

File No. 11023/53/2018-QA
Food Safety and Standards Authority of India
(A statutory Authority established under the Food Safety and Standards Act, 2006)
(Quality Assurance Division)
FDA Bhawan, Kotla Road, New Delhi - 110002

Dated, the 29th June, 2020


ORDER

Subject: Method for the estimation of $\Delta\delta^{13}\text{C}_{\text{fru-glu}}$, $\Delta\delta^{13}\text{C}_{\text{max}}$, and Foreign Oligosaccharides in Honey by Elemental Analysis (EA)/Liquid Chromatography (LC) - Isotopic Ratio Mass Spectrometry (EA/LC-IRMS) - reg.

The Scientific Panel on Methods of Sampling and Analysis and the Competent Authority, FSSAI has approved the Method for the estimation of $\Delta\delta^{13}\text{C}_{\text{fru-glu}}$, $\Delta\delta^{13}\text{C}_{\text{max}}$, and Foreign Oligosaccharides in Honey by Elemental Analysis (EA)/Liquid Chromatography (LC) - Isotopic Ratio Mass Spectrometry (EA/LC-IRMS) [Annexure - I].

2. The food testing laboratories are hereby requested to use the aforesaid method, with immediate effect.

Encl: Method

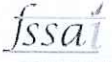

29.6.2020
(Kumar Anil)
Advisor (QA)

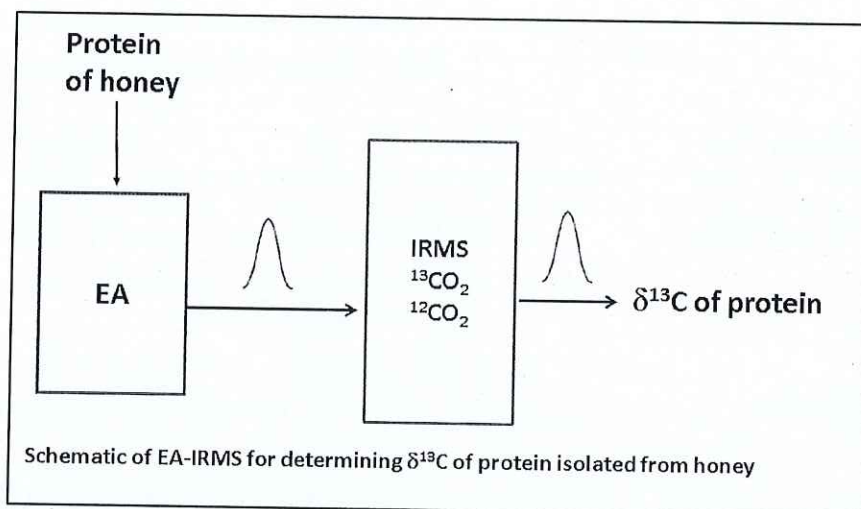
To:

1. All FSSAI Notified Laboratories
2. All State Food Testing Laboratories

Copy to:

