z 564 \rightarrow 485 \rightarrow 426 specific for 3βH,7O-CA, and (D) m/z 580 \rightarrow 501 \rightarrow 442 specific for 3β,24-diH,7O-CA. The upper panels show chromatograms of NPB patient and lower panels NPB carrier plasma. Oxysterols with a the 7-oxo function are labelled in red. Concentrations are given in the right-hand corner of chromatograms. GP derivatised oxysterols give *syn* and *anti* conformers, which may or may not be chromatographically separated. For quantification of 26H,7O-C an extended chromatographic gradient was exploited as described in Griffiths et al. [30].

retention time, exact mass and MS³ spectrum in plasma samples from NPC (0.5–2.5 ng/mL), NPB (2.5–4 ng/mL) and LALD (0.5–4.5 ng/mL, see Supplemental Figure S5Q in Ref. [30]). There was no evidence for 3β ,5 α ,6 β ,24-tetraHCa in control or carrier plasma (Fig. 3E). As with 3β ,5 α ,6 β -triHCa, the concentration of 3β ,5 α ,6 β ,24-tetraHCa reflects that of the triol within separate patient groups, but not between groups where concentrations of the acids are higher in NPB plasma. The ultimate C_{24} bile acid CoA thioester should be formed by beta-oxidation through sterol carrier protein x (SPCx) after dehydrogenation by DBP (HSD17B4). As indicated above, free bile acids are preferentially observed with the current method and 3β ,5 α ,6 β -triHBA was detected in NPC (1–3.5 ng/mL), NPB (1–2 ng/mL) and one of the two LALD (0–1.5 ng/mL) patient samples, but not controls or carriers (\leq 0.1 ng/mL) (Fig. 3B). A peroxisomal CoA thioesterase can catalyse the hydrolysis of bile acid CoA thioesters into free bile acids [38,39].

3.4. Metabolism of 7-OC and 7β-HC

The metabolism of 7-OC has been of interest to many investigators. Lyons and Brown reported that it could be metabolised to 26H,7O-C by CYP27A1 in HepG2 cells, while Heo et al. showed that 26H,7O-C and the down-stream CYP27A1 metabolite 3 β -hydroxy-7-oxocholest-5-en-(25R)26-oic acid (3 β H,7O-CA) could be formed by retinal pigment epithelial cells [40,41]. An alternative route for metabolism of 7-OC is reduction to 7 β -HC by HSD11B1 as shown by Hult et al., Larsson et al., Mitic et al. and Schweizer et al. [5-8]. 7-OC itself can be formed via radical reactions from cholesterol or enzymatically from 7-DHC by CYP7A1 or from 7 β -HC by HSD11B2 oxidation [4,9,34] (Fig. 1). We have reported a metabolic pathway from 7-OC to 7-oxo- and 7 β -hydroxy- Δ ⁵-bile acids in SLOS patients where the 7-DHC concentrations are high in plasma and tissue [22-25]. We next sought to investigate if this pathway is active in other diseases where levels of 7-OC are elevated.

Like 3β,5α,6β-triol, 7-OC is elevated in plasma of patients with lysosomal storage disorders. Patients with NPC and NPB show elevated levels of both 7-OC (NPC, 72.5-135.5 ng/mL; NPB 65.5-112.5 ng/mL, Fig. 5A) and 7β-HC (NPC, 29.5–69 ng/mL; NPB, 19–54 ng/mL, Fig. 3A) in plasma compared to controls and carrier (7-OC, < 8.5 ng/mL; 7β-HC, < 2 ng/mL). In plasma from LALD patients 7-OC (9.5-20 ng/mL) was just elevated but not 7 β -HC (1.5–3.5 ng/mL). The ratio of 7-OC to 7 β -HC was about 2.5 for the NPC (2.2 \pm 0.3, mean \pm standard deviation) and NPB (2.7 \pm 0.7) patients, but much more variable for the carriers (12.9 \pm 5.7) and LALD (6.10) patients. Interestingly, the ratio of 7-OC to 3β , 5α , 6β -triol was also about 2.5 for the NPC (2.2 \pm 0.4), NPB (2.5 \pm 0.2) and LALD (2.7) patients, but more variable for the carriers (3.3 \pm 1.6). 7-OC and 7 β -HC are interconverted by HSD11B1 and HSD11B2 and both may fall into branches of the acidic pathway of bile acid metabolism (Fig. 4), giving the bile acids 3β,7β-dihydroxychol-5-enoic (3 β ,7 β -diH- Δ ⁵-BA) and 3 β -hydroxy-7-oxochol-5-enoic $(3\beta H, 70-\Delta^5-BA)$ acid as metabolic products (see Supplemental Table S1). In fact, we observe almost all the necessary pathway intermediates in NPC, NPB and LALD patient plasma in contrast to controls and carriers where intermediates are absent or minor.

