Customer Application Note

Primary Metabolite Analysis of Plant Material Using a Triple Quadrupole MS Coupled to a Monolith Anion-Exchange Column

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Introduction

A platform using Dionex ion chromatography (IC) coupled to triple quadrupole mass spectrometry (MS) was developed to measure various metabolites in *Brassica napus* seeds. The IC tandem mass spectrometry (IC-MS/MS) technique provides a good tool to identify and quantify metabolites in a complex matrix. The biological pathways include many homologs and structural isomers with the same mass and analog fragmentations, so it is also essential for the analysis to have a good separation of the different compounds. The new lonSwift[™] MAX-100 column (1 × 250 mm) provides good peak shape and separation for a broad spectrum of substances. Comparison with the previous product, the lonPac[®] AS11-HC column (2 × 250 mm), showed a similar separation in a shorter run time and an enhanced sensitivity for several target analytes.

Equipment

Dionex ICS-3000 System including:

- EG Eluent Generator
- DP Dual Pump with Degasser
- AS Autosampler

DC Detector Compartment with Anion Self-Regenerating Suppressor (ASRS®) 300 2 mm Suppressor

API 4000[™] Liquid Chromatography (LC)-MS/MS System (AB SCIEX)

Reagents and Standards

Methanol, chloroform, and other biochemicals were obtained from Sigma-Aldrich Chemie GmbH, (Taufkirchen, Germany) and Roche Diagnostics GmbH (Mannheim, Germany). Water was purified with the Millipore Milli-Q[®] Plus System.

Preparation of Samples

Homogenize approximately 30 mg frozen material at liquid nitrogen temperature in 2 mL microcentrifuge tubes (Eppendorf, Hamburg, Germany) containing two steel beads (ASK, Korntal-Muenchingen, Germany) by grinding in a TissueLyser (Quiagen GmbH, Hilden, Germany) for 45 s at 1800 strokes.¹ After homogenization, extract the samples with 0.5 mL 1:1 (v/v) methanol/chloroform under the same grinding conditions. Extract water-soluble compounds from the CHCl₃ phase by adding 300 μ L water, followed by 45 s shaking, and centrifugation at 2000 g for 2 min. Filter the upper aqueous/ methanol phase with a Vivaclear centrifugal filter (0.8 μ m pore size, SatoriusStedim Biotech, Göttingen, Germany) at 2000 g for 2 min. Re-extract the lower CHCl₃ phase with 200 μ L water and add the aqueous/methanol phase to the first phase. Evaporate the extract to dryness using a centrifugal vacuum dryer at 20 °C and redissolve in 1 mL water.





Chromatographic Conditions

| IonSwift MAX-100, 1 × 250 mm | | | | | |
|------------------------------|------------------------------------------------------|--|--|--|--|
| IonPac AS11-HC, 2 × 250 mm | | | | | |
| Flow Rate: | 150 μL/min | | | | |
| Inj. Volume: | 10 μ L (cleaning, 10 μ L CH ₃ OH) | | | | |
| Column Temp.: | 40 °C | | | | |

| Table 1. KOH Gradient for the Comparison of MAX-100 and AS11-HC Columns | | | | | | |
|---------------------------------------------------------------------------|-------------|---------------|-------------|---------------|--------------|--|
| MAX | -100 | AS11-HC | | Cleani | Cleaning Run | |
| Time (min) | KOH (mM) | Time (min) | KOH (mM) | Time (min) | KOH (mM) | |
| 0 | 5 | 0 | 5 | 0 | 100 | |
| 10 | 5 | 16 | 5 | 9 | 100 | |
| 16 | 12 | 26 | 12 | 9.5 | 5 | |
| 28 | 25 | 41 | 25 | 16 | 5 | |
| 32 | 100 | 48 | 100 | | | |
| 38 | 100 | 67 | 100 | | | |

Electrospray Ionization (ESI)-MS/MS Conditions

Measure the samples in negative mode. Use nitrogen as a curtain gas, nebulizer gas, heater gas, and collision gas. Set the ion spray voltage to -4000 V, the capillary temperature at 450 °C, and the MS parameters as displayed in Table 2.

DP=Declustering Potential, EP=Entrance Potential, CE=Collision Energy, CXP=Collision Cell Exit Potential. *Not the optimum potential to reduce sensitivity due to the high concentrations in seeds.

