Separation of an Amino Acid Mixture in HILIC Mode Using a Luna[™] NH₂ Column

Sergio Guazzotti, Roxana Eggleston-Rangel, Namrata Saxena, Jason Anspach, PhD, and Bryan Tackett, PhD Phenomenex, Inc., 411 Madrid Ave., Torrance, CA 90501 USA

Overview

Amino acids are the building blocks of proteins, making them an extremely well researched compound class. With thousands of naturally occurring amino acids available, separating and quantitating a mixture of amino acids can be challenging. While HPLC is a well characterized method for amino acid analysis, LC-MS/MS analysis of amino acids poses several challenges. Some compounds are very polar and are hard to retain by reversed phase HPLC. Due to the polar and hydrophilic nature of amino acids, HILIC chromatography was utilized to separate a 6 amino acid mixture.

In this application note, a Luna 3 μ m NH₂ column, in HILIC mode, and a robust LC-MS/MS method was used to successfully separate an amino acid mixture. Amino groups bound to the silica surface of the stationary phase particles serve as a weak anion exchanger and offer greater polar selectivity, thus increased separation of highly polar amino acids.

Table 1. Amino Acids Used and Their Respective MRM Transitions.

Q1 (m/z)	Q3 (m/z)
117	47
172	126
138	91
216	155
188	142
94	47
	(m/z) 117 172 138 216 188

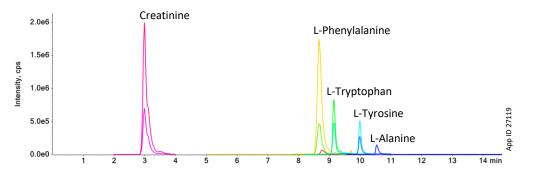
LC Conditions

Column: Luna 3 µm NH₂ Dimensions: 150 x 0.3 mm Part No.: 00F-4377-AC Mobile Phase: A: 10 mM Ammonium Formate in Acetonitrile, pH 3.3 B: 10 mM Ammonium Formate in Water, pH 3.3 Gradient: Time (min) %В 0 3 6 3 11.5 50 12.5 98 18.5 98 19.5 3 32 3 Flow Rate: 10 µL/min Injection: 1 µL Temperature: 40 °C Detector: QTRAP® 6500 System: NanoLC[™] 425 (SCIEX[®]) Detection: MS/MS

MS/MS Conditions

Scan Type:	MRM
Polarity:	Positive
Source:	OptiFlow [®] Turbo V [™]
SG1:	20
SG2:	30
CUR:	30
Temperature:	150 °C
Spray Voltage:	5000 V

Figure 1. Extracted Ion Chromatogram of 6 Amino Acids on a Luna 3 µm NH, Column Under HILIC Conditions.



Have questions or want more details on implementing this method? We would love to help! Visit **www.phenomenex.com/Chat** to get in touch with one of our Technical Specialists

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Japan t: +81 (0) 120-149-262

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Norway t: +47 810 02 005

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t: +64 (0)9-4780951

t: +91 (0)40-3012 2400

indiainfo@phenomenex.com

indoinfo@phenomenex.com

eireinfo@phenomenex.com

italiainfo@phenomenex.com

jpinfo@phenomenex.com

nlinfo@phenomenex.com

nlinfo@phenomenex.com

nzinfo@phenomenex.com

tecnicomx@phenomenex.com

Australia t: +61 (0)2-9428-6444 auinfo@phenomenex.com

Austria t: +43 (0)1-319-1301 anfrage@phenomenex.com

Belaium t: +32 (0)2 503 4015 (French) t: +32 (0)2 511 8666 (Dutch) beinfo@phenomenex.com

Canada t: +1 (800) 543-3681 info@phenomenex.com

China t: +86 400-606-8099 cninfo@phenomenex.com

Czech Republic t: +420 272 017 077 cz-info@phenomenex.com

Denmark t: +45 4824 8048 nordicinfo@phenomenex.com

Finland t: +358 (0)9 4789 0063 nordicinfo@phenomenex.com

France t: +33 (0)1 30 09 21 10 franceinfo@phenomenex.com

Germany t: +49 (0)6021-58830-0 anfrage@phenomenex.com

Hong Kong t: +852 6012 8162 hkinfo@phenomenex.com

www.phenomenex.com

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Sweden t: +46 (0)8 611 6950 nordicinfo@phenomenex.com

Switzerland t: +41 (0)61 692 20 20 swissinfo@phenomenex.com

Taiwan t: +886 (0) 0801-49-1246 twinfo@phenomenex.com

Thailand t: +66 (0) 2 566 0287 thaiinfo@phenomenex.com

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