3.4.1. Acidic pathway:- 7B-HC

Considering the metabolites of 7 β -HC first, authentic standards are available for 7 β -HC, 7 β ,26-diHC, 3 β ,7 β -dihydroxycholest-5-en-(25R) 26-oic acid (3 β ,7 β -diHCA) and 3 β ,7 β -diH- Δ^5 -BA and each are chromatographically resolved from their 7 α -epimers [42]. 7 β ,26-diHC was absent from control and carrier plasma but was present in two of the three NPC (0–1.5 ng/mL) and NPB (0–0.5 ng/mL) and one of the two LALD (0–0.2 ng/mL) plasma samples (Fig. 3C). The further CYP27A1 metabolite, 3 β ,7 β -diHCA, is a normal constituent of plasma from healthy individuals (control and carriers, 3–5 ng/mL), but is greatly elevated in NPC (40.5–229 ng/mL) and NPB (97–174 ng/mL) plasma and to a lesser extent in LALD (11–52.5 ng/mL) plasma (Fig. 3D). We do

not have an authentic standard for 3β , 7β ,24-trihydroxycholest-5-en-26-oic acid $(3\beta$, 7β ,24-triHCA), however, the 7α -epimer $(3\beta$, 7α ,24-triHCA) is commercially available, this allows a presumptive, rather than definitive, identification of 3β , 7β ,24-triHCA based on exact mass, MS³ spectrum and retention time. The 7β -epimer is essentially absent from control and carrier plasma (< 0.1 ng/mL), but present in two of the three NPC (0–1.5 ng/mL), all three of the NPB (2–4.5 ng/mL) and both of the LALD (0.5–2.5 ng/mL) plasma samples (Fig. 3E). The ultimate metabolite of the pathway, 3β , 7β -diH- Δ^5 -BA was present in control and carrier plasma (0.5–2 ng/mL), but at higher levels in NPC (19–50 ng/mL), NPB (14–40 ng/mL) and in LALD (4.5–14 ng/mL) (Fig. 3B).

3.4.2. Acidic pathway:- 7-OC

Considering next the pathway from 7-OC, authentic standards are available for 26H,7O-C and 3βH,7O-CA allowing their definitive identification. 26H,7O-C is present in control and carrier plasma (≤0.5 ng/ mL) and is found to be elevated in NPC (1-2.5 ng/mL), NPB (1-2 ng/ mL) but not LALD (< 0.5 ng/mL) plasma (Fig. 5B). The 7-oxo acid, 26H,7O-CA, is present in plasma from healthy controls and carriers (1-1.5 ng/mL) but greatly elevated in plasma from NPC (13.5-60.5 ng/mL)mL), NPB (39-60 ng/mL) and LALD (9.5-52 ng/mL) (Fig. 5C). An authentic standard is not available for 3β,24-dihydroxy-7-oxocholest-5en-26-oic acid (3β,24-diH,7O-CA), however, based on exact mass, retention time and MS3 spectra it was presumptively identified in two of the three NPC (0-0.5 ng/mL), all three NPB (0.5-1.5 ng/mL) and both LALD (1-3 ng/mL) plasma samples (Fig. 5D, see Fig. 4G in Ref. [30]). 3β,24-diH,7O-CA was absent or of very low abundance in control and carrier plasma samples ($< 0.2\, ng/mL$). No authentic standard was available for $3\beta H,7O-\Delta^5$ -BA. It was presumptively identified in control and carrier (\leq 0.5 ng/mL), NPC (1.5–6.5 ng/mL), NPB (2–6.5 ng/mL) and LALD (1-5.5 ng/mL) plasma (Fig. 5E, see Fig. 5D in Ref. [30]).