| Table 2. F MS/MS | Iltered Transi | Masses tions in | s and P Negat | 'otentia ive Mod | ls of Je | |
|------------------------------|-------------------|--------------------|------------------|---------------------|-------------|-----|
| | Q1 Mass | Q3 Mass | DP | EP | CE | CXP |
| Glycolate | 74.8 | 46.9 | -35 | -10 | -12 | -7 |
| Lactate | 88.9 | 42.9 | -45 | -10 | -18 | -1 |
| Succinate semialdehyde | 100.8 | 56.9 | -45 | -10 | -14 | -1 |
| Pyruvate | 86.9 | 86.9 | -30 | -10 | -14 | -3 |
| Trehalose-6-phosphate | 420.9 | 78.9 | -85 | -10 | -60 | -5 |
| Glucose-1-phosphate | 258.9 | 79.0 | -60 | -10 | -42 | -5 |
| Succinate | 116.8 | 73.1 | -45 | -10 | -16 | -1 |
| Malate* | 132.8 | 70.9 | -80 | -10 | -22 | -3 |
| Glucose-6-phosphate | 258.8 | 96.8 | -50 | -10 | -22 | -5 |
| Glucose-6-phosphate | 258.8 | 198.8 | -50 | -10 | -16 | -19 |
| 2-Oxoglutarate | 144.9 | 100.7 | -55 | -10 | -12 | -17 |
| Fructose-6-phosphate | 258.8 | 78.9 | -55 | -10 | -60 | -5 |
| Fructose-6-phosphate | 258.8 | 70.9 | -50 | -10 | -44 | -3 |
| Fumarate | 114.9 | 70.9 | -40 | -10 | -12 | -11 |
| AMP | 345.9 | 78.8 | -75 | -10 | -66 | -5 |
| NADH | 331.4 | 78.8 | -40 | -10 | -72 | -3 |
| NADH | 331.4 | 133.8 | -40 | -10 | -28 | -11 |
| Oxaloacetate | 130.8 | 86.9 | -25 | -10 | -8 | -5 |
| 6-Phospho-gluconate | 274.9 | 96.9 | -45 | -10 | -28 | -5 |
| 3-PGA | 184.8 | 96.9 | -35 | -10 | -20 | -15 |
| Citrate* | 190.8 | 87.0 | -75 | -10 | -24 | -5 |
| Isocitrate | 190.8 | 72.9 | -20 | -10 | -30 | -5 |
| Phosphoenolpyruvate | 166.7 | 78.9 | -35 | -10 | -14 | -5 |
| ADP-Glucose | 293.4 | 133.9 | -30 | -10 | -34 | -1 |
| ADP-Glucose | 293.4 | 452.8 | -30 | -10 | -12 | -13 |
| UDP-Glucose | 281.9 | 111.0 | -30 | -10 | -22 | -7 |
| UDP-Glucose | 281.9 | 240.8 | -30 | -10 | -20 | -11 |
| Glycerone-phosphate | 168.7 | 78.7 | -40 | -10 | -36 | -3 |
| <i>cis</i> -aconitate | 172.8 | 84.8 | -35 | -10 | -18 | -15 |
| <i>trans</i> -aconitate | 172.8 | 84.8 | -35 | -10 | -18 | -15 |
| ADP | 425.8 | 134.0 | -75 | -10 | -30 | -9 |
| ADP | 425.9 | 79.0 | -75 | -10 | -90 | -5 |
| Pvrophosphate | 177.0 | 79.0 | -35 | -10 | -36 | -5 |
| Fructose-1,6- biphosphate | 338.9 | 240.9 | -50 | -10 | -20 | -15 |
| NADPH | 371.4 | 78.8 | -45 | -10 | -68 | -10 |
| NADPH | 371.4 | 133.8 | -45 | -10 | -30 | -6 |
| UDP | 200.9 | 78.9 | -35 | -10 | -14 | -7 |
| UDP | 200.9 | 323.0 | -35 | -10 | -6 | -9 |
| ATP | 505.9 | 158.7 | -80 | -10 | -40 | -29 |
| UTP | 402.9 | 78.9 | -60 | -10 | -100 | -5 |
| UTP | 240.9 | 403.1 | -30 | -10 | -10 | -27 |



Results and Discussion

In most cases, the IonSwift MAX-100 column showed better sensitivity than the IonPac AS11-HC column (Table 3). The biggest difference between the two columns was seen in pyrophosphate, which had a great signal-to-noise (S/N) ratio on the MAX-100 column. This was significant at low concentration, at which the S/N ratio on the AS11-HC column was below 3. The run time on the AS11-HC column was 30 min longer with the same flow rate. The small inner diameter of the ESI capillary leads to a backpressure of 100 to 110 psi between the 2 mm suppressor and the capillary at a flow rate of 150 µL/min. This value is approximately 75% of the suppressor limit and prevents higher flow rates. For a shorter run time with the AS11-HC column coupled to an ESI-MS/MS, a 4 mm suppressor is recommended.

Oxaloacetate, glycerone-phosphate, and ADP-glucose are not shown in Table 3 due to their instability under IC conditions. Compared to most of the other target analytes with high sensitivity and a low limit of detection, UDP-glucose has a higher limit of detection, possibly due to degradation under IC conditions, even if the calibration showed a good regression factor.

The memory effect of not completely eluted compounds described in the S. Arrivault et al. 2009 study¹ was observed for eight late-eluting compounds (Cit/ICit/c-Aco/t-Aco/ADP/PP/UDP/UTP). A wash run injection of methanol after each sample or standard reduced the problem either completely or to less than 1% (not calculated on the AS11-HC column).

Glucose-6-phosphate (G6P) and Fructose-6-phosphate (F6P) had a retention time gap of 1 min. The high ratio of G6P/F6P in plant material resulted in problems of peak separation. For quantification of G6P and F6P, less intense transitions ($259 \rightarrow 97/79$) were chosen because, for G6P, the $259 \rightarrow 199$ transition was nearly independent of F6P. The $259 \rightarrow 71$ transition for F6P had a slightly higher abundance, compared to that for G6P. Nevertheless, F6P was the only peak area that had to be integrated manually.

