

Gas chromatographic–mass spectrometric determination of leukotriene E₄ in human urine using deuterium-labelled leukotriene E₄ standards

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ABSTRACT

The utility of two deuterium-labelled leukotriene (LT) E₄ analogs, *e.g.* [20,20,20-²H₃]LTE₄ and [14,15,17,17,18,18-²H₆]LTE₄, as internal standards for the determination of LTE₄ in human urine by gas chromatography–mass spectrometry (GC–MS) was investigated. ²H-Exchange during hydrogenation occurred both in [20,20,20-²H₃]LTE₄ and [14,15,17,17,18,18-²H₆]LTE₄ in an extent of 9.4 ± 0.5% and 67.3 ± 0.6% (mean ± S.D., *n* = 6), respectively. The lower extent of ²H-exchange in [20,20,20-²H₃]LTE₄ allowed a more accurate quantitation than the use of [14,15,17,17,18,18-²H₆]LTE₄. Applying [20,20,20-²H₃]LTE₄ as internal standard the coefficients of variation for the intra- and inter-assay determination of LTE₄ in human urine were 5.7% and 6.2% (*n* = 4), respectively. The inter-assay coefficient of variation for [14,15,17,17,18,18-²H₆]LTE₄ was 15%. Using [20,20,20-²H₃]LTE₄ as internal standard and GC–MS, healthy volunteers were found to excrete 17 ± 10 nmol LTE₄ per mol creatinine (mean ± S.D., *n* = 11). Similar excretion rates for LTE₄ in urine of healthy volunteers were found using GC–tandem MS with [1,1-¹⁸O₂]LTE₄ as internal standard. Our results demonstrate that [20,20,20-²H₃]LTE₄ is a suitable internal standard for the GC–MS determination of urinary LTE₄.

INTRODUCTION

Cysteinyl leukotrienes are potent mediators of inflammation [1]. They are thermally labile and non-volatile compounds. Their direct analysis by gas chromatography–mass spectrometry (GC–MS) is therefore impossible. However, catalytic reduction and desulphurization of these

substances to 5-hydroxyeicosanoic acid (5-HEA) enables cysteinyl leukotrienes to be determined by GC–MS [2]. A prerequisite for their quantitation by GC–MS is the availability of stable isotope-labelled cysteinyl leukotrienes analogs. Recently, the synthesis and the utility of [1,1-¹⁸O₂]LTE₄ as internal standard for the GC–MS determination of urinary LTE₄ [3], the index metabolite of cysteinyl leukotrienes in humans [4,5], has been described. In the literature the synthesis of [20,20,20-²H₃]LTE₄ [6] and

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[8,9,10,11- $^{13}\text{C}_4$]LTE₄ [7] has also been described. From these [20,20,20- $^2\text{H}_3$]LTE₄ is commercially available (Biomol, Hamburg, Germany). [14,15,17,17,18,18- $^2\text{H}_6$]LTE₄ (99.7 atom% ^2H) was synthesized in our laboratory starting from [14,15,17,17,18,18- $^2\text{H}_6$]LTA₄ methylester [8] and cysteine methylester hydrochloride. In particular with respect to the common phenomenon of ^2H -exchange by ^1H during catalytic hydrogenation, which has also been reported for some ^2H -labelled cysteinyl leukotrienes [2], it was of interest to investigate whether the commercially available [20,20,20- $^2\text{H}_3$]LTE₄ and the newly synthesized [14,15,17,17,18,18- $^2\text{H}_6$]LTE₄ are useful as internal standards for the determination of LTE₄ in human urine by GC–MS. This study demonstrates that despite $^2\text{H}/\text{H}$ -exchange both [20,20,20- $^2\text{H}_3$]LTE₄ and [14,15,17,17,18,18- $^2\text{H}_6$]LTE₄ are suitable internal standards. However, [20,20,20- $^2\text{H}_3$]LTE₄ permits more accurate quantification of urinary LTE₄ by GC–MS. Using [20,20,20- $^2\text{H}_3$]LTE₄ the urinary excretion rate of LTE₄ in healthy normal volunteers was assessed.