3.4.3. Metabolite ratios: 70/7\beta

Unlike the situation for 7-OC and 7B-HC which are in a concentration ratio of about 2.5 for both NPC and NPB patents, the 7βhydroxy acids, 3β,7β-diHCA and 3β,7β-diH-Δ⁵-BA, are more abundant than their 3-oxo analogues 3 β H,7O-CA and 3 β H,7O- Δ ⁵-BA in the ratio of about 3.5 and 8 for the C_{27} (NPC, 3.5 \pm 0.4; NPB, 3.1 \pm 0.8) and C_{24} (NPC, 9.2 \pm 2.7; NPB, 6.6 \pm 0.2) acids, respectively. Both ratios are also greater than 1 for the LALD patients (C27, 1.1; C24, 3.43) and carriers (C_{27} , 3.1 \pm 0.7; C_{24} , 3.5 \pm 1.9) and controls. In contrast to the situation for 3β , 5α , 6β -triHCa, where concentrations are higher in NPB than NPC plasma (see Supplemental Table S1) the concentration ranges of the acids 3β , 7β -diHCA and 3β H,7O-CA both overlap for the two disorders. In NPB plasma other acids 3β -HCA, 3β , 7α -diHCA and 7αH,3O-CA were higher in NPB than in NPC plasma, as was their precursor 26-HC. An absence of a similar trend for 3β , 7β -diHCA and 3βH,7O-CA suggests that their formation is not dependent on CYP27A1 activity, but rather substrate availability.

4. Discussion

The observation here and elsewhere of elevated levels of 3β , 5α , 6β -triol and 7-OC in the lysosomal storage diseases NPC and NPB points to their endogenous formation [22]. While 7-OC can be formed enzymatically from 7-DHC in SLOS [43], there is little convincing evidence for the enzymatic formation of 5,6-EC, the precursor of 3β , 5α , 6β -triol, hence it is likely that it is formed through free radical reactions *in vivo*. The absence of high levels of 7-DHC in NPC and NPB also points to the formation of 7-OC via *in vivo* free radical reactions.

4.1. Formation of 3β,5α,6β-triHBA

Until recently little was known about the metabolism of 5,6-EC. It is established that 5,6-EC can be enzymatically converted to the 3β ,5 α ,6 β -triol by ChEH [35], but only in 2016 was it shown that the

triol could be converted to the bile acid $3\beta,5\alpha,6\beta$ -triHBA in man [28,29]. A similar product has also been proposed to be generated in rat [44,45], however, a description of the pathway from $3\beta,5\alpha,6\beta$ -triol to $3\beta,5\alpha,6\beta$ -triHBA has not previously been reported.

There are two major pathways of bile acid biosynthesis, the neutral and acidic pathways, and two more minor pathways, the sterol 24-hydroxylase and 25-hydroxylase pathways [37]. The neutral pathway is initiated by 7α -hydroxylation of cholesterol by CYP7A1, but in the metabolism of 3β , 5α , 6β -triol the absence of a 7α -hydroxy group in the ultimate bile acid argues against operation of this pathway. The cholesterol 24-hydroxylase pathway is also an unlikely route for 3β , 5α , 6β -triol metabolism as this starts with a reaction catalysed by CYP46A1, predominantly expressed in brain [37]. This leaves the acidic and 25-hydroxylase pathways for biosynthesis of 3β , 5α , 6β -triHBA from 3β , 5α , 6β -triol. As in the present study we have identified most of the intermediates in a novel branch of the acidic pathway from 3β , 5α , 6β -triol to 3β , 5α , 6β -triHBA in plasma, it is highly likely that this is the pathway followed (Fig. 4).