1. Executive Director (Regulatory Compliance), FSSAI
2. Advisor (Standards), FSSAI
3. Head (Regulations), FSSAI

 <p>FOOD SAFETY AND STANDARDS AUTHORITY OF INDIA Empowering Trust, Assuring Safe & Nutritious Food Ministry of Health and Family Welfare, Government of India</p>	Method for the estimation of $\Delta\delta^{13}\text{C}_{\text{fru-glu}}$, $\Delta\delta^{13}\text{C}_{\text{max}}$, and Foreign Oligosaccharides in Honey by Elemental Analysis(EA)/Liquid Chromatography(LC)-Isotopic Ratio Mass Spectrometry (EA/LC-IRMS)		
Method No.	02	Revision No. & Date	29.06.2020
Abbreviations	IRMS: Isotope Ratio Mass Spectrometer LC: Liquid Chromatography EA: Elemental analyzer CEI: Compact Electron Ionization		
Caution	<ol style="list-style-type: none"> 1. Always wear gloves and mask while doing sample analysis and reference material handling 2. Keep the eluent and reagent bottles under constant He purge to prevent CO_2 contamination from ambient air. 3. Phosphoric acid and Sulphuric acid are highly corrosive. 4. Prepare the oxidation reagents fresh daily, store in dark brown bottle 5. Many of the routine gases for IRMS are hazardous. The laboratory should have an atmospheric monitoring system to warn of dangerous levels of gases. 		
Principle	<p>The method involves the determination of the relative isotopic ratios ($\delta^{13}\text{C}$) of 1) protein isolated from honey by EA-IRMS and 2) the $\delta^{13}\text{C}$ values of every individual sugar present in honey within a single HPLC run by LC-IRMS.</p> <p>Isotopic ratios are measured relative to a working reference gas calibrated using internationally accepted standards and are reported using the delta notation (δ) and expressed as 'per mill (‰)'. The delta notation is defined as</p> <div style="border: 1px solid black; padding: 5px; width: fit-content; margin: 10px auto;"> $\delta^{13}\text{C} (\text{‰}) \text{ sample} = [R(\text{sample}) / R(\text{standard}) - 1] \times 1000$ </div> <p>where R represents the ratio $^{13}\text{CO}_2/^{12}\text{CO}_2$. The $^{13}\text{C}/^{12}\text{C}$ carbon isotope ratios reported as $\delta^{13}\text{C}$ values are related to Vienna Pee Dee Belemnite (VPDB) according to the AOAC Official Method 998.12 δ is the $^{13}\text{C}/^{12}\text{C}$ ratio of the sample related to the $^{13}\text{C}/^{12}\text{C}$ ratio of a reference material to ensure international compatibility of data sets. The unit of expression is, per mill (‰).</p> <p>The CO_2 produced from combustion of the protein fraction isolated from the sample is analyzed to give $\delta^{13}\text{C}_{\text{Protein}}\%$ by IRMS. The schematic for EA-IRMS is shown below</p>		

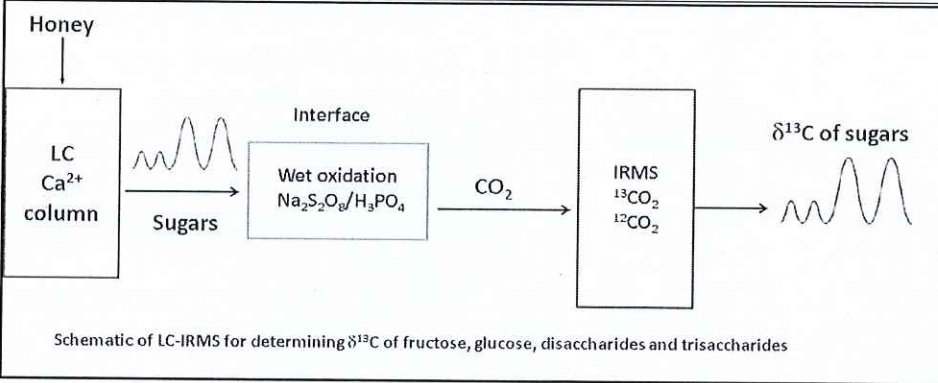


The $\delta^{13}\text{C}$ ‰ values of fructose, glucose, disaccharides, and trisaccharide and any other oligosaccharides present in honey are determined by LC-IRMS. The sugars are separated by LC using a cation exchange column. All individual sugars eluting from the LC column pass into the LC/IRMS interface. Here the carbon from organic samples in the mobile phase is converted into CO_2 by a wet chemical oxidation process using sodium peroxodisulfate either in the presence or absence of phosphoric acid. CO_2 and O_2 both diffuse through, which are subsequently dried in an online gas drying unit. The individual CO_2 peaks are subsequently admitted to the IRMS, which directly gives the $\delta^{13}\text{C}$ values for each individual sugar; $\delta^{13}\text{C}_{\text{fru}}$ ‰, $\delta^{13}\text{C}_{\text{glu}}$ ‰, $\delta^{13}\text{C}_{\text{disaccharide}}$ ‰, $\delta^{13}\text{C}_{\text{trisaccharide}}$ ‰ and $\delta^{13}\text{C}$ ‰ of any other oligosaccharides (see chromatograms below). The schematic of a typical LC-IRMS is shown below.