For the isomers citrate/isocitrate, the transitions $191 \rightarrow 87$ and $191 \rightarrow 73$ were used because they were absolutely independent of each other. The declustering potential of the major compounds citrate (DP -35 reduced to -75) and malate (DP -40 reduced to -80) were decreased relative to their optimum so that the sensitivity was selectively decreased, thus preventing an overload of the detector and allowing the measurement of the entire set of compounds in one dilution step. This technique allows analysis of concentrations of different compounds over 4 to 5 orders of magnitude in one run (see figures).

Table 3. Comparison of Signal-to-Noise Ratios on MAX-100 and AS11-HC Columns with a Standard Mix that Reflects

| the Composition of Plant Samples | | | | | | | |
|----------------------------------|----------------|------------------------------------|-----------------|----------------|--------|--|--|
| Analyte Peak Name | RT IonSwift | Analyte Concentration (pmol) | S/N IonSwift | S/N As11-hc | r | | |
| Glycolate | 2.1 | 0.6 | 1.5 | — | 0.9992 | | |
| Lactate | 2.1 | 2.4 | 15 | 3 | 0.9981 | | |
| Succinate semialdehyde | 2.4 | 0.3 | 3 | — | 0.9972 | | |
| Pyruvate | 2.5 | 0.8 | 7 | 4 | 0.9970 | | |
| Trehalose-6-phosphate | 5.5 | 0.012 | 5 | — | 0.9947 | | |
| Glucose-1-phosphate | 6.2 | 0.36 | 70 | 10 | 0.9970 | | |
| Succinate | 10.2 | 4 | 50 | 40 | 0.9900 | | |
| Malate* | 10.2 | 16 | 100 | 30 | 0.9993 | | |
| Glucose-6-phosphate | 15.1 | 2.4 | 10 | 10 | 0.9945 | | |
| Glucose-6-phosphate | 15.1 | 2.4 | 20 | 7 | 0.9949 | | |
| 2-Oxoglutarate | 16.1 | 1.2 | 13 | 15 | 0.9950 | | |
| Fructose-6-phosphate | 16.2 | 0.28 | 4 | 3 | 0.9856 | | |
| Fructose-6-phosphate | 16.2 | 0.28 | 3 | — | 0.9862 | | |
| Fumerate | 17.6 | 0.32 | 3 | _ | 0.9939 | | |
| AMP | 18.1 | 0.056 | 3 | — | 0.9988 | | |
| NADH | 19.7 | 0.008 | 2 | | 0.9994 | | |
| 6-Phospho-gluconate | 25.2 | 0.032 | 6 | 8 | 0.9960 | | |
| 3-PGA | 27.4 | 0.24 | 11 | 10 | 0.9955 | | |
| Citrate* | 28.8 | 20 | 110 | 150 | 0.9973 | | |
| Isocitrate | 29,9 | 0.24 | 12 | 8 | 0.9954 | | |
| UDP-Glucose | 30.7 | 3 | 2.5 | | 0.9930 | | |
| Phosphoenolpyruvate | 31.0 | 0.1 | 28 | 30 | 0.9952 | | |
| <i>cis</i> -Aconitate | 30.9 | 0.2 | 25 | 13 | 0.9952 | | |
| trans-Aconitate | 32.5 | 0.12 | 14 | 11 | 0.9932 | | |
| ADP | 32.7 | 0.08 | 7 | 5 | 0.9957 | | |
| Pyrophosphate | 33.3 | 0.004 | 20 | 2 | 0.9935 | | |
| Fructose-1,6- biphosphate | 33.4 | 0.012 | 3 | 2 | 0.9932 | | |
| NADPH | 33.8 | 0.012 | 5 | | 0.9954 | | |
| UDP | 34.0 | 0.016 | 8 | 4 | 0.9960 | | |
| ATP | 34.2 | 0.04 | 3 | | 0.9962 | | |
| UTP | 34.8 | 0.012 | 8 | 2 | 0.9934 | | |

*Citrate and malate have poorer limits of detection than are possible due to a DP that was selected to reduce sensistivity due to the high amounts of these compounds in the sample.









*Malate's limit of detection can be lower with an optimal DP.



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*Citrate's limit of detection can be lower with an optimal DP.







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Conclusion

The combination of the 1 mm lonSwift MAX-100 column and the API 4000 ESI-MS/MS system provides a simple, rapid, specific, and sensitive method for the simultaneous detection and quantification of several primary metabolites involved in the TCA cycle and glycolysis pathway. Compared to the previously used lonPac AS11-HC column, the MAX-100 has shorter run times and helps achieve better sensitivity.

Reference

 Arrivault, S.; Guenther, M.; Ivakov, A.; Feil, R.; Vosloh, D.; van Dongen, J.T.; Sulpice, R.; Stitt, M. The Plant Journal. *Plant Physiol.* 2009, *59*, 824–839.

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