

ALMA MATER STUDIORUM Università di Bologna

Comparison of three different methodologies for the determination of hydroxytyrosol and tyrosol in relation to the health claim for olive oils

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HEALTH CLAIM FOR "OLIVE OIL POLYPHENOLS" EU regulation 432/2012

"Olive oil polyphenols contribute to the protection of blood lipids from oxidative stress"

The claim may be used only for olive oil which contains at least 5 mg of hydroxytyrosol and its derivatives (e.g. oleuropein complex and tyrosol) per 20 g of olive oil. In order to bear the claim information shall be given to the consumer that the beneficial effect is obtained with a daily intake of 20 g of olive oil.

OBJECTIVE:

Comparison of three different analytical procedures, already published in the scientific

literature or tested/validated in the IOC, ISO and Oleum project frameworks,

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for the determination of HYTY and TY

Samples

Six different olive oil samples, characterized by a wide range in terms of phenolic compounds content.

Methods by hydrolysis: extraction of HYTY and TY

Method 1. Direct hydrolysis (Romero and Brenes, 2012; J. Agric. Food Chem, 60, 9017–9022; with some modifications)



Methods by hydrolysis: extraction of HYTY and TY

Method 2. Hydrolysis after phenolic extraction (PD ISO/TS 23942:2020)

Sample preparation:

- 2 g of oil
- 5 mL MeOH/H₂O (80:20)
- Vortex 1 min
- Ultrasound 15 min
- Centrifuge 5000 rpm, 25 min
- Filter 1 mL aqueous phase
- Dry at 40 °C under N₂
- Add 1 mL EtOH/ H_2O/H_2SO_4 (50:40:10)
- Bath 40 °C, 1 h
- Ambient T, by a night
- Filter: 0.45 μm, PVDF
- Aproximative sample preparation time: 3 h (overnight)

HPLC conditions:

- Elution by a gradient
- Phase A: H₂O/H₃PO₄ (0.2 %)
- Phase B: MeOH/ACN (1:1)
- Initial conditions: 96% A
- Flow: 1 mL/min
- Column: RP-HPLC C18
 - (4.6 mm i.d. X 250 mm; 5 μm particle size)
- Injection volume : 20 μ L
- Analysis time: 82 min



Methods by hydrolysis: extraction of HYTY and TY

Method 3. Hydrolysis after phenolic extraction (OLEUM Project, Tsimidou et al., 2019; *Molecules*, 24(6): 1044)

Sample preparation:

- 2 g of oil
- 1 mL MeOH/H₂O (80:20), vortex 30 s
- 5 mL MeOH/H₂O (80:20), vortex 1 min
- Ultrasound 15 min
- Centrifuge 2745 rpm, 25 min
- Filter aqueous phase, 0.22 μm, PVDF
- 200 μ L aqueous phase + 200 μ L H₂SO₄ 1 M, vortex 2 s
- X 3 Bath 80 °C, 2 h
 - Dilute with 200 μ L MeOH/H₂O (80:20)
 - Mix 3 extracts
 - Filter: 0.22 μm, PVDF
 - Aproximative sample preparation time: 4 h

UPLC conditions:

- Elution by a gradient
- Phase A: H_2O/H_3PO_4 (0.2%)
- Phase B: MeOH/ACN (1:1)
- Initial conditions: 96% A
- Flow: 0.45 mL/min
- Column: Shim-pack XR-ODS III
 - (2 mm i.d. X 75 mm; 1.6 μm particle size)
- Injection volume : $3 \mu L$
- Analysis time: 20.5 min



Methods by hydrolysis: extraction of HYTY and TY

Method 3. Hydrolysis after phenolic extraction (OLEUM Project, Tsimidou et al., 2019; *Molecules*, 24(6): 1044)

Method 3a

Calculation made by external calibration, without considering the compounds A and B and the correction factors proposed

in the framework of OLEUM project)

Method 3b

Calculation made considering the compounds A and B, and the correction factors established in the OLEUM method



1.8. Calculations and Data Analysis

1.8.1. Expression of Results

The total amount of **Htyr** and **Tyr** is calculated as the sum of their free and bound forms after correction for molecular mass differences between free and bound forms (see scheme of the method). Correction factors are introduced in the quantification of individual total **Htyr** (2.2) and **Tyr** (2.5), which are obtained by dividing the mean molecular mass of the most known bound forms of **Htyr** and **Tyr** (343 amu) by the molecular mass of **Htyr** (154 amu) and **Tyr** (138 amu), respectively*. The results for each of them are expressed as mg/20 g oil (with two decimal points). Their sum is then rounded to the nearest integer. Taking into calculation for the total **Htyr** content the peak areas of peaks A and B, when present in the chromatograms, no correction for recovery is necessary.