The difference in the carbon isotope ratio between $\delta^{13}\text{C}_{\text{fru}}$ ‰ and $\delta^{13}\text{C}_{\text{glu}}$ ‰ gives $\Delta\delta^{13}\text{C}_{\text{fru-glu}}$ ‰.

$\Delta\delta^{13}\text{C}_{\text{max}}$ is the maximum difference observed between all possible isotopic ratios measured ($\Delta\delta^{13}\text{C}_{\text{fru-disaccharides}} / \Delta\delta^{13}\text{C}_{\text{fru-trisaccharides}} / \Delta\delta^{13}\text{C}_{\text{fru-protein}} / \Delta\delta^{13}\text{C}_{\text{glu-disaccharides}} / \Delta\delta^{13}\text{C}_{\text{glu-trisaccharides}} / \Delta\delta^{13}\text{C}_{\text{glu-protein}} / \Delta\delta^{13}\text{C}_{\text{disaccharides-trisaccharides}} / \Delta\delta^{13}\text{C}_{\text{disaccharides-protein}} / \Delta\delta^{13}\text{C}_{\text{trisaccharides-protein}}$)

The peak area (%) for foreign oligosaccharides is calculated from the areas appended in the LC chromatogram.

	 <p style="text-align: center;">Schematic of LC-IRMS for determining $\delta^{13}\text{C}$ of fructose, glucose, disaccharides and trisaccharides</p>
<p>Apparatus</p>	<ol style="list-style-type: none"> 1. An integrated EA-IRMS instrument equipped with an automated combustion system and mass spectrometer designed or modified for isotope ratio measurement at natural abundance 2. An integrated LC-IRMS comprising of a HPLC/UPLC and in line oxidation reactor for aqueous oxidation of LC elute and a mass spectrometer designed or modified for isotope ratio measurement at natural abundance 3. LC comprises of a binary pump, autosampler, column oven (set at 80°C), and cation exchange column (Ca^{2+}, 300 × 7.7 mm, 8μm or equivalent) 4. Analytical microbalance: 0.0001g 5. Micropipette: 10 – 100 μL, 20 – 200 μL and 100 – 1000 μL 6. Volumetric flasks: 10 mL Class A 7. Vortex mixer 8. Sonicator 9. Centrifuge (capable of 10,000 × g) 10. Water bath (80 °C) 11. Convection Oven 12. Centrifuge tubes (50mL, 15mL) 13. Spatula 14. Forceps (blunt end and pointed curved end) 15. Tin capsules 16. Capsule holding tray 17. Nylon stocking material (100-150 mesh) 18. Syringe filters (0.45μm and 0.22 μm) 19. Vacuum concentrator
<p>Chemicals</p>	<ol style="list-style-type: none"> 1. Ultra-pure water (Electrical Resistivity, Min., 18.18 MΩ cm, at 25°C) 2. Phosphoric acid (H_3PO_4) (puriss. P.a. \geq 99%), crystalline 3. Sodium peroxodisulfate ($\text{Na}_2\text{S}_2\text{O}_8$, Sodium persulfate) (purum p.a. \geq 99%) 4. Sodium tungstate dihydrate ($\text{Na}_2\text{WO}_4 \cdot 2\text{H}_2\text{O}$) (puriss. p.a. \geq 99%) 5. Sulfuric acid (p.a. 98%)

