

Aim: One advantage of targeted lipidomics versus untargeted approach is higher sensitivity. This is useful for human plasma samples, in which the presence of highly abundant isobaric sterol compounds (e.g. cholesterol) complicates the separation. Moreover, the issue can be solved by derivatization further lowering detection limits and column load. The aim of this study was method development for analysis of noncholesterol sterols and oxysterols including isobaric species in human plasma.

Methods: Lipids in human plasma were directly saponified or extracted [1], and the isolated SPE fractions [2] were derivatized to carbamates [3] and sterol lipids analysed with LC/MSMS platform. The method was applied on comparison of clinical samples of individuals with metabolic syndrome to control group.

HPLC

column: Hypersil GOLD column type C18 (150 × 2.1 mm, 3 μm); flow 0.4 ml/min, 50 °C

mobile phase: sterols A H₂O (0.1% v/v) HCOOH) B ACN/CH₃OH (7/3 v/v) A/B 25/75 (0 min) - 25/75 (30 min) - 15/85 (59 min) - 25/75 (60 min), equil. 4 min oxysterols A H₂O (0.1% v/v) HCOOH) B ACN/CH₃OH/i-PrOH (400/100/5 v/v/v) A/B 35/65 (0 min) - 28/72 (10 min) - 15/85 (31 min) - 35/65 (33 min), equil. 3 min

MS

TSQ Quantum Access Max with HESI-II probe
 spray voltage +4300 V vaporizer 350 °C
 sheath gas 34 a.u. auxiliary flow 15 a.u.
 sweep gas 1.2 a.u. capillary 320 °C
 skimmer offset not used collision gas Ar, 1.0 mTorr

SRM parameters: parent ion → products (collision E); tube lens voltage
 desmosterol 547.5 → 166(40)/181(29); 117
 lathosterol 549.5 → 166(40)/181(30); 123
 cholesterol 549.5 → 166(42)/181(31); 124
 epicooprostanol(IS) 551.5 → 166(43)/181(32); 128
 5α-cholestanol 551.5 → 166(43)/181(22); 123
 campesterol 563.5 → 166(42)/181(33); 119
 β-sitosterol 577.5 → 166(44)/181(34); 122
 lanosterol 589.5 → 166(46)/181(31); 127

oxysterols with keto group (C4,3b70) 563.5 → 181(18)/883(21); 89
 C4-d7 (IS) 570.5 → 166(18)/390.5(21); 89
 oxysterols with 3OH (TRIOL) 585.5 → 166(42)/181(35); 113
 oxysterols with 2OH 565.5 → 166(42)/181(24); 111
 oxysterols-d4 (IS) 569.5 → 166(42)/181(24); 111
 oxysterols-d7 (IS) 572.5 → 166(42)/181(24); 111

