

Establishing a new quantitative method for *Fusarium* hydrophobins using LC/MS/MS

(Kumiko INOMOTO, Norio DOI, Susumu MASUDA, Masayuki AIZAWA / Asahi Breweries, Ltd., Japan)

[Introduction]

Hydrophobin (HFB), an amphiphilic low-molecular protein produced by filamentous fungi that infect barley and malt, is main cause of beer primary gushing. HFB produced by *Fusarium* fungi are especially known for their tendency to induce gushing. It has been reported that, ~3 µg/L of such HFB in beer can cause gushing¹⁾. Previously, the ELISA method using polyclonal antibody was reported¹⁾. In order to evaluate HFB content more accurately, highly specific quantification of HFB is required. Using UPLC-triple quadrupole tandem mass spectrometry (LC/MS/MS), we established a highly specific and sensitive analysis that quantified the target HFB produced by *Fusarium graminearum*.

[Method]

< HFB analysis using LC/MS/MS >

[The feature]

Since quantification is based on the mass of the peptide fragments specific to the target protein, highly sensitive and specific quantification is possible.

1. Selection of target fragments

Target protein

We chose HFB produced by *Fusarium graminearum*.

<Because>

A typical species that infects barley has strong ability to induce gushing.



Based on the amino acid sequence of the target protein, the peptide fragments with a specific and strong signal were selected.

[Peptide Selection Criteria²⁾]

- Length between 6 and 16 amino acids
- No posttranslational modifications
- No single nucleotide polymorphism
- Containing one of leucine, isoleucine, valine, alanine or proline residue. etc.

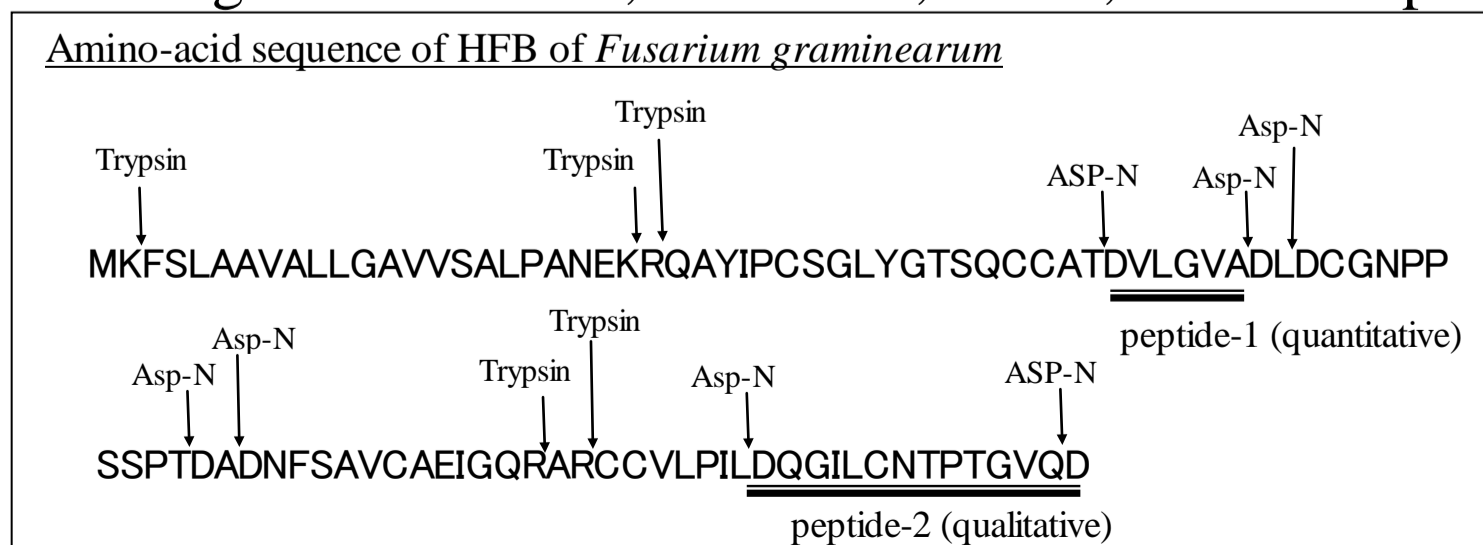


Figure 1 : The estimated cleavage site using Trypsin + Asp-N

2. LC/MS/MS analysis

① Sample preparation

[Beer]