	6. CO ₂ (working standard reference gas): 99.999% Pure 7. O ₂ (flash combustion gas): >99.999% Pure 8. Helium: 99.999% pure																		
Reference standards	<p>Stable isotope reference standards: Any of these certified reference standards defined below can be used as secondary working standards. These are not recommended for daily use as the commercial availability are restricted.</p> <table border="1"> <thead> <tr> <th>Standard</th> <th>$\delta^{13}\text{C}$ (‰)</th> </tr> </thead> <tbody> <tr> <td>Sucrose</td> <td>-10.449</td> </tr> <tr> <td>Casein</td> <td>-26.98</td> </tr> <tr> <td>NBS 22 Oil</td> <td>-30.031</td> </tr> <tr> <td>Beet sugar</td> <td>-26.027</td> </tr> <tr> <td>Galactose</td> <td>-21.415</td> </tr> <tr> <td>Fructose</td> <td>-10.985</td> </tr> <tr> <td>Glucose</td> <td>-10.97</td> </tr> <tr> <td>Cane sugar</td> <td>-11.64</td> </tr> </tbody> </table> <p>The following sugars are used as in-house standards for normalization and verified against any of the listed secondary standards.</p> <ol style="list-style-type: none"> 1. D- (-)-fructose $\geq 99\%$ pure 2. D-(+)-glucose monohydrate $\geq 99.5\%$ pure 3. D-(+)-sucrose $\geq 99\%$ pure 4. D-(+)-maltose monohydrate $\geq 99\%$ pure 5. D-(+)-Raffinose pentahydrate $\geq 99\%$ pure 	Standard	$\delta^{13}\text{C}$ (‰)	Sucrose	-10.449	Casein	-26.98	NBS 22 Oil	-30.031	Beet sugar	-26.027	Galactose	-21.415	Fructose	-10.985	Glucose	-10.97	Cane sugar	-11.64
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Preparation of reagents	<p><i>Reagents for protein isolation</i></p> <ol style="list-style-type: none"> 1. 10% aqueous solution of Sodium tungstate: Dissolve 10 g of Na₂WO₄·2H₂O in 100 mL of pure water. Prepare fresh daily 2. 0.335 M H₂SO₄: Dilute 1.88 mL concentrated H₂SO₄ to 100 mL with ultra-pure water <p><i>Chemical oxidation reagents</i></p> <ol style="list-style-type: none"> 1. 20% Sodium peroxodisulfate: Dissolve 200 g Sodium peroxodisulfate in 1000 mL ultra-pure water in a brown glass bottle using an ultrasonic bath. Use a water-jet pump for vacuum degassing to remove all dissolved CO₂. 2. 1.5 M H₃PO₄ in water: Weigh 147.0 g of crystalline H₃P0₄. Dissolve in ~250 mL of ultra-pure water and make up to 1 L with water. <p><i>LC reagents</i></p> <p>Ultra-pure water: (Electrical Resistivity, Min.18.18 MΩ cm, at 25°C)</p>																		
Preparation of standards for EA IRMS	<ol style="list-style-type: none"> 1. Weigh protein standard (Casein), approximately between 0.1-0.2 mg, with the help of spatula in the tin capsule. 2. Fold the tin capsule with the help of the forceps in such a way so as to remove air. 																		