***NOTE**: Bound forms used for the derived correction factors were: tyrosol glucoside (300.30 amu), aldehydic form of ligstroside aglycone (362.12 amu), dialdehyde form of ligstroside aglycone (304.34 amu), aldehyde form of oleuropein aglycone (378.37 amu), oleuropein aglycone (378.37 amu), (carboxymethylated) dialdehyde form of oleuropein aglycone (378.37 amu), (decarboxymethylated) dialdehyde form of oleuropein aglycone (320.34 amu), (decarboxymethylated) aldehyde form of oleuropein aglycone (320.34 amu), (decarboxymethylated) aldehyde form of oleuropein aglycone (320.34 amu), (decarboxymethylated) aldehyde form of oleuropein aglycone (320.34 amu).

Equation: Total Htyr and Tyr (mg/20 g oil) = $[Htyr_{free^*}] + [Tyr_{free^*}] + 2.2 \times [Htyr_{hydrolysate^-}] + 100 + 1$

The result obtained with two decimals is then rounded to the nearest integer

*NOTE: free Htyr or Tyr refers to the analysis of the PF prior to hydrolysis



MATERIALS AND METHODS Methods without hydrolysis: extraction of the total phenolic fraction

Method A. ISOLATION OF PHENOLIC COMPOUNDS BY SPE-DIOL

Sample preparation:

- 2.5 g of oil
- 500 μL IS
- Shake and evaporate (6-7 min) under N₂
- Activate the cartridge SPE: 6 mL MeOH, 6 mL *n*-hexane
- Dissolve the oil sample (6 mL *n*-hexane)
- Placed sample onto the activated SPE cartridge
- Leave it enter into the cartridge
- Wash the sample container (6 mL *n*-hexane), add it to the cartridge and discard
- Add 4 mL hexane:ethyl acetate (85:15) and discard
- Elute the analytes with 10 mL MeOH and collect
- Evaporate in a rotary evaporator
- Dissolve the residue: 500 μ L, MeOH/H₂O (1:1)
- Filter: 0.45 μm, cellulose acetate
- Aproximative sample preparation time: 3 h

HPLC conditions:

OLIVE

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- Elution by a gradient

INTERNATIONAL

- Phase A: H_2O/H_3PO_4 (95.5:0.5)
- Phase B: MeOH/AcN (1:1)
- Initial conditions: 95% A
- Flow: 1 mL/min
- Column: RP-HPLC C18 column
 (4.0 mm i.d. x 250 mm; 5 μm particle size)
- Injection volume : 20 μ L
- Analysis time: 50 min

COI/T.20/Doc. No 29/Rev.2 June 2022

ENGLISH Original: ITALIAN





Methods without hydrolysis: extraction of the total phenolic fraction

Method B. IOC method (COI/T. 20/Doc No 29)

Sample preparation:

- 2 g of oil
- 1 mL IS (syringic acid), shake 30s
- 5 mL MeOH/H₂O (80:20)
- Vortex 1 min
- Ultrasound 15 min
- Centrifuge 5000 rpm, 25 min
- Take supernatant
- Filter: 0.45 μm, PVDF
- Aproximative sample preparation time: 1 h

HPLC conditions:

- Elution by a gradient
- Phase A: H_2O/H_3PO_4 (0.2 %)
- Phase B: MeOH/ACN (1:1)
- Initial conditions: 96% A
- Flow: 1mL/min
- Column: RP-HPLC C18
 (4.6) mm i.d. X 250 mm; 5 μm particle size)
- Injection volume : 20 μL
- Analysis time: 82 min



Methods without hydrolysis: extraction of the total phenolic fraction

Method C. Folin-Ciocalteu method

Sample preparation (method B without IS):

- 2 g of oil
- 6 mL MeOH/H₂O (80:20)
- Vortex 1 min
- Ultrasound 15 min
- Centrifuge 5000 rpm, 25 min
- Take supernatant
- Filter: 0.45 μm, PVDF

Determination of total phenolic:

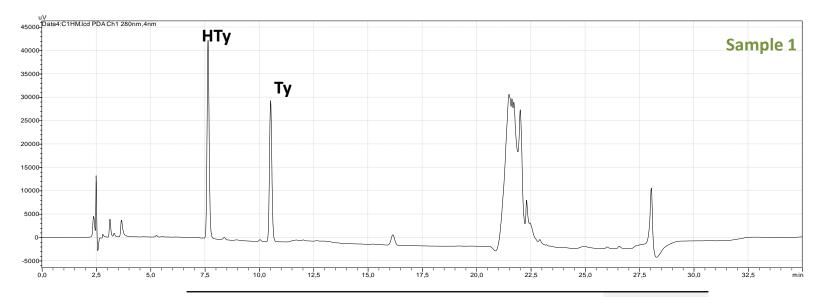
- 0.2 mL of the phenolic extract + 0.5 mL FC reagent +
 2 mL of Na₂CO₃ (15%), in a 10 mL volumetric flask reaching the final volume with purified water
- Store for 2 h at room temperature
- Phenolic compounds were detected at 750 nm and quantified using gallic acid (PM 170) calibration