EXPERIMENTAL

Chemicals and reagents

[20,20,20- $^2\text{H}_3$]LTE₄ (99.8 atom% ^2H) was obtained from Biomol (Hamburg, Germany). [14,15- $^3\text{H}_2$]LTE₄ was obtained from DuPont (Dreieich, Germany). Hydrogen and deuterium gas (99 atom% ^2H) were obtained from Merck (Darmstadt, Germany). The catalyst 5% (w/w) Rh on alumina was obtained from Fluka (Neu-Ulm, Germany) and was washed with ethanol prior to use. Pentafluorobenzyl (PFB) bromide was purchased from Aldrich (Steinheim, Germany). N,O-Bis(trimethylsilyl)trifluoroacetamide (BSTFA) was from Pierce (Rockford, IL, USA). [1,1- $^{18}\text{O}_2$]LTE₄ (90 atom% ^{18}O) was synthesized as described previously [3].

Catalytical reduction/desulphurization

Catalytical reduction/desulphurization of LTE₄ and labelled-LTE₄, and derivatization of labelled and unlabelled 5-HEA to the corre-

sponding PFB ester trimethylsilyl (TMS) ether derivatives using PFB bromide and BSTFA, respectively, were performed in most experiments as described previously [3,9]. For catalytical reduction/desulphurization hydrogen as well as deuterium gas were used.

Solid-phase extraction and reversed-phase high-performance liquid chromatography

Endogenous LTE₄ and the stable isotope-labelled LTE₄ (2 ng in most experiments) added to 20-ml urine samples were solid-phase extracted, purified and isolated by reversed-phase high-performance liquid chromatography (RP-HPLC) as described elsewhere [9–11]. The retention times of unlabelled LTE₄, [20,20,20- $^2\text{H}_3$]LTE₄, [14,15,17,17,18,18- $^2\text{H}_6$]LTE₄ and [1,1- $^{18}\text{O}_2$]LTE₄ in this system were (mean \pm S.D.): 48.5 ± 1.4 ($n = 10$), 48.1 ± 1.5 ($n = 3$), 47.9 ± 1.4 ($n = 3$), and 48.4 ± 1.5 ($n = 10$) min, respectively. RP-HPLC fractions were collected in 1-min intervals between the 45th and 53rd min and aliquots thereof were counted. Radioactivity containing fractions were pooled and treated further for catalytic reduction/desulphurization.

Gas chromatographic–mass spectrometric conditions

GC–MS was performed on a Hewlett–Packard MS Engine 5989A directly connected with a gas chromatograph HP 5890 series II (Waldbronn, Germany). The gas chromatograph was equipped with a fused silica capillary column HP1 (12 m \times 0.25 mm I.D., 0.25 μm film thickness) maintained at a temperature of 100°C. Helium (35 kPa) and methane (200 Pa) were used as carrier and reagent gas for negative-ion chemical ionization (NICI), respectively. The following oven temperature program was used: the column was held for 2 min at 100°C, then the temperature was increased to 250°C at a rate of 25°/min and next to 320°C at a rate of 4°C/min. The interface and ion source were kept at 280°C and 225°C, respectively. Electron energy and electron current were set at 230 eV and 300 μA , respectively. Aliquots of 1 μl were injected in the splitless mode.

GC–tandem MS (GC–MS–MS) was carried

out on a Finnigan MAT gas chromatograph 9611 equipped with a fused silica capillary column OV1, 25 m \times 0.25 mm I.D., film thickness 0.25 μ m (Macherey–Nagel, Düren, Germany) interfaced with a triple-stage quadrupole mass spectrometer Finnigan MAT TSO 45 (San Jose, CA, USA) operating in the NICI mode with methane as reagent gas at a pressure of 65 Pa. Helium was used as carrier gas at a pressure of 50 kPa. The ionization energy was 90 eV and the electron current 300 μ A. Argon was used for collision activated dissociation (CAD) at a cell pressure of 0.2 Pa. The collision energy was 14 eV. Constant temperatures were maintained at the interface (280°C), ion source (130°C) and injector (280°C). The same oven temperature program was used as described above for GC–MS. Aliquots of 1–2.5 μ l were injected in the splitless mode.