	<p>3. Gently fold it from all the sides and place the folded tin capsule in the carousel and start the sequence of operation following the manufacturer's instruction</p>
<p>Sample preparation for EA-IRMS</p>	<p>a) Prepare in triplicate b) Strain honey through 100–150 mesh nylon stocking material to remove insoluble material. c) Add 4 mL H₂O to 10–12 g honey (in triplicate) in a 50 mL centrifuge tube and mix well to get a homogenous solution. d) Prepare fresh by mixing 2.0 mL 10% Na₂WO₄ solution and 2.0 mL 0.335 M H₂SO₄ in a small test tube. e) Add this mixture immediately to the diluted honey solution and mix well. f) Swirl the tube in ca 80°C water bath until a visible flocculant (precipitate) forms with a clear supernatant. <i>Note: If no visible flocculant forms, or if supernatant remains cloudy, add 2 mL aliquots of 0.335 M H₂SO₄ with repeating heating between additions.</i> g) Fill tube with water, mix, centrifuge for 5 min at 6000 × g h) Decant supernatant. i) Repeat washing, mixing, and centrifuging steps nine times with ca 40 mL portions of water, thoroughly dispersing the pellet each time. j) Dry protein at least for 3 h in ca 75°C oven. k) Weigh approximately 0.1-0.2 mg isolated protein in tin capsules. l) Gently fold the tin capsule with the help of forceps and place it on the carousel of EA-IRMS for determining δ¹³C_{protein} ‰.</p> <p>Precautions</p> <p>i. Decant the supernatant immediately after centrifugation to avoid the mixing of pellet with the supernatant. ii. Protein washing must be done very carefully to avoid any loss of pellet with the water. iii. Fold the tin capsules gently to avoid the leakage or loss of sample iv. Be careful during tin capsule folding to avoid air trapping</p>
<p>EA-IRMS chromatography <i>(Instrument tuning varies with make and model. Set parameter as per manufacturer's instructions and optimize for best resolution)</i></p>	<p>Operate the system according to the manufacturer's instructions.</p> <p>Sample analysis</p> <p>1. Placed the isolated protein samples and standard in the tin capsules 2. Completely seal each capsule with tweezers 3. Place sealed capsules in the appropriate place on the tray of the automatic sampler 4. Operate the EA-IRMS following the manufacturer's instructions after calibration with CO₂ reference gas</p>

Instrumental settings for EA (vario ISOTOPE cube, Elementar, UK)

Temperature: Oxidation tube: 950°C

Reduction tube: 650°C

Pressure: 1300-1400 mbar

He flow: 230 mL/min

CO₂ flow: 230 mL/min

O₂ flow: 18 mL/min

Instrumental setting for IRMS (Isoprime IRMS)

Ion Source: CEI

High Vacuum: 5e-6

Turbo Speed: 100%

TCD temperature: 59°C

Focus point: >0.5

Accelerating Voltage: 4000v

Extraction Voltage: 76.00v

Half Plate differential (v): -121.00

Z-Plate Voltage (v): -53.00

Trap Current (μA) : 200.00

Electron Volts (9ev): 75.00

Ion Repellor Voltage (v): -9.00

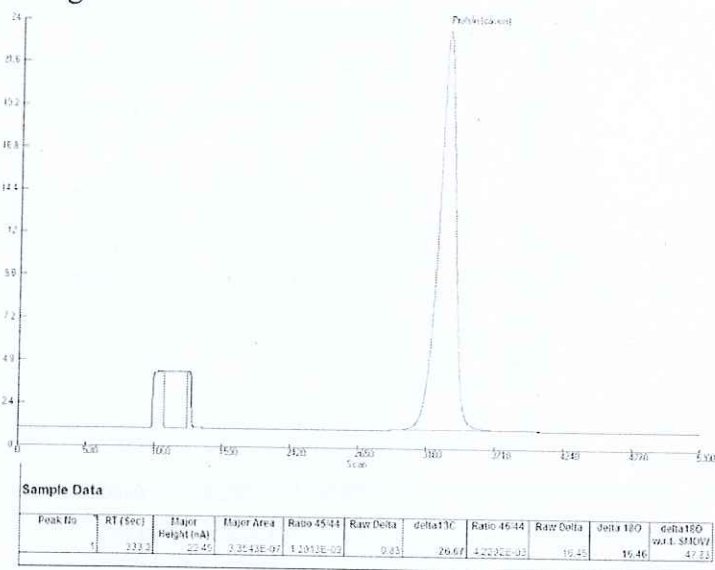
Magnet Current: 4000

The gas cylinders (associated valves etc) which supply working gases to the IRMS must be stored in a temperature-controlled environment.

