

# Application Data Sheet

### No.3

#### **GCMS**

Gas Chromatograph Mass Spectrometer

## Analysis of Amino Acids Contained in a Grain of Rice

: 280°C

Amino acids contained in a grain of rice were treated with EZ:faastTM (Phenomenex, Inc.), which enables easy pretreatment, and then analyzed with a GC-MS system.

#### **Experiment**

#### **Pretreatment**

A grain of rice was treated with EZ:faast. Norvaline was added as an internal standard.

#### Instrument

A GCMS-QP2010 Ultra (with high-power oven) was used for the measurements. Analysis conditions, shown in Table 1, were in conformity with the "Amino Acid Method" in the "GC/MS Metabolic Components Database."

Table 1: Analysis Conditions (GC/MS Metabolic Components Database: Amino Acid Analysis Methods)

GC-MS : GCMS-QP2010 Ultra (with high-power oven)

Column : ZB-AAA (10 mL. × 0.25 mml.D.) (Phenomenex, Inc.) [MS]

[GC] Interface temperature

Injection quantity : 1  $\mu$  L Ion source temperature : 200°C

Vaporization chamber temperature : 280°C Solvent elution time : 0.4 min

Column oven temperature : 110°C -> (30 °C/min) -> 320°C Data sampling time : 0.5 min to 7 min

Control mode : Constant pressure (15 kPa) Measurement mode : Scan

Injection mode: SplitMass range: m/z 45-450 (3,333u/sec)Split ratio: 15Event time: 0.15 secCarrier gas: Helium

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Fig. 1: Total Ion Current Chromatogram for Amino Acid Derivatives in Katsuobushi Extract

The numbers for each component follow the serial numbers in the "GC/MS Metabolic Components Database."

1 Alanine 8 Leucine 16 Asparatic acid 24 Ornithine 28 Hydroxylysine 3 Glycine 10 Isoleucine 17 Methionine 25 Glycine-proline (2 isomers) 4 alpha-aminobutyric acid 11 Threonine 18 4-Hydroxyproline (dipeptide) 29 Tyrosine 5 Valine 12 Serine 19 Glutamic acid 26 Lysine 31 Tryptophan 7 Norvaline (I.S.) 13 Proline 20 Phenylalanine 27 Histidine

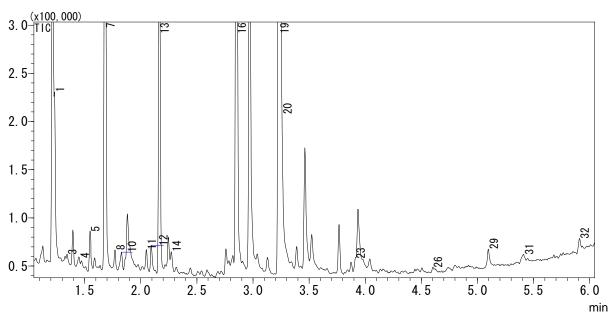


Fig. 2: Total Ion Current Chromatogram for Amino Acid Derivatives in Kombu Extract The numbers for each component follow the serial numbers in the "GC/MS Metabolic Components Database."

1 Alanine	8 Leucine	14 Asparagine	26 Lysine
3 Glycine	10 Isoleucine	16 Aspartic acid	29 Tyrosine
4 alpha-aminobutyric acid	11 Threonine	19 Glutamic acid	31 Tryptophan
5 Valine	12 Serine	20 Phenylalanine	32 Cystathionine
7 Norvaline(I.S.)	13 Proline	23 Glutamine	

#### Summary

Pretreatment using the EZ:faast kit, followed by analysis using the GCMS-QP2010 Ultra, which is equipped with a high-speed scanning function, enabled rapid analysis of amino acids. With this combination, it took only 15 minutes per sample from pretreatment to analysis.

(Reference: Shimadzu Application News No. M246 Analysis of Amino Acids Using Fast-GC/MS and Metabolite Database)

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