RESULTS

Fig. 1 shows NICI mass spectra obtained from the PFB-TMS derivatives of catalytically hydrogenated unlabelled LTE_4 (a), $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ (b), $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ (c), and $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$ (d). Catalytic hydrogenation of unlabelled LTE_4 led to a single compound, *e.g.* 5-HEA, with the most intensive signal at m/z 399 ($[\text{M} - \text{PFB}]^-$) (Fig. 1a). The major mass fragment in the mass spectrum derived from $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ (Fig. 1b) is m/z 402 which corresponds to $[^2\text{H}_3]\text{-5-HEA}$. The less intensive signal at m/z 399 corresponds to 5-HEA and results from exchange of all three ^2H atoms with ^1H atoms during hydrogenation. The mass spectrum derived from catalytically hydrogenated $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ (Fig. 1c) shows a group of intensive ions at m/z 399, 400,

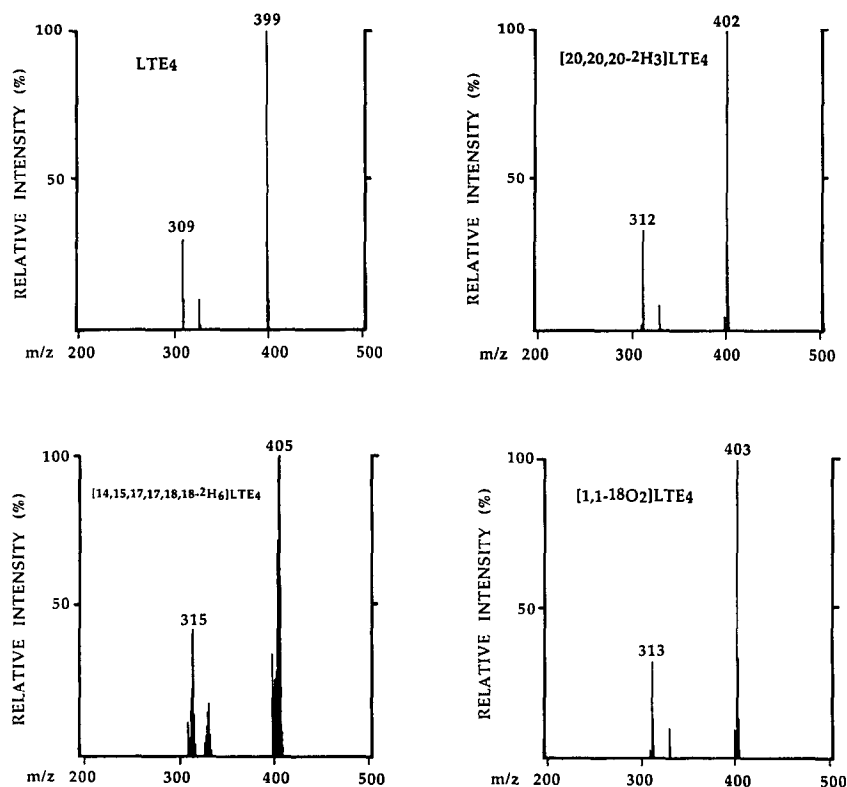


Fig. 1. NICI mass spectra of the PFB-TMS derivatives of reduced/desulphurized LTE_4 (a), $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ (b), $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ (c), and $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$ (d).

401, 402, 403, 404, 405, 406 and 407. These signals, except for m/z 406 and 407, correspond to 5-HEA with various ^2H atoms. The signal at m/z 399 corresponds to 5-HEA from exchange of all six ^2H atoms of $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ with ^1H atoms. The major mass fragments in the mass spectrum derived from $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$ (Fig. 1d) are m/z 403 and m/z 401 corresponding to $[1,1\text{-}^{18}\text{O}_2]\text{-5-HEA}$ and $[1,1\text{-}^{18}\text{O}^{16}\text{O}]\text{-5-HEA}$.

The extent of the $^2\text{H}/\text{H}$ -exchange and its variability in both $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ are of great importance for quantitation of LTE_4 . Six samples of each $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$, $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ and $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$ (0.5 ng, 1 ng, 2 ng, 5 ng, 10 ng and 50 ng) were subjected separately to catalytical reduction/desulphurization using hydrogen gas. GC-MS analysis of the PFB-TMS derivatives was carried out by selected ion monitoring (SIM) on m/z of the ions $[\text{M} - \text{PFB}]^-$, e.g. m/z 399, 400, 401 and 402 for $^2\text{H}_3\text{-LTE}_4$, m/z 399 to m/z 405 for $^2\text{H}_6\text{-LTE}_4$, and m/z 399, 401 and 403 for $^{18}\text{O}_2\text{-LTE}_4$. The ratio m/z 399 to m/z of $[\text{M} - \text{PFB}]^-$ (m/z 402 for $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$, m/z 405 for $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ and m/z 403 for $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$) was determined from the peak areas of the corresponding ions of each LTE_4 . Similarly, the relative contributions of the ion at m/z 399 and of the ions $[\text{M} - \text{PFB}]^-$ were calculated by dividing the peak area of these ions by the sum of the peak areas of all corresponding ions of each LTE_4 , respectively. The results of these