Sample 20 µL

← Add I.S.
(Stable isotope labeled-peptide)

← Dithiothreitol

30 min (56°C)

← Iodoacetamide

30 min

← Trypsin + Asp-N

24 h (37°C)

↓

Clean up (SPE C18)

↓

LC/MS/MS analysis

[Malt]

5 g of the fine flour

↓ ← 60% Ethanol, 50 mL

Extraction (30°C, 200 rpm, over night)

↓

Centrifugation

↓ (Supernatant fraction)

The same method of beer

② LC/MS/MS conditions

[UPLC analytical conditions]

LC : SHIMADZU Nexera X2

Column : L-column2 ODS (2.1 mm I.D. x 50 mm, 2 µm, CERI)

Mobile phase : A : 0.1% Formic acid B : Acetonitrile

Flow rate : 0.3 mL/min

Gradient min (A/B) : 0 (95/5), 30 (50/50), 32 (5/95), 35 (95/5)

injection : 20 µL

[MS analytical conditions]

MS : Sciex API 4000

Ionization : ESI Positive

Table 1 : MS analytical conditions

	Precursor ion (m/z)	Product ion (m/z)	DP (V)*	CE (V)*	CXP (V)*
Peptide-1 (Quantitative)	573.4	484.5	81	23	20
		385.2		29	14
Peptide-2 (Qualitative)	759.8	243.9	96	47	6
		301.2		47	10

*) DP (Declustering potential), CE (Collision energy), CXP (Collision cell exit potential)

< Gushing Test >

We prepared a microbrew beer based on the method proposed by Vaag *et al.*³⁾

and conducted a gushing test according to method-A⁴⁾ proposed by Amaha *et al.*

[Result]

1. Validation of the method for quantitative determination of peptide-1

Table 2: Validation of beer analysis

Linearity	Detection limit* (µg/L)	Quantitation limit* (µg/L)	Repeatability (%)	Interday variation (%)
0.9998	0.8	2.0	4.8	5.1

Table 3: Validation of malt analysis

Linearity	Detection limit* (µg/g malt)	Quantitation limit* (µg/g malt)	Repeatability (%)	Interday variation (%)
0.9990	0.05	0.12	8.0	11.5

*) Converted to HFB concentration

< Repeatability : n=6 Interday variation : n=6, 3days >

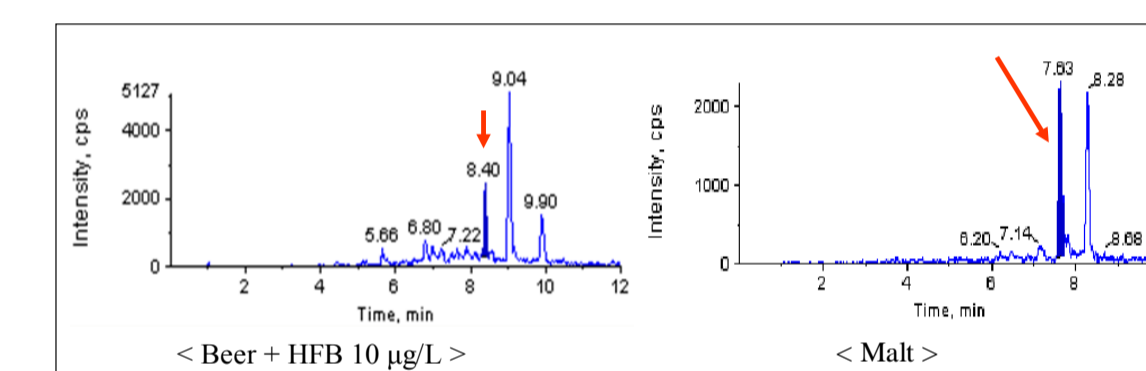


Figure 2: LC/MS/MS chromatogram

- The analytical sensitivity and precision of peptide-1 were satisfactory for both beer and malt.