Isotopic calculation

$\delta^{13}\text{C}_{\text{protein}} \text{‰}$

Isotope ratios are expressed relative to international standard. The $\delta^{13}\text{C}_{\text{protein}} \text{‰}$ is directly read of the chromatogram. See representative chromatogram below



Representative chromatogram: EA-IRMS profile of casein analysis showing $\delta^{13}\text{C}_{\text{protein}} \text{‰} = -26.67$

Preparation of standards for LC-IRMS

Preparation of reference sugar working standards

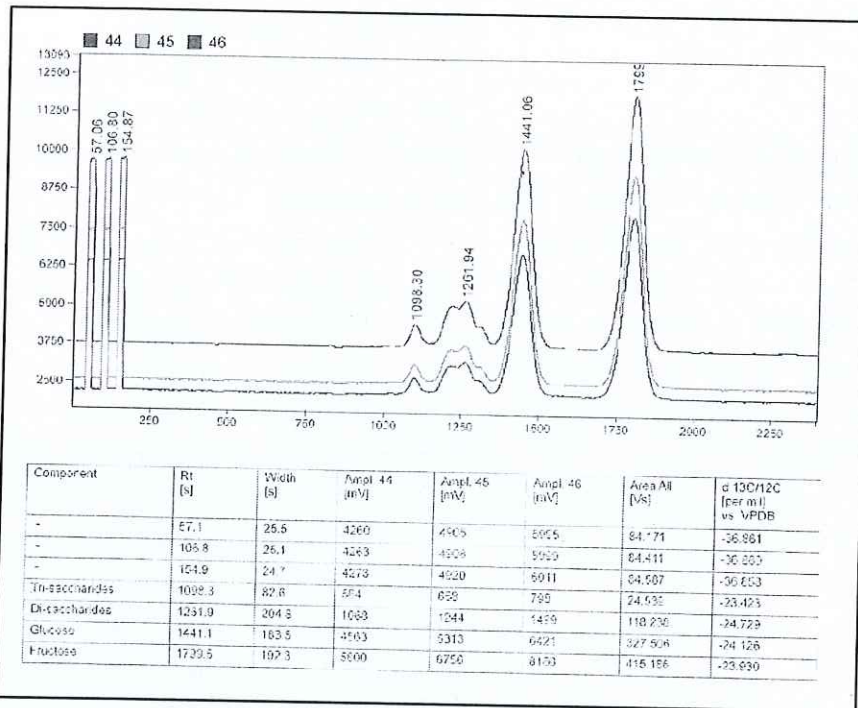
Prepare a solution of Fructose, Glucose, Sucrose and Raffinose containing 250 mg/L of each in ultra-pure water.