experiments are summarized in Table I. This table clearly shows that $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$ is superior with respect to accuracy over the ^2H -analogues. However, the relative low $^2\text{H}/\text{H}$ -exchange in $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and its constant value should allow its use as an internal standard for LTE_4 in GC-MS. The corresponding data for $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ in Table I suggest that this analogue is less suitable as an internal standard.

Linearity was checked by separate catalytic reduction/desulphurization of synthetic LTE_4 up to 10 ng using 2 ng each of $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$, $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$, and $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$. For all three internal standards linear relationships ($r^2 > 0.989$) were obtained by SIMs on m/z 399, 402, 405, and 403, respectively. Furthermore, linearity ($r^2 > 0.957$) was also observed for LTE_4 added (up to 1000 pg/ml) to human urine samples (20 ml) using 2 ng of $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ or 2 ng of $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$, as observed by GC-MS-MS in the selected reaction monitoring (SRM) mode of the daughter ions ($[\text{M} - \text{PFB} - \text{CH}_3\text{CH}_2\text{COO-TMS}]^-$) at m/z 253 for LTE_4 , m/z 256 for $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and m/z 259 for $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ [3].

Based on the data of Table I (1st and 2nd line from the 2nd and 3rd entry) the following equations were developed and used to calculate the concentration of urinary LTE_4 (C_{LTE_4}) from the measured ratio m/z 399/402 or 253/256 for $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ (eqn. 1) and m/z 399/405 or 253/259 for $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$

TABLE I

EXTENT AND VARIABILITY OF THE $^2\text{H}/\text{H}$ -EXCHANGE DURING CATALYTIC ONE-STEP REDUCTION/DESULPHURIZATION OF $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ AND $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$

Values are mean \pm S.D. from six determinations.

LTE_4	Ratio m/z 399 to m/z of $[\text{M} - \text{PFB}]^-$ ^a	Percentage of m/z 399 from total ^b	Percentage of m/z $[\text{M} - \text{PFB}]^-$ ^a from total ^b
$^2\text{H}_3\text{-LTE}_4$	0.059 ± 0.024	4.17 ± 0.14	90.6 ± 4.8
$^2\text{H}_6\text{-LTE}_4$	0.190 ± 0.030	5.32 ± 2.34	32.7 ± 5.6
$^{18}\text{O}_2\text{-LTE}_4$	0.002 ± 0.0007	0.70 ± 0.02	89.8 ± 1.2

^a $[\text{M} - \text{PFB}]^-$ corresponds to the ions at m/z 402, 405 and 403 for $^2\text{H}_3\text{-LTE}_4$, $^2\text{H}_6\text{-LTE}_4$ and $^{18}\text{O}_2\text{-LTE}_4$, respectively.

^b Total is the sum of the peak areas of the $[\text{M} - \text{PFB}]^-$ ions for the corresponding LTE_4 analog as described in the text.

(eqn. 2) and from the concentration in the urine of the weighted internal standard ($C_{[20,20,20-^2H_3]LTE_4}$ or $C_{[14,15,17,17,18,18-^2H_6]LTE_4}$).

$$C_{LTE_4} = \{(m/z\ 399/402S \times 0.906) - 0.042\} \times C_{[20,20,20-^2H_3]LTE_4} \quad (1)$$

$$C_{LTE_4} = \{(m/z\ 399/405 \times 0.327) - 0.053\} \times C_{[14,15,17,17,18,18-^2H_6]LTE_4} \quad (2)$$