✧ The HFB-producing species detected by this method

Table 4: The HFB-producing species estimated by the combination of peptide-1 and peptide-2

	The HFB-producing species
Detection of only peptide-1	<i>Trichoderma reesei</i> , <i>Trichoderma virens</i> , <i>Neurospora crassa</i> , <i>Claviceps fusiformis</i> , <i>Cryphonectria parasitica</i> etc.
Detection of only peptide-2	<i>F. poae</i> , <i>F. oxysporum</i> *
Detection of both peptide-1 and peptide-2	<i>F. graminearum</i> , <i>F. culmorum</i> , <i>F. verticillioides</i> , <i>F. fujikuroi</i> , <i>F. pseudograminearum</i> , <i>F. oxysporum</i> **

* HFB Name : FOQG_13743

** HFB Name : FOVG_15013, FOXB_00066, FOWG_12166, Trihydrophobin

Table 5: Typical species that causes beer gushing not detected by this method

The HFB-producing species
<i>F. avenaceum</i> , <i>Nigrospora</i> sp., <i>Penicillium</i> sp., <i>Aspergillus</i> sp. etc.

BLAST search results for the sequences of species registered in the NCBI database.

Parts in the red : Producing species that strongly induces gushing in beer ¹⁾⁵⁾.

- Detection of peptide-1 indicates the possibility of molds such as *Fusarium* and *Trichoderma*.
- Detection of both peptide-1 and peptide-2 allows for the estimation of six species belonging to the genus *Fusarium*.
- Detection of only peptide-2 allows for the estimation of two species belonging to the genus *Fusarium*.

2. The results of HFB analysis of malt sample

Table 6: The results of HFB analysis of malt sample

Sample No.	Area	Gushing Test (mL/200 mL)	Peptide-1 HFB (µg/g malt)	Peptide-2 Detection
Sample-1	Europe	101	0.57	○
Sample-2		45	trace	○
Sample-3		40	0.97	○
Sample-4		20	0.43	○
Sample-5		11	0.20	
Sample-6		7	N.D.	
Sample-7		7	0.20	
Sample-8		3	0.39	○
Sample-9		1	0.52	○
Sample-10		0	0.14	
Sample-11		0	N.D.	
Sample-12		0	N.D.	
Sample-13		0	trace	
Sample-14	North America	96	N.D.	
Sample-15		71	0.24	
Sample-16		19	N.D.	
Sample-17		0	N.D.	

N.D. : Lower than the detection limit (0.05 µg/g malt)

trace : Lower than the quantitation limit (0.12 µg/g malt)

○ : Detection of peptide-2

- Presence of target HFB was confirmed through detection of peptide-1 and peptide-2 in the malt samples.
- European malts with peptide-1 tended to exhibit gushing. Some types of North American malts exhibited gushing in the absence of target HFB.
- By addition of other target HFBs, this method will be able to detect a larger variety of HFBs produced by fungi that induce gushing.

[Conclusion]

1. Using LC/MS/MS, we established a highly specific and sensitive analysis that quantified the target HFB produced by *Fusarium graminearum*.
2. Measurement of peptide-1 and peptide-2 enabled the detection of HFB-producing species.
3. European malts with peptide-1 tended to exhibit gushing.
4. This method allowed for the quantitative analysis of HFB in beer and malt, and the analytical results aided in the estimation of the HFB-producing species. This method can be used to improve the quality of beer and malt associated with primary gushing.

- [References]
- (1) Sarlin, T., *et al.*; *J. Inst. Brew.*, 111(2), 105–111, 2005
 - (2) Kamiie, I., *et al.*; *Pharm. Res.*, 25, 1469–1483, 2008
 - (3) Vaag, P., *et al.*; *EBC Congress*, 155–162, 1993
 - (4) Amaha, M., *et al.*; *Tech. Quart. MBAA*, 15, 15–22, 1978
 - (5) Sarlin, T.; *VTT Science*, 13, 2012