4.2. Formation of $3\beta H$, $70-\Delta^5$ -BA and 3β , 7β -diH- Δ^5 -BA

The metabolism of 7-OC has been the subject of greater interest than that of $3\beta,5\alpha,6\beta$ -triol [40,41,46], as the former compound is found in atherosclerotic lesions [47]. 7-OC can be metabolised to 26H,7-OC and 3βH,7O-CA [40,41], hence it likely to also follow another branch of the acidic pathway to generate $3\beta H,7O-\Delta^5-BA$. The data reported here confirms this hypothesis as judged by the preponderance of relevant pathway intermediates identified in plasma of NPC and NPB patients. 7-OC can alternatively be converted to 7β -HC [5-8] and this oxysterol may similarly fall into a related branch of the acidic pathway with the formation of 3β , 7β -diH- Δ^5 -BA. These 7-OC and 7β -HC branches of the acidic may be interconvertible through HSD11B1 which can convert 7oxo to 7β-hydroxy groups and HSD11B2 which can catalyse the reverse reactions (Fig. 4). In fact, in an early study Alvelius et al. identified both $3\beta H,7O-\Delta^5$ -BA and $3\beta,7\beta$ -diH- Δ^5 -BA as their sulphate and glycine conjugates in urine of an NPC patient [26], although the plasma pattern of oxysterols was apparently normal. In the present study, the ratio of 7oxo- to 7β-hydroxy-sterols may provide an insight into the activity of HSD11B enzymes. HSD11B1 is the enzyme responsible for reducing 7-OC to 7β-HC in mouse and man [5,7], while HSD11B2 can also catalyse the reverse reaction [9]. The HSD11B1 enzyme also catalyses the reduction of cortisone to cortisol in man and of 11-dehydrocorticosterone to corticosterone in mouse, the reduced metabolites being ligands to the glucocorticoid receptor. On the other hand, HSD11B2 is the enzyme that oxidises cortisol to cortisone and has a similar activity towards 7β-HC and also 7β ,25-dihydroxycholesterol [9,48,49]. Data from the present study indicates that as the acidic pathway proceeds the ratio of 7βhydroxy to 7-oxo metabolites increase, this is true for patients, controls and carriers. This may be explained by the HSD11B reductase having a dominant effect over the oxidase as the pathway descends.

HSD11B2 also has activity towards 3β , 5α , 6β -triol, oxidising it to the oncometabolite 3β , 5α -diHC-6O, while HSD11B1 catalyses the reverse reaction [11]. It appears that the enzymes ChEH, HSD11B1 and HSD11B2 and CYP27A1 sit at a fulcrum balancing the formation from 5α , 6-EC of the tumour suppressor DDA and from 5, 6-EC, through 3β , 5α , 6β -triol, the oncometabolite 3β , 5α -diHC-6O or the bile acid 3β , 5α , 6β -triHBA. It should also be noted that 26H, 70-C activates the Hh signalling pathway, constitutive activation of which is linked to tumorigenesis further connecting the HSD11B and CYP27A1 enzymes to oncology.

In light of the low number of patient samples analysed the diagnostic value of bile acid precursors can only be speculated on. However, the high abundance of the C_{27} acids $3\beta,5\alpha,6\beta$ -triHCa, $3\beta,7\beta$ -diHCA (Figure 3D) and 3β H,7O-CA (Fig. 5C) in NPC and NPB plasma suggests that these three acids in combination may diagnose these disorders. It may not be possible to distinguish these diseases from LALD as the three

acids are elevated in this disorder also.

5. Conclusion

In man, most of the primary bile acids are synthesised in the liver through the neutral pathway of bile acid biosynthesis. The acidic pathway can proceed extrahepatically [37], while quantitatively less important in adult this pathway may be dominant in neonates [50]. We show here that novel branches of the acidic pathway, starting from oxysterols formed non-enzymatically, are important for the formation of unusual bile acids in patients with lysosomal storage disease. In healthy controls, the acidic pathway also proceeds to generate unusual 7β-hydroxy and 7-oxo bile acids but these are quantitatively minor.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.freeradbiomed.2019.04.020.

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