curve



RESULTS Methods by hydrolysis

Method 1. Direct hydrolysis



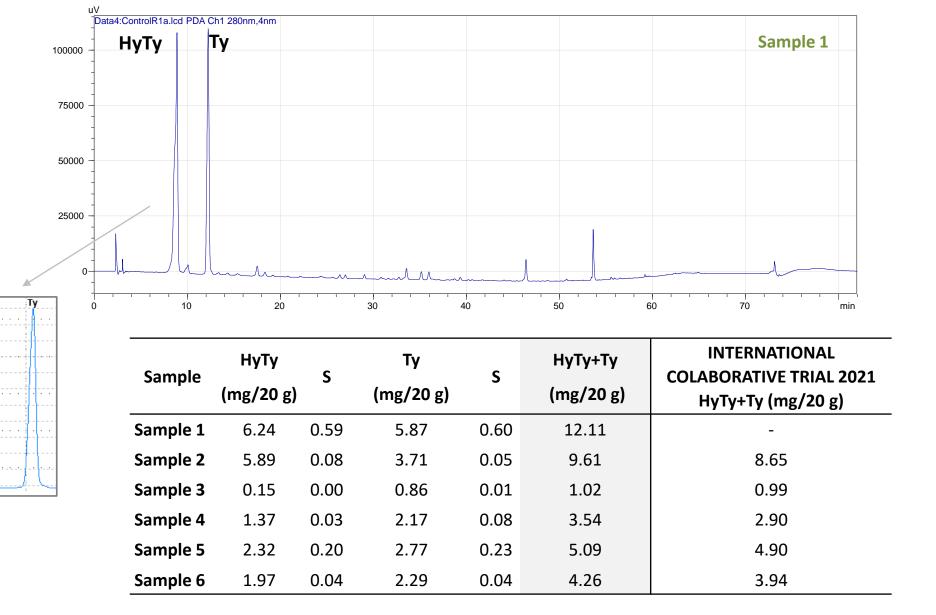
Sample	HyTy (mg/20 g)	S	Ty (mg/20 g)	S	HyTy+Ty (mg/20 g)
Sample 1	6.27	0.64	6.64	0.72	12.91
Sample 2	6.06	0.33	4.35	0.09	10.40
Sample 3	nd	nd	1.16	0.04	1.16
Sample 4	0.76	0.05	2.26	0.04	3.02
Sample 5	2.22	0.15	3.59	0.21	5.81
Sample 6	1.88	0.08	3.07	0.03	4.95



Method 2. Hydrolysis after phenolic extraction (PD ISO/TS 23942:2020)

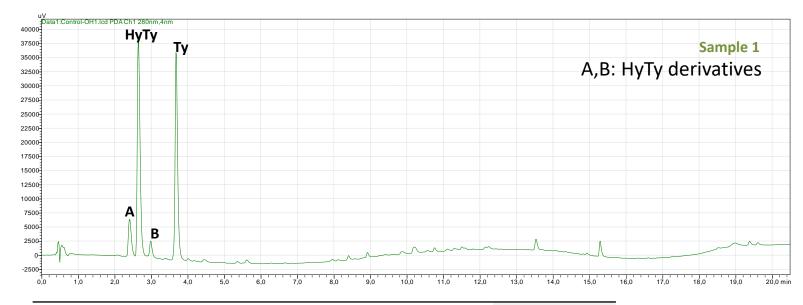
HTy

RESULTS Methods by hydrolysis





Method 3. Hydrolysis after phenolic extraction (OLEUM Project)



RESULTS Methods by hydrolysis

Sample	НуТу	S	Ту	ſ	HTy+Ty
	(mg/20 g)		(mg/20 g)	S	(mg/20 g)
Sample 1	6.42	0.01	7.23	0.01	13.65
Sample 2	6.03	0.01	4.37	0.00	10.40
Sample 3	0.20	0.00	1.11	0.00	1.31
Sample 4	1.20	0.00	2.36	0.00	3.56
Sample 5	2.62	0.00	3.56	0.00	6.18
Sample 6	1.99	0.00	2.92	0.00	4.91

Method 3a: calculation made without considering the compounds A and B and the correction factors established in the oleum method

	А, В, НуТу е Ту	
Sample	(mg/20 g oil)	S
Sample 1	35.24	0.05
Sample 2	25.56	0.01
Sample 3	3.08	0.01
Sample 4	8.52	0.01
Sample 5	14.97	0.02
Sample 6	12.20	0.01

Method 3b: calculation made considering the compounds A and B,and the correction factors established in the oleum method





Conclusions: comparison of three different hydrolytic methods

- The three hydrolytic methods give comparable results
- From the samples analyzed there is no evidence of an underestimation by any of the methods
- All methods quantify as tyrosol and hydroxytyrosol (or gallic acid for the Folin-Ciocalteu) and therefore a correction factor should be applied
- Analyzes are ongoing to compare the results with IOC methods that quantify individual polyphenols





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