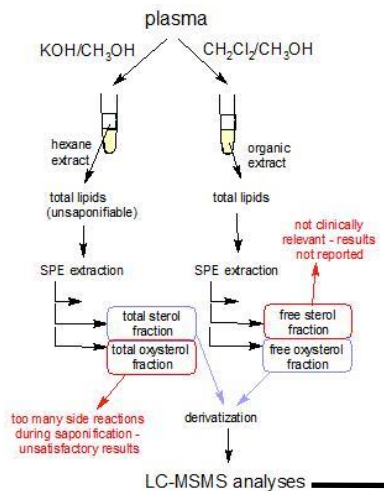


Figure 1 Total sterol fraction analysis

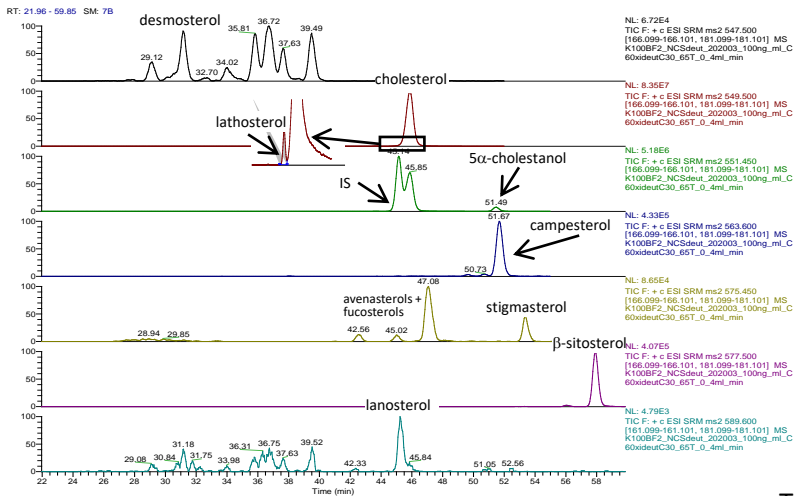
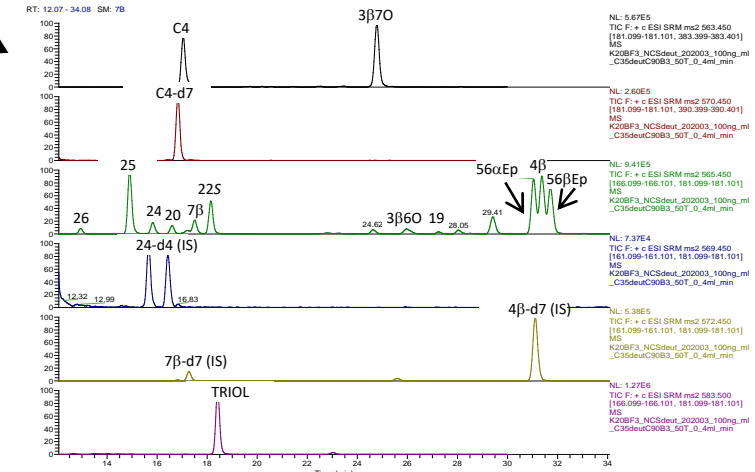


Figure 2 Free oxysterol fraction analysis



IS – internal standard, numbers indicate the location of hydroxyl group in oxysterol molecule, TRIOL – 3β,5α,6β-trihydroxycholesterol

Table 1 Basic characteristics of groups

parameter	control	MetSy
gender (M/F)	7/7	7/7
age (yrs)	33.5 ± 7.1 ^a	37.2 ± 4.9
BMI (kg.m ⁻²)	23.4 ± 2.0	31.4 ± 2.9***
TAG (mmol/L)	0.88 ± 0.30	2.81 ± 2.39*** ^b
TC (mmol/L)	4.65 ± 0.66	5.19 ± 1.29***

^a – mean ± SD; BMI – body mass index, TAG – triacylglycerols, TC – total cholesterol; ***.p < 0.001 (t-test); ^b – transformed

Results: We were able to analyze several noncholesterol sterols including lathosterol, campesterol, and β-sitosterol. From oxygenated derivatives of cholesterol, various mono-, di-, oxo- and epoxy- derivatives of cholesterol were analysed including C4 (metabolic precursor of bile acids). The method was applied for analysis of human plasma.

Table 2 Concentrations of selected plasma sterol lipids in groups

parameter	control	MetSy
total lathosterol (μmol/L)	5.6 ± 1.9 ^a	8.7 ± 4.3*
total campesterol (μmol/L)	8.8 ± 3.7	9.1 ± 3.1
total β-sitosterol (μmol/L)	6.3 ± 2.4	5.7 ± 1.6
free C4 (ng/mL)	23.7 ± 9.1	25.4 ± 14.4
free 24R/SOH cholesterol (ng/mL) ^b	4.3 ± 2.7	5.2 ± 4.3
free 26OH cholesterol (ng/mL) ^b	9.4 ± 9.7	6.4 ± 6.5
free 7αOH cholesterol (ng/mL) ^b	10.1 ± 4.3	15.0 ± 14.7

^a – mean ± SD; *-p < 0.05 (t-test), transformed for analysis

Conclusions: With the presented method, it is possible to gain deeper insight into metabolism of sterols. The implementation of saponification step enables to analyze total concentrations of sterol lipids. Assessment of oxygenated cholesterol derivatives, including C4, is recommended without saponification step.