LTE_4 was determined by GC–MS in 20-ml urine samples of a healthy volunteer spiked separately with 2 ng of each $[1,1-^{18}O_2]LTE_4$, $[20,20,20-^2H_3]LTE_4$ and $[14,15,17,17,18,18-^2H_6]LTE_4$ as the internal standards. Applying eqns. (1) and (2) the following concentrations of LTE_4 (pg/ml) in the urine were observed (mean \pm S.D., $n = 4$): 60 ± 3 using $[1,1-^{18}O_2]LTE_4$, 65 ± 4 using $[20,20,20-^2H_3]LTE_4$ and 58 ± 10 using $[14,15,17,17,18,18-^2H_6]LTE_4$. The corresponding coefficients of variation (C.V.) were 4.8%, 5.7% and 15%, respectively. Using $[20,20,20-^2H_3]LTE_4$ as internal standard the inter-assay C.V. for the determination of LTE_4 in human urine was 6.2% ($n = 4$).

Healthy normal volunteers were found to excrete 17 ± 10 nmol LTE_4 per mol creatinine (mean \pm S.D., $n = 11$) as measured by GC–MS using $[20,20,20-^2H_3]LTE_4$. Representative tracings from the GC–MS and GC–MS–MS analysis of urinary LTE_4 of a healthy volunteer using $[20,20,20-^2H_3]LTE_4$ are shown in Fig. 2. Linear regression analysis of the data obtained by GC–MS and those obtained by GC–MS–MS of the same samples gave a correlation coefficient of 0.958. The lower values from the GC–MS analyses probably result from the fact that the peak for endogenous LTE_4 is not base-line resolved (Fig. 2a) so that integration of the peak area as well as the peak height gave lower values compared to GC–MS–MS (Fig. 2b). The urinary LTE_4 concentrations in healthy volunteers measured by GC–MS are within the range for healthy volunteers as measured by GC–MS–MS using $[1,1-^{18}O_2]LTE_4$ [12] and by radioimmunoassay [9,10]. An example for the utility of $[14,15,17,17,18,18-^2H_6]LTE_4$ as internal standard is shown in Fig. 3.

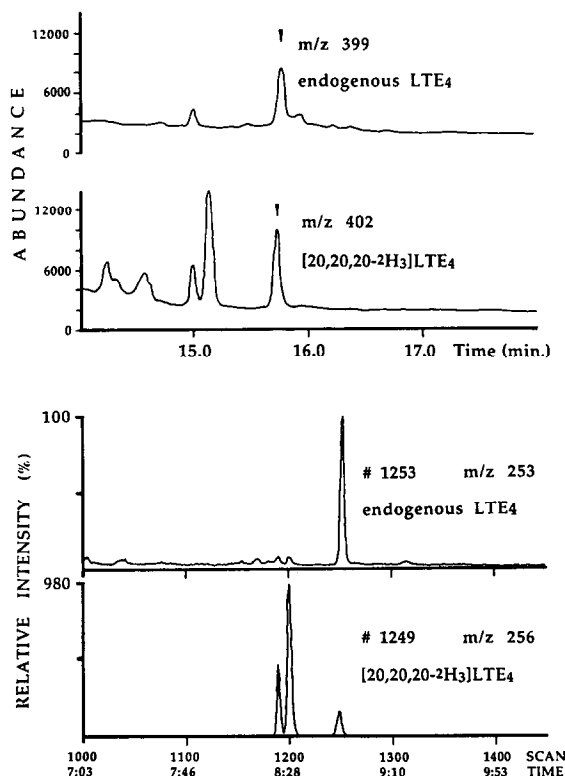


Fig. 2. Partial GC–MS (a) (Hewlett-Packard instrument) and (b) GC–MS–MS (Finnigan MAT instrument) chromatograms from the analysis of urinary LTE_4 in a 20-ml urine sample of a healthy volunteer spiked with 2 ng of $[20,20,20-^2H_3]LTE_4$. In GC–MS the ions at m/z 399 and m/z 402 for endogenous and $[20,20,20-^2H_3]LTE_4$ were detected in the SIM mode. In GC–MS–MS the daughter ions at m/z 253 and m/z 256 derived from the corresponding parent ions at m/z 399 and m/z 402 were detected in the SRM mode. The concentration of LTE_4 in this urine sample was determined to be 49 pg/ml by GC–MS and 50 pg/ml by GC–MS–MS.