Filter through 0.22 μm syringe filter

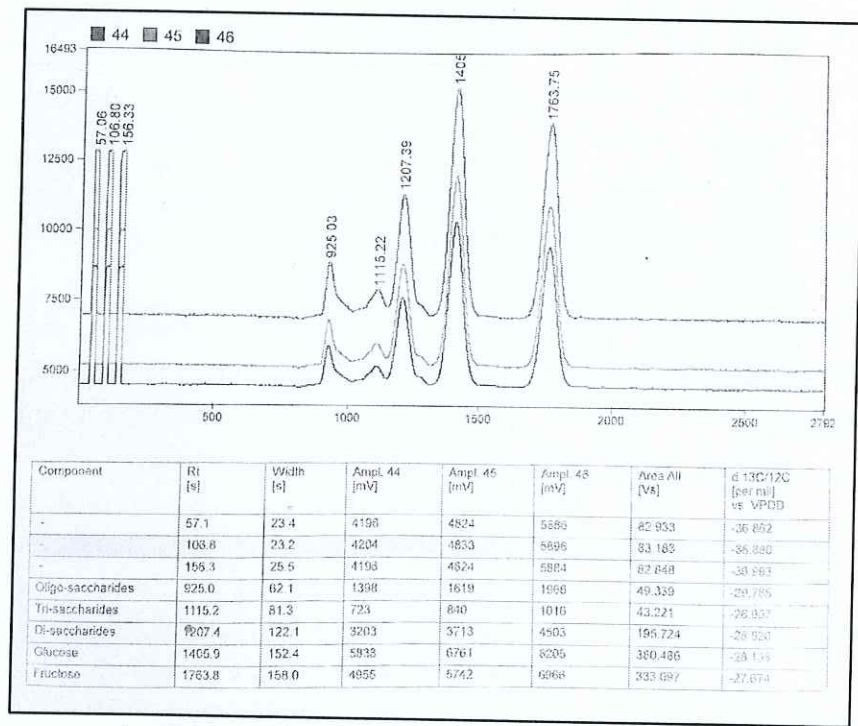
Sample Preparation for LC-IRMS analysis

1. Strain honey through a 100-150 mesh size nylon stocking material
2. In triplicate accurately weigh about 200 mg sample in a 15 ml centrifuge tube. Mix well with 5 ml of Ultra-pure water.

	<p>3. Sonicate the mixture and make the volume up to 10 mL with water in a 10 mL volumetric flask.</p> <p>4. Filter through 0.22 μm syringe filter into the HPLC injection vials.</p> <p><i>Note: Prepare sample solutions fresh everyday</i></p>														
<p>LC-IRMS conditions (Instrument setting and tuning vary depending on make and model. Set parameter and tune as per manufacturer's instructions and optimize for best results)</p>	<p>The conditions shown below have been optimized with the following:</p> <p>LC conditions</p> <ol style="list-style-type: none"> Column: Ca^{2+} (300 \times 7.7 mm, 8μm) Solvent: Ultra-pure Water Flow rate: 0.3 mL/min Column Oven temperature: 80 $^{\circ}\text{C}$ Injection volume: 10 μL <p>Interface for wet oxidation (Isoprime Liquiface, Elementar, UK)</p> <ol style="list-style-type: none"> Reactor temperature: 95$^{\circ}\text{C}$ Oxidation reagent: 20% Sodium peroxodisulfate (Purge the solution with Helium gas before use) Flow rate: 60 $\mu\text{L}/\text{min}$ <p><i>Note: Some instruments use 20% Sodium peroxodisulfate and 1.5 M H_3PO_4 for wet oxidation. Follow the manufacturer's instructions.</i></p> <p>IRMS Parameters (Isoprime IRMS):</p> <table> <tr> <td>Ion Source: CEI</td> <td>High Vacuum: 5e-6</td> </tr> <tr> <td>Turbo Speed: 100%</td> <td>TCD temperature: 59$^{\circ}\text{C}$</td> </tr> <tr> <td>Focus point: >0.5</td> <td>Accelerating Voltage: 4000v</td> </tr> <tr> <td>Extraction Voltage: 76.00v</td> <td>Half Plate differential (v): -121.00</td> </tr> <tr> <td>Z-Plate Voltage (v): -53.00</td> <td>Trap Current (μA): 200.00</td> </tr> <tr> <td>Electron Volts (9ev): 75.00</td> <td>Ion Repellor Voltage (v): -9.00</td> </tr> <tr> <td>Magnet Current: 4000</td> <td></td> </tr> </table>	Ion Source: CEI	High Vacuum: 5e-6	Turbo Speed: 100%	TCD temperature: 59 $^{\circ}\text{C}$	Focus point: >0.5	Accelerating Voltage: 4000v	Extraction Voltage: 76.00v	Half Plate differential (v): -121.00	Z-Plate Voltage (v): -53.00	Trap Current (μA): 200.00	Electron Volts (9ev): 75.00	Ion Repellor Voltage (v): -9.00	Magnet Current: 4000	
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<p>LC-IRMS chromatography of samples</p>	<ol style="list-style-type: none"> Introduce CO_2 reference gas pulse three times (20 s each) at the beginning of each run. The constant flow rate during this period gives the peaks a flattop appearance (see figure below). A level of CO_2 corresponding to 2–5 V (depending on the instrument) at m/z 44 is used to calibrate the system. Inject standard mixture (10 μL) of fructose, glucose, disaccharide and trisaccharide. Repeat 10 times to obtain the mean and standard deviation for the $\delta^{13}\text{C}$ ‰ of individual sugars. Inject Honey sample (10 μL) in triplicate. The IRMS chromatogram provides details of the $\delta^{13}\text{C}$ ‰ of each of the sugars in the sample and the area under the curve of each of the resolved sugars (see representative chromatograms below). The $\Delta\delta^{13}\text{C}_{\text{fru-glu}}$ ‰, $\Delta\delta^{13}\text{C}_{\text{max}}$ ‰, and foreign oligosaccharide content are calculated from the chromatogram data. 														
<p>Representative LC-IRMS profiles</p>															



