

Application

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News

High Performance Liquid Chromatography

Glycerophospholipids Analysis by Comprehensive HPLC Coupled with a Triple Quadrupole Mass Spectrometer

Glycerophospholipids (GPLs) are the major component of biological membranes. They can not only act as a barrier from the external environment, but can also play a key role in a variety of biological processes including membrane trafficking and signal transduction. Thus, analysis of GPLs is one of the most important studies in the metabolomics field. Although reversed phase (RP) HPLC coupled with electrospray ionization (ESI) MS/MS is an effective strategy for lipidomics, there is still room for further improvement of the analytical methods. One drawback to performing determination of GPLs is ion suppression caused by co-eluting compounds. To obtain reliable results, complete separation of target GPLs by comprehensive HPLC with ESI-MS/MS is an effective strategy.

Flow Diagram of Comprehensive HPLC

Fig. 1 shows the flow diagram of the comprehensive HPLC-ESI-MS/MS system. The system comprises 2 flow lines: one for the first dimension separation with a normal phase column and the second dimension separation with a reversed phase column. A mixture of GPLs was roughly classified by normal phase chromatography in the first dimension. All the eluents are trapped into two loops alternatively. Then the entire eluents are introduced into second dimensional reversed phase UHPLC without any risk of sample-loss. The GPLs of interest are separated according to the orthogonal retention selectivity and detected with ESI-MS/MS quantitatively.



Fig. 1 Flow Diagram of the Comprehensive HPLC-ESI-MS/MS System

| Table 1 | Analytical | Conditions |
|---------|------------|------------|
|---------|------------|------------|

| 1D Column | : Nucleosil SIL (150 mm L. × 1.0 mm I.D., 3 μm) |
|------------------------------|--|
| Mobile Phase | : A: Isooctane / Acetone / Ethyl Acetate / Acetic acid = $40/20/20/0.03 (y/y/y/y)$ |
| | B: Isooctane / 2-propanol / Water / Acetic acid / 28 % Ammonia aq.sol. |
| Flowrate | = 40/51/9/0.05/0.05 (V/V/V/V) |
| Time Program Column Temp. | : B Conc. 30 % (0 min) \rightarrow 40 % (25 min) \rightarrow 100 % (40 min) \rightarrow 100 % (55 min) \rightarrow 30 % (55.1 min) \rightarrow STOP (70 min) : 40 °C |
| Injection Vol. | : 5 μL |
| Loop Vol. | : 20 µL |
| 2D Column | : Phenomenex Kinetex C18 (50 mm L. × 4.6 mm I.D., 2.6 μm) |
| Mobile Phase | : A: Methanol / Water / Acetic acid / 28 % Ammonia aq.sol. |
| | = 90/10/0.05/0.05 (WW/W) |
| | = 100/0.05/0.05 (V/V/V) |
| Flowrate | : 3.5 mL/min (50 % split to MS) |
| Time Program | : B Conc. 10 % (0 min) \rightarrow 50 % (0.75 min) \rightarrow 10 % (0.76 min) \rightarrow STOP (1 min) |
| | The initial B Conc. has been changed by a stepwise method |
| Detector | : Shimadzu LCMS-8050 (ESI positive, MRM mode) |

Comprehensive Separation of Glycerophospholipids Comprehensive Separation of GPLs in ESI-positive MRM

mode are shown in Fig. 2.

The GPLs mixture was comprised of 500 ppb each of Phosphatidylglycerol (PG), Phosphatidylethanolamine (PE), Phosphatidylinositol (PI), Phosphatidylserine (PS) and Phosphatidylcholine (PC). The 2D plot of ESI-positive MRM shows the separation of PG, PE, PI, PS and PC. The repeatability (n=5) of retention times and blob areas, which correspond to peak areas in ordinary quantitation and linearity for 50-5000 μ g/L of each 3 PC compounds are is shown in Table 2.

Necessary information for compound-identification is shown in Fig. 2.



Fig. 2 Comprehensive Separation and Calibration Curve of GPLs

| Table 2 D | onostability o | of E Analysos in | % PCD and Linearit | by of EO E000 ug/l | for 2 PC Compounds |
|-----------|----------------|-------------------|---------------------|--------------------|--------------------|
| Iddle Z N | epeatability o | JI 5 Allalyses II | i /0KSD and Lineari | Ly 01 50-5000 μg/L | Tor 5 PC Compounds |

| Compound | MRM transition | Total retention time | Retention time (2D) | Blob Area | Correlation coefficient (R) |
|---------------|--------------------------|----------------------|---------------------|-----------|-----------------------------|
| 16:0-16:0 PC | <i>m/z</i> 734.6 > 184.1 | 0.0072 | 0.9 | 6.8 | 0.999799 |
| 18:0-18:1 PC | <i>m/z</i> 788.6 > 184.1 | 0.013 | 1.1 | 8.9 | 0.999947 |
| 18:0p-18:1 PC | <i>m/z</i> 772.6 > 184.1 | 0.013 | 1.2 | 6.4 | 0.999656 |

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