DISCUSSION

Accurate determination of urinary LTE_4 by GC–MS requires several procedures for its extraction from urine, chromatographic separation from other cysteinyl leukotrienes and interfering compounds, and also a stable isotope-labelled LTE_4 analogue. The synthesis of ^{18}O -, 2H -, and ^{13}C -labelled cysteinyl leukotrienes has been recently reported by various groups [3,6,7]. $[1,1-^{18}O_2]LTE_4$ was demonstrated to fulfill all the criteria for an ideal internal standard for the GC–MS determination of LTE_4 in human urine

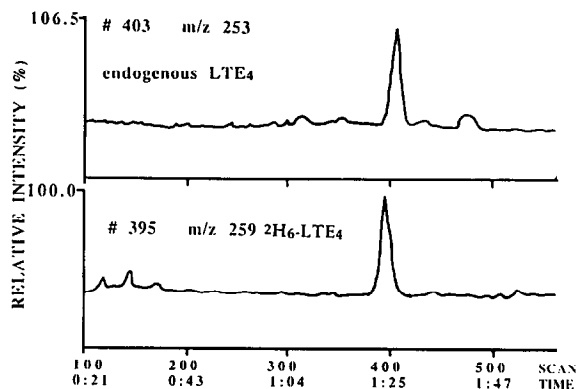


Fig. 3. Partial GC–MS–MS chromatogram from the analysis of urinary LTE_4 in a 20-ml urine sample of a healthy volunteer spiked with 4 ng of $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$. SRM of m/z 253 and 259 for endogenous LTE_4 and the internal standard, respectively. The concentration of LTE_4 in this urine was determined to be 55 pg/ml.

[3]. In particular with respect to the well known $^2\text{H}/\text{H}$ -exchange during catalytic hydrogenation we investigated the usefulness of the commercially available $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and the newly synthesized $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ as internal standards for quantitative measurements by GC–MS. Our study demonstrates that, despite $^2\text{H}/\text{H}$ -exchange in $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ (Figs. 1 and 2; Table I) under the conditions needed for sufficient recovery of 5-HEA [2,3], these compounds can be used as internal standards for LTE_4 in GC–MS (Fig. 2). The $^2\text{H}/\text{H}$ -exchange is more reproducible for $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ than for $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ (Table I). This is also reflected in the C.V. for the quantification by GC–MS of urinary LTE_4 applying these standards: The intra-assay C.V. using $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ (5.7%) is only little higher than the corresponding C.V. value using $[1,1\text{-}^{18}\text{O}_2]\text{LTE}_4$ (4.8%), but significantly lower than that using $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ (15%).

The differences in the extent of the $^2\text{H}/\text{H}$ -exchange in $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ result apparently from the position of ^2H in the molecules of the deuterated cysteinyl leukotrienes. As two ^2H atoms in $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ are olefinic and the other four are in their close proximity, these ^2H atoms seem to be more activated during the

catalytic process than the three end-standing ^2H atoms in $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$. Similar results were observed when $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ were subjected to catalytic reduction/desulphurization using deuterium gas (data not shown). Balazy and Murphy have reported significant loss of ^2H through H-atom scrambling in various ^2H -labelled cysteinyl leukotrienes without however giving the position of the ^2H atoms in these standards [2].

Analogous to LTC_4 , LTD_4 and LTE_4 the metabolites of ω - and β -oxidation such as 20-carboxy- LTE_4 , 18-carboxy-dinor- LTE_4 , 16-carboxy- Δ^{13} -tetranor- LTE_4 and 14-carboxy- LTE_3 , which also appear in human urine [13], can be measured by GC–MS as their monohydroxy dicarboxylic acids [13]. The results from our investigation on the utility of $[20,20,20\text{-}^2\text{H}_3]\text{LTE}_4$ and $[14,15,17,17,18,18\text{-}^2\text{H}_6]\text{LTE}_4$ as internal standards suggest that if ^2H -labelled analogs have to be used for quantitative measurements the label must stay preferably at positions 2, 3 and/or 4. Since cysteinyl leukotrienes are not ω - and β -oxidized from C-1 [14] enzymatic tools could be employed for the preparation of ^2H -labelled ω - and β -oxidized cysteinyl leukotrienes starting from chemically synthesized precursors such as $[2,2,3,3\text{-}^2\text{H}_4]\text{LTC}_4$. Alternatively, ^{18}O labelling could be employed starting from unlabelled materials [3,15].

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