Representative LC-IRMS profile of a pure Honey sample



Representative LC-IRMS profile of a honey sample containing foreign oligosaccharides

Calculations

1. $\Delta\delta^{13}\text{C}_{\text{fru} - \text{flu}} \%$

Subtract the $\delta^{13}\text{C}_{\text{Glu}}$ (‰) value given in the chromatogram from the $\delta^{13}\text{C}_{\text{Fru}}$ (‰) value. Report as $\Delta\delta^{13}\text{C}_{\text{fru} - \text{flu}} \%$

2. $\Delta\delta^{13}\text{C}_{\text{max}} \%$

Extract the $\delta^{13}\text{C}$ ‰ values of fructose, glucose, disaccharides and trisaccharides from the LC-IRMS profile.

Extract the $\delta^{13}\text{C}$ ‰ of protein from EA-IRMS profile and tabulate as shown

A $\delta^{13}\text{C}$ ‰	B $\delta^{13}\text{C}$ ‰	A-B $\Delta\delta^{13}\text{C}$ ‰
Fructose	Disaccharide	
Fructose	Trisaccharide	
Fructose	Protein	
Glucose	Disaccharide	
Glucose	Trisaccharide	
Glucose	Protein	
Disaccharide	Trisaccharide	
Disaccharide	Protein	
Trisaccharide	Protein	

The highest value observed in column three gives $\Delta\delta^{13}\text{C}_{\text{max}}$ ‰

3. Foreign oligosaccharides (% peak area)

Extract the area of individual peaks and calculate using the formula

$$\text{Foreign oligosaccharide (area \%)} = \frac{\text{Sum of the peak area of all peak(s) other than Fructose, Glucose, disaccharides and trisaccharides}}{\text{Total peak area}} \times 100$$

Quality control

- The laboratory working standard must be analysed in each sequence of sample measurements at least once.
- A quality control sample of a chosen pure honey must be analysed in each LC-IRMS measurement sequence.

Reference

- AOAC Official Method 998.12 C-4 Plants Sugars in Honey. Internal Standard Stable Carbon Isotope Ratio Method First Action 1998
- Improved detection of honey adulteration by measuring differences between $^{13}\text{C}/^{12}\text{C}$ stable carbon isotope ratios of protein and sugar compounds with a combination of elemental analyzer - isotope ratio mass spectrometry and liquid chromatography - isotope ratio mass spectrometry ($\delta^{13}\text{C}$ -EA/LCIRMS). Lutz Elflein, Kurt-Peter Ræzke; *Apidologie* 2008, 39 (5), 574–587.
- Liquid chromatography coupled to isotope ratio mass spectrometry: A new perspective on honey adulteration detection. Ana I. Cabanero, Jose L. Recio, Mercedes RupeãRez; *J. Agric. Food Chem.* 2006, 54, 9719-9727.
- LC-IRMS: Authenticity control of honey using the Thermo Scientific LC IsoLink LC-IRMS. Andreas W. Hilker, Michael Krummen, Dieter Juchelka; Thermo application note 30024.
- "Scientific support to the implementation of a Coordinated Control Plan with a view to establishing the prevalence of fraudulent practices in the marketing of honey" N° SANTE/2015/E3/JRC/SI2.706828. E.

	Aries, J. Burton, L. Carrasco, O. De Rudder, and A. Maquet. JRC Technical Report 2016, JRC104749, 38 p.
Approved by	Scientific Panel on Methods of Sampling and Analysis